

Manual for v. 8.0

1 Installation and usage

GENEC_toolBox is a set of python routines designed to exploit the files generated by the Geneva evolution code and the Syclist tool for clusters and isochrones. It processes the files and provides the possibility to plot any variable as a function of any other one. It can be downloaded on our group's webpage: http://obswww.unige.ch/Recherche/evol/Tools or alternatively on GitHub: https://github.com/GESEG/GENEC_toolBox

It requires

- python 2.7,
- matplotlib >= 1.2,
- numpy >= 1.7,
- and scipy >= 0.12.

1.1 Installation

You just have to save the .py file and the data directory in your favorite location.

To use it, open a terminal.

- log into a ipython console: ipython --pylab=auto (--pylab=osx for Mac OSX users);
- declare the directory where the routine sits: sys.path.append('path_to_directory');
- import the set of routines:
 - 1. either with the command from GENEC_toolBox import ★,
 - 2. or import GENEC_toolBox as GtB (or whatever short name you wish);
- use the commands either directly (if loaded with method 1) or with GtB. command.

Note that depending on your system, you might need to import sys first.

1.2 First run

The first time you run it, you will be asked to define the location of the data directory. Leave blank if you installed the package. Enter the path to the data directory in case you use the standalone. After that, you'll be asked to define the default figure directory. Leave blank to use the current location.

Your settings will be saved in a hidden config file ~/.GENEC_toolBox.ini.

1.3 Quick outlook

The philosophy of GENEC_toolBox is to load a file, define the x-axis, and then plot the desired y variable.

Evolution files (.wg, .dat, grids format) are loaded with

- loadE()
- loadEfromList()
- loadEFromDir().

Structure files (*.v*) are loaded with

- loadS()
- loadSFromList()
- loadSFromDir().

Cluster / isochrones files generated by Syclist are loaded with

- loadC()
- loadCFromList()
- loadCFromDir().

The variables available for plotting can be printed with VarEvol(), VarStruc(), or VarCluster(). New variables can be determined by using the command Get_Var() and Set_Var(), see example in the comments of the latter. After Set_Var(), the variable is available for plotting.

The x axis variable is set by defX(). By default, it is t [Myr] ('t6') in EVOL mode, M_r/M ('Mfrac') in STRUC mode, and $T_{\rm eff}$ ('Teff') in CLUSTER mode. The plot command is Plot('y') for normal plots, and Plot_colour('y') for plots with lines or points colour-coded by the value of another variable. In CLUSTER mode, histograms can be plotted with the command Histo().

Several pre-programmed plots exist, as:

- HRD_plot(): HR diagram in L vs $T_{\rm eff}$. With the optional parameter corr=True: L vs $T_{\rm eff,corr}$ ($T_{\rm eff}$ corrected for the wind thickness). With optional parameter spectro=True: sHRD.
- CMD(): colour-magnitude diagram, with a large choice of colours
- rhoT(): T_c vs ρ_c in EVOL mode, and T_r vs ρ_r diagram in struc mode
- YTeff(): T_{eff} vs $X(^{4}\text{He})_{c}$
- gTeff(): $T_{\rm eff}$ vs $g_{\rm surf}$
- Abund(): plot of the abundances. In EVOL mode, with input parameter 's' for surface, 'c' for centre. In STRUC mode, abundance profiles with input parameter 'p'.
- NCNO(): N/C vs N/O diagram.
- eps(): energy generation zones for H, He, C, and neutrinos (STRUC mode only).
- Coeff(): plot of the diffusion coefficients.
- Nablas(): plot of the three Nablas $(\nabla_{ad}, \nabla_{rad}, \nabla_{\mu})$ in struc mode.
- j_profiles(): plots the profile of the specific angular momentum j_r as well as $j_{\text{Schwarzschild}}$, j_{Kerr} , and $j_{\text{Kerr,max}}$.
- Kippen(): Kippenhahn diagram for a given model. With the optional parameter burn=True, addition of the burning zones from an existing *.burn file.
- plotRatio(): plot of the ratio between two variables or one variable and the given index of a second one
- Summary_plots(): set of plots on 4 windows based on Raphael Hirschi's dhr14.py (EVOL) or structure10.py (STRUC), summarising the evolution or structure of a model.

