The PENCIL CODE: multi-purpose and multi-user maintained

http://www.nordita.org/~brandenb/talks/misc/PencilCode04.htm

Figure 1: Check-in patterns as a function of time for different subroutines. The different users are marked by different symbols and different colors.


Contributions to master, excluding merge commits

Figure 2: Check-in frequency on the github page as a function of time; see https://github.com/pencil-code/pencil-code/graphs/contributors
### Contributors to the code

(in inverse alphabetical order according to their user name)

An up to date list of Pencil Code contributors can be found at [GitHub](https://github.com).

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We, the PENCIL CODE community, ask that in publications and presentations the use of the code (or parts of it) be acknowledged with reference to the web site http://www.nordita.org/software/pencil-code/ or (equivalently) to. https://github.com/pencil-code/pencil-code

As a courtesy to the people involved in developing particularly important parts of the program (use svn annotate src/* .f90 to find out who did what!) we suggest to give appropriate reference to one or several of the following (or other appropriate) papers (listed here in temporal order):


This list is not always up-to-date. We therefore ask the developers to check in new relevant papers, avoiding however redundancies.

We are aware of the fact that certain extensions to the code may still be under intense development and no paper can be quoted yet. Again, as a courtesy to those developers, we suggest to contact them, if possible, and ask whether there is anything else that can be quoted instead.
Foreword

This code was originally developed at the Turbulence Summer School of the Helmholtz Institute in Potsdam (2001). While some SPH and PPM codes for hydrodynamics and magnetohydrodynamics are publicly available, this does not generally seem to be the case for higher order finite-difference or spectral codes. Having been approached by people interested in using our code, we decided to make it as flexible as possible and as user-friendly as seems reasonable, and to put it onto a public CVS repository. Since 21 September 2008 it is distributed via https://github.com/pencil-code/pencil-code. The code can certainly not be treated as a black box (no code can), and in order to solve a new problem in an optimal way, users will need to find their own optimal set of parameters. In particular, you need to be careful in choosing the right values of viscosity, magnetic diffusivity, and radiative conductivity.

The PENCIL CODE is primarily designed to deal with weakly compressible turbulent flows, which is why we use high-order first and second derivatives. To achieve good parallelization, we use explicit (as opposed to compact) finite differences. Typical scientific targets include driven MHD turbulence in a periodic box, convection in a slab with non-periodic upper and lower boundaries, a convective star embedded in a fully nonperiodic box, accretion disc turbulence in the shearing sheet approximation, etc. Furthermore, nonlocal radiation transport, inertial particles, dust coagulation, self-gravity, chemical reaction networks, and several other physical components are installed, but this number increases steadily. In addition to Cartesian coordinates, the code can also deal with spherical and cylindrical polar coordinates.

Magnetic fields are implemented in terms of the magnetic vector potential to ensure that the field remains solenoidal (divergence-free). At the same time, having the magnetic vector potential readily available is a big advantage if one wants to monitor the magnetic helicity, for example. The code is therefore particularly well suited for all kinds of dynamo problems.

The code is normally non-conservative; thus, conserved quantities should only be conserved up to the discretization error of the scheme (not to machine accuracy). There is no guarantee that a conservative code is more accurate with respect to quantities that are not explicitly conserved, such as entropy. Another important quantity that is (to our knowledge) not strictly conserved by ordinary flux conserving schemes is magnetic helicity.

There are currently no plans to implement adaptive mesh refinement into the code, which would cause major technical complications. Given that turbulence is generically space-filling, local refinement to smaller scales would often not be very useful anyway. On the other hand, in some geometries turbulence may well be confined to certain regions in space, so one could indeed gain by solving the outer regions with fewer points.

In order to be cache-efficient, we solve the equations along pencils in the $x$ direction. One very convenient side-effect is that auxiliary and derived variables use very little memory, as they are only ever defined on one pencil. The domain can be tiled in the $y$ and $z$ directions. On multiprocessor computers, the code can use MPI (Message Passing Interface) calls to communicate between processors. An easy switching mechanism allows the user to run the code on a machine without MPI libraries (e.g. a notebook computer). Ghost zones are used to implement boundary conditions on physical and processor boundaries.
A high level of flexibility is achieved by encapsulating individual physical processes and variables in individual modules, which can be switched on or off in the file ‘Makefile.local’ in the local ‘src’ directory. This approach avoids the use of difficult-to-read preprocessor directives, at the price of requiring one dummy module for each physics module. For nonmagnetic hydrodynamics, for example, one will use the module ‘nomagnetic.f90’ and specifies

    MAGNETIC = nomagnetic

in ‘Makefile.local’, while for MHD simulations, ‘magnetic.f90’ will be used:

    MAGNETIC = magnetic

Note that the term module as used here is only loosely related to Fortran modules: both ‘magnetic.f90’ and ‘nomagnetic.f90’ define an F90 module named Magnetic — this is the basis of the switching mechanism we are using.

Input parameters (which are set in the files ‘start.in’, ‘run.in’) can be changed without recompilation. Furthermore, one can change the list of variables for monitoring (diagnostic) output on the fly, and there are mechanisms for making the code reload new parameters or exit gracefully at runtime. You may want to check for correctness of these files with the command pc_configtest.

The requirements for using the Pencil-MPI code are modest: you can use it on any Linux or Unix system with a F95 and C compiler suite, like GNU gcc and gfortran, together with the shell CSH, and the Perl interpreter are mandatory requirements.

Although the PENCIL CODE is mainly designed to run on supercomputers, more than 50% of the users run their code also on Macs, and the other half uses either directly Linux on their laptops or they use VirtualBox on their Windows machine on which they install Ubuntu Linux. If you have IDL as well, you will be able to visualize the results (a number of sample procedures are provided), but other tools such as Python, DX (OpenDX, data explorer) can also be used and some relevant tools and routines come with the PENCIL CODE.

If you want to make creative use of the code, this manual will contain far too little information. Its major aim is to give you an idea of the way the code is organized, so you can more efficiently read the source code, which contains a reasonable amount of comments. You might want to read through the various sample directories that are checked in. Choose one that is closest to your application and start modifying. For further enhancements that you may want to add to the code, you can take as an example the lines in the code that deal with related variables, functions, diagnostics, equations etc., which have already been implemented. Just remember: grep is one of your best friends when you want to understand how certain variables or functions are used in the code.

We will be happy to include user-supplied changes and updates to the code in future releases and welcome any feedback.

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Many people have contributed in different ways to the development of this code. We thank first of all Åke Nordlund (Copenhagen Observatory) and Bob Stein (University of Michigan) who introduced us to the idea of using high-order schemes in compressible flows and who taught us a lot about simulations in general.

The calculation of the power spectra, structure functions, the remeshing procedures, routines for changing the number of processors, as well as the shearing sheet approximation and the flux-limited diffusion approximation for radiative transfer were implemented by Nils Erland L. Haugen (University of Trondheim). Tobi Heinemann added the long characteristics method for radiative transfer as well as hydrogen ionization. He also added and/or improved shock diffusion for other variables and improved the resulting timestep control. Anders Johansen, Wladimir (Wlad) Lyra, and Jeff Oishi contributed to the implementation of the dust equations (which now comprises an array of different components). Antony (Tony) Mee (University of Newcastle) implemented shock viscosity and added the interstellar module together with Graeme R. Sarson (also University of Newcastle), who also implemented the geodynamo set-up together with David McMillan (currently also at the University of Newcastle). Tony also included a method for outputting auxiliary variables and enhanced the overall functionality of the code and related idl and dx procedures. He also added, together with Andrew Snodin, the evolution equations for the cosmic ray energy density. Vladimir Pariev (University of Rochester) contributed to the development and testing of the potential field boundary condition at an early stage. The implementation of spherical and cylindrical coordinates is due to Dhrubaditya (Dhruba) Mitra and Wladimir Lyra. Wlad also implemented the global set-up for protoplanetary disks (as opposed to the local shearing sheet formalism). He also added a $N$-body code (based on the particle module coded by Anders Johansen and Tony), and implemented the coupled evolution equations of neutrals and ions for two-fluid models of ambipolar diffusion. Boris Dintrans is in charge of implementing the anelastic and Boussinesq modules. Philippe-A. Bourdin implemented HDF5 support and wrote the optional IO-modules for high-performance computing featuring various communication strategies. He also contributed to the solar-corona module and worked on the IDL GUI, including the IDL routines for reading and working with large amounts of data. Again, this list contains other recent items that are not yet fully documented and acknowledged.

Use of the PPARC supported supercomputers in St Andrews (Mhd) and Leicester (Ukaff) is acknowledged. We also acknowledge the Danish Center for Scientific Computing for granting time on Horseshoe, which is a 512+140 processor Beowulf cluster in Odense (Horseshoe).
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Part I

Using the PENCIL Code

1 System requirements

To use the code, you will need the following:

1. Absolutely needed:
   - *F95* compiler
   - *C* compiler

2. Used heavily (if you don’t have one of these, you will need to adjust many things manually):
   - a Unix/Linux-type system with *make* and *csh*
   - *Perl* (remember: if it doesn’t run Perl, it’s not a computer)

3. The following are dispensable, but enhance functionality in one way or the other:
   - an *MPI* implementation (for parallelization on multiprocessor systems)
   - *DX* alias *OpenDX* or *data explorer* (for 3-D visualization of results)
   - *IDL* (for visualization of results; the 7-minute demo license will do for many applications)
2 Obtaining the code

The code is now distributed via [https://github.com/pencil-code/pencil-code](https://github.com/pencil-code/pencil-code), where you can either download a tarball, or, preferably, download it via `svn` or `git`. In Iran and some other countries, GitHub is not currently available. To alleviate this problem, we have made a recent copy available on [http://www.nordita.org/software/pencil-code/](http://www.nordita.org/software/pencil-code/). If you want us to update this tarball, please contact us.

To ensure at least some level of stability of the `svn/git` versions, a set of test problems (listed in `$PENCIL_HOME/bin/auto-test`) are routinely tested. This includes all problems in `$PENCIL_HOME/samples`. See Sect. [10] for details.

2.1 Obtaining the code via `git` or `svn`

1. Many machines have `svn` installed (try `svn -v` or `which svn`). On Ubuntu, for example, `svn` comes under the package name `subversion`.
2. The code is now saved under Github, `git` can be obtained in Linux by typing `sudo apt-get install git`.
3. Unless you are a privileged users with write access, you can download the code with the command
   ```bash
   git clone https://github.com/pencil-code/pencil-code.git
   ``
   or
   ```bash
   svn checkout https://github.com/pencil-code/pencil-code/trunk/ ...\n   pencil-code --username MY_GITHUB_USERNAME
   ```

   In order to push your changes to the repository, you have to ask the maintainer of pencil code for push access (to become a contributor), or put a pull request to the maintainer of the code.

   Be sure to run `auto-test` before you check anything back in again. It can be very annoying for someone else to figure out what’s wrong, especially if you are just up to something else. At the very least, you should do
   ```bash
   pc_auto-test --level=0 --no-pencil-check -C
   ```
   This allows you to run just 2 of the most essential tests starting with all the no-modules and then most-modules.

2.2 Updating via `svn` or `git`

Independent of how you installed the code in the first place (from tarball or via `svn/git`), you can update your version using `svn/git`. If you have done nontrivial alterations to your version of the code, you ought to be careful about upgrading: although `svn/git` is an excellent tool for distributed programming, conflicts are quite possible, since many of us are going to touch many parts of the code while we develop it further. Thus, despite the
2.3 Getting the last validated version

The fact that the code is under \texttt{svn/git}, you should probably back up your important changes before upgrading.

Here is the upgrading procedure for \texttt{git}:

1. Perform a \texttt{git} update of the tree:
   
   \begin{verbatim}
   unix> git pull
   \end{verbatim}

2. Fix any conflicts you encounter and make sure the examples in the directory \texttt{‘samples/’} are still working.

Here is the upgrading procedure for \texttt{svn}:

1. Perform a \texttt{svn} update of the tree:
   
   \begin{verbatim}
   unix> pc_svnup
   \end{verbatim}

2. Fix any conflicts you encounter and make sure the examples in the directory \texttt{‘samples/’} are still working.

If you have made useful changes, please contact one of the (currently) 10 “Contributors” (listed under \url{https://github.com/pencil-code/pencil-code}) who can give you push or check-in permission. Be sure to have sufficient comments in the code and please follow our standard coding conventions explained in Section 9.1. There is also a script to check and fix the most common stylebreaks, \texttt{pc_codingstyle}.

2.3 Getting the last validated version

The script \texttt{pc_svnup} accepts arguments \texttt{-val} or \texttt{-validated}, which means that the current changes on a user’s machine will be merged into the last working version. This way every user can be sure that any problems with the code must be due to the current changes done by this user since the last check-in.

Examples:

\begin{verbatim}
unix> pc_svnup -src -s -validated
\end{verbatim}

brings all files in \texttt{'$PENCIL_HOME/src'} to the last validated status, and merges all your changes into this version. This allows you to work with this, but in order to check in your changes you have to update everything to the most recent status first, i.e.

\begin{verbatim}
unix> pc_svnup -src
\end{verbatim}

Your own changes will be merged into this latest version as before.

NOTE: The functionality of the head of the trunk should be preserved at all times. However, accidents do happen. For the benefit of all other developers, any errors should be corrected within 1-2 hours. This is the reason why the code comes with a file \texttt{‘pencil-code/license/developers.txt’}, which should contain contact details of all developers. The \texttt{pc_svnup -val} option allows all other people to stay away from any trouble.
2.4 Getting older versions

You may find that the latest \textit{svn} version of the code produces errors. If you have reasons to believe that this is due to changes introduced on 27 November 2008 (to give an example), you can check out the version prior to this by specifying a revision number with \texttt{svn update -r ####}. One reason why one cannot always reproduce exactly the same situation too far back in time is connected with the fact that processor architecture and the compiler were different, resulting e.g. in different rounding errors.
3 Getting started

To get yourself started, you should run one or several examples which are provided in one of the ‘samples/’ subdirectories. Note that you will only be able to fully assess the numerical solutions if you visualize them with IDL, DX or other tools (see Sect. 5.19).

3.1 Setup

3.1.1 Environment settings

The functionality of helper scripts and IDL routines relies on a few environment variables being set correctly. The simplest way to achieve this is to go to the top directory of the code and source one of the two scripts ‘sourceme.csh’ or ‘sourceme.sh’ (depending on the type of shell you are using):

```csh
  csh> cd pencil-code
  csh> source ./sourceme.csh
```

for tcsh or csh users; or

```sh
  sh> cd pencil-code
  sh> . ./sourceme.sh
```

for users of bash, Bourne shell, or similar shells. You should get output similar to

```
PENCIL_HOME = </home/dobler/f90/pencil-code>
Adding /home/dobler/f90/pencil-code/bin to PATH
```

Apart from the PATH variable, the environment variable IDL_PATH is set to something like ./idl:+$PENCIL_HOME/idl:/data:<IDL_DEFAULT>.

**Note 1** The <IDL_DEFAULT> mechanism does not work for IDL versions 5.2 or older. In this case, you will have to edit the path manually, or adapt the ‘sourceme’ scripts.

**Note 2** If you don’t want to rely on the ‘sourceme’ scripts’ (quite heuristic) ability to correctly identify the code’s main directory, you can set the environment variable PENCIL_HOME explicitly before you run the source command.

**Note 3** Do not just source the ‘sourceme’ script from your shell startup file (‘~/cshrc’ or ‘~/bashrc’), because it outputs a few lines of diagnostics for each sub-shell, which will break many applications. To suppress all output, follow the instructions given in the header documentation of ‘sourceme.csh’ and ‘sourceme.sh’. Likewise, output from other files invoked by source should also be suppressed.

**Note 4** The second time you source ‘sourceme’, it will not add anything to your PATH variable. This is on purpose to avoid cluttering of your environment: you can source the file as often as you like (in your shell startup script, then manually and in addition in some script you have written), without thinking twice. If, however, at the first sourcing, the setting of PENCIL_HOME was wrong, this mechanism would keep you from ever adding
the right directory to your PATH. In this case, you need to first undefine the environment variable PENCIL_HOME:

```
csh> unsetenv PENCIL_HOME
```

```
csh> source ./sourceme.csh
```

```
hs> unset PENCIL_HOME
hs> . ./sourceme.sh
```

### 3.1.2 Linking scripts and source files

With your environment set up correctly, you can now go to the directory you want to work in and set up subdirectories and links. This is accomplished by the script `pc_setupsrc`, which is located in `"$PENCIL_HOME/bin"` and is thus now in your executable path.

For concreteness, let us assume you want to use `samples/conv-slab` as your run directory, i.e. you want to run a three-layer slab model of solar convection. You then do the following:

```
unix> cd samples/conv-slab
unix> pc_setupsrc
```

```
src already exists
2 files already exist in src
```

The script has linked a number of scripts from `"$PENCIL_HOME/bin"`, generated a directory `src` for the source code and linked the Fortran source files (plus a few more files) from `"$PENCIL_HOME/src"` to that directory:

```
unix> ls -F
reference.out src/
start.csh@ run.csh@ getconf.csh@
start.in run.in print.in
```

### 3.1.3 Adapting ‘Makefile.src’

This step requires some input from you, but you only have to do this once for each machine you want to run the code on. See Sect. 5.2 for a description of the steps you need to take here.

**Note**: If you are lucky and use compilers similar to the ones we have, you may be able to skip this step; but blame yourself if things don’t compile, then. If not, you can run make with explicit flags, see Sect. 5.2 and in particular Table 1.

### 3.1.4 Running make

Next, you run `make` in the ‘src’ subdirectory of your run directory. Since you are using one of the predefined test problems, the settings in ‘src/Makefile.local’ and ‘src/cparam.local’ are all reasonable, and you just do
3.1 Setup

If you have set up the compiler flags correctly, compilation should complete successfully.

3.1.5 Choosing a data directory

The code will by default write data like snapshot files to the subdirectory ‘data’ of the run directory. Since this will involve a large volume of IO-operations (at least for large grid sizes), one will normally try to avoid writing the data via NFS. The recommended way to set up a ‘data’ data directory is to generate a corresponding directory on the local disc of the computer you are running on and (soft-)link it to ‘./data’. Even if the link is part of an NFS directory, all the IO operations will be local. For example, if you have a local disc ‘/scratch’, you can do the following:

```sh
unix> mkdir -p /scratch/$USER/pencil-data/samples/conv-slab
unix> ln -s /scratch/$USER/pencil-data/samples/conv-slab ./data
```

This is done automatically by the `pc_mkdatadir` command which, in turn, is invoked when making a new run directory with the `pc_newrun` command, for example.

Even if you don’t have an NFS-mounted directory (say, on your notebook computer), it is probably still a good idea to have code and data well separated by a scheme like the one described above.

An alternative to symbolic links, is to provide a file called ‘datadir.in’ in the root of the run directory. This file should contain one line of text specifying the absolute or relative data directory path to use. This facility is useful if one wishes to switch one run directory between different data directories. It is suggested that in such cases symbolic links are again made in the run directory to the various locations, then the ‘datadir.in’ need contain only a short relative path.

3.1.6 Running the code

You are now ready to start the code:

```sh
unix> start.csh
```

Linux cincinnatus 2.4.18-4GB #1 Wed Mar 27 13:57:05 UTC 2002 i686 unknown
Non-MPI version
datadir = data
Fri Aug 8 21:36:43 CEST 2003
```
cvs/start.x
CVS: io_dist.f90 v. 1.61 (brandenb ) 2003/08/03 09:26:55
[...]
CVS: start.in v. 1.4 (dobil ) 2002/10/02 20:11:14
```
nxgrid,nygrid,nzgrid= 32 32 32
thermodynamics: assume cp=1
```
uu: up-down
piecewise polytropic vertical stratification (lnrho)
init_lnrho: cs2bot,cs2top= 1.450000 0.3333330
  e.g. for ionization runs: cs2bot,cs2top not yet set
piecewise polytropic vertical stratification (ss)
The Pencil Code

start.x has completed successfully

Fri Aug 8 21:36:43 CEST 2003

This runs ‘src/start.x’ to construct an initial condition based on the parameters set in ‘start.in’. This initial condition is stored in ‘data/proc0/var.dat’ (and in ‘data/proc1/var.dat’, etc. if you run the multiprocessor version). It is fair to say that this is now a rather primitive routine; see ‘pencil-code/idl/read’ for various reading routines. You can then visualize the data using standard idl language.

If you visualize the profiles using IDL (see below), the result should bear some resemblance to Fig. [3] but with different values in the ghost zones (the correct values are set at run-time only) and a simpler velocity profile.

Now we run the code:

unix> run.csh

This executes ‘src/run.x’ and carries out nt time steps, where nt and other run-time parameters are specified in ‘run.in’. On a decent PC (1.7 GHz), 50 time steps take about 10 seconds.

The relevant part of the code’s output looks like

```
--it----t-------dt-------urms----umax----rhom------ssm-----dtc----dtu---dtnu---dtchi--
0  0.34  6.792E-03  0.0060  0.0452  14.4708  -0.4478  0.978  0.013  0.207  0.346
10  0.41  6.787E-03  0.0062  0.0440  14.4707  -0.4480  0.978  0.013  0.207  0.345
20  0.48  6.781E-03  0.0064  0.0429  14.4705  -0.4481  0.977  0.012  0.207  0.345
30  0.54  6.777E-03  0.0067  0.0408  14.4703  -0.4482  0.977  0.012  0.207  0.345
40  0.61  6.776E-03  0.0069  0.0381  14.4702  -0.4482  0.977  0.011  0.207  0.346
```

and lists

1. the number it of the current time step;
2. the time, t;
3. the time step, dt;
4. the rms velocity, urms = \sqrt{\langle u^2 \rangle};
5. the maximum velocity, umax = \max |u|;
6. the mean density, rhom = \langle \rho \rangle;
7. the mean entropy, ssm = \langle s \rangle / c_p;
8. the time step in units of the acoustic Courant step, dtc = \delta t c_{s0} / \delta x_{\min};
9. the time step in units of the advective time step, dtu = \delta t / (c_{\delta t} \delta x / \max |u|);
10. the time step in units of viscous time step, dtnu = \delta t / (c_{\delta t,v} \delta x^2 / \nu_{\max});
11. the time step in units of the conductive time step, dtchi = \delta t / (c_{\delta t,v} \delta x^2 / \chi_{\max}).

The entries in this list can be added, removed or reformatted in the file ‘print.in’, see Sects [5.5] and [K.3]. The output is also saved in ‘data/time_series.dat’ and should be identical to the content of ‘reference.out’.
3.2 Further tests

Figure 3: Stratification of the three-layer convection model in 'samples/conv-slab' after 50 timesteps ($t = 0.428$). Shown are (from left to right) density $\rho$, vertical velocity $u_z$, entropy $s/c_p$ and temperature $T$ as functions of the vertical coordinate $z$ for about ten different vertical lines in the computational box. The dashed lines denote domain boundaries: $z < -0.68$ is the lower ghost zone (points have no physical significance); $-0.68 < z < 0$ is a stably stratified layer ($ds/dz > 0$); $0 < z < 1$ is the unstable layer ($ds/dz < 0$); $1 < z < 1.32$ is the isothermal top layer; $z > 1.32$ is the upper ghost zone (points have no physical significance).

If you have *IDL*, you can visualize the stratification with (see Sect. 5.19.4 for details)

```
unix > idl
IDL > pc_read_var, obj=var, /trimall
IDL > tvscl, var, uu(*,*,0,0)
```

which shows $u_z$ in the $xy$ plane through the first meshpoint in the $z$ direction. There have been some now outdated specific routines that produce results like that shown in Fig. 3.

**Note:** If you want to run the code with *MPI*, you will probably need to adapt ‘getconf.csh’, which defines the commands and flags used to run MPI jobs (and which is sourced by the scripts ‘start.csh’ and ‘run.csh’). Try

```
csh -v getconf.csh
or
csh -x getconf.csh
```

to see how ‘getconf.csh’ makes its decisions. You would add a section for the host name of your machine with the particular settings. Since ‘getconf.csh’ is linked from the central directory ‘pencil-code/bin’, your changes will be useful for all your other runs too.

3.2 Further tests

There is a number of other tests in the ‘samples/’ directory. You can use the script ‘bin/auto-test’ to automatically run these tests and have the output compared to refer-
ence results.
4 Code structure

4.1 Directory tree

The overall directory structure of the code is shown in Fig. 4. Under `pencil-code`, there are currently the following files and directories:

```
bin/  config/  doc/  idl/  license/  perl/  samples/  sourceme.sh  utils/
bugs/  dx/  lib/  misc/  README  sourceme.csh  src/  www/
```

Almost all of the source code is contained in the directory `src/`, but in order to encapsulate individual applications, the code is compiled separately for each run in a local directory `src` below the individual run directory, like e.g. `samples/conv-slab/src`.

It may be a good idea to keep your own runs also under `svn` or `cvs` (which is older than but similar to `svn`), but this would normally be a different repository. On the machine where you are running the code, you may want to check them out into a subdirectory of
For example, we have our own runs in a repository called `pencil-runs`, so we do

```bash
unix> cd $PENCIL_HOME
unix> svn co runs pencil-runs
```

In this case, `runs` contains individual run directories, grouped in classes (like `spher` for spherical calculations, or `kinematic` for kinematic dynamo simulations). The current list of classes in our own `pencil-runs` repository is

```
1d-tests/  disc/  kinematic/  rings/
2d-tests/  discont/  Misc/  slab_conv/
3d-tests/  discussion/  OLD/  test/
buoy_tube/  forced/  pass_only/
convstar/  interstellar/  radiation/
```

The directory `forced/` contains some forced turbulence runs (both magnetic and non-magnetic); `gravz/` contains runs with vertical gravity; `rings/` contains decaying MHD problems (interlocked flux rings as initial condition, for example); and `kinematic/` contains kinematic dynamo problems where the hydrodynamics is turned off entirely. The file `samples/README` should contain an up-to-date list and short description of the individual classes.

The subdirectory `src` of each run directory contains a few local configuration files (currently these are `Makefile.local` and `cparam.local`) and possibly `ctimeavg.local`. To compile the samples, links the files `.f90`, `.c` and `Makefile.src` need to be linked from the top file[src]/src directory to the local directory `./src`. These links are set up by the script `pc_setupsrc` when used in the root of a run directory.

General-purpose visualization routines for **IDL** or **DX** are in the directories `idl` and `dx`, respectively. There are additional and more specialized **IDL** directories in the different branches under `pencil-runs`.

The directory `doc` contains this manual; `bin` contains a number of utility scripts (mostly written in csh and Perl), and in particular the `start.csh`, `run.csh`, and `getconf.csh` scripts. The `.svn` directory is used (you guessed it) by .svn, and is not normally directly accessed by the user; `bugs`, finally is used by us for internal purposes.

The files `sourceme.csh` and `sourceme.sh` will set up some environment variables — in particular `PATH` — and aliases/shell functions for your convenience. If you do not want to source one of these files, you need to make sure your **IDL** path is set appropriately (provided you want to use **IDL**) and you will need to address the scripts from `bin` with their explicit path name, or adjust your `PATH` manually.

### 4.2 Basic concepts

#### 4.2.1 Data access in pencils

Unlike the CRAY computers that dominated supercomputing in the 80s and early 90s, all modern computers have a cache that constitutes a significant bottleneck for many
codes. This is the case if large three-dimensional arrays are constantly used within each time step, which has the obvious advantage of working on long arrays and allows vectorization of elementary machine operations. This approach also implies conceptual simplicity of the code and allows extensive use of the intuitive F90 array syntax. However, a more cache-efficient way of coding is to calculate an entire time step (or substep of a multi-stage time-stepping scheme) only along a one-dimensional pencil of data within the numerical grid. This technique is more efficient for modern RISC processors: on Linux PCs and SGI workstations, for example, we have found a speed-up by about 60% in some cases. An additional advantage is a drastic reduction in temporary storage for auxiliary variables within each time step.

4.2.2 Modularity

Each run directory has a file `src/Makefile.local` in which you choose certain modules\(^2\), which tell the code whether or not entropy, magnetic fields, hydrodynamics, forcing, etc. should be invoked. For example, the settings for forced turbulent MHD simulations are

\begin{verbatim}
HYDRO = hydro
DENSITY = density
ENTROPY = noentropy
MAGNETIC = magnetic
GRAVITY = nogravity
FORCING = forcing

MPICOMM = nompicomm
GLOBAL = noglobal
IO = io_dist
FOURIER = nofourier
\end{verbatim}

This file will be processed by `make` and the settings are thus assignments of `make` variables. Apart from the physics modules (equation of motion: yes, density [pressure]: yes, entropy equation: no, magnetic fields: yes, gravity: no, forcing: yes), a few technical modules can also be used or deactivated; in the example above, these are MPI (switched off), additional global variables (none), input/output (distributed), and FFT (not used). The table in Sect.\(\text{J}\) in the Appendix lists all currently available modules.

Note that most of these `make` variables must be set, but they will normally obtain reasonable default values in `Makefile` (so you only need to set the non-standard ones in `Makefile.local`). It is by using this switching mechanism through `make` that we achieve high flexibility without resorting to excessive amounts of cryptic preprocessor directives or other switches within the code.

Many possible combinations of modules have already been tested and examples are part of the distribution, but you may be interested in a combination which was never tried before and which may not work yet, since the modules are not fully orthogonal. In such cases, we depend on user feedback for fixing problems and documenting the changes for others.

\(^2\)We stress once more that we are not talking about F90 modules here, although there is some connection, as most of our modules define F90 modules: For example each of the modules `gravity_simple`, `grav_r` and `nogravity` defines a Fortran module `Gravity`.\(\)
4.3 Files in the run directories

4.3.1 ‘start.in’, ‘run.in’, ‘print.in’

These files specify the startup and runtime parameters (see Sects. 5.12 and K.2), and the list of diagnostic variables to print (see 5.5). They specify the setup of a given simulation and are kept under svn in the individual ‘samples’ directories.

You may want to check for the correctness of these configuration files by issuing the command pc_configtest.

4.3.2 ‘datadir.in’

If this file exists, it must contain the name of an existing directory, which will be used as data directory, i.e. the directory where all results are written. If ‘datadir.in’ does not exist, the data directory is ‘data/’.

4.3.3 ‘reference.out’

If present, ‘reference.out’ contains the output you should obtain in the given run directory, provided you have not changed any parameters. To see whether the results of your run are OK, compare ‘time_series.dat’ to ‘reference.out’:

```
unix> diff data/time_series.dat reference.out
```  

4.3.4 ‘start.csh’, ‘run.csh’, ‘getconf.csh’ [obsolete; see Sect. 5.1]

These are links to ‘$PENCIL_HOME/bin’. You will be constantly using the scripts ‘start.csh’ and ‘run.csh’ to initialize the code. Things that are needed by both (like the name of the mpirun executable, MPI options, or the number of processors) are located in ‘getconf.csh’, which is never directly invoked.

4.3.5 ‘src/’

The ‘src’ directory contains two local files, ‘src/Makefile.local’ and ‘src/cparam.local’, which allow the user to choose individual modules (see 4.2.2) and to set parameters like the grid size and the number of processors for each direction. These two files are part of the setup of a given simulation and are kept under svn in the individual ‘samples’ directories.

The file ‘src/cparam.inc’ is automatically generated by the script ‘mkcparam’ and contains the number of fundamental variables for a given setup.

All other files in ‘src/’ are either links to source files (and ‘Makefile.src’) in the ‘$PENCIL_HOME/src’ directory, or object and module files generated by the compiler.
4.3 Files in the run directories

4.3.6 ‘data/’

This directory (the name of which will actually be overwritten by the first line of ‘datadir.in’, if that file is present; see \[4.3.2\]) contains the output from the code:

‘data/dim.dat’ The global array dimensions.

‘data/legend.dat’ The header line specifying the names of the diagnostic variables in ‘time_series.dat’.

‘data/time_series.dat’ Time series of diagnostic variables (also printed to stdout). You can use this file directly for plotting with Gnuplot, IDL, Xmgrace or similar tools (see also \[5.19\]).

‘data/tsnap.dat’, ‘data/tvid.dat’ Time when the next snapshot ‘\texttt{VAR_N}’ or animation slice should be taken.

‘data/params.log’ Keeps a log of all your parameters: ‘\texttt{start.x}’ writes the startup parameters to this file, ‘\texttt{run.x}’ appends the runtime parameters and appends them anew, each time you use the ‘\texttt{RELOAD}’ mechanism (see \[5.10\]).

‘data/param.nml’ Complete set of startup parameters, printed as Fortran namelist. This file is read in by ‘\texttt{run.x}’ (this is how values of startup parameters are propagated to ‘\texttt{run.x}’) and by IDL (if you use it).

‘data/param2.nml’ Complete set of runtime parameters, printed as Fortran namelist. This file is read by IDL (if you use it).

‘data/index.pro’ Can be used as include file in IDL and contains the column in which certain variables appear in the diagnostics file (‘time_series.dat’). It also contains the positions of variables in the ‘\texttt{VAR_N}’ files. These positions depend on whether entropy or noentropy, etc, are invoked. This is a temporary solution and the file may disappear in future releases.

‘data/interstellar.dat’ Unformatted file containing the time at which the next supernova event will occur, under certain supernova schemes. (Only needed by the interstellar module.)

‘data/proc0’, ‘data/proc1’, … These are the directories containing data from the individual processors. So after running an MPI job on two processors, you will have the two directories ‘data/proc0’ and ‘data/proc1’. Each of the directories can contain the following files:

‘var.dat’ binary file containing the latest snapshot;
‘VAR.N’ binary file containing individual snapshot number $N$;
‘dim.dat’ ASCII file containing the array dimensions as seen by the given processor;
‘time.dat’ ASCII file containing the time corresponding to ‘var.dat’ (not actually used by the code, unless you use the io_mpioidist.f90 module);
‘grid.dat’ binary file containing the part of the grid seen by the given processor;
‘seed.dat’ the random seed for the next time step (saved for reasons of reproducibility).
   For multi-processor runs with velocity forcing, the files ‘procN/seed.dat’ must all contain the same numbers, because globally coherent waves of given wavenumber are used;
‘X.xy’, ‘X.xz’, ‘X.yz’ two-dimensional sections of variable $X$, where $X$ stands for the corresponding variable. The current list includes
   bx.xy bx.xz by.xy by.xz bz.xy bz.xz divu.xy lnrho.xz
   ss.xz ux.xy ux.xz uz.xy uz.xz
   Each processor writes its own slice, so these need to be reassembled if one wants to plot a full slice.
5 Using the code

5.1 Configuring the code to compile and run on your computer

Note: We recommend to use the procedure described here, rather than the old method described in Sects. 5.2 and 4.3.4.

Quick instructions: You may compile with a default compiler-specific configuration:

1. Single-processor using the GNU compiler collection:
   
   unix> pc_build -f GNU-GCC

2. Multi-processor using GNU with MPI support:

   unix> pc_build -f GNU-GCC_MPI

Many compilers are supported already, please refer to the available config files in ‘$PENCIL_HOME/config/compilers/*.conf’, e.g. ‘Intel.conf’ and ‘Intel_MPI.conf’.

If you have to set up some compiler options specific to a certain host system you work on, or if you like to create a host-specific configuration file so that you can simply execute pc_build without any options, you can clone an existing host-file, just include an existing compiler configuration, and simply only add the options you need. A good example of a host-file is ‘$PENCIL_HOME/config/hosts/IWF/host-andromeda-GNU_Linux-Linux.conf’. You may save a clone under ‘$PENCIL_HOME/config/hosts/<ID>.conf’, where ‘<ID>’ is to be replaced by the output of pc_build -i. This will be the new default for pc_build.

If you don’t know what this was all about, read on.

In essence, configuration, compiling and running the code work like this:

1. Create a configuration file for your computer’s host ID.
2. Compile the code using pc_build.
3. Run the code using pc_run

In the following, we will discuss the essentials of this scheme. Exhaustive documentation is available with the commands perldoc Pencil::ConfigFinder and perldoc PENCIL::ConfigParser.

5.1.1 Locating the configuration file

Commands like pc_build and pc_run use the Perl module ‘Pencil::ConfigFinder’ to locate an appropriate configuration file and ‘Pencil::ConfigParser’ to read and interpret it. When you use ‘ConfigFinder’ on a given computer, it constructs a host ID for the system it is running on, and looks for a file ‘host_ID.conf’ in any subdirectory of ‘$PENCIL_HOME/config/hosts’.

E.g., if the host ID is “workhorse.pencil.org”, ‘ConfigFinder’ would consider the file ‘$PENCIL_HOME/config/hosts/pencil.org/workhorse.pencil.org.conf’.
**Note 1:** The location in the tree under `hosts/` is irrelevant, which allows you to group related hosts by institution, owner, hardware, etc.

**Note 2:** ‘ConfigFinder’ actually uses the following search path:

1. `./config`
2. `$PENCIL_HOME/config-local`
3. `$HOME/.pencil/config-local`
4. `$PENCIL_HOME/config`

This allows you to override part of the `config/` tree globally on the given file system, or locally for a particular run directory, or for one given copy of the PENCIL CODE. This search path is used both, for locating the configuration file for your host ID, and for locating included files (see below).

The host ID is constructed based on information that is easily available for your system. The algorithm is as follows:

1. Most commands using ‘ConfigFinder’ have a `--host-id` (sometimes abbreviated as `-H`) option to explicitly set the host ID.
2. If the environment variable `PENCIL_HOST_ID` is set, its value is used.
3. If any of the files `./host-ID`, `$PENCIL_HOME/host-ID`, `$HOME/.pencil/host-ID`, exist, its first line is used.
4. If ‘ConfigFinder’ can get hold of a fully qualified host name, that is used as host ID.
5. Else, a combination of host name, operating system name and possibly some other information characterizing the system is used.
6. If no config file for the host ID is found, the current operating system name is tried as fallback host ID.

To see which host IDs are tried (up to the first one for which a configuration file is found), run

```
unix> pc_build --debug-config
```

This command will tell you the `host-ID` of the system that you are using:

```
unix> pc_build -i
```

### 5.1.2 Structure of configuration files

It is strongly recommended to include in a users configuration file one of the preset compiler suite configuration files. Then, only minor options need to be set by a user, e.g. the optimization flags. One of those user configuration files looks rather simple:

```
# Simple config file. Most files don't need more.
%include compilers/GNU-GCC
```

or if you prefer a different compiler:
# Simple Intel compiler suite config file.
%include compilers/Intel

A more complex file (using MPI with additional options) would look like this:

# More complex config file.
%include compilers/GNU-GCC_MPI

%section Makefile
   MAKE_VAR1 = -j4 # joined compilation with four threads
   FFLAGS += -O3 -Wall -fbacktrace # don’t redefine, but append with ‘+=’
%endsection Makefile

%section runtime
   mpiexec = mpirun # some MPI backends need a redefinition of mpiexec
%endsection runtime

%section environment
   SCRATCH_DIR=/var/tmp/$USER
%endsection environment

Adding “_MPI” to a compiler suite’s name is usually sufficient to use MPI.

**Note 3:** We strongly advise *not* to mix Fortran- and C-compilers from different manufacturers or versions by manually including multiple separate compiler configurations.

**Note 4:** We strongly advise to use *at maximum* the optimization levels `-O2` for the Intel compiler and `-O3` for all other compilers. Higher optimization levels implicate an inadequate loss of precision.

The `.conf` files consist of the following elements:

**Comments:** A # sign and any text following it on the same line are ignored.

**Sections:** There are three sections:

- **Makefile** for setting make parameters
- **runtime** for adding compiler flags used by pc_run
- **environment** shell environment variables for compilation and running

**Include statements:** An %include ... statement is recursively replaced by the contents of the files it points to.

The included path gets a `.conf` suffix appended. Included paths are typically “absolute”, e.g.

%include os/Unix

will include the file ‘os/Unix.conf’ in the search path listed above (typically from ‘$PENCIL_HOME/config’). However, if the included path starts with a dot, it is a relative path, so

---

3However, if the include statement is inside a section, only the file’s contents inside that section are inserted.
%include ./Unix

will only search in the directory where the including file is located.

**Assignments:** Statements like FFLAGS += -03 or mpiexec=mpirun are assignments and will set parameters that are used by pc_build/make or pc_run.

Lines ending with a backslash '\' are continuation lines.

If possible, one should always use *incremental assignments*, indicated by using a += sign instead of =, instead of redefining certain flags.

Thus,

```
CFLAGS += -03
CFLAGS += -I../include -Wall
```

is the same as

```
CFLAGS = $(CFLAGS) -O3 -I../include -Wall
```

### 5.1.3 Compiling the code

Use the pc_build command to compile the code:

```
unix> pc_build       # use default compiler suite
unix> pc_build -f Intel_MPI  # specify a compiler suite
unix> pc_build -f os/GNU_Linux,mpi/open-mpi  # explicitly specify config files
unix> pc_build VAR=something  # set variables for the makefile
unix> pc_build --cleanall  # remove generated files
```

The third example circumvents the whole host ID mechanism by explicitly instructing pc_build which configuration files to use. The fourth example shows how to define extra variables (VAR=something) for the execution of the Makefile.

See pc_build --help for a complete list of options.

### 5.1.4 Running the code

Use the pc_run command to run the code:

```
unix> pc_run          # start if necessary, then run
unix> pc_run start
unix> pc_run run

unix> pc_run start run~3 # start, then run 3 times
unix> pc_run start run run run # start, then run 3 times
unix> pc_run ^3        # start if necessary, then run 3 times
```

See pc_run --help for a complete list of options.

### 5.1.5 Testing the code

The pc_auto-test command uses pc_build for compiling and pc_run for running the tests. Here are a few useful options:
unix> pc_auto-test
unix> pc_auto-test --no-pencil-check # suppress pencil consistency check
unix> pc_auto-test --max-level=1  # run only tests in level 0 and 1
unix> pc_auto-test --time-limit=2m  # kill each test after 2 minutes

See pc_auto-test --help for a complete list of options.

The pencil-test script will use pc_auto-test if given the ‘--use-pc_auto-test’ or ‘-b’ option:

unix> pencil-test --use-pc_auto-test
unix> pencil-test -b                # ditto
unix> pencil-test -b -Wa,--max-level=1,--no-pencil-check  # quick pencil

See pencil-test --help for a complete list of options.

5.2 Adapting ’Makefile.src’ [obsolete; see Sect. 5.1]

By default, one should use the above described configuration mechanism for compilation. If for whatever reason one needs to work with a modified ‘Makefile’, there is a mechanism for picking the right set of compiler flags based on the hostname.

To give you an idea of how to add your own machines, let us assume you have several Linux boxes running a compiler f95 that needs the options ‘-O2 -u’, while one of them, Janus, runs a compiler f90 which needs the flags ‘-O3’ and requires the additional options ‘-lmpi -llam’ for MPI.

The ‘Makefile.src’ you need will have the following section:

```makefile
### Begin machine dependent

## IRIX64:
[... ] (leave as it is in the original Makefile)
## OSF1:
[... ] (leave as it is in the original Makefile)

## Linux:
[... ] (leave everything from original Makefile and add:)
#FC=f95
#FFLAGS=-O2 -u
#FC=f90            #(Janus)
#FFLAGS=-O3        #(Janus)
#LDMPI=-lmpi -llam #(Janus)

## SunOS:
[... ] (leave as it is in the original Makefile)
## UNICOS/mk:
[... ] (leave as it is in the original Makefile)
## HI-UX/MPP:
[... ] (leave as it is in the original Makefile)
## AIX:
[... ] (leave as it is in the original Makefile)
```
Table 1: Compiler flags for common compilers. Note that some combinations of OS and compiler require much more elaborate settings; also, if you use MPI, you will have to set LDMPI.

<table>
<thead>
<tr>
<th>Compiler</th>
<th>FC</th>
<th>FFLAGS</th>
<th>CC</th>
<th>CFLAGS</th>
</tr>
</thead>
<tbody>
<tr>
<td>Unix/POSIX:</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>GNU</td>
<td>gfortran</td>
<td>-O3</td>
<td>gcc</td>
<td>-O3 -DFUNDERSC=1</td>
</tr>
<tr>
<td>Intel</td>
<td>ifort</td>
<td>-O2</td>
<td>icc</td>
<td>-O3 -DFUNDERSC=1</td>
</tr>
<tr>
<td>PGI</td>
<td>pgf95</td>
<td>-O3</td>
<td>pgcc</td>
<td>-O3 -DFUNDERSC=1</td>
</tr>
<tr>
<td>G95</td>
<td>g95</td>
<td>-O3 -fno-second-underscore</td>
<td>gcc</td>
<td>-O3 -DFUNDERSC=1</td>
</tr>
<tr>
<td>Absoft</td>
<td>f95</td>
<td>-O3 -N15</td>
<td>gcc</td>
<td>-O3 -DFUNDERSC=1</td>
</tr>
<tr>
<td>IBM XL</td>
<td>xlf95</td>
<td>-qsuffix=f=f90 -O3</td>
<td>xlc</td>
<td>-O3 -DFUNDERSC=1</td>
</tr>
<tr>
<td>outdated:</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>IRIX Mips</td>
<td>f90</td>
<td>-64 -O3 -mips4</td>
<td>cc</td>
<td>-O3 -64 -DFUNDERSC=1</td>
</tr>
<tr>
<td>Compaq</td>
<td>f90</td>
<td>-fast -O3</td>
<td>cc</td>
<td>-O3 -DFUNDERSC=1</td>
</tr>
</tbody>
</table>

### End machine dependent

**Note 1** There is a script for adapting the Makefile: ‘adapt.mkfile’. In the example above, #(Janus) is not a comment, but marks this line to be activated (uncommented) by adapt.mkfile if your hostname (‘uname -n’) matches ‘Janus’ or ‘janus’ (capitalization is irrelevant). You can combine machine names with a vertical bar: a line containing #(onsager|Janus) will be activated on both, Janus and Onsager.

**Note 2** If you want to experiment with compiler flags, or if you want to get things running without setting up the machine-dependent section of the ‘Makefile’, you can set make variables at the command line in the usual manner:

```
src> make FC=f90 FFLAGS='-fast -u'
```

will use the compiler f90 and the flags ‘-fast -u’ for both compilation and linking. Table 1 summarizes flags we use for common compilers.

### 5.3 Changing the resolution

It is advisable to produce a new run directory each time you run a new case. (This does not include restarts from an old run, of course.) If you have a 32\(^3\) run in some directory ‘runA_32a’, then go to its parent directory, i.e.

```
runA_32a> cd ..
forced> pc_newrun runA_32a runA_64a
forced> cd runA_64a/src
forced> vi cparam.local
```

and edit the ‘cparam.local’ for the new resolution.

If you have ever wondered why we don’t do dynamic allocation of the main variable (f) array, the main reason it that with static allocation the compiler can check whether we are out of bounds.
5.4 Using a non-equidistant grid

We introduce a non-equidistant grid $z_i$ (by now, this is also introduced for the other directions) as a function $z(\zeta)$ of an equidistant grid $\zeta$, with grid spacing $\Delta \zeta = 1$.

The way the parameters are handled, the box size and position are not changed when you switch to a non-equidistant grid, i.e., they are still determined by $\text{xyz}0$ and $L_{\text{xyz}}$.

The first and second derivatives can be calculated by

$$\frac{df}{dz} = \frac{df}{d\zeta} \frac{d\zeta}{dz} = \frac{1}{z'} f'(\zeta), \quad \frac{d^2f}{dz^2} = \frac{1}{z'^2} f''(\zeta) - \frac{z''}{z'^3} f'(\zeta),$$

which can be written somewhat more compactly using the inverse function $\zeta(z)$:

$$\frac{df}{dz} = \zeta'(z) f'(\zeta), \quad \frac{d^2f}{dz^2} = \zeta''(z) f''(\zeta) + \zeta''(z) \zeta'(z) f'(\zeta).$$

Internally, the code uses the quantities $dz_1 \equiv 1/z' = \zeta'(z)$ and $dz_{\text{tilde}} \equiv -z''/z'^2 = \zeta''/\zeta'$, and stores them in 'data/procN/grid.dat'.

The parameters $\text{lequidist}$ (a 3-element logical array), $\text{grid func}$, $\text{coeff grid}$ (a $\geq 2$-element real array) are used to choose a non-equidistant grid and define the function $z(\zeta)$. So far, one can choose between grid function='sinh', grid_=function='linear' (which produces an equidistant grid for testing purposes), and grid_function='step-linear'.

**The sinh profile:** For grid_function='sinh', the function $z(\zeta)$ is given by

$$z(\zeta) = z_0 + L_z \frac{\sinh a(\zeta-\zeta_1) + \sinh a(\zeta-\zeta_1)}{\sinh a(\zeta_2-\zeta_1) + \sinh a(\zeta-\zeta_1)},$$

where $z_0$ and $z_0 + L_z$ are the lowest and uppermost levels, $\zeta_1, \zeta_2$ are the $\zeta$ values representing those levels (normally $\zeta_1 = 0, \zeta_2 = N_z - 1$ for a grid of $N_z$ vertical layers [excluding ghost layers]), and $\zeta_*$ is the $\zeta$ value of the inflection point of the sinh function. The $z$ coordinate and $\zeta$ value of the inflection point are related via

$$z_* = z_0 + L_z \frac{\sinh a(\zeta_*-\zeta_1)}{\sinh a(\zeta_2-\zeta_1) + \sinh a(\zeta_*-\zeta_1)},$$

which can be inverted (“after some algebra”) to

$$\zeta_* = \frac{\zeta_1 + \zeta_2}{2} + \frac{1}{a} \text{artanh} \left[ \left( \frac{2 z_* - z_0}{L_z} - 1 \right) \tanh \frac{\zeta_2 - \zeta_1}{2} \right].$$

**General profile:** For a general monotonic function $\psi()$ instead of sinh we get,

$$z(\zeta) = z_0 + L_z \frac{\psi[a(\zeta-\zeta_1)] + \psi[a(\zeta-\zeta_1)]}{\psi[a(\zeta_2-\zeta_1)] + \psi[a(\zeta_*-\zeta_1)]},$$

and the reference point $\zeta_*$ is found by inverting

$$z_* = z_0 + L_z \frac{\psi[a(\zeta_*-\zeta_1)]}{\psi[a(\zeta_2-\zeta_1)] + \psi[a(\zeta_*-\zeta_1)]},$$

numerically.
**Duct flow:** The profile function `grid_function='duct'` generates a grid profile for turbulent flow in ducts. For a duct flow, most gradients are steepest close to the walls, and hence very fine resolution is required near the walls. Here we follow the method of [24] and use a Chebyshev-type grid with a cosine distribution of the grid points such that in the \( y \) direction they are located at

\[
y_j = h \cos \theta_j ,
\]

where

\[
\theta_j = \frac{(N_y-j) \pi}{N_y-1}, \quad j = 1,2,\ldots,N_y
\]

and \( h = L_y/2 \) is the duct half-width.

Currently this method is only adapted for ducts where \( x \) is the stream-wise direction, \( z \) is in the span-wise direction and the walls are at \( y = y_0 \) and \( y = y_0 + L_y \).

In order to have fine enough resolution, the first grid point should be a distance \( \delta = 0.05l_w \) from the wall, where

\[
l_w = \frac{\nu}{u_\tau}, \quad u_\tau = \sqrt{\tau_{\text{wall}}} \rho ,
\]

and \( \tau_{\text{wall}} \) is the shear wall stress. This is accomplished by using at least

\[
N_y \geq N_y^* = \frac{\pi}{\arccos \left( 1 - \frac{\delta}{h} \right)} + 1
\]

\[
= \pi \sqrt{\frac{h}{2 \delta}} + 1 - \frac{\pi}{24} \sqrt{\frac{2 \delta}{h}} + O \left[ \left( \frac{\delta}{h} \right)^{3/2} \right]
\]

grid points in the \( y \)-direction. After rounding up to the next integer value, we find that the truncated condition

\[
N_y \geq \left\lceil \pi \sqrt{\frac{h}{2 \delta}} \right\rceil + 1
\]

(where \( \lceil x \rceil \) denotes the ceiling function, i.e. the smallest integer equal to, or larger than, \( x \)) gives practically identical results.

**Example:** To apply the sinh profile, you can set the following in ‘\texttt{start.in}’ (this example is from ‘\texttt{samples/sound-spherical-noequi}’):

```plaintext
&init_pars
  ...
  xyz0 = -2., -2., -2.  ! first corner of box
  Lxyz = 4., 4., 4.  ! box size
  lperi = F, F, F  ! periodic direction?
  lequidist = F, F, T  ! non-equidistant grid in z
  xyz_star = , , -2.  ! position of inflection point
  grid_func = , , 'sinh'  ! sinh function: linear for small, but
  \hspace{1cm} exponential for large z
  coeff_grid = , , 0.5
/
```

```
The parameter array \textit{coeff\_grid} represents \(z_s\) and \(a\); the bottom height \(z_0\) and the total box height \(L_z\) are taken from \textit{xyz0} and \textit{Lxyz} as in the equidistant case. The inflection point of the \textit{sinh} profile (the part where it is linear) is not in the middle of the box, because we have set \textit{xyz\_star}(3) (i.e. \(z_s\)) to \(-2\).

5.5 Diagnostic output

Every \textit{it1} time steps (\textit{it1} is a runtime parameter, see Sect. \[K.2\]), the code writes monitoring output to \textit{stdout} and, parallel to this, to the file ‘\textit{data/time\_series.dat}’. The variables that appear in this listing and the output format are defined in the file ‘\textit{print.in}’ and can be changed without touching the code (even while the code is running). A simple example of ‘\textit{print.in}’ may look like this:

\begin{verbatim}
t(F10.3)
urms(E13.4)
rhom(F10.5)
oum
\end{verbatim}

which means that the output table will contain time \(t\) in the first column formatted as (F10.3), followed by the mean squared velocity, \(urms\) (i.e. \(\langle u^2 \rangle^{1/2}\)) in the second column with format (E13.4), the average density \(rhom\) (\(\langle \rho \rangle\), which allows to monitor mass conservation) formatted (F10.5) and the kinetic helicity \(oum\) (that is \(\langle \omega \cdot u \rangle\)) in the last column with the default format (E10.2).

The corresponding diagnostic output will look like

\begin{verbatim}
----t---------urms--------rhom------oum----
 0.000  4.9643E-03  14.42457  -8.62E-06
 0.032  3.9423E-03  14.42446  -5.25E-06
 0.063  6.8399E-03  14.42449  -3.50E-06
 0.095  1.1437E-02  14.42455  -2.58E-06
 0.126  1.6980E-02  14.42457  -1.93E-06
\end{verbatim}

5.6 Data files

5.6.1 Snapshot files

Snapshot files contain the values of all evolving variables and are sufficient to restart a run. In the case of an MHD simulation with entropy equation, for example, the snapshot files will contain the values of velocity, logarithmic density, entropy and the magnetic vector potential.

There are two kinds of snapshot files: the current snapshot and permanent snapshots, both of which reside in the directory ‘\textit{data/procN/}’. The parameter \textit{isav} determines the frequency at which the current snapshot ‘\textit{data/procN/var\_dat}’ is written. If you keep this frequency too high, the code will spend a lot of time on I/O, in particular for large jobs; too low a frequency makes it difficult to follow the evolution interactively during test runs. There is also the \textit{ialive} parameter. Setting this to 1 or 10 gives an updated

\footnote{The format specifiers are like in Fortran, apart from the fact that the E format will use standard scientific format, corresponding to the Fortran 1pE syntax. Seasoned Fortran IV programmers may use formats like (0pE13.4) to enjoy nostalgic feelings, or (1pF10.5) if they depend on getting wrong numbers.}
timestep in the files ‘data/proc*/alive.info’. You can put ialive=0 to turn this off to limit the I/O on your machine.

The permanent snapshots ‘data/proc*/VAR’ are written every dsnap time units. These files are numbered consecutively from \( N = 1 \) upward and for long runs they can occupy quite some disk space. On the other hand, if after a run you realize that some additional quantity \( q \) would have been important to print out, these files are the only way to reconstruct the time evolution of \( q \) without re-running the code.

**File structure**  Snapshot files consist of the following *Fortran records*:

1. variable vector \( f [mx \times my \times mz \times nvar] \)

All numbers (apart from the record markers) are single precision (4-byte) floating point numbers, unless you use double precision (see §5.21, in which case all numbers are 8-byte floating point numbers, while the record markers remain 4-byte integers.

The script `pc_tsnap` allows you to determine the time \( t \) of a snapshot file:

```plaintext
unix> pc_tsnap data/proc0/var.dat
data/proc0/var.dat: t = 8.32456
unix> pc_tsnap data/proc0/VAR2
data/proc0/VAR2: t = 2.00603
```

### 5.7 Video files and slices

We use the terms *video files* and *slice files* interchangeably. These files contain a time series of values of one variable in a given plane. The output frequency of these video snapshots is set by the parameter `dvid` (in code time units).

When output to video files is activated by some settings in ‘run.in’ (see example below) and the existence of ‘video.in’, slices are written for four planes:

1. \( x-z \) plane (\( y \) index \( iy \); file suffix `.xz`)
2. \( y-z \) plane (\( y \) index \( ix \); suffix `.yz`)
3. \( x-y \) plane (\( y \) index \( iz \); suffix `.xy`)
4. another slice parallel to the \( x-y \) plane (\( y \) index \( iz2 \); suffix `.xy2`)

You can specify the position of the slice planes by setting the parameters `ix`, `iy`, `iz` and `iz2` in the namelist `run_pars` in ‘run.in’. Alternatively, you can set the input parameter `slice_position` to one of ‘p’ (periphery of box) or ‘m’ (middle of box). Or you can also specify the \( z \)–position in terms of \( z \) using the tags `zbot_slice` and `ztop_slice`. In this case, the `zbot_slice` slice will have the suffix `.xy` and the `ztop_slice` the suffix `.xy2`

In the file ‘video.in’ of your run directory, you can choose for which variables you want to get video files; valid choices are listed in §K.4.

---

5 A *Fortran record* is marked by the 4-byte integer byte count of the data in the record at the beginning and the end, i.e. has the form \((N_{\text{bytes}}, \text{raw data}, N_{\text{bytes}})\).
The slice files are written in each processor directory ‘data/proc*/’ and have a file name indicating the individual variable (e.g. ‘slice_uu1.yz’ for a slice of \( u_x \) in the \( y-z \) plane). Before visualizing slices one normally wants to combine the sub-slices written by each processor into one global slice (for each plane and variable). This is done by running ‘src/read_videofiles.x’, which will prompt for the variable name, read the individual sub-slices and write global slices to ‘data/’ Once all global slices have been assembled you may want to remove the local slices ‘data/proc*/slice*’.

To read all sub-slices demanded in ‘video.in’ at once use ‘src/read_all_videofiles.x’. This program doesn’t expect any user input and can thus be submitted in computing queues.

For visualization of slices, you can use the IDL routines ‘rvid_box.pro’, ‘rvid_-plane.pro’, or ‘rvid_line.pro’ which allows the flag ‘/png’ for writing PNG images that can then be combined into an MPEG movie using mpeg_encode. Based on ‘rvid_box’, you can write your own video routines in IDL.

An example  Suppose you have set up a run using entropy.f90 and magnetic.f90 (most probably together with hydro.f90 and other modules). In order to animate slices of entropy \( s \) and the \( z \)-component \( B_z \) of the magnetic field, in planes passing through the center of the box, do the following:

1. Write the following lines to ‘video.in’ in your run directory:

   \[
   \text{ss} \\
   \text{bb} \\
   \text{divu}
   \]

2. Edit the namelist run_pars in the file ‘run.in’. Request slices by setting write_slices and set dvid and slice_position to reasonable values, say

   !write_slices=T !(no longer works; write requested slices into video.in) 
   dvid=0.05 
   slice_position='m'

3. Run the PENCIL CODE:

   unix> start.csh 
   unix> run.csh

4. Say ‘make read_videofiles’ to compile the routine and then run ‘src/read_videofiles.x’ to assemble global slice files from those scattered across ‘data/proc*/’:

   unix> src/read_videofiles.x 
   enter name of variable (lnrho, uu1, ..., bb3): ss 
   unix> src/read_videofiles.x 
   enter name of variable (lnrho, uu1, ..., bb3): bb3

5. Start IDL and run ‘rvid_box’:

   unix> idl 
   IDL> rvid_box,’bb3’ 
   IDL> rvid_box,’ss’,min=-0.3,max=2.
Another example  Suppose you have set up a run using \textit{magnetic.f90} and some other modules. This run studies some process in a “surface” layer inside the box. This “surface” can represent a sharp change in density or turbulence. So you defined your box setting the $z = 0$ point at the surface. Therefore, your ‘\textit{start.in}’ file will look something similar to:

```
&init_pars
lperi=T,T,F
bcz = 's','s','a','hs','s','s','a'
xyz0 = -3.14159, -3.14159, -3.14159
Lxyz = 6.28319, 6.28319, 9.42478
```

A smarter way of specifying the box size in units of $\pi$ is to write

```
&init_pars
xyz_units = 'pi', 'pi', 'pi'
xyz0 = -1., -1., -1.
Lxyz = 2., 2., 2.
```

Now you can analyze quickly the surface of interest and some other $xy$ slice setting $zbot\_slice$ and $ztop\_slice$ in the ‘\textit{run.in}’ file:

```
&run_pars
slice_position='c'
zbot\_slice=0.
ztop\_slice=0.2
```

In this case, the slices with the suffix .xy will be at the “surface” and the ones with the suffix .xy2 will be at the position $z = 0.2$ above the surface. And you can visualize this slices by:

1. Write the following lines to ‘\textit{video.in}’ in your run directory:

```
bb
```

2. Edit the namelist \textit{run_pars} in the file ‘\textit{run.in}’ to include $zbot\_slice$ and $ztop\_slice$.

3. Run the PENCIL CODE:

```
unix> start.csh
unix> run.csh
```

4. Run ‘\textit{src/read_videofiles.x}’ to assemble global slice files from those scattered across ‘\textit{data/proc/*}’:

```
unix> src/read_videofiles.x
    enter name of variable (lnrho, uu1, ..., bb3):  bb3
```

5. Start IDL, load the slices with ‘\textit{pc\_read_video}’ and plot them at some time:

```
unix> idl
IDL> pc\_read_video, field='bb3',ob=bb3,nt=ntv
IDL> tvsc1,bb3.xy(*,*,100)
IDL> tvsc1,bb3.xy2(*,*,100)
```
etc.

File structure  Slice files consist of one Fortran record (see footnote on page 26) for each slice, which contains the data of the variable (without ghost zones), the time \( t \) of the snapshot and the position of the sliced variable (e.g. the \( x \) position for a \( y-z \) slice):

1. data\(_1\) \([nx \times ny \times nz]\), time \( t_1\) [1], position\(_1\) [1]
2. data\(_2\) \([nx \times ny \times nz]\), time \( t_2\) [1], position\(_2\) [1]
3. data\(_3\) \([nx \times ny \times nz]\), time \( t_3\) [1], position\(_3\) [1]
   etc.

5.8 Averages

5.8.1 One-dimensional output averaged in two dimensions

In the file ‘xyaver.in’, \( z \)-dependent (horizontal) averages are listed. They are written to the file ‘data/xyaverages.dat’. A new line of averages is written every \( it1\)th time steps.

There is the possibility to output two-dimensional averages. The result then depends on the remaining dimension. The averages are listed in the files ‘xyaver.in’, ‘xzaver.in’, and ‘yzaver.in’ where the first letters indicate the averaging directions. The output is then stored to the files ‘data/xyaverages.dat’, ‘data/xzaverages.dat’, and ‘data/yzaverages.dat’. The output is written every \( it1\)th time steps.

The rms values of the so defined mean magnetic fields are referred to as \( bmz \), \( bmy \) and \( bmx \), respectively, and the rms values of the so defined mean velocity fields are referred to as \( umz \), \( umy \), and \( umx \). (The last letter indicates the direction on which the averaged quantity still depends.)

See Sect. 9.2 on how to add new averages.

In \texttt{idl} such \( xy \)-averages can be read using the procedure ‘\texttt{pc_read_xyaver}’.

5.8.2 Two-dimensional output averaged in one dimension

There is the possibility to output one-dimensional averages. The result then depends on the remaining two dimensions. The averages are listed in the files ‘yafer.in’, ‘zafer.in’, and ‘phiaver.in’ where the first letter indicates the averaging direction. The output is then stored to the files ‘data/yaferages.dat’, ‘data/zaferages.dat’, and ‘data/phiaverages.dat’.

See Sect. 9.2 on how to add new averages.

Disadvantage: The output files, e.g. ‘data/zaferages.dat’, can be rather big because each average is just appended to the file.
5.8.3 Azimuthal averages

Azimuthal averages are controlled by the file ‘phiaver.in’, which currently supports the quantities listed in Sect. K.5. In addition, one needs to set lwrite_phiaverages, lwrite_yaverages, or lwrite_zaverages to .true.. For example, if ‘phiaver.in’ contains the single line

\begin{verbatim}
b2mphi
\end{verbatim}

then you will get azimuthal averages of the squared magnetic field \( B^2 \).

Azimuthal averages are written every \( d2davg \) time units to the files ‘data/averages/PHIAVG1’. The file format of azimuthal-average files consists of the following Fortran records:

1. number of radial points \( N_{r,\phi-avg} \) [1], number of vertical points \( N_{z,\phi-avg} \) [1], number of variables \( N_{var,\phi-avg} \) [1], number of processors in \( z \) direction [1]
2. time \( t \) [1], positions of cylindrical radius \( r_{cyl} [N_{r,\phi-avg}] \) and \( z [N_{z,\phi-avg}] \) for the grid, radial spacing \( \delta r_{cyl} \) [1], vertical spacing \( \delta z \) [1]
3. averaged data \( [N_{r,\phi-avg} \times N_{z,\phi-avg}] \)
4. label length [1], labels of averaged variables \( [N_{var,\phi-avg}] \)

All numbers are 4-byte numbers (floating-point numbers or integers), unless you use double precision (see § K.21).

To read and visualize azimuthal averages in IDL, use ‘$PENCIL_HOME/idl/files/pc_read_phiavg.pro’

\begin{verbatim}
IDL> avg = pc_read_phiavg('data/averages/PHIAVG1')
IDL> contour, avg.b2mphi, avg.rcyl, avg.z, TITLE='!17B!U2!N!X'
\end{verbatim}

or have a look at ‘$PENCIL_HOME/idl/phiavg.pro’ for a more sophisticated example.

5.8.4 Time averages

Time averages need to be prepared in the file ‘src/ctimeavg.local’, since they use extra memory. They are controlled by the averaging time \( \tau_{avg} \) (set by the parameter \( tavg \) in ‘run.in’), and by the indices \( idx_{tavg} \) of variables to average.

Currently, averaging is implemented as exponential (memory-less) average\(^6\)

\begin{equation}
\langle f \rangle_{t+\delta t} = \langle f \rangle_t + \frac{\delta t}{\tau_{avg}} [f(t+\delta t) - \langle f \rangle_t],
\end{equation}

\(6\) At some point we may also implement the more straight-forward average

\begin{equation}
\langle f \rangle_{t+\delta t} = \langle f \rangle_t + \frac{\delta t}{t-t_0+\delta t} [f(t+\delta t) - \langle f \rangle_t],
\end{equation}

which is equivalent to

\begin{equation}
\langle f \rangle_t = \frac{1}{t-t_0} \int_{t_0}^{t} f(t') \, dt',
\end{equation}

but we do not expect large differences.
which is equivalent to

\[ \langle f \rangle_t = \int_{t_0}^{t} e^{-(t-t')/\tau_{\text{avg}}} f(t') \, dt'. \]  \hspace{1cm} (17)

Here \( t_0 \) is the time of the snapshot the calculation started with, i.e. the snapshot read by the last \texttt{run.x} command. Note that the implementation (16) will approximate Eq. (17) only to first-order accuracy in \( \delta t \). In practice, however, \( \delta t \) is small enough to make this accuracy suffice.

In ‘\texttt{src/ctimeavg.local}’, you need to set the number of slots used for time averages. Each of these slots uses \( mx \times my \times mz \) floating-point numbers, i.e. half as much memory as each fundamental variable.

For example, if you want to get time averages of all variables, set

\[
\text{integer, parameter :: } \text{mtavg} = \text{mvar}
\]

in ‘\texttt{src/ctimeavg.local}’, and don’t set \texttt{idx_tavg} in ‘\texttt{run.in}’.

If you are only interested in averages of variables 1–3 and 6–8 (say, the velocity vector and the magnetic vector potential in a run with ‘\texttt{hydro.f90}’, ‘\texttt{density.f90}’, ‘\texttt{entropy.f90}’ and ‘\texttt{magnetic.f90}’), then set

\[
\text{integer, parameter :: } \text{mtavg} = 6
\]

in ‘\texttt{src/ctimeavg.local}’, and set

\[
\text{idx_tavg} = 1,2,3,6,7,8 \quad \text{! time-average velocity and vector potential}
\]

in ‘\texttt{run.in}’.

Permanent snapshots of time averages are written every \( \text{tavg} \) time units to the files ‘\texttt{data/proc*/TAV\_N}’. The current time averages are saved periodically in ‘\texttt{data/proc*/timeavg.dat}’ whenever ‘\texttt{data/proc*/var.dat}’ is written. The file format for time averages is equivalent to that of the snapshots; see §5.6.1 above.

5.9 Helper scripts

The ‘\texttt{bin}’ directory contains a collection of utility scripts, some of which are discussed elsewhere. Here is a list of the more important ones.

adapt-mkfile  Activate the settings in a ‘\texttt{Makefile}’ that apply to the given computer, see §5.2

auto-test  Verify that the code compiles and runs in a set of run directories and compare the results to the reference output. These tests are carried out routinely to ensure that the \texttt{svn} version of the code is in a usable state.

cleanf95  Can be use to clean up the output from the Intel x86 Fortran 95 compiler (ifc).

copy-proc-to-proc  Used for restarting in a different directory. Example \texttt{copy-proc-to-proc seed.dat \ldots/hydro256e}.
**copy-snapshots**  Copy snapshots from a processor-local directory to the global directory. To be started in the background before `run.x` is invoked. Used by `start.csh` and `run.csh` on network connected processors.

**pc_copyvar var1 var2 source dest**  Copies snapshot files from one directory (source) to another (dest). See documentation in file.

**pc_copyvar v v dir**  Copies all `var.dat` files from current directory to `var.dat` in 'dir' run directory. Used for restarting in a different directory.

**pc_copyvar N v**  Used to restart a run from a particular snapshot `VARN`. Copies a specified snapshot `VARN` to `var.dat` where N and (optionally) the target run directory are given on the command line.

**cvs-add-rundir**  Add the current run directory to the svn repository.

**cvsci_run**  Similar to cvs-add-rundir, but it also checks in the `*.in` and `src/*.local` files. It also checks in the files `data/time_series.dat`, `data/dim.dat` and `data/index.pro` for subsequent processing in IDL on another machine. This is particularly useful if collaborators want to check each others’ runs.

**dx.***  These script perform several data collection or reformatting exercises required to read particular files into DX. They are called internally by some of the DX macros in the ‘dx/macros/’ directory.

**getconf.csh**  See § [4.3.4]

**gpgrowth**  Plot simple time evolution with Gnuplot’s ASCII graphics for fast orientation via a slow modem line.

**local**  Materialize a symbolic link

**mkcparam**  Based on ‘Makefile’ and ‘Makefile.local’, generate ‘src/cparam.inc’, which specifies the number `mvar` of fundamental variables, and `maux` of auxiliary variables. Called by the ‘Makefile’.

**pc_mkdatadir**  Creates a link to a data directory in a suitable workspace. By default this is on ‘/var/tmp/’, but different locations are specified for different machines.

**mkdotin**  Generate minimal ‘start.in’, ‘run.in’ files based on ‘Makefile’ and ‘Makefile.local’.

**mkinpars**  Wrapper around ‘mkdotin’ — needs proper documentation.

**mkproc-tree**  Generates a multi-processor(`procN/`), directory structure. Useful when copying data files in a processor tree, such as slice files.

**mkwww**  Generates a template HTML file for describing a run of the code, showing input parameters and results.

**move-slice**  Moves all the slice files from a processor tree structure, ‘procN/’, to a new target tree creating directories where necessary.

**nl2idl**  Transform a Fortran namelist (normally the files ‘param.nml’, ‘param2.nml’ written by the code) into an IDL structure. Generates an IDL file that can be sourced from ‘start.pro’ or ‘run.pro’.

**pacx-adapt-makefile**  Version of adapt-makefile for highly distributed runs using PACX MPI.
5.10 RELOAD and STOP files

The code periodically (every \(it\) time steps) checks for the existence of two files, ‘RELOAD’ and ‘STOP’, which can be used to trigger certain behavior.

**pc_newrun** Generates a new run directory from an old one. The new one contains a copy of the old ‘*.local’ files, runs pc_setups src, and makes also a copy of the old ‘*.in’ and ‘k.dat’ files.

**pc_newscan** Generates a new scan directory from an old one. The new one contains a copy of the old, e.g. the one given under ‘samples/parameter_scan’. Look in the ‘README’ file for details.

**pc.inspectrun** Check the execution of the current run: prints legend and the last few lines of the ‘time_series.dat’ file. It also appends this result to a file called ‘SPEED’, which contains also the current wall clock time, so you can work out the speed of the code (without being affected by i/o time).

**read_video files.csh** The script for running read_video files.x.

**remote-top** Create a file ‘top.log’ in the relevant ‘procN/’ directory containing the output of top for the appropriate processor. Used in batch scripts for multi-processor runs.

**run.csh** The script for producing restart files with the initial condition; see §4.3.4

**scpdatadir** Make a tarball of data directory, ‘data/’ and use scp to secure copy to copy it to the specified destination.

**pc_setups src** Link ‘start.csh’, ‘run.csh’ and ‘getconf.csh’ from ‘$PENCIL_HOME/bin’. Generate ‘src/’ if necessary and link the source code files from ‘$PENCIL_HOME/src’ to that directory.

**start.csh** The script for initializing the code; see §4.3.4

**summarize-history** Evaluate ‘params.log’ and print a history of changes.

**timestr** Generate a unique time string that can be appended to file names from shell scripts through the backtick mechanism.

**pc_tsnap** Extract time information from a snapshot file, ‘VAR.N’.

There are several additional scripts on ‘pencil-code/utils’. Some are located in separate folders according to users. There could be redundancies, but it is often just as easy to write your own new script than figuring out how something else works.
Reloading run parameters  In the directory where you started the code, create the file ‘RELOAD’ with

```bash
unix> touch RELOAD
```

to force the code to re-read the runtime parameters from ‘run.in’. This will happen the next time the code is writing monitoring output (the frequency of this happening is controlled by the input parameter $it$, see Sect. 5.12).

Each time the parameters are reloaded, the new set of parameters is appended (in the form of namelists) to the file ‘data/params.log’ together with the time $t$, so you have a full record of your changes. If ‘RELOAD’ contains any text, its first line will be written to ‘data/params.log’ as well, which allows you to annotate changes:

```bash
unix> echo "Reduced eta to get fields growing" > RELOAD
```

Use the command `summarize-history` to print a history of changes.

Stopping the code  In the directory where you started the code, create the file ‘STOP’ with

```bash
unix> touch STOP
```

to stop the code in a controlled manner (it will write the latest snapshot). Again, the action will happen the next time the code is writing monitoring output.

5.11 RERUN and NEWDIR files

After the code finishes (e.g., when the final timestep number is reached or when a ‘STOP’ file is found), the ‘run.csh’ script checks whether there is a ‘RERUN’ file. If so, the code will simply run again, perhaps even after you have recompiled the code. This is useful in the development phase when you changed something in the code, so you don’t need to wait for a new slot in the queue!

Even more naughty, as Tony says, is the ‘NEWDIR’ file, where you can enter a new directory path (relative path is ok, e.g. `../conv-slab`). If nothing is written in this file (e.g. via `touch NEWDIR`) it stays in the same directory. On distributed machines, the ‘NEWDIR’ method will copy all the ‘VAR#’ and ‘var.dat’ files back to and from the server. This can be useful if you want to run with new data files, but you better do it in a separate directory, because with ‘NEWDIR’ the latest data from the code are written back to the server before running again.

Oh, by the way, if you want to be sure that you haven’t messed up the content of the pair of ‘NEWDIR’ files, you may want to try out the `pc_jobtransfer` command. It writes the decisive ‘STOP’ file only after the script has checked that the content of the two ‘NEWDIR’ files points to existing run directory paths, so if the new run crashes, the code returns safely to the old run directory. I’m not sure what Tony would say now, but this is now obviously extremely naughty.

5.12 Start and run parameters

All input parameters in ‘start.in’ and ‘run.in’ are grouped in Fortran namelists. This allows arbitrary order of the parameters (within the given namelist; the namelists need
5.12 Start and run parameters

no longer be in the correct order), as well as enhanced readability through inserted Fortran comments and whitespace. One namelist (init.pars / run.pars) contains general parameters for initialization/running and is always read in. All other namelists are specific to individual modules and will only be read if the corresponding module is used.

The syntax of a namelist (in an input file like ‘start.in’) is

```fortran
&init_pars
  ip=5, Lxyz=2,4,2
/
```

— in this example, the name of the namelist is init.pars, and we read just two variables (all other variables in the namelist retain their previous value): ip, which is set to 5, and Lxyz, which is a vector of length three and is set to (2, 4, 2).

While all parameters from the namelists can be set, in most cases reasonable default values are preset. Thus, the typical file ‘start.in’ will only contain a minimum set of variables or (if you are very minimalistic) none at all. If you want to run a particular problem, it is best to start by modifying an existing example that is close to your application.

Before starting a simulation run, you may want to execute the command `pc_configtest` in order to test the correctness of your changes to these configuration files.

As an example, we give here the start parameters for ‘samples/helical-MHDturb’

```fortran
&init_pars
  cvsid='$Id:$', ! identify version of start.in
  xyz0 = -3.1416, -3.1416, -3.1416, ! first corner of box
  Lxyz = 6.2832, 6.2832, 6.2832, ! box size
  lperi = T, T, T, ! periodic in x, y, z
  random_gen='nr_f90'
/
&hydro_init_pars
/
&density_init_pars
  gamma=1.
/
&magnetic_init_pars
  initaa='gaussian-noise', amplaa=1e-4
/
```

The three entries specifying the location, size and periodicity of the box are just given for demonstration purposes here — in fact a periodic box from $-\pi$ to $-\pi$ in all three directions is the default. In this run, for reproducibility, we use a random number generator from the Numerical Recipes [33], rather than the compiler's built-in generator. The adiabatic index $\gamma$ is set explicitly to 1 (the default would have been 5/3) to achieve an isothermal equation of state. The magnetic vector potential is initialized with uncorrelated, normally distributed random noise of amplitude $10^{-4}$.

The run parameters for ‘samples/helical-MHDturb’ are

```fortran
&run_pars
  cvsid='$Id:$', ! identify version of start.in
  nt=10, it1=2, cdt=0.4, cdtv=0.80, isave=10, itorder=3
```
Here we run for $nt = 10$ timesteps, every second step, we write a line of diagnostic output; we require the time step to keep the advective Courant number $\leq 0.4$ and the diffusive Courant number $\leq 0.8$, save 'var.dat' every 20 time steps, and use the 3-step time-stepping scheme described in Appendix H.4 (the Euler scheme itorder= 1 is only useful for tests). We write permanent snapshot file ‘VAR’ every $dsnap = 50$ time units and 2d slices for animation every $dvid = 0.5$ time units. Again, we use a deterministic random number generator. Viscosity $\nu$ and magnetic diffusivity $\eta$ are set to $5 \times 10^{-3}$ (so the mesh Reynolds number based on the rms velocity is about $u_{rms,0} \delta x / \nu = 0.3 \times (2\pi / 32) / 5 \times 10^{-3} \approx 12$, which is in fact rather a bit to high). The parameters in forcing_run_pars specify fully helical forcing of a certain amplitude.

A full list of input parameters is given in Appendix K.

5.13 Physical units

Many calculations are unit-agnostic, in the sense that all results remain the same independent of the unit system in which you interpret the numbers. E.g. if you simulate a simple hydrodynamical flow in a box of length $L = 1$ and get a maximum velocity of $u_{max} = 0.5$ after $t = 3$ time units, then you may interpret this as $L = 1$ m, $u_{max} = 0.5$ m/s, $t = 3$ s, or as $L = 1$ pc, $u_{max} = 0.5$ pc/Myr, $t = 3$ Myr, depending on the physical system you have in mind. The units you are using must of course be consistent, thus in the second example above, the units for diffusivities would be pc$^2$/Myr, etc.

The units of magnetic field and temperature are determined by the values $\mu_0 = 1$ and $c_p = 1$ used internally by the code. This means that if your units for density and velocity are $[\rho]$ and $[v]$, then magnetic fields will be in

$$[B] = \sqrt{\mu_0 [\rho]} [v]^2,$$

and temperatures are in

$$[T] = \frac{[v]^2}{c_p} = \frac{\gamma - 1}{\gamma} \frac{[v]^2}{R / \mu}.$$

Note that $c_p = 1$ is only assumed if you use the module noionization.f90. If you work with ionization.f90, temperature units are specified by unit_temperature as described below.
5.14 Minimum amount of viscosity

Table 2: Units of magnetic field and temperature for some choices of $[\rho]$ and $[v]$ according to Eqs. (18) and (19). Values are for a monatomic gas ($\gamma = 5/3$) of mean atomic weight $\bar{\mu}_g = \bar{\mu}/1\text{ g}$ in grams.

<table>
<thead>
<tr>
<th>$[\rho]$</th>
<th>$[v]$</th>
<th>$[B]$</th>
<th>$[T]$</th>
</tr>
</thead>
<tbody>
<tr>
<td>1 kg/m$^3$</td>
<td>1 m/s</td>
<td>1.12 mT = 11.2 G</td>
<td>$(\bar{\mu}_g/0.6) \times 2.89 \times 10^{-5}$ K</td>
</tr>
<tr>
<td>1 g/cm$^3$</td>
<td>1 cm/s</td>
<td>$3.54 \times 10^{-4}$ T = 3.54 G</td>
<td>$(\bar{\mu}_g/0.6) \times 2.89$ nK</td>
</tr>
<tr>
<td>1 g/cm$^3$</td>
<td>1 km/s</td>
<td>$35.4$ T = 354 kG</td>
<td>$(\bar{\mu}_g/0.6) \times 28.9$ K</td>
</tr>
<tr>
<td>1 g/cm$^3$</td>
<td>10 km/s</td>
<td>$354$ T = 3.54 MG</td>
<td>$(\bar{\mu}_g/0.6) \times 2890$ K</td>
</tr>
</tbody>
</table>

For some choices of density and velocity units, Table 2 shows the resulting units of magnetic field and temperature.

On the other hand, as soon as material equations are used (e.g. one of the popular parameterizations for radiative losses, Kramers opacity, Spitzer conductivities or ionization, which implies well-defined ionization energies), the corresponding routines in the code need to know the units you are using. This information is specified in ‘start.in’ or ‘run.in’ through the parameters unit_system, unit_length, unit_velocity, unit_density and unit_temperature like e.g.

