MESA tutorial: Session 2 MESA output and Intermediate mass stars

Hand in your answers to all the numbered questions (i.e. 3.1 a, b, c etc.), they will make up your grade for the MESA practicum part of the course.

1 Understanding MESA output

The following is a summary of the official MESA documentation. For a more comprehensive overview of available output options, please refer directly to the documentation. You might have noticed that MESA stores its output as plain text files in the LOGS directory. There are two main types of output:

- history.data: logs each model's evolution in one line per model. The first line contains column indices, the second column names, and the remainder the data. Note: in case of a restart, older models are not removed—new data is appended. As a result the model_numbers are not guaranteed to be monotonically increasing! Existing tools (like MESA Reader below) automatically take care of this, but if you write your own parser, be aware of this.
- Each **profile**.data:** file contains a snapshot of a single model's structure. Each profile includes both global properties of the star, such as its age, and a large set of properties for each point in the model of the star given one line per point. In each case, the lines of data are preceded by a line with column numbers and a line with column names.

Lastly, you might also notice a **profiles.index** file. This maps profile filenames to model numbers. MESA saves profiles only at selected steps. Each line lists a model number, its priority (1 or 2), and the corresponding profile number.

You can customize the MESA output through the history_columns.list and profile_columns.list files. In order to customize the output, just copy these files to your work directory, and uncomment the variables you want to include in the output. You can find default variations of these files in the \$MESA_DIR/star/defaults/ directory. We will control some of the output of your MESA models in section 4.

Using MESA Reader MESA Reader is a Python tool for reading and plotting history and profile data; see MESA Reader for details. Although MESA outputs text files, restarts can duplicate model numbers and large headers make manual parsing tedious—MESA Reader handles all of this for you!

We want to install MESA Reader, but in such a way that we can always activate/access it, regardless of which computer we are using in the Faculty. Turns out that python virtual environments are exactly what we are looking for!

2 Jupyter Hub and virtual environments

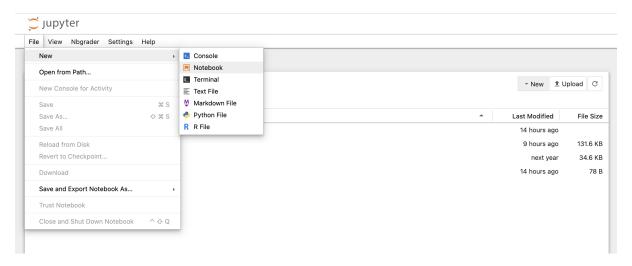
Throughout this course, we will use Jupyter Notebooks to analyse our results. If you have correctly copied your data to your home directory as explained in the previous session, you

can access your this data via the RU JupyterHub: https://jupyterhub.science.ru.nl/ and logging in with your science account.

You can create a Jupyter Notebook by clicking File>New>Notebook. It ask you which kernel to use: Python3 ipykernel will work out of the box for general analysis, but it does not have MESA Reader installed. For this purpose we will create our own virtual environment (venv) in which we install MESA Reader and other useful packages.

Creating a virtual environment

• When you are logged in, open a Jupyter Hub terminal



• Inside your JupyterHub terminal, **Create a virtual environment** by running the following:

```
mkdir -p ~/venvs  # make dir to store venvs

cd ~/venvs  # go to that directory

python -m venv myenv # create a venv called myenv

source ~/venvs/myenv/bin/activate # activate venv
```

You will notice that your terminal prompt changes adding (myenv) to the start, indicating that you are now working inside the virtual environment. To deactivate the virtual environment, simply run: deactivate.

We need to ensure the package manager 'pip install' is up to date and available in our venv:

```
python -m ensurepip --upgrade
python -m pip install --upgrade pip setuptools wheel ipykernel
```

• While your venv is active, you can install Mesa Reader by running:

```
python -m pip install mesa-reader
```

You can install all kinds of useful python packages like this in your venv using python -m pip install chage_name.

¹Note that normally you often use the slightly simpler pip install package_name>, but this can sometimes lead to some issues with the environment not being recognized due to PATH issues.