Examples of more complex plots are given in Sec. 8.

For any questions or suggestions, contact sylvia.ekstrom@unige.ch

2 Reading of files

2.1 Individual loading

2.2

```
• Loading an EVOL file
   Command: loadE('#file',num_★)
   The optional parameters are:
          - num_deb=i, to skip i lines at the beginning of the file (default: 0);
          - num_end=j, to stop before reading j lines at the end (default: -1);
          - format='fmt', where fmt is one of the following:
                    · o2013: standard or reduced .wg files;
                    · tgrids: grids files;
                    · tools: files generated by the interactive online tools;
                    · bin: binary version of the code;
                    · preMS: pre-MS version of the code;
                If omitted, the format will be automatically detected (default).

    forced=True, to skip the verification of the number entered as num_★ (default: False);

          - wa=True, to read an associated .wa file (only possible for .wg files, default: False);
          - quiet=True, to switch off the printings (default: False).
   Usage: loadE('~/calc/Z002/P015z02S0/P015z02S0.wg',1)
           loadE('~/grids/tables/M015z02S0.dat',2)
• Loading a STRUC file
   Command: loadS('#file',num_★)
   The optional parameters are:
          - toread=i, to read a given structure located in the file (default: all);
          - format='fmt', where 'fmt' is one of the following:
                    · o2013: standard .v file;
                    · o2010: old .v file;
                    · full: full StrucData file:
                If omitted, the format will be automatically detected (default).
          - quiet=True, to switch off the printings (default: False).
   Usage: loadS('P015z02S0.v0001001',1)
           loadS('P015z02S0.v0001051.gz',2)
           loadS('P015z02S0_StrucData_0001001.dat',3)
   NB: The structure files don't need to be unzipped before loading.
• Loading a cluster file
   Command: loadC('#file',num_★)
   The optional parameters are:
          - num_deb=i, to skip some lines at the beginning (default: 0);
          - format='fmt', where 'fmt' is one of the following:
              ·cluster:
              · isochr;
              If omitted, the format will be automatically detected (default);
          - random=n, to load only n randomly selected lines from the cluster;
          - quiet=True, to switch off the printings (default: False)
   Usage: loadC('Cluster_z0.014_t07.900.dat',1)
           loadC('Isochr_Z0.006_Vini0.40_t09.240.dat',2)
   Grouped loading
• Loading files from a list
   Command: loadXFromList('file_name')
          where X stands for one of the following: E, S, or C and file_name is a text file containing the list of files to
          be loaded (be sure to write the full path).
   The optional parameters are:
          - ini_index=i, to set the num_★ of the first model (default 1).
                 The numbering of the n models in the list is automatically set from i to i+n
                except in struc mode where more than one structure might exist in the file;
          - format=' format', to force the format definition (default: auto-detection);
          - forced=True, to skip verification of the number entered as num \star (default: False);
          - quiet=True, to switch off the printings (default: False).
```

Usage: loadEFromList('MyLoadedFiles.txt')

• Loading models directly from a directory

Command: loadXFromDir('dir_name')

where X stands for one of the following: E, S, or C

This command loads automatically all suitable files existing in the given directory:

- all .wg, .dat, or .grids files (EVOL);
- all *StrucData*.dat, or .v files (struc);
- all Clu*.dat, or Iso*.dat files (CLUSTER).

The optional parameters are:

- select='string', to restrict the loading to files having string in their name (default empty);
- ini_index, to set the num_★ of the first model (default 1);
- format='fmt', to set the format of the files. Left empty (default), the format is automatically detected;
- forced=True, to skip verification of the number entered as num_★ (default: False);
- quiet=True, to switch off the printings (default: False).