```plaintext
unit_system='SI',
unit_length=3.09e16, unit_velocity=978. ! [l]=1pc, [v]=1pc/Myr
```

Note: The default unit system is unit_system='cgs' which is a synonym for unit_system='Babylonian cubits'.

5.14 Minimum amount of viscosity

We emphasize that, by default, the code works with constant diffusion coefficients (viscosity $\nu$, thermal diffusivity $\chi$, magnetic diffusivity $\eta$, or passive scalar diffusivity $D$). If any of these numbers is too small, you would need to have more meshpoints to get consistent numerical solutions; otherwise the code develops wiggles (‘ringing’) and will eventually crash. A useful criterion is given by the mesh Reynolds number based on the maximum velocity,

$$\text{Re}_{\text{mesh}} = \max(|u|) \max(\delta x,\delta y,\delta z)/\nu,$$

which should not exceed a certain value which can be problem-dependent. Often the largest possible value of $\text{Re}_{\text{mesh}}$ is around 5. Similarly there exist mesh Péclet and mesh magnetic Reynolds numbers that should not be too large.

Note that in some cases, ‘wiggles’ in $\ln \rho$ will develop despite sufficiently large diffusion coefficients, essentially because the continuity equation contains no dissipative term. For convection runs (but not only for these), we have found that this can often be prevented by upwinding, see Sect. H.2.

If the Mach number of the code approaches unity, i.e. if the rms velocity becomes comparable with the speed of sound, shocks may form. In such a case the mesh Reynolds

---

Note: The parameter unit_temperature is currently only used in connection with ionization.f90. If you are working with noionization.f90, the temperature unit is completely determined by Eq. (19) above.
number should be smaller. In order to avoid excessive viscosity in the unshocked regions, one can use the so-called shock viscosity (Sect. 6.6.1) to concentrate the effects of a low mesh Reynolds number to only those areas where it is necessary.

5.15 The time step

5.15.1 The usual RK-2N time step

RK-2N refers to the third order Runge-Kutta scheme by Williamson (1980) with a memory consumption of two chunks. Therefore the 2N in the name.

The time step is normally specified as Courant time step through the coefficients $c dt$ ($c_{dt}$), $c dt v$ ($c_{dt,v}$) and $c dt s$ ($c_{dt,s}$). The resulting Courant step is given by

$$
\delta t = \min\left( \frac{\delta x_{\min}}{U_{\max}}, \frac{\delta x_{\min}^2}{D_{\max}}, \frac{1}{H_{\max}} \right),
$$

where

$$
\delta x_{\min} \equiv \min(\delta x, \delta y, \delta z); \quad U_{\max} \equiv \max(|u| + \sqrt{c_s^2 + v_A^2}),
$$

$c_s$ and $v_A$ denoting sound speed and Alfvén speed, respectively;

$$
D_{\max} = \max(\nu, \gamma \chi, \eta, D),
$$

where $\nu$ denotes kinematic viscosity, $\chi = K/(c_p \rho)$ thermal diffusivity and $\eta$ the magnetic diffusivity; and

$$
H_{\max} = \max\left( \frac{2\nu S^2 + \zeta_{\text{shock}}(\nabla \cdot u)^2 + \ldots}{c_v T} \right),
$$

where dots indicate the presence of other terms on the rhs of the entropy equation.

To fix the time step $\delta t$ to a value independent of velocities and diffusivities, explicitly set the run parameter $dt$, rather than $c dt$ or $c dt v$ (see p. 186).

If the time step exceeds the viscous time step the simulation may actually run ok for quite some time. Inspection of images usually helps to recognize the problem. An example is shown in Fig. 5.

Timestepping is accomplished using the Runge-Kutta 2N scheme. Regarding details of this scheme see Sect. H.4.

5.15.2 The Runge-Kutta-Fehlberg time step

A fifth order Runge-Kutta-Fehlberg time stepping procedure is available. It is used mostly for chemistry application, often together with the double precision option. In order to make this work, you need to compile with

```
TIMESTEP = timestep_rkf
```

in ‘src/Makefile.local’. In addition, you must put $itorder=5$ in ‘run.in’. An example application is ‘samples/1d-tests/H2_flamespeed’. This procedure is still experimental.
5.16 Boundary conditions

5.16.1 Where to specify boundary conditions

In most tests that come with the PENCIL CODE, boundary conditions are set in ‘run.in’, which is a natural choice. However, this may lead to unexpected initial data written by ‘start.x’, since when you start the code (via ‘start.csh’), the boundary conditions are unknown and ‘start.x’ will then fill the ghost zones assuming periodicity (the default boundary condition) in all three directions. These ghost data will never be used in a calculation, as ‘run.x’ will apply the boundary conditions before using any ghost-zone values.

To avoid these periodic conditions in the initial snapshot, you can set the boundary conditions in ‘start.in’ already. In this case, they will be inherited by ‘run.x’, unless you also explicitly set boundary conditions in ‘run.in’.

5.16.2 How to specify boundary conditions

Boundary conditions are implemented through three layers of ghost points on either boundary, which is quite a natural choice for an MPI code that uses ghost zones for representing values located on the neighboring processors anyway. The desired type of boundary condition is set through the parameters bc\{x,y,z\} in ‘run.in’; the nomenclature used is as follows. Set bc\{x,y,z\} to a sequence of letters like

\[
\text{bcx} = \text{'}p\text{','p','p','p','p'}
\]

for periodic boundaries, or

\[
\text{bcz} = \text{'}s\text{','s','a','a2','c1:c2'}
\]

for non-periodic ones. Each element corresponds to one of the variables, which are those of the variables \( u_x, u_y, u_z, \ln \rho, s/c_p, A_x, A_y, A_z, \ln \rho \) that are actually used in this order. The following conditions are available:

‘p’ periodic boundary condition

‘a’ antisymmetric condition w.r.t. the boundary, i.e. vanishing value
‘s’ symmetric condition w. r. t. the boundary, i. e. vanishing first derivative

‘a2’ antisymmetry w. r. t. the arbitrary value on the boundary, i. e. vanishing second derivative

‘c1’ special boundary condition for \( \ln \rho \) and \( s \): constant heat flux through the boundary

‘c2’ special boundary condition for \( s \): constant temperature at the boundary — requires boundary condition a2 for \( \ln \rho \)

‘cT’ special boundary condition for \( s \) or \( \ln T \): constant temperature at the boundary (for arbitrarily set \( \ln \rho \))

‘ce’ special boundary condition for \( s \): set temperature in ghost points to value on boundary (for arbitrarily set \( \ln \rho \))

‘db’ low-order one-sided derivatives (“no boundary condition”) for density

‘she’ shearing-sheet boundary condition (default when the module Shear is used)

‘g’ force the value of the corresponding field on vertical boundaries (should be used in combination with the force_lower_bound and force_upper_bound flags set in the namelist init.pars)

‘hs’ special boundary condition for \( \ln \rho \) and \( s \) which enforces hydrostatic equilibrium on vertical boundaries

The extended syntax \( a:b \) (e. g. ‘c1:c2’) means: use boundary condition \( a \) at the left/lower boundary, but \( b \) at the right/upper one.

If you build a new ‘run.in’ file from another one with a different number of variables (noentropy vs. entropy, for example), you need to remember to adjust the length of the arrays bcx to bcz. The advantage of the present approach is that it is very easy to exchange all boundary conditions by a new set of conditions in a particular direction (for example, make everything periodic, or switch off shearing sheet boundary conditions and have stress-free instead).

5.17 Restarting a simulation

When a run stops at the end of a simulation, you can just resubmit the job again, and it will start from the latest snapshot saved in ‘data/proc*/var.dat’. The value of the latest time is saved in a separate file, ‘data/proc*/time.dat’. On parallel machines it is possible that some (or just one) of the ‘var.dat’ are corrupt; for example after a system crash. Check for file size and date, and restart from a good ‘VAR’ file instead.

If you want to run on a different machine, you just need to copy the ‘data/proc*/var.dat’ (and, just to be sure) ‘data/proc*/time.dat’) files into a new directory tree. You may also need the ‘data/proc*/seed.dat’ files for the random number generator. The easiest way to get all these other files is to run start.csh again on the new machine (or in a new directory) and then to overwrite the ‘data/proc*/var.dat’ files with the correct ones.

For restarting from runs that didn’t have magnetic fields, passive scalar fields, or test fields, see Sect. E.4.
5.18 One- and two-dimensional runs

If you want to run two-dimensional problems, set the number of mesh points in one direction to unity, e.g. \( nygrid = 1 \) or \( nzgrid = 1 \) in `cparam.local`. Remember that the number of mesh points is still divisible by the number of processors. For 2D-runs, it is also possible to write only 2D-snapshots (i.e. VAR files written only in the considered \((x, y)\) or \((x, z)\) plane, with a size seven times smaller as we do not write the third unused direction). To do that, please add the logical flag `lwrite_2d=T` in the namelist `init.pars` in `start.in`.

Similarly, for one-dimensional problems, set, for example, \( nygrid = 1 \) and \( nzgrid = 1 \) in `cparam.local`. You can even do a zero-dimensional run, but then you better set \( dt \) (rather than \( cdt \)), because there is no Courant condition for the time step.

See 0d, 1d, 2d, and 3d tests with examples.

5.19 Visualization

5.19.1 Gnuplot

Simple visualization can easily be done using Gnuplot (http://www.gnuplot.info), an open-source plotting program suitable for two-dimensional plots.

For example, suppose you have the variables

\[ \text{---it-----t-------dt-------urms-----umax-----rhom-----ssm------dtc---} \]

in `time_series.dat` and want to plot \( u_{\text{rms}}(t) \). Just start gnuplot and type

```bash
gnuplot> plot "data/time_series.dat" using 2:4 with lines
```

If you work over a slow line and want to see both \( u_{\text{rms}}(t) \) and \( u_{\text{max}}(t) \), use ASCII graphics:

```bash
gnuplot> set term dump
gnuplot> set logscale y
gnuplot> plot "data/time_series.dat" using 2:4 title "urms",
\n        "data/time_series.dat" using 2:5 title "umax"
```

5.19.2 Data explorer

\( DX \) (data explorer; http://www.opendx.org) is an open-source tool for visualization of three-dimensional data.

The PENCIL CODE provides a few networks for \( DX \). It is quite easy to read in a snapshot file from \( DX \) (the only tricky thing is the four extra bytes at the beginning of the file, representing a Fortran record marker), and whenever you run `start.x`, the code writes a file `var.general` that tells \( DX \) all it needs to know about the data structure.

As a starting point for developing your own \( DX \) programs or networks, you can use a few generic \( DX \) scripts provided in the directory `dx/basic/`. From the run directory, start \( DX \) with

```bash
unix> dx -edit $PENCIL_HOME/dx/basic/lnrho
```
to load the file `dx/basic/lrno.net`, and execute it with $\text{[Ctrl-o]}$ or $\text{Execute} \rightarrow \text{Execute Once}$. You will see a set of iso-surfaces of logarithmic density. If the viewport does not fit to your data, you can reset it with $\text{[Ctrl-f]}$. To rotate the object, drag the mouse over the Image window with the left or right mouse button pressed. Similar networks are provided for entropy (`ss.net`), velocity (`uu.net`) and magnetic field (`bb.net`).

When you expand these simple networks to much more elaborate ones, it is probably a good idea to separate the different tasks (like Importing and Selecting, visualizing velocity, visualizing entropy, and Rendering) onto separate pages through $\text{Edit} \rightarrow \text{Page}$.  

**Note** Currently, DX can only read in data files written by one single processor, so from a multi-processor run, you can only visualize one subregion at a time.

5.19.3 GDL

GDL, also known as *Gnu Data Language* is a free visualization package that can be found at [http://gnudatalanguage.sourceforge.net/](http://gnudatalanguage.sourceforge.net/). It aims at replacing the very expensive IDL package (see S. 5.19.4). For the way we use IDL for the Pencil Code, compatibility is currently not completely sufficient, but you can use GDL for many of the visualization tasks. If you get spurious “Error opening file” messages, you can normally simply ignore them.

This section tells you how to get started with using GDL for visualization.

**Setup** As of GDL 0.9 – at least the version packed with Ubuntu Jaunty (9.10) – you will need to add GDL’s `examples/pro/` directory to your \!PATH variable. So the first call after starting GDL should be

```
GDL> .run setup_gdl
```

**Starting visualization** There are mainly two possibilities for visualization: using a simple GUI or loading the data with `pc_read` and work with it interactively. Please note that the GUI was written and tested only with IDL, see § 5.19.4.

Here, the `pc_read` family of IDL routines to read the data is described. Try

```
GDL> pc_read
```
to get an overview.

To plot a time series, use

```
GDL> pc_read_ts, OBJECT=ts
GDL> help, ts, /STRUCT ;; (to see which slots are available)
GDL> plot, ts.t, ts.umax
GDL> oplot, ts.t, ts.urms
```

Alternatively, you could simply use the `ts.pro` script:

```
GDL> .run ts
```

To work with data from `var.dat` and similar snapshot files, you can e.g. use the following routines:
Having thus read the data structures, we can have a look at them to see what information is available:

```
GDL> help, dim, /STRUCT
GDL> help, par, /STRUCT
GDL> help, grid, /STRUCT
GDL> help, var, /STRUCT
```

To visualize data, we can e.g. do

```
GDL> plot, grid.x, var.ss[*', dim.ny/2, dim.nz/2]
GDL> contourfill, var.ss[*', *, dim.nz/2], grid.x, grid.y
```

```
GDL> ux_slice = var.uu[*', *, dim.nz/2, 0]
GDL> uy_slice = var.uu[*', *, dim.nz/2, 1]
GDL> wdvelovect, ux_slice, uy_slice, grid.x, grid.y
```

```
GDL> surface, var.lnrho[*', *, dim.nz/2, 0]
```

See also Sect. 5.19.4.

### 5.19.4 IDL

IDL is a commercial visualization program for two-dimensional and simple three-dimensional graphics. It allows to access and manipulate numerical data in a fashion quite similar to how Fortran handles them.

In `$PENCIL_HOME/idl` , we provide a number of general-purpose IDL scripts that we are using all the time in connection with the PENCIL CODE. While IDL is quite an expensive software package, it is quite useful for visualizing results from numerical simulations. In fact, for many applications, the 7-minute demo version of IDL is sufficient.

**Visualization in IDL** The Pencil Code GUI is a data post-processing tool for the usage on a day-to-day basis. It allows fast inspection of many physical quantities, as well as advanced features like horizontal averages, streamline tracing, freely orientable 2D-slices, and extraction of cut images and movies. To use the Pencil Code GUI, it is sufficient to run:

```
_unix> idl
IDL> .r pc_gui
```

If you like to load only some subvolume of the data, like any 2D-slices from the given data snapshots, or 3D-subvolumes, it is possible to choose the corresponding options in the file selector dialog. The Pencil Code GUI offers also some options to be set on the command-line, please refer to their description in the source code.

---

9 If `contourfill` produces just contour lines instead of a color-coded plot, your version of GDL is too old. E.g. the version shipped with Ubuntu 9.10 is based on GDL 0.9rc1 and has this problem.
There are also other small GUIs available, e.g. the file ‘time-series.dat’ can easily be analyzed with the command:

```
unix> idl
IDL> pc_show_ts
```

The easiest way to derive physical quantities at the command-line is to use one of the many pc_read_var-variants (pc_read_var_raw is recommended for large datasets because of its high efficiency regarding computation and memory usage) for reading the data. With that, one can make use of pc_get_quantity to derive any implemented physical quantity. Packed in a script, this is the recommended way to get reproducible results, without any chance for accidental errors on the interactive IDL command-line.

Alternatively, by using the command-line to see the time evolution of e.g. velocity and magnetic field (if they are present in you run), start IDL and run ‘ts.pro’:

```
unix> idl
IDL> .run ts
```

The IDL script ‘ts.pro’ reads the time series data from ‘data/time_series.dat’ and sorts the column into the structure ts, with the slot names corresponding to the name of the variables (taken from the header line of ‘data/time_series.dat’). Thus, you can refer to time as ts.t, to the rms velocity as ts.urms, and in order to plot the mean density as a function of time, you would simply type

```
IDL> plot, ts.t, ts.rhom
```

The basic command sequence for working with a snapshot is:

```
unix> idl
IDL> .run start
IDL> .run r
IDL> [specific commands]
```

You run ‘start.pro’ once to initialize (or reinitialize, if the mesh size has changed, for example) the fields and read in the startup parameters from the code. To read in a new snapshot, run ‘r.pro’ (or ‘rall.pro’, see below).

If you are running in parallel on several processors, the data are scattered over different directories. To reassemble everything in IDL, use

---

10 If you run IDL from the command line, you will highly appreciate the following tip: IDL’s command line editing is broken beyond hope. But you can fix it, by running IDL under rlwrap, a wrapper for the excellent GNU readline library.

Download and install rlwrap from http://utopia.knoware.nl/~hlub/uck/rlwrap/ (on some systems you just need to run ‘emerge rlwrap’, or ‘apt-get install rlwrap’), and alias your idl command:

```
csh> alias idl 'rlwrap -a -c idl'
bash> alias idl='rlwrap -a -c idl'
```

From now on, you can

- use long command lines that correctly wrap around;
- type the first letters of a command and then [PageUp] to recall commands starting with these letters;
- capitalize, uppercase or lowercase a word with [Esc]C, [Esc]U, [Esc]L;
- use command line history across IDL sessions (you might need to create ‘~/.idl_history’ for this);
- complete file names with Tab (works to some extent);
- ...use all the other readline features that you are using in bash, octave, bc, gnuplot, ftp, etc.
instead of \texttt{.r r} (here, \texttt{.r} is a shorthand for \texttt{.run}). The procedure ‘\texttt{rall.pro}’ reads (and assembles) the data from all processors and correspondingly requires large amounts of memory for very large runs. If you want to look at just the data from one processor, use ‘\texttt{r.pro}’ instead.

If you need the magnetic field or the current density, you can calculate them in IDL by \footnote{Keep in mind that \texttt{jj=curl(bb)} would use iterated first derivatives instead of the second derivatives and thus be numerically less accurate than \texttt{jj=curl2(aa)}, particularly at small scales.}

\begin{verbatim}
IDL> bb=curl(aa)
IDL> jj=curl2(aa)
\end{verbatim}

By default one is reading always the current snapshot ‘data/proc\texttt{N}/var.dat’; if you want to read one of the permanent snapshots, use (for example)

\begin{verbatim}
IDL> varfile='VAR2'
IDL> .r r (or .r rall)
\end{verbatim}

See Sect. 5.6.1 for details on permanent snapshots.

With ‘\texttt{r.pro}’, you can switch the part of the domain by changing the variable \texttt{datadir}:

\begin{verbatim}
IDL> datadir='data/proc3'
IDL> .r r
\end{verbatim}

will read the data written by processor 3.

\paragraph*{Reading data into IDL arrays or structures} As an alternative to the method described above, there is also the possibility to read the data into structures. This makes some more operations possible, e.g. reading data from an IDL program where the command \texttt{.r} is not allowed.

An efficient and still scriptable way would look like the following:

\begin{verbatim}
IDL> pc_read_var_raw, obj=var, tags=tags
IDL> bb = pc_get_quantity ('B', var, tags)
IDL> jj = pc_get_quantity ('j', var, tags)
\end{verbatim}

This reads the data into an array ‘\texttt{var}’, as well as the array indices of the contained physical variables into a separate structure ‘\texttt{tags}’. To use a caching mechanism within \texttt{pc-get_quantity}, please refer to the documentation and the examples contained in ‘pencil-code/idl/pc_get_quantity.pro’, where you can also start adding not yet implemented physical quantities.

To read a snapshot ‘\texttt{VAR10}’ into the IDL structure \texttt{ff}, type the following command

\begin{verbatim}
IDL> pc_read_var, obj=ff, varfile='VAR10', /trimall
\end{verbatim}

The option /\texttt{trimall} removes ghost zones from the data. A number of other options are documented in the source code of \texttt{pc_read_var}. You can see what data the structure contains by using the command \texttt{tag_names}

\begin{verbatim}
IDL> print, tag_names(ff)
        T X Y Z DX DY DZ UU LNRHO AA
\end{verbatim}
One can access the individual variables by typing `ff.varname`, e.g.

```
IDL> help, ff.aa
<Expression> FLOAT = Array[32, 32, 32, 3]
```

There are a number of files that read different data into structures. They are placed in the directory `\$PENCIL_HOME/idl/files`. Here is a list of files (including suggested options to call them with):

- `pc_read_var_raw`, `obj=var`, `tags=tags`
  Efficiently read a snapshot into an array.
- `pc_read_var`, `obj=ff`, `/trimall`
  Read a snapshot into a structure.
- `pc_read_ts`, `obj=ts`
  Read the time series into a structure.
- `pc_read_xyaver`, `obj=xya`
  `pc_read_xzaver`, `obj=xza`
  `pc_read_yzaver`, `obj=yza`
  Read 1-D time series into a structure.
- `pc_read_const`, `obj=cst`
  Read code constants into a structure.
- `pc_read_pvar`, `obj=fp`
  Read particle data into a structure.
- `pc_read_param`, `obj=par`
  Read startup parameters into a structure.
- `pc_read_param`, `obj=par2`, `/param2`
  Read runtime parameters into a structure.

Other options are documented in the source code of the files.

For some examples on how to use these routines, see Sect. [5.19.3](#).

### 5.19.5 Python

Pencil supports reading, processing and the visualization of data using *python*. A number of scripts are placed in the subfolder `\$PENCIL_HOME/python`. A README file placed under that subfolder contains the information needed to read Pencil output data into python.

**Installation** For modern operating systems, Python is generally installed together with the system. If not, it can be installed via your preferred package manager or downloaded from the website [https://www.python.org/](https://www.python.org/). For convenience, it is strongly recommend to also install IPython, which is a more convenient console for Python. You will also need the NumPy, matplotlib, h5py and Tk library.

Perhaps the easiest way to obtain all the required software mentioned above is to install either Continuum’s Anaconda or Enthought’s Canopy. These Python distributions also provide (or indeed are) integrated graphical development environments.
Another way of installing libraries, particularly on a cluster without root privileges you can use pip or pip3: pip install h5py or pip3 install h5py.

In order for Python to find the Pencil Code commands you will have to add to your .bashrc:

```bash
echo 'export PYTHONPATH=$PENCIL_HOME/python' >> .bashrc
```

**ipythonrc** If you use IPython, for convenience, you should modify your .ipython/ipythonrc (create it if it doesn't exist) and add:

```python
import_all pencil
```

Additional, add to your .ipython/profile_default/startup/init.py the following lines:

```python
import numpy as np
import pylab as plt
import pencil as pc
import matplotlib
from matplotlib import rc

plt.ion()
matplotlib.rcParams['savefig.directory'] = ''
```

**.pythonrc** In case you are on a cluster and don't have access to IPython you can edit your .pythonrc:

```bash
#!/usr/bin/python
import numpy as np
import pylab as plt
import pencil as pc
import atexit
# import readline
import rlcompleter
# enables search with CTR+r in the history
try:
    import readline
except ImportError:
    print "Module readline not available."
else:
    import rlcompleter
    readline.parse_and_bind("tab: complete")
# enables command history
historyPath = os.path.expanduser("~/.pyhistory")
def save_history(historyPath=historyPath):
    import readline
    readline.write_history_file(historyPath)
if os.path.exists(historyPath):
    readline.read_history_file(historyPath)
atexit.register(save_history)
del os, atexit, readline, rlcompleter, save_history, historyPath
```
plt.ion()
create the file .pythonhistory and add to your .bashrc:
export PYTHONSTARTUP=~/.pythonrc

**Pencil Code Commands in General** For a list of all Pencil Code commands start IPython and type pc. <TAB> (as with auto completion). To access the help of any command just type the command followed by a '?' (no spaces), e.g.:

```
In [1]: pc.math.dot?
Signature: pc.math.dot(a, b)
Docstring:
Take dot product of two pencil-code vectors a and b.

call signature:

dot(a, b):

Keyword arguments:

*a*, *b*:
Pencil-code vectors with shape [3, mz, my, mx].
```

File: ~/pencil-code/python/pencil/math/vector_multiplication.py
Type: function

You can also use help(pc.dot) for a more complete documentation of the command.

There are various reading routines for the Pencil Code data. All of them return an object with the data. To store the data into a user defined variable type e.g.

```
In [1]: ts = pc.read.ts()
```

Most commands take some arguments. For most of them there is a default value, e.g.

```
In [1]: pc.read.ts(file_name='time_series.dat', datadir='data').
```

**Reading and Plotting Time Series** Reading the time series file is very easy. Simply type

```
In [1]: ts = pc.read.ts()
```

and Python stores the data in the variable ts. The physical quantities are members of the object ts and can be accessed accordingly, e.g. ts.t, ts.emag. To check which other variables are stored simply do the tab auto completion ts. <TAB>.

Plot the data with the matplotlib commands:

```
In [1]: plt.plot(ts.t, ts.emag)
```

The standard plots are not perfect and need a little polishing. See the online wiki for a few examples on how to make pretty plots ([https://github.com/pencil-code/pencil-code/wiki/PythonForPencil](https://github.com/pencil-code/pencil-code/wiki/PythonForPencil)). You can save the plot into a file using the GUI or with
**5.20 Running on multi-processor computers**

The code is parallelized using *MPI* (*message passing interface*) for a simple domain decomposition (data-parallelism), which is a straightforward and very efficient way of parallelizing finite-difference codes. The current version has a few restrictions, which need to be kept in mind when using the MPI features.

The global number of grid points (but excluding the ghost zones) in a given direction must be an exact multiple of the number of processors you use in that direction. For example, if you have *nprocy*=8 processors for the *y* direction, you can run a job with *nygrid*=64 points in that direction, but if you try to run a problem with *nygrid*=65 or *nygrid*=94, the code will complain about an inconsistency and stop. (So far, this has not been a serious restriction for us.)

**5.20.1 How to run a sample problem in parallel**

To run the sample problem in the directory `samples/conv-slab` on 16 CPUs, you need to do the following (in that directory):

1. Edit `src/Makefile.local` and replace
   
   ```
   MPICOMM    = nompicomm
   
   by
   
   MPICOMM    = mpicomm
   ```

2. Edit `src/cparam.local` and replace
integer, parameter :: ncpus=1, nprocy=1, nprocz=ncpus/nprocy, nprocx=1
integer, parameter :: nxgrid=32, nygrid=nxgrid, nzgrid=nxgrid

by

integer, parameter :: ncpus=16, nprocy=4, nprocz=ncpus/nprocy, nprocx=1
integer, parameter :: nxgrid=128, nygrid=nxgrid, nzgrid=nxgrid

The first line specifies a $4 \times 4$ layout of the data in the $y$ and $z$ direction. The second line increases the resolution of the run because running a problem as small as $32^3$ on 16 CPUs would be wasteful. Even $128^3$ may still be quite small in that respect. For performance timings, one should try and keep the size of the problem per CPU the same, so for example $256^3$ on 16 CPUs should be compared with $128^3$ on 2 CPUs.

3. Recompile the code

unix> (cd src; make)

4. Run it

unix> start.csh
unix> run.csh

Make sure that all CPUs see the same ‘data/’ directory; otherwise things will go wrong. Remember that in order to visualize the full domain with IDL (rather than just the domain processed and written by one processor), you need to use ‘rall.pro’ instead of ‘r.pro’.

5.20.2 Hierarchical networks (e.g. on Beowulf clusters)

On big Beowulf clusters, a group of nodes is often connected with a switch of modest speed, and all these groups are connected via a $n$ times faster uplink switch. When bandwidth-limited, it is important to make sure that consecutive processors are mapped onto the mesh such that the load on the uplink is $\lesssim n$ times larger than the load on the slower switch within each group. On a 512 node cluster, where groups of 24 processors are linked via fast ethernet switches, which in turn are connected via a Gigabit uplink ($\sim 10$ times faster), we found that $nprocy=4$ is optimal. For 128 processors, for example we find that $nprocy \times nprocz = 4 \times 32$ is the optimal layout, while. For comparison, $8 \times 16$ is 3 times slower, and $16 \times 8$ is 17 (!) times slower. These results can be understood from the structure of the network, but the basic message is to watch out for such effects and to try varying $nprocy$ and $nprocz$.

In cases where $nygrid>nzgrid$ it may be advantageous to swap the ordering of processor numbers. This can be done by setting $lprocz\_slowest=F$.

5.20.3 Extra workload caused by the ghost zones

Normally, the workload caused by the ghost zones is negligible. However, if one increases the number of processors, a significant fraction of work is done in the ghost zones. In other words, the effective mesh size becomes much larger than the actual mesh size.
Consider a mesh of size $N_w = N_x \times N_y \times N_z$, and distribute the task over $P_w = P_x \times P_y \times P_z$ processors. If no communication were required, the number of points per processor would be

$$\frac{N_w}{P_w} = \frac{N_x \times N_y \times N_z}{P_x \times P_y \times P_z}. \quad (26)$$

However, for finite difference codes some communication is required, and the amount of communication depends on spatial order of the scheme, $Q$. The PENCIL CODE works by default with sixth order finite differences, so $Q = 6$, i.e. one needs 6 ghost zones, 3 on each end of the mesh. With $Q \neq 0$ the number of points per processor is

$$\frac{N_{w}^{(\text{eff})}}{P_w} = \left( \frac{N_x}{P_x} + Q \right) \times \left( \frac{N_y}{P_y} + Q \right) \times \left( \frac{N_z}{P_z} + Q \right). \quad (27)$$

There is efficient scaling only when

$$\min \left( \frac{N_x}{P_x}, \frac{N_y}{P_y}, \frac{N_z}{P_z} \right) \gg Q. \quad (28)$$

In the special case were $N_x = N_y = N_z \equiv N = N_w^{1/3}$, with $P_x = 1$ and $P_y = P_z \equiv P = P_w^{1/2}$, we have

$$\frac{N_{w}^{(\text{eff})}}{P_w} = (N + Q) \times \left( \frac{N}{P} + Q \right)^2. \quad (29)$$

For $N = 128$ and $Q = 6$ the effective mesh size exceeds the actual mesh size by a factor

$$\frac{N_{w}^{(\text{eff})}}{N_w} = (N + Q) \times \left( \frac{N}{P} + Q \right)^2 \times \frac{P_w}{N_w}. \quad (30)$$

These factors are listed in Table 3.

<table>
<thead>
<tr>
<th>$P$</th>
<th>$N$</th>
<th>128</th>
<th>256</th>
<th>512</th>
<th>1024</th>
<th>2048</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1.15</td>
<td>1.07</td>
<td>1.04</td>
<td>1.02</td>
<td>1.01</td>
<td></td>
</tr>
<tr>
<td>2</td>
<td>1.19</td>
<td>1.09</td>
<td>1.05</td>
<td>1.02</td>
<td>1.01</td>
<td></td>
</tr>
<tr>
<td>4</td>
<td>1.25</td>
<td>1.12</td>
<td>1.06</td>
<td>1.03</td>
<td>1.01</td>
<td></td>
</tr>
<tr>
<td>8</td>
<td>1.34</td>
<td>1.16</td>
<td>1.08</td>
<td>1.04</td>
<td>1.02</td>
<td></td>
</tr>
<tr>
<td>16</td>
<td>1.48</td>
<td>1.22</td>
<td>1.11</td>
<td>1.05</td>
<td>1.03</td>
<td></td>
</tr>
<tr>
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<td>1.68</td>
<td>1.31</td>
<td>1.15</td>
<td>1.07</td>
<td>1.04</td>
<td></td>
</tr>
<tr>
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<td>1.98</td>
<td>1.44</td>
<td>1.21</td>
<td>1.10</td>
<td>1.05</td>
<td></td>
</tr>
<tr>
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<td>2.45</td>
<td>1.64</td>
<td>1.30</td>
<td>1.14</td>
<td>1.07</td>
<td></td>
</tr>
<tr>
<td>256</td>
<td>3.21</td>
<td>1.93</td>
<td>1.43</td>
<td>1.20</td>
<td>1.10</td>
<td></td>
</tr>
<tr>
<td>512</td>
<td>4.45</td>
<td>2.40</td>
<td>1.62</td>
<td>1.29</td>
<td>1.14</td>
<td></td>
</tr>
</tbody>
</table>

Ideally, one wants to keep the work in the ghost zones at a minimum. If one accepts that 20–25% of work are done in the ghost zones, one should use 4 processors for $128^3$ mesh points, 16 processors for $256^3$ mesh points, 64 processors for $512^3$ mesh points, 256 processors for $1024^3$ mesh points, and 512 processors for $1536^3$ mesh points.
5.21 Running in double-precision

With many compilers, you can easily switch to double precision (8-byte floating point numbers) as follows.

Add the lines

```plaintext
# Use double precision
REALPRECISION = double
```

to `src/Makefile.local' and (re-)run `pc_setupsrc'.

If `REALPRECISION' is set to 'double', the flag `FFLAGS_DOUBLE' is appended to the Fortran compile flags. The default for `FFLAGS_DOUBLE' is `-r8', which works for `g95' or `ifort'; for `gfortran', you need to make sure that `FFLAGS_DOUBLE' is set to `-fdefault-real-8'.

You can see the flags in `src/Makefile.inc', which is the first place to check if you have problems compiling for double precision.

Using double precision might be important in turbulence runs where the resolution is 256³ meshpoints and above (although such runs often seem to work fine at single precision). To continue working in double precision, you just say `lread_from_other_prec=T' in `run_pars'; see Sect. 5.1.

5.22 Power spectrum

Given a real variable \( u \), its Fourier transform \( \tilde{u} \) is given by

\[
\tilde{u}(k_x, k_y, k_z) = \mathcal{F}(u(x, y, z)) = \frac{1}{N_x N_y N_z} \sum_{p=0}^{N_x-1} \sum_{q=0}^{N_y-1} \sum_{r=0}^{N_z-1} u(x_p, y_q, z_r) \times \exp(-ik_x x_p) \exp(-ik_y y_q) \exp(-ik_z z_r),
\]  

(31)

where

\[
|k_x| < \frac{\pi N_x}{L_x}, \quad |k_y| < \frac{\pi N_y}{L_y}, \quad |k_z| < \frac{\pi N_z}{L_z},
\]

when \( L \) is the size of the simulation box. The three-dimensional power spectrum \( P(k) \) is defined as

\[
P(k) = \frac{1}{2} \tilde{u} \tilde{u}^*,
\]

(32)

where

\[
k = \sqrt{k_x^2 + k_y^2 + k_z^2}.
\]

(33)

Note that we can only reasonably calculate \( P(k) \) for \( k < \pi N_x / L_x \).

To get power spectra from the code, edit `run.in' and add for example the following lines

```plaintext
dspec=5., ou_spec=T, ab_spec=T !(for energy spectra)
oned=T
```

under `run_pars'. The kinetic (vel_spec) and magnetic (mag_spec) power spectra will now be calculated every 5.0 (dspec) time units. The Fourier spectra is calculated using `fftpack'. In addition to calculating the three-dimensional power spectra also the one-dimensional power spectra will be calculated (oned).
5.23 Structure functions

In addition one must edit `src/Makefile.local` and add the lines

```plaintext
FOURIER = fourier_fftpack
POWER = power_spectrum
```

Running the code will now create the files `powerhel_mag.dat` and `power_kin.dat` containing the three-dimensional magnetic and kinetic power spectra respectively. In addition to these three-dimensional files we will also find the one-dimensional files `powerbx_x.dat`, `powerby_x.dat`, `powerbz_x.dat`, `powerux_x.dat`, `poweruy_x.dat` and `poweruz_x.dat`. In these files the data are stored such that the first line contains the time of the snapshot, the following \( \frac{nxgrid}{2} \) numbers represent the power at each wavenumber, from the smallest to the largest. If several snapshots have been saved, they are being stored immediately following the preceding snapshot.

You can read the results with the idl procure `power`, like this:

```plaintext
power,'_kin','_mag',k=k,spec1=spec1,spec2=spec2,i=n,tt=t,/noplot
power,'hel_kin','hel_mag',k=k,spec1=spec1h,spec2=spec2h,i=n,tt=t,/noplot
```

If powerhel is invoked, krms is written during the first computation. The relevant output file is `power_krms.dat`. This is needed for a correct calculation of \( k \) used in the realizability conditions.

A caveat of the implementation of Fourier transforms in the PENCIL CODE is that, due to the parallelization, the permitted resolution is limited to the case when one direction is an integer multiple of the other. So, it can be done for

\[
Nx = n*Ny
\]

Unfortunately, for some applications one wants \( Nx < Ny \). Wlad experimented with arbitrary resolution by interpolating \( x \) to the same resolution of \( y \) prior to transposing, then transform the interpolated array and then interpolating it back (check `fourier_transform_y` in `fourier_fftpack.f90`).

A feature of our current implementation with \( x \) parallelization is that `fft_xyz_parallel_3D` requires `nygrid` to be an integer multiple of `nprocy*nprocz`. Examples of good mesh layouts are listed in Table 4.

To visualize with IDL just type `power` and you get the last snapshot of the three-dimensional power spectrum. See head of `$PENCIL_HOME/idl/power.pro` for options to `power`.

### 5.23 Structure functions

We define the p-th order longitudinal structure function of \( u \) as

\[
S_{\text{long}}^p(l) = \langle |u_x(x+l,y,z) - u_x(x,y,z)|^p \rangle ,
\]

(34)

while the transverse is

\[
S_{\text{trans}}^p(l) = \langle |u_y(x+l,y,z) - u_y(x,y,z)|^p \rangle + \langle |u_z(x+l,y,z) - u_z(x,y,z)|^p \rangle .
\]

(35)

Edit `run.in` and add for example the following lines
Table 4: Examples of mesh layouts for which Fourier transforms with $x$ parallelization is possible.

<table>
<thead>
<tr>
<th>ny</th>
<th>nprocx</th>
<th>nprocy</th>
<th>nprocz</th>
<th>ncpus</th>
</tr>
</thead>
<tbody>
<tr>
<td>256</td>
<td>1</td>
<td>16</td>
<td>16</td>
<td>256</td>
</tr>
<tr>
<td>256</td>
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<td>16</td>
<td>512</td>
</tr>
<tr>
<td>256</td>
<td>4</td>
<td>16</td>
<td>16</td>
<td>1024</td>
</tr>
<tr>
<td>256</td>
<td>8</td>
<td>16</td>
<td>16</td>
<td>2048</td>
</tr>
<tr>
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<td>2</td>
<td>16</td>
<td>18</td>
<td>576</td>
</tr>
<tr>
<td>512</td>
<td>2</td>
<td>16</td>
<td>32</td>
<td>1024</td>
</tr>
<tr>
<td>512</td>
<td>4</td>
<td>16</td>
<td>16</td>
<td>1024</td>
</tr>
<tr>
<td>512</td>
<td>4</td>
<td>16</td>
<td>32</td>
<td>2048</td>
</tr>
<tr>
<td>576</td>
<td>4</td>
<td>18</td>
<td>32</td>
<td>2304</td>
</tr>
<tr>
<td>576</td>
<td>8</td>
<td>18</td>
<td>32</td>
<td>4608</td>
</tr>
<tr>
<td>576</td>
<td>16</td>
<td>18</td>
<td>32</td>
<td>9216</td>
</tr>
<tr>
<td>1024</td>
<td>4</td>
<td>32</td>
<td>32</td>
<td>4096</td>
</tr>
<tr>
<td>1024</td>
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<td>32</td>
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<tr>
<td>1152</td>
<td>4</td>
<td>36</td>
<td>32</td>
<td>4608</td>
</tr>
<tr>
<td>1152</td>
<td>4</td>
<td>32</td>
<td>36</td>
<td>4608</td>
</tr>
<tr>
<td>2304</td>
<td>2</td>
<td>32</td>
<td>72</td>
<td>4608</td>
</tr>
<tr>
<td>2304</td>
<td>4</td>
<td>36</td>
<td>64</td>
<td>9216</td>
</tr>
<tr>
<td>2304</td>
<td>4</td>
<td>32</td>
<td>72</td>
<td>9216</td>
</tr>
</tbody>
</table>

dspec=2.3,
lsfu=T, lsfb=T, lsfz1=T, lsfz2=T

under run_pars. The velocity (lsfu), magnetic (lsfb) and Elsasser (lsfz1 and lsfz2) structure functions will now be calculated every 2.3 (dspec) time unit.

In addition one must edit ‘src/Makefile.local’ and add the line

```plaintext
STRUCT_FUNC = struct_func
```

In ‘src/cparam.local’, define $lb_nxgrid$ and make sure that

```plaintext
nxgrid = nygrid = nzgrid = 2**lb_nxgrid
```

E.g.

```plaintext
integer, parameter :: lb_nxgrid=5
integer, parameter :: nxgrid=2**lb_nxgrid,nygrid=nxgrid,nzgrid=nxgrid
```

Running the code will now create the files:
’sfu-1.dat’, ‘sfu-2.dat’, ‘sfu-3.dat’ (velocity),
’sfb-1.dat’, ‘sfb-2.dat’, ‘sfb-3.dat’ (magnetic field),
’sfz1-1.dat’, ‘sfz1-2.dat’, ‘sfz1-3.dat’ (first Elsasser variable),
’sfz2-1.dat’, ‘sfz2-2.dat’, ‘sfz2-3.dat’ (second Elsasser variable),
which contains the data of interest. The first line in each file contains the time $t$ and the number $q_{max}$, such that the largest moment calculated is $q_{max} – 1$. The next $imax$
numbers represent the first moment structure function for the first snapshot, here

\[ \text{imax} = 2 \frac{\ln(n_{\text{grid}})}{\ln 2} - 2. \] (36)

The next imax numbers contain the second moment structure function, and so on until qmax – 1. The following imax numbers then contain the data of the signed third order structure function i.e. \( S_3^{\text{long}}(l) = \langle [u_x(x+l, y, z) - u_x(x, y, z)]^3 \rangle \).

The following imax \( \times qmax \times 2 \) numbers are zero if nr\_directions = 1 (default), otherwise they are the same data as above but for the structure functions calculated in the y and z directions.

If the code has been run long enough as to calculate several snapshots, these snapshots will now follow, being stored in the same way as the first snapshot.

To visualize with IDL just type structure and you get the time-average of the first order longitudinal structure function (be sure that `pencil-runs/forced/idl/` is in IDL\_PATH). See head of `pencil-runs/forced/idl/structure.pro` for options to structure.

## 5.24 Particles

The PENCIL CODE has modules for tracer particles and for dust particles (see Sect. 6.16). The particle modules are chosen by setting the value of the variable PARTICLES in Makefile.local to either particles\_dust or particles\_tracers. For the former case each particle has six degrees of freedom, three positions and three velocities. For the latter it suffices to have only three position variables as the velocity of the particles are equal to the instantaneous fluid velocity at that point. In addition one can choose to have several additional internal degrees of freedoms for the particles. For example one can temporally evolve the particles radius by setting PARTICLES\_RADIUS to particles\_radius in Makefile.local.

All particle infrastructure is controlled and organized by the Particles\_main module. This module is automatically selected by Makefile.src if PARTICLES is different from noparticles. Particle modules are compiled as a separate library. This way the main part of the Pencil Code only needs to know about the particles\_main.a library, but not of the individual particle modules.

For a simulation with particles one must in addition define a few parameters in cparam.local. Here is a sample of cparam.local for a parallel run with 2,000,000 particles:

```fortran
integer, parameter :: ncpus=16, nprocy=4, nprocz=4, nprocx=1
integer, parameter :: nxgrid=128, nygrid=256, nzgrid=128
integer, parameter :: npar=2000000, mpar_loc=400000, npar_mig=1000
integer, parameter :: npar\_species=2
```

The parameter npar is the number of particles in the simulation, mpar\_loc is the number of particles that is allowed on each processor and npar\_mig is the number of particles that are allowed to migrate from one processor to another in any time-step. For a non-parallel run it is enough to specify npar. The number of particle species is set through npar\_species (assumed to be one if not set). The particle input parameters are given in start.in and run.in. Here is a sample of the particle part of start.in for dust particles:

```plaintext
...
The initial positions and velocities of the dust particles are set in \texttt{initxxp} and \texttt{initvvp}. The next four input parameters are further specifications of the initial condition. Interaction between the particles and the mesh, e.g. through drag force or self-gravity, require a mapping of the particles on the mesh. The \textsc{Pencil Code} currently supports NGP (Nearest Grid Point, default), CIC (Cloud in Cell, set \texttt{lparticlemesh\_cic=T}) and TSC (Triangular Shaped Cloud, set \texttt{lparticlemesh\_tsc=T}). See Youdin & Johansen (2007) for details.

Here is a sample of the particle part of \texttt{run.in} (also for dust particles):

```plaintext
/*
&particles\_init\_pars
  initxxp='gaussian-z', initvvp='random'
  zp0=0.02, delta_vvp0=0.01, eps_dtog=0.01, taursp=0.1
  lparticlemesh\_tsc=T
*/

The logical \texttt{ldragforce\_gas\_par} determines whether the dust particles influence the gas with a drag force. \texttt{cdtp} tells the code how many friction times should be resolved in a minimum time-step.

The sample run `samples\_sedimentation/' contains the latest setup for dust particles.

### 5.24.1 Particles in parallel

The particle variables (e.g. $x_i$ and $v_i$) are kept in the arrays \texttt{fp} and \texttt{dfp}. For parallel runs, particles must be able to move from processor to processor as they pass out of the $(x,y,z)$-interval of the local processor. Since not all particles are present at the same processor at the same time (hopefully), there is some memory optimization in making \texttt{fp} not big enough to contain all the particles at once. This is achieved by setting the code variable \texttt{mpar\_loc} less than \texttt{npar} in \texttt{cparam.local} for parallel runs. When running with millions of particles, this trick is necessary to keep the memory need of the code down.

The communication of migrating particles between the processors happens as follows (see the subroutine \texttt{redist\_particles\_procs} in \texttt{particles\_sub.f90}):

1. In the beginning of each time-step all processors check if any of their particles have crossed the local $(x,y,z)$-interval. These particles are called migrating particles. A run can have a maximum of \texttt{npar\_mig} migrating particles in each time-step. The value of \texttt{npar\_mig} must be set in \texttt{cparam.local}. The number should (of course) be slightly larger than the maximum number of migrating particles at any time-step during the run. The diagnostic variable \texttt{nmigmax} can be used to output the maximum number of migrating particles at a given time-step. One can set \texttt{lmigration\_redo=T} in \texttt{&particles\_run\_pars} to force the code to redo the migration step if more than \texttt{npar\_mig} want to migrate. This does slow the code down somewhat, but has the benefit that the code does not stop when more than \texttt{npar\_mig} particles want to migrate.
2. The index number of the receiving processor is then calculated. This requires some assumption about the grid on other processors and will currently not work for nonequidistant grids. Particles do not always pass to neighboring processors as the global boundary conditions may send them to the other side of the global domains (periodic or shear periodic boundary conditions).

3. The migrating particle information is copied to the end of fp, and the empty spot left behind is filled up with the particle of the highest index number currently present at the processor.

4. Once the number of migrating particles is known, this information is shared with neighboring processors (including neighbors over periodic boundaries) so that they all know how many particles they have to receive and from which processors.

5. The communication happens as directed MPI communication. That means that processors 0 and 1 can share migrating particles at the same time as processors 2 and 3 do it. The communication happens from a chunk at the end of fp (migrating particles) to a chunk that is present just after the particle of the highest index number that is currently at the receiving processor. Thus the particles are put directly at their final destination, and the migrating particle information at the source processor is simply overwritten by other migrating particles at the next time-step.

6. Each processor keeps track of the number of particles that it is responsible for. This number is stored in the variable npar_loc. It must never be larger than npar_loc (see above). When a particle leaves a processor, npar_loc is reduced by one, and then increased by one at the processor that receives that particle. The maximum number of particles at any processor is stored in the diagnostic variable nparmax. If this value is not close to npar/ncpus, the particles have piled up in such a way that computations are not evenly shared between the processors. One can then try to change the parallelization architecture (nprocy and nprocz) to avoid this problem.

In simulations with many particles (comparable to or more than the number of grid cells), it is crucial that particles are shared relatively evenly among the processors. One can as a first approach attempt to not parallelize directions with strong particle density variations. However, this is often not enough, especially if particles clump locally.

Alternatively one can use Particle Block Domain Decomposition (PBDD, see Johansen et al. 2011). The steps in Particle Block Domain Decomposition scheme are as follows:

1. The fixed mesh points are domain-decomposed in the usual way (with ncpus=nprocx×nprocy×nprocz).

2. Particles on each processor are counted in bricks of size nbx×nby×nbz (typically nbx=nby=nbz=4).

3. Bricks are distributed among the processors so that each processor has approximately the same number of particles

4. Adopted bricks are referred to as blocks.

5. The Pencil Code uses a third order Runge-Kutta time-stepping scheme. In the beginning of each sub-time-step particles are counted in blocks and the block counts communicated to the bricks on the parent processors. The particle density assigned to ghost cells is folded across the grid, and the final particle density (defined on the bricks) is communicated back to the adopted blocks. This step is necessary because
the drag force time-step depends on the particle density, and each particle assigns density not just to the nearest grid point, but also to the neighboring grid points.

6. In the beginning of each sub-time-step the gas density and gas velocity field is communicated from the main grid to the adopted particle blocks.

7. Drag forces are added to particles and back to the gas grid points in the adopted blocks. This partition aims at load balancing the calculation of drag forces.

8. At the end of each sub-time-step the drag force contribution to the gas velocity field is communicated from the adopted blocks back to the main grid.

Particle Block Domain Decomposition is activated by setting `PARTICLES = particles_dust_blocks` and `PARTICLES_MAP = particles_map_blocks` in Makefile.local. A sample of cparam.local for Particle Block Domain Decomposition can be found in samples/sedimentation/blocks:

```fortran
integer, parameter :: ncpus=4, nprocx=2, nprocy=2, nprocz=1
integer, parameter :: nxgrid=32, nygrid=32, nzgrid=32
integer, parameter :: npar=10000, mpar_loc=5000, npar_mig=100
integer, parameter :: npar_species=4
integer, parameter :: nbrickx=4, nbricky=4, nbrickz=4, nblockmax=32
```

The last line defines the number of bricks in the total domain – here we divide the grid into \(4 \times 4 \times 4\) bricks each of size \(8 \times 8 \times 8\) grid points. The parameter `nblockmax` tells the code the maximum number of blocks any processor may adopt. This should not be so low that there is not room for all the bricks with particles, nor so high that the code runs out of memory.

### 5.24.2 Large number of particles

When dealing with large number of particles, one needs to make sure that the number of particles `npar` is less than the maximum integer that the compiler can handle with. The maximum integer can be checked by the Fortran intrinsic function `huge`,

```fortran
program huge_integers
  print *, huge(0_4) ! for default Fortran integer (32 Bit)
  print *, huge(0_8) ! for 64 Bit integer in Fortran
end program huge_integers
```

If the number of particles `npar` is larger than default maximum integer, one can promote the maximum integer to 64 Bit by setting

```fortran
integer(kind=8), parameter :: npar=4294967296
```

in the cparam.local file. This works because the data type of `npar` is only set here. It is worth noting that one **should not** use the flag

```
FFLAGS += -integer-size 64
```

to promote all the integers to 64 Bit. This will break the Fortran-C interface. One should also make sure that `npar_mig<=npar/ncpus`. It is also beneficial to set `mpar_loc=2*npar/ncpus`. Sometimes one may encounter the following error, “additional relocation overflows omitted from the output” due to the 2G memory limit (caused by large
static array). It is \texttt{mpar\_loc} that determines the usage of memory instead of \texttt{npar}. There are two ways to resolve this issue:

- Use a specific memory model to generate code and store data by setting the following for Intel compiler in your configuration file,

  \begin{verbatim}
  FFLAGS += -shared-intel -mcmodel=medium
  \end{verbatim}

  This will, however, affect the performance of the code. This method can handle at least the following setup,

  \begin{verbatim}
  integer, parameter :: ncpus=256, nprocx=4, nprocy=8, nprocz=ncpus/(nprocx*nprocy)
  integer, parameter :: nxgrid=1024, nygrid=nxgrid, nzgrid=nxgrid
  integer(kind=ikind8), parameter :: npar=124999680
  integer, parameter :: npar_stalk=100000, npar_mig=npar/10
  integer, parameter :: mpar_loc=npar/5
  \end{verbatim}

- Increase \texttt{ncpu} and decrease \texttt{mpar\_loc}. For \texttt{npar=124999680}, \texttt{ncpu=1024} is needed.

  \begin{verbatim}
  integer, parameter :: ncpus=1024, nprocx=8, nprocy=8, nprocz=ncpus/(nprocx*nprocy)
  integer, parameter :: nxgrid=1024, nygrid=nxgrid, nzgrid=nxgrid
  integer(kind=ikind8), parameter :: npar=124999680
  integer, parameter :: mpar_loc=5*npar/ncpus
  integer, parameter :: npar_stalk=100000, npar_mig=npar/ncpus
  \end{verbatim}

It is worth noting that even without particles, a simulation with \texttt{1024^3} resolution requires at least 512 CPUs to be compiled.

### 5.24.3 Random number generator

There are several methods to generate random number in the code. It is worth noting that when simulating coagulation with the super-particle approach, one should use the intrinsic random number generator of FORTRAN instead of the one implemented in the code. When invoking \texttt{random\_number\_wrapper}, there will be back-reaction to the gas flow. This unexpected back-reaction can be tracked by inspecting the power spectra, which exhibits the oscillation at the tail. To avoid this, one should set \texttt{luser\_random\_number\_wrapper=F} under the module \texttt{particles\_coag\_run\_pars} in \texttt{run.in}.

### 5.25 Non-cartesian coordinate systems

Spherical coordinates are invoked by adding the following line in the file `start.in`

\begin{verbatim}
&init_pars
  coord_system=’spherical_coords’
\end{verbatim}

One can also invoke cylindrical coordinates by saying \texttt{cylindrical\_coords} instead. In practice, the names \((x,y,z)\) are still used, but they refer then to \((r,\theta,\phi)\) or \((r,\phi,z)\) instead.

When working with curvilinear coordinates it is convenient to use differential operators in the non-coordinate basis, so the derivatives are taken with respect to length, and not in a mixed fashion with respect to length for \(\partial/\partial r\) and with respect to angle for
\(\partial/\partial \theta\) and \(\partial/\partial \phi\). The components in the non-coordinate basis are denoted by hats, see, e.g., [28], p. 213; see also Appendix B of [29]. For spherical polar coordinates the only nonvanishing Christoffel symbols (or connection coefficients) are
\[
\Gamma^{\hat{\theta}}_{\hat{r}\hat{\theta}} = \Gamma^{\hat{\phi}}_{\hat{r}\hat{\phi}} = -\Gamma^{\hat{\phi}}_{\hat{\theta}\hat{\phi}} = -\Gamma^{\hat{\phi}}_{\hat{\phi}\hat{\phi}} = 1/r, \quad (37)
\]
\[
\Gamma^{\hat{\phi}}_{\hat{\theta}\hat{\phi}} = -\Gamma^{\hat{\phi}}_{\hat{\phi}\hat{\theta}} = \cot \theta/r. \quad (38)
\]

The Christoffel symbols enter as correction terms for the various differential operators in addition to a term calculated straightforwardly in the non-coordinate basis. The derivatives of some relevant Christoffel symbols are
\[
\Gamma^{\hat{\theta}}_{\hat{r}\hat{\theta},\hat{\hat{\theta}}} = \Gamma^{\hat{\phi}}_{\hat{r}\hat{\phi},\hat{\hat{\phi}}} = \Gamma^{\hat{\theta}}_{\hat{\phi},\hat{\hat{\phi}}} = \Gamma^{\hat{\phi}}_{\hat{\theta},\hat{\hat{\phi}}} = 0 \quad (39)
\]
\[
\Gamma^{\hat{\phi}}_{\hat{\theta},\hat{\hat{\phi}}} = \Gamma^{\hat{\phi}}_{\hat{\phi},\hat{\hat{\phi}}} = -r^{-2} \quad (40)
\]
\[
\Gamma^{\hat{\phi}}_{\hat{\theta},\hat{\phi}} = -r^{-2} \sin^{-2} \theta \quad (41)
\]

Further details are given in Appendix I.
6. The equations

The equations solved by the PENCIL CODE are basically the standard compressible MHD equations. However, the modular structure allows some variations of the MHD equations, as well as switching off some of the equations or individual terms of the equation (nomagnetic, noentropy, etc.).

In this section the equations are presented in their most complete form. It may be expected that the code can evolve most subsets or simplifications of these equations.

6.1 Continuity equation

In the code the continuity equation, \( \partial \rho / \partial t + \nabla \cdot \rho \mathbf{u} = 0 \), is written in terms of \( \ln \rho \),

\[
\frac{D \ln \rho}{Dt} = -\nabla \cdot \mathbf{u} .
\]  

Here \( \rho \) denotes density, \( \mathbf{u} \) the fluid velocity, \( t \) is time and \( D/Dt \equiv \partial/\partial t + \mathbf{u} \cdot \nabla \) is the convective derivative.

6.2 Equation of motion

In the equation of motion, using a perfect gas, the pressure term, can be expressed as

\[ -\rho^{-1} \nabla p = -c_s^2 (\nabla s/c_p + \nabla \ln \rho) , \]

where the squared sound speed is given by

\[
c_s^2 = \frac{\gamma p}{\rho} = c_0^2 \exp \left[ \frac{\gamma s/c_p + (\gamma-1) \ln \rho}{\rho_0} \right] ,
\]  

and \( \gamma = c_p/c_v \) is the ratio of specific heats, or adiabatic index. Note that \( c_s^2 \) is proportional to the temperature with \( c_s^2 = (\gamma - 1)c_p T \).

The equation of motion is accordingly

\[
\frac{D \mathbf{u}}{Dt} = -c_s^2 \nabla \left( \frac{s}{c_p} + \ln \rho \right) - \nabla \Phi_{grav} + \frac{j \times \mathbf{B}}{\rho} + \nu \left( \nabla^2 \mathbf{u} + \frac{1}{3} \nabla \nabla \cdot \mathbf{u} + 2 \mathbf{S} \cdot \nabla \ln \rho \right) + \zeta (\nabla \nabla \cdot \mathbf{u}) ;
\]  

Here \( \Phi_{grav} \) is the gravity potential, \( j \) the electric current density, \( \mathbf{B} \) the magnetic flux density, \( \nu \) is kinematic viscosity, \( \zeta \) describes a bulk viscosity, and, in Cartesian coordinates

\[
S_{ij} = \frac{1}{2} \left( \frac{\partial u_i}{\partial x_j} + \frac{\partial u_j}{\partial x_i} - \frac{2}{3} \delta_{ij} \nabla \cdot \mathbf{u} \right)
\]  

is the rate-of-shear tensor that is traceless, because it can be written as the generic rate-of-strain tensor minus its trace. In curvilinear coordinates, we have to replace partial differentiation by covariant differentiation (indicated by semicolons), so we write \( S_{ij} = \frac{1}{2} (u_{i,j} + u_{j,i}) - \frac{1}{3} \delta_{ij} \nabla \cdot \mathbf{u} \).

The interpretation of the two viscosity terms varies greatly depending upon the Viscosity module used, and indeed on the parameters given to the module. See [6.6].

For isothermal hydrodynamics, see [6.4] below.
6.3 Induction equation

\[ \frac{\partial A}{\partial t} = u \times B - \eta \mu_0 j. \]  

(46)

Here \( A \) is the magnetic vector potential, \( B = \nabla \times A \) the magnetic flux density, \( \eta = 1/(\mu_0 \sigma) \) is the magnetic diffusivity (\( \sigma \) denoting the electrical conductivity), and \( \mu_0 \) the magnetic vacuum permeability. This form of the induction equation corresponds to the Weyl gauge \( \Phi = 0 \), where \( \Phi \) denotes the scalar potential.

6.4 Entropy equation

The current thermodynamics module entropy formulates the thermal part of the physics in terms of entropy \( s \), rather than thermal energy \( e \), which you may be more familiar with. Thus the two fundamental thermodynamical variables are \( \ln \rho \) and \( s \). The reason for this choice of variables is that entropy is the natural physical variable for (at least) convection processes: the sign of the entropy gradient determines convective (in)stability, the Rayleigh number is proportional to the entropy gradient of the associated hydrostatic reference solution, etc. The equation solved is

\[ \rho T \frac{Ds}{Dt} = \mathcal{H} - C + \nabla \cdot (K \nabla T) + \eta \mu_0 j^2 + 2 \rho \nu S \otimes S + \zeta \rho (\nabla \cdot u)^2. \]

(47)

Here, \( T \) is temperature, \( c_p \) the specific heat at constant pressure, \( \mathcal{H} \) and \( C \) are explicit heating and cooling terms, \( K \) is the radiative (thermal) conductivity, \( \zeta \) describes a bulk viscosity, and \( S \) is the rate-of-shear tensor that is traceless.

In the entropy module we solve for the specific entropy, \( s \). The heat conduction term on the right hand side can be written in the form

\[ \nabla \cdot (K \nabla T) \]

\[ = c_p \chi \left[ \nabla^2 \ln T + \nabla \ln T \cdot \nabla (\ln T + \ln \chi + \ln \rho) \right] \]

(48)

\[ = c_p \chi \left[ \gamma \nabla^2 s/c_p + (\gamma - 1) \nabla^2 \ln \rho \right] \]

\[ + c_p \chi \left[ \gamma \nabla s/c_p + (\gamma - 1) \nabla \ln \rho \right] \cdot \left[ \gamma (\nabla s/c_p + \nabla \ln \rho) + \nabla \ln \chi \right], \]

(49)

where \( \chi = K/(\rho c_p) \) is the thermal diffusivity. The latter equation shows that the diffusivity for \( s \) is \( \gamma \chi \), which is what we have used in Eq. (24).

In an alternative formulation for a constant \( K \), the heat conduction term on the right hand side can also be written in the form

\[ \nabla \cdot (K \nabla T) \]

\[ = \frac{K}{\rho} \left[ \nabla^2 \ln T + (\nabla \ln T)^2 \right] \]

(50)

(51)

Note that by setting \( \gamma = 1 \) and initially \( s = 0 \), one obtains an isothermal equation of state (albeit at some unnecessary expense of memory). Similarly, by switching off the evolution terms of entropy, one immediately gets polytropic behavior (if \( s \) was initially constant) or generalized polytropic behavior (where \( s \) is not uniform, but \( \partial s/\partial t = 0 \)).

A better way to achieve isothermality is to use the noentropy module.
6.4 Entropy equation

6.4.1 Viscous heating

We can write the viscosity as the divergence of a tensor $\tau_{ij,j}$,

$$\frac{\partial u_i}{\partial t} = \ldots + \tau_{ij,j},$$  \hspace{1cm} (52)

where $\tau_{ij} = 2\nu \rho S_{ij}$ is the stress tensor. The viscous power density $P$ is

$$P = u_i \tau_{ij,j}$$  \hspace{1cm} (53)

$$= \frac{\partial}{\partial x_j} (u_i \tau_{ij}) - u_{i,j} \tau_{ij}$$  \hspace{1cm} (54)

The term under the divergence is the viscous energy flux and the other term is the kinetic energy loss due to heating. The heating term $u_{i,j} \tau_{ij}$ is positive definite, because $\tau_{ij}$ is a symmetric tensor and the term only gives a contribution from the symmetric part of $u_{i,j}$, which is $\frac{1}{2}(u_{i,j} + u_{j,i})$, so

$$u_{i,j} \tau_{ij} = \frac{1}{2} \nu \rho (u_{i,j} + u_{j,i})(2S_{ij}).$$  \hspace{1cm} (55)

But, because $S_{ij}$ is traceless, we can add anything proportional to $\delta_{ij}$ and, in particular, we can write

$$u_{i,j} \tau_{ij} = \frac{1}{2} (u_{i,j} + u_{j,i})(2\nu \rho S_{ij})$$  \hspace{1cm} (56)

$$= \frac{1}{2} (u_{i,j} + u_{j,i} - \frac{1}{3} \delta_{ij} \nabla \cdot u)(2\nu \rho S_{ij})$$  \hspace{1cm} (57)

$$= 2\nu \rho S^2,$$  \hspace{1cm} (58)

which is positive definite.

6.4.2 Alternative description

By setting pretend\_lnTT=T in init\_pars or run\_pars (i.e. the general part of the name list) the logarithmic temperature is used instead of the entropy. This has computational advantages when heat conduction (proportional to $K \nabla T$) is important. Another alternative is to use another module, i.e. set ENTROPY=temperature\_idealgas in ‘Makefile.local’.

When pretend\_lnTT=T is set, the entropy equation

$$\frac{\partial s}{\partial t} = -u \cdot \nabla s + \frac{1}{\rho T} \text{RHS}$$  \hspace{1cm} (59)

is replaced by

$$\frac{\partial \ln T}{\partial t} = -u \cdot \nabla \ln T + \frac{1}{\rho c_u T} \text{RHS} - (\gamma - 1) \nabla \cdot u,$$  \hspace{1cm} (60)

where RHS is the right hand side of equation (47).
6.5 Transport equation for a passive scalar

In conservative form, the equation for a passive scalar is
\[ \frac{\partial}{\partial t}(\rho c) + \nabla \cdot [\rho c \mathbf{u} - \rho D \nabla c] = 0. \] (61)

Here \( c \) denotes the concentration (per unit mass) of the passive scalar and \( D \) its diffusion constant (assumed constant). In the code this equation is solved in terms of \( \ln c \),

\[ \frac{D \ln c}{Dt} = D \left[ \nabla^2 \ln c + \left( \nabla \ln \rho + \nabla \ln c \right) \cdot \nabla \ln c \right]. \] (62)

Using \( \ln c \) instead of \( c \) has the advantage that it enforces \( c > 0 \) for all times. However, the disadvantage is that one cannot have \( c = 0 \). For this reason we ended up using the non-logarithmic version by invoking PSCALAR=pscalar_nolog.

6.6 Bulk viscosity

For a monatomic gas it can be shown that the bulk viscosity vanishes. We therefore don't use it in most of our runs. However, for supersonic flows, or even otherwise, one might want to add a shock viscosity which, in its simplest formulation, take the form of a bulk viscosity.

6.6.1 Shock viscosity

Shock viscosity, as it is used here and also in the Stagger Code of Åke Nordlund, is proportional to positive flow convergence, maximum over five zones, and smoothed to second order,

\[ \zeta_{\text{shock}} = c_{\text{shock}} \left( \left( \nabla \cdot \mathbf{u} \right)_{\text{max}} \right) \left( \min(\delta x, \delta y, \delta z) \right)^2, \] (63)

where \( c_{\text{shock}} \) is a constant defining the strength of the shock viscosity. In the code this dimensionless coefficient is called nu_shock, and it is usually chosen to be around unity.

Assume that the shock viscosity only enters as a bulk viscosity, so the whole stress tensor is then

\[ \tau_{ij} = 2 \nu S_{ij} + \rho \zeta_{\text{shock}} \delta_{ij} \nabla \cdot \mathbf{u}. \] (64)

Assume \( \nu = \text{const}, \) but \( \zeta \neq \text{const}, \) so

\[ \rho^{-1} F_{\text{visc}} = \nu \left[ \nabla^2 \mathbf{u} + \frac{1}{3} \nabla \nabla \cdot \mathbf{u} + 2 \mathbf{S} \cdot \nabla \ln \rho \right] + \zeta_{\text{shock}} \left[ \nabla \nabla \cdot \mathbf{u} + \left( \nabla \ln \rho + \nabla \ln \zeta_{\text{shock}} \right) \nabla \cdot \mathbf{u} \right]. \] (65)

and

\[ \rho^{-1} \Gamma_{\text{visc}} = 2 \nu S^2 + \zeta_{\text{shock}} (\nabla \cdot \mathbf{u})^2. \] (66)

In the special case with periodic boundary conditions, we have

\[ 2 \langle S^2 \rangle = \langle \omega^2 \rangle + \frac{4}{3} \langle (\nabla \cdot \mathbf{u})^2 \rangle. \]

6.7 Equation of state

In its present configuration only hydrogen ionization is explicitly included. Other constituents (currently He and H_2) can have fixed values. The pressure is proportional to the total number of particles, i.e.

\[ p = (n_{\text{HI}} + n_{\text{HII}} + n_{\text{H}_2} + n_e + n_{\text{He}} + ...) k_B T. \] (67)
It is convenient to normalize to the total number of H both in atomic and in molecular hydrogen, \( n_{\text{Htot}} \equiv n_\text{H} + 2n_\text{H}_2 \), where \( n_\text{H} + n_\text{HII} = n_\text{H} \), and define \( x_\text{e} \equiv n_\text{e}/n_{\text{Htot}} \), \( x_\text{He} \equiv n_\text{He}/n_{\text{Htot}} \), and \( x_\text{H}_2 \equiv n_\text{H}_2/n_{\text{Htot}} \). Substituting \( n_\text{H} = n_{\text{Htot}} - 2n_\text{H}_2 \), we have

\[
p = (1 - x_\text{H}_2 + x_\text{e} + x_\text{He} + \ldots)n_{\text{Htot}} k_B T. \tag{68}
\]

This can be written in the more familiar form

\[
p = \frac{\mathcal{R}}{\mu} \rho T, \tag{69}
\]

where \( \mathcal{R} = k_B/m_u \) and \( m_u \) is the atomic mass unit (which is for all practical purposes the same as \( m_{\text{Htot}} \)) and \( \mu = n_\text{H} + 2n_\text{H}_2 + n_\text{e} + 4n_\text{He} \)

\[
\mu = \frac{n_\text{H} + 2n_\text{H}_2 + n_\text{e} + 4n_\text{He}}{n_\text{H} + n_\text{H}_2 + n_\text{e} + n_\text{He}} = \frac{1 + 4x_\text{He}}{1 - x_\text{H}_2 + x_\text{e} + x_\text{He}}. \tag{70}
\]

is the mean molecular weight (which is here dimensionless; see Kippenhahn & Weigert 1990, p. 102). The factor 4 is really to be substituted for 3.9715 3. Some of the familiar relations take still the usual form, in particular \( e = c_v T \) and \( h = c_p T \) with \( c_v = \frac{3}{2} \mathcal{R}/\mu \) and \( c_p = \frac{5}{2} \mathcal{R}/\mu \).

The number ratio, \( x_\text{He} \), is more commonly expressed as the mass ratio, \( Y = m_\text{He}/(m_\text{Htot} + m_\text{He}) \), or \( Y = 4x_\text{He}/(1 + 4x_\text{He}) \), or \( 4x_\text{He} = (1/Y - 1)^{-1} \). For example, \( Y = 0.27 \) corresponds to \( x_\text{He} = 0.092 \) and \( Y = 0.25 \) to \( x_\text{He} = 0.083 \). Note also that for 100% H2 abundance, \( x_\text{H}_2 = 1/2 \).

In the following, the ionization fraction is given as \( y = n_\text{e}/n_\text{H} \), which can be different from \( x_\text{e} \) if there is H2. Substituting for \( n_\text{H} \) in terms of \( n_{\text{Htot}} \) yields \( y = x_\text{e}/(1 - 2x_\text{H}_2) \).

### 6.8 Ionization

This part of the code can be invoked by setting EOS=eos_ionization (or EOS=eos_temperature_ionization) in the ‘Makefile.local’ file. The equation of state described below works for variable ionization, and the entropy offset is different from that used in Eq. (43), which is now no longer valid. As a replacement, one can use EOS=eos_fixed_ionization, where the degree of ionization can be given by hand. Here the normalization of the entropy is the same as for EOS=eos_ionization. This case is described in more detail below.

We treat the gas as being composed of partially ionized hydrogen and neutral helium. These are four different particle species, each of which regarded as a perfect gas.

The ionization fraction \( y \), which gives the ratio of ionized hydrogen to the total amount of hydrogen \( n_\text{H} \), is obtained from the Saha equation which, in this case, may be written as

\[
\frac{y^2}{1 - y} = \frac{1}{n_\text{H}} \left( \frac{m_\text{e}k_BT}{2\pi\hbar^2} \right)^{3/2} \exp \left( -\frac{\chi_\text{H}}{k_BT} \right). \tag{71}
\]

The temperature \( T \) cannot be obtained directly from the PENCIL CODE’s independent variables \( \ln \rho \) and \( s \), but is itself dependent on \( y \). Hence, the calculation of \( y \) essentially becomes a root finding problem.

---

12We omit here the contribution of H2.
The entropy of a perfect gas consisting of particles of type $i$ is known from the Sackur-Tetrode equation

$$S_i = k_B N_i \left( \ln \left[ \frac{1}{n_{\text{tot}}} \left( \frac{m_i k_B T}{2\pi \hbar^2} \right)^{3/2} \right] + 5/2 \right).$$  \hspace{1cm} (72)$$

Here $N_i$ is the number of particles of a single species and $n_{\text{tot}}$ is the total number density of all particle species.

In addition to the individual entropies we also have to take the entropy of mixing, $S_{\text{mix}} = -N_{\text{tot}} k_B \sum_i p_i \ln p_i$, into account. Summing up everything, we can get a closed expression for the specific entropy $s$ in terms of $y$, $\ln \rho$ and $T$, which may be solved for $T$.

For given $\ln \rho$ and $s$ we are then able to calculate the ionization fraction $y$ by finding the root of

$$f(y) = \ln \left[ \frac{1 - y}{y^2} \frac{1}{n_{\text{H}}} \left( \frac{m_e k_B T(y)}{2\pi \hbar^2} \right)^{3/2} \right] - \frac{\chi_{\text{H}}}{k_B T(y)}.$$  \hspace{1cm} (73)$$

In the ionized case, several thermodynamic quantities of the gas become dependent on the ionization fraction $y$ such as its pressure, $P = (1 + y + x_{\text{He}}) n_{\text{H}} k_B T$, and its internal energy, $E = \frac{3}{2}(1 + y + x_{\text{He}}) n_{\text{H}} k_B T + y \chi_{\text{H}}$, where $x_{\text{He}}$ gives the ratio of neutral helium to the total amount of hydrogen. The dependence of temperature on entropy is shown in Fig. 6 for different values of the density.

For further details regarding the procedure of solving for the entropy see Sect. 4.6 in the appendix.

6.8.1 Ambipolar diffusion

Another way of dealing with ionization in the PENCIL CODE is through use of the neutrals module. That module solves the coupled equations of neutral and ionized gas, in a two-fluid model.
\[ \frac{\partial \rho_i}{\partial t} = -\nabla \cdot (\rho_i u_i) + G \]  
\[ \frac{\partial \rho_n}{\partial t} = -\nabla \cdot (\rho_n u_n) - G \]  
\[ \frac{\partial (\rho_i u_i)}{\partial t} = -\nabla \cdot (\rho_i u_i : u_i) - \nabla \left( p_i + p_e + \frac{B^2}{2\mu_0} \right) + F \]  
\[ \frac{\partial (\rho_n u_n)}{\partial t} = -\nabla \cdot (\rho_n u_n : u_n) - \nabla p_n - F \]  
\[ \frac{\partial A}{\partial t} = u_i \times B \]  

where the subscripts \( n \) and \( i \) are for neutral and ionized, respectively. The terms \( G \) and \( F \), through which the two fluids exchange mass and momentum, are given by

\[ G = \zeta \rho_n - \alpha \rho_i^2 \]  
\[ F = \zeta \rho_n u_n - \alpha \rho_i^2 u_i + \gamma \rho_i \rho_n (u_n - u_i) \]

In the above equations, \( \zeta \) is the ionization coefficient, \( \alpha \) is the recombination coefficient, and \( \gamma \) the collisional drag strength. By the time of writing (spring 2009), these three quantities are supposed constant. The electron pressure \( p_e \) is also assumed equal to the ion pressure. Only isothermal neutrals are supported so far.

In the code, Eq. (74) and Eq. (76) are solved in `density.f90` and `hydro.f90` respectively. Equation (75) is solved in `neutraldensity.f90` and Eq. (77) in `neutralvelocity.f90`. The sample ‘1d-test/ambipolar-diffusion’ has the current setup for a two-fluid simulation with ions and neutrals.

6.9 Combustion

The easiest entry into the subject of simulating combustion is through samples/0d-tests/chemistry_H2_ignition_rkf or samples/1d-tests/H2_flamespeed. The former case is studying H2 ignition delay, while the second one is focusing on H2 flamespeed. If you want to study turbulent premixed flames, the recommended cases are samples/2d-tests/2d_methane_flame and samples/turbulent_flame. Here, the first of these examples is not really realistic since its only 2D, but this does of course make it faster to try out. Also, this case is easier to set up. The most realistic test case is samples/turbulent_flame, which studies a full 3D hydrogen flame. This test case requires that a set of smaller pre-runs are finalized before the main simulation can be run (see the associated README file for more details).

The chemistry_H2_ignition_rkf directory, for example, has the file ‘tran.dat’, which contains the parameters characterizing the transport properties, and ‘chem.inp’, which contains the NASA polynomials; see Eq. (18) of Ref. [2].

6.10 Radiative transfer

Here we only state the basic equations. A full description about the implementation is given in Sect. [H.7] and in the original paper by Heinemann et al. (2006).
The basic equation for radiative transfer is
\[
\frac{dI}{d\tau} = -I + S ,
\]
where
\[
\tau \equiv \int_0^s \chi(s') \, ds'
\]
is the optical depth (\(s\) is the geometrical coordinate along the ray).

Note that radiative transfer is called also in 'start.csh', and again each time a snapshot is being written, provided the output of auxiliary variables is being requested \texttt{Iwrite_aux=T}. (Also, of course, the pencil check runs radiative transfer 7 times, unless you put \texttt{pencil_check_small=F}.)

### 6.11 Self-gravity

The \texttt{PENCIL CODE} can consider the self-gravity of the fluid in the simulation box by adding the term
\[
\frac{\partial u}{\partial t} = \ldots - \nabla \phi_{\text{self}}
\]
to the equation of motion. The self-potential \(\phi_{\text{self}}\) (or just \(\phi\) for simplicity) satisfies Poisson's equation
\[
\nabla^2 \phi = 4\pi G \rho .
\]
The solution for a single Fourier component at scale \(k\) is
\[
\phi_k = -\frac{4\pi G \rho_k}{k^2} .
\]
Here we have assumed periodic boundary conditions. The potential is obtained by Fourier-transforming the density, then finding the corresponding potential at that scale, and finally Fourier-transforming back to real space.

The \(x\)-direction in the shearing sheet is not strictly periodic, but is rather shear periodic with two connected points at the inner and outer boundary separated by the distance \(\Delta y(t) = \mod[(3/2)\Omega_0 L_x t, L_y]\) in the \(y\)-direction. We follow here the method from [19] to allow for shear-periodic boundaries in the Fourier method for self-gravity. First we take the Fourier transform along the periodic \(y\)-direction. We then shift the entire \(y\)-direction by the amount \(\delta y(x) = \Delta y(t) x / L_x\) to make the \(x\)-direction periodic. Then we proceed with Fourier transforms along \(x\) and then \(z\). After solving the Poisson equation in Fourier space, we transform back to real space in the opposite order. We differ here from the method by [19] in that we shift in Fourier space rather than in real space.[13]

The Fourier interpolation formula has the advantage over polynomial interpolation in that it is continuous and smooth in all its derivatives.

### 6.12 Incompressible and anelastic equations

This part has not yet been documented and is still under development.

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[13] We were kindly made aware of the possibility of interpolating in Fourier space by C. McNally on his website.
6.13 Dust equations

The code treats gas and dust as two separate fluids. The dust and the gas interact through a drag force. This force can most generally be written as an additional term to the equation of motion as

$$\frac{D\mathbf{u}_d}{Dt} = \ldots - \frac{1}{\tau_s} (\mathbf{u}_d - \mathbf{u}). \quad (86)$$

Here $\tau_s$ is the so-called stopping time of the considered dust species. This measures the coupling strength between dust and gas. In the Epstein drag regime

$$\tau_s = \frac{a_d \rho_s}{c_s \rho}, \quad (87)$$

where $a_d$ is the radius of the dust grain and $\rho_s$ is the solid density of the dust grain.

Two other important effects work on the dust. The first is coagulation controlled by the discrete coagulation equation

$$\frac{dn_k}{dt} = \frac{1}{2} \sum_{i+j=k} A_{ij} n_i n_j - n_k \sum_{i=1}^{\infty} A_{ik} n_i. \quad (88)$$

In the code $N$ discrete dust species are considered. Also the bins are logarithmically spaced in order to give better mass resolution. It is also possible to keep track of both number density and mass density of each bin, corresponding to having a variable grain mass in each bin.

Dust condensation is controlled by the equation

$$\frac{dN}{dt} = \frac{1}{\tau_{\text{cond}}} N \frac{d+1}{d} \cdot (90)$$

Here $N$ is the number of monomers in the dust grain (such as water molecules) and $d$ is the physical dimension of the dust grain. The condensation time $\tau_{\text{cond}}$ is calculated from

$$\frac{1}{\tau_{\text{cond}}} = A_1 v_{\text{th}} \alpha n_{\text{mon}} \left\{ 1 - \frac{1}{S_{\text{mon}}} \right\}, \quad (90)$$

where $A_1$ is the surface area of a monomer, $\alpha$ is the condensation efficiency, $n_{\text{mon}}$ is the number density of monomers in the gas and $S_{\text{mon}}$ is the saturation level of the monomer given by

$$S_{\text{mon}} = \frac{P_{\text{mon}}}{P_{\text{sat}}}. \quad (91)$$

Here $P_{\text{sat}}$ is the saturated vapor pressure of the monomer. Currently only water ice has been implemented in the code.

All dust species fulfill the continuity equation

$$\frac{\partial \rho_d}{\partial t} + \nabla \cdot (\rho_d \mathbf{u}_d) = 0. \quad (92)$$

See master’s thesis of A. Johansen (can be downloaded from [http://www.mpia.de/homes/johansen/research_en.php](http://www.mpia.de/homes/johansen/research_en.php))
Cosmic rays are treated in the diffusion approximation. The equation of state is $p_c = (\gamma_c)c$ where the value of $\gamma_c$ is usually somewhere between $14/9$ and $4/3$. In the momentum equation (44) the cosmic ray pressure force, $-\rho^{-1}\nabla p_c$ is added on the right hand side, and $e_c$ satisfies the evolution equation

$$\frac{\partial e_c}{\partial t} + \nabla \cdot (e_c \mathbf{u}) + p_c \nabla \cdot \mathbf{u} = \partial_i (K_{ij} \partial_j e_c) + Q_c,$$

where $Q_c$ is a source term and

$$K_{ij} = K_\perp \delta_{ij} + (K_\parallel - K_\perp) \hat{B}_i \hat{B}_j$$

is an anisotropic diffusivity tensor.

In the non-conservative formulation of this code it is advantageous to expand the diffusion term using the product rule, i.e.

$$\partial_i (K_{ij} \partial_j e_c) = -\mathbf{U}_c \cdot \nabla e_c + K_{ij} \partial_i \partial_j e_c.$$

where $\mathbf{U}_c = -\partial K_{ij}/\partial x_j$ acts like an extra velocity trying to straighten magnetic field lines. We can write this term also as $\mathbf{U}_c = -(K_\parallel - K_\perp) \nabla \cdot (\mathbf{B} \mathbf{B})$, where the last term is a divergence of the dyadic product of unit vectors. However, near magnetic nulls, this term can becomes infinite. In order to avoid this problem we are forced to limit $\nabla \cdot (\mathbf{B} \mathbf{B})$, and hence $|\mathbf{U}_c|$, to the maximum possible value that can be resolved at a given resolution.

A physically appealing way of limiting the maximum propagation speed is to restore an explicit time dependence in the equation for the cosmic ray flux, and to replace the diffusion term in Eq. (93) by a divergence of a flux that in turn obeys the equation

$$\frac{\partial \mathbf{F}_{ci}}{\partial t} = -\mathbf{K}_{ij} \nabla_j e_c - \frac{\mathbf{F}_{ci}}{\tau} \quad \text{ (non-Fickian diffusion)},$$

where $K_{ij} = \tau \mathbf{K}_{ij}$ would be the original diffusion tensor of Eq. (94), if the time derivative were negligible. Further details are described in Snodin et al. (2006).

**6.15 Chiral MHD**

At high energies, $k_B T \gtrsim 10$ MeV, a quantum effect called the *chiral magnetic effect* (CME) can modify the MHD equations. The CME occurs in magnetized relativistic plasmas, in which the number density of left-handed fermions, $n_L$, differs from the one of right-handed fermions, $n_R$ (see e.g., Kharzeev et al. (2013) for a review). This asymmetry is described by the chiral chemical potential $\mu_5 \equiv 6 (n_L - n_R) (\hbar c)^3/(k_B T)^2$, where $T$ is the temperature, $k_B$ is the Boltzmann constant, $c$ is the speed of light, and $\hbar$ is the reduced Planck constant. In the presence of a magnetic field, $\mu_5$ leads to the occurrence of the current

$$J_{\text{CME}} = \frac{\alpha_{\text{em}}}{\pi \hbar} \mu_5 \mathbf{B},$$

where

$$\partial_j (\hat{B}_i \hat{B}_j) = (\delta_{ij} - 2\hat{B}_i \hat{B}_j) \hat{B}_j B_{k,j} / |\mathbf{B}|,$$

where derivatives of $\mathbf{B}$ are calculated as $B_{i,j} = \epsilon_{ijk} A_{i,j,k}$. 

\[\text{[15] In practice, we calculate } \partial_j (\hat{B}_i \hat{B}_j) = (\delta_{ij} - 2\hat{B}_i \hat{B}_j) \hat{B}_j B_{k,j} / |\mathbf{B}|, \text{ where derivatives of } \mathbf{B} \text{ are calculated as } B_{i,j} = \epsilon_{ijk} A_{i,j,k}.\]
where $\alpha_{em} \approx 1/137$ is the fine structure constant.

The chiral current (97) adds to the classical Ohmic current, leading to a modification of the Maxwell equations. As a result, the induction equation is extended by one additional term:

$$\frac{\partial A}{\partial t} = U \times B - \eta (\nabla \times B - \mu B),$$

(98)

where the chiral chemical potential $\mu_5$ is normalized such that $\mu = (4\alpha_{em}/hc)\mu_5$. The latter is determined by the evolution equation

$$\frac{D\mu}{Dt} = D_5 \Delta \mu + \lambda \eta [B \cdot (\nabla \times B) - \mu B^2] - \Gamma_\mu \mu,$$

(99)

where $D_5$ is a chiral diffusion coefficient, $\lambda$ the chiral feedback parameter, and $\Gamma_\mu$ the rate of chiral flipping reactions. All remaining evolution equations are the same as in classical MHD. Details on the derivation of the chiral MHD equations can be found in Boyarsky et al. (2015) and Rogachevskii et al. (2017).

In the Pencil Code, the chiral MHD equations can be solved by adding the line