If for some reason you encounter trouble with the pip install, you can alternatively directly git clone (download) the code from the project's Github repository somewhere in your home directory (e.g. ~/codes)² by running:

```
git clone https://github.com/wmwolf/py_mesa_reader.git ~/codes/
py_mesa_reader
```

Then run python setup.py install inside the cloned directory.

Next, we need to register our venv as a Jupyter kernel:

```
pip install ipykernel # makes venv able to run noteboooks
python -m ipykernel install --user --name=myenv --display-name "
Python (myenv)"
```

this should give you a message like: "Installed kernelspec myenv in ..."

- Open a new notebook, you should now see your newly created Python (myenv) as an option in the kernel dropdown menu. Note that you can always switch kernels by clicking on the kernel name in the top right of your notebook.
- You should now be able to import MESA Reader without any errors, type:

```
import mesa_reader as mr
```

and run your cell (Shift + Enter)

Note some other useful imports for calculating and plotting purposes could be:

```
import numpy as np
import matplotlib.pyplot as plt
import astropy.units as u
import astropy.constants as const
```

All of these packages can be installed in your venv using pip install <package_name> in your jupyter terminal as explained above. We leave it to you to figure out which pip install commands do you need to run to get all the aforementioned packages.

3 Overshooting in an intermediate-mass star

We will now investigate the effect of overshooting. In order to do this, we evolve an intermediatemass star for different levels of convective overshooting.

- 3.1 **Download and set up the work folder.** Instead of starting from the beginning and copying the model work directory like last time to edit, a MESA model has largely been constructed for you already.
 - Download the tar file from Brightspace and unpack it. This can be done using the archive manager by extracting the contents to your desired location on the scratch disc, or using the terminal by typing tar -xvf session2.tar in the directory where you downloaded the tar file and then moving the directory to your desired location on the scratch disc using a cp -R command. This file contains a MESA work folder in which we will work.

² ~ is an alias for your home directory.

- Once you open the inlist_project file, you will find a number of controls available for the model. First, choose a mass for the star between 2.5 M_☉ and 10 M_☉, and modify the inlist accordingly. Note that the stopping condition allows the star to evolve up to the end of core helium burning. You may further notice several lines commented out concerning Convective overshooting. We will run several models for different values of the overshooting parameter.
- First, compile (./mk) and run (./rn) the model without any overshooting. Use the various PGSTAR windows to follow and understand the evolution of this star. One of the PGSTAR windows is a Kippenhahn diagram, which shows information about the structure of the star as it evolves. Also try to understand this plot.
- After the model finishes, create a new copy of the work folder and rename the **old** folder to an appropriate name (e.g. identify it by the mass and overshoot value used, like M3_0ov0 for mass 3 and overshoot 0). In the new work folder, uncomment the lines relating to convective overshooting in your inlist and change the overshooting parameter overshoot_f(1) to 0.25. Look for the meaning of this and other overshooting parameters in the file \$MESA_DIR/star/defaults/controls.defaults or by checking the documentation online. ³ Then compile and run the code again. Repeat this process for an overshooting parameter of 0.5.

Pro Tip

It is a good idea to organise your MESA work folders in a logical directory structure, in order to not make your home directory a mess. Just make sure you change the path in your Python script, such that it points to the right file.

3.2 **Inspecting the results.** We can now analyse the data in the 3 history files. For this import MESA Reader into your python notebook, as explained above.

To read in the data, you need to move the history.data files from your scratch directory to somewhere in your home foler, for example inside a folder called data and there inside a folder called after its run (e.g. M3.0ov0).

For this example, you use the following line in your Python script:

```
f0_hist_data = mr.MesaData('data/M3.0ov0/LOGS/history.data')
```

To read a specific columns you want from the fo_hist_data, use fo_hist_data.X, where X is the name of one of the columns inside history.data. The names of the columns can be found inside history.data, or you can print the available column names by typing:

```
print(f0_hist_data.bulk_names)
```

- (a) Make an HR diagram containing the 3 models. What changes do you see in the main-sequence evolution? What changes appear in the evolution *after* the main sequence? Can you explain these changes?
 - Make a plot of ρ_c vs T_c . How do the evolution tracks in this diagram change for different levels of overshooting?