Usage: loadEFromDir('Grids2010/tables/Z014/')

loadEFromDir('Grids2010/wgred/Z014',select='S0',ini_index=12)

loadCFromDir('SYCLIST_Clusters/sylviaegn','v0.50',3)

2.3 Various commands

• Adding a column for the reading of modified files

Command: add_column(['varName',col_num],'label','category')

Back to normal with the command standard_columns().

• Reloading a file

Command: reloadX(num_★)

where X stands for one of the following: E, S, or C

NB: the options when loading the file are remembered, so only the num_★ is needed; to reload the file with different options, use the standard loadX() command with forced=True

3 Models selection

• Switching from one mode to another

Command: switch('mode')

where mode is one of the following: evol, struc, or cluster.

• Selection:

Command: select_model(num_★)

This command defines entirely the selected models (erases the previous selection)

Usage: select_model(2)

select_model([1,2,3,4,5])

select_model(range(1,6)) (same result as above)

Command: select_all()

This command selects all the loaded models of the current mode.

• Addition of a model:

Command: $add_model(num_*)$

This commands keeps the previous selection of models and adds the num \star one(s)

• Removal of a model:

Command: del_model(num_★)

• Recall of loaded models:

Command: Loaded('mode')

where mode is one of the following: evol, struc, cluster. Called without argument, it refers to the current mode.

4 Variables

• List of available variables

Command: VarEvol() (EVOL)
Command: VarStruc() (STRUC)
Command: VarCluster() (CLUSTER)
(see also section 7)

• Getting the values of a variable for a model

Command: Get_Var('variable',num_★)

• Creation of a new variable

• Deletion of a new variable

Command: Del_Var('varName')

with the optional argument num \star . Called without it, it deletes the variable for all loaded models.

Getting the derivative of variable 1 by variable 2

Command: Deriv('var1', 'var2')

Called with a number, or a list of numbers, restricts the derivative calculation to the models number given.

Usage: Deriv('M','t',[1,2])

• Splitting of a vector into its positive and negative components

Command: Vector_split('variable',num_*)
Creates vectors variable_pos and variable_neg.

• Computation of the colours in EVOL mode

Command: colours_calc(num_★)

with the optional argument num_★. Called without it, it performs the correction for all the clusters loaded.

• Correction of M_V and B-V for a distance modulus and reddening excess (only in Cluster mode)

Command: colour_corr(excess,dist_modulus)

It creates new variables called 'M_V_corr' and 'B-V_corr'.

The optional argument num_★ restricts the correction to the cluster(s)/isochrone(s) specified.

• Addition of noise on a variable (only in Cluster mode)

Command: add_noise('varName',errorValue)

It creates new variables called $`varName_noised$.

The optional argument num_★ restricts the correction to the cluster(s) specified.

5 Graphics

The philosophy of plotting with Origin_Tools is to first define which variable will sit on the *x*-axis, and then to do the plot with a commande giving the *y*-axis variable.

5.1 Basics

• Definition of the x axis

Command: defX('variable')

As default, the axis is t [Myr] in evol mode, M_r in struc mode, and T_{eff} in cluster mode.

• Basic command

Command: Plot('variable_y')

• Plot variable y only if a condition is fulfilled

```
Command: Plot('y',plotif=['var_cond','condition'])
Usage: Plot('He4s',plotif=['H1c','>0.'])
    Plot('gsurf',plotif=[['Vsurf','Vsurf'],['>200.','<=300']])</pre>
```

• Graph with variable(s) in log

Command: logVar('axis')

with axis=x,y,z,xy,xz,yz, then Plot('variable'). Back to real values with no_logVar('axis').

NB: to plot the real value but on a logarithmic scale, use logScale('axis') (described in section 6.1)

• Ratio between two variables

Command: plotRatio('var_1','var_2')

The optional argument index=#line allows to divide variable 1 by the value of variable 2 at that line. Useful for example to plot the relative evolution of a variable with respect to its initial value: plotRatio('var', 'var', index=0).