```
SPECIAL = special/chiral_mhd
```

to `src/Makefile.local`. Further, for running the chiral MHD module, one needs to add

```make
&special_init_pars
  initspecial='const', mu5_const=10.
```

to `start.in` and

```make
&special_run_pars
  diffmu5=1e-4, lambda5=1e3, cdtchiral=1.0
```

to `run.in`, where we have chosen exemplary values for the chiral parameters.

Caution should be taken when solving the chiral MHD equations numerically, since the evolution of the plasma can be strongly affected by the CME. In particular, the initial value of $\mu$ is related to a small-scale dynamo instability. In order to resolve this instability in a domain of size $(2\pi)^3$, the minimum number of grid points is given as:

$$N_{grid} \gtrsim \left( \frac{\mu_0}{\lambda} \right)^{1/2} \frac{2\pi}{\nu \text{Re}_{\text{mesh, crit}}},$$

(100)

Also the value of $\lambda$ should not be chosen too small, since it scales inversely with the saturation magnetic helicity produced by a chiral dynamo. Hence, for a very small $\lambda$ parameter, the Alfvén time step becomes extremely small in the non-linear stage which can lead to a code crash. More details on the numerical modeling of chiral MHD can be found in Schober et al. (2018).

6.16 Particles

Particles are entities that each have a space coordinate and a velocity vector, where a fluid only has a velocity vector field (the continuity equation of a fluid in some way corresponds to the space coordinate of particles). In the code particles are present either as tracer particles or as dust particles.
6.16.1 Tracer particles

Tracer particles always have the local velocity of the gas. The dynamical equations are thus
\[
\frac{\partial x_i}{\partial t} = u,
\]  
(101)
where the index \(i\) runs over all particles. Here \(u\) is the gas velocity at the position of the particle. One can choose between a first order (default) and a second order spline interpolation scheme (set \texttt{numberic_interpolation=T} in \texttt{&particles_initpars}) to calculate the gas velocity at the position of a tracer particle.

The sample run ‘\texttt{samples/dust-vortex}’ contains the latest setup for tracer particles.

6.16.2 Dust particles

Dust particles are allowed to have a velocity that is not similar to the gas,
\[
\frac{dx_i}{dt} = v_i. 
\]  
(102)

The particle velocity follows an equation of motion similar to a fluid, only there is no advection term. Dust particles also experience a drag force from the gas (proportional to the velocity difference between a particle and the gas).
\[
\frac{dv_i}{dt} = \ldots - \frac{1}{\tau_s} (v_i - u). 
\]  
(103)

Here \(\tau_s\) is the stopping time of the dust particle. The interpolation of the gas velocity to the position of a particle is done using one of three possible particle-mesh schemes,

- NGP (Nearest Grid Point, default)
  The gas velocity at the nearest grid point is used.
- CIC (Cloud in Cell, set \texttt{particlemesh_cic=T})
  A first order interpolation is used to obtain the gas velocity field at the position of a particle. Affects 8 grid points.
- TSC (Triangular Shaped Cloud, set \texttt{particlemesh_tsc=T})
  A second order spline interpolation is used to obtain the gas velocity field at the position of a particle. Affects 27 grid points.

The particle description is the proper description of dust grains, since they do not feel any pressure forces (too low number density). Thus there is no guarantee that the grains present within a given volume will be equilibrated with each other, although drag force may work for small grains to achieve that. Larger grains (meter-sized in protoplanetary discs) must be treated as individual particles.

To conserve momentum the dust particles must affect the gas with a friction force as well. The strength of this force depends on the dust-to-gas ratio \(\epsilon_d\), and it can be safely ignored when there is much more gas than there is dust, e.g. when \(\epsilon_d = 0.01\). The friction force on the gas appears in the equation of motion as
\[
\frac{\partial u}{\partial t} = \ldots - \frac{\rho_p}{\rho} \left( \frac{\partial v_p}{\partial t} \right)_\text{drag}. 
\]  
(104)
Here $\rho_p^{(i)}$ is the dust density that particle $i$ represents. This can be set through the parameter $\text{eps\_todt}$ in $\text{particle\_init\_pars}$. The drag force is assigned from the particles onto the mesh using either NGP, CIC or TSC assignment. The same scheme is used both for interpolation and for assignment to avoid any risk of a particle accelerating itself (see Hockney & Eastwood 1981).

6.17 $N$-body solver

The $N$-body code takes advantage of the existing Particles module, which was coded with the initial intent of treating solid particles whose radius $a_*$ is comparable to the mean free path $\lambda$ of the gas, for which a fluid description is not valid. A $N$-body implementation based on that module only needed to include mass as extra state for the particles, solve for the $N^2$ gravitational pair interactions and distinguish between the $N$-body and the small bodies that are mapped into the grid as a $\rho_p$ density field.

The particles of the $N$-body ensemble evolve due to their mutual gravity and by interacting with the gas and the swarm of small bodies. The equation of motion for particle $i$ is

$$\frac{dv_{p_i}}{dt} = F_{g_i} - \sum_{j \neq i}^{N} \frac{GM_j}{R_{ij}^2} \hat{R}_{ij}$$

(105)

where $R_{ij} = |r_{p_i} - r_{p_j}|$ is the distance between particles $i$ and $j$, and $\hat{R}_{ij}$ is the unit vector pointing from particle $j$ to particle $i$. The first term of the R.H.S. is the combined gravity of the gas and of the dust particles onto the particle $i$, solved via

$$F_{g_i} = -G \int_V \frac{[\rho_g(r) + \rho_p(r)]\hat{R}_i}{(R_i^2 + b_i^2)^{3/2}} dV,$$

(106)

where the integration is carried out over the whole disk. The smoothing distance $b_i$ is taken to be as small as possible (a few grid cells). For few particles ($<10$), calculating the integral for every particle is practical. For larger ensembles one would prefer to solve the Poisson equation to calculate their combined gravitational potential.

The evolution of the particles is done with the same third-order Runge-Kutta time-stepping routine used for the gas. The particles define the timestep also by the Courant condition that they should not move more than one cell at a time. For pure particle runs, where the grid is absent, one can adopt a fixed time-step $t_p \ll 2\pi\Omega_{fp}^{-1}$ where $\Omega_{fp}$ is the angular frequency of the fastest particle.

By now (spring 2009), no inertial accelerations are included in the $N$-body module, so only the inertial frame - with origin at the barycenter of the $N$-body ensemble - is available. For a simulation of the circular restricted three-body problem with mass ratio $q=10^{-3}$, the Jacobi constant of a test particle initially placed at position $(x,y)=(2,0)$ was found to be conserved up to one part in $10^5$ within the time span of 100 orbits.

We stress that the level of conservation is poor when compared to integrators designed to specifically deal with long-term $N$-body problems. These integrators are usually symplectic, unlike the Runge-Kutta scheme of the PENCIL CODE. As such, PENCIL should not be used to deal with evolution over millions of years. But for the time-span typical of astrophysical hydrodynamical simulations, this degree of conservation of the Jacobi constant can be deemed acceptable.
Table 5: Scale factor and conformal Hubble parameter for different values of \( n \).

<table>
<thead>
<tr>
<th>( n )</th>
<th>( a )</th>
<th>( \mathcal{H} )</th>
<th>( H )</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>1</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>1/2</td>
<td>( \eta/2 )</td>
<td>1/( \eta )</td>
<td>1/( \eta )</td>
</tr>
<tr>
<td>2/3</td>
<td>( \eta^2/3 )</td>
<td>2/( \eta )</td>
<td>6/( \eta^2 )</td>
</tr>
</tbody>
</table>

As an extension of the particle’s module, the \( N \)-body is fully compatible with the parallel optimization of PENCIL, which further speeds up the calculations. Parallelization, however, is not yet possible for pure particle runs, since it relies on splitting the grid between the processors. At the time of writing (spring 2009), the \( N \)-body code does not allow the particles to have a time-evolving mass.

6.18 Test-field equations

The test-field method is used to calculate turbulent transport coefficients for magneto-hydrodynamics. This is a rapidly evolving field and we refer the interested reader to recent papers in this field, e.g. by Sur et al. (2008) or Brandenburg et al. (2008). For technical details; see also Sect. F.4.

6.19 Gravitational wave equations

The expansion of the universe with time is described by the scale factor \( a(\tau) \), where \( \tau \) is the physical time. Using conformal time, \( t(\tau) = \int_0^\tau d\tau'/a(\tau') \), and dependent variables that are appropriately scaled with powers of \( a \), the hydromagnetic equations can be expressed completely without scale factor [11, 18]. This is not true, however, for the gravitational wave (GW) equations, where a dependence on \( a \) remains [18]. The time dependence of \( a \) can be modeled as a power law, \( a \propto \tau^n \), where \( n = 1/2 \) applies to the radiation-dominated era; see Table 5 the general relationship. To compare with cases where the expansion is ignored, we put \( n = 0 \).

In the transverse traceless (TT) gauge, the six components of the spatial part of the symmetric tensor characterizing the linearized evolution of the metric perturbations \( h_{ij} \), reduce to two components which, in the linear polarization basis, are the \( + \) and \( \times \) polarizations. However, the projection onto that basis is computationally intensive, because it requires nonlocal operations involving Fourier transformations. It is therefore advantageous to evolve instead the perturbation of the metric tensor, \( h_{ij} \), in an arbitrary gauge, compute then \( h_{ij}^{\text{TT}} \) in the TT gauge, and perform then the decomposition into the linear polarization basis whenever we compute diagnostic quantities such as averages or spectra. Thus, we solve the linearized GW equation in the form [18]

\[
\frac{\partial^2 h_{ij}}{\partial t^2} = -2\mathcal{H} \frac{\partial h_{ij}}{\partial t} + c^2 \nabla^2 h_{ij} + \frac{16\pi G}{a^2 c^2} T_{ij}
\]  (107)

for the six components \( 1 \leq i \leq j \leq 3 \), where \( t \) is comoving time, \( a \) is the scale factor, \( \mathcal{H} = \dot{a}/a \) is the comoving Hubble parameter, \( T_{ij} \) is the source term, \( c \) is the speed of light, and \( G \) is Newton’s constant. For \( n = 0 \), when the cosmic expansion is ignored, we have
\[ a = 1 \text{ and } \mathcal{H} = 0. \text{ In practice, we solve the GW equation for the scaled variable } \hat{h}_{ij} = a h_{ij}, \]

\[
\frac{\partial^2 \hat{h}_{ij}}{\partial t^2} = c^2 \nabla^2 \hat{h}_{ij} + \frac{16\pi G}{ac^2} T_{ij}. \tag{108}
\]

For the numerical treatment of Eq. (107) or Eq. (108) and equations (110)–(112).

The source term is chosen to be the traceless part of the stress tensor,

\[
T_{ij}(x, t) = \rho u_i u_j - B_i B_j - \frac{1}{3} \delta_{ij} (\rho u^2 - B^2). \tag{109}
\]

The removal of the trace is in principle not necessary, but it helps preventing a continuous build-up of a large trace, which would be numerically disadvantageous. We have ignored here the viscous stress, which is usually small.

We compute \( T_{ij} \) by solving the energy, momentum, and induction equations for an ultra-relativistic gas in the form \cite{11, 13}:

\[
\frac{\partial \ln \rho}{\partial t} = -\frac{4}{3} (\nabla \cdot u + u \cdot \nabla \ln \rho) + \frac{1}{\rho} \left[ u \cdot (J \times B) + \eta J^2 \right], \tag{110}
\]

\[
\frac{D u}{D t} = \frac{u}{3} (\nabla \cdot u + u \cdot \nabla \ln \rho) - \frac{u}{\rho} \left[ u \cdot (J \times B) + \eta J^2 \right]
- \frac{1}{4} \nabla \ln \rho + \frac{3}{4\rho} J \times B + \frac{2}{\rho} \nabla \cdot (\rho \nu S) + f, \tag{111}
\]

\[
\frac{\partial B}{\partial t} = \nabla \times (u \times B - \eta J), \tag{112}
\]

where \( B = \nabla \times A \) is the magnetic field expressed in terms of the magnetic vector potential to ensure that \( \nabla \cdot B = 0 \), \( J = \nabla \times B \) is the current density, \( D/Dt = \partial/\partial t + u \cdot \nabla \) is the advective derivative, \( S_{ij} = \frac{1}{2}(u_{ij,j} + u_{ji,j}) - \frac{1}{2} \delta_{ij} u_{k,k} \) is the trace-free rate of strain tensor, and \( p = \rho c_s^2 \) is the pressure, where \( c_s = c/\sqrt{3} \) is the sound speed for an ultra-relativistic gas. Lorentz-Heaviside units for the magnetic field are used.

We are interested in the rms value of the metric tensor perturbations and the GW energy density in the linear polarization basis. To compute \( h_{ij}^{TT} \) from \( h_{ij} \), we Fourier transform the six components of \( h_{ij} \) and \( \hat{h}_{ij} \),

\[
\hat{h}_{ij}(k, t) = \int h_{ij}(x, t) e^{-ik \cdot x} \, d^3x \quad \text{for } 1 \leq i \leq j \leq 3 \tag{113}
\]

and compute the components in the TT gauge as

\[
\hat{h}_{ij}^{TT}(k, t) = (P_i P_j - \frac{1}{2} P_{ij} P_{lm}) \hat{h}_{lm}(k, t), \tag{114}
\]

where \( P_{ij} = \delta_{ij} - \hat{k}_i \hat{k}_j \) is the projection operator, and \( \hat{k} = k/|k| \) is the unit vector of \( k \), with \( k = |k| \) being the modulus. Next, we compute the linear polarization bases

\[
e_i^+ = e_i^1 e_j^1 - e_i^2 e_j^2, \quad e_i^\times = e_i^1 e_j^2 + e_i^2 e_j^1, \tag{115}
\]

where \( e^1 \) and \( e^2 \) are unit vectors perpendicular to \( k \). Thus

\[
\tilde{h}_+(k, t) = \frac{1}{2} e^+_{ij}(k) \hat{h}_{ij}(k, t), \tag{116}
\]

\[
\tilde{h}_\times(k, t) = \frac{1}{2} e^\times_{ij}(k) \hat{h}_{ij}(k, t). \tag{117}
\]
We then return into real space and compute
\[ h_{+/\times}(x, t) = \int \hat{h}_{+/\times}(k, t) e^{i k \cdot x} \, d^3k/(2\pi)^3. \] (118)

Analogous calculations are performed for \( \dot{h}_{+/\times}(x, t) \), which are used to compute the GW energy via
\[ E_{\text{GW}}(t) = c^2/(32\pi G) \left( \langle \dot{h}_{+}^2 \rangle + \langle \dot{h}_{\times}^2 \rangle \right), \] (119)
where angle brackets denote volume averages.

Analogously to kinetic and magnetic energy and helicity spectra, it is convenient to compute the GW energy and polarization spectra integrated over concentric shells of surface \( \int_{4\pi} k^2 \, d\Omega_k \) in \( k \) space, defined by
\[ S_{\dot{h}}(k) = \int_{4\pi} \left( |\dot{h}_+|^2 + |\dot{h}_\times|^2 \right) k^2 \, d\Omega_k, \] (120)
\[ A_{\dot{h}}(k) = \int_{4\pi} 2 \text{Im} \left( \dot{h}_+ \dot{h}_\times^* \right) k^2 \, d\Omega_k, \] (121)
and normalized such that \( \int_0^\infty S_{\dot{h}}(k) \, dk = \langle \dot{h}_+^2 \rangle + \langle \dot{h}_\times^2 \rangle \) is proportional to the energy density and \( \int_0^\infty A_{\dot{h}}(k) \, dk \) is proportional to the polarized energy density. The \( A_{\dot{h}}(k) \) spectra are not to be confused with the magnetic vector potential \( A(x, t) \). The corresponding GW energy spectra are noted by
\[ E_{\text{GW}}(k) = (c^2/32\pi G) S_{\dot{h}}(k), \] (122)
\[ H_{\text{GW}}(k) = (c^2/32\pi G) A_{\dot{h}}(k). \] (123)

We also define spectra for the metric tensor perturbation,
\[ S_{h}(k) = \int_{4\pi} \left( |\tilde{h}_+|^2 + |\tilde{h}_\times|^2 \right) k^2 \, d\Omega_k, \] (124)
\[ A_{h}(k) = \int_{4\pi} 2 \text{Im} \left( \tilde{h}_+ \tilde{h}_\times^* \right) k^2 \, d\Omega_k, \] (125)
which are normalized such that \( \int_0^\infty S_{h}(k) \, dk = h_{\text{rms}}^2 \) is the mean squared metric tensor perturbation.
7 Troubleshooting / Frequently Asked Questions

7.1 Download and setup

7.1.1 Download forbidden


7.1.2 When sourcing the ‘sourceme.sh’/‘sourceme.csh’ file or running pc_setups, I get error messages from the shell, like ‘if: Expression Syntax.’ or ‘set: Variable name must begin with a letter.’

A: This sounds like a buggy shell setup, either by yourself or your system administrator — or a shell that is even more idiosyncratic than the ones we have been working with.

To better diagnose the problem, collect the following information before filing a bug report to us:

1. `uname -a`
2. `/bin/csh -v`
3. `echo $version`
4. `echo $SHELL`
5. `ps -p $$`
6. If you have problems while sourcing the ‘sourceme’ script,
   
   (a) unset the PENCIL_HOME variable:
   
   **for csh and similar:** unsetenv PENCIL_HOME
   **for bash and similar:** unexport PENCIL_HOME; unset PENCIL_HOME

   (b) switch your shell in verbose mode,
   
   **for csh and similar:** set verbose; set echo
   **for bash and similar:** set -v; set -x

   then source again.

7. If you have problems with pc_setups, run it with csh in verbose mode:

   `/bin/csh -v -x $PENCIL_HOME/bin/pc_setupsrc`
7.2 Compilation

7.2.1 Error: ‘relocation truncated to fit’

If you get errors while compiling and linking that are similar to:

density.f90:(.text+0x5e0): relocation truncated to fit: R_X86_64_PC32
against symbol ‘cdata_mp_m_’ defined in COMMON section in cdata.o
density.f90:(.text+0x644): additional relocation overflows omitted from the
output
make[2]: *** [start.x] Error 1

A: Your setup is probably too large to fit a ‘normal’ memory model. Please choose a
‘medium’ or ‘large’ memory model by adding one of these compiler options to your con-
figuration: ‘-mcmodel=medium’ or ‘-mcmodel=large’. See Sect. 5.1 for configuration details. Alternatively, if you use pc_build, you may simply add the respective extension:

pc_build -f GNU-GCC_MPI,GNU-GCC_medium

or for the Intel compiler and a ‘large’ memory model you would use:

pc_build -f Intel_MPI,Intel_large

7.2.2 Linker can’t find the syscalls functions:

ld: 0711-317 ERROR: Undefined symbol: .is_nan_c
ld: 0711-317 ERROR: Undefined symbol: .sizeof_real_c
ld: 0711-317 ERROR: Undefined symbol: .system_c
ld: 0711-317 ERROR: Undefined symbol: .get_env_var_c
ld: 0711-317 ERROR: Undefined symbol: .get_pid_c
ld: 0711-317 ERROR: Undefined symbol: .file_size_c

A: The Pencil Code needs a working combination of a Fortran- and a C-compiler. If this
is not correctly set up, usually the linker won’t find the functions inside the syscalls
module. If that happens, either the combination of C- and Fortran-compiler is inappro-
priate (e.g. ifort needs icc), or the compiler needs additional flags, like g95 might need
the option ‘-fno-second-underscore’ and xlf might need the option ‘-qextname’. Please
refer to Sect. 5.2 Table 1.

7.2.3 Make gives the following error now:

PGF90-S-0017-Unable to open include file: chemistry.h (nochemistry.f90: 43)
  0 inform, 0 warnings, 1 severes, 0 fatal for chemistry

Line 43 of the nochemistry routine, only has 'contains'.

A: This is because somebody added a new module (together with a corresponding nomod-
ule.f90 and a module.h file (chemistry in this case). These files didn’t exist before, so you
need to say:

pc_setupsirc
If this does not help, say first make clean and then pc_setupsrcc.

7.2.4 How do I compile the PENCIL CODE with the Intel (ifc) compiler under Linux?

A: The PENCIL CODE should compile successfully with ifc 6.x, ifc 7.0, sufficiently recent versions of ifc 7.1 (you should get the latest version; if yours is too old, you will typically get an ‘internal compiler error’ during compilation of ‘src/hydro.f90’), as well as with recent versions of ifort 8.1 (8.0 may also work).


On many current (as of November 2003) Linux systems, there is a mismatch between the glibc versions used by the compiler and the linker. To work around this, use the following flag for compiling

```
FC=ifc -i_dynamic
```

and set the environment variable

```
LD_ASSUME_KERNEL=2.4.1; export LD_ASSUME_KERNEL
```

or

```
setenv LD_ASSUME_KERNEL 2.4.1
```

This has solved the problems e.g. on a system with glibc-2.3.2 and kernel 2.4.22.

Thanks to Leonardo J. Milano ([http://udel.edu/~lmilano/](http://udel.edu/~lmilano/)) for part of this info.

7.2.5 I keep getting segmentation faults with ‘start.x’ when compiling with ifort 8.0

A: There was/is a number of issues with ifort 8.0. Make sure you have the latest patches applied to the compiler. A number of things to consider or try are:

1. Compile with the the ‘-static -nothreads’ flags.
2. Set your stacksize to a large value (but a far too large value may be problematic, too), e.g.
   
   ```
   limit stacksize 256m
   ulimit -s 256000
   
   3. Set the environment variable KMP_STACKSIZE to a large value (like 100M)
   

7.2.6 When compiling with MPI on a Linux system, the linker complains:

```plaintext
mpicomm.o: In function ‘mpicomm_mpicomm_init_’:
mpicomm.o(.text+0x36): undefined reference to ‘mpi_init_’
mpicomm.o(.text+0x55): undefined reference to ‘mpi_comm_size_’
mpicomm.o(.text+0x6f): undefined reference to ‘mpi_comm_rank_’
[...]
```
A: This is the infamous underscore problem. Your MPI libraries have been compiled with G77 without the option `-fno-second-underscore`, which makes the MPI symbol names incompatible with other Fortran compilers.

As a workaround, use

\[ \text{MPICOMM = mpicomm} \_ \]

in `Makefile.local`. Or, even better, you can set this globally (for the given computer) by inserting that line into the file `~/.adapt-mkfile.inc` (see perldoc adapt-mkfile for more details).

7.2.7 Compilation stops with the cryptic error message:

```bash
f95 -O3 -u -c .f90.f90
Error: Could not open sourcefile .f90.f90
compilation aborted for .f90.f90 (code 1)
make[1]: *** [.f90.o] Error 1
```

What is the problem?

A: There are two possibilities:

1. One of the variables for make has not been set, so make expands it to the empty string. Most probably you forgot to specify a module in `src/Makefile.local`. One possibility is that you have upgraded from an older version of the code that did not have some of the modules the new version has.

   Compare your `src/Makefile.local` to one of the examples that work.

2. One of the variables for make has a space appended to it, e.g. if you use the line

   \[ \text{MPICOMM = mpicomm} \_ \]

   (see §7.2.6) with a trailing blank, you will encounter this error message. Remove the blank. This problem can also occur if you added a new module (and have an empty space after the module name in `src/Makefile.src`, i.e. `CHIRAL=nochiral`), in which case the compiler will talk about “circular dependence” for the file `nochiral`.

7.2.8 The code doesn’t compile,

...there is a problem with mvar:

```bash
make start.x run.x
f95 -O3 -u -c cdata.f90
Error: cdata.f90, line 71: Implicit type for MVAR
      detected at MVAR@)
[f95 terminated - errors found by pass 1]
make[1]: *** [cdata.o] Error 2
```

A: Check and make sure that `mkcparam` (directory `$PENCIL_HOME/bin`) is in your path. If this doesn’t help, there may be an empty `cparam.inc` file in your `src` directory. Remove
‘cparam.inc’ and try again (Note that ‘cparam.inc’ is automatically generated from the ‘Makefile’).

7.2.9 Some samples don’t even compile,

as you can see on the web, http://www.nordita.org/software/pencil-code/tests.html

samples/helical-MHDturb:

  Compiling... not ok:
  make start.x run.x read_videos.x
make[1]: Entering directory `/home/dobler/f90/pencil-code/samples/helical-MHDturb/src'
/usr/lib/lam/bin/mpif95 -O3 -c initcond.f90
/usr/lib/lam/bin/mpif95 -O3 -c density.f90

use Gravity, only: gravz, nu_epicycle

Error 208 at (467:density.f90) : No such entity in the module
Error 355 : In procedure INIT_LNRHO variable NU_EPICYCLE has not been given a type
Error 355 : In procedure POLYTROPIC_LNRHO_DISC variable NU_EPICYCLE has not been given
3 Errors
compilation aborted for density.f90 (code 1)
make[1]: *** [density.o] Error 1
make[1]: Leaving directory `/home/dobler/f90/pencil-code/samples/helical-MHDturb/src'
make: *** [code] Error 2

A: Somebody may have checked in something without having run auto-test beforehand. The problem here is that something has been added in one module, but not in the corresponding no-module. You can of course check with svn who it was...

7.2.10 Internal compiler error with Compaq/Dec F90

The Dec Fortran optimizer has occasional problems with ‘nompicomm.f90’:

  make start.x run.x read_videos.x
  f90 -fast -O3 -tune ev6 -arch ev6 -c cparam.f90
[...]
  f90 -fast -O3 -tune ev6 -arch ev6 -c nompicomm.f90
  otal vm 2755568  otal vm 2765296  otal vm 2775024
  otal vm 2784752  otal...
Assertion failure: Compiler internal error - please submit problem r...
  GEM ASSERTION, Compiler internal error - please submit problem report
Fatal error in: /usr/lib/cmplrs/fort90_540/decfort90 Terminated
*** Exit 3
Stop.
*** Exit 1
Stop.

A: The occurrence of this problem depends upon the grid size; and the problem never seems to occur with ‘mpicomm.f90’, except when ncpus=1. The problem can be avoided by switching off the loop transformation optimization (part of the ‘-O3’ optimization), via:

#OPTFLAGS=-fast -O3 -notransform_loops
This is currently the default compiler setting in `Makefile`, although it has a measurable performance impact (some 8% slowdown).

### 7.2.11 Assertion failure under SunOS

Under SunOS, I get an error message like

```
user@sun> f90 -c param_io.f90
Assertion failed: at_handle_table[at_idx].tag == VAR_TAG,
    file ../srcfw/FWcvrt.c, line 4018
f90: Fatal error in f90comp: Abort
```

**A:** This is a compiler bug that we find at least with Sun's WorkShop Compiler version '5.0 00/05/17 FORTRAN 90 2.0 Patch 107356-05'. Upgrade the compiler version (and possibly also the operating system): we find that the code compiles and works with version 'Sun WorkShop 6 update 2 Fortran 95 6.2 Patch 111690-05 2002/01/17' under SunOS version '5.8 Generic 108528-11'.

### 7.2.12 After some dirty tricks I got pencil code to compile with MPI, ...

> Before that i installed lam-7.1.4 from source.

Goodness gracious me, you shouldn't have to compile your own MPI library.

**A:** Then don't use the old LAM-MPI. It is long superseded by open-mpi now. Open-mpi doesn't need a daemon to be running. I am using the version that ships with Ubuntu (e.g. 9.04):

```
frenesi:~> aptitude -w 210 search openmpi | grep '^i'
```

```
i libopenmpi-dev - high performance message passing library -- header files
i A libopenmpi1 - high performance message passing library -- shared library
i openmpi-bin - high performance message passing library -- binaries
i openmpi-common - high performance message passing library -- common files
i openmpi-doc - high performance message passing library -- man pages
```

Install that and keep your configuration (Makefile.src and getconf.csh) close to that for 'frenesi' or 'norlx50'. That should work.

### 7.2.13 Error: Symbol 'mpi_comm_world' at (1) has no IMPLICIT type

I installed the pencil code on Ubuntu system and tested "run.csh" in ...\samples\conv-slab. Here the code worked pretty well. Nevertheless, running (auto-test), I found there are some errors.

The messages are,

```
Error: Symbol 'mpi_comm_world' at (1) has no IMPLICIT type
Fatal Error: Error count reached limit of 25.
make[2]: *** [mpicomm_double.o] Error 1
```
make[2]: Leaving directory
`/home/pkiwan/Desktop/pencil-code/samples/2d-tests/selfgravitating-shearwave/src`
make[1]: *** [code] Error 2
make[1]: Leaving directory
`/home/pkiwan/Desktop/pencil-code/samples/2d-tests/selfgravitating-shearwave/src`
make: *** [default] Error 2

Finally, ### auto-test failed ###

Will it be OK? Or, how can I fix this?

A: Thanks for letting me know about the status, and congratulations on your progress! Those tests that fail are those that use MPI. If your machine is a dual or multi core machine, you could run faster by running under MPI. But this is probably not crucial for you at this point. (I just noticed that there is a ToDo listed in the auto-test command to implement the option not to run the MPI tests, but this hasn't been done yet. So I guess you can start with the science next.

7.2.14 Error: Can’t open included file ‘mpif.h’

It always worked, but now, after some systems upgrade, I get

```
gfortran -O3 -o mpicomm.o -c mpicomm.f90
Error: Can’t open included file ’mpif.h’
```

When I say `locate mpif.h` I only get things like

```
/scratch/ntest/1.2.7p1-intel/include/mpif.h
```

But since I use `FC=mpif90` I thought I don’t need to worry.

A: Since you use `FC=mpif90` there must definitely be something wrong with their setup. Try `mpif90 -showme` or `mpif90 -show`; the `-I` option should say where it looks for `mpif.h`. If those directories don’t exist, it’s no wonder that it doesn’t work, and it is time to complain.

7.3 Pencil check

7.3.1 The pencil check complains for no reason.

A: The pencil check only complains for a reason.

7.3.2 The pencil check reports MISSING PENCILS and quits

A: This could point to a serious problem in the code. Check where the missing pencil is used in the code. Request the right pencils, likely based on input parameters, by adapting one or more of the `pencil_criteria_MODULE` subroutines.
7.3.3 The pencil check reports unnecessary pencils

The pencil check reports possible overcalculation... pencil rho (43) is requested, but does not appear to be required!

A: Such warnings show that your simulation is possibly running too slowly because it is calculating pencils that are not actually needed. Check in the code where the unnecessary pencils are used and adapt one or more of the pencil_criteria_MODULE subroutines to request pencils only when they are actually needed.

7.3.4 The pencil check reports that most or all pencils are missing

A: This is typically a thing that can happen when testing new code development for the first time. It is usually an indication that the reference df changes every time you call pde. Check whether any newly implemented subroutines or functionality has a “memory”, i.e. if calling the subroutine twice with the same f gives different output df.

7.3.5 Running the pencil check triggers mathematical errors in the code

A: The pencil check puts random numbers in f before checking the dependence of df on the chosen set of pencils. Sometimes these random numbers are inconsistent with the physics and cause errors. In that case you can set lrandom_f_pencil_check=F in &run_-pars in ‘run.in’. The initial condition may contain many idealized states (zeros or ones) which then do not trigger pencil check errors when lrandom_f_pencil_check=F, even if pencils are missing. But it does prevent mathematical inconsistencies.

7.3.6 The pencil check still complains

A: Then you need to look into the how the code and the pencil check operate. Reduce the problem in size and dimensions to find the smallest problem that makes the pencil check fail (e.g. 1x1x8 grid points). At the line of ‘pencil_check.f90’ when a difference is found between df_ref and df, add some debug lines telling you which variable is inconsistent and in what place. Often you will be surprised that the pencil check has correctly found a problem in the simulation.

7.3.7 The pencil check is annoying so I turned it off

A: Then you are taking a major risk. If one or more pencils are not calculated properly, then the results will be wrong.

7.4 Running

7.4.1 Why does ‘start.x’ / ‘start.csh’ write data with periodic boundary conditions?

A: Because you are setting the boundary conditions in ‘run.in’, not in ‘start.in’; see Sect. 5.16.1 There is nothing wrong with the initial data — the ghost-zone values will
be re-calculated during the very first time step.

### 7.4.2 csh problem?

**Q:** On some rare occasions we have problems with csh not being supported on other machines. (We hope to fix this by contacting the responsible person, but may not be that trivial today!) Oliver says this is a well known bug of some years ago, etc. But maybe in the long run it would be good to avoid csh.

**A:** These occasions will become increasingly frequent, and eventually for some architectures, there may not even be a csh variant that can be installed.

We never pushed people to use `pc_run` and friends (and to report corresponding bugs and get them fixed), but if we don’t spend a bit of effort (or annoy users) now, we create a future emergency, where someone needs to run on some machine, but there is no csh and he or she just gets stuck.

We don’t have that many csh files, and for years now it should be possible to compile run without csh (using `bin/pc_run`) — except that people still fall back on the old way of doing things. This is both cause and consequence of the ‘new’ way not being tested that much, at least for the corner cases like ‘rerun’, ‘newdir’, ‘scratch_dir’.

### 7.4.3 ‘run.csh’ doesn’t work:

```
Invalid character ' ' in NAMELIST input
Program terminated by fatal I/O error
Abort
```

**A:** The string array for the boundary condition, e.g. `bcx` or `bcz` is too long. Make sure it has exactly as many elements as `nvar` is big.

### 7.4.4 Code crashes after restarting

> > removing `mu_r` from the namelist just ‘like that’ makes the code
> > backwards incompatible.
> >
> > That means that we can never get rid of a parameter in `start.in` once we have introduced it, right?

**A:** In the current implementation, without a corresponding cleaning procedure, unfortunately yes.

Of course, this does not affect users’ private changes outside the central svn tree.

### 7.4.5 auto-test gone mad...?

**Q:** Have you ever seen this before:

```
giga01:/home/pg/n7026413/cvs-src/pencil-code/samples/conv-slab> auto-test
```

/home/pg/n7026413/cvs-src/pencil-code/samples/conv-slab:
  Compiling.. ok
  No data directory; generating data -> /var/tmp/pencil-tmp-25318
  Starting.. ok
  Running.. ok
  Validating results.. Malformed UTF-8 character (unexpected continuation byte 0x80, with no preceding start byte) in split at
  /home/pg/n7026413/cvs-src/pencil-code/bin/auto-testline 263.
  Malformed UTF-8 character (unexpected continuation byte 0x80, with no preceding start byte) in split at
  /home/pg/n7026413/cvs-src/pencil-code/bin/auto-testline 263.

A: You are running on a RedHat 8 or 9 system, right?
Set LANG=POSIX in your shell's startup script and life will be much better.

7.4.6 Can I restart with a different number of cpus?

Q: I am running a simulation of nonhelical turbulence on the cluster using MPI. Suppose if I am running a $128^3$ simulation on 32 cpus/cores i.e.

```plaintext
integer, parameter :: ncpus=32, nprocy=2, nprocz=ncpus/nprocy, nprocx=1
integer, parameter :: nxgrid=128, nygrid=nxgrid, nzgrid=nxgrid
```

And I stop the run after a bit. Is there a way to resume this run with different number of cpus like this:

```plaintext
integer, parameter :: ncpus=16, nprocy=2, nprocz=ncpus/nprocy, nprocx=1
integer, parameter :: nxgrid=128, nygrid=nxgrid, nzgrid=nxgrid
```

I understand it has to be so in a new directory but making sure that the run starts from where I left it off in the previous directory.

A: The answer is no, if you use the standard distributed io. There is also parallel io, but I never used it. That would write the data in a single file, and then you could use the data for restart in another processor layout.

7.4.7 Can I restart with a different number of cpus?

Q: Is it right that once the simulation is resumed, pencil-code takes the last data from var.dat (which is the current snapshot of the fields)? If that is true, then, is it not possible to give that as the initial condition for the run in the second directory (with changed "ncpus")? Is there a mechanism already in place for that?

A: Yes, the code restarts from the last var.dat. It is written after a successful completion of the run, but it crashes or you hit a time-out, there will be a var.dat that is overwritten every isave timesteps. If the system stops during writing, some var.dat files may be corrupt or have the wrong time. In that case you could restart from a good VAR file, if you have one, using, e.g.,

```plaintext
restart-new-dir-VAR 46
```

where 46 is the number of your VAR file, i.e., VAR46 in this case. To restart in another directory, you say, from the old run directory,
Hope this helps. Look into pencil-code/bin/restart-new-dir to see what it is doing.

7.4.8 fft_xyz_parallel_3D: nygrid needs to be an integer multiple...

Q: I just got an:

```
fft_xyz_parallel_3D: nygrid needs to be an integer multiple of nprocy*nprocz
```

In my case, nygrid=2048, nprocy=32, and nprocz=128, so nprocy*nprocz=4096. In other words, 2048 needs to be a multiple of 4096. But isn't this the case then?

A: No, because 2048 = 0.5 * 4096 and 0.5 is not an integer. Maybe try either setting nprocz=64 or nprocy=64. You could compensate the change of ncpus with the \( x \)-direction. For 2048\(^3\) simulations, nprocy=32 and nprocz=64 would be good. A list of good meshes is given in Table [4].

7.4.9 Unit-agnostic calculations?

Q: The manual speaks about unit-agnostic calculations, stating that one may choose to interpret the results in any (consistent) units, depending on the problem that is solved at hand. So, for example, if I chose to run the ‘2d-tests/battery_term’ simulation for an arbitrary number of time-steps and then choose to examine the diagnostics, am I correct in assuming the following:

1) \([Brms]\) = Gauss (as output by unit_magnetic, before the run begins)
2) \([t]\) = s (since the default unit system is left as CGS)
3) \([urms]\) = cm/s (again, as output by unit_velocity, before the run begins)
4) and etc. for the units of the other diagnostics

A: Detailed correspondence on this item can be found on: https://groups.google.com/forum/?fromgroups#!topic/pencil-code-discuss/zek-uYNbgXI

There is also working material on unit systems under http://www.nordita.org/~brandenb/teach/PencilCode/MixedTopics.html with a link to http://www.nordita.org/~brandenb/teach/PencilCode/material/AlfvenWave_SIunits/

Below is a pedagogical response from Wlad Lyra:

In the sample battery-term, the sound speed \( cs0=1 \) sets the unit of velocity. Together with the unit of length, that sets your unit of time. The unit of magnetic field follows from the unit of velocity, density, and your choice of magnetic permittivity, according to the definition of the Alfven velocity.

If you are assuming cgs, you are saying that your sound speed \( cs0=1 \) actually means \([U]=1\) cm/s. Your unit of length is equivalently 1 cm, and therefore the unit of time is \([t]= [L]/[U]=1\) s. The unit of density is \([\rho]=1\) g/cm\(^3\). Since in cgs \( vA=B/sqrt(4*pi * \rho) \), your unit of magnetic field is \([B]=[U]*sqrt([\rho] * 4*pi) \approx 3.5\) sqrt(g/cm) / s = 3.5 Gauss.

If instead you are assuming SI, you have \( cs0=1 \) assuming that means \([U]=1\) m/s and \( \rho0=1\) assuming that to mean \([\rho]=1\) kg/m\(^3\). Using \([L]=1\)
m, you have still \([t]=1\) s, but now what appears as \(B=1\) in your output is actually \([B] = \frac{[U]}{[\mu] \cdot \sqrt{[\rho] \cdot 4\pi}} = 1 \text{ m/s} \cdot \sqrt{4\pi \cdot 1 \times 10^{-7} \text{ N/A}^{-2} \cdot 1 \text{ kg/m}^3} \approx 0.0011210 \text{ kg/(s}^2\text{A)} \approx 11 \text{ Gauss.}

You can make it more interesting and use units relevant to the problem. Say you are at the photosphere of the Sun. You may want to use dimensionless \(c_0=1\) meaning a sound speed of 10 km/s. Your appropriate length can be a megameter. Now your time unit is \([t]=\frac{[L]}{[U]} = 1 \times 10^3 \text{ km/} \cdot 10 \text{ km/s} = 10^2 \text{ s, i.e., roughly 1.5 minute. For density, assume \(\rho=2 \times 10^{-4} \text{ kg/m}^3\), typical of the solar photosphere. Your unit of magnetic field is therefore \([B] = \frac{[U]}{[\mu] \cdot \sqrt{[\rho] \cdot 4\pi}} = 1 \times 10^6 \text{ cm/s} \cdot \sqrt{4\pi \cdot 2 \times 10^{-7} \text{ g/cm}^3} \approx 1585.33 \text{ Gauss.}

Notice that for \(\mu_0=1\) and \(\rho_0=1\) you simply have \(v_A=B\). Then you can conveniently set the field strength by your choice of plasma beta (= \(2 \times c_0^2/v_A^2\)). There’s a reason why we like dimensionless quantities!

7.5 Visualization

7.5.1 ‘start.pro’ doesn’t work:

\begin{verbatim}
Reading grid.dat..
Reading param.nml.
\% Expression must be a structure in this context: PAR.
\% Execution halted at: \$MAIN\$
/home/brandenb/pencil-code/runs/forced/hel1/../../../idl/start.pro
\end{verbatim}

A: You don’t have the subdirectory ‘data’ in your IDL variable \!’path. Make sure you source ‘sourceme.csh’/‘sourceme.sh’ or set a sufficient IDL path otherwise.

7.5.2 ‘start.pro’ doesn’t work:

Isn’t there some clever (or even trivial) way that one can avoid the annoying error messages that one gets, when running e.g. ”.r rall” after a new variable has been introduced in ”idl/varcontent.pro”? Ever so often there’s a new variable that can’t be found in my param2.nml – this time it was IECR, IGG, and ILNTT that I had to circumvent…

A: The simplest solution is to invoke ‘NOERASE’, i.e. say

```
touch NOERASE
start.csh
```
or, alternatively, \texttt{start.run.csh}. What it does is that it reruns src/start.x with a new version of the code; this then produces all the necessary auxiliary files, but it doesn’t overwrite or erase the ‘var.dat’ and other ‘VAR’ and ‘slice’ files.

7.5.3 Something about tag name undefined:

Q: In one of my older run directories I can’t read the data with idl anymore. What should I do? Is says something like
Reading param.nml..
% Tag name LEQUIDIST is undefined for structure <Anonymous>.
% Execution halted at: $MAIN$ 182
 /people/disk2/brandenb/pencil-code/idl/start.pro

A: Go into `data/param.nml` and add `, LEQUIDIST=T` anywhere in the file (but before the last slash).

### 7.5.4 Something INC in start.pro

Q: start doesn’t even work:

% Compiled module: $MAIN$.
nname= 11
Reading grid.dat..
Reading param.nml..
Can’t locate Namelist.pm in INC (INC contains: /etc/perl /usr/local/lib/perl/5.8.4 /usr/local/share/perl/5.8.4
BEGIN failed--compilation aborted at /home/brandenb/pencil-code/bin/nl2idl line 49.

A: Go into `$PENCIL_HOME` and say svn up sourceme.csh and/or svn up sourceme.sh.
(They were just out of date.)

### 7.5.5 nl2idl problem when reading param2.nml

Q: Does anybody encounter a backward problem with nl2idl? The file param*.nml files are checked in under `pencil-code/axel/couette/SStrat128a_mu0.20_g2` and the problem is below.

```
    at /people/disk2/brandenb/pencil-code/bin/nl2idl line 120
HCOND0= 0.0,HCOND1= 1.000000,HCOND2= 1.000000,WIDTHSS= 1.192093E-06,MPOLY0=
^------ HERE
    at /people/disk2/brandenb/pencil-code/bin/nl2idl line 120
```

A: The problem is the stupid ifc compiler writing the following into the namelist file:

```
COOLING_PROFILE=’gaussian ’,COOLTYPE=’Temp
 ’,COOL= 0.0,CS2COOL= 0.0,RCOOL= 1.000000,WCool= 0.1000000,FBOT= 0.0,CHI_T= 0.0
```

If you add a comma after the closing quote:

```
COOLING_PROFILE=’gaussian ’,COOLTYPE=’Temp
 ’,COOL= 0.0,CS2COOL= 0.0,RCOOL= 1.000000,WCool= 0.1000000,FBOT= 0.0,CHI_T= 0.0
```

things will work.

Note that ifc cannot even itself read what it is writing here, so if this happened to occur in param.nml, the code would require manual intervention after each start.csh.

### 7.5.6 Spurious dots in the time series file

Q: Wolfgang, you explained it to me once, but I forget. How can one remove spurious dots after the timestep number if the time format overflows?
A: I don’t know whether it exists anywhere, but it’s easy. In Perl you’d say

```perl
perl -pe 's/^\s*([-0-9]+)\.([-0-9eEdD])/$1 $2/g'
```

and in sed (but that’s harder to read)

```sed
s/^\s*([-0-9]+)\.([-0-9eEdD])/$1 $2/g'
```

7.5.7 Problems with `pc_varcontent.pro`

Q:

% Subscript range values of the form low:high must be >= 0, < size, with low <= high: VARCONTENT.
% Error occurred at: PC_VARCONTENT 391
/home/brandenb/pencil-code/idl/read/pc_varcontent.pro
% PC_READ_VAR 318
/home/brandenb/pencil-code/idl/read/pc_read_var.pro
% $MAIN$

A: Make sure you don’t have any unused items in your `src/cparam.local` such as

```
! MAUX CONTRIBUTION 3
! COMMUNICATED AUXILIARIES 3
```

They would leave gaps in the counting of entries in your `data/index.pro` file.

7.6 General questions

7.6.1 “Installation” procedure

Why don’t you use GNU `autoconf/automake` for installation of the PENCIL CODE?

A: What do you mean by “installation”? Unlike the applications that normally use `autoconf`, the Pencil Code is neither a binary executable, nor a library that you compile once and then dump somewhere in the system tree. `Autoconf` is the right tool for these applications, but not for numerical codes, where the typical compilation and usage pattern is very different:

You have different directories with different ‘`Makefile.local`’ settings, recompile after introducing that shiny new term in your equations, etc. Moreover, you want to sometimes switch to a different compiler (but just for that run directory) or another `MPI` implementation. Our `adapt-mkfile` approach gives you this flexibility in a reasonably convenient way, while doing the same thing with `autoconf` would be using that system against most of its design principles.

Besides, it would really get on my (WD’s) nerves if I had to wait two minutes for `autoconf` to finish before I can start compiling (or maybe 5–10 minutes if I worked on a NEC machine...).

Finally, if you have ever tried to figure out what a ‘`configure`’ script does, you will appreciate a comprehensible configuration system.
7.6.2 Small numbers in the code

What is actually the difference between epsi, tini and tiny?

A:

F90 has two functions epsilon() and tiny(), with

\[
\text{epsilon}(x) = 1.1920929\times 10^{-7} \\
\text{tiny}(x) = 1.1754944\times 10^{-38} \\
\text{(and then there is huge}(x) = 3.4028235\times 10^{38})
\]

for a single-precision number x.

epsilon(x) is the smallest number that satisfies

\[1+\text{epsilon}(1.) /= 1,\]

while tiny(x) is the smallest number that can be represented without precision loss.

In the code we have variants hereof,

\[
\begin{align*}
\text{epsi} &= 5 \times \text{epsilon}(1.0) \\
\text{tini} &= 5 \times \text{tiny}(1.0) \\
\text{huge1} &= 0.2 \times \text{huge}(1.0)
\end{align*}
\]

that have added safety margins, so we don’t have to think about doing things like 1/tini.

So in sub.f90,

\[- \quad \text{evr = evr} /= \text{spread(r_mn+epsi,2,3)}\]

did (minimally) affect the result for r_mn=0(1), while the correct version

\[+ \quad \text{evr = evr} /= \text{spread(r_mn+tini,2,3)}\]

only avoids overflow.

7.6.3 Why do we need a /lphysics/namelist in the first place?

Wolfgang answered on 29 July 2010: “cdata.f90’ has the explanation”

! Constant ‘parameters’ cannot occur in namelists, so in order to get the
! now constant module logicals into the lphysics name list...
! We have some proxies that are used to initialize private local variables
! called lhydro etc, in the lphysics namelist!

So the situation is this: we want to write parameters like ldensity to param.nml so
IDL (and potentially octave, python, etc.) can know whether density was on or not. To
avoid confusion, we want them to have exactly their original names. But we cannot
assemble the original ldensity etc. constants in a namelist, so we have to define a local
ldensity variable. And to provide it with the value of the original cdata.ldensity, we need
to transfer the value via ldensity.var. That’s pretty scary, although it seems to work
fine. I can track the code back to the big eos_merger commit, so it may originate from
that branch. One obvious problem is that you have to add code in a number of places
(the ldensity \rightarrow ldensity.var assignment and the local definition of ldensity) to really
get what you need. And when adding a new boolean of that sort to ‘cdata.f90’, you may
not even have a clue that you need all the other voodoo.
There may be a cleaner solution involving generated code. Maybe something like

```
logical :: ldensity ! INCLUDE_IN_LPHYSICS
```

could later generate code (in some param_io_extra.inc file) that looks like this:

```
write(unit, *) 'ldensity = ', ldensity
```
i.e. we can manually write in namelist format. But maybe there are even simpler solutions?

### 7.6.4 Can I run the code on a Mac?

**A**: Macs work well for Linux stuff, except that the file structure is slightly different. Problems when following Linux installs can usually be traced to the PATH. For general reference, if you need to set an environment variable for an entire OS-X login session, google environment.plist. That won’t be needed here.

For a Mac install, the following should work:

a) Install Dev Tools (an optional install on the MacOS install disks). Unfortunately, last time I checked the svn version that comes with DevTools is obsolete. So:

b) Install MacPorts (download from web). Note that MacPorts installs to a non-standard location, and will need to be sourced. The installation normally drops an appropriate line in .profile. If it does so, make sure that that line gets sourced. Otherwise

```
export PATH=/opt/local/bin:/opt/local/sbin:$PATH
export MANPATH=/opt/local/share/man:$MANPATH
```

c) Install g95 (download from web). Make sure it is linked in /bin.

d) execute macports svn install
e) download the pencil-code and enjoy.

Note: the above way to get svn works. It takes a while however, so there are certainly faster ways out there. If you already have a non-obsolete svn version, use that instead.

### 7.6.5 Pencil Code discussion forum

Do I just need to send an email somewhere to subscribe or what?

**A**: The answer is yes; just go to:

```
http://groups.google.com/group/pencil-code-discuss
```

### 7.6.6 The manual

It would be a good idea to add this useful information in the manual, no?

**A**: When you have added new stuff to the code, don’t forget to mention this in the ‘pencil-code/doc/manual.tex’ file.
Again, the answer is yes; just go to:

```bash
  cd pencil-code/doc/
  vi manual.tex
  svn ci -m "explanations about a new module in the code"
```
Part II

Programming the PENCIL Code

All developers are supposed to have an up-to-date entry in the file ‘pencil-code/license/developers.txt’ so that they can be contacted in case a code change breaks an auto-test or other code functionality.

Several PENCIL Code committers have done several hundred check-ins, but many of the currently 68 registered people on the repository have hardly done anything. To put a number to this, one can define an $h$ index, which gives the number of users, who have done at least as many as that number of check-ins. This $h$ index is currently 32, i.e., 32 users have done at least 32 check-ins; see Figure 7.

![Figure 7: The $h$ index of PENCIL Code check-ins.](image)

The PENCIL Code has expanded approximately linearly in the number of lines of code and the number of subroutines (Fig. 8). The increase in the functionality of the code is documented by the rise in the number of sample problems (Fig. 9). It is important to monitor the performance of the code as well. Figure 10 shows that for most of the runs the run time has not changed much.

Before making changes to the code, it is important that you verify that you can run the pc_auto-test successfully. Don’t do this when you have already modified the code, because then you cannot be sure that any problems are caused by your changes, or because it wouldn’t have worked anyway. Also, keep in mind that the code is public, so your changes should make sense from a broader perspective and should not only be intended for yourself. Regarding more general aspects about coding standards see Sect. B.2.

In order to keep the development of the code going, it is important that the users are able to understand and modify (program!) the code. In this section we explain first how
to orient yourself in the code and to understand what is in it, and then to modify it according to your needs.

The Pencil Code check-ins occur regularly all the time. By the Pencil Code User Meeting 2010 we have arrived at a revision number of 15,000. In February 2017, the number of check-ins has risen to 26,804; see [https://github.com/pencil-code/pencil-code](https://github.com/pencil-code/pencil-code). Major code changes are nowadays being discussed by the Pencil Code Steering Committee ([https://www.nordita.org/~brandenb/pencil-code/PCSC/](https://www.nordita.org/~brandenb/pencil-code/PCSC/)). The increase of the revision number with time is depicted in Figure 11. The number of Pencil Code developers increases too (Figure 12), but the really active ones are getting rare. This may indicate that new users can produce new science with the code as it is, but it may also indicate that it is getting harder to understand the code. How to understand the code will be discussed in the next section.

**Figure 8:** Number of lines of code and the number of subroutines since the end of 2001. The jump in the Summer of 2005 was the moment when the developments on the side branch (eos branch) were merged with the main trunk of the code. Note the approximately linear scaling with time.

**Figure 9:** Number of tests in the sample directory that are used in the nightly auto tests. Note again the approximately linear scaling with time.
Figure 10: Run time of the daily auto-tests since August 17, 2008. For most of the runs the run time has not changed much. The occasional spikes are the results of additional load on the machine.
Figure 11: Number of check-ins since 2002. Note again the linear increase with time, although in the last part of the time series there is a notable speed-up.

Figure 12: Check-ins since 2002 per user. Users with more than 100 check-ins are color coded.
8 Understanding the code

Understanding the code means looking through the code. This is not normally done by just printing out the entire code, but by searching your way through the code in order to address your questions. The general concept will be illustrated here with an example.

8.1 Example: how is the continuity equation being solved?

All the physics modules are solved in the routine pde, which is located in the file and module ‘Equ’. Somewhere in the pde subroutine you find the line

```fortran
    call dlnrho_dt(f,df,p)
```

This means that here the part belonging to $\partial \ln \rho / \partial t$ is being assembled. Using the grep command you will find that this routine is located in the module density, so look in there and try to understand the pieces in this routine. We quickly arrive at the following crucial part of code,

```fortran
! Continuity equation.
!
    if (lcontinuity_gas) then
        if (ldensity_nolog) then
            df(l1:l2,m,n,irho) = df(l1:l2,m,n,irho) - p%ugrho - p%rho*p%divu
        else
            df(l1:l2,m,n,ilnrho) = df(l1:l2,m,n,ilnrho) - p%uglnrho - p%divu
        endif
    endif
```

where, depending on some logica that tell you whether the continuity equation should indeed be solved and whether we do want to solve for the logarithmic density and not the actual density, the correct right hand side is being assembled. Note that all these routines always only add to the existing $df(l1:l2,m,n,ilnrho)$ array and never reset it. Resetting $df$ is only done by the time-stepping routine. Next, the pieces $p%uglnrho$ and $p%divu$ are being subtracted. These are pencils that are organized in the structure with the name $p$. The meaning of their names is obvious: $uglnrho$ refers to $u \cdot \nabla \ln \rho$ and $divu$ refers to $\nabla \cdot u$. In the subroutine pencil_criteria_density you find under which conditions these pencils are requested. Using grep, you also find where they are calculated. For example $p%uglnrho$ is calculated in ‘density.f90’; see

```fortran
    call u_dot_grad(f,ilnrho,p%glnrho,p%uu,p%uglnrho,UPWIND=lupw_lnrho)
```

So this is a call to a subroutine that calculates the $u \cdot \nabla$ operator, where there is the possibility of upwinding, but this is not the default. The piece $divu$ is calculated in ‘hydro.f90’ in the line

```fortran
! Calculate uiuj and divu, if requested.
!
    if (lpencil(i_uuij)) call gij(f,iuu,p%uij,1)
    if (lpencil(i_divu)) call div_mn(p%uij,p%divu,p%uu)
```
Note that the divergence calculation uses the velocity gradient matrix as input, so no new derivatives are recalculated. Again, using grep, you will find that this calculation and many other ones happen in the module and file 'sub.f90'. The various derivatives that enter here have been calculated using the \( gij \) routine, which calls the \( \text{der} \) routine, e.g., like so

\[
k1=k-1
\]
\[
do \ i=1,3
\]
\[
do \ j=1,3
\]
\[
\text{if (nder==1) then }
\]
\[
\text{call der}(f, k1+i, \text{tmp}, j)
\]

For all further details you just have to follow the trail. So if you want to know how the derivatives are calculated, you have to look in deriv.f90, and only here is it where the indices of the \( f \) array are being addressed.

If you are interested in magnetic fields, you have to look in the file 'magnetic.f90'. The right hand side of the equation is assembled in the routine

```fortran
subroutine daa_dt(f,df,p)
!
! Magnetic field evolution.
!
! Calculate \( \frac{dA}{dt}=uxB+3/2 \Omega_y x_{\text{dir}} -\eta \mu_0 J \). 
! For mean field calculations one can also add \( \frac{dA}{dt}=...+\alpha*bb+\delta*WXJ \). 
! Add \( jxb/rho \) to momentum equation. 
! Add \( \eta \mu_0 j2/rho \) to entropy equation. 
!
where the header tells you already a little bit of what comes below. It is also here where ohmic heating effects and other possible effects on other equations are included, e.g.

```fortran
if (lentropy .and. lohmic_heat) then
    df(l1:l2,m,n,iss) = df(l1:l2,m,n,iss) 
    + etatotal*mu0*p%j2*p%rho1*p%TT1
endif
```

We leave it at this and encourage the user to do similar inspection work on a number of other examples. If you think you find an error, file a ticket at [http://code.google.com/p/pencil-code/issues/list](http://code.google.com/p/pencil-code/issues/list). You can of course also repair it!
9 Adapting the code

9.1 The PENCIL CODE coding standard

As with any code longer than a few lines the appearance and layout of the source code is of the utmost importance. Well laid out code is more easy to read and understand and as such is less prone to errors.

A consistent coding style has evolved in the PENCIL CODE and we ask that those contributing try to be consistent for everybody’s benefit. In particular, it would be appreciated if those committing changes of existing code via svn follow the given coding style.

There are not terribly many rules and using existing code as a template is usually the easiest way to proceed. In short the most important rules are:

- tab characters do not occur anywhere in the code (in fact the use of tab character is an extension to the Fortran standard).
- Code in any delimited block, e.g. if statements, do loops, subroutines etc., is indented be precisely 2 spaces. E.g.

  ```fortran
  if (l_cylindrical) then
      call fatal_error(’del2fjv’,’del2fjv not implemented’)
  endif
  ```

- continuation lines (i.e. the continuation part of a logical line that is split using the & sign) are indented by 4 spaces. E.g. (note the difference from the previous example)

  ```fortran
  if (l_cylindrical) &
      call fatal_error(’del2fjv’,’del2fjv not implemented’)
  ```

- There is always one space separation between ’if’ and the criterion following in parenthesis:

  ```fortran
  if (ldensity_nolog) then
      rho=f(l1:l2,m,n,irho)
  endif
  ```

  This is wrong:

  ```fortran
  if(ldensity_nolog) then ! WRONG
      rho=f(l1:l2,m,n,irho)
  endif
  ```

- In general, try to follow common practice used elsewhere in the code. For example, in the code fragment above there are no empty spaces within the mathematical expressions programmed in the code. A unique convention helps in finding certain expressions and patterns in the code. However, empty spaces are often used after commas and semicolons, for examples in name lists.

- Relational operators are written with symbols (==, /=, <, <=, >, >=), not with characters (.eq., .ne., .lt., .le., .gt., .ge.).

- In general all comments are placed on their own lines with the ’!’ appearing in the first column.
• All subroutine/functions begin with a standard comment block describing what they do, when and by whom they were created and when and by whom any non-trivial modifications were made.

• Lines longer that 78 characters should be explicitly wrapped using the & character, unless there is a block of longer lines that can only be read easily when they are not wrapped. Always add one whitespace before the & character.

These and other issues are discussed in more depth and with examples in Appendix B, and in particular in Sect. B.2.

9.2 Adding new output diagnostics

With the implementation of new physics and the development of new procedures it will become necessary to monitor new diagnostic quantities that have not yet been implemented in the code. In the following, we describe the steps necessary to set up a new diagnostic variable.

This is nontrivial as, in order to keep latency effects low on multi-processor machines, the code minimizes the number of global reduction operations by assembling all quantities that need the maximum taken in \( f_{\text{max}} \), and those that need to be summed up over all processors (mostly for calculating mean quantities) in \( f_{\text{sum}} \) (see subroutine diagnostic in file ‘src/equ.f90’).

As a sample variable, let us consider \( jbm \) (the volume average \( \langle j \cdot B \rangle \)). Only the module magnetic will be affected, as you can see (the diagnostic quantity \( jbm \) is already implemented) with

```
unix> grep -i jbm src/*.f90
```

If we pretend for the sake of the exercise that no trace of \( jbm \) was in the code, and we were only now adding it, we would need to do the following

1. add the variable \( \text{idiag}_jbm \) to the module variables of Magnetic in both ‘magnetic.f90’ and ‘nomagnetic.f90’:

   ```
   integer :: idiag_jbm=0
   ```

   The variable \( \text{idiag}_jbm \) is needed for matching the position of \( jbm \) with the list of diagnostic variables specified in ‘print.in’.

2. in the subroutine daa_dt in ‘magnetic.f90’, declare and calculate the quantity \( jb \) (the average of which will be \( jbm \)), and call \( \text{sum}_mn\_name \)

   ```
   real, dimension (nx) :: jb  ! jj.BB
   [...] 
   if (l diagnos) then  ! only calculate if diagnostics is required
      if (idiag\_jbm/=0) then  ! anybody asked for jbm?
         call dot\_mn(jj,bb,jb)  ! assuming jj and bb are known
         call sum\_mn\_name(jb,i\_jbm)
      endif
   endif
   ```
3. in the subroutine `rprint_magnetic` in both `magnetic.f90`, add the following:

```fortran
! reset everything in case of RELOAD
! (this needs to be consistent with what is defined above!)
!
if (lreset) then  ! need to reset list of diagnostic variables?
    [...]
    idiag_jbm=0
    [...]
endif
!
! check for those quantities that we want to evaluate online
!
do iname=1,nname
    [...]
    call parse_name(iname,cname(iname),cform(iname),'jbm',idiag_jbm)
    [...]
enddo
![...]
!
! write column, i_XYZ, where our variable XYZ is stored
!
write(3,*) 'i_jbm=',idiag_jbm
![...]
```

4. in the subroutine `rprint_magnetic` in `nomagnetic.f90`, add the following (newer versions of the code may not require this any more):

```fortran
!
! write column, i_jbm, where our variable jbm is stored
! idl needs this even if everything is zero
!
write(3,*) 'i_jbm=',idiag_jbm
![...]
```

5. and don’t forget to add your new variable to `print.in`:

```fortran
jbm(f10.5)
```

If, instead of a mean value, you want a new maximum quantity, you need to replace `sum_mn_name()` by `max_mn_name()`.

Sect. 5.8.1 describes how to output horizontal averages of the magnetic and velocity fields. New such averages can be added to the code by using the existing averaging procedures `calc_bmz()` or `calc_jmz()` as examples.
9.3 The f-array

The ‘f’ array is the largest array in the PENCIL CODE and its primary role is to store the current state of the timestepped PDE variables. The f-array and its slightly smaller counterpart (the df-array; see below) are the only full size 3D arrays in the code. The f-array is of type real but PDEs for a complex variable may be solved by using two slots in the f-array. The actual size of the f-array is \( mx \times my \times mz \times m_{\text{farray}} \). Here, \( m_{\text{farray}} = m_{\text{var}} + m_{\text{aux}} + m_{\text{global}} + m_{\text{scratch}} \) where \( m_{\text{var}} \) refers to the number of real PDE variables.

As an example, we describe here how to put the time-integrated velocity, \( u_{\text{ut}} \), into the f-array (see ‘hydro.f90’). If this is to be invoked, there must be the following call somewhere in the code:

```fortran
call farray_register_auxiliary('uut',iuut,vector=3)
```

Here, \( iuut \) is the index of the variable \( u_{\text{ut}} \) in the f-array. Of course, this requires that \( m_{\text{aux}} \) is increased by 3, but in order to do this for a particular run only one must write a corresponding entry in the ‘cparam.local’ file,

```plaintext
! *** AUTOMATIC CPARAM.INC GENERATION ****************************************
! Declare (for generation of cpam.inc) the number of f array variables and auxiliary variables added by this module
!
! MAUX CONTRIBUTION 3
!
! Local settings concerning grid size and number of CPUs.
! This file is included by cpam.f90
!
integer, parameter :: ncpus=1,nprocy=1,nprocz=ncpus/nprocy,nprocx=1
integer, parameter :: nxgrid=16,nygrid=nxgrid,nzgrid=nxgrid
```

This way such a change does not affect the memory usage for other applications where this addition to ‘cparam.local’ is not made. In order to output this part of the f-array, one must write \( \text{lw} \)rite\_aux=T in the init\_pars of ‘start.in’. (Technically, \( [\text{lw} \)rite\_aux]=T can also be invoked in run\_pars of ‘run.in’, but this does not work at the moment.)

9.4 The df-array

The ‘df’ array is the second largest chunk of data in the PENCIL CODE. By using a 2N storage scheme (see H.4) after Williamson [35] the code only needs one more storage area for each timestepped variable on top of the current state stored in the f-array. As such, and in contrast to the f-array, the df-array is of size \( mx \times my \times mz \times m_{\text{var}} \). Like the df-array it is of type real. In fact the ghost zones of df are not required or calculated but having f- and df-arrays of the same size make the coding more transparent. For \( mx, my \) and \( mz \) large the wasted storage becomes negligible.
9.5 The fp-array

Similar to the ‘f’ array the code also has a ‘fp’ array which contains current states of all the particles. Like the f-array the fp-array also has a time derivative part, the dfp-array. The dimension of the fp-array is \( mpar_{\text{local}} \times mpar_{\text{var}} \) where \( mpar_{\text{local}} \) is the number of particles in the local processor (for serial runs this is the total number of particles) and \( mpar_{\text{var}} \) depends on the problem at hand. For example if we are solving for only tracer particles then \( mpar_{\text{var}} = 3 \), for dust particles \( mpar_{\text{var}} = 6 \) The sequence in which the slots in the fp-array are filled up depends on the sequence in which different particle modules are called from the particles_main.f90. The following are the relevant lines from particles_main.f90.

