

Pencil Code: Quick Start guide

Illa R. Losada, Michiel Lambrechts, Elizabeth Cole, Philippe Bourdin

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1 Download the Pencil Code

The Pencil Code is an open source code written mainly in Fortran and available under GPL. General information can be found at our official homepage:

`http://pencil-code.nordita.org/`.

The latest version of the code can be downloaded with `svn`. In the directory where you want to put the code, type:

```
svn checkout http://pencil-code.googlecode.com/svn/trunk/ pencil-code
```

The downloaded `pencil-code` directory contains several sub-directories

1. `doc`: you may build the latest manual as PDF by issuing the command `make` inside this directory
2. `samples`: contains many sample problems
3. `config`: has all the configuration files
4. `src`: the actual source code
5. `bin` and `lib`: supplemental scripts
6. `idl`, `python`, `julia`, etc.: data processing for diverse languages

2 Configure the shell

For a quick start, you need to load some environment variable into your shell. First, you enter to the freshly downloaded directory:

```
cd pencil-code
```

Depending on which shell you use, you can do that by a simple command:

```
. source.sh
```

that will work for `bash` and all `sh`-compatible shells, while this command:

```
source source.csh
```

is for `tcsh` and any `csh`-compatible shell.

3 Fortran

A Fortran and a C compiler are needed to compile the code. Both compilers should belong to the same distribution package and version (e.g. GNU GCC 4.8.3, 64 bit Linux).

3.1 Fortran on a mac machine

For Mac, you first need to install Xcode from the AppleDeveloper site <http://developer.apple.com/>. This requires you to first register as a member. An easy to install gfortran can be found at <http://gcc.gnu.org/wiki/GFortranBinaries>. Just download it and it comes with an installer. It installs in the directory `/usr/local/gfortran` with a symbolic link in `/usr/local/bin/gfortran`. It might be necessary to add the following line in the `.cshrc`-file in the home folder:

```
setenv PATH /usr/local/bin:\$PATH
```

4 Try a sample

Go to a folder that contains one of the many available samples, e.g.:

```
cd samples/ld-tests/jeans-x
```

You may also start with a fresh directory and copy over the files from one of the samples.

4.1 Setting up...

One command sets up all needed symbolic links to the original Pencil Code directory:

```
pc_setupsrc
```

4.2 Makefile

Two basic configuration files define a simulation setup: `src/Makefile.local` contains a list of modules that are being used, and `src/cparam.local` defines the grid size and the number of processors to be used. Take a quick look at these files...

4.2.1 Single-processor

An example using the module for only one processor would look like:

```
MPICOMM=nompicomm
```

For most modules there is also a `no`-variant which switches that functionality off.

In `src/cparam.local` the number of processors needs to be set to 1 accordingly:

```
integer, parameter :: ncpus=1,nprocx=1,nprocy=1,nprocz=ncpus/(nprocx*nprocy)
integer, parameter :: nxgrid=128,nygrid=1,nzgrid=128
```

4.2.2 Multi-processor

If you like to use MPI for multi-processors simulations, be sure that you have a MPI library installed and change `src/Makefile.local` to use MPI:

```
MPICOMM=mpicomm
```

Change the `ncpus` setting in `src/cparam.local`. Think about how you want to distribute the volume on the processors — usually, you should have 128 grid points in the x-direction to take advantage of the SIMD processor unit. For compilation, you have to use a configuration file that includes the `_MPI` suffix, see below.

4.3 Compiling...

In order to compile the code, you can use a pre-defined configuration file corresponding to your compiler package. E.g. the default compilers are `gfortran` together with `gcc` and the code is being built with default options by issuing the command:

```
pc_build
```

4.3.1 Using a different compiler (optional)

If you prefer to use a different compiler package (e.g. using `ifort` or `MPI`), you may try:

```
pc_build -f compilers/Intel
pc_build -f compilers/GNU-GCC_MPI
```

More pre-defined configurations are found in the directory `pencil-code/config/compilers/*.conf`.

4.3.2 Changing compiler options (optional)

Of course you can also create a configuration file in any subdirectory of `pencil-code/config/hosts/`. By default, `pc_build` looks for a config file that is based on your `host-ID`, which you may see with the command:

```
pc_build -i
```

You may add your modified configuration with the filename `host-ID.conf`, where you can change compiler options according to the Pencil Code manual.

4.4 Running...

The initial conditions are set in `start.in` and the parameters for the main simulation run can be found in `run.in`. In `print.in` you can choose which physical quantities are written to the file `data/time_series.dat`.

It is time to run the code with — be sure you have an empty `data` directory:

```
mkdir data
pc_run
```

Welcome to the world of Pencil Code! Visualizing the output can be done with `IDL` or `Python`, see below.

4.5 IDL visualization (optional)

4.5.1 GUI-based visualization

The most simple approach to visualize a cartesian grid setup is to run the Pencil Code GUI and to select the files and physical quantities you want to see:

```
IDL> .r pc_gui
```

If you miss some physical quantities, you might want to extend the two IDL routines `pc_get_quantity` and `pc_check_quantities`. Anything implemented there will be available in the GUI, too.

4.5.2 Command-line based and scripting

Several `idl`-procedures have been written (see in `pencil-code/idl`) to facilitate inspecting the data that can be found in raw format in `jeans-x/data` directory. For example, let us inspect the time series data

```
IDL> pc_read_ts, obj=ts
```

The structure `ts` contains several variables that can be inspected by

```
IDL> print, tag_names(ts)
IT T UMAX RHOMAX
```

The diagnostic `UMAX`, the maximal velocity, is available since it was set in `jeans-x/print.in`. Please check manual for more information.

We can now plot the evolution of the maximal velocity after the initial perturbation we inserted in `start.in`:

```
IDL> plot, ts.t, alog(ts.umax)
```

The complete state of the simulation is saved as snapshots in `jeans-x/data/proc0/VAR*` every `dsnap` time units, as defined in `jeans-x/run.in`. These states are loaded with, for example:

```
IDL> pc_read_var, obj=ff, ivar=1, /trimall
```

Similarly `tag_names` will provide us with the available variables:

```
IDL> print, tag_names(ff)
T X Y Z DX DY DZ UU LNRHO POTSELF
```

The logarithm of the density can be inspected by using a GUI:

```
IDL> cslice, ff.lnrho
```

Of course, for scripting one might use any quantity from the `ff` structure, like calculating the average density:

```
IDL> print, mean(exp(ff.lnrho))
```

Also one should check the documentation inside:

<code>pencil-code/idl/read/pc_read_var_raw.pro</code>	efficient reading of raw data
<code>pencil-code/idl/read/pc_read_subvol_raw.pro</code>	reading of sub-volumes
<code>pencil-code/idl/pc_get_quantity.pro</code>	compute physical quantities out of raw data

in order to read data efficiently and compute quantities in physical units.

4.6 Python visualization (optional)

4.6.1 Python module requirements

For this example we use the modules: `numpy` and `matplotlib`.

4.6.2 Using the 'pencil' module

After executing the `sourceme.sh` script (see above), you should be able to import the `pencil` module:

```
import pencil as pc
```

Some useful functions:

<code>pc.read_ts</code>	read <code>time_series.dat</code> file. Parameters are added as members of the class
<code>pc.read_slices</code>	read 2D slice files and return two arrays: (<code>nslices</code> , <code>vsize</code> , <code>hsize</code>) and (<code>time</code>)
<code>pc.animate_interactive</code>	assemble a 2D animation from a 3D array