• 3rd variable in colour code over the curve

Command: Plot_colour('var_y','var_z')

The optional arguments are:

- binz=n: to limit the number of colours in the map
- s='var_s': to resize the points according to the value of 'var_s' (automatic switch to Points(True));
- logs=True: to take the log of this 'var_s';
- plotif=['var_cond', 'condition']: to limit the plot to a given condition on a variable;
- ticks=[tick list]: to define the location of the colour bar ticks.

The colour map can be changed with the command set_colourMap('cmap') (see Sec. 6.2, page 12)

5.2 Predefined graphs

Several predefined plots can be drawn with the following commands, without the need to define the *x*-axis beforehands. Depending on the mode (evol, struc, or cluster), different options can be available.

5.2.1 Graphs in EVOL mode

• HR diagram

Command: HRD_plot()

The following optional arguments are available:

- spectro=True: for the sHRD $(T_{\text{eff}}^4/g \text{ vs } T_{\text{eff}})$;
- corr=True: to use of $T_{\text{eff,corr}}$ instead of T_{eff} ;
- zcol='z_var': to colour-code the line with the value of z_var;
- binz=n: to limit the number of colours when using zcol (default 256);
- plotif=['var_cond', 'condition']: to limit the plot to a given condition on a variable;
- ticks=[tick list]: to define the location of the colour bar ticks.

• Colour-magnitude diagram

Command: CMD()

Entered without argument, plots the M_V versus B-V diagram. To plot other colours, enter a 3-character string

like 'BUB' (for M_B versus U-B) or 'VVI' (for M_V versus V-I).

The following optional arguments are available:

- zcol='z_var': to colour-code the line with the value of z_var;
- binz=n: to limit the number of colours when using zcol (default 256);
- plotif=['var_cond', 'condition']: to limit the plot to a given condition on a variable;
- ticks=[tick list]: to define the location of the colour bar ticks.

• Kippenhahn diagram

Command: Kippen(num_★)

The following optional arguments are available:

- hatch='/','\','-','|','+','x','0','o','.','*': to hatch the convective zones
 (repeat the symbol to get a denser hatching: '++','//');
- noshade=True to remove the grey shading.

• $\log T_{\rm c}$ – $\log \rho_{\rm c}$ diagram

Command: rhoT()

The following optional arguments are available:

- deg=True: to draw the degeneracy line (actually True by default);
- PISN=True: to shade the Γ_1 < 4/3 zone;

- zcol='z_var': to colour-code the line with the value of z_var;
- binz==n: to limit the number of colours when using zcol (default 256);
- plotif=['var_cond', 'condition']: to limit the plot to a given condition on a variable;
- ticks=[tick list]: to define the location of the colour bar ticks.

• $\log g - \log T_{\rm eff}$ diagram

Command: gTeff()

The following optional arguments are available:

- corr=True: to use $T_{\text{eff,corr}}$ instead of T_{eff} ;
- zcol='z_var': to colour-code the line with the value of z_var;
- binz=n: to limit the number of colours when using zcol (default 256);
- plotif=['var_cond', 'condition']: to limit the plot to a given condition on a variable;
- ticks=[tick list]: to define the location of the colour bar ticks.

• $Y - \log T_{\text{eff}}$ diagram

Command: YTeff()

The following optional arguments are available:

- corr=True: to use $T_{\text{eff,corr}}$ instead of T_{eff} ;
- zcol='z_var': to colour-code the line with the value of z_var;
- binz=n: to limit the number of colours when using zcol (default 256);
- plotif=['var_cond', 'condition']: to limit the plot to a given condition on a variable;
- ticks=[tick list]: to define the location of the colour bar ticks.