```fortran
!***********************************************************************
subroutine particles_register_modules()
!
! Register particle modules.
!
! 07-jan-05/anders: coded
!
   call register_particles   ()
   call register_particles_radius ()
   call register_particles_spin  ()
   call register_particles_number ()
   call register_particles_mass   ()
   call register_particles_selfgrav ()
   call register_particles_nbody   ()
   call register_particles_viscosity ()
   call register_particles_mass   ()
!
endsubroutine particles_register_modules
!***********************************************************************
```

The subroutine register_particles can mean either the tracer particles or dust particles. For the former the first three slots of the fp-array are the three spatial coordinates. For the latter the first six slots of the fp-array are the three spatial coordinates followed by the three velocity components. The seventh slot (or the fourth if we are use tracer particles) is the radius of the particle which can also change as a function of time as particles collide and fuse together to form bigger particles.

9.6 The pencil case

Variables that are derived from the basic physical variables of the code are stored in one-dimensional pencils of length \( nx \). All the pencils that are defined for a given set of physics modules are in turn bundled up in a Fortran structure called \( \mathbf{p} \) (or, more illustrative, the pencil case). Access to individual pencils happens through the variable \( \mathbf{p}(\text{name}) \), where \text{name} is the name of a pencil, e.g. \text{rho} that is a derived variable of the logarithmic density \text{lnrho}.

The pencils provided by a given physics module are declared in the header of the file, e.g. in the Density module:
! PENCILS PROVIDED lnrho; rho; rho1; glnrho(3); grho(3); uglnrho; ugrho

Notice that the pencil names are separated with a semi-colon and that vector pencils are declared with “(3)” after the name. Before compiling the code, the script `mkcparam` collects the names of all pencils that are provided by the chosen physics modules. It then defines the structure `p` with slots for every single of these pencils. The definition of the pencil case `p` is written in the include file `cparam_pencils.inc`. When the code is run, the actual pencils that are needed for the run are chosen based on the input parameters. This is done in the subroutines `pencil_criteria_modulename` that are present in each physics module. They are all called once before entering the time loop. In the `pencil_criteria` subroutines the logical arrays `lpenc_requested`, `lpenc_diagnos`, `lpenc_diagnos2d`, and `lpenc_video` are set according to the pencils that are needed for the given run. Some pencils depend on each other, e.g. `uglnrho` depends on `uu` and `glnrho`. Such interdependencies are sorted out in the subroutines `pencil_interdep_modulename` that are called after `pencil_criteria_modulename`.

In each time-step the values of the pencil logicals `lpenc_requested`, `lpenc_diagnos`, `lpenc_diagnos2d`, and `lpenc_video` are combined to one single pencil array `lpencil` which is different from time-step to time-step depending on e.g. whether diagnostics or video output are done in that time-step. The pencils are then calculated in the subroutines `calc_pencils_modulename`. This is done before calculating the time evolution of the physical variables, as this depends very often on derived variables in pencils.

The centralized pencil calculation scheme is a guarantee that

- All pencils are only calculated once
- Pencils are always calculated by the proper physics module

Since the PENCIL CODE is a multipurpose code that has many different physics modules, it can lead to big problems if a module tries to calculate a derived variable that actually belongs to another module, because different input parameters can influence how the derived variables are calculated. One example is that the Density module can consider both logarithmic and non-logarithmic density, so if the Magnetic module calculates

\[ \rho = \exp(f(l1:l2,m,n,ilnrho)) \]

it is wrong if the Density module works with non-logarithmic density! The proper way for the Magnetic module to get to know the density is to request the pencil `rho` in `pencil_criteria_magnetic`.

9.6.1 Pencil check

To check that the correct pencils have been requested for a given run, one can run a pencil consistency check in the beginning of a run by setting the logical `lpencil_check` in `&run_pars`. The check is meant to see if

- All needed pencils have been requested
- All requested pencils are needed

The consistency check first calculates the value of `df` with all the requested pencils. Then the pencil requests are flipped one at a time – requested to not requested, not requested to requested. The following combination of events can occur:
• not requested $\rightarrow$ requested, $df$ not changed
  The pencil is not requested and is not needed.
• not requested $\rightarrow$ requested, $df$ changed
  The pencil is not requested, but is needed. The code stops.
• requested $\rightarrow$ not requested, $df$ not changed
  The pencil is requested, but is not needed. The code gives a warning.
• requested $\rightarrow$ not requested, $df$ changed
  The pencil is requested and is needed.

### 9.6.2 Adding new pencils

Adding a new pencil to the pencil case is trivial but requires a few steps.

- Declare the name of the pencil in the header of the proper physics module. Pencils
  names must appear come in a ";" separated list, with dimensions in parenthesis
  after the name [(3) for vector, (3,3) for matrix, etc.].
- Set interdependency of the new pencil (i.e. what other pencils does it depend on) in
  the subroutine pencil_interdep_modulename
- Make rule for calculating the pencil in calc_pencils_modulename
- Request the new pencil based on the input parameters in any relevant physics
  module

Remember that the centralized pencilation scheme is partially there to force the users
of the code to think in general terms when implementing new physics. Any derived
variable can be useful for a number of different physics problems, and it is important
that a pencil is accessible in a transparent way to all modules.

### 9.7 Adding new physics: the Special module

If you want to add new physics to the code, you will in many cases want to add a new
Special module. Doing so is relatively straightforward and there is even a special direc-
tory for such additions.

To create your own special module, copy ‘nospecial.f90’ from the src/ directory to a new
name in the src/special/ directory. In many cases, users may want to put all new bits of
physics, needed for the specific problem at hand, into a single special module. The name
chosen for it should then relate to that problem. It is also possible to employ several (at
present up to five) different special modules at a time in a single setup which allows to
let naming follow the specific physics being implemented (for technicalities in this case,
see the end of this section).

The first thing to do in your new module is to change the lspecial=.false. header to say
lspecial=.true.

The file is heavily commented though all such comments can be removed as you go. You
may implement any of the subroutines/function that exist in nospecial.f90 and those
routines must have the names and parameters as in nospecial.f90. You do not how-
ever need to implement all routines, and you may either leave the dummy routines
copied from nospecial.f90 or delete them all together (provided the "include 'special_dummy.inc'" is kept intact at the end of the file. Beyond that, and data and subroutines can be added to a special module as required, though only for use within that module.

There are routines in the special interface to allow you to add new equations, modify the existing equation, add diagnostics, add slices, and many more things. If you feel there is something missing extra hooks can easily be added - please contact the PENCIL CODE team for assistance.

You are encouraged to submit/commit your special modules to the Pencil Code source. When you have added new stuff to the code, don’t forget to mention this in the ‘pencil-code/doc/manual.tex’ file.

Using more than one special module at a time requires that the environment variables $MODULE_PREFIX, $MODULE_INFIX and $MODULE_SUFFIX are set properly at runtime. They can be derived from the qualified names of module functions which have in general the form ⟨prefix⟩⟨module name⟩⟨infix⟩⟨function name⟩⟨suffix⟩ with its details depending on the Fortran compiler used. These can be learned by employing the nm command, say, by

```
unix> nm src/general.o | more .
```

The environment variables are most conveniently set in the user’s .bashrc, .cshrc or a proper configuration file of the PENCIL CODE (section environment). In the Makefile.local file, the requested special modules are simply specified as a list of names: SPECIAL = special/⟨module 1⟩ special/⟨module 2⟩.... In contrast to the case with only a single special module, where the namelists’ names are special_init_pars and special_run_pars, these are individualized for multiple special modules, viz. ⟨module name⟩_init_pars etc.

9.8 Adding switchable modules

In some cases where a piece of physics is thought to be more fundamental, useful in many situations or simply more flexibility is required it may be necessary to add a new module newphysics together with the corresponding nonewphysics module. The special modules follow the same structure as the rest of the switchable modules and so using a special module to prototype new ideas can make writing a new switchable module much easier.

For an example of module involving a new variable (and PDE), the pscalar module is a good prototype. The grep command

```
unix> grep -i pscalar src /*
```
gives you a good overview of which files you need to edit or add.

9.9 Adding your initial conditions: the InitialCondition module

Although the code has many initial conditions implemented, we now discourage such practice. We aim to eventually removed most of them. The recommended course of action is to make use of the InitialCondition module.
InitialCondition works pretty much like the Special module. To implement your own custom initial conditions, copy the file 'noinitial_condition.f90' from the src/ to src/initial_condition, with a new, descriptive, name.

The first thing to do in your new module is to change the linitialcondition=.false. header to say linitialcondition=.true. Also, don’t forget to add ../ in front of the file names in include statements.

This file has hooks to implement a custom initial condition to most variables. After implementing your initial condition, add the line INITIAL_CONDITION=initial_condition/myinitialcondition to your 'src/Makefile.local' file. Here, myinitialcondition is the name you gave to your initial condition file. Add also initial_condition_pars to the 'start.in' file, just below init_pars. This is a namelist, which you can use to add whichever quantity your initial condition needs defined, or passed. You must also un-comment the relevant lines in the subroutines for reading and writing the namelists. For compiling reasons, these subroutines in 'noinitial_condition.f90' are dummies. The lines are easily identifiable in the code.

Check e.g. the samples ‘2d-tests/baroclinic’, ‘2d-tests/spherical_viscous_ring’, or ‘interlocked-fluxrings’, for examples of how the module is used.
10 Testing the code

To maintain reproducibility despite sometimes quite rapid development, the PENCIL CODE is tested nightly on various architectures. The front end for testing are the scripts pc_auto-test and (possibly) pencil-test.

To see which samples would be tested, run

```
unix> pc_auto-test -l
```

to actually run the tests, use

```
unix> pc_auto-test
```
or

```
unix> pc_auto-test --clean
```

The latter compiles every test sample from scratch and currently (September 2009) takes about 2 hours on a mid-end Linux PC.

The pencil-test script is useful for cron jobs and allows the actual test to run on a remote computer. See Sect. 10.1 below.

For a complete list of options, run pc_auto-test --help and/or pencil-test --help.

10.1 How to set up periodic tests (auto-tests)

To set up a nightly test of the PENCIL CODE, carry out the following steps.

1. Identify a host for running the actual tests (the work host) and one to initiate the tests and collect the results (the scheduling host). On the scheduling host, you should be able to
   (a) run cron jobs,
   (b) ssh to the work host without password,
   (c) publish HTML files (optional, but recommended),
   (d) send e-mail (optional, but recommended).

   Work host and scheduling host can be the same (in this case, use pencil-test's '-l' option, see below), but often they will be two different computers.

2. [Recommended, but optional:] On the work host, check out a separate copy of the PENCIL CODE to reduce the risk that you start coding in the auto-test tree. In the following, we will assume that you checked out the code as '~/pencil-auto-test'.

3. On the work host, make sure that the code finds the correct configuration file for the tests you want to carry out. [Elaborate on that: PENCIL_HOME/local_config and '-f' option; give explicit example]

   Remember that you can set up a custom host ID file for your auto-test tree under '${PENCIL_HOME}/config-local/hosts/'.

4. On the scheduling host, use crontab -e to set up a cron job similar to the following:
30 02 * * * $HOME/pencil-auto-test/bin/pencil-test \ 
  -D $HOME/pencil-auto-test \ 
  --use-pc_auto_test \ 
  -N15 -Uc -rs \ 
  -T $HOME/public_html/pencil-code/tests/timings.txt \ 
  -t 15m \ 
  -m <email1@inter.net,email2@inter.net,...> \ 
  <work-host.inter.net> \ 
  -H > $HOME/public_html/pencil-code/tests/nightly-tests.html

**Note 1:** This has to be one long line. The backslash characters are written only for formatting purposes for this manual *you cannot use them in a crontab file.*

**Note 2:** You will have to adapt some parameters listed here and may want to modify a few more:

- `-D <dir>`: Sets the directory (on the work host) to run in.
- `-T <file>`: If this option is given, append a timing statistics line for each test to the given *file*.
- `--use-pc`: You want this option (and at some point, it will be the default).
- `-t 15m`: Limit the time for `start.x` and `run.x` to 15 minutes.
- `-N 15`: Run the tests at nice level 15 (may not have an effect for MPI tests).
- `-Uc`: Do `svn update` and `pc_build --cleanall` before compiling.
- `-m <email-list>`: If this option is given, send e-mails to everybody in the (comma-separated) list of e-mail addresses if any test fails. As soon as this option is set, the maintainers (as specified in the `README` file) of failed tests will also receive an e-mail.

`work-host.inter.net|-l`: Replace this with the remote host that is to run the tests. If you want to run locally, write `-l` instead.

- `-H`: Output HTML.


If you want to run fewer or more tests, you can use the `-Wa,--max-level` option:

```
-Wa,--max-level=3
```

will run all tests up to (and including) level 3. The default corresponds to `-Wa,--max-level=2`.

For a complete listing of `pencil-test` options, run

```
unix> pencil-test --help
```
11 Useful internals

11.1 Global variables

The following variables are defined in ‘cdata.f90’ and are available in any routine that uses the module Cdata.

<table>
<thead>
<tr>
<th>Variable</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>real</td>
<td></td>
</tr>
<tr>
<td>$t$</td>
<td>simulated time $t$.</td>
</tr>
<tr>
<td>integer</td>
<td></td>
</tr>
<tr>
<td>$n[xyz]grid$</td>
<td>global number of grid points (excluding ghost cells) in $x$, $y$ and $z$ direction.</td>
</tr>
<tr>
<td>$nx$, $ny$, $nz$</td>
<td>number of grid points (excluding ghost cells) as seen by the current processor, i.e. $ny=nygrid/nprocy$, etc.</td>
</tr>
<tr>
<td>$mx$, $my$, $mz$</td>
<td>number of grid points seen by the current processor, but including ghost cells. Thus, the total box for the $ivar$th variable (on the given processor) is given by $f(1:mx,1:my,1:mz,ivar)$.</td>
</tr>
<tr>
<td>$l1$, $l2$</td>
<td>smallest and largest $x$-index for the physical domain (i.e. excluding ghost cells) on the given processor.</td>
</tr>
<tr>
<td>$m1$, $m2$</td>
<td>smallest and largest $y$-index for physical domain.</td>
</tr>
<tr>
<td>$n1$, $n2$</td>
<td>smallest and largest $z$-index for physical domain, i.e. the physical part of the $ivar$th variable is given by $f(l1:l2,m1:m2,n1:n2,ivar)$.</td>
</tr>
<tr>
<td>$m$, $n$</td>
<td>pencil indexing variables: During each time-substep the box is traversed in $x$-pencils of length $mx$ such that the current pencil of the $ivar$th variable is $f(l1:l2,m,n,ivar)$.</td>
</tr>
<tr>
<td>logical</td>
<td></td>
</tr>
<tr>
<td>$lroot$</td>
<td>true only for MPI root processor.</td>
</tr>
<tr>
<td>$lfirst$</td>
<td>true only during first time-substep of each time step.</td>
</tr>
<tr>
<td>$headt$</td>
<td>true only for very first full time step (comprising 3 substeps for the 3rd-order Runge–Kutta scheme) on root processor.</td>
</tr>
<tr>
<td>$headtt$</td>
<td>= ($lfirst$.and.$lroot$): true only during very first time-substep on root processor.</td>
</tr>
<tr>
<td>$lfirstpoint$</td>
<td>true only when the very first pencil for a given time-substep is processed, i.e. for the first set of $(m,n)$, which is probably $(3,3)$.</td>
</tr>
<tr>
<td>$lout$</td>
<td>true when diagnostic output is about to be written.</td>
</tr>
</tbody>
</table>

11.2 Subroutines and functions

\texttt{output(file,a,nv)} (module IO): Write (in each ‘procN’ directory) the content of the global array $a$ to a file called $file$, where $a$ has dimensions $mx \times my \times mz \times nv$, or
output_pencil(file,a,nv) (module IO): Same as output(), but for a pencil variable, i.e. an auxiliary variable that only ever exists on a pencil (e.g. the magnetic field strength \( bb \) in ‘magnetic.f90’, or the squared sound speed \( cs^2 \) in ‘entropy.f90’). The file has the same structure as those written by output(), because the values of \( a \) on the different pencils are accumulated in the file. This involves a quite non-trivial access pattern to the file and has thus been coded in C (‘src/debug_c.c’).

cross(a,b,c) (module Sub): Calculate the cross product of two vectors \( a \) and \( b \) and store in \( c \). The vectors must either all be of size \( mx \times my \times mz \times 3 \) (global arrays), or of size \( nx \times 3 \) (pencil arrays).

dot(a,b,c) (module Sub): Calculate the dot product of two vectors \( a \) and \( b \) and store in \( c \). The vectors must either be of size \( mx \times my \times mz \times 3 \) (\( a \) and \( b \)) and \( mx \times my \times mz \) (\( c \)), or of size \( nx \times 3 \) (\( a \) and \( b \)) and \( nx \) (\( c \)).

dot2(a,c) (module Sub): Same as dot(a,a,c).
Part III

Appendix

APPENDIX

A Timings

In the following table we list the results of timings of the code on different machines. Shown is (among other quantities) the wall clock time per mesh point (excluding the ghost zones) and per full 3-stage time step, a quantity that is printed by the code at the end of a run.

As these results were assembled during the development phase of the code (that hasn’t really finished yet, . . .), you may not get the same numbers, but they should give some orientation of what to expect for your specific application on your specific hardware.

The code will output the timing (in microseconds per grid point per time-step) at the end of a run. You can also specify `walltime` in `print.in` to have the code continuously output the physical time it took to reach the time-steps where diagnostics is done. The time-dependent code speed can then be calculated by differentiating, e.g. in IDL with

```
IDL> pc_read_ts, obj=ts
IDL> plot, ts.it, 1/nw*deriv(ts.it,ts.walltime/1.0e-6), psym=2
```

where `nw=nx*ny*nz`.

<table>
<thead>
<tr>
<th>proc</th>
<th>machine</th>
<th>µs</th>
<th>resol.</th>
<th>what</th>
<th>mem/proc</th>
<th>when</th>
<th>who</th>
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<td>1</td>
<td>Nl3</td>
<td>19</td>
<td>64³</td>
<td>kinematic</td>
<td>10 MB</td>
<td>20-may-02</td>
<td>AB</td>
</tr>
<tr>
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<td>30</td>
<td>64³</td>
<td>magn/noentro</td>
<td>20 MB</td>
<td>20-may-02</td>
<td>AB</td>
</tr>
<tr>
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<td>10</td>
<td>64³</td>
<td>magn/noentro</td>
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<td>32³</td>
<td>magn/noentro</td>
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<td>BD</td>
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<td>128³</td>
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<td>AB</td>
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<td>2</td>
<td>Nq5+5</td>
<td>3.7</td>
<td>128³</td>
<td>magn/noentro</td>
<td>8-oct-02</td>
<td>AB</td>
<td></td>
</tr>
</tbody>
</table>

Note that when using `nompicomm.f90`, the timer currently used will overflow on some machines, so you should not blindly trust the timings given by the code.
<table>
<thead>
<tr>
<th>Name</th>
<th>Size</th>
<th>Format</th>
<th>Date</th>
<th>Author</th>
</tr>
</thead>
<tbody>
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<td>64^3</td>
<td>nomag/noentro</td>
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<td>magn/noentro</td>
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A. Timings

1024 Hermit 0.00707 512×1024×512 spherical conv/magn 2×32×16 22-aug-13 PJK
1024 Hermit 0.00698 1024×2048×1024 spherical conv/magn 4×16×16 22-aug-13 PJK
1024 Hermit 0.00630 512×1024×512 spherical conv/magn 4×16×16 22-aug-13 PJK
1024 Triolith 0.00236 512×1024×512 magn/noentro 4×16×16 1-mar-14 AB
1024 Triolith 0.00126 512×1024×512 magn/noentro 2×16×32 1-mar-14 AB
1024 Triolith 0.00129 512×1024×512 magn/noentro 4×16×16 1-mar-14 AB
1024 Sisu 0.00225 1024×2048×1024 spherical conv/magn 22-aug-13 PJK
1024 Sisu 0.00148 512×1024×512 spherical conv/magn 2×32×16 22-aug-13 PJK
1512 Kraken 0.0212 192×384×64 magn/noentro 12×24×4 13-jan-12 WL
1512 Kraken 0.00856 384×768×128 magn/noentro 12×24×4 17-jan-12 WL
1512 Kraken 0.00549 768×1536×256 magn/noentro 12×24×4 17-jan-12 WL
1512 Lindgren 0.016 512×512 magn/rad 1×36×32 17-mar-14 AB
1512 Lindgren 0.0066 1152×1152 magn/noentro 1×32×36 25-nov-14 AB
1512 Beskow 0.0055 1152×1152 GWO/magn/noentro 1×32×36 27-aug-17 AB
1512 Beskow 0.0024 1152×1152 magn/noentro 1×32×36 20-jan-15 AB
1512 Beskow 0.00098 1152×1152 magn/noentro 1×32×36 18-jan-16 AB-gnu
1512 Beskow 0.00090 1152×1152 magn/noentro 1×32×36 30-mar-17 AB
1512 Beskow 0.0060 1152×1152 GWO/magn/noentro 1×32×36 31-mar-18 AB
1512 Beskow 0.0063 576×576 magn/noentro 1×32×36 17-feb-18 AB
1512 Beskow 0.0030 1152×1152 GWN/magn/noentro 1×32×36 30-jul-20 AB
1512 Beskow 0.0017 1152×1152 GWO/magn/noentro 1×32×36 30-jul-20 AB
1536 Lindgren 0.00171 512×384 magn/noentro 2×32×24 15-jun-13 AB
2048 Hermit 0.00451 1024×2048×1024 spherical conv/magn 2×32×32 22-aug-13 PJK
2048 Hermit 0.00380 512×1024×512 spherical conv/magn 8×16×16 22-aug-13 PJK
2048 Hermit 0.00355 512×1024×512 spherical conv/magn 4×32×16 22-aug-13 PJK
2048 Hermit 0.00350 1024×2048×1024 spherical conv/magn 4×32×16 22-aug-13 PJK
2048 Beskow 0.0022 1024×1024 GWN/magn/noentro 8×16×16 16-aug-20 AB
2048 Lindgren 0.00129 512×1024 magn/noentro 32×64 20-apr-13 AB
2048 Lindgren 0.00129 1024×2048 magn/noentro 32×64 31-jul-12 AB
2048 Triolith 9.3×10⁻⁴ 512×1024 magn/noentro 4×16×32 1-mar-14 AB
2048 Sisu 0.00120 1024×2048×1024 spherical conv/magn 22-aug-13 PJK
2048 Sisu 9.2×10⁻⁴ 512×1024×512 spherical conv/magn 4×32×16 22-aug-13 PJK
2304 Triolith 1.07×10⁻³ 576×576 magn/noentro 4×18×32 1-mar-14 AB
2304 Kraken 0.02267 192×384×64 magn/noentro 12×24×8 13-jan-12 WL
2304 Kraken 0.01233 192×768×64 magn/noentro 12×48×4 13-jan-12 WL
2304 Kraken 0.00300 768×3072×256 magn/noentro 12×48×4 18-jan-12 WL
4096 Hermit 0.00193 1024×2048×1024 spherical conv/magn 4×32×32 22-aug-13 PJK
4096 Triolith 3.6×10⁻⁴ 1024×1024 magn/noentro 8×16×32 1-mar-14 AB
4096 Triolith 3.8×10⁻⁴ 1024×1024 magn/noentro 8×16×32 1-mar-14 AB
4096 Triolith 4.2×10⁻⁴ 1024×1024 magn/noentro 4×16×64 1-mar-14 AB
4096 Lindgren 4.6×10⁻⁴ 2048×1024 magn/noentro 4×16×64 26-mar-13 AB
4096 Sisu 6.7×10⁻⁴ 1024×2048×1024 spherical conv/magn 22-aug-13 PJK
4608 Triolith 7.4×10⁻⁴ 576×576 magn/noentro 8×18×32 1-mar-14 AB
4608 Triolith 2.7×10⁻⁴ 1152×1152 magn/noentro 4×32×36 1-mar-14 AB
4608 Triolith 3.0×10⁻⁴ 1152×1152 magn/noentro 4×36×32 1-mar-14 AB
4608 Triolith 3.7×10⁻⁴ 1152×1152 magn/noentro 4×18×64 1-mar-14 AB
4608 Triolith 2.36×10⁻⁴ 2304×1024 magn/noentro 2×32×72 1-mar-14 AB
4608 Kraken 0.00764 192×768×128 magn/noentro 12×48×8 13-jan-12 WL
4608 Kraken 0.00144 768×3072×512 magn/noentro 12×48×8 18-jan-12 WL
6144 Lindgren 4.2×10⁻⁴ 1024×1536 magn/noentro 4×16×64 21-oct-13 AB
6144 Lindgren 8.9×10⁻⁴ 256×512 magn/noentro/sph 2×48×64 6-jan-15 SJ
8192 Hermit 0.00101 1024×2048×1024 spherical conv/magn 8×32×32 22-aug-13 PJK
8192 Sisu 4.1×10⁻⁴ 1024×2048×1024 spherical conv/magn 22-aug-13 PJK
8192 Triolith 1.48×10⁻⁴ 2048×1024 magn/noentro 4×32×64 1-mar-14 AB
9216 Kraken 0.00485 192×768×256 magn/noentro 24×48×8 13-jan-12 WL
9216 Kraken 0.00158 768×1536×256 magn/noentro 24×48×8 17-jan-12 WL
9216 Kraken 8.0×10⁻⁴ 1536×3072×512 magn/noentro 24×48×8 18-jan-12 WL
9216 Lindgren 2.36×10⁻⁴ 2304×1024 magn/noentro 4×48×48 15-feb-14 AB
9216 Triolith 1.04×10⁻¹ 576×1152 magn/noentro 16×18×32 1-mar-14 AB
9216 Triolith 1.28×10⁻⁴ 2304×1024 magn/noentro 4×36×64 1-mar-14 AB
The machines we have used can be characterized as follows:

**NI3:** 500 MHz Pentium III single CPU; RedHat Linux 6.2; 256 MB memory

**Nq0:** 931 MHz Pentium III single CPU; RedHat Linux 7.3; 0.5 GB memory

**Nq[1-4]:** 869 MHz Pentium III dual-CPU cluster; RedHat Linux 7.3; 0.77 GB memory per (dual) node

**Nq[5-6]:** 1.2 GHz Athlon dual-CPU cluster; RedHat Linux 7.3; 1 GB memory per (dual) node

**Kabul:** 1.9 GHz Athlon dual-CPU cluster; 1 GB memory per (dual) node; 256 kB cache per CPU; Gigabit ethernet; SuSE Linux 8.0; LAM-MPI

**Cincinnatus:** 1.7 GHz Pentium 4 single CPU; 1 GB memory; 256 kB cache per CPU; SuSE Linux 7.3

**Horseshoe (fe1, giga, and giga2):** consists of different subclusters. The old one (queue name: workq, referred to as fe1) 2.0 GHz Pentium 512 single CPU; 25x 24-port fast ethernet switches with gigabit ethernet uplink; 1 30-port gigabit ethernet switch; 1 GB memory. The next generation has gigabit switches directly between nodes, and 2.6 GHz processors. The third generation (giga2) has 3.2 GHz processors (most of which have 1 GB, some 2 GB), is organized in 2 blocks interconnected with 2 Gb links, with 10 Gb uplinks within each block.

**Ukaff:** SGI Origin 3000; 400 MHz IP35 CPUs; IRIX 6.5; native MPI

**Mhd:** EV6 Compaq cluster with 4 CPUs per node; 4 GB memory per node (i.e. 1 GB per CPU) OSF1 4.0; native MPI

**Sander and Rasmussen:** Origin 3000

**Steno** 118 node IBM cluster with dual node AMD Opteron processors with 10 Gb infiniband network, compiled with pgf90 -fastsse -tp k8-64e (Copenhagen).

**Gridur:** Origin 3000

**Luci:** (full name Lucidor) is an HP Itanium cluster, each of the 90 nodes has two 900 MHz Itanium 2 "McKinley" processors and 6 GB of main memory. The interconnect is myrinet.

**Lenn:** (full name Lenngren) is a Dell Xeon cluster with 442 nodes. Each node has two 3.4GHz “Nocona” Xeon processors and 8GB of main memory. A high performance Infiniband network from Mellanox is used for MPI traffic.
Kraken: Cray Linux Environment (CLE) 3.1, with a peak performance of 1.17 PetaFLOP; the cluster has 112,896 cores, 147 TB of memory, in 9,408 nodes. Each node has two 2.6 GHz six-core AMD Opteron processors (Istanbul), 12 cores, and 16 GB of memory. Connection via Cray SeaStar2+ router.

Hermit: Cray XE6 with 7104 2.3 GHz AMD Interlagos 16 core processors (113,664 cores in total), nodes with either 1 or 2 GB of memory per core.

Sisu: Cray XC30 with 1472 2.6 GHz Intel (Xeon) Sandy Bridge 8 core (E5-2670) processors (11,776 cores in total), 2 GB of memory per core.

Beskow: Cray XC40 with 2.3 GHz Intel (Xeon) Haswell 16 core (E5-2698v3) processors (67,456 cores in total), 2 GB of memory per core. Theoretical peak performance 2.43 pflops.

Table 8 shows a similar list, but for a few well-defined sample problems. The svn check-in patterns are displayed graphically in Fig 1.

Figure 13: Scaling results on three different machines. The thin straight line denotes perfectly linear scaling.

Table 8: Like previous table, but for the versions from the 'samples' directory.

<table>
<thead>
<tr>
<th>proc(s)</th>
<th>machine</th>
<th>$\mu$s pt step</th>
<th>resol.</th>
<th>mem./proc</th>
<th>when</th>
<th>who</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>conv-slab</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>1 Mhd</td>
<td>6.45</td>
<td>$32^3$</td>
<td>4 MB</td>
<td>23-jul-02</td>
<td>wd</td>
<td></td>
</tr>
<tr>
<td>1 Cincinnatus</td>
<td>4.82</td>
<td>$32^3$</td>
<td>3 MB</td>
<td>23-jul-02</td>
<td>wd</td>
<td></td>
</tr>
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<td>11.6</td>
<td>$64^3$</td>
<td>14 MB</td>
<td>23-jul-02</td>
<td>wd</td>
<td></td>
</tr>
<tr>
<td>1 Cincinnatus</td>
<td>20.8</td>
<td>$128^3$</td>
<td>93 MB</td>
<td>23-jul-02</td>
<td>wd</td>
<td></td>
</tr>
<tr>
<td>1 Kabul</td>
<td>3.91</td>
<td>$32^3$</td>
<td>23-jul-02</td>
<td>wd</td>
<td></td>
<td></td>
</tr>
<tr>
<td>1 Kabul</td>
<td>3.88</td>
<td>$64^3$</td>
<td>23-jul-02</td>
<td>wd</td>
<td></td>
<td></td>
</tr>
<tr>
<td>1 Kabul</td>
<td>4.16</td>
<td>$128^3$</td>
<td>93 MB</td>
<td>23-jul-02</td>
<td>wd</td>
<td></td>
</tr>
<tr>
<td>conv-slab-flat</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>1 Kabul</td>
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<td>29 MB</td>
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<td>wd</td>
<td></td>
</tr>
<tr>
<td>2 Kabul</td>
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<td>18 MB</td>
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<td>wd</td>
<td></td>
</tr>
<tr>
<td>4 Kabul</td>
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<td>$128^2 \times 32$</td>
<td>11 MB</td>
<td>23-jul-02</td>
<td>wd</td>
<td></td>
</tr>
<tr>
<td>8 Kabul</td>
<td>0.87</td>
<td>$128^2 \times 32$</td>
<td>9 MB</td>
<td>23-jul-02</td>
<td>wd</td>
<td></td>
</tr>
</tbody>
</table>
Figure 14: Scaling results of particle-mesh problem on Blue Gene/P on up to 4096 cores. The different lines denote different particle-mesh schemes (NGP=Nearest Grid Point, TSC=Triangular Shaped Cloud) and whether self-gravity is included (FFT).

A.1 Test case

In the following test samples, we run isothermal magnetohydrodynamics in a periodic domain.\(^{17}\) Power spectra are computed during the run, but our current parallelization of the Fourier transform requires that the meshpoint number is an integer multiple of the product of processor numbers in the \(y\) and \(z\) directions and the product of processor


Figure 15: Scaling results on Kraken at fixed problem size, for a magnetized disk model in cylindrical coordinates. The black line shows ideal scaling from 32 cores. The blue line is the best second-order fit to the data points. A load of \(16^3\) mesh points per processor marks the best strong scaling.
A.2 Running the code

To run the code, get one of the sample run directories, e.g., [http://norlx51.nordita.org/~brandenb/pencil-code/timings/bforced/512_4x16x32](http://norlx51.nordita.org/~brandenb/pencil-code/timings/bforced/512_4x16x32). The relevant file to be changed is `src/cparam.local` in particular the values of ncpus, nprocx, nprocy, and nxgrid. Once they are chosen, say make, and submit start_run.csh.

A.3 Triolith

On Triolith, strong scaling tests have been performed for three mesh sizes. The time per time step and mesh point is given for different processor numbers and layouts. Generally, it is advantageous to keep the number of processors in the $x$ direction small.

Comments. Although on Triolith the number of processors per node is 16, resolutions with one or two powers of 3 (such as 576) still work well. Furthermore, the number of processors above which the scaling becomes poor increases quadratically with the number of mesh points. This implies that the RAM per processor increases linearly with the problem size per direction. However, this will not be a limitation, because even for $2304^3$ meshes, the required RAM is still below 100 MB.
A.4 Lindgren

On Lindgren, we have performed weak scaling tests and compare with weak scaling results for Triolith. Triolith is about twice as fast as Lindgren.

Figure 17: Strong scaling on Triolith (2014).

Figure 18: Comparison Triolith (black, plus signs) and Lindgren (red, triangles). Weak scaling (2014).
Table 9: Triolith timings

<table>
<thead>
<tr>
<th>proc</th>
<th>$\mu$s</th>
<th>pt step</th>
<th>resol.</th>
<th>layout</th>
</tr>
</thead>
<tbody>
<tr>
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<td>128$^3$</td>
<td>2x2x4</td>
<td></td>
</tr>
<tr>
<td>16</td>
<td>0.065</td>
<td>128$^3$</td>
<td>1x4x4</td>
<td></td>
</tr>
<tr>
<td>16</td>
<td>0.0544</td>
<td>256$^3$</td>
<td>1x4x4</td>
<td></td>
</tr>
<tr>
<td>64</td>
<td>0.0146</td>
<td>256$^3$</td>
<td>1x8x8</td>
<td></td>
</tr>
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</tr>
<tr>
<td>1024</td>
<td>0.00129</td>
<td>512$^3$</td>
<td>4x16x16</td>
<td></td>
</tr>
<tr>
<td>2048</td>
<td>9.34x10^{-4}</td>
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<td>4x16x32</td>
<td></td>
</tr>
<tr>
<td>2304</td>
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<td>4x18x32</td>
<td></td>
</tr>
<tr>
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<td>4x32x32</td>
<td></td>
</tr>
<tr>
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</tr>
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<tr>
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<td>4x36x64</td>
<td></td>
</tr>
<tr>
<td>9216</td>
<td>1.30x10^{-4}</td>
<td>2304$^3$</td>
<td>4x32x72</td>
<td></td>
</tr>
</tbody>
</table>

Table 10: Lindgren timings

<table>
<thead>
<tr>
<th>proc</th>
<th>$\mu$s</th>
<th>pt step</th>
<th>resol.</th>
<th>layout</th>
</tr>
</thead>
<tbody>
<tr>
<td>1536</td>
<td>0.00171</td>
<td>512$^2$x384</td>
<td>2x32x24</td>
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</tr>
<tr>
<td>2048</td>
<td>0.00129</td>
<td>512$^2$x1024</td>
<td>1x32x64</td>
<td></td>
</tr>
<tr>
<td>2048</td>
<td>0.00129</td>
<td>1024$^2$x2048</td>
<td>1x32x64</td>
<td></td>
</tr>
<tr>
<td>4096</td>
<td>4.6x10^{-4}</td>
<td>2048$^3$</td>
<td>4x16x64</td>
<td></td>
</tr>
<tr>
<td>9216</td>
<td>2.36x10^{-4}</td>
<td>2304$^3$</td>
<td>4x48x48</td>
<td></td>
</tr>
</tbody>
</table>
B Coding standard

The numerous elements that make up the PENCIL CODE are written in a consistent style that has evolved since it was first created. Many people have contributed their knowledge and experience with in this and the result is what we believe is and extremely readable and manageable code.

As well as improving the readability of the code, by having some naming conventions for example aids greatly in understanding what the code does.

There is a standard for all aspects of the code, be it Fortran source, shell scripts, Perl scripts, LaTeX source, Makefiles, or otherwise. Where nothing has been explicitly stated it is recommended that similar existing examples found in the code are used as a template.

B.1 File naming conventions

All files with the exception of the ‘Makefile’ s are given lowercase filenames.

Fortran source files all have the ‘.f90’ extension. Files that contain ‘non-executable code’ i.e. declarations that are included into other files are given the extension ‘.h’ and those that are generated dynamically at compile time have an ‘.inc’ extension.

Fortran source code defining a module is placed in files whose names begin with the Fortran module name in all lowercase. Where there exist multiple implementations of a specific module, the filenames are extended using an underscore and a brief name relating to what they do.

Text files containing parameters to be read by the code at run time are placed in files with the extension ‘.in’

B.2 Fortran Code

The code should remain fully compatible with the Fortran90 standard. This ensures that the code will run on all platforms. Indeed, an important aspect of PENCIL CODE philosophy is to be maximally flexible. This also means that useful non-standard extensions to the code should be hidden in and be made accessible through suitable non-default modules.

Fortran is not case-sensitive but in almost all instances we prescribe some form of capitalization for readability.

In general all Fortran code including keywords, variable names etc. are written in lowercase. Some of the coding standard has already been discussed in Sect. 9.1. Here we discuss and amplify some remaining matters.

B.2.1 Indenting and whitespace

Whitespace should be removed from the end of lines.

Blank lines are kept to a minimum, and when occurring in subroutines or functions are replaced by a single ‘!’ in the first column.
Tab characters are not used anywhere in the code. Tab characters are not in fact allowed by the Fortran standard and compilers that accept them do so as an extension.

All lines are kept to be not more than 80 characters long. Where lines are longer they must be explicitly wrapped using the Fortran continuation character ‘&’. Longer lines (up to 132 characters) and additional spaces are allowed in cases where the readability of the code is enhanced, e.g. when one line is followed by a similar one with minor differences in some places.

Code in syntactic blocks such as if–endif, do–endo, subroutine–endsubroutine etc. is always indented by precisely two spaces. The exception to this is that nested loops where only the innermost loop contains executable code should be written with the do–endo pairs at the same level of indentation,

```
   do n=n1,n2
   do m=m1,m2
      [...]
   enddo
   enddo
```

Alternatively nested loops may be written on a single line, i.e.

```
   do n=n1,n2; do m=m1,m2
      [...]
   enddo; enddo
```

### B.2.2 Comments

Descriptive comments are written on their own lines unless there is a strong reason to do otherwise. Comments are never indented and the ‘!’ should appear in the first column followed by two spaces and then the text of the comment. Extremely short comments may follow at the end of a line of code, provided there is space.

Comments also must not exceed the 78 character line length and should be wrapped onto more lines as needed.

Typically comments should appear with a blank commented line above and below the wrapped text of the comment.

All subroutine/functions begin with a standard comment block describing what they do, when and by whom they were created and when and by whom any non-trivial modifications were made.

Comments should be written in sentences using the usual capitalization and punctuation of English, similar to how text is formatted in an e-mail or a journal article.

For example:

```
some fortran code
some more fortran code
!
! A descriptive comment explaining what the following few lines
! of code do.
!
   the fortran code being described
```
Subroutines and functions are started with a comment block describing what they do, when and by whom they were created and when and by whom any non-trivial modifications were made. The layout of this comment block is a standard, for example:

```fortran
!***********************************************************************
subroutine initialize_density(f,lstarting)
!
! Perform any post-parameter-read initialization i.e. calculate derived
! parameters.
!
! For compatibility with other applications, we keep the possibility
! of giving diffrho units of dmin*cs0, but cs0 is not well defined general.
!
! 24-nov-02/tony: coded
! 1-aug-03/axel: normally, diffrho should be given in absolute units
!
```

where dates are written in dd-mmm-yy format as shown and names appearing after the ‘/’ are either the users cvs login name or, where such exists amongst the PENCIL CODE community, the accepted short form (∼ 4 characters) of the authors name.

### B.2.3 Module names

The names of modules are written with initial letter capitalization of each word and the multiple words written consecutively without any separator.

### B.2.4 Variable names

Variable are given short but meaningful names and written in all lowercase. Single character names are avoided except for commonly used loop indices and the two code data structures of the PENCIL CODE: ‘f’ the main state array (see [9.3]) and ‘p’ the pencil case structure (see [9.6]).

Quantities commonly represented by a particular single character in mathematics are typically given names formed by repeating the character (usually in lowercase), e.g. the velocity \( u \) becomes ‘uu’, specific entropy \( s \) becomes ‘ss’ etc.

Temperature in variable names is denoted with a capital T so as not to be confused with time as represented by a lowercase \( t \). Note however the since Fortran is not case sensitive the variables for example ‘TT’ and ‘tt’ are the same so distinct names must be used. For this reason time is usually represented by a single \( t \) contrary to the above guideline.
The natural log of a quantity is represented by using adding ‘ln’ to its name, for example log of temperature would be ‘lnTT’.

There are some standard prefixes used to help identify the type and nature of variables they are as follows:

- i – Denotes integer variables typically used as array indices.
- i_ – Denotes pencil case array indices.
- idiag_ – Denotes diagnostic indices.
- l – Denotes logical/boolean flags
- cdt – Denotes timestep constraint parameters.
- unit_ – Denotes conversion code/physics unit conversion parameters.

### B.2.5 Emacs settings

Here are some settings from wd’s ‘~/.emacs’ file:

```lisp
;; ~/.f90.emacs
;;; Set up indentation and similar things for coding the \sc Pencil Code\. Most of this can probably be set through Emacs’ Customize interface
;;; as well.
;;; To automatically load this file, put the lines
;;; (if (file-readable-p "~/.f90\.emacs")
;;; (load-file "~/.f90\.emacs"))
;;; into your ~/.emacs file.

;;; F90-mode indentation widths
(setq f90-beginning-ampersand nil) ; no 2nd ampersand at continuation line
(setq f90-do-indent 2)
(setq f90-if-indent 2)
(setq f90-type-indent 2)
(setq f90-continuation-indent 4)

;;; Don’t use any tabs for indentation (with TAB key).
;;; This is actually already set for F90-mode.
(setq-default indent-tabs-mode nil)

;;; Ensure Emacs uses F90-mode (and not Fortran-mode) for F90 files:
(setq auto-mode-alist
  (append
   '("\.[fF]90\$" . f90-mode)
   ('"\.[iI]nc\$" . f90-mode)
  )
  auto-mode-alist))
```

;; Make M-Backspace behave in Xemacs as it does in GNU Emacs. The default
;; behavior is apparently a long-known bug the fix for which wasn’t
;; propagated from fortran.el to f90.el.
;; (http://list-archive.xemacs.org/xemacs-patches/200109/msg00026.html):
(add-hook 'f90-mode-hook
  (function (lambda ()
    (define-key f90-mode-map [(meta backspace)] 'backward-kill-word)
  )))

B.3 Other best practices

When implementing IF or SELECT blocks always write code for all cases – including the default or else case. This should be done even when that code is only a call to raise an error that the case should not have been reached. If you see a missing case anywhere then do add it. These failsafes are essential in a large multi-purpose multi-user code like the PENCIL CODE.

If a case is supposed to do nothing and it may be unclear that the coder has recognized this fact then make it explicit by adding the default case with a comment like ! Do Nothing. The compiler will clean away any such empty blocks.

B.4 General changes to the code

It is sometimes necessary to do major changes to the code. Since this may affect many people and may even be controversial among the developers, such changes are restricted to the time of the next Pencil Code User Meeting. Such meetings are advertised on http://www.nordita.org/software/pencil-code/ under the news section. Notes about previous such meetings can be found under http://www.nordita.org/software/pencil-code/UserMeetings/.

Major changes can affect those developers who have not checked in their latest changes for some time. Before doing such changes it is therefore useful to contact the people who have contributed to the latest developments on that module. If it is not functional or otherwise in bad shape, it should be moved to ‘experimental’, i.e. one says svn mv file.f90 experimental/file.f90. However, any such directory change constitutes a major change in itself and should be performed in agreement with those involved in the development. Otherwise any file that has been changed in the mean time will end up being outside revision control, which is to be avoided at all cost.
C Some specific initial conditions

C.1 Random velocity or magnetic fields

Obtained with inituu='gaussian-noise' (or initaa='gaussian-noise'). The vector $u$ (or $A$) is set to normally distributed, uncorrelated random numbers in all meshpoints for all three components. The power spectrum of $u$ ($A$) increases then quadratically with wavenumber $k$ (without cutoff) and the power spectrum of $\omega$ ($B$) increases like $k^4$.

Note that a random initial condition contains significant power at the Nyquist wavenumber $k_{Ny} = \pi/\delta x$, where $\delta x$ is the mesh spacing. In a decay calculation, because of the discretization error, such power decays slower than it ought to; see Fig. [19] where we show the evolution for a random initial velocity field for $64^3$ meshpoints, $\nu = 5 \times 10^{-2}$ (fairly large!), and nfilter=30.

It is clearly a good idea to filter the initial condition to prevent excess power at $k_{Ny}$. On the other hand, such excess power is weak by comparison with the power at the energy carrying scale, so one does not see it in visualizations in real space. Furthermore, as seen from Fig. [19] for $k < k_{Ny}/2$ the power spectra for filtered and unfiltered initial conditions is almost the same.

Figure 19: Velocity power spectra at three different times with and without filtering of the initial condition.

C.2 Turbulent initial with given spectrum

The most general procedure for producing an initial condition with a turbulent spectrum is inituu='power_randomphase_hel', which allows one to set two different slopes, together with an exponential cutoff as well as a Gaussian peak within the spectrum. By default, the field is solenoidal unless one puts lskip_projection=.true. and can have fractional helicity by setting relhel_uu to a value between $-1$ and $1$. By default it is 0, which means it is nonhelical.
The spectral indices \textit{initpower} and \textit{initpower2} refer to energy spectral indices. By default, \textit{initpower2}=-5/3, corresponding to a Kolmogorov spectrum. For a delta-correlated spectrum, we have to put \textit{initpower}=2, corresponding to a $k^2$ energy spectrum for kinetic energy. This would be suitable for the subinertial range from $k = 1$ to $k = k_p$ (corresponding to the variable \textit{kpeak}).

If \textit{cutoff}=0, no cutoff will be imposed. Otherwise, the spectrum will be multiplied by an exponential function with $\exp(-k^{2n})$, where $n = 'ncutoff=1'$ by default.

Example, for \textit{ampluu}=1e-1, \textit{initpower}=4., and \textit{kpeak}=3., we get \textit{urms}=3.981E-01 when \textit{relhel_uu}=1 and \textit{urms}=3.981E-01 when \textit{relhel_uu}=0 and \textit{urms}=5.560E-01 when \textit{relhel_uu}=1. The \textit{urms} values scale linearly with \textit{ampluu} and for \textit{initpower}=2 also approximately linearly with \textit{kpeak}. For the magnetic field, we initialize the magnetic vector potential, so to get a $k^4$ spectrum, we have to put \textit{initpower_aa}=2. Everything else is analogous; see, e.g.,

\begin{verbatim}
&hydro_init_pars
  inituu='power_randomphase_hel', ampluu=1e-1, initpower=4., kpeak=3.
  relhel_uu=0., cutoff=30.
/
&magnetic_init_pars
  initaa='power_randomphase_hel', amplaa=1e-1, initpower_aa=2., kpeak_aa=3.
  relhel_aa=0., cutoff_aa=30.
/
\end{verbatim}

for which we get \textit{urms}=3.981E-01 and \textit{brms}=3.871E-01.

\section{C.3 Beltrami fields}

Obtained with \textit{inituu}='Beltrami-z' or \textit{initaa}='Beltrami-z'.

\begin{equation}
A = (\cos z, \sin z, 0), \quad \text{or} \quad u = (\cos z, \sin z, 0)
\end{equation}

\section{C.4 Magnetic flux rings: \textit{initaa}='fluxrings'}

This initial condition sets up two interlocked thin magnetic tori (i.e. thin, torus-shaped magnetic flux tubes). One torus of radius $R$ lying in the plane $z = 0$ can be described in cylindrical coordinates, $(r, \varphi, z)$, by the vector potential

\begin{equation}
A = \Phi_m \begin{pmatrix} 0 \\ 0 \\ -\theta(r-R)\delta(z) \end{pmatrix},
\end{equation}

resulting in a magnetic field

\begin{equation}
B = \Phi_m \begin{pmatrix} 0 \\ \delta(r-R)\delta(z) \\ 0 \end{pmatrix}.
\end{equation}

Here $\Phi_m$ is the magnetic flux through the tube, $\theta(x)$ denotes the Heaviside function, and $\delta(x) = \theta'(x)$.
is Dirac’s delta function.

Any smoothed versions of \( \theta(x) \) and \( \delta(x) \) will do, as long as the consistency condition (129) is satisfied. E.g. the pairs

\[
\delta_\varepsilon(x) = \frac{1}{\sqrt{2\pi\varepsilon^2}} e^{-\frac{x^2}{2\varepsilon^2}}, \quad \theta_\varepsilon(x) = \frac{1}{2} \left( 1 + \text{erf} \frac{x}{\sqrt{2\varepsilon}} \right)
\]

or

\[
\delta_\varepsilon(x) = \frac{1}{2\varepsilon} e^{-\frac{x^2}{2\varepsilon}}, \quad \theta_\varepsilon(x) = \frac{1}{2} \left( 1 + \tanh \frac{x}{\varepsilon} \right)
\]

are quite popular. Another possibility is a constant or box-like profile with

\[
\delta_\varepsilon(x) = \frac{1}{2\varepsilon} \theta(|x| - \varepsilon), \quad \theta_\varepsilon(x) = \frac{1}{2} \left\{ 1 + \max\{-1, \min(x/\varepsilon), 1\} \right\}
\]

Note, however, that the Gaussian profile (130) is the only one that yields a radially symmetric (with respect to the distance from the central line of the torus) magnetic field profile

\[
B_\varphi = B_\phi \left( \sqrt{(r-R)^2 + z^2} \right)
\]

if \( \varepsilon \) is sufficiently small.

In Cartesian coordinates, the vector potential (127) takes the form

\[
A = \Phi_m \begin{pmatrix} 0 \\ 0 \\ -\theta \left( \sqrt{x^2 + y^2} - R \right) \delta(z) \end{pmatrix}.
\]

### C.5 Vertical stratification

Gravity, \( g = -\nabla \Phi \), is specified in terms of a potential \( \Phi \). In slab geometry, \( \Phi = \Phi(z) \), we have \( g = (0, 0, g_z) \) and \( g_z = -d\Phi/dz \).

Use `grav_profile='const'` together with `gravz=-1` to get

\[
\Phi = (z - z_\infty)(-g_z), \quad (-g_z) > 0.
\]

Use `grav_profile='linear'` to get

\[
\Phi = \frac{1}{2}(z^2 - z_\infty^2)\nu_g^2, \quad g_z = -\nu_g^2 z
\]

where \( \nu_g \) is the vertical epicyclic frequency. For a Keplerian accretion disc, \( \nu_g = \Omega \). For galactic discs, \( \nu_g = 0.5\Omega \) is representative of the solar neighborhood.

The value of \( z_\infty \) is determined such that \( \rho = \rho_0 \) and \( c_s^2 = c_{s0}^2 \) at \( z = z_{\text{ref}} \). This depends on the values of \( \gamma \) and the polytropic index \( m \) (see below).

#### C.5.1 Isothermal atmosphere

Here we want \( c_s = c_{s0} = \text{const} \). Using `initlnrho='isothermal'` means

\[
\ln \frac{\rho}{\rho_0} = -\gamma \frac{\Phi}{c_{s0}^2}.
\]

The entropy is then initialized to

\[
\frac{s}{c_p} = (\gamma - 1) \frac{\Phi}{c_{s0}^2}.
\]

In order that \( \rho = \rho_0 \) and \( c_s^2 = c_{s0}^2 \) at \( z = z_{\text{ref}} \), we have to choose \( z_\infty = z_{\text{ref}} \).
C.5.2 Polytropic atmosphere

For a polytropic equation of state, \( p = K \rho^\Gamma \), where generally \( \Gamma \neq \gamma \), we can write

\[
- \nabla h + T \nabla s = - \frac{1}{\rho} \nabla p = - \nabla \left( \frac{\Gamma K}{\Gamma - 1} \rho^{\Gamma - 1} \right) = - \nabla \hat{h},
\]

(138)

where we have introduced a pseudo enthalpy \( \hat{h} \) as

\[
\hat{h} = \frac{\Gamma K}{\Gamma - 1} \rho^{\Gamma - 1} = \left[ \frac{1 - \frac{1}{\gamma}}{1 - \frac{1}{\Gamma}} \right] h.
\]

(139)

Obviously, for \( \Gamma = \gamma \), the pseudo enthalpy \( \hat{h} \) is identical to \( h \) itself. Instead of specifying \( \Gamma \), one usually defines the polytropic index \( m = \frac{1}{\Gamma - 1} \). Thus, \( \Gamma = 1 + \frac{1}{m} \), and

\[
\hat{h} = (m+1) \left( 1 - \frac{1}{\gamma} \right) h
\]

(140)

This is consistent with a fixed entropy dependence, where \( s \) only depends on \( \rho \) like

\[
\frac{s}{c_p} = \left( \frac{\Gamma}{\gamma} - 1 \right) \ln \frac{\rho}{\rho_0},
\]

(141)

and implies that

\[
\ln \left( \frac{c_s^2}{c_{s0}^2} \right) = (\Gamma - 1) \ln \frac{\rho}{\rho_0}.
\]

(142)

For hydrostatic equilibrium we require \( \hat{h} + \Phi = \hat{h}_0 = \text{const} \). For gravity potentials that vanish at infinity, we can have \( \hat{h}_0 \neq 0 \), i.e. a finite pseudo enthalpy at infinity. For \( g_z = -1 \) or \( g_z = -z \), this is not the case, so we put \( h_0 = 0 \), and therefore \( h = -\Phi \). Using \( c_s^2 = (\gamma - 1)h \) together with (140) we find

\[
c_s^2 = -\frac{\gamma}{m+1} \Phi.
\]

(143)

In order that \( \rho = \rho_0 \) and \( c_s^2 = c_{s0}^2 \) at \( z = z_{\text{ref}} \), we have to choose (remember that \( g_z \) is normally negative!)

\[
z_\infty = z_{\text{ref}} + (m+1) \frac{c_{s0}^2}{\gamma(-g_z)} \quad \text{for grav\_profile}='\text{const}',
\]

(144)

and

\[
z_{\infty}^2 = z_{\text{ref}}^2 + (m+1) \frac{c_{s0}^2}{\frac{2}{\gamma} \nu_g \nu_s^2} \quad \text{for grav\_profile}='\text{linear}'.
\]

(145)

Thus, when using initlnrho='polytropic\_simple' we calculate

\[
\ln \left( \frac{c_s^2}{c_{s0}^2} \right) = \ln \left[ -\frac{\gamma \Phi}{(m+1)c_{s0}^2} \right]
\]

(146)

and so the stratification is given by

\[
\ln \frac{\rho}{\rho_0} = m \ln \frac{c_s^2}{c_{s0}^2}, \quad \frac{s}{c_p} = \left( \frac{\Gamma}{\gamma} - 1 \right) m \ln \frac{c_s^2}{c_{s0}^2}.
\]

(147)
C.5.3 Changing the stratification

Natural: measure length in units of $c_s^2/g_z$. Can increase stratification by moving $z_{\text{top}}$ close to $z_\infty$ or, better still, keeping $z_{\text{top}} = 0$ and moving $z_{\text{bot}} \to -\infty$. Disadvantage: in the limit of weak stratification, the box size will be very small (in nondimensional units).

Box units: measure length in units of $d$. Can increase stratification by increasing $g_z$ to $g_{\text{max}}$, which can be obtained by putting $z_{\text{top}} = z_\infty$ in (144), so

$$g_{\text{max}} = \frac{m+1}{\gamma} \frac{c_{s0}^2}{z_{\text{top}} - z_{\text{ref}}}.$$ (148)

For $m = 1$, $\gamma = 5/3$, $z_{\text{top}} = 1$, and $z_{\text{ref}} = 0$, for example, we have $g_{\text{max}} = 6/5 = 1.2$.

Gravitational box units: measure speed in units of $\sqrt{g_z d}$. The limit of vanishing stratification corresponds to $c_s \to \infty$. This seems optimal if we want to approach the Boussinesq case.

In Hurlburt et al. (1984), $z$ increased downward and the atmosphere always terminated at $z = 0$. In order to reproduce their case most directly, we put $z_{\infty} = 0$ and consider only negative values of $z$. To reproduce their case with a density stratification of 1:1.5, we place the top of the model at $z = -2$ and the bottom at $z = -3$. In addition, the reference height, $z_{\text{ref}}$, is chosen to be at the top of the box, i.e. $z_{\text{ref}} = -2$. From Eq. (144) we have $c_{s0}^2 = \gamma (-g_z) (-z_{\text{ref}})/(m + 1)$. Using $(-g_z) = 1$ and $m = 1$ we find $c_{s0}^2 = \gamma$, so $c_{s0} = 1.291$ (for $\gamma = 5/3$). Values for other combinations are listed in Table 11.

<table>
<thead>
<tr>
<th>$\rho_{\text{bot}}/\rho_{\text{top}}$</th>
<th>$z_{\text{bot}}$</th>
<th>$z_{\text{top}}$</th>
<th>$c_{s0}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.5</td>
<td>3</td>
<td>2</td>
<td>1.291</td>
</tr>
<tr>
<td>3</td>
<td>1.5</td>
<td>0.5</td>
<td>0.645</td>
</tr>
<tr>
<td>6</td>
<td>1.2</td>
<td>0.2</td>
<td>0.408</td>
</tr>
<tr>
<td>11</td>
<td>1.1</td>
<td>0.1</td>
<td>0.289</td>
</tr>
<tr>
<td>21</td>
<td>1.05</td>
<td>0.05</td>
<td>0.204</td>
</tr>
</tbody>
</table>

C.5.4 The Rayleigh number

In Ref. [12] the Rayleigh number is defined as

$$Ra = \frac{gd^4}{\bar{\nu} \bar{\chi}} \left( -\frac{ds/c_p}{dz} \right)_{\text{hydrostat}},$$ (149)

where the (negative) entropy gradient was evaluated in the middle of the box for the associated hydrostatic reference solution, and $\bar{\chi} = K/(\bar{\rho}c_p)$ and either $\bar{\nu} = \nu$ (if $\nu$ was assumed constant) or $\bar{\nu} = \mu/\bar{\rho}$ (if $\mu$ was assumed constant). Note that $\bar{\rho}$ is the average mass in the box per volume, which is conserved. For a polytrope we have

$$\left( -\frac{ds/c_p}{dz} \right)_{\text{hydrostat}} = \left[ 1 - (m + 1) \left( 1 - \frac{1}{\gamma} \right) \right] \frac{1}{z_{\infty} - z_m},$$ (150)

where $z_m = (z_1 + z_2)/2$. This factor was also present in the definition of Hurlburt et al. [23], but their definition differs slightly from Eq. (149), because they normalized the
density not with respect to the average value (which is constant for all times), but with respect to the value at the top of the initial hydrostatic solution. Since the Rayleigh number is proportional to $\rho^2$, their definition included the extra factor $[(z_\infty - z_m)/d]^2$. Therefore

$$Ra_{HTM} = \left(\frac{z_\infty - z_m}{d}\right)^2 \left(\frac{\rho_{\text{top}}}{\bar{\rho}}\right)^2 Ra$$  \hspace{1cm} (151)

In the first model of Hurlburt et al. (1984), the Rayleigh number, $Ra_{HTM}$, was chosen to be 310 times supercritical, and the critical Rayleigh number was around 400, so $Ra_{HTM} = 1.25 \times 10^5$. In their model the density contrast was 1:1.5 and $m = 1$. This turns out to correspond to $Ra = 4.9 \times 10^4$, $F_{\text{bot}} = 0.0025$, and $K = 0.002$.

Another model that was considered by Hurlburt & Toomre (1988) had $Ra_{HTM} = 10^5$, a density contrast of 11, and had a vertical imposed magnetic field (Chandrasekhar number $Q = 72$). This corresponds to $Ra = 3.6 \times 10^8$, $K = 0.0011$, $F_{\text{bot}} = 0.0014$.

C.5.5 Entropy boundary condition

This discussion only applies to the case of convection in a slab. A commonly used lower boundary condition is to prescribe the radiative flux at the bottom, i.e. $F_{\text{bot}} = -KdT/dz$. Assuming that the density in the ghost zones has already been updated, we can calculate the entropy gradient from

$$F_{\text{bot}} = -K \frac{c_s^2}{c_p \gamma - 1} \left(\gamma - 1\right) \left(\frac{d \ln \rho}{dz} + \frac{d s/c_p}{dz}\right),$$  \hspace{1cm} (152)

which gives

$$\frac{d s/c_p}{dz} = -\frac{\gamma - 1}{\gamma} \left(\frac{F_{\text{bot}} c_p}{K c_s^2} + \frac{d \ln \rho}{dz}\right)$$  \hspace{1cm} (153)

for the derivative of the entropy at the bottom. This is implemented as the ‘c1’ boundary condition at the bottom.

C.5.6 Temperature boundary condition at the top

In earlier papers the temperature at the top was set in terms of the quantity $\xi_0$, which is the ratio of the pressure scale height relative to the depth of the unstable layer. Expressed in terms of the sound speed at the top we have

$$c_{s,\text{top}}^2 = \gamma \xi_0 gd.$$  \hspace{1cm} (154)

$$\xi_{s,\text{bot}}^2 = \left(\xi_0 + \frac{1}{m + 1}\right) \gamma gd.$$  \hspace{1cm} (155)

C.6 Potential-field boundary condition

The ‘pot’ [or currently rather the ‘pwd’] boundary condition for the magnetic vector potential implements a potential-field boundary condition in $z$ for the case of an $x$-$y$-periodic box. In this section, we discuss the relevant formulas and their implementation in the Pencil Code.
Table 12: Correspondence between $\xi_0$ and $c_{s,\text{bot}}^2$ in single layer polytropes.

<table>
<thead>
<tr>
<th>$\xi_0$</th>
<th>$c_{s,\text{bot}}^2$</th>
</tr>
</thead>
<tbody>
<tr>
<td>10.00</td>
<td>17.500</td>
</tr>
<tr>
<td>0.20</td>
<td>1.167</td>
</tr>
<tr>
<td>0.10</td>
<td>1.000</td>
</tr>
<tr>
<td>0.05</td>
<td>0.917</td>
</tr>
<tr>
<td>0.02</td>
<td>0.867</td>
</tr>
</tbody>
</table>

If the top boundary is at $z = 0$, the relevant potential field for $z > 0$ is given by

\[
\tilde{A}_x(k_x, k_y, z) = C_x(k_{xy}) e^{-\kappa z},
\]

\[
\tilde{A}_y(k_x, k_y, z) = C_y(k_{xy}) e^{-\kappa z},
\]

\[
\tilde{A}_x(k_x, k_y, z) = C_z(k_{xy}) e^{-\kappa z},
\]

where

\[
\tilde{A}_i(k_x, k_y, z) \equiv \int e^{-ik_{xy} \cdot x} A_i(x, y, z) \, dx \, dy
\]

is the horizontal Fourier transform with $k_{xy} = (k_x, k_y, 0)$, and $k = |k_{xy}|$. Note that this implies a certain gauge and generally speaking the $z$ dependence in Eq. (158) is completely arbitrary, but the form used here works well in terms of numerical stability.

At the very boundary, the potential field (156)–(158) implies

\[
\frac{\partial \tilde{A}}{\partial z} + \kappa \tilde{A} = 0,
\]

and, due to natural continuity requirements on the vector potential, these conditions also hold for the interior field at the boundary.

**Robin boundary conditions and ghost points** To implement a homogeneous Robin boundary condition, i.e. a condition of the form

\[
\frac{df}{dz} + \kappa f = 0
\]

using ghost points, we first write it as

\[
\frac{d}{dz} (f e^{\kappa z}) = 0
\]

and implement this as symmetry condition for the variable $\phi(z) \equiv f(z) e^{\kappa z}$:

\[
\phi_{N-j} = \phi_{N+j}, \quad j = 1, 2, 3
\]

(where $z_N$ is the position of the top boundary and $z_{N+1}, \ldots$ are the boundary points). In terms of $f$, this becomes

\[
f_{N+j} = f_{N-j} e^{-\kappa(z_{N+j}-z_{N-j})}.
\]

Note that although the exponential term in Eq. (164) looks very much like the exterior potential field (156)–(158), our ghost-zone values do not represent the exterior field – they are rather made-up values that allow us to implement a local boundary condition at $z = 0$. 
C.7 Planet solution in the shearing box

In order to test the setup for accretion discs and the sliding periodic shearing sheet boundary condition, a useful initial condition is the so-called planet solution of Goodman, Narayan, & Goldreich [20].

Assume $s = 0$ (isentropy), so the equations in 2-D are

\begin{align}
  u_x u_{x,x} + (u_y^{(0)} + u_y) u_{x,y} &= 2\Omega u_y - h_{,x} \quad (165) \\
  u_x u_{y,x} + (u_y^{(0)} + u_y) u_{y,y} &= -(2 - q)\Omega u_x - h_{,y} \quad (166)
\end{align}

where $u_y^{(0)} = -q\Omega x$. Express $u$ in terms of a stream function, so $u = \nabla \times (\psi \hat{z})$, or

\begin{align}
  u_x = \psi_{,y}, \quad u_y = -\psi_{,x}. \quad (167)
\end{align}

**Ansatz for enthalpy**

\begin{align}
  h &= \frac{1}{2}\delta^2 \Omega^2 \left( R^2 - x^2 - \epsilon^2 y^2 - z^2 / \delta^2 \right) \quad (168) \\
  \psi &= -\frac{1}{2}\sigma \Omega \left( R^2 - x^2 - \epsilon^2 y^2 \right) - \frac{1}{2}q\Omega x^2 \quad (169)
\end{align}

This implies

\begin{align}
  u_x = \sigma \Omega \epsilon^2 y, \quad u_y = (q - \sigma)\Omega x \quad (170)
\end{align}

and $u_{x,x} = u_{y,y} = 0$. Inserting into Eqs (165) and (166) yields

\begin{align}
  (-q + q - \sigma)\sigma \epsilon^2 &= 2(q - \sigma) + \delta^2 \quad (171) \\
  \sigma(q - \sigma) &= -(2 - q)\sigma + \delta^2 \quad (172)
\end{align}

where we have already canceled common $\Omega^2$ factors in both equations and common $\epsilon^2$ factors in the last equation. Simplifying both equations yields

\begin{align}
  -\sigma^2 \epsilon^2 &= 2(q - \sigma) + \delta^2 \quad (173) \\
  -\sigma^2 &= -2\sigma + \delta^2 \quad (174)
\end{align}

The second equation yields

\begin{align}
  \delta^2 &= (2 - \sigma)\sigma \quad (175)
\end{align}

and subtracting the two yields

\begin{align}
  \sigma^2 &= 2q / (1 - \epsilon^2) \quad (176)
\end{align}

**Table 13:** Dependence of $\epsilon$ and $\delta$ on $\epsilon$.

<table>
<thead>
<tr>
<th>$\epsilon$</th>
<th>$\sigma$</th>
<th>$\delta$</th>
</tr>
</thead>
<tbody>
<tr>
<td>0.1</td>
<td>1.74</td>
<td>0.67</td>
</tr>
<tr>
<td>0.2</td>
<td>1.77</td>
<td>0.64</td>
</tr>
<tr>
<td>0.3</td>
<td>1.82</td>
<td>0.58</td>
</tr>
<tr>
<td>0.4</td>
<td>1.89</td>
<td>0.46</td>
</tr>
<tr>
<td>0.48</td>
<td>1.97</td>
<td>0.22</td>
</tr>
<tr>
<td>0.5</td>
<td>2.00</td>
<td>0.00</td>
</tr>
</tbody>
</table>
D Some specific boundary conditions

In this section, we formulate and discuss the implementation of some common boundary conditions in spherical and cylindrical coordinates.

D.1 Perfect-conductor boundary condition

This is a popular boundary condition for the magnetic field; it implies that

\[ B_n = 0 \]  \hspace{1cm} (177)

and

\[ E_t = 0 \]  \hspace{1cm} (178)

on the boundary, where the subscript \( n \) denotes the normal component, and \( E_t \) denotes the tangential components of the electric field.

In Cartesian geometry, these conditions can be implemented by setting the two tangential components of the vector potential \( A \) to zero on the boundary. It is easy to see that this also works in arbitrary curvilinear coordinates.

In particular, for spherical coordinates on a radial boundary we must have

\[ r \sin \theta B_r = \partial_\theta (\sin \theta A_\phi) - \partial_\phi A_\theta = 0 \].  \hspace{1cm} (179)

This can be achieved by setting

\[ A_\phi = A_\theta = 0 \]  \hspace{1cm} (180)

everywhere on the boundary. Note that this does not impose any condition on the radial component of the vector potential.

Next, in spherical coordinates on a boundary with constant \( \theta \), we must have

\[ B_\theta = \frac{1}{r \sin \theta} \partial_\phi A_r - \frac{1}{r} \partial_r (r A_\phi) = 0 \].  \hspace{1cm} (181)

Again this can be achieved by \( A_r = A_\phi = 0 \).

D.2 Stress-free boundary condition

On an impenetrable, stress-free boundary, we have

\[ u_n = 0 \],  \hspace{1cm} (182)

and the shear stress components \( S_{nt} \) must vanish for any tangential direction \( t \). At the radial boundary, the relevant components of the strain tensor (required to vanish at the boundary) are:

\[ S_{r\theta} = \frac{1}{r} \partial_\theta u_r + r \partial_r \left( \frac{u_\theta}{r} \right) \],  \hspace{1cm} (183)

\[ S_{r\phi} = \frac{1}{r \sin \theta} \partial_\phi u_r + r \partial_r \left( \frac{u_\phi}{r} \right) \].  \hspace{1cm} (184)

Both of them vanish if we require

\[ u_r = 0, \quad \partial_r (u_\theta/r) = 0, \quad \partial_r (u_\phi/r) = 0 \].  \hspace{1cm} (185)
We implement this by requiring \( u_r \) to be antisymmetric and \( u_\theta/r \) and \( u_\phi/r \) to be symmetric with respect to the boundary.

The more general condition

\[
r^\alpha \partial_r (u_\theta/r^\alpha) = \partial_r u_\theta - \frac{\alpha}{r} u_\theta = 0
\]  

(186)

(where \( \alpha \) is a constant) can be implemented by requiring \( u_\theta/r^\alpha \) to be symmetric.

At a boundary \( \theta = \text{const} \), the stress-free boundary condition will take the form

\[
S_{r\theta} = \frac{1}{r} \partial_\theta u_r + r \partial_r \left( \frac{u_\theta}{r} \right) = 0 ,
\]

(187)

\[
S_{\theta\phi} = \frac{1}{r \sin \theta} \partial_\phi u_\theta + \sin \theta \partial_\theta \left( \frac{u_\phi}{r \sin \theta} \right) = 0 .
\]

(188)

With \( u_\theta = 0 \), the first condition gives \( \partial_\theta u_r = 0 \), i.e. we require \( u_r \) to be symmetric with respect to the boundary. The second condition requires

\[
\frac{\sin \theta}{r} \partial_\theta \left( \frac{u_\phi}{\sin \theta} \right) = 0
\]

(189)

and is implemented by requiring \( u_\phi/\sin \theta \) to be symmetric.

### D.3 Normal-field-radial boundary condition

While unphysical, this boundary condition is often used as a cheap replacement for a potential-field condition for the magnetic field. It implies that the two tangential components of the magnetic field are zero at the boundary, while the normal component is left unconstrained.

At a radial boundary, this gives:

\[
B_{\theta} = \frac{1}{r \sin \theta} \partial_\phi A_r - \frac{1}{r} \partial_r (r A_\phi) = 0 ,
\]

(190)

\[
B_{\phi} = \frac{1}{r} \partial_r (r A_\theta) - \frac{1}{r} \partial_\theta A_r = 0 .
\]

(191)

Which are satisfied by setting

\[
A_r = 0 , \quad \partial_r (r A_\theta) = 0 , \quad \partial_r (r A_\phi) = 0 ,
\]

(192)

and these are implemented by requiring \( A_r \) to be antisymmetric, and \( r A_\theta \) and \( r A_\phi \) to be symmetric.

On a boundary \( \theta = \text{const} \), we have

\[
r \sin \theta B_r = \partial_\theta (\sin \theta A_\phi) - \partial_\phi A_\theta = 0 ,
\]

(193)

\[
r B_\phi = \partial_r (r A_\theta) - \partial_\theta A_r = 0
\]

(194)

which can be achieved by setting

\[
\partial_\theta A_r = 0 , \quad A_\theta = 0 , \quad \partial_\theta (\sin \theta A_\phi) = 0 .
\]

(195)

We thus require \( A_r \) and \( \sin \theta A_\phi \) to be symmetric, and \( A_\theta \) to be antisymmetric.
High-frequency filters

Being high order, PENCIL CODE has much reduced numerical dissipation. In order to perform inviscid simulations, high-frequency filters can be used to provide extra dissipation for modes approaching the Nyquist frequency. Usual Laplacian viscosity $\nu \nabla^2 u$ is equivalent to a multiplication by $k^2$ in Fourier space, where $k$ is the wavenumber. Another tool is hyperviscosity, which replaces the $k^2$ dependency by a higher power-law, $k^n$, $n \geq 2$. The idea behind it is to provide large dissipation only where it is needed, at the grid scale (high $k$), while minimizing it at the largest scales of the box (small $k$). In principle, one can use as high $n$ as desired, but in practice we are limited by the order of the code. A multiplication by $k^n$ is equivalent to an operator $\nabla^n$ in real space. As PENCIL CODE is of sixth order, three ghost cells are available in each direction, thus the sixth-order derivative is the highest we can compute. The hyperdissipation we use is therefore $\nabla^6$, or $k^6$ is Fourier space. Figure 20 illustrates how such tool maximizes the inertial range of a simulation.

Simplified hyperdiffusivity has been implemented for many dynamical variables and can be found in the respective modules. A strict generalization of viscosity and resistivity to higher order is implemented in the modules 'hypervisc_strict_2nd' and 'hyperresi_strict_2nd'.

Hyperdiffusivity is meant purely as a numerical tool to dissipate energy at small scales and comes with no guarantee that results are convergent with regular second order dissipation. See Haugen & Brandenburg (2004) for a discussion. In fact, large-scale dynamo action is known to be seriously altered in simulations of closed systems where magnetic helicity is conserved: this results in prolonged saturation times and enhanced saturation amplitudes (Brandenburg & Sarson 2002).

E.1 Conservative hyperdissipation

It is desirable to have this high-frequency filter obeying the conservation laws. So, for density we want a mass conserving term, for velocities we want a momentum conserving term, for magnetic fields we want a term conserving magnetic flux, and for entropy we want an energy conserving term. These enter as hyperdiffusion, hyperviscosity, hyperresistivity, and hyper heat conductivity terms in the evolution equations. To ensure conservation under transport, they must take the form of the divergence of the flux $\mathbf{J}$ of the quantity $\psi$, so that Gauss theorem applies and we have

$$\frac{\partial \psi}{\partial t} + \nabla \cdot \mathbf{J} = 0 \quad (196)$$

For density, the flow due to mass diffusion is usually taken as the phenomenological Fick’s Law

$$\mathbf{J} = -D \nabla \rho \quad (197)$$

i.e., proportional to the density gradient, in the opposite direction. This leads to the usual Laplacian diffusion

$$\frac{\partial \rho}{\partial t} = D \nabla^2 \rho \quad (198)$$
under the assumption that the diffusion coefficient $D$ is isotropic. Higher order hyperdiffusion of order $2n$ involves a generalization of Eq. (197), to

$$\mathbf{J}^{(n)} = (-1)^n D^{(n)} \nabla^{2n-1} \rho.$$  \hspace{1cm} (199)

In our case, we are interested in the case $n = 3$, so that the hyperdiffusion term is

$$\frac{\partial \rho}{\partial t} = D^{(3)} \nabla^{6} \rho.$$  \hspace{1cm} (200)

The hyperdiffusion coefficient $D^{(3)}$ can be calculated from $D$ assuming that at the Nyquist frequency the two formulations (198) and (200) yield the same quenching. Considering a wave as a Fourier series in one dimension ($x$), one element of the series is expressed as

$$\psi_k = A e^{i(kx+\omega t)}.$$  \hspace{1cm} (201)
Plugging it into the second order diffusion equation (198) we have the dispersion condition \(i\omega = -Dk^2\). The sixth order version (200) yields \(i\omega = -D^{(3)}k^6\). Equating both we have \(D^{(3)} = Dk^{-4}\). This condition should hold at the grid scale, where \(k = \pi/\Delta x\), therefore

\[
D^{(3)} = D\left(\frac{\Delta x}{\pi}\right)^4
\]

(202)

For the magnetic potential, resistivity has the same formulation as mass diffusion

\[
\frac{\partial A}{\partial t} = -\eta \nabla \times B = \eta \nabla^2 A,
\]

(203)

where we used the Coulomb gauge \(\nabla \cdot A = 0\). The algebra is the same as above, also yielding \(\eta^{(3)} = \eta(\Delta x/\pi)^4\). For entropy, the heat conduction term is

\[
\frac{\partial S}{\partial t} = \frac{1}{\rho T} \nabla \cdot (K\nabla T),
\]

(204)

and requiring that \(K\) be constant, we substitute it by

\[
\frac{\partial S}{\partial t} = \frac{K^{(3)}}{\rho T} \nabla^6 T.
\]

(205)

also with \(K^{(3)} = K(\Delta x/\pi)^4\).

### E.2 Hyperviscosity

Viscosity has some caveats where subtleties apply. The difference is that the momentum flux due to viscosity is not proportional to the velocity gradient, but to the rate-of-strain tensor

\[
S_{ij} = \frac{1}{2} \left( \frac{\partial u_i}{\partial x_j} + \frac{\partial u_j}{\partial x_i} - \frac{2}{3} \delta_{ij} \nabla \cdot \mathbf{u} \right),
\]

(206)

which only allows the viscous acceleration to be reduced to the simple formulation \(\nu \nabla^2 \mathbf{u}\) under the condition of incompressibility and constant dynamical viscosity \(\mu = \nu \rho\). Due to this, the general expression for conservative hyperviscosity involves more terms. In some cases, it is no great overhead, but for others, simpler formulations can be applied.

#### E.2.1 Conservative case

In the general case, the viscous acceleration is

\[
f_{\text{visc}} = \rho^{-1} \nabla \cdot (2\rho \nu \mathbf{S})
\]

(207)

So, for the hypertensive force, we must replace the rate-of-strain tensor by a high order version

\[
f_{\text{visc}}^{\text{(hyper)}} = \rho^{-1} \nabla \cdot \left(2\rho \nu_n \mathbf{S}^{(n)}\right)
\]

(208)

where the \(n\)th-order rate of strain tensor is

\[
\mathbf{S}^{(n)} = (-\nabla^2)^{n-1} \mathbf{S}.
\]

(209)
For the $n = 3$ case it is

$$S^{(3)}_{ij} = \frac{1}{2} \left( \frac{\partial^5 u_j}{\partial x_i^5} + \frac{\partial^4 u_i}{\partial x_j^4} \right) - \frac{1}{3} \frac{\partial^4}{\partial x_i^4} \left( \nabla \cdot u \right).$$  \hspace{1cm} (210)

Plugging it into Eq. (208), and assuming $\mu_3 = \rho \nu_3 = \text{const}$

$$f_{\text{visc}}^{(\text{hyper})} = \nu_3 \left( \nabla^6 u + \frac{1}{3} \nabla^4 (\nabla (\nabla \cdot u)) \right).$$  \hspace{1cm} (211)

For $\nu_3 = \text{const}$, we have to take derivatives of density as well

$$f_{\text{visc}}^{(\text{hyper})} = \nu_3 \left( \nabla^6 u + \frac{1}{3} \nabla^4 (\nabla (\nabla \cdot u)) + 2 S^{(3)} \cdot \nabla \ln \rho \right)$$  \hspace{1cm} (212)

E.2.2 Non-conservative cases

Equations (211) and (212) explicitly conserve linear and angular momentum. Although desirable properties, such expressions are cumbersome and numerically expensive, due to the fourth order derivatives of $\nabla (\nabla \cdot u)$.

This term, however, is only important when high compressibility is present (since it depends on the divergence of $u$). In practice we drop this term and use a simple hyper-viscosity

$$f_{\text{visc}} = \begin{cases} 
\nu_3 \nabla^6 u & \text{if } \mu = \text{const} \\
\nu_3 \left( \nabla^6 u + 2 S^{(3)} \cdot \nabla \ln \rho \right) & \text{if } \nu = \text{const}
\end{cases}$$  \hspace{1cm} (213)

Notice that this can indeed be expressed as the divergence of a simple rate-of-strain tensor

$$S^{(3)}_{ij} = \frac{\partial^5 u_i}{\partial x_j^5},$$  \hspace{1cm} (214)

so it does conserve linear momentum. It does not, however, conserve angular momentum, since the symmetry of the rate-of-strain tensor was dropped. Thus, vorticity sinks and sources may be spuriously generated at the grid scale.

A symmetric tensor can be computed, that conserves angular momentum and can be easily implemented

$$S_{ij} = \frac{1}{2} \left( \frac{\partial^5 u_i}{\partial x_j^5} + \frac{\partial^5 u_j}{\partial x_i^5} \right)$$  \hspace{1cm} (215)

This tensor, however, is not traceless, and therefore accurate only for weak compressibility. It should work well if the turbulence is subsonic. Major differences are not expected, since the spectral range in which hyperviscosity operates is very limited: as a numerical tool, only its performance as a high-frequency filter is needed. This also supports the usage of the highest order terms only, since these are the ones that provide quenching at high $k$. Momentum conservation is a cheap bonus. Angular momentum conservation is perhaps playing it too safe, at great computational expense.
E.2.3 Choosing the coefficient

When changing the resolution, one wants to keep the grid Reynolds number, here defined as

\[ \text{Re}_{\text{grid}} = \frac{u_{\text{rms}}}{(\nu_n k_{N_Y}^{2n-1})} \]  \hfill (216)

approximately constant. Here, \( k_{N_Y} = \pi/\delta x \) is the Nyquist wavenumber and \( \delta x \) is the mesh spacing. Thus, when doubling the number of meshpoints, we can decrease the viscosity by a factor of about \( 2^5 = 32 \) (Haugen & Brandenburg 2004). This shows that hyperviscosity can allow a dramatic increase of the Reynolds number based on the scale of the box.

By choosing \( \text{idiff='hyper3\_mesh'} \) in \text{density\_run\_pars} the hyperdiffusion for density is being set automatically in a mesh-independent way. A hyper-mesh Reynolds number of 30 corresponds to a coefficient \( \text{diffrho\_hyper3\_mesh=2} \) if \text{maxadvec} is about 1, but in practice we need a bit more (5 is currently the default).

E.2.4 Turbulence with hyperviscosity

When comparing hyperviscous simulations with non-hyperviscous ones, it turns out that the Reynolds number at half the Nyquist frequency is usually in the range 5–7, i.e.

\[ \text{Re}_{\text{half-grid}} = \frac{u_{\text{rms}}}{[\nu_n (k_{N_Y}/2)^{2n-1}]} \approx 5–7 \]  \hfill (217)

The following table gives some typical values used in simulations with forcing wavenumber \( k_f = 1.5 \) and a forcing amplitude of \( f_0 = 0.02 \). If hyperdiffusion \( D_3 \) is used in the continuity equation, the corresponding values are about 30 times smaller than those of \( \nu_3 \); see Table 14.

<table>
<thead>
<tr>
<th>( N )</th>
<th>( \nu_1 )</th>
<th>( \nu_2 )</th>
<th>( \nu_3 )</th>
<th>( D_3 )</th>
<th>( \kappa_2^{\text{CFL}=0.4} )</th>
<th>( \kappa_2^{\text{CFL}=0.9} )</th>
</tr>
</thead>
<tbody>
<tr>
<td>16</td>
<td>( 1 \times 10^{-2} )</td>
<td>( 3 \times 10^{-4} )</td>
<td>( 2 \times 10^{-5} )</td>
<td>( 6 \times 10^{-7} )</td>
<td>( 7 \times 10^{-4} )</td>
<td>( 1 \times 10^{-4} )</td>
</tr>
<tr>
<td>32</td>
<td>( 5 \times 10^{-3} )</td>
<td>( 4 \times 10^{-5} )</td>
<td>( 6 \times 10^{-7} )</td>
<td>( 2 \times 10^{-8} )</td>
<td>( 1 \times 10^{-6} )</td>
<td>( 2 \times 10^{-5} )</td>
</tr>
<tr>
<td>64</td>
<td>( 2 \times 10^{-3} )</td>
<td>( 5 \times 10^{-6} )</td>
<td>( 2 \times 10^{-8} )</td>
<td>( 6 \times 10^{-10} )</td>
<td>( 2 \times 10^{-7} )</td>
<td>( 3 \times 10^{-6} )</td>
</tr>
<tr>
<td>128</td>
<td>( 1 \times 10^{-3} )</td>
<td>( 6 \times 10^{-7} )</td>
<td>( 6 \times 10^{-10} )</td>
<td>( 2 \times 10^{-11} )</td>
<td>( 3 \times 10^{-8} )</td>
<td>( 4 \times 10^{-7} )</td>
</tr>
<tr>
<td>256</td>
<td>( 5 \times 10^{-4} )</td>
<td>( 8 \times 10^{-8} )</td>
<td>( 2 \times 10^{-11} )</td>
<td>( 6 \times 10^{-13} )</td>
<td>( 4 \times 10^{-9} )</td>
<td>( 5 \times 10^{-8} )</td>
</tr>
<tr>
<td>512</td>
<td>( 2 \times 10^{-4} )</td>
<td>( 1 \times 10^{-8} )</td>
<td>( 6 \times 10^{-13} )</td>
<td>( 2 \times 10^{-14} )</td>
<td>( 5 \times 10^{-10} )</td>
<td>( 6 \times 10^{-9} )</td>
</tr>
<tr>
<td>1024</td>
<td>( 1 \times 10^{-4} )</td>
<td>( 1 \times 10^{-9} )</td>
<td>( 2 \times 10^{-14} )</td>
<td>( 6 \times 10^{-16} )</td>
<td>( 6 \times 10^{-11} )</td>
<td>( 8 \times 10^{-10} )</td>
</tr>
</tbody>
</table>

For comparison, we give in Table 14 estimates of the numerical 4th order hyperdiffusion resulting from a third order time step, for which we have

\[ \kappa_2^{\text{CFL}} = \frac{1}{24} \frac{u_{\text{rms}}}{(C_{\text{CFL}} \delta x)^3} \]  \hfill (218)

where \( C_{\text{CFL}} \) is the CFL parameter which is either 0.4 in the conservative case or 0.9 in the more progressive case.
E.3 Anisotropic hyperdissipation

As we want quenching primarily at the Nyquist frequency, hyperdissipation depends intrinsically on the resolution, according to Eq. (202). Because of this, isotropic hyperdissipation only gives equal quenching in all spatial directions if $\Delta x=\Delta y=\Delta z$, i.e., if the cells are cubic. For non-cubic cells, anisotropic dissipation is required as different directions may be better/worse sampled, thus needing less/more numerical smoothing. Such generalization is straightforward. For that, we replace Eq. (199) by

$$
\mathbf{J} = \left( \frac{\partial^5 \rho}{\partial x^5}, \frac{\partial^5 \rho}{\partial y^5}, \frac{\partial^5 \rho}{\partial z^5} \right),
$$

so that different diffusion operates in different directions. Since $D_x$, $D_y$, and $D_z$ are constants, the divergence of this vector is

$$
\nabla \cdot \mathbf{J} = D_x \frac{\partial^6 \rho}{\partial x^6} + D_y \frac{\partial^6 \rho}{\partial y^6} + D_z \frac{\partial^6 \rho}{\partial z^6}.
$$

The formulation for resistivity and heat conductivity are strictly the same. For viscosity it also assumes the same form if we consider the simple non-conservative rate-of-strain tensor (214).

Mathematically, these operations can be written compactly by noticing that the coefficients in Eq. (220) transform like diagonal tensors $\chi_{ij}^{(3)} = \chi_k^{(3)} \delta_{ijk}$, where $\delta_{ijk}$ is the unit diagonal third order tensor, $\chi^{(3)}$ is the vector containing the dissipative coefficients (diffusion, viscosity, resistivity, or heat conductivity) in $x$, $y$, and $z$, and summation over repeated indices applies.

Therefore, for a scalar quantity $\psi$ (density, any of the three components of the velocity or magnetic potential), we can write

$$
\frac{\partial \psi}{\partial t} = -\chi_{ij}^{(3)} \partial_i \partial_j^5 \psi = -\sum_q \chi_q^{(3)} \frac{\partial^6}{\partial x_q^6} \psi.
$$

E.4 Hyperviscosity in Burgers shock

![Figure 21: Left: Burgers shock from teach/PencilCode/material/BurgersShock (in the teaching material) with $-20 \leq x \leq 20$, $n_x = 64$ mesh points, $u_x = \pm 1$ on the two ends, $\nu = 0.4$ and either $\nu_3 = 0$ (solid line) or $\nu_3 = 0.05$ (dashed line). Right: similar to the left hand side, but with $\nu = 0$ and $\nu_3 = 0.05$ (dashed line), compared with the case $\nu = 0.4$ and $\nu_3 = 0$ (solid line).](image)
Hyperviscosity has the unfortunate property of introducing (numerically stable) wiggles, even if one just adds a little bit of hyperviscosity to a run with normal viscosity; see left hand side of Fig. [21]. Running with just hyperviscosity give strong wiggles.

E.5 Time-dependent viscosity and magnetic diffusivity

In connection with decaying hydrodynamic and MHD turbulence studies, [32] noted that the equations are invariant under rescaling of space and time coordinates, along with a corresponding rescaling of the other dependent variables.

\[
\begin{align*}
& t = \tau t', \quad x = \tau^q x', \quad \nu = \tau^{2q-1}\nu', \quad \eta = \tau^{2q-1}\eta', \\
& u = \tau^{q-1}u', \quad B = \tau^{q-1}B'.
\end{align*}
\] (222)

Inserting these variables into equation (44) and (46), the resulting equation in the primed quantities has the same form as the equations in their original formulation. This requires that \( \nu \propto \eta \propto t^r \) where \( r = 2q - 1 \). For \( q < 1/2 \), \( r \) is negative so \( t^r \) becomes singular for \( t \to 0 \). Therefore, we use in such cases

\[
\begin{align*}
\nu(t) &= \nu_0[\max(1, t/t_0)]^r, \\
\eta(t) &= \eta_0[\max(1, t/t_0)]^r,
\end{align*}
\] (223)

where \( t_0 \) is the time below which \( \nu \) and \( \eta \) are assumed fixed; see Ref. [8] for details. This can be accomplished by putting