³Note that the documentation online is for the latest version of MESA, which may differ slightly from the version you are using. The bottom right of the documentation page shows which version of the docs you are viewing, but note this only dates back to version r15140, which is when MESA was migrated to GitHub. \$MESA_DIR/star/defaults/controls.defaults will always show the information for the correct version for your installation.

- (b) Construct a plot of the central helium abundance vs age for all 3 models and explain your findings.
- (c) By what fraction is the main sequence lifetime increased for an overshooting parameter of 0.25 compared to the model without overshooting? By what fraction does the *helium burning* lifetime change? Compare your findings with your neighbours who, hopefully, have chosen a star of different mass.

Pro Tip

As explained in section 2.2 of the first tutorial, it is a very good idea to copy your work folder from the /scratch directory to your home directory, after the MESA run has finished. This allows you to analyse your results using MESA Reader from any computer in the Faculty, not just the computer you ran your MESA models on!

4 Make your own Kippenhahn diagram

Kippenhahn diagrams (KHDs) show a star's internal structure over time and are useful for your final MESA project reports. Rather than taking screenshots from PGSTAR, you can use the Python script mkipp⁴ to generate clearer, customizable plots. Here we will enable the necessary MESA output columns, help you set up mkipp, and get started with the scripts.

4.1 Setting MESA output

We will use the mkipp. tool from https://github.com/orlox/mkipp. As you can read on the github page, to use mkipp you need to have the following MESA output available:

IMPORTANT!! your history_columns.list needs to have mixing_regions and mix_relr_regions specified for MESA to output the necessary data of the mixing regions in terms of mass and radius respectively. Also, newer versions of MESA include significantly less output in the profile files, so values such as eps_nuc need to be added to profile_columns.list as well.

- Pick one star that you like, and make sure that all these variables are uncommented. (i.e., remove the '!' in front of the variable) for both history_columns.list and profile_columns.list files of that star's work folder.
- When you uncomment mix_relr_regions, it will most likely say <integer> after it. You should change this to 10, so that it has the same value as for mixing_regions.

For MESA to use the columns you have specified in these 2 files instead of the default set of columns, we have to make sure MESA knows that it needs to read these 2 files. To do this, we have to open inlist_project and include the following under &star_job:

```
! to specify which output columns we want in history.data and profile.data
history_columns_file = 'history_columns.list'
profile_columns_file = 'profile_columns.list'
```

Now recompile and rerun your model.

• Next, actually cloning the mkipp repository:

⁴https://github.com/orlox/mkipp

```
cd ~/codes # or wherever you want to store the code git clone git@github.com:orlox/mkipp.git
```

• Unfortunately, mkipp is not available via pip install, so we have to make sure Python can find it. You can do this by adding the path to mkipp to your PYTHONPATH environment variable. To make your Jupyter notebook recognize and import a script or module from , you need to add that path to Python's search path at runtime.

```
import sys
import os
sys.path.append(os.path.expanduser("~/codes/mkipp"))
```

Your notebook will now look for python functions in the /codes/mkipp directory as well, and so you can import the three python modules that live in there:

```
import mkipp
import kipp_data
import mesa_data
```

a) To run a Kippenhahn diagram, you need to define the logs_dir argument of the KippArgs function as a list containing the path to your MESA output directory. For example, you can create a Kippenhahn diagram using the following command:

```
mkipp.kipp_plot(mkipp.Kipp_Args(logs_dirs = ['data/M3.0ov0/LOG']))
```

Describe what you see in your Kippenhahn plot, please explain all the colors and hatching (hint: look inside mkipp.py).

b) Play around with the various options of KippArgs to customize your KHD. Have a look at the mkipp/example.py script for inspiration. Create at least one variation of the KHD that you like, and include it in your hand in.