• Abundances evolution

Command: Abund('c') (centre)
Command: Abund('s') (surface)

• N/C vs N/O diagram

Command: NCNO()

The following optional parameter is available:

- plotif=['var_cond', 'condition']: to limit the plot to a given condition on a variable

• Set of plots summarising a model

Command: Summary_plots(num_★)

This command automatically generates the following plots:

- window 1: HRD, Abund(c), Kippenhahn, L and $T_{\rm eff}$ vs time
- window 2 : V_{surf} , $\Omega/\Omega_{\text{crit}}$, \dot{M} , $\Omega_{\text{c}}/\Omega_{\text{s}}$
- window 3: Abund(s), N/O, N/H, N/C
- window 4 : $T_c \rho_c$, T_c and ρ_c vs time

With the optional argument 'legend', a legend is added to the plots.

5.2.2 Graphs in STRUC mode

$\bullet \log T - \log \rho \operatorname{diagram}$

Command: rhoT()

The following optional parameter is available:

- plotif=['var_cond', 'condition']: to limit the plot to a given condition on a variable

• Energy generation zones

Command: eps(mode)

with mode defining the x-axis variable:

- -1: M_r/M_{tot} (default);
- -2: $r[R_{☉}]$;
- -3: free x variable defined by defX().

The optional parameter conv=False avoids the shading of convective zones (True by default if only one structure selected).

• Gradients (∇_{ad} - ∇_{rad} - ∇_{μ})

Command: Nablas(num_★)

• Abundances profile

Command: Abund('p')

ullet Diffusion coefficients ($D_{ m conv}$ - $D_{ m shear}$ - $D_{ m h}$ - $D_{ m eff}$ - $K_{ m ther}$)

Command: Coeff()

• Addition of the convective zones on the plot

Command: convZones(num_★)

The optional parameter colour='colour' changes the default colour (grey).

• Set of plots summarising a structure

Command: Summary_plots(num_★)

This command automatically generates the following plots:

- window 1 : Abund(p)
- window $2:T,r,P,\rho$
- window 3 : L and ϵ , T vs ρ , $\epsilon_{\rm reac}$ and $\epsilon_{\rm grav}$, nablas and kappa
- window 4 : D, N^2 , c_{sound} and V_{MLT} , magn. field variables
- window 5 : Ω and $\Omega/\Omega_{\rm crit}$, $V_{\rm eq}$, j, U and V

With the optional argument 'legend', a legend is added to the plots.

5.2.3 Graphs in CLUSTER mode

• HR diagram

Command: HRD_plot()

The following optional parameters are available:

- spectro=True: for the sHRD $(T_{\rm eff}^4/g~{\rm vs}~T_{\rm eff})$
- corr=True: to use of $T_{\rm eff,corr}$ instead of $T_{\rm eff}$
- dark=True: to use of L and T_{eff} corrected for the gravity(+limb) darkening
- zcol='z_var': to colour-code the line with the value of z_var
- binz=n: to limit the number of colours when using zcol (default 256)
- plotif=['var_cond', 'condition']: to limit the plot to a given condition on a variable

• Colour-magnitude diagram

Command: CMD()

Entered without argument, plots the M_V versus B-V diagram. To plot other colours, enter a 3-character string like 'BUB' (for M_B versus U-B) or 'VVI' (for M_V versus V-I).

The following optional arguments are available:

- noised='xy': to plot the noised variables (created with add_noise())
- zcol='z_var': to colour-code the line with the value of z_var
- binz=n: to limit the number of colours when using zcol (default 256)
- noised='xy': to plot the noised variable (created by add_noise());
- plotif=['var_cond', 'condition']: to limit the plot to a given condition on a variable

• $\log g - \log T_{\text{eff}}$ diagram

Command: gTeff()

The following optional arguments are available:

- dark=True: to use $T_{\text{eff,lgd}}$ instead of T_{eff}
- corr=True: to use $T_{\rm eff,corr}$ instead of $T_{\rm eff}$
- surf=True: to use g_{surf} instead of g_{pol}
- mean=True: to use g_{mean} instead of g_{pol}
- noised='xy': to plot the noised variables (created with add_noise())
- zcol='z_var': to colour-code the line with the value of z_var
- binz=n: to limit the number of colours when using zcol (default 256)
- plotif=['var_cond', 'condition']: to limit the plot to a given condition on a variable

• Abundances evolution

Command: Abund('s')

• N/C vs N/O diagram

Command: NCNO()

The following optional parameter is available:

-plotif=['var_cond', 'condition']: to limit the plot to a given condition on a variable

• Histogram of variable var

Command: Histo(var,bin_number)

The optional parameter cum=True makes a cumulative histogram.