```bash
&viscosity_run_pars
  ivisc='nu-tdep', nu=1e-3, nu_tdep_t0=.1, nu_tdep_exponent=-.43
  lvisc_nu_tdep_t0_norm=T
/
```

and

```bash
&magnetic_run_pars
  iresistivity='eta-tdep', eta=1e-3, eta_tdep_t0=.1, eta_tdep_exponent=-.43
  lresi_eta_tdep_t0_norm=T
/
```

For backward compatibility, the default is \( lvisc_nu_tdep_t0_norm=F \) and likewise \( lresi_eta_tdep_t0_norm=F \). This means that instead of Eq. (223), we use \( \nu(t) = \nu_0[\max(t, t_0)]^r \) and \( \eta(t) = \eta_0[\max(t, t_0)]^r \), which has the disadvantage that then \( \nu_0 \) and \( \eta_0 \) have funny dimensions.
F Special techniques

F.1 After changing REALPRECISION

To continue working in double precision (REALPRECISION=double), you just say lread_from_other_prec=T in run_pars. Afterwards, remember to put lread_from_other_prec=F. If continuation is done in a new run directory, first execute start.csh there and then copy the files var.dat (and if present global.dat) from the old to the new directory, using pc-copyvar.

F.2 Remeshing (regridding)

Currently (29.07.19) two options are available for taking an existing run with a given resolution and processor layout and continuing with a changed resolution or different layout of processors. These apply for the original fortran binary data format used by the Pencil Code. The original version is described in section F.2.3, while a recent more versatile option employing Python to perform the task is described in section F.2.2.

The parallel hdf5 format for storing Pencil Code data has now been implemented to replace the less portable fortran unformatted data. As the hdf5 data is stored in a single file, the processor layout can be revised at any time during a run, without the need to change the data files at all. Changing resolution or grid dimensions is also more convenient, using appropriate interpolation tools. When scripts to automatically resize the grid and physical data for the full f-array become available, instructions on their use shall be included in section F.2.1.

F.2.1 Remeshing hdf5-formatted data

Instructions shall follow shortly . . .

F.2.2 Remeshing unformatted fortran binary data using Python

Ensure that $PENCIL_HOME/python has been added to your $PYTHON_PATH. Let us assume you have an existing run and you would like to continue an experiment from a mature state, but with

- higher (or lower) resolution,
- changes to the size of the numerical domain,
- change to the processor layout to improve efficiency and/or speed of the calculation,
- added/reduced variables included in the model,

or any combination involving at least one of the first three options in the list. The following procedure can be used to handle the first three options. The remeshed model with the ‘var.dat’ files obtained from an existing run can then also be used to advance with additional physics, following section F.4. As part of the remeshing procedure reduced physics could be obtained by omitting unwanted variables (not yet implemented).
1. First set up the new run. One option will be to navigate to the path of the existing run $\text{path-to-old-run}$ and apply the command

   $\text{path-to-old-run}> \text{pc_newrun \ path-to-new-run}$

$\text{path-to-new-run}$ is the full path of the new run or its relative path assuming it will be prefixed by ‘./’. Then navigate to $\text{path-to-new-run}$. Revise as required the files ‘\text{start.in}’, ‘\text{src/cparam.local}’ and ‘\text{src/Makefile.local}’ to match the parameters for the remeshed run. Note, any changes involving domain size will need to ensure continuity across any periodic boundaries. Then compile and start the new run

   $\text{path-to-new-run}> \text{pc_build}$
   $\text{path-to-new-run}> \text{pc_run \ start}$

or otherwise submit a batch script containing the call to ‘\text{start.csh}’.

2. Once the new run has been started successfully, create in the new run directory a python script called ‘\text{local_remesh.py}’ or similar, as described in the notes at the end of ‘\text{$\text{PENCIL_HOME/python/pencil/files/remesh.py}$}’. Add \text{import pencil as pc} at the start of the file.

3. To read the old data and then interpolate this onto the new grid as a complete f-array object \text{fnew}, add to the ‘\text{local_remesh.py}’ the lines

   \begin{verbatim}
   fnew=pc.interp_var(
       source_path=\text{path-to-old-run},
       target_path=\text{path-to-new-run},
       arrs=['uu','rho','lnrho','ss','aa',
             'shock','netheat','cooling'],
   )
   \end{verbatim}

By default this will read in the data from the old run ‘\text{var.dat}’ files. If another source file is required add the line \text{oldvar=$\text{VAR}$,} between the brackets, where $\text{VAR}$ is one of the snapshots with suffix ‘\text{VAR}’. Replace the default list of variables \text{arrs} with the variables used in the old run listed in f-array index order. For other options inspect ‘\text{$\text{PENCIL_HOME/python/pencil/files/remesh.py}$}’. Handling particles is not yet implemented, but should be possible with minor edits.

4. To write \text{fnew} to the new run ‘\text{var.dat}’ files add to ‘\text{local_remesh.py}’ the lines

   \begin{verbatim}
   pc.distribute_fort(
       fnew,
       \text{path-to-new-run},
   )
   \end{verbatim}

5. To execute the script run

   $\text{path-to-new-run}> \text{python \ local_remesh.py}$

To preserve long term the remeshed data

   $\text{path-to-new-run}> \text{pc_copyvar v 0 -e}$
6. Once the remeshing is completed edit as required ‘run.in’ and execute

```
$path-to-new-run> pc_run run
```

or submit batch script as appropriate.

**F.2.3 Remeshing unformatted fortran binary - original method**

[This should be written up in a better way and put somewhere else. But currently, remeshing is only available for the Pencil developers anyway.]

Suppose you have a directory run_64 with a 64³ run (running on \( N_0 = ny \times nz = 2 \times 1 \) CPUs) that you want to continue from ‘VAR1’ at 128³ (on \( ny \times nz = 4 \times 4 \) CPUs).

1. The remeshing code is part of the PENCIL CODE repository.

2. Create another run directory with current ‘VAR1’ as ‘var.dat’ (remesh.csh so far only works with ‘var.dat’):

```
unix> cd run_64
run_64> pc_newrun ../tmp_64 or new tmp_64
run_64> mkdir -p ../tmp_64/data or (cd ../tmp_64/; crtmp)
run_64> (cd ../tmp_64/data ; mkproc-tree \( N_0 \))
run_64> restart-new-dir-VAR ../tmp_64 1
```

3. Create the new run directory (linking the executables with -s):

```
run_64> cd ../tmp_64
tmp_64> pc_newrun -s ../run_128 or new run_128
tmp_64> vi ../run_128/src/cparam.local
  # set nxgrid=128, ncpus=16, nprocy=4
tmp_64> (cd ../run_128; crtmp; pc_setupsrc; make)
```

4. Setup and do remeshing

```
tmp_64> setup-remesh
tmp_64> vi remesh/common.local
  # set muly=2, mulz=4, remesh_par=2
tmp_64> (cd remesh; make)
tmp_64> vi remesh.in
  # Replace line by ../run_128
  # Answer ‘yes’
```

**F.3 Restarting with different I/O strategy**

One might want to switch the I/O strategy for a run, ongoingly to be continued by restarts, typically from one of the more traditional schemes, reading and writing FORTRAN binary or formatted data, to using the HDF5 data format. For this, include a
definition for IO_IN in Makefile.local, say IO_IN=io_dist, change the definition of IO, say into IO=io_hdf5, and recompile. The restarted run will write all data with the new I/O scheme and write an (empty) control file IO_LOCK which prevents the code from trying to read the data at the next restart still with the old I/O scheme, specified by IO_IN. A backswitch from HDF5 to binary format is also possible, but note that averages and slices will be continued to be written in HDF5 unless you recompile the code once more with IO_IN removed (to be improved).

F.4 Restarting from a run with less physics

First, prepare a new run directory with the new physics included. By new physics, we mean that the new run wants to read in more fields (e.g. magnetic fields, if the old run didn’t have magnetic fields).

Example for test fields:

1. Prepare ‘src/cparam.local’

   Add the following 2 fragments into the ‘cparam.local’ file. The first piece comes in the beginning and the second in the end of the file.

   ** AUTOMATIC CPARAM.INC GENERATION ****************************************
   ! Declare (for generation of cparam.inc) the number of f array
   ! variables and auxiliary variables added by this module
   ! Use MVAR to reserve the appropriate workspace for testfield_z.f90
   ! The MAUX number must be equally big and is used for uxb in the f-array.
   ! At the end of this file, njtest must be set such that 3*njtest=MVAR.
   !
   ! MVAR CONTRIBUTION 12
   ! MAUX CONTRIBUTION 12
   !
   !***************************************************************************
   !
   ! note that MVAR=MAUX=3*njtest must be obeyed
   !
   integer, parameter :: njtest=4
   !

2. Prepare ‘src/Makefile.local’

   Add the line TESTFIELD=testfield_z to the file. Finally, compile the code.

3. Prepare restart data

   Go into data directory of the new run and prepare the directory tree using, e.g., the command pc_mkproctree 16. [In principle this could be automatized, but it isn’t yet.]

   Next, go into old run directory and say restart-new-dir ../32c, if ‘../32c’ is the name of the new run directory. This procedure copies all the files from the processor tree, plus files like ‘param.nml’, but this file may need some manual modification (or you could just us one from another runs with the new physics included, which is definitely the simplest!!)
4. Prepare ‘run.in’

Set lread_oldsnap_notestfield=T in run_pars. This means (as the name says) that one reads an old snapshot that did not have test fields in it.

Reset boundary conditions and add stuff for the newly added fields, e.g., 
bcz='a:s', 'a', 'a:s', 'a2', 'a', 'a', 's', 'a', 'a', 's', 'a', 's', 'a', 's', 'a', 's', 'a', 's' in run_pars. If you don’t do this, you would effectively use periodic boundary conditions for the response to the test field, which is hardly correct once you set non-periodic boundary conditions for the other variables.

Add something like the following text fragments in the right position (after grav-run_pars and magnetic_run_pars, but before shear_run_pars and viscosity_run_pars).

&testfield_run_pars
  !linit_aatest=T, daainit=100.
  etestfield='B11-B22' 
  etatest=1e-4
  lsoca=F 
/

Make sure that the data above are correct. You may want to change the values of daainit or etatest.

If you now run, and if you didn’t fix the file ‘data/param.nml’ you might get something like the following error:

forrtl: severe (24): end-of-file during read, unit 1, file /wkspace/brandenb/pencil-code/axel/Poiseuille/2d/32c/data/pa

The reason for this is that it reads the old boundary data, but the corresponding array is too short. This includes stuff like FBCX1 to FBCX2_2, but it is still not enough. Therefore it is easiest to use the ‘data/param.nml’ file from another run. You may well just use one from a single processor run with a different mesh. But remember to fix the ‘start.in’ file by correcting the boundary conditions and adding things like

&testfield_init_pars
  luxb_as_aux=T
/

5. Prepare ‘print.in’, ‘xyaver.in’, and other obvious files such as ‘video.in’.

6. Once it works and is running, you must say explicitly

&run_pars

        lread_oldsnap_notestfield=F
/

because otherwise you won’t read in your precious test field data next time you restart the code! (If you instead just remove this line, it will remember lread_oldsnap_notestfield=T from the previous run, which is of course wrong!)

Comments: For large magnetic Reynolds numbers the solutions to the test-field equations can show a linear instability, which can introduce large fluctuations. In that case
it is best to reset the dependent test-field variable to zero in regular intervals. This is done by setting limit.aatest=T. Note that daainit=100 sets the reset interval to 100.

F.5 Restarting with particles from a run without them

If you want to restart from a run without particles to a run with them, you need to

1) Compile a run with particles,

RunWithParticles$ pc_build

2) Copy a VARN or var.dat into this new run directory. Say the old run is at the directory OldRun and you want to copy the var.dat of that run and restart with particles. You do

RunWithParticles$ pc_copyvar v v ../OldRun . -e

3) In the start.in init_pars of the new run, add the lines

lread_oldsnap = T
ireset_tstart = 0

4) Remove all calls to initial conditions (make all initlnrho, inituu, initss, initaa, etc ‘nothing’), and

5) Run pc_start.

The variable lread_oldsnap makes the code on start time read from the f-array of the snapshot, instead of the default, which is to initialize it with zeros. The necessity in step (4) to remove all calls to initial conditions is because otherwise the code would rewrite the content of the f-array with these initial conditions, or add them on top of the existing values of the snapshot.

The variable ireset_tstart when set to zero makes it read the timestamp of the old snapshot and restart from that time. The default is 2, which sets the timestamp back to zero.
G Runs and reference data

For reference purposes we document here some results obtained with various samples of the code.

G.1 Shock tests

G.1.1 Sod shock tube problem

Table 15: Combinations of $\rho$, $p$, and $s/c_p$ that are relevant for the Sod shock tube problem with constant temperature and different pressure ratios on the left and right hand sides of the shock.

<table>
<thead>
<tr>
<th>$\rho$</th>
<th>$p$</th>
<th>$s$</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.0</td>
<td>1.0</td>
<td>0.3065</td>
</tr>
<tr>
<td>0.1</td>
<td>0.1</td>
<td>1.2275</td>
</tr>
<tr>
<td>0.01</td>
<td>0.01</td>
<td>2.1486</td>
</tr>
</tbody>
</table>

G.1.2 Temperature jump

Table 16: Combinations of $c_s^2$, $p$, and $s/c_p$ that are relevant for the temperature shock problem with constant density, $\rho = 1$, and different temperature ratios on the left and right hand sides of the shock.

<table>
<thead>
<tr>
<th>$c_s^2$</th>
<th>$s$</th>
</tr>
</thead>
<tbody>
<tr>
<td>1.0</td>
<td>0.0</td>
</tr>
<tr>
<td>0.1</td>
<td>-2.3</td>
</tr>
<tr>
<td>0.01</td>
<td>-4.6</td>
</tr>
<tr>
<td>$10^{-4}$</td>
<td>-9.2</td>
</tr>
</tbody>
</table>

G.2 Random forcing function

A solenoidal random forcing function $f$ can be invoked by putting iforce='helical' in the forcing_run_pars namelist. This produces the forcing function $f$ of the form

$$f(x, t) = \text{Re}\{N f_{k(t)} \exp[i k(t) \cdot x + i \phi(t)]\},$$

(224)

where $k(t) = (k_x, k_y, k_z)$ is a time dependent wave vector, $x = (x, y, z)$ is position, and $\phi(t)$ with $|\phi| < \pi$ is a random phase. On dimensional grounds the normalization factor is chosen to be $N = f_0 c_s (k c_s / \delta t)^{1/2}$, where $f_0$ is a nondimensional factor, $k = |k|$, and $\delta t$ is the length of the timestep. The $\delta t^{-1/2}$ dependence ensures that the forcing, which is delta-correlated in time, is properly normalized such that the correlator of the forcing function is independent of the length of the time step, $\delta t$. We focus on the case where $|k|$ is around 5, and select at each timestep randomly one of the 350 possible vectors in $4.5 < |k| < 5.5$. We force the system with eigenfunctions of the curl operator,

$$f_k = \frac{ik \times (k \times e) - \sigma |k|(k \times e)}{\sqrt{1 + \sigma^2} k^2 \sqrt{1 - (k \cdot e)^2 / k^2}},$$

(225)
where \( e \) is an arbitrary unit vector needed in order to generate a vector \( k \times e \) that is perpendicular to \( k \). Note that \( |f_k|^2 = 1 \) and, for \( \sigma = 1 \), \( ik \times f_k = |k|f_k \), so the helicity density of this forcing function satisfies

\[
f \cdot \nabla \times f = |k|f^2 > 0 \quad \text{(for } \sigma = 1) \tag{226}\]

at each point in space. We note that since the forcing function is like a delta-function in \( k \)-space, this means that all points of \( f \) are correlated at any instant in time, but are different at the next timestep. Thus, the forcing function is delta-correlated in time (but the velocity is not). This is the forcing function used in Brandenburg (2001), Brandenburg & Dobler (2001), and other papers in that series.

For \( \sigma = 0 \), the forcing function is completely nonhelical and reduces to the simpler form

\[
f_k = (k \times e) / \sqrt{k^2 - (k \cdot e)^2}. \tag{227}\]

For \( 0 < |\sigma| < 1 \), the forcing function has fractional helicity, where \( \sigma \approx \langle \omega \cdot u \rangle / (k_f \langle u^2 \rangle) \); see Sect. 4.5 of Ref. [10]. In the code and the forcing_run_pars namelist, \( \sigma \) is called relhel.

In the code, the possible wavevectors are pre-calculated and stored in ‘k.dat’, which is being read in the beginning the code runs. To change the wavevectors (e.g. the typical value of \( k_f \), you need to change the file. In the directory ‘$PENCIL_HOME/samples/helical-MHDturb/K_VECTORS/’ there are several such files prepared:

\[
k10.dat \quad k1.dat \quad k2.dat \quad k3.dat \quad k5.dat \quad k15.dat \quad k27.dat \quad k30.dat \quad k4.dat \quad k8.dat
\]

and more can be prepared in IDL with the procedure ‘$PENCIL_HOME/samples/helical-MHDturb/idl/generate_kvectors.pro’ There is also more help in the ‘README’ file in ‘helical-MHDturb’.

In forcing_hel: if lcrosshel_forcing=T and if ltestfield_forcing=T.and.ltestflow_forcing=T, uu and aa (uu0 and aa0 for testfield*) are simultaneously forced, using the same filek.dat. Relative scaling by force1_scl for vector potentials and force2_scl for velocities (default: 1). Simplified “cross helicity forcing” is now activated by hydro_forcing.and.magnetic_forcing=.true. or by ltestfield_forcing.and.ltestflow_forcing=.true.; lcrosshel_forcing is now obsolete.

### G.3 Three-layered convection model

In Sect. 3 we have shown the early stages of the convection model located in ‘samples/conv-slab’. To arrive at fully developed convection, you will need to run the code for many more time steps. Figure 22 shows the vertical profiles of four basic quantities at time \( t = 50 \). Figure 23 shows the time evolution of rms and maximum velocity for the model for \( 0 < t < 50 \).

Figures 24 and 25 show vertical and horizontal sections for time \( t = 50 \).

### G.4 Magnetic helicity in the shearing sheet

To test magnetic helicity evolution in the shearing shear, we can choose as initial condition initaa='Beltrami-y' with amplaa=1. in magnetic_init_pars together with Sshear=-1. in shear_run_pars.
Figure 22: Like in Fig. 3 but at time $t = 50$.

Figure 23: Time evolution of rms and maximum velocity for the model 'samples/conv-slab'. Similar plots can be produced by running the IDL script 'ts.pro'.

Thus, in 'src/Makefile.local' we just use

```plaintext
MAGNETIC=magnetic
HYDRO=nohydro
EOS=noeos
DENSITY=nodensity
SHEAR=shear
VISCOSITY=noviscosity
```

and put

```plaintext
&init_pars
   cvsid='$Id$',
/
&magnetic_init_pars
   initaa='Beltrami-y', amplaa=1.
/
&shear_init_pars
```
Figure 24: Horizontal sections for $t = 50$. Top: velocity field. Bottom: entropy (color coded) and density (isocontours). Plots of this type can be produced by running the IDL script 'hsections.pro'.

in 'start.in' and, for example,

```plaintext
&run_pars
  cvsid='$Id$'
  nt=150000, it1=10, cdt=0.9, isave=50, itorder=3
  dsnap=100. dvid=5., ialive=1
/
&magnetic_run_pars
  eta=0.
/
&shear_run_pars
  Sshear=-1.
/
```

in 'run.in'. The output includes, among other things

- $\text{arms}(f10.7)$
- $\text{brms}(f12.7)$
- $\text{jrms}(f14.7)$
- $\text{abm}(f14.11)$
- $\text{jbm}(f14.7)$

The result is shown in Figure 26, where we show the wind-up of the magnetic field, which leads to a linear increase in the rms magnetic field strength until Ohmic diffusion begins to become important (top panel). During this time the magnetic helicity is conserved. With Ohmic diffusion, the decay of $\langle A \cdot B \rangle$ is well described by integrating $-2\eta\langle J \cdot B \rangle$ (indicated by “from rhs” in the second panel).
Figure 25: Vertical section $y = 0.516$ at $t = 50$. Top: velocity field. Bottom: entropy (color coded) and density (isocontours). Plots of this type can be produced by running the IDL scripts 'vsections.pro' or 'vsections2.pro').
Figure 26: Wind-up of the magnetic field leads to a linear increase in the rms magnetic field strength until Ohmic diffusion begins to become important (top panel). During this time the magnetic helicity is conserved. With Ohmic diffusion, the decay of $\langle A \cdot B \rangle$ is well described by integrating $-2\eta \langle J \cdot B \rangle$ (indicated by “from rhs” in the second panel).
H Numerical methods

H.1 Sixth-order spatial derivatives

Spectral methods are commonly used in almost all studies of ordinary (usually incompressible) turbulence. The use of this method is justified mainly by the high numerical accuracy of spectral schemes. Alternatively, one may use high order finite differences that are faster to compute and that can possess almost spectral accuracy. Nordlund & Stein [31] and Brandenburg et al. [15] use high order finite difference methods, for example fourth and sixth order compact schemes [27].

The sixth order first and second derivative schemes are given by

\[ f'_i = \frac{(-f_{i-3} + 9f_{i-2} - 45f_{i-1} + 45f_{i+1} - 9f_{i+2} + f_{i+3})}{(60\delta x)}, \]  
\[ f''_i = \frac{(2f_{i-3} - 27f_{i-2} + 270f_{i-1} - 490f_i + 270f_{i+1} - 27f_{i+2} + 2f_{i+3})}{(180\delta x^2)}, \]

(228)
(229)

In Fig. 27 we plot effective wavenumbers for different schemes. Apart from the different explicit finite difference schemes given above, we also consider a compact scheme of 6th order, which can be written in the form

\[ \frac{1}{3}f'_{i-1} + f'_i + \frac{1}{3}f'_{i+1} = \frac{(f_{i-2} - 28f_{i-1} + 28f_{i+1} - f_{i+2})}{(36\delta x)}, \]  
\[ \frac{2}{\Pi}f''_{i-1} + f''_i + \frac{2}{\Pi}f''_{i+1} = \frac{(3f_{i-2} + 48f_{i-1} - 102f_i + 48f_{i+1} + 3f_{i+2})}{(44\delta x^2)}. \]

(230)
(231)

for the first derivative, and

for the second derivative. As we have already mentioned in the introduction, this scheme involves obviously solving tridiagonal matrix equations and is therefore effectively non-local.

In the second panel of Fig. 27 we have plotted effective wavenumbers for second derivatives, which were calculated as

\[ (\cos kx)'_{\text{num}} = -k^2_{\text{eff}} \cos kx. \]

(232)

Of particular interest is the behavior of the second derivative at the Nyquist frequency, because that is relevant for damping zig-zag modes. For a second-order finite difference scheme \( k^2_{\text{eff}} \) is only 4, which is less than half the theoretical value of \( \pi^2 = 9.87 \). For fourth, sixth, and tenth order schemes this value is respectively 5.33, 6.04, 6.83. The last value is almost the same as for the 6th order compact scheme, which is 6.86. Significantly stronger damping at the Nyquist frequency can be obtained by using hyperviscosity, which Nordlund & Galsgaard (1995) treat as a quenching factor that diminishes the value of the second derivative for wavenumbers that are small compared with the Nyquist frequency. Accurate high order second derivatives (with no quenching factors) are important when calculating the current \( J \) in the Lorentz force \( J \times B \) from a vector potential \( A \) using \(-\mu_0J = \nabla^2A - \nabla\nabla \cdot A\). This will be important in the MHD calculations presented below.

---

\[18\] The fourth order compact scheme is really identical to calculating derivatives from a cubic spline, as was done in Ref. [31]. In the book by Collatz [16] the compact methods are also referred to as Hermitian methods or as Mehrstellen-Verfahren, because the derivative in one point is calculated using the derivatives in neighboring points.
High-order centered-difference convection simulations often show “wiggles” (Nyquist zigzag pattern) in $\ln \rho$, which are apparently caused by a velocity profile where the velocity approaches zero on the boundary or inside the box. This causes the density profile to be squeezed into a boundary layer where eventually numerical resolution is insufficient and, for centered differences, a spurious Nyquist signal is generated that almost instantaneously propagates into much of the whole box.

Even if the stagnation point is on the boundary (and enforced by the boundary conditions), this behavior is hardly influenced by the boundary conditions on $\ln \rho$ at all. A solution, however, is to apply upwinded derivative operators. The simplest upwind derivative is a finite-difference derivative operator where the point furthest downwind is excluded from the stencil. For $u > 0$, that means that instead of

$$f'_0 = -f_{-3} + 9f_{-2} - 45f_{-1} + 45f_{1} - 9f_{2} + f_{3} - \frac{\delta x^6 f^{(7)}}{140} = D^{(\text{cent},6)} + O(\delta x^6),$$

one takes

$$f'_0 = -2f_{-3} + 15f_{-2} - 60f_{-1} + 20f_{0} + 30f_{1} - 3f_{2} + \frac{\delta x^5 f^{(6)}}{60} = D^{(\text{up},5)} + O(\delta x^5).$$

A fourth-order upwind scheme (excluding two downwind points) would be

$$f'_0 = -f_{-3} + 6f_{-2} - 18f_{-1} + 10f_{0} + 3f_{1} - \frac{\delta x^4 f^{(5)}}{20} = D^{(\text{up},4)} + O(\delta x^4).$$

A simple one-dimensional test profile would be $u(x) = 1 - x^2$ on $x \in [-1, 1]$, which will accumulate more and more mass near the right boundary $x = 1$.

In two- or three-dimensional settings, the presence of stagnation points of X-type leads to the same configuration, this time with the possibility of a steady state (i.e. without accumulation of mass). Such stagnation points occur e.g. at the top of an upwelling, or at the bottom of a downdraft in convection simulations, where locally $u_z \propto z_X - z$. 

---

**Figure 27:** Effective wave numbers for first and second derivatives using different schemes. Note that for the second derivatives the sixth order compact scheme is almost equivalent to the tenth order explicit scheme. For the first derivative the sixth order compact scheme is still superior to the tenth order explicit scheme.
The effect of upwinding is mostly to stop the Nyquist perturbations from spreading away from the boundary or stagnation point. With the fourth-order formula they actually hardly ever develop.

The difference between central and fifth-order upwind derivative is

$$\left[D^{(up,5)} - D^{(cent,6)}\right]f_0 = \frac{-f_{-3} + 6f_{-2} - 15f_{-1} + 20f_0 - 15f_1 + 6f_2 - f_3}{60 \delta x} = -\frac{\delta x^5}{60}f^{(6)}_0. \quad (236)$$

In other words, 5th-order upwinding can be represented for any sign of $u$ as hyperdiffusion (Dobler et al. 2006):

$$-uf'_{(up,5th)} = -uf'_{(centr,6th)} + \frac{|u| \delta x^5}{60}f^{(6)}_0. \quad (237)$$

The advantage over adopting constant hyperdiffusion is that in the upwinding scheme hyperdiffusion is only applied where it is needed (i.e. where advection is happening, hence the factor $|u|$).

The form (237) also suggests an easy way to get ‘stronger’ upwinding: Rather than excluding more points in the downwind direction, we can simply treat the weight of the hyperdiffusion term as a free parameter $\alpha$:

$$-uf'_{(up,5th,\alpha)} = -uf'_{(centr,6th)} + \alpha |u| \delta x^5 f^{(6)}_0. \quad (238)$$

If $\alpha$ is large, this may affect the time step, but for $\alpha = 1/60$, the stability requirement for the hyperdiffusive term should always be weaker than the advective Courant criterion.

### H.3 The bidiagonal scheme for cross-derivatives

The old default scheme used for cross-derivatives of type $\partial^2 / (\partial x \partial y)$ used to read as follows:

```matlab
df=fac*( &
  270.*( f(11+1:12+1,m+1,n,k)-f(11-1:12-1,m+1,n,k) &
    +f(11-1:12-1,m-1,n,k)-f(11+1:12+1,m-1,n,k)) &
  -27.*( f(11+2:12+2,m+2,n,k)-f(11-2:12-2,m+2,n,k) &
     +f(11-2:12-2,m-2,n,k)-f(11+2:12+2,m-2,n,k)) &
  +2.*( f(11+3:12+3,m+3,n,k)-f(11-3:12-3,m+3,n,k) &
    +f(11-3:12-3,m-3,n,k)-f(11+3:12+3,m-3,n,k)) &
)
```

and is “visualized” in the left part of Fig. 28. It is way more efficient than the straightforward approach of first taking the $x$ and the $y$ derivative consecutively. (shown in the right part of Fig. 28).

Off-diagonal terms enter not only the diffusion terms through $\nabla \cdot \nabla \cdot u$ and $\nabla \cdot A$ terms, but also through the $J = \nabla \times \nabla \times A$ operator. The general formula is $J_i = A_{j,ij} - A_{i,jj}$, so in 2-D in the $xy$-plane we have

$$J_x = A_{x,xx} + A_{y,xy} - A_{x,xy} - A_{x,yy} = A_{y,xy} - A_{x,yy}, \quad (239)$$

$$J_y = A_{x,yx} + A_{y,yy} - A_{y,xy} - A_{y,yy} = A_{x,yx} - A_{y,xx}. \quad (240)$$

Figure 29 shows how the two schemes perform for the propagation of Alfvén waves,
H.4 The 2N-scheme for time-stepping

For time stepping, higher-order schemes are necessary in order to reduce the amplitude and phase errors of the scheme and, to some extent, to allow longer time steps. Usually

The initial condition (as implemented in subroutine alfven_xy) is

\[ u_z \sim \cos(k_x x + k_y y - \omega t) , \]
\[ A_x \sim +B_{0y} \sin(k_x x + k_y y - \omega t)/\omega , \]
\[ A_y \sim -B_{0x} \sin(k_x x + k_y y - \omega t)/\omega , \]

where \( \omega = k \cdot B_0 \). The figure shows that there is no clear advantage of either scheme, so the code uses the more efficient bidiagonal one.

Figure 28: Weights of bidiagonal scheme (left) and consecutive scheme (right) for mixed derivatives \( \partial^2/\partial x \partial y \). The numbers shown need to be divided by \( 720 \delta x \delta y \) for the bidiagonal and by \( 3600 \delta x \delta y \) for the consecutive scheme.

Figure 29: Alfvén wave for \( B_0 = (1, 2, 0) \) and \( k = (1, 2, 0) \) after \( t = 2\pi \). The wave travels in the direction of \( k \). Red symbols are for the bidiagonal scheme, black symbols show results obtained using the consecutive scheme. Already for 16\(^2\) mesh points there are no clear differences. For 8\(^2\) mesh points both schemes are equally imprecise regarding the phase error, but the amplitude error is still quite small (but this is mainly a property of the time stepping scheme).

\[
\begin{array}{cccccccc}
-2 & 0 & 0 & 0 & 0 & 0 & +2 \\
0 & +27 & 0 & 0 & 0 & -27 & 0 \\
0 & 0 & -270 & 0 & +270 & 0 & 0 \\
0 & 0 & 0 & 0 & 0 & 0 & 0 \\
0 & 0 & +270 & 0 & -270 & 0 & 0 \\
0 & -27 & 0 & 0 & 0 & +27 & 0 \\
+2 & 0 & 0 & 0 & 0 & 0 & -2
\end{array}
\]
such schemes require large amounts of memory. However, we here use the memory-effective $2N$-schemes that require only two sets of variables to be held in memory. Such schemes work for arbitrarily high order, although not all Runge-Kutta schemes can be written as $2N$-schemes \[35, 34\]. Consider the ordinary differential equation (ODE)

$$\dot{u} = F(u, t),$$

which can also be used as a prototype for a system of ODEs to be solved, like the ones obtained by spatial discretization of PDEs. The $2N$-schemes construct an approximation to the solution

$$u^{(n)} \equiv u(t_n)$$

according to the iterative procedure

$$w_i = \alpha_i w_{i-1} + \delta t F(u_{i-1}, t_{i-1}),$$

$$u_i = u_{i-1} + \beta_i w_i.$$  \hspace{1cm} (249)

(250)

For a three-step (RK3-2N) scheme we have $i = 1, \ldots, 3$. In order to advance the variable $u$ from $u^{(n)}$ at time $t^{(n)}$ to $u^{(n+1)}$ at time $t^{(n+1)} = t^{(n)} + \delta t$ we set in Eq. (250)

$$u_0 = u^{(n)} \quad \text{and, after the last step,} \quad u^{(n+1)} = u_3,$$  \hspace{1cm} (251)

with $u_1$ and $u_2$ being intermediate steps. In order to be able to calculate the first step, $i = 1$, for which no $w_{i-1} \equiv w_0$ exists, we have to require $\alpha_1 = 0$. Thus, we are left with 5 unknowns, $\alpha_2, \alpha_3, \beta_1, \beta_2,$ and $\beta_3$. Three conditions follow from the fact that the scheme be third order for linear equations, so we have to have two more conditions. One possibility is to choose the fractional times at which the right hand side is evaluated, for example $(0, 1/3, 2/3)$ or even $(0, 1/2, 1)$. Yet another possibility is to require that inhomogeneous equations of the form $\dot{u} = t^n$ with $n = 1$ and $2$ are solved exactly. The corresponding coefficients are listed in Table 17 and compared with those given by Williamson \[35\]. In practice all of them are about equally good when it comes to real applications, although we found the first one in Table 17 (‘symmetric’) marginally better in some simple test problems where an analytic solution was known. In Ref. \[5\] the accuracy of some nonlinear equations is tested.

Table 17: Coefficients for different $2N$-type third-order Runge-Kutta schemes. The coefficients $c_i$ (which are determined by the $\alpha_i, \beta_i$) give the time for each substep, $t_i = t_0 + c_i \delta t$

<table>
<thead>
<tr>
<th>scheme</th>
<th>$c_1$</th>
<th>$c_2$</th>
<th>$c_3$</th>
<th>$\alpha_2$</th>
<th>$\alpha_3$</th>
<th>$\beta_1$</th>
<th>$\beta_2$</th>
<th>$\beta_3$</th>
</tr>
</thead>
<tbody>
<tr>
<td>symmetric</td>
<td>0</td>
<td>1/3</td>
<td>2/3</td>
<td>-2/3</td>
<td>-1</td>
<td>1/3</td>
<td>1</td>
<td>1/2</td>
</tr>
<tr>
<td>(predictor/corrector)</td>
<td>0</td>
<td>1/2</td>
<td>1</td>
<td>-1/4</td>
<td>-4/3</td>
<td>1/2</td>
<td>2/3</td>
<td>1/2</td>
</tr>
<tr>
<td>inhomogeneous</td>
<td>0</td>
<td>15/32</td>
<td>4/9</td>
<td>-17/32</td>
<td>-32/27</td>
<td>1/4</td>
<td>8/9</td>
<td>3/4</td>
</tr>
</tbody>
</table>

H.5 Diffusive error from the time-stepping

In many cases we use centered first derivatives for the advection operator, so the resulting discretization errors are only of dispersive nature (proportional to odd derivatives). A diffusive error can be obtained from the discretization error of the time-stepping scheme. For the RK3-2N scheme we have

$$\left( \frac{df}{dt} \right)^{\text{exact}}_{\text{nth order}} = \left( \frac{df}{dt} \right)^{\text{nth order}}_{\text{exact}} + a_n \delta t^n \left( \frac{df^{n+1}}{dt^{n+1}} \right) + \ldots,$$  \hspace{1cm} (252)
where \( a_n = 1/(n+1)! = 0.5 \). In particular, for \( n = 1 \) we have \( a_1 = 1/2 = 0.2 \) and for \( n = 3 \) we have \( a_3 = 1/24 \approx 0.04 \). The advection operator leads to a diffusive error equal to \( a_1 \delta t(u \cdot \nabla)^2 \) for \( n = 1 \) and a hyperdiffusive error equal to \( a_3 \delta t^3(u \cdot \nabla)^4 \) for \( n = 3 \).

Substituting \( \delta t = c_{\text{CFL}} \delta x / |u| \), where \( c_{\text{CFL}} \) is Courant-Friedrich-Lewy constant, we have a diffusive error \( \nu \nabla^2 \) with negative \( \nu = -a_1 c_{\text{CFL}} |u| \delta x \) for \( n = 1 \), and a hyperdiffusive error \(-\nu_{\text{hyp}} \nabla^4\) with positive \( \nu_{\text{hyp}} = a_3 c_{\text{CFL}}^3 |u| \delta x^3 \) for \( n = 3 \). The fact that the hyperdiffusive error has a positive effective hyperdiffusivity is an important factor in the choice of this scheme.

To decide whether the effective hyperdiffusivity from the diffusive error is significant, we can compare with the error that would occur had we used a third-order upwinding scheme (Sect. H.2). In that case we would have an effective hyperdiffusive coefficient \( |u| \delta x / 12 \) that is \( 1/(12a_3 c_{\text{CFL}}^3) \approx 5.8 \) times larger than that from the time stepping scheme. In this sense, the hyperdiffusive error can be regarded as small.

Since the hyperdiffusive error is proportional to \(-\nabla^4\), we cannot directly compare with the physical diffusion which is proportional to \( \nabla^2 \). Therefore we define an effective viscosity as \( \nu_{\text{eff}} = \nu_{\text{hyp}} k^2_{\text{Ny}} \) with \( k_{\text{Ny}} = \pi / \delta x \) being the Nyquist wavenumber of the mesh of the domain covered by \( N \) mesh points. We define Reynolds number based on the Nyquist wavenumber as \( \text{Re}_{\text{Ny}} = |u| / \nu_{\text{eff}} k_{\text{Ny}} \), and find \( \text{Re} = -24/(\pi c_{\text{CFL}})^3 \approx 2.3 \) for our favorite choice \( c_{\text{CFL}} = 0.7 \). Thus, at the scale of the mesh, the effective Reynolds number is comparable to the value often obtained in simulations. However, in turbulence simulations the viscous cutoff wavenumber is usually 5–10 times smaller than \( k_{\text{Ny}} \), so the relevant Reynolds number at the viscous scale is then another 2–3 orders of magnitude larger and does therefore not impose a constraint in view of the physical viscosity that is applied in such calculations.

### H.6 Ionization

The specific entropy of each particle species (neutral hydrogen, electrons, protons and neutral helium) may be written as

\[
\frac{s_i}{s_0} = x_i \left( \ln \left[ \frac{1}{x_{\text{tot}}} \frac{\rho_i}{\rho} \left( \frac{T}{T_0} \right)^{3/2} \right] + \frac{5}{2} \right),
\]

where

\[
x_H = 1 - y, \quad x_e = x_p = y, \quad x_{\text{tot}} = 1 + y + x_{\text{He}}
\]

\[
s_0 = \frac{k_B}{\mu_{m_H}}, \quad T_0 = \frac{\chi_H}{k_B},
\]

and

\[
\rho_i = \mu_{m_H} \left( \frac{m_i \chi_H}{2\pi\hbar^2} \right)^{3/2}.
\]

The specific entropy of mixing is

\[
\frac{s_{\text{mix}}}{s_0} = -\sum_i x_i \ln \frac{x_i}{x_{\text{tot}}},
\]
Summing up everything, we get the total specific entropy

\[
\frac{s}{s_0} = \sum_i \frac{s_i}{s_0} + \frac{s_{\text{mix}}}{s_0} = \sum_i x_i \left( \ln \left( \frac{1}{x_i} \frac{\rho_i}{\rho} \right) \left( \frac{T}{T_0} \right)^{3/2} + \frac{5}{2} \right)
\]  
(258)

\[
= \sum_i x_i \ln \frac{\rho_i}{x_i} + x_{\text{tot}} \left( \ln \left( \frac{1}{\rho} \left( \frac{T}{T_0} \right)^{3/2} \right) + \frac{5}{2} \right) .
\]  
(259)

Solving for \( T \) gives

\[
\frac{3}{2} \ln \frac{T}{T_0} = \frac{s}{s_0} + \sum_i x_i \ln \frac{x_i}{\rho_i} + \ln \rho - \frac{5}{2} .
\]  
(260)

Using this expression and the constants defined above, we may obtain the ionization fraction \( y \) for given \( \ln \rho \) and \( s \) by finding the root of

\[
F = \ln \left[ \left( \frac{\rho_{\text{He}}}{\rho} \right) \left( \frac{T}{T_0} \right)^{3/2} \frac{1 - y}{y^2} \right] - \frac{T_0}{T} .
\]  
(261)

The derivative with respect to \( y \) for Newton-Raphson is

\[
\frac{\partial F}{\partial y} = \left( \frac{3}{2} + \frac{T_0}{T} \right) \frac{\partial \ln T / T_0}{\partial y} \frac{1}{1 - y} - \frac{2}{y} ,
\]  
(262)

where

\[
\frac{\partial \ln T / T_0}{\partial y} = \frac{2}{3} \frac{\left( \ln \frac{\rho_{\text{He}}}{\rho_0} - F - \frac{T_0}{T} \right) - 1}{1 + y + x_{\text{He}}} .
\]  
(263)

In order to compute the pressure gradient in the momentum equation, the derivative of \( y \) with respect to \( \ln \rho \) and \( s \) needs to be known:

\[
\frac{\partial \ln P}{\partial \ln \rho} = \frac{1}{1 + y + x_{\text{He}}} \frac{\partial y}{\partial \ln \rho} + \frac{\partial \ln T}{\partial \ln \rho} + \frac{\partial \ln T}{\partial y} \frac{\partial y}{\partial \ln \rho} + 1 ,
\]  
(264)

\[
\frac{\partial \ln P}{\partial s} = \frac{1}{1 + y + x_{\text{He}}} \frac{\partial y}{\partial s} + \frac{\partial \ln T}{\partial s} + \frac{\partial \ln T}{\partial y} \frac{\partial y}{\partial s} .
\]  
(265)

Since \( F = 0 \) for all desired solutions \((y, \ln \rho, s)\) we also have

\[
dF = \frac{\partial F}{\partial \ln \rho} d \ln \rho + \frac{\partial F}{\partial s} ds + \frac{\partial F}{\partial y} dy = 0 ,
\]  
(266)

and thus

\[
\frac{\partial y}{\partial \ln \rho} = \left( \frac{dy}{d \ln \rho} \right)_{s=0} = - \frac{\partial F / \partial \ln \rho}{\partial F / \partial y}
\]  
(267)

and

\[
\frac{\partial y}{\partial s} = \left( \frac{dy}{ds} \right)_{\ln \rho=0} = - \frac{\partial F / \partial s}{\partial F / \partial y} .
\]  
(268)

**H.7 Radiative transfer**

**H.7.1 Solving the radiative transfer equation**

A formal solution of Eq. (81) is given by

\[
I(\tau) = I(\tau_0) e^{-(\tau-\tau_0)} + \int_{\tau_0}^{\tau} e^{-(\tau-\tau')} S(\tau') d\tau' .
\]  
(269)
Using a generalization of the trapezoidal rule,

\[
\int_{\tau_0}^{\tau} e^{-(\tau - \tau')} f(\tau') \, d\tau' \approx \int_{\tau_0}^{\tau} e^{-(\tau - \tau_0)} \left[ f(\tau_0) + \frac{f(\tau) - f(\tau_0)}{\tau - \tau_0} (\tau' - \tau_0) \right] \, d\tau' \tag{270}
\]

which is exact for linear functions \( S(\tau) \), we discretize this as

\[
I_{k+1} = I_k e^{-\delta \tau} + (1 - e^{-\delta \tau}) S_{k+1} - \frac{1}{\delta \tau} (S_{k+1} - S_k) \tag{271}
\]

where the simplest way to calculate \( \delta \tau \) is

\[
\delta \tau = \frac{\chi_{k+1} + \chi_k}{2} \delta x \tag{272}
\]

more accurate alternatives are

\[
\delta \tau = \sqrt{\chi_k \chi_{k+1}} \delta x \tag{273}
\]

or

\[
\delta \tau = \chi_{k+1} \ln \frac{\chi_{k+1}}{\ln \chi_k} \delta x \tag{274}
\]

### H.7.2 Angular integration

#### Table 18: Sums \( \sqrt{4\pi} Y_l^m(\theta, \phi) \) for special sets of directions. For all degrees and orders up to \( l = 8 \) not mentioned in this table, the sums are 0. The label ‘Non-h. f-d.’ stands for ‘non-horizontal face-diagonals’, i.e. the eight face diagonals that are not in the horizontal plane.

<table>
<thead>
<tr>
<th>Directions</th>
<th>( Y_0^0 )</th>
<th>( Y_2^0 )</th>
<th>( Y_4^0 )</th>
<th>( Y_6^0 )</th>
<th>( Y_8^0 )</th>
<th>( Y_0^{+4} )</th>
<th>( Y_2^{+4} )</th>
<th>( Y_4^{+4} )</th>
<th>( Y_6^{+4} )</th>
<th>( Y_8^{+4} )</th>
</tr>
</thead>
<tbody>
<tr>
<td>Coord.</td>
<td>6</td>
<td>0</td>
<td>( \frac{21}{2} )</td>
<td>3</td>
<td>( \frac{1}{4} )</td>
<td>( \sqrt{70} )</td>
<td>( \frac{1}{4} )</td>
<td>( \sqrt{13} )</td>
<td>( -\frac{3}{8} )</td>
<td>( \sqrt{182} )</td>
</tr>
<tr>
<td>Face diag.</td>
<td>12</td>
<td>0</td>
<td>( \frac{21}{4} )</td>
<td>-3</td>
<td>( \frac{1}{8} )</td>
<td>( \sqrt{70} )</td>
<td>-3</td>
<td>( \frac{1}{16} )</td>
<td>( \sqrt{13} )</td>
<td>( \frac{39}{32} )</td>
</tr>
<tr>
<td>Space diag.</td>
<td>8</td>
<td>0</td>
<td>( \frac{28}{3} )</td>
<td>-2</td>
<td>( \frac{1}{3} )</td>
<td>( \sqrt{70} )</td>
<td>( \frac{16}{3} )</td>
<td>( \sqrt{13} )</td>
<td>-8</td>
<td>( \sqrt{182} )</td>
</tr>
<tr>
<td>Non-h. f-d.</td>
<td>8</td>
<td>2( \sqrt{5} )</td>
<td>( \frac{39}{4} )</td>
<td>-3</td>
<td>( \frac{3}{8} )</td>
<td>( \sqrt{70} )</td>
<td>-19</td>
<td>( \frac{1}{16} )</td>
<td>( \sqrt{13} )</td>
<td>( \frac{27}{32} )</td>
</tr>
<tr>
<td>Coord. ( x ), ( y )</td>
<td>4</td>
<td>2( \sqrt{5} )</td>
<td>( \frac{9}{2} )</td>
<td>4</td>
<td>( \frac{3}{8} )</td>
<td>( \sqrt{70} )</td>
<td>-5</td>
<td>( \frac{1}{4} )</td>
<td>( \sqrt{13} )</td>
<td>-( \frac{3}{8} )</td>
</tr>
<tr>
<td>Coord. ( z )</td>
<td>2</td>
<td>2( \sqrt{5} )</td>
<td>6</td>
<td>0</td>
<td>2( \sqrt{13} )</td>
<td>0</td>
<td>2( \sqrt{17} )</td>
<td>0</td>
<td>0</td>
<td></td>
</tr>
</tbody>
</table>

For angular integration over the full solid angle, we make the ansatz

\[
\int 4\pi f(\theta, \phi) \frac{d\omega}{4\pi} = \sum_{i=1}^{N} w_i f(\theta_i, \phi_i) + R_N \tag{277}
\]
Table 18 shows the sums $\sqrt{4\pi} Y^m_l(\theta_i, \phi_i)$ over special sets of directions $(\theta_i, \phi_i)$. Using these numbers and requiring that angular integration is exact for $l \leq l_{\text{max}}$, we find the following weights $w_i$ for different sets of directions (see also [1], §25.4.65).

Cooling times have been determined numerically in [3]. Comparing with analytic expressions obtained in the Eddington approximation, the proposed suitable switches in 1-D and 2-D problems.

1. Axes
   Coordinate axes: 1/6
   $l_{\text{max}} = 3$

2. Face diagonals
   Face diagonals: 1/12
   $l_{\text{max}} = 3$

3. Space diagonals
   Space diagonals: 1/8
   $l_{\text{max}} = 3$

4. Axes + face diagonals
   Coordinate axes: 1/30
   Face diagonals: 1/15
   $l_{\text{max}} = 5$

5. Axes + space diagonals
   Coordinate axes: 1/15
   Space diagonals: 3/40
   $l_{\text{max}} = 5$

6. Face + space diagonals
   Face diagonals: 2/15
   Space diagonals: -3/40
   $l_{\text{max}} = 5$

7. Axes, face + space diagonals
   Coordinate axes: 1/21
   Face diagonals: 4/105
   Space diagonals: 9/280
   $l_{\text{max}} = 7$

8. Axes, non-horizontal face diagonals
   Coordinate axes $x, y$: 1/10
   Coordinate axes $z$: 1/30
   Non-hor. face diagonals: 1/15
   $l_{\text{max}} = 3$

9. Axes, non-horizontal face diagonals, space diagonals
   Coordinate axes $x, y$: 12/215
   Coordinate axes $z$: 10/129
   Non-hor. face diagonals: -14/645
   Space diagonals: 171/1720
   $l_{\text{max}} = 5$
I Curvilinear coordinates

The use and implementation of non-cartesian coordinate systems is briefly explained in Section 5.25. All differential operators look like their cartesian counterparts, except that all derivatives are now replaced by covariant derivatives. The relevant reference for the PENCIL CODE is [29]; see their Appendix B. Here some details.

I.1 Covariant derivatives

\( A_{\alpha;\beta} = A_{\alpha,\beta} - \Gamma^\sigma_{\alpha\beta} A_\sigma, \) \hspace{1cm} (278)

\( A_{\alpha;\beta;\gamma} = A_{\alpha,\beta,\gamma} - \Gamma^\delta_{\alpha\beta} A_{\delta,\gamma} - \Gamma^\delta_{\alpha\gamma} A_{\delta,\beta}. \) \hspace{1cm} (279)

Second derivative tensor

\[ A_{\alpha;\beta;\gamma} = A_{\alpha;\beta;\gamma} - \Gamma^\delta_{\alpha\beta} A_{\delta,\gamma} - \Gamma^\delta_{\alpha\gamma} A_{\delta,\beta} - \Gamma^\delta_{\beta\delta} A_{\delta,\alpha} + \Gamma^\delta_{\beta\gamma} \Gamma^\delta_{\delta\gamma} A_{\delta,\alpha}. \]

Elements of the first derivative tensor

\( A_{\hat{r},\hat{r}} = A_{\hat{r},\hat{r}}, \quad A_{\hat{\theta},\hat{\theta}} = A_{\hat{\theta},\hat{\theta}} + r^{-1} A_{\hat{r}}, \quad A_{\hat{\phi},\hat{\phi}} = A_{\hat{\phi},\hat{\phi}} + r^{-1} A_{\hat{r}} + r^{-1} \cot \theta A_{\hat{\theta}}. \) \hspace{1cm} (280)

\( A_{\hat{r},\hat{\theta}} = A_{\hat{\theta},\hat{r}} = A_{\hat{\phi},\hat{\phi}} = A_{\hat{\phi},\hat{r}} - r^{-1} A_{\hat{\phi}}, \quad A_{\hat{\theta},\hat{\phi}} = A_{\hat{\phi},\hat{\theta}} - r^{-1} \cot \theta A_{\hat{\phi}}. \) \hspace{1cm} (281)

I.2 Differential operators

All differential operators look like their cartesian counterparts, except that all derivatives are now replaced by covariant derivatives.

I.2.1 Gradient

For the gradient operator the covariant and partial derivatives are the same, i.e.

\[ \nabla \Psi = \Psi_{,\hat{r}} = \Psi_{,\theta} = \left( \begin{array}{c} \frac{\partial \Psi}{\partial \hat{r}} \\ \frac{\partial \Psi}{\partial \hat{\theta}} \\ \frac{\partial \Psi}{\partial \hat{\phi}} \end{array} \right) \] \hspace{1cm} (282)

where

\[ \frac{\partial}{\partial \hat{r}} = \partial_r \] \hspace{1cm} (283)

\[ \frac{\partial}{\partial \hat{\theta}} = r^{-1} \partial_\theta \] \hspace{1cm} (284)

\[ \frac{\partial}{\partial \hat{\phi}} = \varpi^{-1} \partial_\phi \] \hspace{1cm} (285)

and \( \varpi = r \sin \theta \) is the cylindrical radius. Thus,

\[ \nabla \Psi = (\nabla \Psi)^{(0)}, \] \hspace{1cm} (286)
where the superscript \((0)\) indicated the straightforward calculation in the non-coordinate basis. The coordinate and non-coordinate bases are related to each other via

\[
(\nabla \Psi)^{(0)} \equiv \begin{pmatrix}
\Psi_\rho \\
\Psi_{\rho \theta} \\
\Psi_{\rho \phi}
\end{pmatrix} = \begin{pmatrix}
1 & 0 & 0 \\
0 & r^{-1} & 0 \\
0 & 0 & \omega^{-1}
\end{pmatrix} (\nabla \Psi)^{(\text{coord})}.
\] (287)

Here, the result in the coordinate basis is just what one would get by computing as if one had cartesian coordinates. In the PENCIL CODE the output or the subroutine \text{der} is now in this non-coordinate basis.

\section*{I.2.2 Divergence}

For the divergence operator we have a ‘correction term’, i.e.

\[
\nabla \cdot A = A_\alpha \frac{\partial}{\partial \alpha} = A_\alpha \hat{\alpha} + \Gamma_{\beta \gamma} A_\beta \hat{\gamma},
\] (288)

(289)

where the only non-vanishing contributions to the last term are

\[
\Gamma_{\beta \gamma} A_\beta \hat{\gamma} = \Gamma_{\rho \theta} A^\rho \hat{\rho} + \Gamma_{\rho \phi} A^\rho \hat{\phi} + \Gamma_{\theta \phi} A^\theta \hat{\phi} = 2r^{-1} A^\rho + r^{-1}\cot \theta A^\theta.
\] (290)

Thus,

\[
\nabla \cdot A = (\nabla \cdot A)^{(0)} + M^{(\text{div})}_\alpha A_\alpha,
\] (292)

where

\[
M^{(\text{div})}_\alpha = \begin{pmatrix}
2r^{-1} \\
r^{-1}\cot \theta \\
0
\end{pmatrix}
\] (293)

represents the correction term.

Alternatively:

\[
A_\alpha \hat{\alpha} = A_\alpha \hat{\alpha} - \Gamma_{\alpha \beta} A^\beta
\] (294)

(295)

\section*{I.2.3 Curl}

The curl operator is given by

\[
(\nabla \times B)^\alpha = \epsilon^{\hat{\alpha} \hat{\beta} \hat{\gamma}} B_{\gamma, \hat{\beta}}.
\] (296)

So

\[
\nabla \times A = \begin{pmatrix}
A_{\rho, \theta} - A_{\theta, \rho} \\
A_{\rho, \phi} - A_{\phi, \rho} \\
A_{\theta, \phi} - A_{\phi, \theta}
\end{pmatrix}
\] (297)

\[
\nabla \times A = \begin{pmatrix}
A_{\rho, \theta} - A_{\theta, \rho} + \Gamma_{\theta \phi} A^\phi \\
A_{\rho, \phi} - A_{\phi, \rho} + \Gamma_{\phi \theta} A^\theta \\
A_{\theta, \phi} - A_{\phi, \theta} + \Gamma_{\phi \rho} A^\rho
\end{pmatrix}
\] (298)
Thus,
\[ \nabla \times A = (\nabla \times A)^{(0)} + M^{(\text{curl})}_{\dot{\alpha}\dot{\beta}} A_{\dot{\beta}}, \]
where
\[ M^{(\text{curl})} = \begin{pmatrix} 0 & 0 & r^{-1} \cot \theta \\ 0 & 0 & -r^{-1} \\ 0 & r^{-1} & 0 \end{pmatrix} \]
is the correction term. In the PENCIL CODE we use the subroutine \text{curl}$mn(aij,bb,aa)$, which uses $aij = A_{\dot{\alpha}j}$ and $aa = A_{\dot{\alpha}}$ as input pencils and produces $bb = B_{\dot{\alpha}}$ as output.

### I.2.4 Advection operator

[The usage of roman indices here is insignificant.]

\[ u \cdot \nabla u = \omega \times u + \frac{1}{2} \nabla u^2 = -u \times \omega + \frac{1}{2} \nabla u^2 \]  
(301)

\[ (u \cdot \nabla u)_i = -\epsilon_{ijk}\epsilon_{klm}u_j u_{m;l} + u_j u_{j,i} = (-\delta_{ij}\delta_{jm} + \delta_{im}\delta_{jl})u_j u_{m;l} + u_j u_{j,i} = -u_j u_{j;i} + u_j u_{i;j} + u_j u_{j,i} = u_j u_{i;j} + \Gamma^k_{ji} u_j u_k = u_j u_{i,j} + (-\Gamma^k_{ij} + \Gamma^k_{ji}) u_j u_k \]  
(302)

Note that the terms with $\Gamma^\kappa_{\dot{\theta}\dot{\phi}}$, $\Gamma^\kappa_{\dot{\phi}\dot{\theta}}$, and $\Gamma^\phi_{\dot{\phi}\dot{\phi}}$ drop out. Thus, we have

\[ (u \cdot \nabla u)_\dot{\tau} = u_j u_{\dot{\tau},j} + (-\Gamma^k_{\dot{\tau}j} + \Gamma^k_{j\dot{\tau}}) u_j u_k = u_j u_{\dot{\tau},j} - \Gamma^\phi_{\dot{\phi}\dot{\theta}} u_\phi^2 - \Gamma^\phi_{\dot{\phi}\dot{\phi}} u_\phi^2 = u_j u_{\dot{\tau},j} - r^{-1}(u_\phi^2 + u_\phi^2) \]  
(303)

\[ (u \cdot \nabla u)_\dot{\theta} = u_j u_{\dot{\theta},j} + (-\Gamma^k_{\dot{\theta}j} + \Gamma^k_{j\dot{\theta}}) u_j u_k = u_j u_{\dot{\theta},j} - \Gamma^\phi_{\dot{\phi}\dot{\theta}} u_\phi^2 + \Gamma^\phi_{\dot{\phi}\dot{\phi}} u_\phi^2 = u_j u_{\dot{\theta},j} - r^{-1} \cot \theta u_\phi^2 + r^{-1} u_\phi u_\phi \]  
(304)

\[ (u \cdot \nabla u)_\dot{\phi} = u_j u_{\dot{\phi},j} + (-\Gamma^k_{\dot{\phi}j} + \Gamma^k_{j\dot{\phi}}) u_j u_k = u_j u_{\dot{\phi},j} + \Gamma^\phi_{\dot{\phi}\dot{\theta}} u_\phi u_\phi + \Gamma^\phi_{\dot{\phi}\dot{\phi}} u_\phi u_\phi = u_j u_{\dot{\phi},j} + r^{-1} u_\phi u_\phi + r^{-1} \cot \theta u_\phi u_\phi \]  
(305)

Note that the formulation above is slightly misleading, because

\[ \Gamma^k_{\dot{\tau}j} u_j u_k = \Gamma^k_{\dot{\phi}\dot{\theta}} u_\phi u_\phi + \Gamma^\phi_{\dot{\phi}\dot{\phi}} u_\phi u_\phi = 0 \]  
(306)

\[ \Gamma^k_{\dot{\phi}j} u_j u_k = \Gamma^k_{\dot{\phi}\dot{\phi}} u_\phi u_\phi + \Gamma^\phi_{\dot{\phi}\dot{\phi}} u_\phi u_\phi + \Gamma^\phi_{\dot{\phi}\dot{\phi}} u_\phi u_\phi + \Gamma^\phi_{\dot{\phi}\dot{\phi}} u_\phi u_\phi = 0 \]  
(307)

### I.2.5 Mixed advection operator

\[ u \times B = u \times \nabla B = u_j \epsilon_{ijk}\epsilon_{klm}A_{m;l} = u_j (\delta_{ij}\delta_{jm} - \delta_{im}\delta_{jl})A_{m;l} = u_j A_{j;l} - u_j A_{i;j} \]  
(308)
I.2.6 Shear term

\[ u_j A_{j,i} = (u_j A_j)_i - u_{j,i} A_j = (u_j A_j)_i - u_{j,i} A_j \]  \hspace{1cm} (309)

\[ u_{j,i} A_j = u_{j,i} A_j - \Gamma^k_{ji} u_k A_j \]  \hspace{1cm} (310)

So

\[ u_{\phi,\phi} A_{\phi} = u_{\phi,\phi,\phi} A_{\phi} - \Gamma^k_{\phi,\phi} u_k A_{\phi} = u_{\phi,\phi,\phi} A_{\phi} \]  \hspace{1cm} (311)

\[ u_{\phi,\phi} A_{\phi} = u_{\phi,\phi,\phi} A_{\phi} - \Gamma^k_{\phi,\phi} u_k A_{\phi} \]  \hspace{1cm} (312)

I.2.7 Another mixed advection operator

\[ u_{\beta} A_{\alpha,\beta} = u_{\beta} A_{\alpha,\beta} - \Gamma^\gamma_{\alpha,\beta} u_{\beta} A_{\gamma} \]  \hspace{1cm} (313)

Write out

\[ (u \cdot \nabla A)_{\phi} = u_{\phi} A_{\phi,\phi} + u_{\theta,\phi} A_{\phi,\phi} + u_{\phi,\phi} A_{\phi,\phi} \]

\[ (u \cdot \nabla A)_{\theta} = u_{\theta} A_{\theta,\theta} + u_{\theta,\theta} A_{\theta,\theta} + u_{\phi,\theta} A_{\theta,\theta} \]

\[ (u \cdot \nabla A)_{\phi} = u_{\phi} A_{\phi,\phi} + u_{\theta,\phi} A_{\phi,\phi} + u_{\phi,\phi} A_{\phi,\phi} \]

Reorder

\[ (u \cdot \nabla A)_{\phi} = u_{\phi} A_{\phi,\phi} + u_{\theta,\phi} A_{\phi,\phi} + u_{\phi,\phi} A_{\phi,\phi} - r^{-1} u_{\phi} A_{\phi} - r^{-1} u_{\phi} A_{\phi} \]

\[ (u \cdot \nabla A)_{\theta} = u_{\phi} A_{\theta,\theta} + u_{\theta,\theta} A_{\theta,\theta} + u_{\phi,\theta} A_{\theta,\theta} + r^{-1} u_{\phi} A_{\phi} - r^{-1} \cot \theta u_{\phi} A_{\phi} \]

\[ (u \cdot \nabla A)_{\phi} = u_{\phi} A_{\phi,\phi} + u_{\theta,\phi} A_{\phi,\phi} + u_{\phi,\phi} A_{\phi,\phi} + r^{-1} u_{\phi} A_{\phi} + r^{-1} \cot \theta u_{\phi} A_{\phi} \]  \hspace{1cm} (314)

I.2.8 Strain Matrix

The strain matrix takes the form \[ 2s_{\alpha,\beta} = u_{\alpha,\beta} + u_{\beta,\alpha} \]. The components of the covariant derivative tensor were given in Eqs. (280) and (281), so the final result reads

\[ 2s = \begin{pmatrix}
2u_{\phi,\phi} & u_{\phi,\theta} + u_{\theta,\phi} - u_{\phi}/r & u_{\phi,\phi} - u_{\phi}/r \\
u_{\phi,\phi} - u_{\phi}/r & 2u_{\phi,\phi} + 2u_{\phi}/r & u_{\phi,\phi} - u_{\phi}/r \\
u_{\phi,\phi} - u_{\phi}/r & u_{\phi,\phi} - u_{\phi}/r & 2u_{\phi,\phi} + 2u_{\phi}/r + 2u_{\phi,\phi} \end{pmatrix} \]  \hspace{1cm} (315)

The trace-less rate of strain matrix is given by

\[ S_{ij} = s_{ij} - \frac{1}{3} \delta_{ij} \nabla \cdot u. \]  \hspace{1cm} (316)

I.2.9 Lambda effect

\[ Q_{ij} = \begin{pmatrix}
0 & 0 & \Lambda_V \Omega \\
0 & 0 & \Lambda_H \Omega \\
\Lambda_V \Omega & \Lambda_H \Omega & 0 \\
\end{pmatrix} \]  \hspace{1cm} (317)

where \[ \Omega = u_{\phi}/r \sin \theta \]. Next, compute \[ Q_{ij;j} \].

\[ Q_{\alpha,\beta,\gamma} = Q_{\alpha,\beta,\gamma} - \Gamma^\gamma_{\alpha,\gamma} Q_{\alpha,\beta} - \Gamma^\gamma_{\beta,\gamma} Q_{\alpha,\beta}. \]  \hspace{1cm} (318)
\[ Q_{\hat{\alpha} r;\hat{\beta}} = Q_{\hat{\alpha} r;\hat{\beta}} - \Gamma^\sigma_{\hat{\alpha} \hat{\beta}} Q_{\sigma \hat{r}} - \Gamma^\sigma_{\hat{r} \hat{\alpha}} Q_{\hat{\beta} \sigma} . \]
\[ = 0 \tag{319} \]

\[ Q_{\hat{\beta} \hat{\phi};\hat{\phi}} = Q_{\hat{\beta} \hat{\phi};\hat{\phi}} - \Gamma^\sigma_{\hat{\beta} \hat{\phi}} Q_{\sigma \phi} - \Gamma^\sigma_{\phi \hat{\beta}} Q_{\phi \sigma} . \]
\[ = -\Gamma^\phi_{\hat{\beta} \hat{\phi}} Q_{\hat{\phi} \sigma} - \Gamma^\phi_{\phi \hat{\beta}} Q_{\phi \hat{\phi}} . \]
\[ = 0 \tag{320} \]

\[ Q_{\hat{\beta} \hat{\phi} ;\hat{\phi}} = Q_{\hat{\beta} \hat{\phi} ;\hat{\phi}} - \Gamma^\phi_{\hat{\beta} \hat{\phi}} Q_{\phi \phi} - \Gamma^\phi_{\phi \hat{\beta}} Q_{\phi \phi} . \]
\[ = Q_{\hat{\beta} \hat{\phi} ;\hat{\phi}} + r^{-1} Q_{\phi \hat{r}} \tag{323} \]

\[ Q_{\phi ;\phi} = Q_{\phi ;\phi} - \Gamma^\phi_{\phi} Q_{\phi \phi} - \Gamma^\phi_{\phi} Q_{\phi \phi} . \]
\[ = -\Gamma^\phi_{\phi} Q_{\phi \phi} - \Gamma^\phi_{\phi} Q_{\phi \phi} . \]
\[ = +2r^{-1} \Lambda_\Omega + 2 \cot \theta r^{-1} \Lambda_\Omega \tag{324} \]

So, the \( \phi \) component of the divergence of the Lambda tensor is then
\[ Q_{\hat{\phi} \hat{\phi} ;\hat{\phi}} = Q_{\phi \hat{r} ;\hat{\phi}} + 3r^{-1} \Lambda_\Omega + 2 \cot \theta r^{-1} \Lambda_\Omega \tag{325} \]

### I.2.10 Laplacian of a scalar

Let \( E_{\hat{\alpha}} = (\nabla \Psi)_{\hat{\alpha}} = \partial_{\hat{\alpha}} \Psi \) Then
\[ \Delta \Psi = E_{\hat{\beta} ;\hat{\beta}} = (\partial_{\hat{\beta}} \Psi)_{\hat{\beta}} + \frac{2}{r} \Psi_{,\hat{r}} + \frac{\partial \theta}{r} \Psi_{,\hat{\theta}} \tag{326} \]

### I.2.11 Hessian of a scalar

\[ s_{,\hat{\alpha}} = s_{,\hat{\alpha}} \tag{327} \]
\[ s_{,\hat{r} r} = s_{,\hat{r} r} - \Gamma^\phi_{,\hat{r} \hat{r}} s_{,\hat{r} \phi} = s_{,\hat{r} \phi} \tag{328} \]
\[ s_{,\hat{r} \hat{\phi}} = s_{,\hat{r} \hat{\phi}} - \Gamma^\phi_{,\hat{r} \hat{\phi}} s_{,\hat{r} \phi} = s_{,\hat{r} \hat{\phi}} + r^{-1} s_{,\phi} \tag{329} \]
\[ s_{,\phi \phi} = s_{,\phi \phi} - \Gamma^\phi_{,\phi \phi} s_{,\phi \phi} \tag{330} \]
\[ s_{,\phi \phi} = s_{,\phi \phi} - \Gamma^\phi_{,\phi \phi} s_{,\phi \phi} \]
\[ s_{\hat{\vartheta}\hat{\vartheta}} = s_{\hat{\vartheta}\hat{\vartheta}} - \Gamma^\vartheta_{\hat{\vartheta}\hat{\vartheta}} s_{\vartheta\vartheta} = s_{\hat{\vartheta}\hat{\vartheta}} - r^{-1} s_{\vartheta\vartheta} \] (331)

\[ s_{\hat{\varphi}\hat{\varphi}} = s_{\hat{\varphi}\hat{\varphi}} - \Gamma^\varphi_{\hat{\varphi}\hat{\varphi}} s_{\varphi\varphi} = s_{\hat{\varphi}\hat{\varphi}} - r^{-1} s_{\varphi\varphi} \] (332)

\[ s_{\hat{\vartheta}\hat{\varphi}} = s_{\hat{\varphi}\hat{\vartheta}} - \Gamma^\varphi_{\hat{\vartheta}\hat{\varphi}} s_{\varphi\vartheta} = s_{\hat{\vartheta}\hat{\varphi}} - \Gamma^\vartheta_{\hat{\varphi}\hat{\vartheta}} s_{\vartheta\varphi} \] (333)

So,

\[ s_{\hat{\alpha}\hat{\beta}} = s_{\hat{\alpha}\hat{\beta}} + \begin{pmatrix} 0 & -r^{-1} s_{\hat{\vartheta}} & -r^{-1} s_{\hat{\varphi}} \\ 0 & +r^{-1} s_{\hat{\vartheta}} & -r^{-1} \cot \theta s_{\hat{\varphi}} \\ 0 & 0 & -\Gamma^\varphi_{\hat{\varphi}\hat{\varphi}} s_{\varphi} - \Gamma^\vartheta_{\hat{\vartheta}\hat{\vartheta}} s_{\vartheta} \end{pmatrix} \] (337)

**I.2.12 Double curl**

For the calculation of \( J = \nabla \times B \) we use the same curl routine, but with different arguments, \( \text{bij} = B_{\alpha,\beta} \) and \( \text{bb} = B_{\alpha} \) as input pencils and \( \text{jj} = J_{\alpha} \) as output, i.e. \text{curl mn(bij, jj, bb)}, so that

\[ J = \begin{pmatrix} B_{\hat{\alpha},\hat{\beta}} - B_{\hat{\beta},\hat{\alpha}} + \Gamma^\vartheta_{\hat{\alpha},\hat{\beta}} B^\vartheta \\
B_{\hat{\alpha},\hat{\vartheta}} - B_{\hat{\vartheta},\hat{\alpha}} - \Gamma^\varphi_{\hat{\alpha},\hat{\vartheta}} B^\varphi \\
B_{\hat{\alpha},\hat{\varphi}} - B_{\hat{\varphi},\hat{\alpha}} + \Gamma^\varphi_{\hat{\alpha},\hat{\varphi}} B^\varphi \end{pmatrix} = (\nabla \times B)^{(0)} + M^{(\text{curl})} B_{\beta}, \] (338)

where

\[ (\nabla \times B)^{(0)} = \epsilon_{\hat{\alpha}\hat{\beta}\hat{\gamma}} B_{\hat{\gamma},\hat{\beta}}. \] (339)

Expressing the components of \((\nabla \times B)^{(0)}\) in terms of \( A_{\alpha} \) we have

\[ B_{\hat{\alpha},\hat{\beta}} = (A_{\hat{\alpha},\hat{\beta}} - A_{\hat{\beta},\hat{\alpha}} + \Gamma^\vartheta_{\hat{\alpha},\hat{\beta}} A^\vartheta)_{\hat{\alpha}} \] \[ B_{\hat{\alpha},\hat{\vartheta}} = (A_{\hat{\alpha},\hat{\vartheta}} - A_{\hat{\vartheta},\hat{\alpha}} + \Gamma^\varphi_{\hat{\alpha},\hat{\vartheta}} A^\varphi)_{\hat{\alpha}} \] \[ B_{\hat{\alpha},\hat{\varphi}} = (A_{\hat{\alpha},\hat{\varphi}} - A_{\hat{\varphi},\hat{\alpha}} - \Gamma^\varphi_{\hat{\alpha},\hat{\varphi}} A^\varphi)_{\hat{\alpha}} \] (340)

or

\[ B_{\hat{\alpha},\hat{\beta}} = A_{\hat{\alpha},\hat{\beta}} - A_{\hat{\alpha},\hat{\beta}} + (\Gamma^\vartheta_{\hat{\alpha},\hat{\beta}} A^\vartheta)_{\hat{\beta}} \] \[ B_{\hat{\alpha},\hat{\vartheta}} = A_{\hat{\alpha},\hat{\vartheta}} - A_{\hat{\vartheta},\hat{\alpha}} + (\Gamma^\varphi_{\hat{\alpha},\hat{\vartheta}} A^\varphi)_{\hat{\beta}} \] \[ B_{\hat{\alpha},\hat{\varphi}} = A_{\hat{\alpha},\hat{\varphi}} - A_{\hat{\varphi},\hat{\alpha}} - (\Gamma^\varphi_{\hat{\alpha},\hat{\varphi}} A^\varphi)_{\hat{\beta}} \] (341)

Thus,

\[ B_{\hat{\alpha},\hat{\beta}} = (B_{\hat{\alpha},\hat{\beta}})^{(0)} + M^{(2\text{curl}^2)} A_{\hat{\beta},\hat{\alpha}} + M^{(1\text{curl}^2)} A_{\hat{\alpha},\hat{\beta}}, \] (342)

where

\[ (B_{\hat{\alpha},\hat{\beta}})^{(0)} = \epsilon_{\hat{\alpha}\hat{\beta}\hat{\gamma}} A_{\hat{\gamma},\hat{\beta}}. \] (343)
I.2.13 Gradient of a divergence

Note that some derivatives of Christoffel symbols vanish, so we are left with

\[ B_\phi,\dot{\theta} = \ldots + \Gamma^\phi_\theta^\alpha A^\phi_\theta \quad B_{\theta,\phi} = \ldots - \Gamma^\phi_\theta^\alpha A^\phi_\theta - \Gamma^\phi_\theta^\phi A^\phi_\theta \]
\[ B_{\dot{r},\phi} = \ldots + \Gamma^\phi_\theta^\alpha A^\phi_\theta \quad B_{\phi,\dot{r}} = \ldots + \Gamma^\phi_\theta^\phi A^\phi_\theta + \Gamma^\phi_\theta^\alpha A^\phi_\theta \]
\[ B_{\dot{r},\dot{\theta}} = \ldots - \Gamma^\phi_\theta^\alpha A^\phi_\theta - \Gamma^\phi_\theta^\alpha A^\phi_\theta \]

Note that some derivatives of Christoffel symbols vanish, so we are left with

\[ B_\phi,\dot{\theta} = r^{-1} A^\phi_\theta \quad B_{\theta,\phi} = -r^{-1} A^\phi_\theta \]
\[ B_{\dot{r},\phi} = r \cot \theta A^\phi_\phi \quad B_{\phi,\dot{r}} = r^{-1} A^\phi_\phi - r^{-2} \sin^{-2} \theta A^\phi_\phi \]

\[ \dot{r} \text{ component:} \]

\[ A_{\dot{r}} = A_{\dot{r}} = \Gamma^\sigma_\alpha A_{\dot{r}} - \Gamma^\sigma_{\dot{\alpha}} A_{\dot{r}} - \Gamma^\sigma_{\dot{\alpha} \dot{\gamma}} A_{\dot{r}} = \Gamma^\sigma_\alpha A_{\dot{r}} + \Gamma^\sigma_{\dot{\alpha} \dot{\gamma}} A_{\dot{r}} + \Gamma^\sigma_{\alpha \dot{\gamma}} A_{\dot{r}} \]

\[ \dot{\theta} \text{ component:} \]

\[ A_{\dot{\theta}} = A_{\dot{\theta}} = \Gamma^\sigma_\alpha A_{\dot{\theta}} - \Gamma^\sigma_{\dot{\alpha}} A_{\dot{\theta}} - \Gamma^\sigma_{\dot{\alpha} \dot{\theta}} A_{\dot{\theta}} = \Gamma^\sigma_\alpha A_{\dot{\theta}} + \Gamma^\sigma_{\dot{\alpha} \dot{\theta}} A_{\dot{\theta}} \]

Note that the last four terms in the above expression canceled, because

\[ -\Gamma^\sigma_{\dot{\alpha}} A_{\dot{\theta}} + \Gamma^\sigma_{\dot{\alpha} \dot{\theta}} A_{\dot{\theta}} - \Gamma^\sigma_{\dot{\alpha} \dot{\theta}} A_{\dot{\theta}} = -r^{-1} A_{\dot{\theta}} + r^{-1} A_{\dot{\theta}} - r^{-2} A_{\dot{\theta}} - r^{-2} A_{\dot{\theta}} = 0 \]

\[ \dot{\phi} \text{ component:} \]

\[ A_{\dot{\phi}} = A_{\dot{\phi}} = \Gamma^\sigma_\alpha A_{\dot{\phi}} - \Gamma^\sigma_{\dot{\alpha}} A_{\dot{\phi}} - \Gamma^\sigma_{\dot{\alpha} \dot{\phi}} A_{\dot{\phi}} = \Gamma^\sigma_\alpha A_{\dot{\phi}} + \Gamma^\sigma_{\dot{\alpha} \dot{\phi}} A_{\dot{\phi}} \]
\[
\begin{align*}
   &= A_{\dot{a},\dot{a}} - \Gamma^\sigma_{\dot{a}\dot{a}} A_{\dot{\sigma},\dot{\sigma}} \\
   &= A_{\dot{a},\dot{a}} - \Gamma^\rho_{\dot{\theta}\dot{\theta}} A_{\dot{r},\dot{\phi}} - \Gamma^\phi_{\dot{\theta}\dot{\phi}} A_{\dot{\rho},\dot{\phi}} - \Gamma^\phi_{\dot{\phi}\dot{\phi}} A_{\dot{\theta},\dot{\phi}} \\
   &= A_{\dot{a},\dot{a}} + r^{-1} A_{\dot{r},\dot{\phi}} + r^{-1} A_{\dot{\rho},\dot{\phi}} + r^{-1} \cot \theta A_{\dot{\theta},\dot{\phi}} \\
   &= A_{\dot{a},\dot{a}} + 2r^{-1} A_{\dot{r},\dot{\phi}} + r^{-1} \cot \theta A_{\dot{\theta},\dot{\phi}}
\end{align*}
\]

(357)

In the first line of the expression above, the last four terms vanish\(^{21}\) and the \(\phi\) derivative of any Christoffel symbol (term before that) also vanishes.

\(^{21}\) The following four terms vanish because

\[
\begin{align*}
   &-\Gamma^\sigma_{\dot{a}\dot{a}} A_{\dot{\sigma},\dot{\sigma}} + \Gamma^\rho_{\dot{\theta}\dot{\theta}} \Gamma^\phi_{\dot{\phi}\dot{\phi}} A_{\dot{\rho},\dot{\phi}} - \Gamma^\phi_{\dot{\theta}\dot{\theta}} A_{\dot{\phi},\dot{\phi}} + \Gamma^\phi_{\dot{\phi}\dot{\phi}} A_{\dot{\theta},\dot{\phi}} \\
   &= \Gamma^\rho_{\dot{\theta}\dot{\theta}} (-A_{\dot{\theta},\dot{\rho}} + \Gamma^\phi_{\dot{\theta}\dot{\theta}} A_{\dot{\phi},\dot{\rho}} - A_{\dot{\rho},\dot{\phi}} + \Gamma^\phi_{\dot{\phi}\dot{\phi}} A_{\dot{\theta},\dot{\phi}}) \\
   &= \Gamma^\phi_{\dot{\theta}\dot{\phi}} (-A_{\dot{\phi},\dot{\theta}} + \Gamma^\rho_{\dot{\theta}\dot{\theta}} A_{\dot{\theta},\dot{\rho}} - A_{\dot{\phi},\dot{\theta}} + \Gamma^\phi_{\dot{\phi}\dot{\phi}} A_{\dot{\phi},\dot{\rho}}) \\
   &= r^{-1}(-A_{\dot{\phi},\dot{\theta}} - A_{\dot{\theta},\dot{\phi}} + \Gamma^\phi_{\dot{\phi}\dot{\phi}} A_{\dot{\phi},\dot{\rho}}) \\
   &= r^{-1} \cot \theta (-A_{\dot{\phi},\dot{\theta}} - A_{\dot{\theta},\dot{\phi}} + \Gamma^\phi_{\dot{\phi}\dot{\phi}} A_{\dot{\phi},\dot{\rho}}) \\
   &- r^{-1} \cot \theta (-A_{\dot{\phi},\dot{\theta}} - A_{\dot{\theta},\dot{\phi}} + \Gamma^\phi_{\dot{\phi}\dot{\phi}} A_{\dot{\phi},\dot{\rho}}) = 0
\end{align*}
\]

(358)
J Switchable modules

The material in this section is being assembled from the file ‘inlinedoc-modules.tex’, which is automatically assembled. However, it is currently incomplete and contains only a small number of the available modules.

<table>
<thead>
<tr>
<th>Module</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>hydro.f90</td>
<td>This module takes care of most of the things related to velocity. Pressure, for example, is added in the energy (entropy) module.</td>
</tr>
<tr>
<td>chemistry.f90</td>
<td>This module adds chemical species and reactions. The units used in the chem.in files are cm3,mole,sec,kcal and K</td>
</tr>
<tr>
<td>geometrical_types.f90</td>
<td>Collection of geometrical object types. (Presently only rectangular toroid)</td>
</tr>
<tr>
<td>gpu_astaroth.f90</td>
<td>This module contains GPU related types and functions to be used in ASTAROTH nucleus.</td>
</tr>
<tr>
<td>hydro_potential.f90</td>
<td>This module takes care of most of the things related to velocity. Pressure, for example, is added in the energy (entropy) module.</td>
</tr>
<tr>
<td>noentropy.f90</td>
<td>Calculates pressure gradient term for polytropic equation of state $p = \text{const} \rho^\Gamma$.</td>
</tr>
<tr>
<td>nogpu.f90</td>
<td>This module contains GPU related dummy types and functions.</td>
</tr>
<tr>
<td>nohydro.f90</td>
<td>no variable $u$: useful for kinematic dynamo runs.</td>
</tr>
<tr>
<td>nopower_spectrum.f90</td>
<td>reads in full snapshot and calculates power spectrum of $u$</td>
</tr>
<tr>
<td>noyinyang.f90</td>
<td>This module contains Yin-Yang related dummy types and functions.</td>
</tr>
<tr>
<td>noyinyang_mpi.f90</td>
<td>This module contains Yin-Yang related dummy types and functions.</td>
</tr>
<tr>
<td>particles_adsorbed.f90</td>
<td>This module takes care of the evolution of adsorbed species on the particle surface for reactive particles</td>
</tr>
<tr>
<td>particles_chemistry.f90</td>
<td>This module implements reactive particles.</td>
</tr>
<tr>
<td>particles_surfspec.f90</td>
<td>immediate vicinity of reactive particles.</td>
</tr>
<tr>
<td>power_spectrum.f90</td>
<td>reads in full snapshot and calculates power spectrum of $u$</td>
</tr>
<tr>
<td>test_chemistry.f90</td>
<td>This module adds chemical species and reactions. The units used in the chem.in files are cm3,mole,sec,kcal and K</td>
</tr>
<tr>
<td>timestep.f90</td>
<td>Runge-Kutta time advance, accurate to order itorder. At the moment, itorder can be 1, 2, or 3.</td>
</tr>
<tr>
<td>timestep_strang.f90</td>
<td>Runge-Kutta time advance, accurate to order itorder. At the moment, itorder can be 1, 2, or 3.</td>
</tr>
<tr>
<td>timestep_subcycle.f90</td>
<td>This is a highly specified timestep module currently only working together with the special module coronae.f90.</td>
</tr>
<tr>
<td>yinyang.f90</td>
<td>This module contains Yin-Yang related types and functions which are incompatible with FORTRAN 95.</td>
</tr>
<tr>
<td>yinyang_mpi.f90</td>
<td>This module contains Yin-Yang related types and functions which are incompatible with FORTRAN 95.</td>
</tr>
</tbody>
</table>
K  Startup and run-time parameters

K.1  List of startup parameters for ‘start.in’

The following table lists all (at the time of writing, September 2002) namelists used in ‘start.in’, with the corresponding parameters and their default values (in square brackets). Any variable referred to as a flag can be set to any nonzero value to switch the corresponding feature on. Not all parameters are used for a given scenario. This list is not necessarily up to date; also, in many cases it can only give an idea of the corresponding initial state; to get more insight and the latest set of parameters, you need to look at the code.

The value $\varepsilon$ corresponds to 5 times the smallest number larger than zero. For single precision, this is typically about $\varepsilon \approx 5 \times 1.2 \times 10^{-7} = 6 \times 10^{-7}$; for double precision, $\varepsilon \approx 10^{-15}$.

<table>
<thead>
<tr>
<th>Variable [default value]</th>
<th>Namelist init_pars</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>cvsid [’’]</td>
<td></td>
<td>the svn identification string, which allows you to keep track of the version of ‘start.in’.</td>
</tr>
<tr>
<td>xyz0 [($-\pi, -\pi, -\pi$)]</td>
<td></td>
<td>determine the geometry of the box. All three are vectors of the form $(x$-comp., $y$-comp., $z$-comp.); $xyz0$ describes the left (lower) corner of the box, $Lxyz$ the box size. $lperi$ specifies whether a direction is considered periodic (in which case the last point is omitted) or not. In all cases, three ghost zones will be added.</td>
</tr>
<tr>
<td>$Lxyz$ [(2$\pi$, 2$\pi$, 2$\pi$)],</td>
<td></td>
<td></td>
</tr>
<tr>
<td>$lperi$ [(T,T,T)]</td>
<td></td>
<td></td>
</tr>
<tr>
<td>lprocz_slowest [T]</td>
<td></td>
<td>if set to F, the ordering of processor numbers is changed, so the z processors are now in the inner loop. Since $nprocy=4$ is optimal (see Sect. 5.20.2), you may want to put $lprocz_slowest=T$ when $nygrid&gt;nzgrid$.</td>
</tr>
<tr>
<td>lwrite_ic [F]</td>
<td></td>
<td>if set T, the initial data are written into the file ‘VAR0’. This is generally useful, but doing this all the time uses up plenty of disk space.</td>
</tr>
<tr>
<td>lnwrite [F]</td>
<td></td>
<td>if set T, all initialization files are written, including the param.nml file, except ‘var.dat’. This option allows you to use old file var.dat files, but updates all other initialization files. This could be useful after having changed the code and, in particular, when the ‘var.dat’ files will be overwritten by ‘remesh.csh’.</td>
</tr>
<tr>
<td>lwrite_aux [F]</td>
<td></td>
<td>if set T, auxiliary variables (those calculated at each step, but not evolved mathematically) to ‘var.dat’ after the evolved quantities.</td>
</tr>
<tr>
<td>lwrite_2d [F]</td>
<td></td>
<td>if set T, only 2D-snapshots are written into VAR files in the case of 2D-runs with $nygrid=1$ or $nzgrid=1$.</td>
</tr>
</tbody>
</table>
K.1 Startup parameters for ‘start.in’

lread_oldsnap [F] if set T, the old snapshot will be read in before producing (overwriting) initial conditions. For example, if you just want to add a perturbation to the magnetic field, you’d give no initial condition for density and velocity (so you keep the data from a hopefully relaxed run), and just add whatever you need for the magnetic field. In this connection you may want to touch NOERASE, so as not to erase the previous data. Also, put: ireset_tstart=0,lread_oldsnap=T,lwrite_varAnyway=T

lread_oldsnap_nomag [F] if set T, the old snapshot from a non-magnetic run will be read in before producing (overwriting) initial conditions. This allows one to let a hydrodynamic run relax before adding a magnetic field. However, for this to work one has to modify manually ‘data/param.nml’ by adding an entry for MAGNETIC_INIT.PARS or PSCALAR_INIT.PARS. In addition, for idl to read correctly after the first restarted run, you must adjust the value of mvar in ‘data/dim.dat’

lread_oldsnap_nopscalar [F] if set T, the old snapshot from a run without passive scalar will be read in before producing (overwriting) initial conditions. This allows one to let a hydrodynamic run relax before adding a passive scalar.

lshift_origin [F,F,F] if set T for any or some of the three directions, the mesh is shifted by 1/2 meshpoint in that or those directions so that the mesh goes through the origin.

unit_system ['cgs'] you can set this character string to ‘SI’, which means that you can give physical dimensions in SI units. The default is cgs units.

unit_length [1] allows you to set the unit length. Suppose you want the unit length to be 1 kpc, then you would say unit_length=’3e21’. (Of course, politically correct would be to say unit_system=’SI’ in which case you say unit_length=’3e19’.)

unit_velocity [1] Example: if you want km/s you say unit_length=’1e5’.

unit_density [1] Example: if you want your unit density to be \(10^{-24} \text{ g/cm}^3\) you say unit_density=’1e-24’.

unit_temperature [1] Example: unit_temperature=’1e6’ if you want mega-Kelvin.

random_gen [system] choose random number generator; currently valid choices are

‘system’ (your compiler’s generator),
‘min_std’ (the ‘minimal standard’ generator ran0() from ‘Numerical Recipes’),
‘nr_f90’ (the Parker-Miller-Marsaglia generator ran() from ‘Numerical Recipes for F90’).

bcx (‘p’, ’p’, …),
boundary conditions. See Sect. 5.16 for a discussion of where and how to set these.

selects $\ln T$ as fundamental thermodynamic variable in the entropy module.

---

**Namelist `hydro_init_pars`**

**inituu ["zero"]**

initialization of velocity. Currently valid choices are
- ‘zero’ ($u = 0$),
- ‘gaussian-noise’ (random, normally-distributed $u_x, u_z$),
- ‘gaussian-noise-x’ (random, normally-distributed $u_x$),
- ‘sound-wave’ (sound wave in $x$ direction),
- ‘shock-tube’ (polytropic standing shock),
- ‘bullets’ (blob-like velocity perturbations),
- ‘Alfven-circ-x’ (circularly polarized Alfven wave in $x$ direction),
- ‘const-ux’ (constant $x$-velocity),
- ‘const-uy’ (constant $y$-velocity),
- ‘tang-discont-z’ (tangential discontinuity: velocity is directed along $x$, jump is at $z = 0$),
- ‘Fourier-trunc’ (truncated Fourier series),
- ‘up-down’ (flow upward in one spot, downward in another; not solenoidal).

**ampluu [0.]**

amplitude for some types of initial velocities.

**widthuu [0.1]**

width for some types of initial velocities.

**urand [0.]**

additional random perturbation of $u$. If $\text{urand}>0$, the perturbation is additive, $u_i \mapsto u_i + \text{urand}\mathcal{U}_{[0,0.5,0.5]}$; if $\text{urand}<0$, it is multiplicative, $u_i \mapsto u_i \times \text{urand}\mathcal{U}_{[0,0.5,0.5]}$; in both cases, $\mathcal{U}_{[0,0.5,0.5]}$ is a uniformly distributed random variable on the interval $[-0.5, 0.5]$.

**uu_left [0.], uu_right [0.]**

needed for inituu=’shock-tube’.

---

**Namelist `density_init_pars`**
K.1 Startup parameters for ‘start.in’

initlnrho ['zero']

initialization of density. Currently valid choices are
‘zero’ ($\ln \rho = 0$),
‘isothermal’ (isothermal stratification),
‘polytropic_simple’ (polytropic stratification),
‘hydrostatic-z-2’ (hydrostatic vertical stratification for isentropic atmosphere),
‘xjump’ (density jump in $x$ of width widthlnrho),
‘rho-jump-z’ (density jump in $z$ of width widthlnrho),
‘piecew-poly’ (piecewise polytropic vertical stratification for solar convection),
‘polytropic’ (polytropic vertical stratification),
‘sound-wave’ (sound wave),
‘shock-tube’ (polytropic standing shock),
‘gaussian-noise’ (Gaussian-distributed, uncorrelated noise),
‘gaussian-noise’ (Gaussian-distributed, uncorrelated noise in $x$, but uniform in $y$ and $z$),
‘hydrostatic-r’ (hydrostatic radial density stratification for isentropic or isothermal sphere),
‘sin-xy’ (sine profile in $x$ and $y$),
‘sin-xy-rho’ (sine profile in $x$ and $y$, but in \(\rho\), not $\ln \rho$),
‘linear’ (linear profile in $k \cdot x$),
‘planet’ (planet solution; see [C.7]).

gamma [5./3]

adiabatic index $\gamma = c_p/c_v$.

cs0 [1.]

can be used to set the dimension of velocity; larger values can be used to decrease stratification

rho0 [1.]

reference values of sound speed and density, i.e. values at height $z_{ref}$.

ampllnrho [0.],
widthlnrho [0.1]

amplitude and width for some types of initial densities.

rho_left [1.],
rho_right [1.]

taken for initlnrho=’shock-tube’.

cs2bot [1.],
cs2top [1.]

sound speed at bottom and top. Needed for some types of stratification.

Namelist grav_init_pars

zref [0.]

reference height where in the initial stratification
$c_z^2 = c_{s0}^2$ and $\ln \rho = \ln \rho_0$.

grvz [-1.]

grvz_profile
['const']

vertical gravity component $g_z$.

constant gravity $g_z = \text{grvz (grvz_profile='const')}$

gravity or linear profile $g_z = \text{grvz} \cdot z$ (grvz_profile='linear', for accretion discs and similar).

z1 [0.],
z2 [1.] specific to the solar convection case

initlnrh0='piecew-poly'. The stable layer is

\( z_0 < z < z_1 \), the unstable layer \( z_1 < z < z_2 \), and the
top (isothermal) layer is \( z_2 < z < z_{\text{top}} \).

nu_epicycle [1.] vertical epicyclic frequency; for accretion discs it

should be equal to \( \Omega \), but not for galactic discs;
see Eq. (135) in Sect. C.5.

g grav_amp [0.], grav_tilt [0.] specific to the tilted gravity case (amplitude and an-
gle wrt the vertical direction).

<table>
<thead>
<tr>
<th>Namelist</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>initss ['nothing']</td>
<td>initialization of entropy. Currently valid choices are</td>
</tr>
<tr>
<td></td>
<td>'nothing' (leaves the initialization done in the density module unchanged),</td>
</tr>
<tr>
<td></td>
<td>'zero' (put ( s = 0 ) explicitly; this may overwrite the</td>
</tr>
<tr>
<td></td>
<td>initialization done in the density module),</td>
</tr>
<tr>
<td></td>
<td>'isothermal' (isothermal stratification, ( T = \text{const} )),</td>
</tr>
<tr>
<td></td>
<td>'isobaric' (isobaric, ( p = \text{const} )),</td>
</tr>
<tr>
<td></td>
<td>'isentropic' (isentropic with superimposed hot [or cool] bubble),</td>
</tr>
<tr>
<td></td>
<td>'linprof' (linear entropy profile in ( z )),</td>
</tr>
<tr>
<td></td>
<td>'piecew-poly' (piecewise polytropic stratification for convection),</td>
</tr>
<tr>
<td></td>
<td>'polytropic' (polytropic stratification, polytropic exponent is ( m_{\text{poly0}} )),</td>
</tr>
<tr>
<td></td>
<td>'blob' (puts a gaussian blob in entropy for buoyancy experiments; see Ref. [7] for details)</td>
</tr>
<tr>
<td></td>
<td>'xjump' (jump in ( x ) direction),</td>
</tr>
<tr>
<td></td>
<td>'hor-tube' (horizontal flux tube in entropy, oriented in the ( y )-direction).</td>
</tr>
<tr>
<td>pertss ['zero']</td>
<td>additional perturbation to entropy. Currently valid choices are</td>
</tr>
<tr>
<td></td>
<td>'zero' (no perturbation)</td>
</tr>
<tr>
<td></td>
<td>'hexagonal' (hexagonal perturbation for convection).</td>
</tr>
<tr>
<td>ampl_ss [0.],</td>
<td>amplitude and width for some types of initial entropy.</td>
</tr>
<tr>
<td>widthss [2e]</td>
<td>initial entropy gradient for initss=linprof.</td>
</tr>
<tr>
<td>radius_ss [0.1]</td>
<td>radius of bubble for initss=isentropic.</td>
</tr>
<tr>
<td>mpoly0 [1.5],</td>
<td></td>
</tr>
<tr>
<td>mpoly1 [1.5],</td>
<td></td>
</tr>
</tbody>
</table>
K.1 Startup parameters for `start.in`

**mpoly2 [1.5]** specific to the solar convection case

- `inittss=piecew-poly`: polytropic indices of unstable (mpoly0), stable (mpoly1) and top layer (mpoly2). If the flag `isothtop` is set, the top layer is initialized to be isothermal, otherwise thermal (plus hydrostatic) equilibrium is assumed for all three layers, which results in a piecewise polytropic stratification.

**isothtop [0]** flag for isothermal top layer for `inittss=piecew-poly`.

**khor_ss [1.]** horizontal wave number for `pertss=hexagonal`.

<table>
<thead>
<tr>
<th>Namelist</th>
<th>magnetic_init_pars</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>initaa</strong> ['zero']</td>
<td>initialization of magnetic field (vector potential). Currently valid choices are</td>
</tr>
<tr>
<td><code>Alfven-x</code> (Alfvén wave traveling in the $x$-direction; this also sets the velocity),</td>
<td></td>
</tr>
<tr>
<td><code>Alfven-z</code> (Alfvén wave traveling in the $z$-direction; this also sets the velocity),</td>
<td></td>
</tr>
<tr>
<td><code>Alfvenz-rot</code> (same as <code>Alfven-z</code>, but with rotation),</td>
<td></td>
</tr>
<tr>
<td><code>Alfven-circ-x</code> (circularly polarized Alfvén wave in x direction),</td>
<td></td>
</tr>
<tr>
<td><code>Beltrami-x</code> ($x$-dependent Beltrami wave),</td>
<td></td>
</tr>
<tr>
<td><code>Beltrami-y</code> ($y$-dependent Beltrami wave),</td>
<td></td>
</tr>
<tr>
<td><code>Beltrami-z</code> ($z$-dependent Beltrami wave),</td>
<td></td>
</tr>
<tr>
<td><code>Bz(x)</code> ($B_z \propto \cos(kx)$),</td>
<td></td>
</tr>
<tr>
<td><code>crazy</code> (for testing purposes).</td>
<td></td>
</tr>
<tr>
<td><code>diffrot</code> ([needs to be documented]),</td>
<td></td>
</tr>
<tr>
<td><code>fluxrings</code> (two interlocked magnetic fluxrings; see §C.4),</td>
<td></td>
</tr>
<tr>
<td><code>gaussian-noise</code> (white noise),</td>
<td></td>
</tr>
<tr>
<td><code>halfcos-Bx</code> ([needs to be documented]),</td>
<td></td>
</tr>
<tr>
<td><code>hor-tube</code> (horizontal flux tube in $B$, oriented in the $y$-direction).</td>
<td></td>
</tr>
<tr>
<td><code>hor-fluxlayer</code> (horizontal flux layer),</td>
<td></td>
</tr>
<tr>
<td><code>mag-support</code> ([needs to be documented]),</td>
<td></td>
</tr>
<tr>
<td><code>mode</code> ([needs to be documented]),</td>
<td></td>
</tr>
<tr>
<td><code>modeb</code> ([needs to be documented]),</td>
<td></td>
</tr>
<tr>
<td><code>propto-ux</code> ([needs to be documented]),</td>
<td></td>
</tr>
<tr>
<td><code>propto-uy</code> ([needs to be documented]),</td>
<td></td>
</tr>
<tr>
<td><code>propto-uz</code> ([needs to be documented]),</td>
<td></td>
</tr>
<tr>
<td><code>sinxsinz</code> ($\sin x \sin z$),</td>
<td></td>
</tr>
<tr>
<td><code>uniform-Bx</code> (uniform field in $x$ direction),</td>
<td></td>
</tr>
<tr>
<td><code>uniform-By</code> (uniform field in $y$ direction),</td>
<td></td>
</tr>
<tr>
<td><code>uniform-Bz</code> (uniform field in $z$ direction),</td>
<td></td>
</tr>
<tr>
<td><code>zero</code> (zero field),</td>
<td></td>
</tr>
</tbody>
</table>
**initaa2 ['zero']**

additional perturbation of magnetic field. Currently valid choices are
- ‘zero’ (zero perturbation),
- ‘Beltrami-x’ (x-dependent Beltrami wave),
- ‘Beltrami-y’ (y-dependent Beltrami wave),
- ‘Beltrami-z’ (z-dependent Beltrami wave).

**amplaa [0.]**

amplitude for some types of initial magnetic fields.

**amplaa2 [0.]**

amplitude for some types of magnetic field perturbation.

**fring{1,2} [0.],**

**Iring{1,2} [0.],**

**Rring{1,2} [1.],**

**wr{1,2} [0.3]**

flux, current, outer and inner radius of flux ring 1/2; see Sect. C.4.

**radius [0.1]**

used by some initial fields.

**epsilonaa [10^{-2}]**

used by some initial fields.

**widthaa [0.5]**

used by some initial fields.

**z0aa [0.]**

used by some initial fields.

**kx_aa [1.],**

**ky_aa [1.],**

**kz_aa [1.]**

cnumbers used by some initial fields.

**lpress_equil [F]**

flag for pressure equilibrium (can be used in connection with all initial fields)

---

**Namelist pscalar_init_pars**

**initLncc ['zero']**

initialization of passive scalar (concentration per unit mass, c). Currently valid choices (for \( \ln c \)) are
- ‘zero’ (\( \ln c = 0. \)),
- ‘gaussian-noise’ (white noise),
- ‘wave-x’ (wave in x direction),
- ‘wave-y’ (wave in y direction),
- ‘wave-z’ (wave in z direction),
- ‘tang-discont-z’ (Kelvin-Helmholtz instability),
- ‘hor-tube’ (horizontal tube in concentration; used as a marker for magnetic flux tubes).

**initLncc2 ['zero']**

additional perturbation of passive scalar concentration c. Currently valid choices are
- ‘zero’ (\( \delta \ln c = 0. \)),
- ‘wave-x’ (add x-directed wave to \( \ln c \)).

**amplLncc [0.1]**

amplitude for some types of initial concentration.

**amplLncc2 [0.]**

amplitude for some types of concentration perturbation.

**kx_Lncc [1.],**

**ky_Lncc [1.],**

**kz_Lncc [1.]**

wave numbers for some types of initial concentration.

---

**Namelist shear_init_pars**
K.2 Runtime parameters for ‘run.in’

<table>
<thead>
<tr>
<th>Variable</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>qshear [0.]</td>
<td>degree of shear for shearing-box simulations (the shearing-periodic boundaries are the $x$-boundaries and are sheared in the $y$-direction). The shear velocity is $U = -q\Omega x \hat{y}$.</td>
</tr>
<tr>
<td>init_ads_mol_frac [0.]</td>
<td>initial adsorbed mole fraction</td>
</tr>
<tr>
<td>init_surf_mol_frac [0.]</td>
<td>initial surface mole fraction</td>
</tr>
<tr>
<td>total_carbon_sites [1.08e-8]</td>
<td>carbon sites per surface area [mol/cm]$^2$</td>
</tr>
<tr>
<td>dstalk [0.1]</td>
<td>times between printout of stalker data</td>
</tr>
<tr>
<td>lstalk_xx [F]</td>
<td>particles position</td>
</tr>
<tr>
<td>lstalk_vv [F]</td>
<td>particles velocity</td>
</tr>
<tr>
<td>lstalk_uu [F]</td>
<td>gas velocity at particles position</td>
</tr>
<tr>
<td>lstalk_guu [F]</td>
<td>gas velocity gradient at particles position</td>
</tr>
<tr>
<td>lstalk_rho [F]</td>
<td>gas density at particles position</td>
</tr>
<tr>
<td>lstalk_grho [F]</td>
<td>gas density gradient at particles position</td>
</tr>
<tr>
<td>lstalk_ap [F]</td>
<td>particles diameter</td>
</tr>
<tr>
<td>lstalk_bb [T]</td>
<td>magnetic field at particles position</td>
</tr>
<tr>
<td>lstalk_relvel [F]</td>
<td>particles relative velocity to gas</td>
</tr>
</tbody>
</table>

The following table lists all (at the time of writing, September 2002) namelists used in file ‘run.in’, with the corresponding parameters and their default values (in square brackets). Default values marked as [start] are taken from ‘start.in’. Any variable referred to as a flag can be set to any nonzero value to switch the corresponding feature on. Not all parameters are used for a given scenario. This list is not necessarily up to date; also, in many cases it can only give an idea of the corresponding setup; to get more insight and the latest set of parameters, you need to look at the code.

Once you have changed any of the ‘*.in’ files, you may want to first execute the command `pc_configtest` in order to test the correctness of these configuration files, before you apply them in an active simulation run.

<table>
<thead>
<tr>
<th>Variable [default value]</th>
<th>Namelist run_pars</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>cvsdr [''']</td>
<td></td>
<td>svn identification string, which allows you to keep track of the version of ‘run.in’.</td>
</tr>
<tr>
<td>nt [0]</td>
<td></td>
<td>number of time steps to run. This number can be increased or decreased during the run by touch RELOAD.</td>
</tr>
</tbody>
</table>
**it1 [10]** write diagnostic output every it1 time steps (see Sect. 5.5).

**it1d [it1]** write averages every it1d time steps (see Sect. 5.8.1). it1d has to be greater than or equal to it1.

**cdt [0.9]** Courant coefficient for advective time step; see §5.15.

**cdtv [0.25]** Courant coefficient for diffusive time step; see §5.15.

**dt [0.0]** time step; if ≠ 0, this overwrites the Courant time step. See §5.15 for a discussion of the latter.

**dtmin [10^{-6}]** abort if time step δt ≤ δtmin.

**tmax [10^{53}]** don’t run time steps beyond this time. Useful if you want to run for a given amount of time, but don’t know the necessary number of time steps.

**isave [100]** update current snapshot ‘var.dat’ every isave time steps.

**itorder [3]** order of time step (1 for Euler; 2 for 3rd-order, 3 for 3rd-order Runge–Kutta).

**dsnap [100.0]** save permanent snapshot every dsnap time units to files ‘VARN’, where N counts from N = 1 upward. (This information is stored in the file ‘data/tsnap.dat’; see the module wsnaps.f90, which in turn uses the subroutines out1 and out2).

**dvid [100.0]** write two-dimensional sections for generation of videos every dvid time units (not timesteps; see the subroutines out1 and out2 in the code).

**iwig [0]** if ≠ 0, apply a Nyquist filter (a filter eliminating any signal at the Nyquist frequency, but affecting large scales as little as possible) every iwig time steps to logarithmic density (sometimes necessary with convection simulations).

**ix [-1], iy [-1], iz [-1], iz2 [-1]** position of slice planes for video files. Any negative value of some of these variables will be overwritten according to the value of slice_position. See §5.7 for details.

**slice_position [‘p’]** symbolic specification of slice position. Currently valid choices are

- ‘p’ (periphery of the box)
- ‘m’ (middle of the box)
- ‘e’ (equator for half-sphere calculations, i.e. x, y centered, z bottom)

These settings are overridden by explicitly setting ix, iy, iz or iz2. See §5.7 for details.

**zbot_slice [value]** z position of slice xy-plane. The value can be any float number inside the z domain. These settings are overridden by explicitly setting ix, iy, iz or iz2. Saved as slice with the suffix xy. See §5.7 for details.

**ztop_slice [value]** z position of slice xy-plane. The value can be any float number inside the z domain. These settings are overridden by explicitly setting ix, iy, iz or iz2. Saved as slice with the suffix xy2. See §5.7 for details.
Runtime parameters for ‘run.in’

- **tavg [0]**: Averaging time \( \tau_{\text{avg}} \) for time averages (if \( \neq 0 \)); at the same time, time interval for writing time averages. See §5.8.4 for details.

- **idx_tavg \([0, 0, \ldots, 0] \)**: Indices of variables to time-average. See §5.8.4 for details.

- **d2davg [100.]**: Time interval for azimuthal and \( z \)-averages, i.e. the averages that produce 2d data. See §5.8.3 for details.

- **ialive [0]**: If \( \neq 0 \), each processor writes the current time step to ‘alive.info’ every \( ialive \) time steps. This provides the best test that the job is still alive. (This can be used to find out which node has crashed if there is a problem and the run is hanging.)

- **bcx \([\text{'p', 'p', } \ldots \text{ }] \)**, **bcy \([\text{'p', 'p', } \ldots \text{ }] \)**, **bcz \([\text{'p', 'p', } \ldots \text{ }] \)**: Boundary conditions. See Sect. 5.16 for a discussion of where and how to set these.

- **random_gen [start]**: See start parameters, p. 179

- **lwrite_aux [start]**: If set \( T \), auxiliary variables (those calculated at each step, but not evolved mathematically) to ‘var.dat’ and ‘VAR’ files after the evolved quantities.

**Namelist hydro_run_pars**

- **Omega [0.]**: Magnitude of angular velocity for Coriolis force (note: the centrifugal force is turned off by default, unless lcentrifugal_force=T is set).

- **theta [0.]**: Direction of angular velocity in degrees (\( \theta = 0 \) for \( z \)-direction, \( \theta = 90 \) for the negative \( x \)-direction, corresponding to a box located at the equator of a rotating sphere. Thus, e.g., \( \theta = 60 \) corresponds to 30° latitude. (Note: prior to April 29, 2007, there was a minus sign in the definition of \( \theta \).)

- **ttransient [0.]**: Initial time span for which to do something special (transient). Currently just used to smoothly switch on heating [Should be in run_pars, rather than here].

- **dampu [0.], tdamp [0.], ldamp_fade [F]**: Damp motions during the initial time interval \( 0 < t < t_{\text{damp}} \) with a damping term \(-dampu(u)\). If ldamp_fade is set, smoothly reduce damping to zero over the second half of the time interval tdamp. Initial velocity damping is useful for situations where initial conditions are far from equilibrium.

- **dampuint [0.], dampuext [0.], rdampint [0.], rdampext [impossible]**: Weighting of damping external to spherical region (see wdamp, damp below).

- **radius of internal damping region

- **radius of external damping region, used in place of former variable rdamp**
wdamp [0.2], permanently damp motions in $|x| < r_{\text{dampint}}$
with damping term $-damp_{\text{int}} u \chi(r-r_{\text{dampint}})$
or $|x| > r_{\text{dampext}}$ with damping term
$-damp_{\text{ext}} u \chi(r-r_{\text{dampext}})$, where $\chi(\cdot)$ is a smooth
profile of width wdamp.

ampl_forc [0.], amplitude of the ux-forcing or uy-forcing on the vertical
boundaries that is of the form $u_x(t) = ampl_{\text{forc}}\sin(k_{\text{forc}} x) \times \cos(w_{\text{forc}} t)$
[must be used in connection with bcx='g' or bcz='g' and force_lower_bound='vel_time' or force_upper_bound='vel_time']

k_forc [0.], corresponding horizontal wavenumber
w_forc [0.], corresponding frequency

density run pars

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>cs0 [start],</td>
<td>see start parameters, p. [181]</td>
</tr>
<tr>
<td>rho0 [start],</td>
<td></td>
</tr>
<tr>
<td>gamma [start]</td>
<td>Coefficient for mass diffusion (diffusion term will be $c_{\text{diffrho}} \delta x c_{s0}$).</td>
</tr>
<tr>
<td>cdiffrho [0.]</td>
<td></td>
</tr>
<tr>
<td>cs2bot [start],</td>
<td>squared sound speed at bottom and top for boundary condition 'c2'.</td>
</tr>
<tr>
<td>cs2top [start]</td>
<td></td>
</tr>
<tr>
<td>lupw_lnrho [false.]</td>
<td>use 5th-order upwind derivative operator for the advection term $u \cdot \nabla \ln \rho$ to avoid spurious Nyquist signal ('wiggles'); see §[H.2]</td>
</tr>
</tbody>
</table>

entropy run pars

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>hcond0 [0.],</td>
<td>specific to the solar convection case initss=piecew-poly; heat conductivities $K$ in the individual layers. $hcond0$ is the value $K_{\text{unst}}$ in the unstable layer, $hcond1$ is the ratio $K_{\text{stab}}/K_{\text{unst}}$ for the stable layer, and $hcond2$ is the ratio $K_{\text{top}}/K_{\text{unst}}$ for the top layer. The function $K(z)$ is not discontinuous, as the transition between the different values is smoothed over the width widthss. If $hcond1$ or $hcond2$ are not set, they are calculated according to the polytropic indices of the initial profile, $K \propto m+1$.</td>
</tr>
<tr>
<td>hcond1 [start],</td>
<td>select type of heat conduction. Currently valid choices are 'K-const' (constant heat conductivity), 'K-profile' (vertical or radial profile), 'chi-const' (constant thermal diffusivity), 'magnetic' (heat conduction by electrons in magnetic field – currently still experimental).</td>
</tr>
<tr>
<td>hcond2 [start]</td>
<td></td>
</tr>
<tr>
<td>iheatcond ['K-const']</td>
<td>flag for assuming thermal diffusivity $\chi = K/(c_p \rho) = const$, rather than $K = const$ (which is the default). This is currently only correct with 'noionization.f90'. Superseded by iheatcond.</td>
</tr>
</tbody>
</table>
chi [0.]
widthss [start]
isothtop [start]
luminosity [0.],
wheat [0.1]
cooltype ['Temp']

value of \( \chi \) when lcalc_heatcond_constchi=T.
width of transition region between layers. See start
parameters, p. 183
flag for isothermal top layer for solar convection case.
See start parameters, p. 183
strength and width of heating region.
type of cooling; currently only implemented for spherical
gallery. Currently valid choices are
'Temp','cs2' (cool temperature toward \( c_s^2 =
cool \) with a cooling term
\[ -C = -c_{cool} \frac{c_s^2 - c_{cool}^2}{c_{cool}^2} \]
)'Temp-rho','cs2-rho (cool temperature toward \( c_s^2 =
cool \) with a cooling term
\[ -C = -c_{cool} \rho \frac{c_s^2 - c_{cool}^2}{c_{cool}^2} \]
— this avoids numerical instabilities in low-
density regions [currently, the cooling coeffi-
cient \( c_{cool} = cool \) is not taken into account when
the time step is calculated])
'entropy' (cool entropy toward 0.).

cool [0.],
wcool [0.1]
rcool [1.]
Fbot [start]

strength \( c_{cool} \) and smoothing width of cooling region.
radius of cooling region: cool for \( |x| \geq rcool \).
heat flux for bottom boundary condition 'c1'. For
polytropic atmospheres, if Fbot is not set, it will be
calculated from the value of hcond0 in 'start.x', pro-
vided the entropy boundary condition is set to 'c1'.
entropy diffusion coefficient for diffusive term
\( \partial s / \partial t = \ldots + \chi_t \nabla^2 s \) in the entropy equation, that can
represent some kind of turbulent (sub-grid) mixing.
It is probably a bad idea to combine this with heat
conduction \( hcond0 \neq 0 \).

lupw_ss [.false.]

tauheat_buffer [0.]
zeheat_buffer [0.]
dheat_buffer1 [0.]
TTheat_buffer [0.]

use 5th-order upwind derivative operator for the ad-
vection term \( u \cdot \nabla s \) to avoid spurious Nyquist signal
('wiggles'); see §H.2
time scale for heating to target temperature
(=TTheat_buffer); zero disables the buffer zone.
\( z \) coordinate of the thermal buffer zone. Buffering is
active in \( |z| > TTheat_buffer \).
Inverse thickness of transition to buffered layer.
target temperature in thermal buffer zone (z direc-
tion only).
lhcond_global [F] flag for calculating the heat conductivity $K$ (and also $\nabla \log K$) globally using the global arrays facility. Only valid when $iheatcond='K$-profile'.

lread_hcond [F] flag for reading the conductivity profile and its derivative from the file “hcond_glhc.dat” or “hcond-_glhc.ascii”.

---

**Namelist magnetic_run_pars**

$B_{\text{ext}} [(0.,0.,0.)]$ uniform background magnetic field (for fully periodic boundary conditions, uniform fields need to be explicitly added, since otherwise the vector potential $A$ has a linear $x$-dependence which is incompatible with periodicity).

$lignore_{B_{\text{ext}}\text{ in } b2}$ [F] or $luse_{B_{\text{ext}}\text{ in } b2}$ [T] add uniform background magnetic field when computing $b^2$ pencils

eeta $[0.]$ magnetic diffusivity $\eta = 1/(\mu_0 \sigma)$, where $\sigma$ is the electric conductivity.

$\text{height}_\text{eeta} [0.,]$ used to add extra diffusivity in a halo region.

$\text{eta} \text{ out} [0.]$ used to add extra diffusivity inside sphere of radius $r_{\text{int}}$.

$\text{eta} \text{ ext} [0.]$ used to add extra diffusivity outside sphere of radius $r_{\text{ext}}$.

$\text{kinflow} ['']$ set type of flow fixed with ‘nohydro’. Currently the only recognized value is ‘ABC’ for an $ABC$ flow; all other values lead to $u=0$.

$kx [1.]$, $ky [1.]$, $kz [1.]$ wave numbers for $ABC$ flow.