5.3 Saving the figures

The python window allows to save the current figure in png format, with a clic on ...



For a vectorised format, the command MyFig('#file_name') can be used. The default format is pdf, but it might be customised with the optional argument format='#format'.

5.4 Extra informations on the graphs

• Superimposing plots

Command: plot2var('mode')

By default, the mode is 'same' that draws all the variables on the same y axis.

The mode 'double' allows to draw the 2nd variable with an independent axis on the right.

NB: back to normal with plot1var()

• Keeping the plot to draw something else on it

Command: keep_plot(True)

Back to normal with keep_plot(False)

Addition of marks at a given time step

Command: timesteps(True)

This command must precede the plotting command.

The time step can be determined by the command set_deltat(value). By default, this command acts on all stars selected for plotting. It is possible to limit it to some specific stars by entering a number, or a list of numbers after the value.

Example: set_deltat(5.e5,[2,3])

The type of marker can be chosen by timestep_marker('marker').

• Marking the beginning and end of a given burning phase

Command: mark_phase('fuel')

with fuel being one of the following: H, He, C, Ne, O, or Si.

The following optional parameters are accepted:

- marker=[' ',' ']: changes the marker style (default ['o','x']);
- colour: changes the marker colour (default black).

• Drawing iso-radius lines in an HRD

Command: isoRadius()

The following optional parameters are accepted:

- colour='colour': sets the colour of the line (default grey 0.80);
- -line='style': sets the line style. It can be any of the matplotlib styles (see Sec. 6.2, default '-');
- fontsize=size: sets the fontsize for the radius labels.

• Drawing a vertical or horizontal line at a given value

Command: xline(value) (vertical)

Command: yline(value) (horizontal)

The following optional parameters are accepted:

- colour='colour': sets the colour of the line (default grey 0.80);
- line='style': sets the line style. It can be any of the matplotlib styles (see Sec. 6.2, default'-').

• Drawing a line from (x_1,y_1) to (x_2,y_2)

Command: line(x_1, x_2, y_1, y_2)

The following optional parameters are accepted:

- colour='colour': sets the colour of the line (default grey 0.80);
- -line='style': sets the line style. It can be any of the matplotlib styles (see Sec. 6.2, default '-').

• Drawing a line with a given slope

Command: slope(value)

The following optional parameters are accepted:

- centre=[x,y] allows to centre the line at point (x,y), default (0,0);
- colour='colour': sets the colour of the line (default grey 0.80);
- -line='style': sets the line style. It can be any of the matplotlib styles (see Sec. 6.2, default '-').

• Drawing a dot at the coordinate (x,y)

Command: dotxy(x,y)

The following optional parameters are accepted:

- 'style' allows to enter directly the style of the point: dotxy(x,y,'ro');
 - Otherwise, the style is defined by the values entered with set_pointStyle() and set_colourFlag().
- err=[xerr, yerr] allows to draw error bars;
 - For asymmetric errorbars, write err=[[xerr_left,xerr_right],[yerr_down,yerr_up]].
- label='string' writes string besides the point;
- ha='pos' allows to put this string left or right of the point;
- fontsize=int sets the size of the font.

• Shading a zone defined by vectors x and y

Command: shade(x,y)

The following optional parameters are accepted:

- colour='colour': sets the colour of the shaded zone (grey 80% by default);
- alpha=value: sets the transparency (20% by default);
- hatch='/', '\', '-', '|', '+', 'x', '0', 'o', '.', '*' (repeat symbol to get a denser hatching: '++', '///').