---

**Namelist pscalar_run_pars**

$pscalar\_diff [0.]$ diffusion for passive scalar concentration $c$.

tensor_pscalar_diff [0.] coefficient for non-isotropic diffusion of passive scalar.

reinitialize_Incc [F] reinitialize the passive scalar to the value of $cc_{\text{const}}$ in start.in at next run

---

**Namelist forcing_run_pars**
K.2 Runtime parameters for ‘run.in’

**iforce [2]**
select form of forcing in the equation of motion; currently valid choices are
'zero’ (no forcing),
'irrotational’ (irrotational forcing),
'helical’ (helical forcing),
'fountain’ (forcing of “fountain flow”; see Ref. [14]),
'horizontal-shear’ (forcing localized horizontal sinusoidal shear).
'vevariable_gravz’ (time-dependent vertical gravity for forcing internal waves),

**iforce2 [0]**
select form of additional forcing in the equation of motion; valid choices are as for iforce.

**force [0.]**
amplitude of forcing.

**relhel [1.]**
helicity of forcing. The parameter relhel corresponds to $\sigma$ introduced in Sect. G.2 ($\sigma = \pm 1$ corresponds to maximum helicity of either sign).

**height_ff [0.]**
multiply forcing by $z$-dependent profile of width $height_ff$ (if $\neq 0$).

**r_ff [0.]**
if $\neq 0$, multiply forcing by spherical cutoff profile (of radius $r_ff$) and flip signs of helicity at equatorial plane.

**width_ff [0.5]**
width of vertical and radial profiles for modifying forcing.

**kfountain [5]**
horizontal wavenumber of the fountain flow.

**fountain [1.]**
amplitude of the fountain flow.

**omega_ff [1.]**
frequency of the cos or sin forcing [e.g. $\cos(omega_ff*t)$].

**ampl_ff [1.]**
amplitude of forcing in front of cos or sin [e.g. $ampl_ff*\cos(omega_ff*t)$].

---

**Namelist grav_run_pars**

**zref [start],**
**gravz [start],**
**grav_profile [start]**
see p. 181
**nu_epicycle [start]**
see Eq. (135) in Sect. C.5

---

**Namelist viscosity_run_pars**

**nu [0.]**
kinematic viscosity.

**nu_hyper2 [0.]**
kineamic hyperviscosity (with $\nabla^4 u$).

**nu_hyper3 [0.]**
kineamic hyperviscosity (with $\nabla^6 u$).

**zeta [0.]**
bulk viscosity.
ivisc ['nu-const']

select form of viscous term (see §6.2); currently valid choices are

'nu-const' – viscous force for \( \nu = \text{const} \), \( F_{\text{visc}} = \nu(\nabla^2 u + \frac{1}{3} \nabla \nabla \cdot u + 2S \cdot \nabla \ln \rho) \)

'rho_nu-const' – viscous force for \( \mu = \rho \nu = \text{const} \), \( F_{\text{visc}} = (\mu/\rho)(\nabla^2 u + \frac{1}{3} \nabla \nabla \cdot u) \). With this option, the input parameter \( \nu \) actually sets the value of \( \mu/\rho \) (\( \rho_0 = \rho \) is another input parameter, see pp. 181 and 188)

'simplified' – simplified viscous force \( F_{\text{visc}} = \nu \nabla^2 u \)

K.3 List of parameters for ‘print.in’

The following table lists all possible inputs to the file ‘print.in’ that are documented.

<table>
<thead>
<tr>
<th>Variable</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>it</td>
<td>number of time step (since beginning of job only)</td>
</tr>
<tr>
<td>t</td>
<td>time ( t ) (since start.csh)</td>
</tr>
<tr>
<td>dt</td>
<td>time step ( \delta t )</td>
</tr>
<tr>
<td>walltime</td>
<td>wall clock time since start of run.x, in seconds</td>
</tr>
<tr>
<td>Rmesh</td>
<td>( R_{\text{mesh}} )</td>
</tr>
<tr>
<td>Rmesh3</td>
<td>( R_{\text{mesh}}^{(3)} )</td>
</tr>
<tr>
<td>maxadvec</td>
<td>maxadvec</td>
</tr>
</tbody>
</table>

Module ‘cdat.f90’

Module ‘hydro.f90’
K.3 Parameters for ‘print.in’

\[ u_{2tm} = \langle u(t) \cdot \int_0^t u(t')dt' \rangle \]
\[ u_{otm} = \langle u(t) \cdot \int_0^t \omega(t')dt' \rangle \]
\[ outm = \langle \omega(t) \cdot \int_0^t u(t')dt' \rangle \]
\[ f_{kinzm} = \frac{1}{2} \langle \delta \mathbf{u}^2 \mathbf{u}_z \rangle \]
\[ u_{2m} = \langle \mathbf{u}^2 \rangle \]
\[ u_{xpt} = u_x(x_1, y_1, z_1, t) \]
\[ u_{ypt} = u_y(x_1, y_1, z_1, t) \]
\[ u_{zpt} = u_z(x_1, y_1, z_1, t) \]
\[ u_{xp2} = u_x(x_2, y_2, z_2, t) \]
\[ u_{yp2} = u_y(x_2, y_2, z_2, t) \]
\[ u_{zp2} = u_z(x_2, y_2, z_2, t) \]
\[ u_{uxuypt} = (u_x u_y)(x_1, y_1, z_1, t) \]
\[ u_{uyuzpt} = (u_y u_z)(x_1, y_1, z_1, t) \]
\[ u_{uzuxpt} = (u_z u_x)(x_1, y_1, z_1, t) \]
\[ u_{rms} = \langle \mathbf{u}^2 \rangle^{1/2} \]
\[ u_{rmsx} = \langle \mathbf{u}^2 \rangle^{1/2} \text{ for the hydro}_{xaver}\text{ range} \]
\[ u_{rmsz} = \langle \mathbf{u}^2 \rangle^{1/2} \text{ for the hydro}_{zaver}\text{ range} \]
\[ u_{durms} = \langle \delta \mathbf{u}^2 \rangle^{1/2} \]
\[ u_{max} = \max(||\mathbf{u}||) \]
\[ u_{min} = \min(||\mathbf{u}||) \]
\[ u_{xrms} = \langle u_x^2 \rangle^{1/2} \]
\[ u_{yrms} = \langle u_y^2 \rangle^{1/2} \]
\[ u_{zrms} = \langle u_z^2 \rangle^{1/2} \]
\[ u_{xmin} = \min(u_x) \]
\[ u_{ymin} = \min(u_y) \]
\[ u_{zmin} = \min(u_z) \]
\[ u_{xmax} = \max(u_x) \]
\[ u_{ymax} = \max(u_y) \]
\[ u_{zmax} = \max(u_z) \]
\[ u_{xm} = \langle u_x \rangle \]
\[ u_{ym} = \langle u_y \rangle \]
\[ u_{zm} = \langle u_z \rangle \]
\[ u_{uduum} = \langle \mathbf{u} \cdot \mathbf{u} \rangle \]
\[ u_{x2m} = \langle u_x^2 \rangle \]
\[ u_{y2m} = \langle u_y^2 \rangle \]
\[ u_{z2m} = \langle u_z^2 \rangle \]
\[ u_{x4m} = \langle u_x^4 \rangle \]
\[ u_{y4m} = \langle u_y^4 \rangle \]
\[ u_{z4m} = \langle u_z^4 \rangle \]
\[ u_{uxuy2m} = \langle u_x^2 u_y^2 \rangle \]
\[ u_{uyuz2m} = \langle u_y^2 u_z^2 \rangle \]
\[ u_{uxux2m} = \langle u_x^2 u_z^2 \rangle \]
\[ u_{ux2ccm} = \langle u_x^2 \cos^2 k_z \rangle \]
\[ u_{ux2ssm} = \langle u_x^2 \sin^2 k_z \rangle \]
\[ u_{uy2ccm} = \langle u_y^2 \cos^2 k_z \rangle \]
\[ u_{uy2ssm} = \langle u_y^2 \sin^2 k_z \rangle \]
\[ u_{uxuycsm} = \langle u_x u_y \cos k_z \sin k_z \rangle \]
\[ u_{uxuym} = \langle u_x u_y \rangle \]
uxuzm \langle u_x u_z \rangle
uyuzm \langle u_y u_z \rangle
umx \langle u_x \rangle
umy \langle u_y \rangle
umz \langle u_z \rangle
omumz \left\langle \langle W \rangle_x y \cdot \langle U \rangle_x y \right\rangle \quad (x y\text{-averaged mean cross helicity production})
umamz \left\langle \langle u \rangle_x y \cdot \langle A \rangle_x y \right\rangle
umbmz \left\langle \langle U \rangle_x y \cdot \langle B \rangle_x y \right\rangle \quad (x y\text{-averaged mean cross helicity production})
umxbmz \left\langle \langle U \rangle_x y \times \langle B \rangle_x y \right\rangle_z \quad (x y\text{-averaged mean emf})
rux2m \langle \rho u_x^2 \rangle \quad \text{(mean } x\text{-momentum density)}
ruy2m \langle \rho u_y^2 \rangle \quad \text{(mean } y\text{-momentum density)}
rutz2m \langle \rho u_z^2 \rangle \quad \text{(mean } z\text{-momentum density)}
divum \langle \text{div}u \rangle
rdivum \langle (\text{div}u)^2 \rangle
divu2m \langle (\text{grad} \text{div}u)^2 \rangle
u3u21m \langle u_3 u_{2,1} \rangle
u1u32m \langle u_1 u_{3,2} \rangle
u2u13m \langle u_2 u_{1,3} \rangle
u2u31m \langle u_{2,1} u_{3,1} \rangle
u3u12m \langle u_{3,1} u_{1,2} \rangle
u1u23m \langle u_{1,2,3} \rangle
ruxm \langle \rho u_x \rangle \quad \text{(mean } x\text{-momentum density)}
ruym \langle \rho u_y \rangle \quad \text{(mean } y\text{-momentum density)}
ruzm \langle \rho u_z \rangle \quad \text{(mean } z\text{-momentum density)}
ruxtot \langle \rho |u| \rangle \quad \text{(mean absolute } x\text{-momentum density)}
rumax \text{max}\langle \rho |u| \rangle \quad \text{(maximum modulus of momentum)}
ruxuyym \langle \rho u_x u_y \rangle \quad \text{(mean Reynolds stress)}
ruxuzm \langle \rho u_x u_z \rangle \quad \text{(mean Reynolds stress)}
ruyuzm \langle \rho u_y u_z \rangle \quad \text{(mean Reynolds stress)}
divrhourms \langle \nabla \cdot \langle \rho u \rangle \rangle_{\text{rms}}
divrhoumax \langle \nabla \cdot \langle \rho u \rangle \rangle_{\text{max}}
rlx2m \langle \rho u_z - z u_y \rangle
rlnym \langle \rho z u_x - x u_z \rangle
rlzm \langle \rho x u_y - y u_x \rangle
rlx2m \langle \rho u_z - z u_y \rangle^2
rl2m \langle \rho u_z - x u_x \rangle^2
rly2m \langle \rho u_z - x u_x \rangle^2
rll2m \langle \rho x u_y - y u_x \rangle^2
\text{tot\_ang\_mom} \quad \text{Total angular momentum in spherical coordinates about the axis.}
dtu \delta t / \langle c_s t \delta x / \text{max}|u| \rangle \quad \text{(time step relative to advective time step; see §5.15)}
oum \langle \omega \cdot u \rangle
ou_int \int V \omega \cdot u \, dV
fum \langle f \cdot u \rangle
odel2um \langle \omega \nabla^2 u \rangle
**K.3 Parameters for ‘print.in’**

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Formula</th>
</tr>
</thead>
<tbody>
<tr>
<td>( o2m )</td>
<td>( (\omega^2) \equiv \langle (\nabla \times u)^2 \rangle )</td>
</tr>
<tr>
<td>( orms )</td>
<td>( (\omega^2)^{1/2} )</td>
</tr>
<tr>
<td>( omax )</td>
<td>( \max(</td>
</tr>
<tr>
<td>( ox2m )</td>
<td>( (\omega_x^2) )</td>
</tr>
<tr>
<td>( oy2m )</td>
<td>( (\omega_y^2) )</td>
</tr>
<tr>
<td>( oz2m )</td>
<td>( (\omega_z^2) )</td>
</tr>
<tr>
<td>( oxuzxm )</td>
<td>( (\omega_x u_{z,x}) )</td>
</tr>
<tr>
<td>( oyuzym )</td>
<td>( (\omega_y u_{z,y}) )</td>
</tr>
<tr>
<td>( oxoym )</td>
<td>( (\omega_x \omega_y) )</td>
</tr>
<tr>
<td>( oxozm )</td>
<td>( (\omega_x \omega_z) )</td>
</tr>
<tr>
<td>( oyozm )</td>
<td>( (\omega_y \omega_z) )</td>
</tr>
<tr>
<td>( qfm )</td>
<td>( (q \cdot f) )</td>
</tr>
<tr>
<td>( q2m )</td>
<td>( (q^2) )</td>
</tr>
<tr>
<td>( qrms )</td>
<td>( (q^2)^{1/2} )</td>
</tr>
<tr>
<td>( qmax )</td>
<td>( \max(</td>
</tr>
<tr>
<td>( qom )</td>
<td>( (q \cdot \omega) )</td>
</tr>
<tr>
<td>( quxom )</td>
<td>( (q \cdot (u \times \omega)) )</td>
</tr>
<tr>
<td>( pvzm )</td>
<td>( (\omega_z + 2\Omega/\rho) ) (z component of potential vorticity)</td>
</tr>
<tr>
<td>( oumphi )</td>
<td>( (\omega \cdot u) \omega )</td>
</tr>
<tr>
<td>( ugsrjsxm )</td>
<td>( \langle (u \nabla u)^2 \rangle ) for the hydro_xaver_range</td>
</tr>
<tr>
<td>( ugu2m )</td>
<td>( u \nabla u^2 )</td>
</tr>
<tr>
<td>( ddx )</td>
<td>( \langle u_{xx} \rangle )</td>
</tr>
<tr>
<td>( Mrms )</td>
<td>( (u^2/c_s^2)^{1/2} ) (rms Mach number)</td>
</tr>
<tr>
<td>( Mamax )</td>
<td>( \max(</td>
</tr>
<tr>
<td>( EEK )</td>
<td>( \langle q u^2 \rangle /2 )</td>
</tr>
<tr>
<td>( ekin )</td>
<td>( \langle \frac{1}{2} \rho u^2 \rangle )</td>
</tr>
<tr>
<td>( ekin\text{tot} )</td>
<td>( \int V \frac{1}{2} \rho u^2 dV )</td>
</tr>
<tr>
<td>( uzxglnrym )</td>
<td>( \langle u_x \partial_y \ln \rho \rangle )</td>
</tr>
<tr>
<td>( uyglrxm )</td>
<td>( \langle u_y \partial_x \ln \rho \rangle )</td>
</tr>
<tr>
<td>( uzdivum )</td>
<td>( \langle u_z \nabla \cdot u \rangle )</td>
</tr>
<tr>
<td>( uxyydvium )</td>
<td>( \langle u_x u_y \nabla \cdot u \rangle )</td>
</tr>
<tr>
<td>( divuHrms )</td>
<td>( \langle (\nabla \cdot u_{H})^2 \rangle )</td>
</tr>
<tr>
<td>( uxxrms )</td>
<td>( u_{xx} )</td>
</tr>
<tr>
<td>( uyyrms )</td>
<td>( u_{yy} )</td>
</tr>
<tr>
<td>( uzxrms )</td>
<td>( u_{xz} )</td>
</tr>
<tr>
<td>( uyzrms )</td>
<td>( u_{yz} )</td>
</tr>
<tr>
<td>( uzyrms )</td>
<td>( u_{zy} )</td>
</tr>
</tbody>
</table>
| \( dtF \)          | \( \delta t/[c_{
acute{s}} \delta x/\max |F|] \) (time step relative to max force time step; see §5.15) |
| \( udpxxm \)       | components of symmetric tensor \( \langle u_i \partial_j p + u_j \partial_i p \rangle \) |

Module ‘density.f90’

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Formula</th>
</tr>
</thead>
<tbody>
<tr>
<td>( rhom )</td>
<td>( \langle \rho \rangle ) (mean density)</td>
</tr>
<tr>
<td>( rhomxmask )</td>
<td>( \langle \rho \rangle ) for the density_xaver_range</td>
</tr>
<tr>
<td>( rhomzmask )</td>
<td>( \langle \rho \rangle ) for the density_xaver_range</td>
</tr>
<tr>
<td>( rho2m )</td>
<td>( \langle \rho^2 \rangle )</td>
</tr>
<tr>
<td>( rho2m )</td>
<td>( \langle \rho^2 \rangle )</td>
</tr>
<tr>
<td>( drho2m )</td>
<td>( &lt; (\rho - \rho_0)^2 &gt; )</td>
</tr>
<tr>
<td>( drhom )</td>
<td>( &lt; \rho - \rho_0 &gt; )</td>
</tr>
<tr>
<td>( rhomin )</td>
<td>( \min(\rho) )</td>
</tr>
</tbody>
</table>
rhomax $\max(\rho)$
lnrhomin $\min(\log \rho)$
lnrho $\max(\log \rho)$
ugrhom $\langle u \cdot \nabla \phi \rangle$
uglnrhomin $\langle u \cdot \nabla \ln \rho \rangle$
totmass $\int \rho \, dV$
mass $\int \rho \, dV$
inertia $xx$ component of the inertia tensor (spherical coordinates)
inertia $yy$ component of the inertia tensor (spherical coordinates)
inertia $zz$ component of the inertia tensor (spherical coordinates)
inertia $xx$ component of the inertia tensor (Cartesian coordinates)
inertia $yy$ component of the inertia tensor (Cartesian coordinates)
inertia $zz$ component of the inertia tensor (Cartesian coordinates)
vol $\int dV$ (volume)
grhomax $\max(|\nabla \phi|)$

Module ‘entropy.f90’
dtc $\delta t /[c_{st}, \delta x / \max c_s]$ (time step relative to acoustic time step; see §5.15)
ethm $\langle \rho e \rangle$ (mean thermal [=internal] energy)
ssruzm $\langle s u_z / c_p \rangle$
ssuzm $\langle s u_z / c_p \rangle$
ssm $\langle s / c_p \rangle$ (mean entropy)
ss2m $\langle (s / c_p)^2 \rangle$ (mean squared entropy)
eem $\langle e \rangle$
ppm $\langle p \rangle$
csm $\langle c_s \rangle$
cmax $\max(c_s)$
cgam $\langle c_\gamma \rangle$
pdivum $\langle p \nabla \cdot u \rangle$
fradbot $\int F_{\text{bot}} \cdot dS$
fradtop $\int F_{\text{top}} \cdot dS$
TTtop $\int T_{\text{top}} dS$
ethtot $\int \rho e \, dV$ (total thermal [=internal] energy)
dtchi $\delta t / [c_{st,v} \delta x^2 / \chi_{\max}]$ (time step relative to time step based on heat conductivity; see §5.15)
Hmax $H_{\max}$ (net heat sources summed see §5.15)
tauhmin $H_{\max}$ (net heat sources summed see §5.15)
dtH $\delta t / [c_{st,s} c_v T / H_{\max}]$ (time step relative to time step based on heat sources; see §5.15)
yHm mean hydrogen ionization
yHmax max of hydrogen ionization
TTm $\langle T \rangle$
TTmax $T_{\max}$
TTmin $T_{\min}$
gTmax $\max(|\nabla T|)$
ssmax $s_{\max}$
ssmin $s_{\min}$
gTrms $(\nabla T)_{\text{rms}}$
gsrms $(\nabla s)_{\text{rms}}$
K.3 Parameters for ‘print.in’

\( g_{Txgsrms} \) \( (\nabla T \times \nabla s)_{rms} \)

\( f_{convm} \) \( (c_p\rho u_z T) \)

\( u_{fresm} \) \( (-u/\rho \nabla p) \)

\( K_{Kramersm} \) \( \langle K_{kramers} \rangle \)

\( ch_{krammin} \) \( \min(\chi_{kramers}) \)

\( ch_{krammax} \) \( \max(\chi_{kramers}) \)

Module ‘magnetic.f90’

\( \text{eta}_{tdep} \) \( t\)-dependent \( \eta \)

\( \text{ab}_{int} \) \( \int A \cdot B \, dV \)

\( \text{jb}_{int} \) \( \int j \cdot B \, dV \)

\( b_{2tm} \) \( \langle b(t) \cdot \int_0^t b(t')dt' \rangle \)

\( b_{jtm} \) \( \langle b(t) \cdot \int_0^t j(t')dt' \rangle \)

\( j_{btm} \) \( \langle j(t) \cdot \int_0^t b(t')dt' \rangle \)

\( b_{2ruzm} \) \( B^2 \rho u_z \)

\( b_{2uzm} \) \( B^2 u_z \)

\( u_{bbzm} \) \( (u \cdot B) B_z \)

\( b_{1m} \) \( \|B\| \)

\( b_{2m} \) \( B^2 \)

\( E_{EM} \) \( \langle B^2 \rangle / 2 \)

\( b_{4m} \) \( B^4 \)

\( b_{m2} \) \( \max(B^2) \)

\( j_{2m} \) \( \langle j^2 \rangle \)

\( j_{m2} \) \( \max(j^2) \)

\( a_{bm} \) \( \langle A \cdot B \rangle \)

\( a_{bumx} \) \( \langle u_x A \cdot B \rangle \)

\( a_{bumy} \) \( \langle u_y A \cdot B \rangle \)

\( a_{bumz} \) \( \langle u_z A \cdot B \rangle \)

\( a_{bhmh} \) \( \langle A \cdot B \rangle \) (temp)

\( a_{bmn} \) \( \langle A \cdot B \rangle \) (north)

\( a_{bms} \) \( \langle A \cdot B \rangle \) (south)

\( a_{brms} \) \( \langle (A \cdot B)^2 \rangle^{1/2} \)

\( j_{brms} \) \( \langle (j \cdot B)^2 \rangle^{1/2} \)

\( a_{jm} \) \( \langle j \cdot A \rangle \)

\( j_{bm} \) \( \langle j \cdot B \rangle \)

\( j_{bhmh} \) \( \langle J \cdot B \rangle \) (temp)

\( j_{bmn} \) \( \langle J \cdot B \rangle \) (north)

\( j_{bms} \) \( \langle J \cdot B \rangle \) (south)

\( u_{bm} \) \( \langle u \cdot B \rangle \)

\( u_{dubrms} \) \( \langle (u - B)^2 \rangle^{1/2} \)

\( d_{obrms} \) \( \langle (\omega - B)^2 \rangle^{1/2} \)

\( u_{xbxm} \) \( \langle u_x B_x \rangle \)

\( u_{ybxm} \) \( \langle u_y B_x \rangle \)

\( u_{zbxm} \) \( \langle u_z B_x \rangle \)

\( u_{xbym} \) \( \langle u_x B_y \rangle \)

\( u_{ybym} \) \( \langle u_y B_y \rangle \)

\( u_{zbym} \) \( \langle u_z B_y \rangle \)

\( u_{xbzm} \) \( \langle u_x B_z \rangle \)
uybzm  \langle u_y B_z \rangle \\
uzbzm  \langle u_z B_z \rangle \\
uxjxm  \langle u_x J_z \rangle \\
uuxjm  \langle u_x J_y \rangle \\
uxjzm  \langle u_x J_z \rangle \\
uyjxm  \langle u_y J_z \rangle \\
uyjym  \langle u_y J_y \rangle \\
uyjzm  \langle u_y J_z \rangle \\
uzjxm  \langle u_z J_z \rangle \\
uzjym  \langle u_z J_y \rangle \\
uzjzm  \langle u_z J_z \rangle \\
cosubm  \langle \mathbf{U} \cdot \mathbf{B} / (|\mathbf{U}| |\mathbf{B}|) \rangle \\
jxbxm  \langle j_x B_x \rangle \\
jybxm  \langle j_y B_x \rangle \\
jzbxm  \langle j_z B_x \rangle \\
jxbym  \langle j_x B_y \rangle \\
jybym  \langle j_y B_y \rangle \\
jzbym  \langle j_z B_y \rangle \\
jxbzm  \langle j_x B_z \rangle \\
jybzm  \langle j_y B_z \rangle \\
jzbzm  \langle j_z B_z \rangle \\
uam  \langle \mathbf{u} \cdot \mathbf{A} \rangle \\
ujm  \langle \mathbf{u} \cdot \mathbf{J} \rangle \\
fbm  \langle \mathbf{f} \cdot \mathbf{B} \rangle \\
fxbxm  \langle f_x B_x \rangle \\
epsM  \langle \eta \mu_0 J^2 \rangle \\
epsAD  \langle \rho^{-1} t_{AD} (\mathbf{J} \times \mathbf{B})^2 \rangle \text{ (heating by ion-neutrals friction)} \\
bxpt  B_x(x_1, y_1, z_1, t) \\
bypt  B_y(x_1, y_1, z_1, t) \\
.bzpt  B_z(x_1, y_1, z_1, t) \\
bxbypt  (B_x B_y)(x_1, y_1, z_1, t) \\
bybzpt  (B_y B_z)(x_1, y_1, z_1, t) \\
zbzpxpt  (B_z B_z)(x_1, y_1, z_1, t) \\
jxpt  J_x(x_1, y_1, z_1, t) \\
jypt  J_y(x_1, y_1, z_1, t) \\
jzpt  J_z(x_1, y_1, z_1, t) \\
Exp  \mathcal{E}_x(x_1, y_1, z_1, t) \\
Eyp  \mathcal{E}_y(x_1, y_1, z_1, t) \\
Ezp  \mathcal{E}_z(x_1, y_1, z_1, t) \\
axpt  A_x(x_1, y_1, z_1, t) \\
aypt  A_y(x_1, y_1, z_1, t) \\
azpt  A_z(x_1, y_1, z_1, t) \\
bxp2  B_x(x_2, y_2, z_2, t) \\
bypt  B_y(x_2, y_2, z_2, t) \\
.bzpt  B_z(x_2, y_2, z_2, t) \\
jxp2  J_x(x_2, y_2, z_2, t) \\
jypt  J_y(x_2, y_2, z_2, t) \\
jzpt  J_z(x_2, y_2, z_2, t) \\
Exp2  \mathcal{E}_x(x_2, y_2, z_2, t) \\
Eyp2  \mathcal{E}_y(x_2, y_2, z_2, t) \\
Ezp2  \mathcal{E}_z(x_2, y_2, z_2, t)
K.3 Parameters for ‘print.in’

\( E_{zp2} \) \( \mathcal{E}_z(x_2, y_2, z_2, t) \)
\( axp2 \) \( A_x(x_2, y_2, z_2, t) \)
\( ayp2 \) \( A_y(x_2, y_2, z_2, t) \)
\( azp2 \) \( A_z(x_2, y_2, z_2, t) \)
\( exabot \) \( \int \mathbf{E} \times \mathbf{A} \, dS \big|_{\text{bot}} \)
\( exatop \) \( \int \mathbf{E} \times \mathbf{A} \, dS \big|_{\text{top}} \)
\( emag \) \( \int_V \frac{1}{2\mu_0} B^2 \, dV \)
\( brms \) \( \left\langle B^2 \right\rangle^{1/2} \)
\( bfrms \) \( \left\langle B'^2 \right\rangle^{1/2} \)
\( bf2m \) \( \left\langle B'^2 \right\rangle \)
\( bf4m \) \( \left\langle B'^4 \right\rangle \)
\( bmax \) \( \max(\|B\|) \)
\( bxmin \) \( \min(\|B_x\|) \)
\( bymin \) \( \min(\|B_y\|) \)
\( bzmmin \) \( \min(\|B_z\|) \)
\( bmax \) \( \max(\|B_x\|) \)
\( bymax \) \( \max(\|B_y\|) \)
\( bzmmax \) \( \max(\|B_z\|) \)
\( bbxmax \) \( \max(\|B_x\|) \) \( \text{excluding } B_v \text{ ext} \)
\( bbymax \) \( \max(\|B_y\|) \) \( \text{excluding } B_v \text{ ext} \)
\( bbzmax \) \( \max(\|B_z\|) \) \( \text{excluding } B_v \text{ ext} \)
\( jxmax \) \( \max(\|j_v\|) \)
\( jymax \) \( \max(\|j_v\|) \)
\( jzmax \) \( \max(\|j_v\|) \)
\( jrms \) \( \left\langle J^2 \right\rangle^{1/2} \)
\( hjrms \) \( \left\langle J'^2 \right\rangle^{1/2} \)
\( jmax \) \( \max(\|j\|) \)
\( vArms \) \( \left\langle B^2/\theta \right\rangle^{1/2} \)
\( vAmax \) \( \max(\|B'^2/\theta\|^{1/2}) \)
\( dtb \) \( \delta t/[c_{sil} \delta x/v_{A,max}] \) \( \text{ (time step relative to Alfvén time step; see §5.15) } \)
\( dteta \) \( \delta t/[c_{sil,v} \delta x^2/\eta_{max}] \) \( \text{ (time step relative to resistive time step; see §5.15) } \)
\( dteta3 \) \( \delta t/[c_{sil,v} \delta x^6/\eta_{hyper}^2] \) \( \text{ (time step relative to hyper resistive time step; see §5.15) } \)
\( a2m \) \( \left\langle A^2 \right\rangle \)
\( arms \) \( \left\langle A^2 \right\rangle^{1/2} \)
\( amax \) \( \max(\|A\|) \)
\( divarms \) \( \left\langle (\nabla \cdot A)^2 \right\rangle^{1/2} \)
\( beta1m \) \( \left\langle B^2/(2\mu_0 p) \right\rangle \) \( \text{ (mean inverse plasma beta) } \)
\( beta1max \) \( \max[B^2/(2\mu_0 p)] \) \( \text{ (maximum inverse plasma beta) } \)
\( betam \) \( \langle \beta \rangle \)
\( betamax \) \( \max \beta \)
\( betamin \) \( \min \beta \)
\( bxm \) \( \langle B_x \rangle \)
\( bym \) \( \langle B_y \rangle \)
\( bzm \) \( \langle B_z \rangle \)
bxbym \langle B_xB_y \rangle \\

bm{x} \langle B_{yz}^2 \rangle^{1/2} \text{ (energy of } yz\text{-averaged mean field) } \\
bm{y} \langle B_{xz}^2 \rangle^{1/2} \text{ (energy of } xz\text{-averaged mean field) } \\
bm{z} \langle B_{xy}^2 \rangle^{1/2} \text{ (energy of } xy\text{-averaged mean field) } \\

bmzS2 \langle B_{xy}^2 \rangle \\

bmzA2 \langle B_{xy}^2 \rangle \\

jmx \langle J_{yz}^2 \rangle^{1/2} \text{ (energy of } yz\text{-averaged mean current density) } \\

jmy \langle J_{xz}^2 \rangle^{1/2} \text{ (energy of } xz\text{-averaged mean current density) } \\

jnz \langle J_{xy}^2 \rangle^{1/2} \text{ (energy of } xy\text{-averaged mean current density) } \\

bmzph \text{ Phase of a Beltrami field } \\

bmzphe \text{ Error of phase of a Beltrami field } \\
bsinphz \text{ sine of phase of a Beltrami field } \\
bcosphz \text{ cosine of phase of a Beltrami field } \\
emxamz3 \langle E_{xy} \times \langle A_{xy} \rangle \rangle \text{ (} xy\text{-averaged mean field helicity flux) } \\

embmz \langle E_{xy} \cdot \langle B_{xy} \rangle \rangle \text{ (} xy\text{-averaged mean field helicity production) } \\

ambmz \langle \langle A_{xy} \rangle \cdot \langle B_{xy} \rangle \rangle \text{ (magnetic helicity of } xy\text{-averaged mean field) } \\

ambmznh \langle \langle A_{xy} \rangle \cdot \langle B_{xy} \rangle \rangle \text{ (magnetic helicity of } xy\text{-averaged mean field, temp) } \\

ambmzn \langle \langle A_{xy} \rangle \cdot \langle B_{xy} \rangle \rangle \text{ (magnetic helicity of } xy\text{-averaged mean field, north) } \\

ambmzns \langle \langle A_{xy} \rangle \cdot \langle B_{xy} \rangle \rangle \text{ (magnetic helicity of } xy\text{-averaged mean field, south) } \\

jmbmz \langle \langle J_{xy} \rangle \cdot \langle B_{xy} \rangle \rangle \text{ (current helicity of } xy\text{-averaged mean field) } \\

Rmz \langle \frac{u \times B}{\|u \times J\|} \rangle_{xy} \\

kx_\text{aa} k_x \\
kx \langle J_{xy} \rangle \cdot \langle B_{xy} \rangle / \langle B_{xy}^2 \rangle \\
bm{x2} \langle B_{x}^2 \rangle \\
bm{y2} \langle B_{y}^2 \rangle \\
bm{z2} \langle B_{z}^2 \rangle \\
bm{x4} \langle B_{x}^4 \rangle \\
bm{y4} \langle B_{y}^4 \rangle \\
bm{z4} \langle B_{z}^4 \rangle \\
jx_{2} \langle J_{x}^2 \rangle \\
jy_{2} \langle J_{y}^2 \rangle \\
jz_{2} \langle J_{z}^2 \rangle \\
jx_{4} \langle J_{x}^4 \rangle \\
jy_{4} \langle J_{y}^4 \rangle \\
jz_{4} \langle J_{z}^4 \rangle \\
uxb \langle u \times B \rangle \cdot B_0/B_0^2
K.3 Parameters for ‘print.in’

\( j \times B \cdot B_0/B_0^2 \)
\( \nu_{\text{mag}} \max(1/|j \times B/B^2|) \)
\( \nu_{\text{mag}} \left\langle \frac{1}{|j \times B/B^2|} \right\rangle^{1/2} \)

\( B_{3B_{21}} \)
\( B_{3B_{12}} \)
\( B_{B_{13}2} \)
\( B_{2B_{13}} \)
\( B_{B_{23}1} \)
\( B_{B_{31}2} \)
\( (u \times B)_x \)
\( (u \times B)_y \)
\( (u \times B)_z \)
\( (j \times B)_x \)
\( (j \times B)_y \)
\( (j \times B)_z \)
\( (E \times A)_x \)
\( (E \times A)_y \)
\( (E \times A)_z \)
\( (E \times J)_x \)
\( (E \times J)_y \)
\( (E \times J)_z \)
\( \nabla \times E \times B \big|_x \)
\( \nabla \times E \times B \big|_y \)
\( \nabla \times E \times B \big|_z \)
\( \phi B \big|_x \)
\( \phi B \big|_y \)
\( \phi B \big|_z \)
\( B^2 \nabla \cdot u \)
\( \left( J \cdot \nabla^2 A \right) \)
\( \left( u \cdot (J \times B) \right) \)
\( \max(|J \times B/\rho|) \)
\( \left( J \times B/\rho \right)^2 \)
\( \sqrt{[(b_x)_z(x,y)]^2 + [(b_y)_z(x,y)]^2 + [(b_z)_z(x,y)]^2} \)
\( \text{Mean of Smagorinsky resistivity} \)
\( \text{Min of Smagorinsky resistivity} \)
\( \text{Max of Smagorinsky resistivity} \)
\( \text{Max of artificial resistivity } \eta \sim v_A \)
\( \text{Max of artificial resistivity } \eta \sim J/\sqrt{\rho} \)
\( \text{Max of artificial resistivity } \eta \sim J^2/\rho \)
\( \eta_1 \)
\( \left( J \cdot B/(|J||B|) \right) \)
\( \text{Mean value of the component of J parallel to B} \)
\( \text{Mean value of the component of J perpendicular to B} \)
\( \text{Mean value of the component of J_hyper parallel to B} \)
\( \text{Mean value of the component of J_hyper perpendicular to B} \)
\( \langle B^2 \rangle^{1/2} \text{ for the magnetic_xaver_range} \)
\( \langle B^2 \rangle^{1/2} \text{ for the magnetic_zaver_range} \)
\( \langle \mathcal{E}_x \rangle_z \)
The Pencil Code

\[ E_{\text{ymxy}} \quad \langle \mathcal{E}_y \rangle_z \]
\[ E_{\text{zmxy}} \quad \langle \mathcal{E}_z \rangle_z \]

**Module ‘pscalar.f90’**

- rhoccm \( \langle \rho \rangle \)
- ccmax \( \max(c) \)
- ceglnrm \( \langle c \nabla_{z} \rangle \)

**Module ‘1D_loop.f90’**

- dtchi2 heat conduction
- dtrad radiative loss from RTV
- dtspitzer Spitzer heat conduction time step
- qmax max of heat flux vector
- qrms rms of heat flux vector

**Module ‘advective_gauge.f90’**

- Lamm \( \langle \Lambda \rangle \)
- Lamp \( \Lambda(x_1, y_1, z_1) \)
- Lam2 \( \Lambda(x_2, y_2, z_2) \)
- Lamrms \( \langle \Lambda^2 \rangle^{1/2} \)
- Lambzm \( \langle \Lambda B_z \rangle \)
- Lambzmz \( \langle \Lambda B_z \rangle_{xy} \)
- gLambm \( \langle \Lambda B \rangle \)
- apbrms \( \langle (A' B) \rangle^{1/2} \)
- jxarms \( \langle (J \times A) \rangle^{1/2} \)
- jxaprms \( \langle (J \times A') \rangle^{1/2} \)
- jxglamrms \( \langle (J \times \nabla \Lambda) \rangle^{1/2} \)
- glamrms \( \langle (\nabla \Lambda) \rangle^{1/2} \)
- divabrms \( \langle (\nabla \cdot A) \rangle B \rangle^{1/2} \)
- divapbrms \( \langle (\nabla \cdot A') \rangle B \rangle^{1/2} \)
- d2Lambrms \( \langle (\nabla^2 \Lambda) \rangle B \rangle^{1/2} \)
- d2Lamrms \( \langle (\nabla^2 \Lambda) \rangle^{1/2} \)

**Module ‘anelastic.f90’**

- rhom \( \langle \rho \rangle \) (mean density)
- ugrhm \( \langle u \cdot \nabla \rho \rangle \)
- mass \( \int \rho \, dV \)
- divrhom \( \langle \nabla \cdot (\rho u) \rangle \)
- divrhoum \( \nabla \cdot (\rho u) \rangle_{\text{rms}} \)
- divrhoumax \( \nabla \cdot (\rho u) \rangle_{\text{max}} \)

**Module ‘bfield.f90’**

- bmax \( \max B \)
- bmin \( \min B \)
- brms \( \langle B^2 \rangle^{1/2} \)
- bm \( \langle B \rangle \)
- b2m \( \langle B^2 \rangle \)
- bxmax \( \max |B_x| \)
- bymax \( \max |B_y| \)
K.3 Parameters for ‘print.in’

\begin{align*}
    \text{bzmax} & \quad \max |B_z| \\
    \text{bzm} & \quad \langle B_z \rangle \\
    \text{bym} & \quad \langle B_y \rangle \\
    \text{bzm} & \quad \langle B_z \rangle \\
    \text{bx2m} & \quad \langle B_x^2 \rangle \\
    \text{by2m} & \quad \langle B_y^2 \rangle \\
    \text{bz2m} & \quad \langle B_z^2 \rangle \\
    \text{bxym} & \quad \langle B_x B_y \rangle \\
    \text{bxzm} & \quad \langle B_x B_z \rangle \\
    \text{bym} & \quad \langle B_y \rangle \\
    \text{dbx} & \quad \text{max} |B_x - B_{\text{ext},x}| \\
    \text{dbym} & \quad \text{max} |B_y - B_{\text{ext},y}| \\
    \text{dbzmax} & \quad \text{max} |B_z - B_{\text{ext},z}| \\
    \text{dxb} & \quad \langle B_x - B_{\text{ext},x} \rangle \\
    \text{dbym} & \quad \langle B_y - B_{\text{ext},y} \rangle \\
    \text{dbzm} & \quad \langle B_z - B_{\text{ext},z} \rangle \\
    \text{dbx2m} & \quad \langle (B_x - B_{\text{ext},x})^2 \rangle \\
    \text{dby2m} & \quad \langle (B_y - B_{\text{ext},y})^2 \rangle \\
    \text{dbz2m} & \quad \langle (B_z - B_{\text{ext},z})^2 \rangle \\
    \text{jmax} & \quad \text{max} J \\
    \text{jmin} & \quad \text{min} J \\
    \text{jrms} & \quad \langle J^2 \rangle^{1/2} \\
    \text{jm} & \quad \langle J \rangle \\
    \text{jm} & \quad \langle J^2 \rangle \\
    \text{jxmax} & \quad \text{max} |J_x| \\
    \text{jymax} & \quad \text{max} |J_y| \\
    \text{jzmax} & \quad \text{max} |J_z| \\
    \text{jx} & \quad \langle J_x \rangle \\
    \text{jy} & \quad \langle J_y \rangle \\
    \text{jm} & \quad \langle J_z \rangle \\
    \text{jx2m} & \quad \langle J_x^2 \rangle \\
    \text{jy2m} & \quad \langle J_y^2 \rangle \\
    \text{jjzm} & \quad \langle J_z^2 \rangle \\
    \text{divbmax} & \quad \text{max} |\nabla \cdot B| \\
    \text{divbrms} & \quad \langle (\nabla \cdot B)^2 \rangle^{1/2} \\
    \text{betamax} & \quad \text{max} \beta \\
    \text{betamin} & \quad \text{min} \beta \\
    \text{betam} & \quad \langle \beta \rangle \\
    \text{vAm} & \quad \text{max} v_A \\
    \text{vAmin} & \quad \text{min} v_A \\
    \text{vAm} & \quad \langle v_A \rangle \\
\end{align*}

Module ‘chemistry.f90’

\textit{dtchem} \quad \textit{dt\_chem}

Module ‘chemistry\_simple.f90’

\textit{dtchem} \quad \textit{dt\_chem}

Module ‘chiral\_mhd.f90’

\textit{muSm} \quad \langle \mu_S \rangle
\[ \mu_{5\text{rms}} \langle \mu_5^2 \rangle^{1/2} \]
\[ \mu_5 \langle \mu_5 \rangle \]
\[ \mu_{5\text{lm}} \langle |\mu_5| \rangle \]
\[ \mu_{5\text{rms}} \langle \mu_5^2 \rangle^{1/2} \]
\[ \mu_{5\text{min}} \min \mu_5 \]
\[ \mu_{5\text{max}} \max \mu_5 \]
\[ \mu_{5\text{abs}} \max |\mu_5| \]
\[ g\mu_{5\text{rms}} \langle (\nabla \mu_5)^2 \rangle^{1/2} \]
\[ g\mu_{5\text{Srms}} \langle (\nabla \mu_5)^2 \rangle^{1/2} \]
\[ g\mu_{5\text{mx}} \langle \nabla \mu_5 \rangle_x \]
\[ g\mu_{5\text{my}} \langle \nabla \mu_5 \rangle_y \]
\[ g\mu_{5\text{mz}} \langle \nabla \mu_5 \rangle_z \]
\[ b\mu_{5\text{rms}} \langle (B \cdot \nabla \mu_5)^2 \rangle^{1/2} \]
\[ b\mu_{5\text{Srms}} \langle (B \cdot \nabla \mu_5)^2 \rangle^{1/2} \]
\[ b\mu_{5\text{bjm}} \langle \mu_5 ((\nabla \times B) \cdot B) \rangle \]
\[ b\mu_{5\text{bjrms}} \langle \mu_5 ((\nabla \times B) \cdot B) \rangle^{1/2} \]
\[ dt_\lambda \min(\mu_5 / B) \]
\[ dt_\lambda \min(B^2) \]
\[ dt_D \min(\mu_5 / B) \]
\[ dt_{\text{CMW}} \delta x / \left( (C_p C_s)^{1/2} \max(|B|) \right) \]
\[ dt_{\text{Dmu}} \min(\mu_5 / B^2) \delta x / (\lambda \eta) \]
\[ dt_\gamma \min(B^2) \]
\[ dt_{\text{chiral}} \] total time-step contribution from chiral MHD
\[ \mu_{5\text{bxm}} \langle \mu_5 B_x \rangle \]
\[ \mu_{5\text{b2m}} \langle \mu_5 B^2 \rangle \]
\[ j_{\text{xm}} \langle J_x \rangle \]

---

**Module ‘coronae.f90’**

\[ dt_{\text{chi2}} \delta t / [c_{\text{St},v} \delta x^2 / \lambda_{\text{max}}] \] (time step relative to time step based on heat conductivity; see §5.15)

\[ dt_{\text{spitzer}} \] Spitzer heat conduction time step

\[ dt_{\text{trad}} \] radiative loss from RTV

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**Module ‘cosmicray_current.f90’**

\[ e_{\text{kin}} \langle 1/2 \rho u_{\text{cr}}^2 \rangle \]

---

**Module ‘density_stratified.f90’**

\[ \int \rho \, d^3x \]

\[ \rho_{\text{min}} \min |\rho| \]

\[ \rho_{\text{max}} \max |\rho| \]

\[ d\rho_{\text{m}} \langle \Delta \rho / \rho_0 \rangle \]

\[ d\rho_{\text{rms}} \langle (\Delta \rho / \rho_0)^2 \rangle \]

\[ d\rho_{\text{homs}} \langle \Delta \rho / \rho_0 \rangle_{\text{rms}} \]

---

**Module ‘detonate.f90’**

\[ \text{detn} \] Number of detonated sites (summed over time steps between adjacent outputs)
<table>
<thead>
<tr>
<th>Parameter</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>\textit{dettot}</td>
<td>Total energy input (summed over time steps between adjacent outputs)</td>
</tr>
</tbody>
</table>

**Module `dustdensity.f90`**

- \(K\text{Km}\) \(\sum T_k^{\text{coag}}\)
- \(K\text{K2m}\) \(\sum T_k^{\text{coag}}\)
- \(M\text{Mxm}\) \(\sum M_x^{\text{coag}}\)
- \(M\text{Mym}\) \(\sum M_y^{\text{coag}}\)
- \(M\text{Mzm}\) \(\sum M_z^{\text{coag}}\)

**Module `entropy_anelastic.f90`**

- \(dtc\) \(\delta t / [c_\text{d}, \delta x / \text{max} c_\text{s}]\) (time step relative to acoustic time step; see \(\S 5.15\))
- \(\text{ethm}\) \(\langle \rho e \rangle\) (mean thermal [=internal] energy)
- \(\text{ssm}\) \(\langle s/c_p \rangle\) (mean entropy)
- \(\text{ss2m}\) \(\langle (s/c_p)^2 \rangle\) (mean squared entropy)
- \(\text{eem}\) \(\langle e \rangle\)
- \(\text{ppm}\) \(\langle p \rangle\)
- \(\text{csm}\) \(\langle c_s \rangle\)
- \(\text{pdivum}\) \(\langle p \nabla u \rangle\)
- \(\text{fradbot}\) \(\int F_{\text{bot}} \cdot dS\)
- \(\text{fradtop}\) \(\int F_{\text{top}} \cdot dS\)
- \(\text{TTtop}\) \(\int T_{\text{top}} dS\)
- \(\text{ethtot}\) \(\int V \rho e dV\) (total thermal [=internal] energy)
- \(dtchi\) \(\delta t / [c_\text{d}, \delta x^2 / \chi_{\text{max}}]\) (time step relative to time step based on heat conductivity; see \(\S 5.15\))
- \(\text{ssmxy}\) \(\langle s \rangle_z\)
- \(\text{ssmxz}\) \(\langle s \rangle_y\)

**Module `gravitational_waves.f90`**

- \(\text{hhT2m}\) \(\langle h_T^2 \rangle\)
- \(\text{hhX2m}\) \(\langle h_X^2 \rangle\)
- \(\text{hhThhXm}\) \(\langle h_T h_X \rangle\)
- \(\text{ggTpt}\) \(g_T(x_1, y_1, z_1, t)\)
- \(\text{strTpt}\) \(S_T(x_1, y_1, z_1, t)\)
- \(\text{strXpt}\) \(S_X(x_1, y_1, z_1, t)\)

**Module `gravitational_waves_hTXk.f90`**

- \(\text{g11pt}\) \(g_{11}(x_1, y_1, z_1, t)\)
- \(\text{g22pt}\) \(g_{22}(x_1, y_1, z_1, t)\)
- \(\text{g33pt}\) \(g_{33}(x_1, y_1, z_1, t)\)
- \(\text{g12pt}\) \(g_{12}(x_1, y_1, z_1, t)\)
- \(\text{g23pt}\) \(g_{23}(x_1, y_1, z_1, t)\)
- \(\text{g31pt}\) \(g_{31}(x_1, y_1, z_1, t)\)
- \(\text{hhTpt}\) \(h_T(x_1, y_1, z_1, t)\)
- \(\text{hhXpt}\) \(h_X(x_1, y_1, z_1, t)\)
- \(\text{ggTpt}\) \(\dot{h}_T(x_1, y_1, z_1, t)\)
- \(\text{ggXpt}\) \(\dot{h}_X(x_1, y_1, z_1, t)\)
- \(\text{hhTp2}\) \(h_T(x_1, y_1, z_1, t)\)
- \(\text{hhXp2}\) \(h_X(x_1, y_1, z_1, t)\)
\[ \begin{align*}
  & \text{ggTp2} & \hat{h}_T(x_1, y_1, z_1, t) \\
  & \text{ggXp2} & \hat{h}_X(x_1, y_1, z_1, t) \\
  & \text{hrms} & \langle h_T^2 + h_X^2 \rangle^{1/2} \\
  & \text{EEGW} & \langle g_T^2 + g_X^2 \rangle c^2 / (32\pi G) \\
  & \text{gg2m} & \langle g_T^2 \rangle \\
  & \text{hhT2m} & \langle h_T^2 \rangle \\
  & \text{hhX2m} & \langle h_X^2 \rangle \\
  & \text{hhTXm} & \langle h_T h_X \rangle \\
  & \text{ggT2m} & \langle g_T^2 \rangle \\
  & \text{ggX2m} & \langle g_X^2 \rangle \\
  & \text{ggTXm} & \langle g_T g_X \rangle \\
\end{align*} \]

Module 'gravitational_waves_hTXk_no_xpara.f90'

\[ \begin{align*}
  & \text{g11pt} & g_{11}(x_1, y_1, z_1, t) \\
  & \text{g22pt} & g_{22}(x_1, y_1, z_1, t) \\
  & \text{g33pt} & g_{33}(x_1, y_1, z_1, t) \\
  & \text{g12pt} & g_{12}(x_1, y_1, z_1, t) \\
  & \text{g23pt} & g_{23}(x_1, y_1, z_1, t) \\
  & \text{g31pt} & g_{31}(x_1, y_1, z_1, t) \\
  & \text{hhTpt} & h_T(x_1, y_1, z_1, t) \\
  & \text{hhXpt} & h_X(x_1, y_1, z_1, t) \\
  & \text{ggTpt} & \hat{h}_T(x_1, y_1, z_1, t) \\
  & \text{ggXpt} & \hat{h}_X(x_1, y_1, z_1, t) \\
  & \text{hhT2m} & \langle h_T^2 \rangle \\
  & \text{hhX2m} & \langle h_X^2 \rangle \\
  & \text{hhTXm} & \langle h_T h_X \rangle \\
  & \text{gg2m} & \langle g_T^2 \rangle \\
  & \text{ggX2m} & \langle g_X^2 \rangle \\
  & \text{ggTXm} & \langle g_T g_X \rangle \\
\end{align*} \]

Module 'gravitational_waves_hij6.f90'

\[ \begin{align*}
  & \text{h22rms} & \xi_{22}^{\text{rms}} \\
  & \text{h33rms} & \xi_{33}^{\text{rms}} \\
  & \text{h23rms} & \xi_{23}^{\text{rms}} \\
  & \text{g11pt} & g_{11}(x_1, y_1, z_1, t) \\
  & \text{g22pt} & g_{22}(x_1, y_1, z_1, t) \\
  & \text{g33pt} & g_{33}(x_1, y_1, z_1, t) \\
  & \text{g12pt} & g_{12}(x_1, y_1, z_1, t) \\
  & \text{g23pt} & g_{23}(x_1, y_1, z_1, t) \\
  & \text{g31pt} & g_{31}(x_1, y_1, z_1, t) \\
  & \text{hhTpt} & h_T(x_1, y_1, z_1, t) \\
  & \text{hhXpt} & h_X(x_1, y_1, z_1, t) \\
  & \text{ggTpt} & \hat{h}_T(x_1, y_1, z_1, t) \\
\end{align*} \]
K.3 Parameters for ‘print.in’

\[ \dot{h}_X(x_1, y_1, z_1, t) \]

\[ h_T(x_1, y_1, z_1, t) \]

\[ h_X(x_1, y_1, z_1, t) \]

\[ \dot{h}_X(x_1, y_1, z_1, t) \]

\[ \dot{h}_X(x_1, y_1, z_1, t) \]

\[ (g_T^2 + g_X^2) c^2 / (32 \pi G) \]

\[ (g_T^2 + g_X^2) \]

\[ (g_T^2) \]

\[ (g_X^2) \]

\[ (g_T^2) \]

\[ (g_X^2) \]

\[ (g_T^2) \]

\[ (g_X^2) \]

\[ (h^2_{ij,ij}) \]

\[ (u \cdot g) \]

Module ‘gravity_simple.f90’

\[ \langle \phi_{\text{grav}} \rangle \quad \text{(mean potential energy)} \]

\[ \int_V \phi_{\text{grav}} dV \quad \text{(total potential energy)} \]

\[ \langle u \cdot g \rangle \]

Module ‘heatflux.f90’

\[ dt_{\text{spitzer}} \quad \text{Spitzer heat conduction time step} \]

\[ dtq \quad \text{heatflux time step} \]

\[ dtq2 \quad \text{heatflux time step due to \( \tau \)} \]

\[ q_{\text{max}} \quad \text{max(\( |q| \))} \]

\[ \tau_{\text{max}} \quad \text{max(\( |\tau_{\text{Spitzer}}| \))} \]

\[ q_{\text{xmin}} \quad \text{min(\( |q_x| \))} \]

\[ q_{ymin} \quad \text{min(\( |q_y| \))} \]

\[ q_{zmin} \quad \text{min(\( |q_z| \))} \]

\[ q_{\text{xmax}} \quad \text{max(\( |q_x| \))} \]

\[ q_{ymax} \quad \text{max(\( |q_y| \))} \]

\[ q_{zmax} \quad \text{max(\( |q_z| \))} \]

\[ q_{\text{rms}} \quad \sqrt{\langle |q|^2 \rangle} \]

\[ q_{\text{satmin}} \quad \text{minimum of qsat/abs} \]

\[ q_{\text{satrms}} \quad \text{rms of qsat/abs} \]

Module ‘hydro_potential.f90’

\[ \langle u(t) \cdot \int_0^t u(t') dt' \rangle \]

\[ \frac{1}{2} \langle uu_z \rangle \]

\[ \langle u_z \rangle \]

\[ u_x(x_1, y_1, z_1, t) \]

\[ u_y(x_1, y_1, z_1, t) \]

\[ u_z(x_1, y_1, z_1, t) \]

\[ u_x(x_2, y_2, z_2, t) \]

\[ u_y(x_2, y_2, z_2, t) \]

\[ u_z(x_2, y_2, z_2, t) \]
urms $\langle u^2 \rangle^{1/2}$
urmsx $\langle u^2 \rangle^{1/2}$ for the hydro_xaver_range
urmsz $\langle u^2 \rangle^{1/2}$ for the hydro_zaver_range
durms $\langle \delta u^2 \rangle^{1/2}$
umax $\max(|u|)$
umin $\min(|u|)$
uxrms $\langle u_x^2 \rangle^{1/2}$
uyrms $\langle u_y^2 \rangle^{1/2}$
uzrms $\langle u_z^2 \rangle^{1/2}$
uxmin $\min(|u_x|)$
uymin $\min(|u_y|)$
uzmin $\min(|u_z|)$
uxmax $\max(|u_x|)$
uymax $\max(|u_y|)$
uzmax $\max(|u_z|)$
ux $\langle u_x \rangle$
uy $\langle u_y \rangle$
uz $\langle u_z \rangle$
ux2m $\langle u_x^2 \rangle$
uy2m $\langle u_y^2 \rangle$
uz2m $\langle u_z^2 \rangle$
ux2ccm $\langle u_x^2 \cos^2 k z \rangle$
ux2ssm $\langle u_x^2 \sin^2 k z \rangle$
uy2ccm $\langle u_y^2 \cos^2 k z \rangle$
uy2ssm $\langle u_y^2 \sin^2 k z \rangle$
uxuyccsm $\langle u_x u_y \cos k z \sin k z \rangle$
uxuy $\langle u_x u_y \rangle$
uxuzm $\langle u_x u_z \rangle$
uyuzm $\langle u_y u_z \rangle$
uxm $\langle u_x \rangle$
umy $\langle u_y \rangle$
umz $\langle u_z \rangle$
omumz $\langle \langle W \rangle_{xy} \cdot \langle U \rangle_{xy} \rangle$ (xy-averaged mean cross helicity production)

umamz $\langle \langle u \rangle_{xy} \cdot \langle A \rangle_{xy} \rangle$

umbmz $\langle \langle U \rangle_{xy} \cdot \langle B \rangle_{xy} \rangle$ (xy-averaged mean cross helicity production)

umxbmz $\langle \langle U \rangle_{xy} \times \langle B \rangle_{xy} \rangle_z$ (xy-averaged mean emf)
rux2m $\langle \rho u_x^2 \rangle$
ruy2m $\langle \rho u_y^2 \rangle$
ruz2m $\langle \rho u_z^2 \rangle$
divum $\langle \text{div}(u) \rangle$
rdivum $\langle \text{div}(u) \rangle$
divu2m $\langle \text{div}(u)^2 \rangle$
gdivu2m $\langle \text{grad} \text{div}(u)^2 \rangle$
u3u21m $\langle u_3 u_{2,1} \rangle$
u1u32m $\langle u_1 u_{3,2} \rangle$
K.3 Parameters for ‘print.in’

\begin{align*}
\mathbf{u}^{2} \mathbf{u}^{1} \mathbf{m} & (u_{2} u_{1,3}) \\
\mathbf{u}^{2} \mathbf{u}^{3} \mathbf{m} & (u_{2} u_{3,1}) \\
\mathbf{u}^{3} \mathbf{u}^{1} \mathbf{m} & (u_{3} u_{1,2}) \\
\mathbf{u}^{1} \mathbf{u}^{2} \mathbf{3} \mathbf{m} & (u_{1} u_{2,3}) \\
\mathbf{ruxm} & (\langle u \rangle) \quad \text{(mean } x \text{-momentum density)} \\
\mathbf{ruym} & (\langle u \rangle) \quad \text{(mean } y \text{-momentum density)} \\
\mathbf{ruzm} & (\langle u \rangle) \quad \text{(mean } z \text{-momentum density)} \\
\mathbf{ruxtot} & (\langle u \rangle) \quad \text{(mean absolute } x \text{-momentum density)} \\
\mathbf{rumax} & \max(\langle |u| \rangle) \quad \text{(maximum modulus of momentum)} \\
\mathbf{ruxym} & \langle u \rangle \quad \text{(mean Reynolds stress)} \\
\mathbf{ruzsm} & \langle u \rangle \quad \text{(mean Reynolds stress)} \\
\mathbf{ruyzm} & \langle u \rangle \quad \text{(mean Reynolds stress)} \\
\mathbf{divrhourms} & \langle \nabla \cdot (u) \rangle_{\text{rms}} \\
\mathbf{divrhoumax} & \langle \nabla \cdot (u) \rangle_{\text{max}} \\
\mathbf{rlx}m & \langle \rho u_{z} - z u_{y} \rangle \\
\mathbf{rly}m & \langle \rho u_{z} - x u_{z} \rangle \\
\mathbf{rlz}m & \langle \rho u_{y} - y u_{x} \rangle \\
\mathbf{rlx2m} & \langle (\rho u_{z} - z u_{y})^{2} \rangle \\
\mathbf{rly2m} & \langle (\rho u_{z} - x u_{z})^{2} \rangle \\
\mathbf{rlz2m} & \langle (\rho u_{y} - y u_{x})^{2} \rangle \\
\mathbf{tot.ang.mom} & \text{Total angular momentum in spherical coordinates about the} \\
& \text{axis.} \\
\mathbf{dtu} & \delta t / [c_{s} \delta x / \max |u|] \quad \text{(time step relative to advective time step; see §5.15)} \\
\mathbf{oum} & \langle \omega \cdot u \rangle \\
\mathbf{ou.int} & \int_{V} \omega \cdot u \, dV \\
\mathbf{fum} & \langle f \cdot u \rangle \\
\mathbf{odel2um} & \langle \nabla \nabla u \rangle \\
\mathbf{o2m} & \langle \omega^{2} \rangle \equiv \langle (\nabla \times u)^{2} \rangle \\
\mathbf{orms} & \langle \omega^{2} \rangle_{1/2} \\
\mathbf{omax} & \max(\langle |\omega| \rangle) \\
\mathbf{ox2m} & \langle \omega_{x}^{2} \rangle \\
\mathbf{oy2m} & \langle \omega_{y}^{2} \rangle \\
\mathbf{oz2m} & \langle \omega_{z}^{2} \rangle \\
\mathbf{oxuzxm} & \langle \omega_{x} u_{z, x} \rangle \\
\mathbf{oyuzym} & \langle \omega_{y} u_{z, y} \rangle \\
\mathbf{oxoym} & \langle \omega_{x} \omega_{y} \rangle \\
\mathbf{oxozm} & \langle \omega_{x} \omega_{z} \rangle \\
\mathbf{oyozm} & \langle \omega_{y} \omega_{z} \rangle \\
\mathbf{qfm} & \langle q \cdot f \rangle \\
\mathbf{q2m} & \langle q^{2} \rangle \\
\mathbf{qrms} & \langle q^{2} \rangle_{1/2} \\
\mathbf{qmax} & \max(\langle |q| \rangle) \\
\mathbf{qom} & \langle q \cdot \omega \rangle \\
\mathbf{quxom} & \langle q \cdot (u \times \omega) \rangle \\
\mathbf{oumphi} & \langle \omega \cdot u \rangle_{\varphi} \\
\mathbf{dudu} & \langle \frac{u^{2} u}{c_{s}^{2}} \rangle \\
\mathbf{Marms} & \langle u^{2} / c_{s}^{2} \rangle \quad \text{(rms Mach number)} \\
\mathbf{Mamax} & \max |u| / c_{s} \quad \text{(maximum Mach number)} \\
\mathbf{EEK} & \langle \rho u^{2} \rangle / 2
\end{align*}
### THE PENCIL CODE

**ekin** \( \langle \frac{1}{2} u^2 \rangle \)

**ekintot** \( \int_V \frac{1}{2} u^2 \, dV \)

**uxglnrym** \( \langle u_x \partial_y \ln \rho \rangle \)

**uyglnrmx** \( \langle u_y \partial_x \ln \rho \rangle \)

**uzdivum** \( \langle u_z \nabla \cdot u \rangle \)

**uxdrydivum** \( \langle u_x u_y \nabla \cdot u \rangle \)

**divuHrms** \( (\nabla_H \cdot u_H)^{\text{rms}} \)

**uxxrms** \( \bar{u}_{x,x}^{\text{rms}} \)

**uuyrms** \( \bar{u}_{y,y}^{\text{rms}} \)

**uxzrms** \( \bar{u}_{x,z}^{\text{rms}} \)

**uyzrms** \( \bar{u}_{y,z}^{\text{rms}} \)

**uzyrms** \( \bar{u}_{z,y}^{\text{rms}} \)

**udpxxm** components of symmetric tensor \( \langle u_i \partial_j p + u_j \partial_i p \rangle \)

---

**Module ‘lorenz_gauge.f90’**

**phim** \( \langle \phi \rangle \)

**phipt** \( \phi(x_1, y_1, z_1) \)

**phip2** \( \phi(x_2, y_2, z_2) \)

**phibzm** \( \langle \phi B_z \rangle \)

**phibzmz** \( \langle \phi B_z \rangle_{xy} \)

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**Module ‘lucky_droplet.f90’**

**tt1m** \( \langle T \rangle \)

**qq1m** \( \langle \ln T \rangle \)

**qq2m** \( \langle \ln T^2 \rangle \)

**qq3m** \( \langle \ln T^3 \rangle \)

**qq4m** \( \langle \ln T^4 \rangle \)

---

**Module ‘magnetic_shearboxJ.f90’**

**ab_int** \( \int \mathbf{A} \cdot \mathbf{B} \, dV \)

**jb_int** \( \int j \cdot \mathbf{B} \, dV \)

**b2tm** \( \langle b(t) \cdot \int_0^t b(t') dt' \rangle \)

**bjtm** \( \langle b(t) \cdot \int_0^t j(t') dt' \rangle \)

**jbtm** \( \langle j(t) \cdot \int_0^t b(t') dt' \rangle \)

**b2ruzm** \( \langle B^2_{u_z} \rangle \)

**b2uzm** \( \langle B^2_{u_z} \rangle \)

**ubbzm** \( \langle (\mathbf{u} \cdot \mathbf{B}) B_z \rangle \)

**b1m** \( \langle B \rangle \)

**b2m** \( \langle B^2 \rangle \)

**bm2** \( \text{max}(B^2) \)

**j2m** \( \langle j^2 \rangle \)

**jm2** \( \text{max}(j^2) \)

**abm** \( \langle \mathbf{A} \cdot \mathbf{B} \rangle \)

**abumx** \( \langle u_x \mathbf{A} \cdot \mathbf{B} \rangle \)

**abumy** \( \langle u_y \mathbf{A} \cdot \mathbf{B} \rangle \)

**abumz** \( \langle u_z \mathbf{A} \cdot \mathbf{B} \rangle \)

**abmh** \( \langle \mathbf{A} \cdot \mathbf{B} \rangle \text{ (temp)} \)

**abmn** \( \langle \mathbf{A} \cdot \mathbf{B} \rangle \text{ (north)} \)

**abms** \( \langle \mathbf{A} \cdot \mathbf{B} \rangle \text{ (south)} \)
\[ \text{Abrms} = \left( \langle A \cdot B \rangle^2 \right)^{1/2} \]
\[ \text{Jbrms} = \left( \langle j \cdot B \rangle^2 \right)^{1/2} \]
\[ \text{Ajm} = \langle j \cdot A \rangle \]
\[ \text{Jbm} = \langle j \cdot B \rangle \]
\[ \text{Jbmh} = \langle j \cdot B \rangle \text{ (temp)} \]
\[ \text{Jbmn} = \langle j \cdot B \rangle \text{ (north)} \]
\[ \text{Jbms} = \langle j \cdot B \rangle \text{ (south)} \]
\[ \text{Ubm} = \langle u \cdot B \rangle \]
\[ \text{Dubrms} = \left( \langle u - B \rangle^2 \right)^{1/2} \]
\[ \text{Dobrms} = \left( \langle \omega - B \rangle^2 \right)^{1/2} \]
\[ \text{Uxbxm} = \langle u_x B_x \rangle \]
\[ \text{Uybxm} = \langle u_y B_x \rangle \]
\[ \text{Uzbxm} = \langle u_z B_x \rangle \]
\[ \text{Uxbym} = \langle u_x B_y \rangle \]
\[ \text{Uybxm} = \langle u_y B_y \rangle \]
\[ \text{Uzbxm} = \langle u_z B_y \rangle \]
\[ \text{Uxbzm} = \langle u_x B_z \rangle \]
\[ \text{Uybxm} = \langle u_y B_z \rangle \]
\[ \text{Uzbxm} = \langle u_z B_z \rangle \]
\[ \text{Cosubm} = \langle U \cdot B \rangle \left/ \left( \langle |U| \right| B \rangle \right\rangle \]
\[ \text{Jxbxm} = \langle j_x B_x \rangle \]
\[ \text{Jybxm} = \langle j_y B_x \rangle \]
\[ \text{Jzbxm} = \langle j_z B_x \rangle \]
\[ \text{Jxbym} = \langle j_x B_y \rangle \]
\[ \text{Jybxm} = \langle j_y B_y \rangle \]
\[ \text{Jzbxm} = \langle j_z B_y \rangle \]
\[ \text{Jxbzm} = \langle j_x B_z \rangle \]
\[ \text{Jybxm} = \langle j_y B_z \rangle \]
\[ \text{Jzbxm} = \langle j_z B_z \rangle \]
\[ \text{Uam} = \langle u \cdot A \rangle \]
\[ \text{Ujm} = \langle u \cdot J \rangle \]
\[ \text{Fbm} = \langle f \cdot B \rangle \]
\[ \text{Fxbxm} = \langle f_x B_x \rangle \]
\[ \text{Epsm} = \left( \eta \mu_0 \right)^2 \]
\[ \text{Epsad} = \left( \rho^{-1} t_{AD} (J \times B)^2 \right) \text{ (heating by ion-neutrals friction)} \]
\[ \text{Bxpt} = B_x(x_1, y_1, z_1, t) \]
\[ \text{Bypt} = B_y(x_1, y_1, z_1, t) \]
\[ \text{Bzpt} = B_z(x_1, y_1, z_1, t) \]
\[ \text{Jxpt} = J_x(x_1, y_1, z_1, t) \]
\[ \text{Jypt} = J_y(x_1, y_1, z_1, t) \]
\[ \text{Jzpt} = J_z(x_1, y_1, z_1, t) \]
\[ \text{Expt} = E_x(x_1, y_1, z_1, t) \]
\[ \text{Eypt} = E_y(x_1, y_1, z_1, t) \]
\[ \text{Ezpt} = E_z(x_1, y_1, z_1, t) \]
\[ \text{Axpt} = A_x(x_1, y_1, z_1, t) \]
\[ \text{Aypt} = A_y(x_1, y_1, z_1, t) \]
\[ \text{Azpt} = A_z(x_1, y_1, z_1, t) \]
\[ \text{Bxpt} = B_x(x_2, y_2, z_2, t) \]
\[ \text{Bypt} = B_y(x_2, y_2, z_2, t) \]
\[ \text{Bzpt} = B_z(x_2, y_2, z_2, t) \]
\begin{align*}
\text{jxp2} & \quad J_x(x_2, y_2, z_2, t) \\
\text{jyp2} & \quad J_y(x_2, y_2, z_2, t) \\
\text{jzp2} & \quad J_z(x_2, y_2, z_2, t) \\
\text{Exp2} & \quad \mathcal{E}_x(x_2, y_2, z_2, t) \\
\text{Eyp2} & \quad \mathcal{E}_y(x_2, y_2, z_2, t) \\
\text{Ezp2} & \quad \mathcal{E}_z(x_2, y_2, z_2, t) \\
\text{axp2} & \quad A_x(x_2, y_2, z_2, t) \\
\text{ayp2} & \quad A_y(x_2, y_2, z_2, t) \\
\text{azp2} & \quad A_z(x_2, y_2, z_2, t) \\
\text{exabot} & \quad \int \mathbf{E} \times \mathbf{A} \, dS \big|_{\text{bot}} \\
\text{exatop} & \quad \int \mathbf{E} \times \mathbf{A} \, dS \big|_{\text{top}} \\
\text{emag} & \quad \int V 1/2 \mathbf{B}^2 \, dV \\
\text{brms} & \quad \langle \mathbf{B}^2 \rangle^{1/2} \\
\text{bfrms} & \quad \langle \mathbf{B}^2 \rangle^{1/2} \\
\text{bmax} & \quad \max(|\mathbf{B}|) \\
\text{bxmin} & \quad \min(|\mathbf{B}_x|) \\
\text{bymin} & \quad \min(|\mathbf{B}_y|) \\
\text{bzmin} & \quad \min(|\mathbf{B}_z|) \\
\text{bxmax} & \quad \max(|\mathbf{B}_x|) \\
\text{bymax} & \quad \max(|\mathbf{B}_y|) \\
\text{bzmax} & \quad \max(|\mathbf{B}_z|) \\
\text{bbxmax} & \quad \max(|\mathbf{B}_x|)_{\text{excludingBv}} \\
\text{bymax} & \quad \max(|\mathbf{B}_y|)_{\text{excludingBv}} \\
\text{bbzmax} & \quad \max(|\mathbf{B}_z|)_{\text{excludingBv}} \\
\text{jxmax} & \quad \max(|jv_x|) \\
\text{jymax} & \quad \max(|jv_y|) \\
\text{jzmax} & \quad \max(|jv_z|) \\
\text{jrms} & \quad \langle j^2 \rangle^{1/2} \\
\text{hjrms} & \quad \langle j^2 \rangle^{1/2} \\
\text{jmax} & \quad \max(|j|) \\
\text{vArms} & \quad \langle \mathbf{B}^2 / \rho \rangle^{1/2} \\
\text{vAmax} & \quad \max(\langle \mathbf{B}^2 / \rho \rangle^{1/2}) \\
\text{dtb} & \quad \delta t / [c_s, \delta x / v_{A,\text{max}}] \quad \text{(time step relative to Alfvén time step; see \S\ 5.15)} \\
\text{dteta} & \quad \delta t / [c_s, \delta x / \delta x / \eta_{\text{max}}] \quad \text{(time step relative to resistive time step; see \S\ 5.15)} \\
\text{a2m} & \quad \langle \mathbf{A}^2 \rangle \\
\text{arms} & \quad \langle \mathbf{A}^2 \rangle^{1/2} \\
\text{amax} & \quad \max(|\mathbf{A}|) \\
\text{divarms} & \quad \langle (\nabla \cdot \mathbf{A})^2 \rangle^{1/2} \\
\text{beta1m} & \quad \langle \mathbf{B}^2 / (2\mu_0 p) \rangle \quad \text{(mean inverse plasma beta)} \\
\text{beta1max} & \quad \max[\mathbf{B}^2 / (2\mu_0 p)] \quad \text{(maximum inverse plasma beta)} \\
\text{betam} & \quad \langle \beta \rangle \\
\text{betamax} & \quad \max \beta \\
\text{betamin} & \quad \min \beta \\
\text{bxm} & \quad \langle B_x \rangle \\
\text{bym} & \quad \langle B_y \rangle \\
\text{bzm} & \quad \langle B_z \rangle 
\end{align*}
K.3 Parameters for ‘print.in’

bxbym \( \langle B_x B_y \rangle \)

bmx \( \langle B_{y z}^2 \rangle^{1/2} \) (energy of \( yz \)-averaged mean field)

bmy \( \langle B_{x z}^2 \rangle^{1/2} \) (energy of \( xz \)-averaged mean field)

bmz \( \langle B_{x y}^2 \rangle^{1/2} \) (energy of \( xy \)-averaged mean field)

bmzS2 \( \langle B_{S y x}^2 \rangle \)

bmzA2 \( \langle B_{A y x}^2 \rangle \)

jmx \( \langle J_{y z}^2 \rangle^{1/2} \) (energy of \( yz \)-averaged mean current density)

jmy \( \langle J_{x z}^2 \rangle^{1/2} \) (energy of \( xz \)-averaged mean current density)

jmxz \( \langle J_{x y}^2 \rangle^{1/2} \) (energy of \( xy \)-averaged mean current density)

bmzph Phase of a Beltrami field

bmzphe Error of phase of a Beltrami field

bsinphz sine of phase of a Beltrami field

bcosphz cosine of phase of a Beltrami field

emxamz3 \( \langle E \rangle_{xy} \times \langle A \rangle_{xy} \) (xy-averaged mean field helicity flux)

embmz \( \langle E \rangle_{xy} \cdot \langle B \rangle_{xy} \) (xy-averaged mean field helicity production)

ambmz \( \langle A \rangle_{xy} \cdot \langle B \rangle_{xy} \) (magnetic helicity of \( xy \)-averaged mean field)

ambmzh \( \langle A \rangle_{xy} \cdot \langle B \rangle_{xy} \) (magnetic helicity of \( xy \)-averaged mean field, temp)

ambmzn \( \langle A \rangle_{xy} \cdot \langle B \rangle_{xy} \) (magnetic helicity of \( xy \)-averaged mean field, north)

ambmzs \( \langle A \rangle_{xy} \cdot \langle B \rangle_{xy} \) (magnetic helicity of \( xy \)-averaged mean field, south)

jmbmz \( \langle J \rangle_{xy} \cdot \langle B \rangle_{xy} \) (current helicity of \( xy \)-averaged mean field)

kx\_aa

kmz \( \langle J \rangle_{xy} \cdot \langle B \rangle_{xy} / \langle B^2 \rangle_{xy} \)

bx2m \( \langle B_x^2 \rangle \)

by2m \( \langle B_y^2 \rangle \)

bz2m \( \langle B_z^2 \rangle \)

uxbm \( \langle u \times B \rangle \cdot B_0 / B_0^2 \)

jxbm \( \langle j \times B \rangle \cdot B_0 / B_0^2 \)

magfricmax Magneto-Frictional velocity \( \langle j \times B \rangle \cdot B^2 \)

b3b21m \( \langle B_3 B_{2,1} \rangle \)

b3b12m \( \langle B_3 B_{1,2} \rangle \)

b1b32m \( \langle B_1 B_{3,2} \rangle \)

b1b23m \( \langle B_1 B_{2,3} \rangle \)

b2b13m \( \langle B_2 B_{1,3} \rangle \)

b2b31m \( \langle B_2 B_{3,1} \rangle \)

uxbmx \( \langle (u \times B) \rangle_x \)

uxbmy \( \langle (u \times B) \rangle_y \)

uxbmez \( \langle (u \times B) \rangle_z \)
\begin{align*}
\text{jxbmx} & = \langle (j \times B)_x \rangle \\
\text{jxbmy} & = \langle (j \times B)_y \rangle \\
\text{jxbmz} & = \langle (j \times B)_z \rangle \\
\text{examx} & = \langle (E \times A) \rangle_x \\
\text{examy} & = \langle (E \times A) \rangle_y \\
\text{examz} & = \langle (E \times A) \rangle_z \\
\text{exjmx} & = \langle (E \times J) \rangle_x \\
\text{exjmy} & = \langle (E \times J) \rangle_y \\
\text{exjmz} & = \langle (E \times J) \rangle_z \\
\text{dexbmx} & = \langle \nabla \times E \times B \rangle_x \\
\text{dexbmy} & = \langle \nabla \times E \times B \rangle_y \\
\text{dexbmz} & = \langle \nabla \times E \times B \rangle_z \\
\text{phibmx} & = \langle \phi B \rangle_x \\
\text{phibmy} & = \langle \phi B \rangle_y \\
\text{phibmz} & = \langle \phi B \rangle_z \\
\text{b2divum} & = \langle B^2 \nabla \cdot u \rangle \\
\text{ujxbm} & = \langle u \cdot (J \times B) \rangle \\
\text{jxbmax} & = \max \|(J \times B) / \rho\| \\
\text{jxbr2m} & = \langle (J \times B) / \rho\rangle^2 \\
\text{bmx_rms} & = \sqrt{\|b_x\|^2 + \|b_y\|^2 + \|b_z\|^2} \\
\text{etasmagm} & = \text{Mean of Smagorinsky resistivity} \\
\text{etasmagmin} & = \text{Min of Smagorinsky resistivity} \\
\text{etasmagmax} & = \text{Max of Smagorinsky resistivity} \\
\text{etavamax} & = \text{Max of artificial resistivity } \eta \sim v \rho \\
\text{etajmax} & = \text{Max of artificial resistivity } \eta \sim J / \sqrt{\rho} \\
\text{etaj2max} & = \text{Max of artificial resistivity } \eta \sim J^2 / \rho \\
\text{etajrhomax} & = \text{Max of artificial resistivity } \eta \sim J / \rho \\
\text{cosjbm} & = \langle J \cdot B / (|J| |B|) \rangle \\
\text{jparallelm} & = \text{Mean value of the component of } J \text{ parallel to } B \\
\text{jperpm} & = \text{Mean value of the component of } J \text{ perpendicular to } B \\
\text{hiparallelm} & = \text{Mean value of the component of } J_{\text{hyper parallel to } B} \\
\text{hiperpm} & = \text{Mean value of the component of } J_{\text{hyper perpendicular to } B} \\
\text{brmsx} & = \langle B^2 \rangle_1^{1/2} \text{ for the magnetic_xaver_range} \\
\text{brmsz} & = \langle B^2 \rangle_1^{1/2} \text{ for the magnetic_zaver_range} \\
\text{Exmxy} & = \langle \mathcal{E}_x \rangle_z \\
\text{Eymxy} & = \langle \mathcal{E}_y \rangle_z \\
\text{Ezmxy} & = \langle \mathcal{E}_z \rangle_z \\
\text{Module 'meanfield.f90'}
\end{align*}
### Module ‘meanfield_demfdt.f90’

- **EMFrms**: \( \langle \langle E \rangle \rangle_{\text{rms}} \)
- **EMFmax**: \( \text{max}(\langle E \rangle) \)
- **EMFmin**: \( \text{min}(\langle E \rangle) \)

### Module ‘neutralvelocity.f90’

- **epsKn**: \( \langle 2\nu_n\theta_nS_n^2 \rangle \)

### Module ‘noentropy.f90’

- **dtc**: \( \delta t/[c_{\delta t} \delta \gamma / \text{max} c_s] \) (time step relative to acoustic time step; see §5.15)
- **ethm**: \( \langle \rho e \rangle \) (mean thermal [=internal] energy)
- **pdivism**: \( \langle p\nabla u \rangle \)

### Module ‘particles_caustics.f90’

- **TrSigmapm**: \( \langle \text{Tr}[\sigma] \rangle \)
- **blowupm**: Mean no. of times \( \sigma \) falls below cutoff
- **lnVpm**: Mean of (logarithm of) Volume around an inertial particle

### Module ‘particles_chemistry.f90’

- **Shchm**: mean particle Sherwood number

### Module ‘particles_dust.f90’

- **xpm**: \( x_{\text{part}} \)
- **xpmin**: \( x_{\text{part}} \)
- **xmax**: \( x_{\text{part}} \)
- **xp2m**: \( x_{\text{part}}^2 \)
- **vrelpabsm**: Absolute value of mean relative velocity
- **vpxm**: \( u_{\text{part}} \)
- **vp2xm**: \( u_{\text{part}}^2 \)
- **ekinp**: \( E_{\text{kin,part}} \)
- **vpxmax**: \( \text{MAX}(u_{\text{part}}) \)
- **vpxmin**: \( \text{MIN}(u_{\text{part}}) \)
- **npm**: mean particle number density

### Module ‘particles_dust_brdeplete.f90’

- **xpm**: \( x_{\text{part}} \)
- **xp2m**: \( x_{\text{part}}^2 \)
- **vrelpabsm**: Absolute value of mean relative velocity
- **vpxm**: \( u_{\text{part}} \)
- **vp2xm**: \( u_{\text{part}}^2 \)
- **ekinp**: \( E_{\text{kin,part}} \)
- **vpxmax**: \( \text{MAX}(u_{\text{part}}) \)
- **vpxmin**: \( \text{MIN}(u_{\text{part}}) \)
- **npm**: mean particle number density

### Module ‘particles_lagrangian.f90’

- **xpm**: \( x_{\text{part}} \)
- **xp2m**: \( x_{\text{part}}^2 \)
- **vrelpabsm**: Absolute value of mean relative velocity
vpxm \quad u_{part} \\
vpx2m \quad u_{part} \\
ekin \quad E_{kin,part} \\
vpxmax \quad MAX(u_{part}) \\
vpxmin \quad MIN(u_{part}) \\
npm \quad \text{mean particle number density} \\

Module ‘particles_mass_swarm.f90’

mpm \quad \bar{m}_p \\
mpmin \quad \min_j m_{p,j} \\
mpmax \quad \max_j m_{p,j} \\

Module ‘particles_surfspec.f90’

dtpchem \quad dt_{\text{particle,chemistry}}

Module ‘particles_tetrad.f90’

TVolm \quad \text{Mean absolute volume of the tetrads} \\
TVolpm \quad \text{Mean of positive volume of the tetrads} \\
TVolnm \quad \text{Mean of negative volume of the tetrads} \\
VelVolm \quad \text{Mean absolute volume of the tetrads in velocity space} \\
VelVolpm \quad \text{Mean of positive volume of the tetrads in velocity space} \\
VelVolnm \quad \text{Mean of negative volume of the tetrads in velocity space.} \\

Module ‘polymer.f90’

polytrm \quad \langle Tr[C_{ij}] \rangle \\
frmax \quad \max(f(r))

Module ‘reaction_0D.f90’

eem \quad \langle \eta \rangle \\
ee1m \quad \langle |\eta| \rangle \\
ee2m \quad \langle \eta^2 \rangle \\
ee3m \quad \langle \eta^3 \rangle \\
ee4m \quad \langle \eta^4 \rangle \\
ee10 \quad \langle \eta_{10\%} \rangle \\
ee50 \quad \langle \eta_{50\%} \rangle \\
ee90 \quad \langle \eta_{90\%} \rangle \\
ee99 \quad \langle \eta_{99\%} \rangle \\
AAm \quad \langle [A] \rangle \\
DDm \quad \langle [D] \rangle \\
LLm \quad \langle [L] \rangle \\
DLm \quad \langle [D] + [L] \rangle \\
kC \quad k_C \\
A1 \quad A_1 \\
A2 \quad A_2 \\
A3 \quad A_3 \\
A4 \quad A_4 \\
A5 \quad A_5 \\
D1 \quad D_1 \\
D2 \quad D_2 \\
D3 \quad D_3 \\
D4 \quad D_4
K.3 Parameters for ‘print.in’

<table>
<thead>
<tr>
<th>Module ‘shear.f90’</th>
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<tbody>
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<td><code>deltay</code></td>
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<th>Module ‘temperature_idealgas.f90’</th>
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<tr>
<td>$\langle u_y T \rangle$</td>
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<td>$\langle u \cdot \nabla (u_x T) \rangle$</td>
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<td>$\langle \nabla u_y \cdot \nabla u_y + u_y u_z \cdot \nabla T \rangle$</td>
<td>$\langle u \cdot \nabla (u_y T) \rangle$</td>
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<tr>
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<td>$\langle u \cdot \nabla (u_z T) \rangle$</td>
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</tbody>
</table>
\(T_{dxpm}\) \(\langle T dp/dx \rangle\)
\(T_{dypm}\) \(\langle T dp/dy \rangle\)
\(T_{dzpm}\) \(\langle T dp/dz \rangle\)

\(frad\)top \(< -K \frac{dT}{dy}\)top \(\) (top radiative flux)
\(frad\)bot \(< -K \frac{dT}{dz}\)bot \(\) (bottom radiative flux)

\(y \)H\(m\)ax \(\) DOCUMENT ME
\(y \)H\(m\)in \(\) DOCUMENT ME

\(ethm\) \(\langle e_{th} \rangle = \langle c_v \rho T \rangle\) \(\) (mean thermal energy)
\(eem\) \(\langle e \rangle = \langle c_v T \rangle\) \(\) (mean internal energy)

\(ethtot\) \(\int_V \langle e \rangle dV\) \(\) (total thermal energy)

\(ssm\) \(\) DOCUMENT ME

\(th\)cool \(\tau_{cool}\)
\(ppm\) \(\overline{P}\)
\(cs\)m \(\overline{c}_s\)
\(cs\)max \(\max(c_s)\)

\(dt\)c \(\delta t/[c_{st} \delta x/ \max c_s]\) \(\) (time step relative to acoustic time step; see \(\S\) 5.15)

\(dt\)chi \(\delta t/[c_{st,v} \delta x^2/\chi_{\text{max}}]\) \(\) (time step relative to time step based on heat conductivity; see \(\S\) 5.15)

---

Module ‘temperature_ionization.f90’

\(TT\)max \(\max(T)\)
\(TT\)min \(\min(T)\)
\(TT\)m \(\langle T \rangle\)

\(ethm\) \(\langle e_{th} \rangle = \langle c_v \rho T \rangle\) \(\) (mean thermal energy)
\(eem\) \(\langle e \rangle\) \(\) (mean internal energy)

\(ppm\) \(\langle p \rangle\)

\(Tppm\) \(\langle \max(p_{\text{thresh}} - p, 0)_{\text{norm}} \rangle\)

---

Module ‘test_chemistry.f90’

\(dt\)chem \(dt_{\text{chem}}\)

---

Module ‘testfield_axisym.f90’

\(alp\)PERP \(\alpha_\perp\)
\(alp\)PARA \(\alpha_\parallel\)
\(gam\) \(\gamma\)
\(bet\)PERP \(\beta_\perp\)
\(bet\)PARA \(\beta_\parallel\)
\(del\) \(\delta\)

\(kap\)PERP \(\kappa_\perp\)
\(kap\)PARA \(\kappa_\parallel\)
\(mu\) \(\mu\)

\(alp\)PERPz \(\alpha_\perp(z)\)
\(alp\)PARAz \(\alpha_\parallel(z)\)
\(gamz\) \(\gamma(z)\)
\(bet\)PERPz \(\beta_\perp(z)\)
\(bet\)PARAz \(\beta_\parallel(z)\)
\(delz\) \(\delta(z)\)
\(kap\)PERPz \(\kappa_\perp(z)\)
\(kap\)PARAz \(\kappa_\parallel(z)\)
### Parameters for 'print.in'

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<td>gam</td>
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\begin{align*}
  bx2pt &= b_x^2 \\
  bx3pt &= b_x^3 \\
  b1rms &= \langle b_1^2 \rangle^{1/2} \\
  b2rms &= \langle b_2^2 \rangle^{1/2} \\
  b3rms &= \langle b_3^2 \rangle^{1/2} \end{align*}

Module ‘testfield_compress_z.f90’

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K.3 Parameters for 'print.in'

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- **M22cc**: $\mathcal{M}_{22} \cos^2 k z$
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- **M12cs**: $\mathcal{M}_{12} \cos k z \sin k z$

- $b_{x11pt}$: $b_{x11}^1$
- $b_{x21pt}$: $b_{x21}^1$
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- $b_{y11pt}$: $b_{y11}^1$
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- $Ex_{11pt}$: $E_x^{11}$
- $Ex_{21pt}$: $E_x^{21}$
- $Ex_{12pt}$: $E_x^{12}$
- $Ex_{22pt}$: $E_x^{22}$
\( \mathcal{E} \cdot B^{pq} \)
\( \mathcal{E} \cdot U \)
\( \mathcal{E} \cdot W \)
\( \langle b_x \rangle_{xy} \)
\( \langle b_y \rangle_{xy} \)
\( \langle b_z \rangle_{xy} \)
\( \langle M_{11} \rangle_{xy} \)
\( \langle M_{22} \rangle_{xy} \)
\( \langle M_{33} \rangle_{xy} \)

Module ‘testfield_meri.f90’

\( E_{11xy} \)
\( E_{12xy} \)
\( E_{13xy} \)
\( E_{21xy} \)
\( E_{22xy} \)
\( E_{23xy} \)
\( E_{31xy} \)
\( E_{32xy} \)
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Module 'testfield_nonlin_z.f90'

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\text{alpMK} & \quad \alpha^{MK} \\
\text{phi11} & \quad \phi_{11} \\
\text{phi21} & \quad \phi_{21} \\
\text{phi12} & \quad \phi_{12} \\
\text{phi22} & \quad \phi_{22} \\
\text{phi32} & \quad \phi_{32} \\
\text{psi11} & \quad \psi_{11} k \\
\text{psi21} & \quad \psi_{21} k \\
\text{psi12} & \quad \psi_{12} k \\
\text{psi22} & \quad \psi_{22} k \\
\text{phiK} & \quad \phi^K \\
\phiM & \quad \phi^M \\
\phiMK & \quad \phi^{MK} \\
\text{alp11cc} & \quad \alpha_{11} \cos^2 k z \\
\text{alp21sc} & \quad \alpha_{21} \sin k z \cos k z \\
\text{alp12cs} & \quad \alpha_{12} \cos k z \sin k z \\
\text{alp22ss} & \quad \alpha_{22} \sin^2 k z \\
\text{eta11cc} & \quad \eta_{11} \cos^2 k z \\
\text{eta21sc} & \quad \eta_{21} \sin k z \cos k z \\
\text{eta12cs} & \quad \eta_{12} \cos k z \sin k z \\
\text{eta22ss} & \quad \eta_{22} \sin^2 k z \\
\text{s2kzDFm} & \quad \langle \sin 2 k z \nabla \cdot F \rangle \\
\text{M11} & \quad M_{11} \\
\text{M22} & \quad M_{22} \\
\text{M33} & \quad M_{33} \\
\text{M11cc} & \quad M_{11} \cos^2 k z \\
\text{M11ss} & \quad M_{11} \sin^2 k z \\
\text{M22cc} & \quad M_{22} \cos^2 k z \\
\text{M22ss} & \quad M_{22} \sin^2 k z \\
\text{M12cs} & \quad M_{12} \cos k z \sin k z \\
\text{bx11pt} & \quad b^1_x \\
\text{bx21pt} & \quad b^1_x \\
\text{bx12pt} & \quad b^2_x \\
\text{bx22pt} & \quad b^2_x \\
\text{bx0pt} & \quad b^0_x \\
\text{by11pt} & \quad b^1_y \\
\text{by21pt} & \quad b^1_y \\
\text{by12pt} & \quad b^2_y \\
\text{by22pt} & \quad b^2_y \\
\text{by0pt} & \quad b^0_y \\
\text{u11rms} & \quad \langle u_{11}^2 \rangle^{1/2} \\
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\text{u22rms} & \quad \langle u_{22}^2 \rangle^{1/2} \\
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\text{b12rms} & \quad \langle b_{12}^2 \rangle^{1/2}
\end{align*}
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K.3 Parameters for ‘print.in’ & 225 \\
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b22rms & $\langle b_{22}^2 \rangle^{1/2}$ 
ux0m & $\langle u_{0x} \rangle$ 
uy0m & $\langle u_{0y} \rangle$ 
ux11m & $\langle u_{11x} \rangle$ 
uy11m & $\langle u_{11y} \rangle$ 
u0rms & $\langle u_0^2 \rangle^{1/2}$ 
b0rms & $\langle b_0^2 \rangle^{1/2}$ 
u0max & $\max |u_0|$ 
b0max & $\max |b_0|$ 
jb0m & $\langle jb_0 \rangle$
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E11rms & $\langle \mathcal{E}_{11}^2 \rangle^{1/2}$ 
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Ex22pt & $\mathcal{E}_{x22}$ 
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Ey21pt & $\mathcal{E}_{y21}$ 
Ey12pt & $\mathcal{E}_{y12}$ 
Ey22pt & $\mathcal{E}_{y22}$ 
Ey0pt & $\mathcal{E}_y$
bamp & bamp
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E111z & $\mathcal{E}_{111}$ 
E211z & $\mathcal{E}_{211}$ 
E311z & $\mathcal{E}_{311}$ 
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E322z & $\mathcal{E}_{322}$ 
E10z & $\mathcal{E}_{10}$ 
E20z & $\mathcal{E}_{20}$ 
E30z & $\mathcal{E}_{30}$ 
EBpq & $\mathcal{E} \cdot B_{eq}$ 
E0Um & $\mathcal{E} \cdot U$
E0Wm & $\mathcal{E} \cdot W$
bx0mz & $\langle b_{x} \rangle_{xy}$ 
by0mz & $\langle b_{y} \rangle_{xy}$ 
bz0mz & $\langle b_{z} \rangle_{xy}$ 
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\[ E221z \quad \mathcal{E}^{21}_2 \]
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\[ E322z \quad \mathcal{E}^{22}_3 \]
\[ E10z \quad \mathcal{E}^0_1 \]
\[ E20z \quad \mathcal{E}^0_2 \]
\[ E30z \quad \mathcal{E}^0_3 \]
\[ EBpq \quad \mathcal{E} \cdot B^{eq} \]
\[ bx0mz \quad \langle b_x \rangle_{xy} \]
\[ by0mz \quad \langle b_y \rangle_{xy} \]
\[ bz0mz \quad \langle b_z \rangle_{xy} \]
\[ alp11 \quad \alpha_{11}(x, t) \]
\[ alp21 \quad \alpha_{21}(x, t) \]
\[ alp12 \quad \alpha_{12}(x, t) \]
\[ alp22 \quad \alpha_{22}(x, t) \]
\[ eta11 \quad \eta_{11}(x, t) \]
\[ eta21 \quad \eta_{21}(x, t) \]
\[ eta12 \quad \eta_{12}(x, t) \]
\[ eta22 \quad \eta_{22}(x, t) \]

---

Module ‘testfield_xz.f90’

\[ E111z \quad \mathcal{E}_1^{11} \]
\[ E211z \quad \mathcal{E}_2^{11} \]
\[ E311z \quad \mathcal{E}_3^{11} \]
\[ E121z \quad \mathcal{E}_1^{21} \]
\[ E221z \quad \mathcal{E}_2^{21} \]
\[ E321z \quad \mathcal{E}_3^{21} \]
\[ alp1 \quad \alpha_{11} \]
\[ alp2 \quad \alpha_{21} \]
\[ eta1 \quad \eta_{11} \]
\[ eta2 \quad \eta_{21} \]
\[ b11rms \quad \langle b_{11} \rangle \]
\[ b21rms \quad \langle b_{21} \rangle \]

---

Module ‘testfield_z.f90’

\[ alp1 \quad \alpha_{11} \]
\[ alp2 \quad \alpha_{21} \]
\[ alp3 \quad \alpha_{31} \]
\[ alp12 \quad \alpha_{12} \]
\[ alp22 \quad \alpha_{22} \]
\[ alp32 \quad \alpha_{32} \]
\[ alp13 \quad \alpha_{13} \]
\[ alp23 \quad \alpha_{23} \]
\[ eta1 \quad \eta_{11} \]

\eta_{11} or \eta_{11} \text{ if leta_rank2=T}
eta21 \quad \eta_{213k} \text{ or } \eta_{21k} \text{ if } \text{leta}_\text{rank2}=T
eta31 \quad \eta_{313k}
eta12 \quad \eta_{123k} \text{ or } \eta_{12k} \text{ if } \text{leta}_\text{rank2}=T
eta22 \quad \eta_{223k} \text{ or } \eta_{22k} \text{ if } \text{leta}_\text{rank2}=T
eta32 \quad \eta_{323k}
\text{alp11cc} \quad \alpha_{11} \cos^2 k z
\text{alp21sc} \quad \alpha_{21} \sin k z \cos k z
\text{alp12cs} \quad \alpha_{12} \cos k z \sin k z
\text{alp22ss} \quad \alpha_{22} \sin^2 k z
\text{eta11cc} \quad \eta_{11} \cos^2 k z
\text{eta21sc} \quad \eta_{21} \sin k z \cos k z
\text{eta12cs} \quad \eta_{12} \cos k z \sin k z
\text{eta22ss} \quad \eta_{22} \sin^2 k z
s2kzDFm \quad \langle \sin 2k z \nabla \cdot F \rangle
M11 \quad M_{11}
M22 \quad M_{22}
M33 \quad M_{33}
M11cc \quad M_{11} \cos^2 k z
M11ss \quad M_{11} \sin^2 k z
M22cc \quad M_{22} \cos^2 k z
M22ss \quad M_{22} \sin^2 k z
M12cs \quad M_{12} \cos k z \sin k z
b_{11} \quad b_{11}^x
b_{21} \quad b_{21}^x
b_{12} \quad b_{12}^x
b_{22} \quad b_{22}^x
b_0 \quad b_0^x
b_{11}^y \quad b_{11}^y
b_{21}^y \quad b_{21}^y
b_{12}^y \quad b_{12}^y
b_{22}^y \quad b_{22}^y
b_{0}^y \quad b_0^y
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\langle b_{21}^2 \rangle^{1/2}
\langle b_{12}^2 \rangle^{1/2}
\langle b_{22}^2 \rangle^{1/2}
\langle b_0^2 \rangle^{1/2}
\langle j b_0 \rangle
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E21rms \quad \langle \mathcal{E}_{21}^2 \rangle^{1/2}
E12rms \quad \langle \mathcal{E}_{12}^2 \rangle^{1/2}
E22rms \quad \langle \mathcal{E}_{22}^2 \rangle^{1/2}
E0rms \quad \langle \mathcal{E}_0^2 \rangle^{1/2}
Ex11pt \quad \mathcal{E}_{x}^{11}
Ex21pt \quad \mathcal{E}_{x}^{21}
Ex12pt \quad \mathcal{E}_{x}^{12}
Ex22pt \quad \mathcal{E}_{x}^{22}
Ex0pt \quad \mathcal{E}_{x}^{0}
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\(E_{y12pt}\)  \(E_{y}^{12}\)
\(E_{y22pt}\)  \(E_{y}^{22}\)
\(E_{y0pt}\)  \(E_{y}^{0}\)
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\(alp_{21z}\)  \(\alpha_{21}(z, t)\)
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\(alp_{23z}\)  \(\alpha_{23}(z, t)\)
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\(eta_{21z}\)  \(\eta_{21}(z, t)\)
\(eta_{12z}\)  \(\eta_{22}(z, t)\)
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\(uzjy_{1z}\)  \(u_{z}j_{y}^{11}\)
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\(uzjy_{3z}\)  \(u_{z}j_{y}^{12}\)
\(uzjz_{3z}\)  \(u_{z}j_{z}^{12}\)
\(uzjx_{4z}\)  \(u_{z}j_{x}^{22}\)
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\(E_{10z}\)  \(E_{0}^{11}\)
\(E_{20z}\)  \(E_{2}^{0}\)
\(E_{30z}\)  \(E_{3}^{0}\)
\(EBpq\)  \(\mathbf{E} \cdot \mathbf{B}^{eq}\)
\(E0Um\)  \(E_{0}^{0} \cdot \mathbf{U}\)
\(E0Wm\)  \(E_{0}^{0} \cdot \mathbf{W}\)
\(bx0mz\)  \(\langle b_{x} \rangle_{xy}\)
\(by0mz\)  \(\langle b_{y} \rangle_{xy}\)
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<td>$\nu$-tensor, couples $\overline{\mathcal{F}}$ and $\partial^2 \overline{\mathcal{U}} / \partial z^2$</td>
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**Module ‘testflow_z.f90’**

- $\langle M_{33z} \rangle_{xy}$
- $\langle u_x \rangle_{xy}$
- $\langle u_y \rangle_{xy}$
- $\langle u_z \rangle_{xy}$

**Module ‘testperturb.f90’**

- $\alpha_{11}$
- $\alpha_{21}$
- $\alpha_{31}$
- $\alpha_{12}$
- $\alpha_{22}$
- $\alpha_{32}$
- $\eta_{113k}$
- $\eta_{213k}$
- $\eta_{313k}$
- $\eta_{123k}$
- $\eta_{223k}$
- $\eta_{323k}$

**Module ‘testscalar.f90’**

- $\gamma_1^{(1)}$
- $\gamma_2^{(1)}$
- $\gamma_3^{(1)}$
- $\gamma_1^{(2)}$
- $\gamma_2^{(2)}$
- $\gamma_3^{(2)}$
- $\gamma_1^{(3)}$
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### Module ‘testscalar_axisym.f90’

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K.3 Parameters for ‘print.in’

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Module ‘testscalar_simple.f90’

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\[ \gamma_{12}^{(3)}(z,t) \]
\[
\gamma^{(3)}_3(z, t), \quad \kappa_{11}(z, t), \quad \kappa_{21}(z, t), \quad \kappa_{31}(z, t), \quad \kappa_{12}(z, t), \quad \kappa_{22}(z, t), \quad \kappa_{32}(z, t), \quad \kappa_{13}(z, t), \quad \kappa_{23}(z, t), \quad \kappa_{33}(z, t),
\]
\[
\tilde{\gamma}_{33}, \quad \tilde{\kappa}_{33}, \quad \hat{\gamma}_{33}, \quad \hat{\kappa}_{33},
\]
\[
c_{1\text{rms}} \langle c_1^2 \rangle^{1/2}, \quad c_{2\text{rms}} \langle c_2^2 \rangle^{1/2}, \quad c_{3\text{rms}} \langle c_3^2 \rangle^{1/2}, \quad c_{4\text{rms}} \langle c_4^2 \rangle^{1/2}, \quad c_{5\text{rms}} \langle c_5^2 \rangle^{1/2}, \quad c_{6\text{rms}} \langle c_6^2 \rangle^{1/2},
\]
\[
c_1 p_t, \quad c_2 p_t, \quad c_3 p_t, \quad c_4 p_t, \quad c_5 p_t, \quad c_6 p_t,
\]
\[
F_{11z} \mathcal{F}_{11}, \quad F_{21z} \mathcal{F}_{12}, \quad F_{31z} \mathcal{F}_{13}, \quad F_{12z} \mathcal{F}_{21}, \quad F_{22z} \mathcal{F}_{22}, \quad F_{32z} \mathcal{F}_{32},
\]
\[
\text{Module 'thermal_energy.f90'}
\]
\[
TT_{\text{max}} = \max(T), \quad TT_{\text{min}} = \min(T),
\]
\[
ppm = \langle p \rangle, \quad TTm = \langle T \rangle,
\]
\[
ethm = \langle e_{th} \rangle = \langle c_v \rho T \rangle \quad \text{(mean thermal energy)}, \quad ethtot = \int_V e_{th} \, dV \quad \text{(total thermal energy)},
\]
\[
ethm = \min e_{th}, \quad ethmax = \max e_{th}, \quad eem = \langle e \rangle = \langle c_v T \rangle \quad \text{(mean internal energy)}, \quad etot = \langle e_{th} + \rho u^2 / 2 \rangle
\]

\[
\text{Module 'visc_smagorinsky.f90'}
\]
\[
uu_{\text{LES}} \quad \text{Mean value of Smagorinsky viscosity}
\]

\[
\text{Module 'viscosity.f90'}
\]
\[
uu_{t\text{dep}} \quad \text{time-dependent viscosity}
\]
The following table lists all (at the time of writing, September 8, 2020) possible inputs to the file ‘video.in’.

<table>
<thead>
<tr>
<th>Variable</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Module ‘hydro.f90’</strong></td>
<td></td>
</tr>
<tr>
<td>uu</td>
<td>velocity vector (u); writes all three components separately to files 'u[xyz].{xz,yz,xy,xy2}'</td>
</tr>
<tr>
<td>u2</td>
<td>kinetic energy density (u^2); writes 'u2.{xz,yz,xy,xy2}'</td>
</tr>
<tr>
<td>oo</td>
<td>vorticity vector (\omega = \nabla \times u); writes all three components separately to files 'oo[xyz].{xz,yz,xy,xy2}'</td>
</tr>
<tr>
<td>o2</td>
<td>enstrophy (\omega^2 =</td>
</tr>
<tr>
<td>divu</td>
<td>(\nabla \cdot u); writes 'divu.xz,yz,xy,xy2'</td>
</tr>
<tr>
<td>mach</td>
<td>Mach number squared (Ma^2); writes 'mach.xz,yz,xy,xy2'</td>
</tr>
<tr>
<td><strong>Module ‘density.f90’</strong></td>
<td></td>
</tr>
<tr>
<td>lnrho</td>
<td>logarithmic density (\ln \rho); writes 'lnrho.xz,yz,xy,xy2'</td>
</tr>
<tr>
<td>rho</td>
<td>density (\rho); writes 'rho.xz,yz,xy,xy2'</td>
</tr>
<tr>
<td><strong>Module ‘entropy.f90’</strong></td>
<td></td>
</tr>
<tr>
<td>ss</td>
<td>entropy (s); writes 'ss.xz,yz,xy,xy2'</td>
</tr>
<tr>
<td>pp</td>
<td>pressure (p); writes 'pp.xz,yz,xy,xy2'</td>
</tr>
<tr>
<td><strong>Module ‘temperature_idealgas.f90’</strong></td>
<td></td>
</tr>
<tr>
<td>lnTT</td>
<td>logarithmic temperature (\ln T); writes 'lnTT.xz,yz,xy,xy2'</td>
</tr>
<tr>
<td>TT</td>
<td>temperature (T); writes 'TT.xz,yz,xy,xy2'</td>
</tr>
<tr>
<td><strong>Module ‘shock.f90’</strong></td>
<td></td>
</tr>
</tbody>
</table>
shock shock viscosity $\nu_{\text{shock}}$; writes 'shock.{xz, yz, xy, xy2}'

Module 'eos_ionization.f90'

yH ionization fraction $y_H$; writes 'yH.{xz, yz, xy, xy2}'

Module 'radiation_ray.f90'

$Q_{\text{rad}}$ radiative heating rate $Q_{\text{rad}}$; writes 'Qrad.{xz, yz, xy, xy2}'

Isurf surface intensity $I_{\text{surf}}$ (?); writes 'Isurf.xz'

Module 'magnetic.f90'

$aa$ magnetic vector potential $A$; writes 'aa[xyz].{xz, yz, xy, xy2}'

$bb$ magnetic flux density $B$; writes 'bb[xyz].{xz, yz, xy, xy2}'

$b2$ magnetic energy density $B^2$; writes 'b2.{xz, yz, xy, xy2}'

$jj$ current density $j$; writes 'jj[xyz].{xz, yz, xy, xy2}'

$j2$ current density squared $j^2$; writes 'j2.{xz, yz, xy, xy2}'

$jb$ $jB$; writes 'jb.{xz, yz, xy, xy2}'

$\beta_1$ inverse plasma beta $B^2/(2\mu_0\rho)$; writes 'beta1.{xz, yz, xy, xy2}'

Poynting Poynting vector $\eta j \times B - (u \times B) \times B/\mu_0$; writes 'Poynting[xyz].{xz, yz, xy, xy2}'

$ab$ magnetic helicity density $A \cdot B$; writes 'ab[xyz].{xz, yz, xy, xy2}'

Module 'pscalar.f90'

$\lncc$ logarithmic density of passive scalar $\ln c$; writes 'lncc.{xz, yz, xy, xy2}'

Module 'cosmicray.f90'

$ecr$ energy $e_{cr}$ of cosmic rays (?); writes 'ec.{xz, yz, xy, xy2}'

K.5 List of parameters for 'phiaver.in'

The following table lists all (at the time of writing, November 2003) possible inputs to the file 'phiaver.in'.

<table>
<thead>
<tr>
<th>Variable</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>rcylmphi</td>
<td>cylindrical radius $\varpi = \sqrt{x^2 + y^2}$ (useful for debugging azimuthal averages)</td>
</tr>
<tr>
<td>phimphi</td>
<td>azimuthal angle $\varphi = \arctan \frac{y}{x}$ (useful for debugging)</td>
</tr>
<tr>
<td>zmphi</td>
<td>$z$-coordinate (useful for debugging)</td>
</tr>
<tr>
<td>rmphi</td>
<td>spherical radius $r = \sqrt{\varpi^2 + z^2}$ (useful for debugging)</td>
</tr>
</tbody>
</table>

Module 'hydro.f90'

| urmphi   | $\langle u_{\varpi} \rangle_{\varphi}$ [cyl. polar coords ($\varpi, \varphi, z$)] |
| upmphi   | $\langle u_{\varphi} \rangle_{\varphi}$ |
| uzmphi   | $\langle u_z \rangle_{\varphi}$ |
| rurmpphi | $\langle \rho u_{\varpi} \rangle_{\varphi}$ |
### K.5 Parameters for ‘phiaver.in’

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Expression</th>
</tr>
</thead>
<tbody>
<tr>
<td>rupmphi</td>
<td>$\langle \rho u_\varphi \rangle_{\varphi}$</td>
</tr>
<tr>
<td>ruzmphi</td>
<td>$\langle \rho u_\varphi \rangle_{\varphi}$</td>
</tr>
<tr>
<td>ur2mphi</td>
<td>$\langle u_\varphi^2 \rangle_{\varphi}$</td>
</tr>
<tr>
<td>up2mphi</td>
<td>$\langle u_\varphi^2 \rangle_{\varphi}$</td>
</tr>
<tr>
<td>uz2mphi</td>
<td>$\langle u_z^2 \rangle_{\varphi}$</td>
</tr>
<tr>
<td>urupmphi</td>
<td>$\langle \rho u_\varphi u_\varphi \rangle_{\varphi}$</td>
</tr>
<tr>
<td>uruzmphi</td>
<td>$\langle u_\varphi u_z \rangle_{\varphi}$</td>
</tr>
<tr>
<td>upuzmphi</td>
<td>$\langle u_\varphi u_z \rangle_{\varphi}$</td>
</tr>
<tr>
<td>rurupmphi</td>
<td>$\langle \rho u_\varphi u_\varphi \rangle_{\varphi}$</td>
</tr>
<tr>
<td>ruruzmphi</td>
<td>$\langle \rho u_\varphi u_z \rangle_{\varphi}$</td>
</tr>
<tr>
<td>rupuzmphi</td>
<td>$\langle \rho u_z u_\varphi \rangle_{\varphi}$</td>
</tr>
<tr>
<td>ursphmphi</td>
<td>$\langle u_r \rangle_{\varphi}$</td>
</tr>
<tr>
<td>uthmphi</td>
<td>$\langle u_\varphi \rangle_{\varphi}$</td>
</tr>
<tr>
<td>rursphmphi</td>
<td>$\langle \rho u_r \rangle_{\varphi}$</td>
</tr>
<tr>
<td>ruthmphi</td>
<td>$\langle \rho u_\varphi \rangle_{\varphi}$</td>
</tr>
<tr>
<td>uumphi</td>
<td>shorthand for urmph, upmphi and uzmphi together</td>
</tr>
<tr>
<td>uusphmphi</td>
<td>shorthand for urspmphi, uthmphi and upmphi together</td>
</tr>
<tr>
<td>u2mphi</td>
<td>$\langle u^2 \rangle_{\varphi}$</td>
</tr>
<tr>
<td>fkinrsphmphi</td>
<td>$\langle \frac{1}{2} \rho u^2 u_z \rangle_{\varphi}$</td>
</tr>
</tbody>
</table>

#### Module ‘density.f90’

| Inrhomphi                        | $\langle \ln \rho \rangle_{\varphi}$ |
| rhomphi                          | $\langle \rho \rangle_{\varphi}$ |

#### Module ‘entropy.f90’

| ssmphi                           | $\langle s \rangle_{\varphi}$ |
| ss2mphi                          | $\langle s^2 \rangle_{\varphi}$ |
| cs2mphi                          | $\langle \epsilon_s^2 \rangle_{\varphi}$ |
| TTmphi                           | $\langle T \rangle_{\varphi}$ |
| dcoolmphi                        | surface cooling flux |
| fradrsphmphi.kramers             | $F_{\text{rad}}$ ($\varphi$-averaged, from Kramers’ opacity) |
| fconvrspshmphi                   | $\langle c_p \rho u_r T \rangle_{\varphi}$ |
| fconvthsphmphi                   | $\langle c_p \rho u_\varphi T \rangle_{\varphi}$ |
| fconvpsphmphi                    | $\langle c_p \rho u_z T \rangle_{\varphi}$ |
| ursphTTmphi                      | $\langle u_r T \rangle_{\varphi}$ |

#### Module ‘magnetic.f90’

| jbmphi                           | $\langle J \cdot B \rangle_{\varphi}$ |
| brmphi                           | $\langle B_\varphi \rangle_{\varphi}$ [cyl. polar coords ($\varphi, \varphi, z$)] |
| bpmphi                           | $\langle B_\varphi \rangle_{\varphi}$ |
| bzmphi                           | $\langle B_z \rangle_{\varphi}$ |
| bbbmphi                          | shorthand for brmph, bpmphi and bzmphi together |
| bbsphmphi                        | shorthand for brspmphi, bthmphi and bpmphi together |
| b2mphi                           | $\langle B^2 \rangle_{\varphi}$ |
| brshmphi                         | $\langle B_r \rangle_{\varphi}$ |
| bthmphi                          | $\langle B_\varphi \rangle_{\varphi}$ |

#### Module ‘anelastic.f90’

| Inrhomphi                        | $\langle \ln \rho \rangle_{\varphi}$ |
K.6 List of parameters for ‘xyaver.in’

The following table lists possible inputs to the file ‘xyaver.in’. This list is not complete and maybe outdated.

<table>
<thead>
<tr>
<th>Variable</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>( u2mz )</td>
<td>( \langle u^2 \rangle_{xy} )</td>
</tr>
<tr>
<td>( o2mz )</td>
<td>( \langle W^2 \rangle_{xy} )</td>
</tr>
<tr>
<td>( \text{div}u2mz )</td>
<td>( \langle (\nabla \cdot \mathbf{u})^2 \rangle_{xy} )</td>
</tr>
<tr>
<td>( \text{curl}lru2mz )</td>
<td>( \langle (\nabla \times g\mathbf{U})^2 \rangle_{xy} )</td>
</tr>
<tr>
<td>( \text{div}r2mz )</td>
<td>( \langle (\nabla \cdot g\mathbf{u})^2 \rangle_{xy} )</td>
</tr>
<tr>
<td>( \text{f}^{\text{mass}}zmz )</td>
<td>( \langle gu_z \rangle_{xy} )</td>
</tr>
<tr>
<td>( \text{f}^{\text{kin}}zmz )</td>
<td>( \frac{1}{2} \langle gu^2u_z \rangle_{xy} )</td>
</tr>
<tr>
<td>( \text{f}^{\text{kinz}}upmz )</td>
<td>( \frac{1}{2} \langle gu^2u_{z^1} \rangle_{xy} )</td>
</tr>
<tr>
<td>( \text{f}^{\text{kinz}}downmz )</td>
<td>( \frac{1}{2} \langle gu^2u_{z^\downarrow} \rangle_{xy} )</td>
</tr>
</tbody>
</table>

The table lists possible inputs to the file ‘xyaver.in’. This list is not complete and maybe outdated.
uxmz \langle u_x \rangle_{xy} \quad \text{(horiz. averaged } x \text{ velocity)}

uymz \langle u_y \rangle_{xy}

uzmz \langle u_z \rangle_{xy}

ffdownmz \quad \text{Filling factor of downflows}

uzupmz \langle u_z \rangle_{xy}^{↑}

uzdownmz \langle u_z \rangle_{xy}^{↓}

ruzupmz \langle \rho u_z \rangle_{xy}^{↑}

ruzdownmz \langle \rho u_z \rangle_{xy}^{↓}

divumz \langle \text{div } u \rangle_{xy}

uzdivumz \langle u_z \text{div } u \rangle_{xy}

oxmz \langle \omega_x \rangle_{xy}

oymz \langle \omega_y \rangle_{xy}

ozmz \langle \omega_z \rangle_{xy}

ux2mz \langle u_x^2 \rangle_{xy}

uy2mz \langle u_y^2 \rangle_{xy}

uz2mz \langle u_z^2 \rangle_{xy}

uz2upmz \langle u_z^2 \rangle_{xy}^{↑}

uz2downmz \langle u_z^2 \rangle_{xy}^{↓}

ox2mz \langle \omega_x^2 \rangle_{xy}

oy2mz \langle \omega_y^2 \rangle_{xy}

oz2mz \langle \omega_z^2 \rangle_{xy}

ruxmz \langle u_x u_y \rangle_{xy}

ruxymz \langle u_x u_y \rangle_{xy}

ruymz \langle u_y u_z \rangle_{xy}

ruzmz \langle u_z u_y \rangle_{xy}

rux2mz \langle u_x^2 u_y \rangle_{xy}

ruy2mz \langle u_y^2 u_x \rangle_{xy}

ruz2mz \langle u_z^2 u_y \rangle_{xy}

uxuymz \langle u_x u_y \rangle_{xy}

uxuzmz \langle u_x u_z \rangle_{xy}

uyuzmz \langle u_y u_z \rangle_{xy}

Rxymz \langle u_x'u_y' \rangle_{xy}

Rxyupmz \langle u_x'u_y' \rangle_{xy}^{↑}

Rxydownmz \langle u_x'u_y' \rangle_{xy}^{↓}

Rxzmz \langle u_x'u_z' \rangle_{xy}

Rxzupmz \langle u_x'u_z' \rangle_{xy}^{↑}

Rxzdownmz \langle u_x'u_z' \rangle_{xy}^{↓}

Ryzmz \langle u_y'u_z' \rangle_{xy}

Ryzupmz \langle u_y'u_z' \rangle_{xy}^{↑}

Ryzdownmz \langle u_y'u_z' \rangle_{xy}^{↓}

ruxuymz \langle pu_x u_y \rangle_{xy}

ruxuzmz \langle pu_x u_z \rangle_{xy}

ruyuzmz \langle pu_y u_z \rangle_{xy}

ruxuy2mz \langle pu_x u_y \rangle_{xy}

ruxuz2mz \langle pu_x u_z \rangle_{xy}

ruyuz2mz \langle pu_y u_z \rangle_{xy}

oxuxxmz \langle \omega_x u_x,u_z \rangle_{xy}
oyuxymz $\langle \omega y u_{x,y} \rangle_{xy}$
oxuxymz $\langle \omega z u_{y,x} \rangle_{xy}$
oyuyymz $\langle \omega y u_{y,x} \rangle_{xy}$
oxuzxymz $\langle \omega x u_{z,x} \rangle_{xy}$
oyuzymz $\langle \omega y u_{z,y} \rangle_{xy}$
uuyuxzmxz $\langle u_{y,x} u_{z,x} \rangle_{xy}$
uuyyuzmxz $\langle u_{y,y} u_{z,y} \rangle_{xy}$
uuyzzzmz $\langle u_{y,z} u_{z,z} \rangle_{xy}$

ekinmxz $\frac{1}{2} \langle u^2 \rangle_{xy}$
oumxz $\langle \omega \cdot u \rangle_{xy}$
Remz $\left[ \frac{\partial}{\partial x_j} (\nu S_{ij}) \right]_{xy}$
oguxmxz $\langle (\omega \cdot \nabla u)_{x} \rangle_{xy}$
ogauymz $\langle (\omega \cdot \nabla u)_{y} \rangle_{xy}$
oguzymz $\langle (\omega \cdot \nabla u)_{z} \rangle_{xy}$
ogux2mxz $\langle (\omega \cdot \nabla u)^2 \rangle_{xy}$
oguy2mxz $\langle (\omega \cdot \nabla u)^2 \rangle_{xy}$
oguz2mxz $\langle (\omega \cdot \nabla u)^2 \rangle_{xy}$
oxdvimxz $\langle \omega x \nabla \cdot u \rangle_{xy}$
oydvimxz $\langle \omega y \nabla \cdot u \rangle_{xy}$
ozdvimxz $\langle \omega z \nabla \cdot u \rangle_{xy}$

oxdvimx2mxz $\langle (\omega x \nabla \cdot u)^2 \rangle_{xy}$
oydvimx2mxz $\langle (\omega y \nabla \cdot u)^2 \rangle_{xy}$
ozdvimx2mxz $\langle (\omega z \nabla \cdot u)^2 \rangle_{xy}$
accpowzmz $\langle (u_z D u_z / Dt)^2 \rangle_{xy}$
accpowzupzmz $\langle (u_z D u_z / Dt)^2 \rangle_{xy+}$
accpowzdownzmz $\langle (u_z D u_z / Dt)^2 \rangle_{xy-}$

fkinmxmz $\frac{1}{2} \langle \rho u^2 u_x \rangle_{yz}$

---

**Module 'density.f90'**

rhomz $\langle \rho \rangle_{xy}$
houpmz $\langle \rho^+ \rangle_{xy}$
 rhodownzmz $\langle \rho^- \rangle_{xy}$
rho2mxz $\langle \rho^2 \rangle_{xy}$
rho2upmz $\langle \rho^2 \rangle_{xy}$
rho2downmz $\langle \rho^2 \rangle_{xy}$
rho2upmz $\langle \rho^2 \rangle_{xy}$
rho2downmz $\langle \rho^2 \rangle_{xy}$
gzlnrhomz $\langle \nabla z \ln \rho \rangle_{xy}$
uglnrhomz $\langle u \cdot \nabla \ln \rho \rangle_{xy}$
ugrhomz $\langle u \cdot \nabla \rho \rangle_{xy}$
uygznhrhomz $\langle u_y \nabla z \ln \rho \rangle_{xy}$
uzgynhromz $\langle u_z \nabla y \ln \rho \rangle_{xy}$
rho2mx $\langle \rho^2 \rangle_{yz}$

---

**Module 'entropy.f90'**
fradz \langle F_{rad}\rangle_{xy} \\
fcnvz \langle c_p u_z T\rangle_{xy} \\
Fenthz \langle c_p (\rho u_z)' T'\rangle_{xy} \\
Fenthupz \langle (c_p (\rho u_z)' T')_{\uparrow}\rangle_{xy} \\
Fenthdownz \langle (c_p (\rho u_z)' T')_{\downarrow}\rangle_{xy} \\
ssmz \langle s\rangle_{xy} \\
ssupmz \langle s_{\uparrow}\rangle_{xy} \\
ssdownmz \langle s_{\downarrow}\rangle_{xy} \\
ss2mz \langle s^2\rangle_{xy} \\
ss2upmz \langle s^2_{\uparrow}\rangle_{xy} \\
ss2downmz \langle s^2_{\downarrow}\rangle_{xy} \\
ssf2mz \langle s^2\prime\rangle_{xy} \\
ssf2upmz \langle s^2\prime_{\uparrow}\rangle_{xy} \\
ssf2downmz \langle s^2\prime_{\downarrow}\rangle_{xy} \\
ppmz \langle p\rangle_{xy} \\
TTmz \langle T\rangle_{xy} \\
TTdownmz \langle T_{\downarrow}\rangle_{xy} \\
TTupmz \langle T_{\uparrow}\rangle_{xy} \\
TT2mz \langle T^2\rangle_{xy} \\
TT2upmz \langle T^2_{\uparrow}\rangle_{xy} \\
TT2downmz \langle T^2_{\downarrow}\rangle_{xy} \\
TTf2mz \langle T^2\prime\rangle_{xy} \\
TTf2upmz \langle T^2\prime_{\uparrow}\rangle_{xy} \\
TTf2downmz \langle T^2\prime_{\downarrow}\rangle_{xy} \\
uxTTmz \langle u_x T\rangle_{xy} \\
uyTTmz \langle u_y T\rangle_{xy} \\
uzTTmz \langle u_z T\rangle_{xy} \\
uzTTupmz \langle (u_z T)_{\uparrow}\rangle_{xy} \\
uzTTdownmz \langle (u_z T)_{\downarrow}\rangle_{xy} \\
gTxgsxmz \langle \nabla T \times \nabla s\rangle_{xy} \\
gTxgsymz \langle \nabla T \times \nabla s\rangle_{y}\rangle_{xy} \\
gTxgszmz \langle \nabla T \times \nabla s\rangle_{z}\rangle_{xy} \\
gTxgsx2mz \langle \nabla T \times \nabla s^2\rangle_{xy} \\
gTxgsy2mz \langle \nabla T \times \nabla s^2\rangle_{y}\rangle_{xy} \\
gTxgsz2mz \langle \nabla T \times \nabla s^2\rangle_{z}\rangle_{xy} \\
fradz_kramers F_{rad} (from Kramers' opacity) \\
fradz_Kprof F_{rad} (from Kprof) \\
fradz_constchi F_{rad} (from \chi_{const}) \\
fturbz \langle \rho T \chi_0 (\nabla z s)\rangle_{xy} (turbulent heat flux) \\
fturbtz \langle \rho T \chi_0 (\nabla z s)'\rangle_{xy} (turbulent heat flux) \\
fturbmz \langle \rho T \chi_0 (\nabla z s)_{\uparrow}\rangle_{xy} (turbulent heat flux) \\
fturbfz \langle \rho T \chi_0 (\nabla z s)'_{\uparrow}\rangle_{xy} (turbulent heat flux) \\
dcoolz \text{surface cooling flux} \\
heatmz \text{heating} \\
Kkramersmz \langle K_0 T^{(0-b)}/(\rho^{(a+1)})\rangle_{xy} \\
ethmz \langle \rho e\rangle_{xy}
Module 'magnetic.f90'

<table>
<thead>
<tr>
<th>Symbol</th>
<th>Expression</th>
</tr>
</thead>
<tbody>
<tr>
<td>axmz</td>
<td>$\langle A_x \rangle_{xy}$</td>
</tr>
<tr>
<td>aymz</td>
<td>$\langle A_y \rangle_{xy}$</td>
</tr>
<tr>
<td>azmz</td>
<td>$\langle A_z \rangle_{xy}$</td>
</tr>
<tr>
<td>abuxmz</td>
<td>$\langle (A \cdot B) u_x \rangle_{xy}$</td>
</tr>
<tr>
<td>abuymz</td>
<td>$\langle (A \cdot B) u_y \rangle_{xy}$</td>
</tr>
<tr>
<td>abuzmz</td>
<td>$\langle (A \cdot B) u_z \rangle_{xy}$</td>
</tr>
<tr>
<td>uabxmz</td>
<td>$\langle (u \cdot A) B_x \rangle_{xy}$</td>
</tr>
<tr>
<td>uabymz</td>
<td>$\langle (u \cdot A) B_y \rangle_{xy}$</td>
</tr>
<tr>
<td>uabzmz</td>
<td>$\langle (u \cdot A) B_z \rangle_{xy}$</td>
</tr>
<tr>
<td>bbxmz</td>
<td>$\langle B'<em>x \rangle</em>{xy}$</td>
</tr>
<tr>
<td>bbymz</td>
<td>$\langle B'<em>y \rangle</em>{xy}$</td>
</tr>
<tr>
<td>bbzmz</td>
<td>$\langle B'<em>z \rangle</em>{xy}$</td>
</tr>
<tr>
<td>bxmz</td>
<td>$\langle B_x \rangle_{xy}$</td>
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<tr>
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<tr>
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<tr>
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<td>$\langle u_z b_z \rangle_{xy}$</td>
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<td>$\langle E \times A \rangle_{xy} \mid x$</td>
</tr>
<tr>
<td>examz2</td>
<td>$\langle E \times A \rangle_{xy} \mid y$</td>
</tr>
<tr>
<td>examz3</td>
<td>$\langle E \times A \rangle_{xy} \mid z$</td>
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<tr>
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<td>$\langle E_{\text{hyperl}3} \times A \rangle_{xy} \mid y$</td>
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<td>$\langle E_{\text{hyperl}3} \times A \rangle_{xy} \mid z$</td>
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<tr>
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<tr>
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**Module ‘bfield.f90’**

<table>
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<tbody>
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<tr>
<td>b2mz</td>
<td>$\langle B^2 \rangle_{xy}$</td>
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<tr>
<td>bmxz</td>
<td>$\langle B_x \rangle_{xy}$</td>
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<tr>
<td>bmxz</td>
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<tr>
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<tr>
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<tr>
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<tr>
<td>by2mz</td>
<td>$\langle B^2_y \rangle_{xy}$</td>
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<tr>
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**Module ‘density_stratified.f90’**

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<tbody>
<tr>
<td>drhomz</td>
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<tr>
<td>drho2mz</td>
<td>$\langle (\Delta \rho / \rho_0)^2 \rangle_{xy}$</td>
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**Module ‘gravity_simple.f90’**

<table>
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<tbody>
<tr>
<td>epotmz</td>
<td>$\langle \Phi_{\text{grav}} \rangle_{xy}$</td>
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<tr>
<td>epotuzmz</td>
<td>$\langle \Phi_{\text{grav} u_z} \rangle_{xy}$ (potential energy flux)</td>
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**Module ‘hydro_potential.f90’**

<table>
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<tr>
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<tbody>
<tr>
<td>u2mz</td>
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<tr>
<td>o2mz</td>
<td>$\langle W^2 \rangle_{xy}$</td>
</tr>
<tr>
<td>divu2mz</td>
<td>$\langle (\nabla \cdot u)^2 \rangle_{xy}$</td>
</tr>
<tr>
<td>curlru2mz</td>
<td>$\langle (\nabla \times \Phi U)^2 \rangle_{xy}$</td>
</tr>
<tr>
<td>divru2mz</td>
<td>$\langle (\nabla \cdot \Phi u)^2 \rangle_{xy}$</td>
</tr>
<tr>
<td>fmasszmz</td>
<td>$\langle \rho u_z \rangle_{xy}$</td>
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</table>
\begin{align*}
  f\text{kin}zmz & \quad \langle \frac{1}{2} \rho u^2 u_z \rangle_{xy} \\
  uxzmz & \quad \langle u_x \rangle_{xy} \quad \text{(horiz. averaged } x \text{ velocity)} \\
  uymz & \quad \langle u_y \rangle_{xy} \\
  uzmz & \quad \langle u_z \rangle_{xy} \\
  uzupmz & \quad \langle u_z^+ \rangle_{xy} \\
  uzdownmz & \quad \langle u_z^- \rangle_{xy} \\
  ruzupmz & \quad \langle \rho u_z^+ \rangle_{xy} \\
  ruzdownmz & \quad \langle \rho u_z^- \rangle_{xy} \\
  divumz & \quad \langle \text{div } u \rangle_{xy} \\
  uzdivumz & \quad \langle u_z \text{div } u \rangle_{xy} \\
  oxmz & \quad \langle \omega_x \rangle_{xy} \\
  oymz & \quad \langle \omega_y \rangle_{xy} \\
  ozmz & \quad \langle \omega_z \rangle_{xy} \\
  ux2mz & \quad \langle u^2_x \rangle_{xy} \\
  uymz & \quad \langle u^2_y \rangle_{xy} \\
  uz2mz & \quad \langle u^2_z \rangle_{xy} \\
  ox2mz & \quad \langle \omega^2_x \rangle_{xy} \\
  oy2mz & \quad \langle \omega^2_y \rangle_{xy} \\
  oz2mz & \quad \langle \omega^2_z \rangle_{xy} \\
  ruxmz & \quad \langle \rho u_x \rangle_{xy} \\
  ruymz & \quad \langle \rho u_y \rangle_{xy} \\
  ruzmz & \quad \langle \rho u_z \rangle_{xy} \\
  rux2mz & \quad \langle \rho u^2_x \rangle_{xy} \\
  ruy2mz & \quad \langle \rho u^2_y \rangle_{xy} \\
  ruz2mz & \quad \langle \rho u^2_z \rangle_{xy} \\
  uxuymz & \quad \langle u_x u_y \rangle_{xy} \\
  uxuzmz & \quad \langle u_x u_z \rangle_{xy} \\
  uyuzmz & \quad \langle u_y u_z \rangle_{xy} \\
  ruxuymz & \quad \langle \rho u_x u_y \rangle_{xy} \\
  ruxuzmz & \quad \langle \rho u_x u_z \rangle_{xy} \\
  ruyuzmz & \quad \langle \rho u_y u_z \rangle_{xy} \\
  ruxuy2mz & \quad \langle \rho u_x u_y \rangle_{xy} \\
  ruxuz2mz & \quad \langle \rho u_x u_z \rangle_{xy} \\
  ruyuz2mz & \quad \langle \rho u_y u_z \rangle_{xy} \\
  oxuxxmz & \quad \langle \omega_x u_{x,x} \rangle_{xy} \\
  oyuxymz & \quad \langle \omega_y u_{x,y} \rangle_{xy} \\
  oxuuxmz & \quad \langle \omega_z u_{x,x} \rangle_{xy} \\
  oyuyymz & \quad \langle \omega_y u_{y,y} \rangle_{xy} \\
  oxuyxmz & \quad \langle \omega_z u_{y,y} \rangle_{xy} \\
  oyuyzmx & \quad \langle \omega_y u_{z,y} \rangle_{xy} \\
  uyyuzymz & \quad \langle u_{y,y} u_{z,y} \rangle_{xy} \\
  uyyuzymz & \quad \langle u_{y,y} u_{z,y} \rangle_{xy} \\
  uyyuzymz & \quad \langle u_{y,y} u_{z,y} \rangle_{xy} \\
  ekinmz & \quad \langle \frac{1}{2} \rho u^2 \rangle_{xy} \\
  oumz & \quad \langle \omega \cdot u \rangle_{xy} \\
  Remz & \quad \langle \frac{\partial u \cdot u}{\partial x_j (\nu S_{ij})} \rangle_{xy}
\end{align*}
<table>
<thead>
<tr>
<th>Variable</th>
<th>Expression</th>
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</thead>
<tbody>
<tr>
<td>oguxmz</td>
<td>$\langle (\omega \cdot \nabla u)<em>{x/y} \rangle</em>{xy}$</td>
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<tr>
<td>oguymz</td>
<td>$\langle (\omega \cdot \nabla u)<em>{y/y} \rangle</em>{xy}$</td>
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<tr>
<td>oguzmz</td>
<td>$\langle (\omega \cdot \nabla u)<em>{z/z} \rangle</em>{xy}$</td>
</tr>
<tr>
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<td>$\langle (\omega \cdot \nabla u)<em>{x/z} \rangle</em>{xy}$</td>
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<tr>
<td>oguy2mz</td>
<td>$\langle (\omega \cdot \nabla u)<em>{y/z} \rangle</em>{xy}$</td>
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<tr>
<td>oguz2mz</td>
<td>$\langle (\omega \cdot \nabla u)<em>{z/y} \rangle</em>{xy}$</td>
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<td>$\langle \omega_x \nabla \cdot u \rangle_{xy}$</td>
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<tr>
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<td>$\langle \omega_y \nabla \cdot u \rangle_{xy}$</td>
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<tr>
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<td>$\langle (\omega_y \nabla \cdot u)^2 \rangle_{xy}$</td>
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Module ‘magnetic_shearboxJ.f90’

<table>
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<tr>
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<td>$\langle (A \cdot B) u_x \rangle_{xy}$</td>
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<tr>
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<td>$\langle (A \cdot B) u_y \rangle_{xy}$</td>
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<td>$\langle (A \cdot B) u_z \rangle_{xy}$</td>
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<td>$\langle (u \cdot A) B_x \rangle_{xy}$</td>
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<tr>
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<td>$\langle (u \cdot A) B_y \rangle_{xy}$</td>
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<td>$\langle (u \cdot A) B_z \rangle_{xy}$</td>
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<td>$\langle B_x' \rangle_{xy}$</td>
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<tr>
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<td>$\langle B_y' \rangle_{xy}$</td>
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<tr>
<td>bbzmz</td>
<td>$\langle B_z' \rangle_{xy}$</td>
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<td>$\langle B_z^2/\theta \rangle_{xy}$</td>
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<tr>
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<td>$\langle (B_z^2/2\mu_0) \rangle_{xy}$</td>
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<td>betamz</td>
<td>$\langle \beta \rangle_{xy}$</td>
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</table>
$\langle \beta^2 \rangle_{xy}$

$\langle J \cdot B \rangle |_{xy}$

$\langle \nabla^6 A \cdot B \rangle |_{xy}$

$\langle \nabla^6 A \rangle |_{xy}$

$\langle \nabla^6 A \rangle |_{xy}$

$\langle A \cdot B \rangle |_{xy}$

$\langle u \cdot B \rangle |_{xy}$

$\langle u \cdot A \rangle |_{xy}$

$\langle u_x b_x \rangle |_{xy}$

$\langle u_y b_x \rangle |_{xy}$

$\langle u_z b_x \rangle |_{xy}$

$\langle u_x b_y \rangle |_{xy}$

$\langle u_y b_y \rangle |_{xy}$

$\langle u_z b_y \rangle |_{xy}$

$\langle u_x b_z \rangle |_{xy}$

$\langle u_y b_z \rangle |_{xy}$

$\langle u_z b_z \rangle |_{xy}$

$\langle E \times A \rangle |_{xy}$

$\langle E \times A \rangle |_{xy}$

$\langle E \times A \rangle |_{xy}$

$\langle E_{hyper 3} \times A \rangle |_{xy}$

$\langle E_{hyper 3} \times A \rangle |_{xy}$

$\langle E_{hyper 3} \times A \rangle |_{xy}$

$\langle \eta \rangle |_{xy}$

$\langle B_x B_y \rangle |_{xy}$

$\langle B_x B_z \rangle |_{xy}$

$\langle B_y B_z \rangle |_{xy}$

$\langle B^2 \rangle |_{xy}$

$\langle B^2 \rangle |_{xy}$

$\langle j^2 \rangle |_{xy}$

Average poynting flux in z direction

$\langle \eta \mu_0 j^2 \rangle |_{xy}$

Module 'meanfield.f90'

$\langle q_p \rangle |_{xy}$

Module 'shock_highorder.f90'

Module 'temperature_idealgas.f90'

$\langle p \rangle |_{xy}$

$\langle T \rangle |_{xy}$

$\langle e_{th} \rangle |_{xy}$

$\langle \nabla p \rangle |_{xy}$

$\langle \nabla p \rangle |_{xy}$

$\langle \nabla p \rangle |_{xy}$

$\langle T^2 \rangle |_{xy}$

$\langle u_x T \rangle |_{xy}$

$\langle u_y T \rangle |_{xy}$
K.7 Parameters for ‘xzaver.in’

The following table lists possible inputs to the file ‘xzaver.in’. This list is not complete and maybe outdated.

<table>
<thead>
<tr>
<th>Variable</th>
<th>Meaning</th>
</tr>
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<tbody>
<tr>
<td><strong>Module ‘hydro.f90’</strong></td>
<td></td>
</tr>
<tr>
<td><code>uxmy</code></td>
<td>( \langle u_x \rangle_{xz} )</td>
</tr>
<tr>
<td><code>uemy</code></td>
<td>( \langle u_y \rangle_{xz} )</td>
</tr>
<tr>
<td><code>uzmy</code></td>
<td>( \langle u_z \rangle_{xz} )</td>
</tr>
<tr>
<td><code>oumy</code></td>
<td>( \langle \omega \cdot u \rangle_{xz} )</td>
</tr>
<tr>
<td><strong>Module ‘density.f90’</strong></td>
<td></td>
</tr>
<tr>
<td><code>rhomy</code></td>
<td>( \langle \rho \rangle_{xz} )</td>
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<tr>
<td><strong>Module ‘entropy.f90’</strong></td>
<td></td>
</tr>
<tr>
<td><code>ssmy</code></td>
<td>( \langle s \rangle_{xz} )</td>
</tr>
<tr>
<td><code>ppmy</code></td>
<td>( \langle p \rangle_{xz} )</td>
</tr>
<tr>
<td><code>TTmy</code></td>
<td>( \langle T \rangle_{xz} )</td>
</tr>
<tr>
<td><strong>Module ‘magnetic.f90’</strong></td>
<td></td>
</tr>
<tr>
<td><code>bxmy</code></td>
<td>( \langle B_x \rangle_{xz} )</td>
</tr>
<tr>
<td><code>bymy</code></td>
<td>( \langle B_y \rangle_{xz} )</td>
</tr>
</tbody>
</table>
K.8 List of parameters for ‘yzaver.in’

The following table lists possible inputs to the file ‘yzaver.in’. This list is not complete and maybe outdated.

<table>
<thead>
<tr>
<th>Variable</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Module ‘hydro.f90’</td>
</tr>
</tbody>
</table>
### Parameters for ‘yzaver.in’

**Module ‘density.f90’**

- $\langle \rho \rangle_{yz}$

**Module ‘entropy.f90’**

- $\langle s \rangle_{yz}$
- $\langle s^2 \rangle_{yz}$
- $\langle p \rangle_{yz}$
- $\langle T \rangle_{yz}$
- $\langle T^2 \rangle_{yz}$
- $\langle u_T \rangle_{yz}$
- $\langle u_T \rangle_{yz}$
- $\langle c_p \partial u_T \rangle_{yz}$

**Module ‘magnetic.f90’**

- $\langle B^2 \rangle_{yz}$
- $\langle B_x \rangle_{yz}$
- $\langle B_y \rangle_{yz}$
- $\langle B_z \rangle_{yz}$

K.8 Parameters for ‘yzaver.in’ 249
### The Pencil Code

<table>
<thead>
<tr>
<th>Description</th>
<th>Formula</th>
</tr>
</thead>
<tbody>
<tr>
<td>$bx2mx$</td>
<td>$\langle B_x^2 \rangle_{yz}$</td>
</tr>
<tr>
<td>$by2mx$</td>
<td>$\langle B_y^2 \rangle_{yz}$</td>
</tr>
<tr>
<td>$bz2mx$</td>
<td>$\langle B_z^2 \rangle_{yz}$</td>
</tr>
<tr>
<td>$bxbymx$</td>
<td>$\langle B_x B_y \rangle_{yz}$</td>
</tr>
<tr>
<td>$bxbzmx$</td>
<td>$\langle B_x B_z \rangle_{yz}$</td>
</tr>
<tr>
<td>$bybzmx$</td>
<td>$\langle B_y B_z \rangle_{yz}$</td>
</tr>
<tr>
<td>$betamx$</td>
<td>$\langle \beta \rangle_{yz}$</td>
</tr>
<tr>
<td>$beta2mx$</td>
<td>$\langle \beta^2 \rangle_{yz}$</td>
</tr>
<tr>
<td>$etatotalmx$</td>
<td>$\langle \eta \rangle_{yz}$</td>
</tr>
</tbody>
</table>

**Module 'bfield.f90'**

<table>
<thead>
<tr>
<th>Description</th>
<th>Formula</th>
</tr>
</thead>
<tbody>
<tr>
<td>$bmx$</td>
<td>$\langle B \rangle_{yz}$</td>
</tr>
<tr>
<td>$b2mx$</td>
<td>$\langle B^2 \rangle_{yz}$</td>
</tr>
<tr>
<td>$bxmx$</td>
<td>$\langle B_x \rangle_{yz}$</td>
</tr>
<tr>
<td>$bymx$</td>
<td>$\langle B_y \rangle_{yz}$</td>
</tr>
<tr>
<td>$bzmx$</td>
<td>$\langle B_z \rangle_{yz}$</td>
</tr>
<tr>
<td>$bx2mx$</td>
<td>$\langle B_x^2 \rangle_{yz}$</td>
</tr>
<tr>
<td>$by2mx$</td>
<td>$\langle B_y^2 \rangle_{yz}$</td>
</tr>
<tr>
<td>$bz2mx$</td>
<td>$\langle B_z^2 \rangle_{yz}$</td>
</tr>
<tr>
<td>$bxbymx$</td>
<td>$\langle B_x B_y \rangle_{yz}$</td>
</tr>
<tr>
<td>$bxbzmx$</td>
<td>$\langle B_x B_z \rangle_{yz}$</td>
</tr>
<tr>
<td>$bybzmx$</td>
<td>$\langle B_y B_z \rangle_{yz}$</td>
</tr>
<tr>
<td>$betamx$</td>
<td>$\langle \beta \rangle_{yz}$</td>
</tr>
<tr>
<td>$beta2mx$</td>
<td>$\langle \beta^2 \rangle_{yz}$</td>
</tr>
</tbody>
</table>

**Module 'density_stratified.f90'**

<table>
<thead>
<tr>
<th>Description</th>
<th>Formula</th>
</tr>
</thead>
<tbody>
<tr>
<td>$drhomx$</td>
<td>$\langle \Delta \rho/\rho_0 \rangle_{yz}$</td>
</tr>
<tr>
<td>$drho2mx$</td>
<td>$\langle (\Delta \rho/\rho_0)^2 \rangle_{yz}$</td>
</tr>
</tbody>
</table>

**Module 'gravity_simple.f90'**

<table>
<thead>
<tr>
<th>Description</th>
<th>Formula</th>
</tr>
</thead>
<tbody>
<tr>
<td>$epotmx$</td>
<td>$\langle \Phi_{grav} \rangle_{yz}$</td>
</tr>
<tr>
<td>$epotuxmx$</td>
<td>$\langle \Phi_{grav u_x} \rangle_{yz}$ (potential energy flux)</td>
</tr>
</tbody>
</table>

**Module 'hydro_potential.f90'**

<table>
<thead>
<tr>
<th>Description</th>
<th>Formula</th>
</tr>
</thead>
<tbody>
<tr>
<td>$uxmx$</td>
<td>$\langle u_x \rangle_{yz}$</td>
</tr>
<tr>
<td>$uymx$</td>
<td>$\langle u_y \rangle_{yz}$</td>
</tr>
<tr>
<td>$uzmx$</td>
<td>$\langle u_z \rangle_{yz}$</td>
</tr>
<tr>
<td>$ruxmx$</td>
<td>$\langle u_u x \rangle_{yz}$</td>
</tr>
<tr>
<td>$ruymx$</td>
<td>$\langle u_u y \rangle_{yz}$</td>
</tr>
<tr>
<td>$ruzmx$</td>
<td>$\langle u_u z \rangle_{yz}$</td>
</tr>
<tr>
<td>$rux2mx$</td>
<td>$\langle u^2_u x \rangle_{yz}$</td>
</tr>
<tr>
<td>$ruy2mx$</td>
<td>$\langle u^2_u y \rangle_{yz}$</td>
</tr>
<tr>
<td>$ruz2mx$</td>
<td>$\langle u^2_u z \rangle_{yz}$</td>
</tr>
<tr>
<td>$ruxumx$</td>
<td>$\langle u_x u_y \rangle_{yz}$</td>
</tr>
<tr>
<td>$ruxuzmx$</td>
<td>$\langle u_x u_z \rangle_{yz}$</td>
</tr>
<tr>
<td>$ruyuzmx$</td>
<td>$\langle u_y u_z \rangle_{yz}$</td>
</tr>
<tr>
<td>$ux2mx$</td>
<td>$\langle u^2_x \rangle_{yz}$</td>
</tr>
<tr>
<td>$uy2mx$</td>
<td>$\langle u^2_y \rangle_{yz}$</td>
</tr>
<tr>
<td>$uz2mx$</td>
<td>$\langle u^2_z \rangle_{yz}$</td>
</tr>
<tr>
<td>$ox2mx$</td>
<td>$\langle \omega^2_x \rangle_{yz}$</td>
</tr>
</tbody>
</table>
K.9 Parameters for ‘yaver.in’

\begin{align*}
oy2mx & = \langle \omega_y^2 \rangle_{yz} \\
oz2mx & = \langle \omega_z^2 \rangle_{yz} \\
uxuymx & = \langle u_x u_y \rangle_{yz} \\
uxuzmx & = \langle u_x u_z \rangle_{yz} \\
uyuzmx & = \langle u_y u_z \rangle_{yz} \\
oumx & = \langle \omega \cdot u \rangle_{yz} \\
ekinmx & = \langle \frac{1}{2} \rho u^2 \rangle_{xy}
\end{align*}

Module ‘magnetic_shearboxJ.f90’

\begin{align*}
b2mx & = \langle B^2 \rangle_{yz} \\
bxmx & = \langle B_x \rangle_{yz} \\
bymx & = \langle B_y \rangle_{yz} \\
bzmx & = \langle B_z \rangle_{yz} \\
bx2mx & = \langle B_x^2 \rangle_{yz} \\
by2mx & = \langle B_y^2 \rangle_{yz} \\
bz2mx & = \langle B_z^2 \rangle_{yz} \\
ouxzmxy & = \langle B_x B_y \rangle_{yz} \\
oubzmxy & = \langle B_x B_z \rangle_{yz} \\
betamx & = \langle \beta \rangle_{yz} \\
beta2mx & = \langle \beta^2 \rangle_{yz} \\
etatotalmx & = \langle \eta \rangle_{yz}
\end{align*}

Module ‘shock_highorder.f90’

Module ‘temperature_idealgas.f90’

\begin{align*}
ppmx & = \langle p \rangle_{yz} \\
TTmx & = \langle T \rangle_{yz}
\end{align*}

Module ‘thermal_energy.f90’

\begin{align*}
ppmx & = \langle p \rangle_{yz} \\
TTmx & = \langle T \rangle_{yz}
\end{align*}

Module ‘viscosity.f90’

\begin{align*}
fviscmx & = \langle 2 \nu u_i S_{ix} \rangle_{yz} (x\text{-component of viscous flux}) \\
umx & = \langle \nu \rangle_{yz} (yz\text{-averaged viscosity})
\end{align*}

K.9 List of parameters for ‘yaver.in’

The following table lists possible inputs to the file ‘yaver.in’. This list is not complete and maybe outdated.

<table>
<thead>
<tr>
<th>Variable</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>uxmxz</td>
<td>\langle u_x \rangle_y</td>
</tr>
<tr>
<td>uymxz</td>
<td>\langle u_y \rangle_y</td>
</tr>
<tr>
<td>uzmxz</td>
<td>\langle u_z \rangle_y</td>
</tr>
</tbody>
</table>

Module ‘hydro.f90’
ux2mxz \( \langle u_x^2 \rangle_y \)
uy2mxz \( \langle u_y^2 \rangle_y \)
uz2mxz \( \langle u_z^2 \rangle_y \)
uxuymxz \( \langle u_x u_y \rangle_y \)
uxuzmxz \( \langle u_x u_z \rangle_y \)
uyuzmxz \( \langle u_y u_z \rangle_y \)
oumzx \( \langle \omega \cdot u \rangle_y \)

\underline{Module 'density.f90'}

\( \rho \text{x} \)z \( \langle \rho \rangle_y \)

\underline{Module 'entropy.f90'}

\( T \text{T} \)zx \( \langle T \rangle_y \)
ssmxz \( \langle s \rangle_y \)

\underline{Module 'magnetic.f90'}

b2mxz \( \langle B^2 \rangle_y \)
axmxz \( \langle A_x \rangle_y \)
aymxz \( \langle A_y \rangle_y \)
azmxz \( \langle A_z \rangle_y \)
bx1mxz \( \langle B_x \rangle_y \)
by1mxz \( \langle B_y \rangle_y \)
bz1mxz \( \langle B_z \rangle_y \)
bzxmxz \( \langle B_x \rangle_y \)
bymxz \( \langle B_y \rangle_y \)
bzmxz \( \langle B_z \rangle_y \)
jxmxz \( \langle J_x \rangle_y \)
jymxz \( \langle J_y \rangle_y \)
jzmxz \( \langle J_z \rangle_y \)
bx2mxz \( \langle B_x^2 \rangle_y \)
by2mxz \( \langle B_y^2 \rangle_y \)
bz2mxz \( \langle B_z^2 \rangle_y \)
 bxbymxz \( \langle B_x B_y \rangle_y \)
 bxbzmxz \( \langle B_x B_z \rangle_y \)
 bybzmxx \( \langle B_y B_z \rangle_y \)
 uybmxz \( \langle U_y B_x \rangle_y \)
 uybzmxz \( \langle U_y B_z \rangle_y \)
 Exmxz \( \langle E_x \rangle_y \)
 Eymxz \( \langle E_y \rangle_y \)
 Ezmxz \( \langle E_z \rangle_y \)
vAmxz \( \langle v_A \rangle_y \)

\underline{Module 'density_stratified.f90'}

\( \Delta \rho \rho_0 \text{x} \)z \( \langle \Delta \rho / \rho_0 \rangle_y \)
\( \Delta \rho \rho_0 \text{x} \)z \( \langle (\Delta \rho / \rho_0)^2 \rangle_y \)

\underline{Module 'hydro_potential.f90'}

uxmxz \( \langle u_x \rangle_y \)
uymxz \( \langle u_y \rangle_y \)
K.9 Parameters for 'yaver.in'

uzmxz \(\langle u_z \rangle_y\)
ux2mxz \(\langle u_x^2 \rangle_y\)
uy2mxz \(\langle u_y^2 \rangle_y\)
uz2mxz \(\langle u_z^2 \rangle_y\)
uxuymxz \(\langle u_x u_y \rangle_y\)
uxuzmxz \(\langle u_x u_z \rangle_y\)
uyuzmxz \(\langle u_y u_z \rangle_y\)
oumxz \(\langle \omega \cdot u \rangle_y\)

Module 'magnetic_shearboxJ.f90'

b2mxz \(\langle B^2 \rangle_y\)
axmxz \(\langle A_x \rangle_y\)
aymxz \(\langle A_y \rangle_y\)
azmxz \(\langle A_z \rangle_y\)
bx1mxz \(\langle |B_x| \rangle_y\)
by1mxz \(\langle |B_y| \rangle_y\)
bz1mxz \(\langle |B_z| \rangle_y\)
bxmxz \(\langle B_x \rangle_y\)
bymxz \(\langle B_y \rangle_y\)
bfmxz \(\langle B_z \rangle_y\)
jxmxz \(\langle J_x \rangle_y\)
jymxz \(\langle J_y \rangle_y\)
jzmxz \(\langle J_z \rangle_y\)
bx2mxz \(\langle B_x^2 \rangle_y\)
by2mxz \(\langle B_y^2 \rangle_y\)
bz2mxz \(\langle B_z^2 \rangle_y\)
bxbymxz \(\langle B_x B_y \rangle_y\)
xbzbmxz \(\langle B_x B_z \rangle_y\)
bybzmxz \(\langle B_y B_z \rangle_y\)
uybxmxz \(\langle U_y B_x \rangle_y\)
uybzmxz \(\langle U_y B_z \rangle_y\)
Exmxz \(\langle E_x \rangle_y\)
Eymxz \(\langle E_y \rangle_y\)
Ezmxz \(\langle E_z \rangle_y\)
vAmxz \(\langle v^2_A \rangle_y\)

Module 'meanfield.f90'

peffmxz \(\langle P_{\text{eff}} \rangle_y\)
alpmxz \(\langle \alpha \rangle_y\)

Module 'temperature_idealgas.f90'

TTmxz \(\langle T \rangle_y\)
EmAIA94mxz Emission off AIA 94 channel integrated over y direction
EmAIA131mxz Emission off AIA 131 channel integrated over y direction
EmAIA171mxz Emission off AIA 171 channel integrated over y direction
EmAIA193mxz Emission off AIA 193 channel integrated over y direction
EmAIA211mxz Emission off AIA 211 channel integrated over y direction
EmAIA304mxz Emission off AIA 304 channel integrated over y direction
\textit{EmAIA335mxz} \quad \text{Emission off AIA 335 channel integrated over y direction}
\textit{EmXRTmxz} \quad \text{Emission off XRT Al-poly channel integrated over y direction}

\textbf{Module ‘thermal\_energy.f90’}

\begin{center}
\begin{tabular}{ll}
\texttt{TTmxz} & \langle (T) \rangle_y
\end{tabular}
\end{center}

\section*{K.10 List of parameters for ‘zaver.in’}

The following table lists possible inputs to the file ‘zaver.in’. This list is not complete and maybe outdated.

\begin{center}
\begin{tabular}{ll}
\textbf{Variable} & \textbf{Meaning} \\
\hline
uxmxy & \langle u_x \rangle_z \\
uymxy & \langle u_y \rangle_z \\
uzmxy & \langle u_z \rangle_z \\
xupmxy & \langle u_{x1} \rangle_z \\
exdownmxy & \langle u_{x1} \rangle_z \\
ruxupmxy & \langle \rho u_{x1} \rangle_z \\
ruxdownmxy & \langle \rho u_{x1} \rangle_z \\
x2upmxy & \langle u_{x1}^2 \rangle_z \\
x2downmxy & \langle u_{x1}^2 \rangle_z \\
ffdownmxy & Filling factor of downflows \\
uxuymxy & \langle u_x u_y \rangle_z \\
uxuzmxy & \langle u_x u_z \rangle_z \\
uuyuzmxy & \langle u_y u_z \rangle_z \\
rxymxy & \langle u_x' u_y' \rangle_z \\
rxyupmxy & \langle (u_x' u_y')^\uparrow \rangle_z \\
rxydownmxy & \langle (u_x' u_y')^\downarrow \rangle_z \\
rzmxy & \langle u_y' u_z' \rangle_z \\
rzupmxy & \langle (u_y' u_x')^\uparrow \rangle_z \\
rzdownmxy & \langle (u_y' u_x')^\downarrow \rangle_z \\
orxmx & \langle \omega_x \rangle_z \\
orymxy & \langle \omega_y \rangle_z \\
orzmx & \langle \omega_z \rangle_z \\
oumxy & \langle \omega \cdot u \rangle_z \\
pxzmx & \langle (\omega + 2\Omega) \rangle_z \quad \text{(z component of potential vorticity)} \\
uguxmxy & \langle u \cdot \nabla u \rangle_z \\
uguyymxy & \langle u \cdot \nabla u \rangle_y \\
uguzmxy & \langle u \cdot \nabla u \rangle_z \\
ruxmxy & \langle \rho u_x \rangle_z \\
ruxymxy & \langle \rho u_y \rangle_z \\
ruezmxy & \langle \rho u_z \rangle_z \\
x2mx & \langle u_x^2 \rangle_z \\
\end{tabular}
\end{center}
K.10 Parameters for 'zaver.in'

\[ u_{y2mxy} \langle u_y^2 \rangle_z \]
\[ u_{z2mxy} \langle u_z^2 \rangle_z \]
\[ r_{ux2mxy} \langle pu_x^2 \rangle_z \]
\[ r_{uy2mxy} \langle pu_y^2 \rangle_z \]
\[ r_{uz2mxy} \langle pu_z^2 \rangle_z \]
\[ r_{uxymxy} \langle pu_x u_y \rangle_z \]
\[ r_{uxzmxy} \langle pu_x u_z \rangle_z \]
\[ r_{uyzmxy} \langle pu_y u_z \rangle_z \]
\[ f_{kinmxy} \langle \frac{1}{2} \rho u_x^2 \rangle_z \]
\[ f_{kinymxy} \langle \frac{1}{2} \rho u_y^2 \rangle_z \]
\[ f_{kinxupmxy} \langle \frac{1}{2} \rho u_x^2 u_y \rangle_z \]
\[ f_{kinxdownmxy} \langle \frac{1}{2} \rho u_x^2 u_z \rangle_z \]

Module 'density.f90'

\[ \rho_{omxy} \langle \rho \rangle_z \]

Module 'entropy.f90'

\[ TT_{mxy} \langle T \rangle_z \]
\[ ss_{mxy} \langle s \rangle_z \]
\[ uu_{TT_{mxy}} \langle u_x T \rangle_z \]
\[ uu_{TT_{mxy}} \langle u_y T \rangle_z \]
\[ uu_{TT_{mxy}} \langle u_z T \rangle_z \]
\[ g_{Txmxy} \langle \nabla_x T \rangle_z \]
\[ g_{Tymxy} \langle \nabla_y T \rangle_z \]
\[ g_{Tzmxy} \langle \nabla_z T \rangle_z \]
\[ gs_{xmx} \langle \nabla_x s \rangle_z \]
\[ gs_{ymxy} \langle \nabla_y s \rangle_z \]
\[ gs_{zmx} \langle \nabla_z s \rangle_z \]
\[ g_{Txgsxmx} \langle (\nabla T \times \nabla s)_x \rangle_z \]
\[ g_{Txgsymxy} \langle (\nabla T \times \nabla s)_y \rangle_z \]
\[ g_{Txgszmx} \langle (\nabla T \times \nabla s)_z \rangle_z \]
\[ g_{Txgsx2mxy} \langle (\nabla T \times \nabla s)_{2x} \rangle_z \]
\[ g_{Txgsy2mxy} \langle (\nabla T \times \nabla s)_{2y} \rangle_z \]
\[ f_{convx} \langle c_p u_x T \rangle_z \]
\[ f_{convy} \langle c_p u_y T \rangle_z \]
\[ f_{convx} \langle c_p u_z T \rangle_z \]
\[ fradjx_{Kprof} F_{rad}^x (x\text{-component of radiative flux, } z\text{-averaged, from Kprof}) \]
\[ frady_{Kprof} F_{rad}^y (y\text{-component of radiative flux, } z\text{-averaged, from Kprof}) \]
\[ fradr_{Kramers} F_{rad} (z\text{-averaged, from Kramers' opacity}) \]
\[ fradr_{constchikappa} F_{rad} (from \chi_{const}) \]
\[ f_turb_{x} \langle gT_x \nabla_x s \rangle_z \]
\[ f_turb_{y} \langle gT_y \nabla_y s \rangle_z \]
\[ f_turb_{rx} \langle gT_{rx} \nabla_{r} s \rangle_z \]
\[ f_turb_{ry} \langle gT_{ry} \nabla_{r} s \rangle_z \]
\[ f_turb_{thx} \langle gT_{th} \nabla_{th} s \rangle_z \]
\[ f_turb_{thy} \langle gT_{th} \nabla_{thy} s \rangle_z \]
\[ dcool_{mx} \text{ surface cooling flux} \]

Module 'magnetic.f90'
\[
\begin{align*}
bxmxy & \langle B_x \rangle_z \\
bymxy & \langle B_y \rangle_z \\
zbmxy & \langle B_z \rangle_z \\
jjxmxy & \langle J_x \rangle_z \\
jjymxy & \langle J_y \rangle_z \\
jjzmxy & \langle J_z \rangle_z \\
abxmxy & \langle A_x \rangle_z \\
abymxy & \langle A_y \rangle_z \\
azmxy & \langle A_z \rangle_z \\
bbxmxy & \langle B_2^x \rangle_z \\
bbymxy & \langle B_2^y \rangle_z \\
bzbmxy & \langle B_2^z \rangle_z \\
bbxbymxy & \langle B_x B_y \rangle_z \\
bbxbzmxy & \langle B_x B_z \rangle_z \\
bbybzmyxy & \langle B_y B_z \rangle_z \\
poynxmxmxy & \langle E \times B \rangle_x \vert_z \\
poynymxy & \langle E \times B \rangle_y \vert_z \\
poynzmxy & \langle E \times B \rangle_z \vert_z \\
etatotalmxy & \langle \eta \rangle_z \\
jbmxy & \langle J \cdot B \rangle_z \\
abmxy & \langle A \cdot B \rangle_z \\
ubmxy & \langle U \cdot B \rangle_z \\
examxy1 & \langle E \times A \rangle_x \vert_z \\
examxy2 & \langle E \times A \rangle_y \vert_z \\
examxy3 & \langle E \times A \rangle_z \vert_z \\
StokesImxy & \langle \epsilon_{B \perp} \rangle_z \\
StokesQmxy & -\langle \epsilon_{B \perp} \cos 2\chi \rangle_z \\
StokesUmxy & -\langle \epsilon_{B \perp} \sin 2\chi \rangle_z \\
StokesQ1mxy & +\langle F \epsilon_{B \perp} \sin 2\chi \rangle_z \\
StokesU1mxy & -\langle F \epsilon_{B \perp} \cos 2\chi \rangle_z \\
beta1mxy & \langle B^2 / (2\mu_0 p) \rangle_z \\
\end{align*}
\]

---

**Module 'density_stratified.f90'**

| \(drhomxy\) | \(\langle \Delta \rho / \rho_0 \rangle_z\) |
| \(drho2mxy\) | \(\langle (\Delta \rho / \rho_0)^2 \rangle_z\) |

---

**Module 'gravity_simple.f90'**

| \(epotmxy\) | \(\langle \rho \Phi_{\text{grav}} \rangle_z\) |
| \(epotuxmxy\) | \(\langle \rho \Phi_{\text{grav}} u_x \rangle_z\) (potential energy flux) |

---

**Module 'hydro_potential.f90'**

| \(uxmxy\) | \(\langle u_x \rangle_z\) |
| \(uymxy\) | \(\langle u_y \rangle_z\) |
| \(uzmxy\) | \(\langle u_z \rangle_z\) |
| \(uxuymxy\) | \(\langle u_x u_y \rangle_z\) |
| \(uxuzmxy\) | \(\langle u_x u_z \rangle_z\) |
| \(uyuzmxy\) | \(\langle u_y u_z \rangle_z\) |
| \(oxmxy\) | \(\langle \omega_x \rangle_z\) |
| \(oymxy\) | \(\langle \omega_y \rangle_z\) |
| \(ozmxy\) | \(\langle \omega_z \rangle_z\) |
K.10 Parameters for ‘zaver.in’

\[ \langle \omega \cdot u \rangle_z \]
\[ \langle pu_x \rangle_z \]
\[ \langle pu_y \rangle_z \]
\[ \langle pu_z \rangle_z \]
\[ \langle u_x^2 \rangle_z \]
\[ \langle u_y^2 \rangle_z \]
\[ \langle u_z^2 \rangle_z \]
\[ \langle pu_x u_y \rangle_z \]
\[ \langle pu_x u_z \rangle_z \]
\[ \langle pu_y u_z \rangle_z \]
\[ \langle \rho u^2_x \rangle_z \]
\[ \langle \rho u^2_y \rangle_z \]
\[ \langle \rho u^2_z \rangle_z \]
\[ f_{k1nxmxy} \langle \frac{1}{2} \rho u^2_x \rangle_z \]
\[ f_{k1nymxy} \langle \frac{1}{2} \rho u^2_y \rangle_z \]

Module ‘magnetic_shearboxJ.f90’

\[ \langle B_x \rangle_z \]
\[ \langle B_y \rangle_z \]
\[ \langle B_z \rangle_z \]
\[ \langle J_x \rangle_z \]
\[ \langle J_y \rangle_z \]
\[ \langle J_z \rangle_z \]
\[ \langle A_x \rangle_z \]
\[ \langle A_y \rangle_z \]
\[ \langle A_z \rangle_z \]
\[ \langle B_x^2 \rangle_z \]
\[ \langle B_y^2 \rangle_z \]
\[ \langle B_z^2 \rangle_z \]
\[ \langle B_x B_y \rangle_z \]
\[ \langle B_x B_z \rangle_z \]
\[ \langle B_y B_z \rangle_z \]
\[ \langle E \times B \rangle_x \]
\[ \langle E \times B \rangle_y \]
\[ \langle E \times B \rangle_z \]
\[ \langle J \cdot B \rangle_z \]
\[ \langle A \cdot B \rangle_z \]
\[ \langle E \times A \rangle_z \]
\[ \langle E \times A \rangle_z \]
\[ \langle E \times A \rangle_z \]
\[ \langle \epsilon_{B1L} \rangle_z \]
\[ \langle F \epsilon_{B1L} \cos 2\chi \rangle_z \]
\[ \langle F \epsilon_{B1L} \sin 2\chi \rangle_z \]
\[ \langle F \epsilon_{B1L} \cos 2\chi \rangle_z \]
\[ \langle F \epsilon_{B1L} \sin 2\chi \rangle_z \]
\[ \langle B^2 / (2\mu_0 p) \rangle_z \]

Module ‘temperature_idealgas.f90’

\[ \langle T \rangle_z \]
\[ \langle T \rangle_z \]
\[ \langle T \rangle_z \]
\[ \langle T \rangle_z \]
\[ \langle T \rangle_z \]
\[ \langle T \rangle_z \]
\[ \langle T \rangle_z \]
\[ \langle T \rangle_z \]
\[ \langle T \rangle_z \]

EmAlA94mxy Emission off AIA 94 channel integrated over z direction
The following tables list all possible boundary condition labels that are documented.

### K.11 Boundary conditions

The following tables list all possible boundary condition labels that are documented.

#### K.11.1 Boundary condition bcx

<table>
<thead>
<tr>
<th>Variable</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>zero value in ghost zones, free value on boundary</td>
</tr>
<tr>
<td>p</td>
<td>periodic</td>
</tr>
<tr>
<td>s</td>
<td>symmetry, ( f_{N+i} = f_{N-i} ); implies ( f'(x_N) = f''(x_0) = 0 )</td>
</tr>
<tr>
<td>sf</td>
<td>symmetry with respect to interface</td>
</tr>
<tr>
<td>ss</td>
<td>symmetry, plus function value given</td>
</tr>
<tr>
<td>s0d</td>
<td>symmetry, function value such that ( df/dx=0 )</td>
</tr>
<tr>
<td>a</td>
<td>antisymmetry, ( f_{N+i} = -f_{N-i} ); implies ( f(x_N) = f''(x_0) = 0 )</td>
</tr>
<tr>
<td>af</td>
<td>antisymmetry with respect to interface</td>
</tr>
<tr>
<td>a2</td>
<td>antisymmetry relative to boundary value, ( f_{N+i} = 2f_N - f_{N-i} ); implies ( f''(x_0) = 0 )</td>
</tr>
<tr>
<td>a2v</td>
<td>set boundary value and antisymmetry relative to it ( f_{N+i} = 2f_N - f_{N-i} ); implies ( f''(x_0) = 0 )</td>
</tr>
<tr>
<td>a2r</td>
<td>sets ( d^2f/dr^2 + 2df/dr - 2f/r^2 = 0 ) This is the replacement of zero second derivative in spherical coordinates, in radial direction.</td>
</tr>
<tr>
<td>cpc</td>
<td>cylindrical perfect conductor implies ( f'' + f'/R = 0 )</td>
</tr>
<tr>
<td>cpp</td>
<td>cylindrical perfect conductor implies ( f'' + f'/R = 0 )</td>
</tr>
<tr>
<td>cpz</td>
<td>cylindrical perfect conductor implies ( f'' + f'/R = 0 )</td>
</tr>
<tr>
<td>spr</td>
<td>spherical perfect conductor implies ( f'' + 2f'/R = 0 ) and ( f(x_N) = 0 )</td>
</tr>
<tr>
<td>v</td>
<td>vanishing third derivative</td>
</tr>
<tr>
<td>cop</td>
<td>copy value of last physical point to all ghost cells</td>
</tr>
<tr>
<td>1s</td>
<td>onesided</td>
</tr>
<tr>
<td>d1s</td>
<td>onesided for 1st/2nd derivative in two first inner points, Dirichlet in boundary point</td>
</tr>
</tbody>
</table>
Boundary conditions

11s onesided for 1st/2nd derivative in two first inner points, Neumann in boundary point

so onesided

CT constant temperature (implemented as condition for entropy s or temperature T)

cl constant conductive flux

FGs \[ F_{\text{conv}} = -\chi_t \rho \frac{\partial s}{\partial T} \]

Fct \[ F_{\text{bot}} = -K \frac{\partial T}{\partial n} - \chi_t \rho \frac{\partial T}{\partial s} \]

Fcm \[ F_{\text{bot}} = -K \frac{\partial T}{\partial n} - \chi_t \rho \frac{\partial T}{\partial s} \]

sT symmetric temperature, \( T_{N-i} = T_{N+i}; \) implies \( T'(x_N) = T''(x_0) = 0 \)

asT select entropy for uniform ghost temperature matching fluctuating boundary value, \( T_{N-i} = T_N = \); implies \( T'(x_N) = T'(x_0) = 0 \)

f “freeze” value, i.e. maintain initial

fg “freeze” value, i.e. maintain initial

1 \( f = 1 \) (for debugging)

set set boundary value to fbcx

der set derivative on boundary to fbcx

sl set slope at the boundary = fbcx

slp set slope at the boundary and in ghost cells = fbcx

shx set shearing boundary proportional to x with slope=fbcx and abscissa=fbcx2

shy set shearing boundary proportional to y with slope=fbcx and abscissa=fbcx2

shz set shearing boundary proportional to z with slope=fbcx and abscissa=fbcx2

dr0 set boundary value [really??]

ovr overshoot boundary condition ie \( (d/dx - 1/\text{dist})f = 0. \)

out allow outflow, but no inflow forces ghost cells and boundary to not point inwards

e1o allow outflow, but no inflow uses the e1 extrapolation scheme

ant stops and prompts for adding documentation

e1 extrapolation [describe]

e2 extrapolation [describe]

e3 extrapolation in log [maintain a power law]

el linear extrapolation from last two active cells

hat top hat jet profile in spherical coordinate.

jet top hat jet profile in cartesian coordinate.

spd sets \( \frac{d}{dr}(rA_\phi) = \frac{fbcx(j)}{dr} \)

sfr stress-free boundary condition for spherical coordinate system.

sr1 Stress-free bc for spherical coordinate system. Implementation with one-sided derivative.

nfr Normal-field bc for spherical coordinate system. Some people call this the “(angry) hedgehog bc”.

nr1 Normal-field bc for spherical coordinate system. Some people call this the “(angry) hedgehog bc”. Implementation with one-sided derivative.

sa2 \( (d/dr)(rB_\phi) = 0 \) imposes boundary condition on 2nd derivative of \( rA_\phi \).

Same applies to \( \theta \) component.

pfc perfect-conductor in spherical coordinate: \( d/dr(A_r) + 2/r = 0. \)

fix set boundary value [really??]

fil set boundary value from a file
Module `boundcond_alt.f90`

<table>
<thead>
<tr>
<th>NAME</th>
<th>DESCRIPTION</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>zero value in ghost zones, free value on boundary</td>
</tr>
<tr>
<td>p</td>
<td>periodic</td>
</tr>
<tr>
<td>s</td>
<td>symmetry, ( f_{N+i} = f_{N-i} ); implies ( f'(x_N) = f''(x_0) = 0 )</td>
</tr>
<tr>
<td>ss</td>
<td>symmetry, plus function value given</td>
</tr>
<tr>
<td>s0d</td>
<td>symmetry, function value such that ( df/dx=0 )</td>
</tr>
<tr>
<td>a</td>
<td>antisymmetry, ( f_{N+i} = -f_{N-i} ); implies ( f(x_N) = f''(x_0) = 0 )</td>
</tr>
<tr>
<td>a2</td>
<td>antisymmetry relative to boundary value, ( f_{N+i} = 2f_N - f_{N-i} ); implies ( f''(x_0) = 0 )</td>
</tr>
<tr>
<td>a2r</td>
<td>sets ( d^2f/dr^2 + 2df/dr - 2f/r^2 = 0 ) This is the replacement of zero second derivative in spherical coordinates, in radial direction.</td>
</tr>
<tr>
<td>cpc</td>
<td>cylindrical perfect conductor implies ( f'' + f'/R = 0 )</td>
</tr>
<tr>
<td>cpp</td>
<td>cylindrical perfect conductor implies ( f'' + f'/R = 0 )</td>
</tr>
<tr>
<td>cpz</td>
<td>cylindrical perfect conductor implies ( f'' + f'/R = 0 )</td>
</tr>
<tr>
<td>spr</td>
<td>spherical perfect conductor implies ( f'' + 2f'/R = 0 ) and ( f(x_N) = 0 )</td>
</tr>
<tr>
<td>v</td>
<td>vanishing third derivative</td>
</tr>
<tr>
<td>cop</td>
<td>copy value of last physical point to all ghost cells</td>
</tr>
<tr>
<td>1s</td>
<td>onesided</td>
</tr>
<tr>
<td>1so</td>
<td>onesided</td>
</tr>
<tr>
<td>cT</td>
<td>constant temperature (implemented as condition for entropy ( s ) or temperature ( T ))</td>
</tr>
<tr>
<td>c1</td>
<td>constant temperature (or maybe rather constant conductive flux??)</td>
</tr>
<tr>
<td>Fgs</td>
<td>( F_{\text{conv}} = -\chi_t \rho T \text{grad}(s) )</td>
</tr>
<tr>
<td>Fct</td>
<td>( F_{\text{bot}} = -K \text{grad}(T) - \chi_t \rho T \text{grad}(s) )</td>
</tr>
<tr>
<td>Fcm</td>
<td>( F_{\text{bot}} = -K \text{grad}(T) - \chi_t \rho T \text{grad}(s) )</td>
</tr>
<tr>
<td>sT</td>
<td>symmetric temperature, ( T_{N-i} = T_{N+i} ); implies ( T'(x_N) = T''(x_0) = 0 )</td>
</tr>
<tr>
<td>asT</td>
<td>select entropy for uniform ghost temperature matching fluctuating boundary value, ( T_{N-i} = T_N ); implies ( T'(x_N) = T'(x_0) = 0 )</td>
</tr>
<tr>
<td>f</td>
<td>“freeze” value, i.e. maintain initial</td>
</tr>
<tr>
<td>fg</td>
<td>“freeze” value, i.e. maintain initial</td>
</tr>
<tr>
<td>1</td>
<td>( f = 1 ) (for debugging)</td>
</tr>
<tr>
<td>set</td>
<td>set boundary value to ( fbcx12 )</td>
</tr>
<tr>
<td>der</td>
<td>set derivative on boundary to ( fbcx12 )</td>
</tr>
<tr>
<td>slo</td>
<td>set slope at the boundary = ( fbcx12 )</td>
</tr>
<tr>
<td>dr0</td>
<td>set boundary value [really??]</td>
</tr>
<tr>
<td>ovr</td>
<td>overshoot boundary condition ie ((d/dx - 1/\text{dist})f = 0).</td>
</tr>
<tr>
<td>out</td>
<td>allow outflow, but no inflow forces ghost cells and boundary to not point inwards</td>
</tr>
<tr>
<td>e1o</td>
<td>allow outflow, but no inflow uses the e1 extrapolation scheme</td>
</tr>
<tr>
<td>ant</td>
<td>stops and prompts for adding documentation</td>
</tr>
<tr>
<td>e1</td>
<td>extrapolation [describe]</td>
</tr>
<tr>
<td>e2</td>
<td>extrapolation [describe]</td>
</tr>
<tr>
<td>e3</td>
<td>extrapolation in log [maintain a power law]</td>
</tr>
</tbody>
</table>
top hat jet profile in spherical coordinate.

*jet* top hat jet profile in cartesian coordinate.

**spd** sets \(d(rA_\alpha)/dr = fbcx12(j)\)

**sfr** stress-free boundary condition for spherical coordinate system.

**nfr** Normal-field bc for spherical coordinate system. Some people call this the “(angry) hedgehog bc”.

\[(d/dr)(rB_\phi) = 0\] imposes boundary condition on 2nd derivative of \(rA_\phi\). Same applies to \(\theta\) component.

**pfc** perfect-conductor in spherical coordinate: \(d/dr(A_r) + 2/r = 0\).

**fix** set boundary value [really??]

**fil** set boundary value from a file

**g** set to given value(s) or function

**nil** do nothing; assume that everything is set

**ioc** inlet/outlet on western/eastern hemisphere in cylindrical coordinates do nothing; assume that everything is set

\(s\) implies \(f'(y_N) = f'''(y_0) = 0\)

---

### K.11.2 Boundary condition bcy

<table>
<thead>
<tr>
<th>Variable</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>sds</strong></td>
<td>symmetric-derivative-set</td>
</tr>
<tr>
<td>0</td>
<td>zero value in ghost zones, free value on boundary</td>
</tr>
<tr>
<td><strong>p</strong></td>
<td>periodic</td>
</tr>
<tr>
<td><strong>pp</strong></td>
<td>periodic across the pole</td>
</tr>
<tr>
<td><strong>yy</strong></td>
<td>Yin-Yang grid</td>
</tr>
<tr>
<td><strong>ap</strong></td>
<td>anti-periodic across the pole</td>
</tr>
<tr>
<td><strong>s</strong></td>
<td>symmetry symmetry, (f_{N+i} = f_{N-i};)</td>
</tr>
<tr>
<td><strong>sf</strong></td>
<td>symmetry with respect to interface</td>
</tr>
<tr>
<td><strong>ss</strong></td>
<td>symmetry, plus function value given</td>
</tr>
<tr>
<td><strong>sds</strong></td>
<td>symmetric-derivative-set</td>
</tr>
<tr>
<td><strong>cds</strong></td>
<td>complex symmetric-derivative-set</td>
</tr>
<tr>
<td><strong>s0d</strong></td>
<td>symmetry, function value such that (df/dy=0)</td>
</tr>
<tr>
<td><strong>a</strong></td>
<td>antisymmetry</td>
</tr>
<tr>
<td><strong>af</strong></td>
<td>antisymmetry with respect to interface</td>
</tr>
<tr>
<td><strong>a2</strong></td>
<td>antisymmetry relative to boundary value</td>
</tr>
<tr>
<td><strong>v</strong></td>
<td>vanishing third derivative</td>
</tr>
<tr>
<td><strong>v3</strong></td>
<td>vanishing third derivative</td>
</tr>
<tr>
<td><strong>out</strong></td>
<td>allow outflow, but no inflow forces ghost cells and boundary to not point inwards</td>
</tr>
<tr>
<td><strong>1s</strong></td>
<td>onesided</td>
</tr>
<tr>
<td><strong>d1s</strong></td>
<td>onesided for 1st and 2nd derivative in two first inner points, Dirichlet in boundary point</td>
</tr>
<tr>
<td><strong>n1s</strong></td>
<td>onesided for 1st and 2nd derivative in two first inner points, Neumann in boundary point</td>
</tr>
<tr>
<td><strong>cT</strong></td>
<td>constant temp.</td>
</tr>
<tr>
<td><strong>sT</strong></td>
<td>symmetric temp.</td>
</tr>
<tr>
<td><strong>asT</strong></td>
<td>select entropy for uniform ghost temperature matching fluctuating boundary value, (T_{N-i} = T_N =;) implies (T'(x_N) = T'(x_0) = 0)</td>
</tr>
</tbody>
</table>
f  freeze value
s+f  freeze value
fg  “freeze” value, i.e. maintain initial
fBs  frozen-in B-field (s)
fB  frozen-in B-field (a2)
l  f=1 (for debugging)
set  set boundary value
sse  symmetry, set boundary value
sep  set boundary value
e1  extrapolation
e2  extrapolation
e3  extrapolation in log [maintain a power law]
der  set derivative on the boundary
cop  outflow: copy value of last physical point to all ghost cells
c+k  no-inflow: copy value of last physical point to all ghost cells, but suppressing any inflow
sfr  stress-free boundary condition for spherical coordinate system.
nfr  Normal-field bc for spherical coordinate system. Some people call this the “(angry) hedgehog bc”.
spt  spherical perfect conducting boundary condition along \( \theta \) boundary \( f'' + \cot \theta f' = 0 \) and \( f(x_N) = 0 \)
pfc  perfect conducting boundary condition along \( \theta \) boundary
nil’,’  do nothing; assume that everything is set
sep  set boundary value
crk  no-inflow: copy value of last physical point to all ghost cells, but suppressing any inflow

Module ‘boundcond_alt.f90’

sds  symmetric-derivative-set
0  zero value in ghost zones, free value on boundary
p  periodic
pp  periodic across the pole
ap  anti-periodic across the pole
s  symmetry symmetry, \( f_{N+i} = f_{N-i} \);
ss  symmetry, plus function value given
sds  symmetric-derivative-set
cds  complex symmetric-derivative-set
s0d  symmetry, function value such that \( df/dy=0 \)
a  antisymmetry
a2  antisymmetry relative to boundary value
v  vanishing third derivative
v3  vanishing third derivative
out  allow outflow, but no inflow forces ghost cells and boundary to not point inwards
ls  onesided
cT  constant temp.
sT  symmetric temp.
asT  select entropy for uniform ghost temperature matching fluctuating boundary value, \( T_{N-i} = T_N =; \) implies \( T'(x_N) = T'(x_0) = 0 \)
f  freeze value
s+f
freeze value

fg
“freeze” value, i.e. maintain initial

1
f=1 (for debugging)

set
set boundary value

sse
symmetry, set boundary value

sep
set boundary value

e1
extrapolation

e2
extrapolation

e3
extrapolation in log [maintain a power law]

der
set derivative on the boundary

cop
outflow: copy value of last physical point to all ghost cells

c+k
no-inflow: copy value of last physical point to all ghost cells, but suppressing any inflow

sfr
stress-free boundary condition for spherical coordinate system.
nfr
Normal-field bc for spherical coordinate system. Some people call this the “(angry) hedgehog bc”.

spt
spherical perfect conducting boundary condition along $\theta$ boundary $f'' + \cot \theta f' = 0$ and $f(x_N) = 0$

pfc
perfect conducting boundary condition along $\theta$ boundary

nil',
do nothing; assume that everything is set

sep
set boundary value

---

K.11.3 Boundary condition bcz

<table>
<thead>
<tr>
<th>Variable</th>
<th>Meaning</th>
</tr>
</thead>
<tbody>
<tr>
<td>0</td>
<td>zero value in ghost zones, free value on boundary</td>
</tr>
<tr>
<td>p</td>
<td>periodic</td>
</tr>
<tr>
<td>yy</td>
<td>Yin-Yang grid</td>
</tr>
<tr>
<td>s</td>
<td>symmetry</td>
</tr>
<tr>
<td>sf</td>
<td>symmetry with respect to interface</td>
</tr>
<tr>
<td>s0d</td>
<td>symmetry, function value such that df/dz=0</td>
</tr>
<tr>
<td>0ds</td>
<td>symmetry, function value such that df/dz=0</td>
</tr>
<tr>
<td>a</td>
<td>antisymmetry</td>
</tr>
<tr>
<td>a2</td>
<td>antisymmetry relative to boundary value</td>
</tr>
<tr>
<td>a2v</td>
<td>set boundary value and antisymmetry relative to it</td>
</tr>
<tr>
<td>af</td>
<td>antisymmetry with respect to interface</td>
</tr>
<tr>
<td>a0d</td>
<td>antisymmetry with zero derivative</td>
</tr>
<tr>
<td>v</td>
<td>vanishing third derivative</td>
</tr>
<tr>
<td>v3</td>
<td>vanishing third derivative</td>
</tr>
<tr>
<td>1s</td>
<td>one-sided</td>
</tr>
<tr>
<td>d1s</td>
<td>onesided for 1st and 2nd derivative in two first inner points, Dirichlet in boundary point</td>
</tr>
<tr>
<td>n1s</td>
<td>onesided for 1st and 2nd derivative in two first inner points, Neumann in boundary point</td>
</tr>
<tr>
<td>a1s</td>
<td>special for perfect conductor with const alpha and etaT when A considered as B; one-sided for 1st and 2nd derivative in two first inner points</td>
</tr>
<tr>
<td>fg</td>
<td>“freeze” value, i.e. maintain initial</td>
</tr>
<tr>
<td>Symbol</td>
<td>Description</td>
</tr>
<tr>
<td>--------</td>
<td>-------------</td>
</tr>
<tr>
<td>$c_1$</td>
<td>complex</td>
</tr>
<tr>
<td>$c_{1s}$</td>
<td>complex</td>
</tr>
<tr>
<td>$F_{gs}$</td>
<td>$F_{\text{conv}} = -\chi_{t} \rho \tau_{T} \nabla s$</td>
</tr>
<tr>
<td>$F_{ct}$</td>
<td>$F_{\text{bot}} = -K \nabla T - \chi_{t} \rho \tau_{T} \nabla s$</td>
</tr>
<tr>
<td>$c_3$</td>
<td>constant flux at the bottom with a variable $h_{\text{cond}}$</td>
</tr>
<tr>
<td>$p_{fe}$</td>
<td>potential field extrapolation</td>
</tr>
<tr>
<td>$p_{1D}$</td>
<td>potential field extrapolation in 1D</td>
</tr>
<tr>
<td>$p_{wd}$</td>
<td>a variant of 'pot' for nprocx=1</td>
</tr>
<tr>
<td>$h_{ds}$</td>
<td>hydrostatic equilibrium with a high-frequency filter</td>
</tr>
<tr>
<td>$cT$</td>
<td>constant temp.</td>
</tr>
<tr>
<td>$cT_1$</td>
<td>constant temp.</td>
</tr>
<tr>
<td>$cT_2$</td>
<td>constant temp. (keep lnrho)</td>
</tr>
<tr>
<td>$cT_3$</td>
<td>constant temp. (keep lnrho)</td>
</tr>
<tr>
<td>$h_{s}$</td>
<td>hydrostatic equilibrium</td>
</tr>
<tr>
<td>$h_{se}$</td>
<td>hydrostatic extrapolation rho or lnrho is extrapolated linearly and the temperature is calculated in hydrostatic equilibrium.</td>
</tr>
<tr>
<td>$c_{p}$</td>
<td>constant pressure</td>
</tr>
<tr>
<td>$sT$</td>
<td>symmetric temp.</td>
</tr>
<tr>
<td>$c_{tz}$</td>
<td>for interstellar runs copy $T$</td>
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<tr>
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**Boundary conditions**

*in* allow inflow, but no outflow forces ghost cells and boundary to not point outwards

*out* allow outflow, but no inflow forces ghost cells and boundary to not point inwards

*in0* allow inflow, but no outflow forces ghost cells and boundary to not point outwards relaxes to vanishing 1st derivative at boundary

*ou0* allow outflow, but no inflow forces ghost cells and boundary to not point inwards relaxes to vanishing 1st derivative at boundary

*ind* allow inflow, but no outflow forces ghost cells and boundary to not point outwards creates inwards pointing or zero 1st derivative at boundary

*oud* allow outflow, but no inflow forces ghost cells and boundary to not point inwards creates outwards pointing or zero 1st derivative at boundary

*ubs* copy boundary outflow,

*win* forces massflux given as $\Sigma \rho_i (u_i + u_0) = fbcz1/2(\rho)$

*cop* copy value of last physical point to all ghost cells

*nil* do nothing; assume that everything is set

---

**Module 'boundcond_alt.f90'**

- **cfb** radial centrifugal balance
- **fBs** frozen-in B-field (s)
- **fB** frozen-in B-field (a2)
- **0** zero value in ghost zones, free value on boundary
- **p** periodic
- **s** symmetry
- **sf** symmetry with respect to interface
- **s0d** symmetry, function value such that $df/dz=0$
- **0ds** symmetry, function value such that $df/dz=0$
- **a** antisymmetry
- **a2** antisymmetry relative to boundary value
- **af** antisymmetry with respect to interface
- **a0d** antisymmetry with zero derivative
- **v** vanishing third derivative
- **v3** vanishing third derivative
- **1s** one-sided
- **fg** “freeze” value, i.e. maintain initial
- **c1** complex
- **Fgs** $F_{\text{conv}} = - \chi_t \rho T \cdot \text{grad}(s)$
- **Fct** $F_{\text{bot}} = - K \cdot \text{grad}(T) - \chi_t \rho T \cdot \text{grad}(s)$
- **c3** constant flux at the bottom with a variable hcond
- **pfe** potential field extrapolation
- **p1D** potential field extrapolation in 1D
- **pot** potential magnetic field
- **pwd** a variant of ‘pot’ for nprocx=1
- **hds** hydrostatic equilibrium with a high-frequency filter
- **cT** constant temp.
- **cT2** constant temp. (keep lnrho)
- **cT3** constant temp. (keep lnrho)
- **hs** hydrostatic equilibrium
- **hse** hydrostatic extrapolation rho or lnrho is extrapolated linearly and the temperature is calculated in hydrostatic equilibrium.
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<tr>
<th>Symbol</th>
<th>Description</th>
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<td>allow inflow, but no outflow forces ghost cells and boundary to not point outwards</td>
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<td>allow outflow, but no inflow forces ghost cells and boundary to not point inwards</td>
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<tr>
<td>$in0$</td>
<td>allow inflow, but no outflow forces ghost cells and boundary to not point outwards relaxes to vanishing 1st derivative at boundary</td>
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<td>$ou0$</td>
<td>allow outflow, but no inflow forces ghost cells and boundary to not point inwards relaxes to vanishing 1st derivative at boundary</td>
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<tr>
<td>$ind$</td>
<td>allow outflow, but no outflow forces ghost cells and boundary to not point outwards creates inwards pointing or zero 1st derivative at boundary</td>
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<td>$oud$</td>
<td>allow outflow, but no inflow forces ghost cells and boundary to not point inwards creates outwards pointing or zero 1st derivative at boundary</td>
</tr>
<tr>
<td>$ubs$</td>
<td>copy boundary outflow,</td>
</tr>
<tr>
<td>$win$</td>
<td>forces massflux given as $\Sigma \rho_i (u_i + u_0) = fbcz1/2(\rho)$</td>
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<td>$cop$</td>
<td>copy value of last physical point to all ghost cells</td>
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<tr>
<td>$nil$</td>
<td>do nothing; assume that everything is set</td>
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## K.12 Initial condition parameter dependence

The following tables list which parameters from each Namelist are required (●), optional (○) or irrelevant (blank). The distinction is made between required and optional where by a parameter requires a setting if the default value would give an invalid or degenerate case for the initial condition.

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initss

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