• Shading a vertical or horizontal region around x1 and x2, or y1 and y2

Command: shade_x(x1,x2) ((vertical))

Command: shade_y(y1,y2) ((horizontal))

The following optional parameters are accepted:

- colour='colour': sets the colour of the shaded zone (grey 80% by default);
- alpha=value: sets the transparency (20% by default);
- hatch='/','\','-','|','+','x','0','o','.','*' (repeat symbol to get a denser hatching: '++','///').

• Finding a polynomial fit of degree N to the curve

Command: fit_poly(N)

The following optional arguments are accepted:

- y=var: to change the target curve (default: last curve drawn);
- colour=colour: to change the colour of the fit curve (default grey 0.80).

• Addition of external data on a graph

Command: plotExternal('#File_name',x_column,y_column)

The following optional arguments are accepted:

- skip=n: to skip n lines at the beginning the file #File_name.
 - Note that header lines starting by # are automatically skipped;
- last=n: to stop at line n;
- style='style': to define the points/lines style (example: 'ro', 'b+', 'g.', 'm-').

By default, the general style (defined by set_colourFlag() and either set_pointStyle() or set_lineStyle()) will be taken;

- colz=n: to define the column of the variable that will colour-code the points (works only with Points(True));
- cols=n: to define the column of the variable that will give the size of the points (works only with Points(True));
- $-\log=$ 'x', 'y', 'z', 's' or any combination 'xy', 'xz',...: to take the log of the variable(s);
- zlabel=str: to define the label of the z-axis;
- clim='new'/'old': to set new limits to the colour bar or to retrieve the existing ones;
- new: to create a new figure rather than adding the data to the current figure.

5.5 Text and legends

• Setting the fontsize

Command: set_fontSize()

This command accepts either the size directly (set_fontSize(12)),

or a factor to apply to the actual fontsize (set_fontSize(f=0.5)).

Called with 'default' as argument (set_fontSize('default')), it recovers the default fontsize (24).

• Addition of the legend

Command: put_legend()

The following optional arguments are accepted:

- pos=int: sets the position of the legend box on the plot. The value can be entered directly as first argument.

The integer coding the positions are:

- · 1: top right (default);
- · 2: top left;
- · 3: bottom left;
- · 4: bottom right;
- · 5: middle right;
- · 6: middle left;

```
· 7: middle right again (don't ask me why);
```

- · 8: bottom centre;
- · 9: top centre;
- · 10: middle centre.
- -label=['line 1', 'line 2']: to define the label (default: variable y);
- fontsize=int: sets the fontsize (default: axis fontsize / 1.5).

• Addition of text in the graph

```
Command: add_label(x,y,'string')
```

The following optional parameters are accepted:

- colour='colour': sets the colour of the line (default black);
- fontsize=value modifies the fontsize (default: 24);
- ha='pos': sets the horizontal position relative to (x,y). 'pos' can be any of the following: 'left', 'center', or 'right' (default: 'left').
- va='pos': sets the vertical position relative to (x,y). 'pos' can be any of the following: 'baseline', 'bottom', 'center', or 'top' (default: 'bottom').

• Addition of a title at the top of the window

Command: top_label('string')

The optional argument fontsize=value modifies the fontsize only for this command.

6 Adaptations to specific needs

The following set of commands allows you to modified many settings for the rendering of your plots.

NB: At any time, you can go back to the default ones (as at launch) with the command default_settings().

6.1 Limits and axes

• Change of the limits

```
Command: Limits(xmin=,xmax=,ymin=,ymax=)
```

Back to normal with noLimits('x', 'y' or 'xy'). noLimits() without argument corresponds to 'xy'.

Command: CBLimits(min=,max=)

This command sets the limits for var_z in Plot_colour(). Back to normal with noCBLimits().

• Retrieving the limits of the current axes

Command: get_limits()

• Keeping the actual limits

Command: keep_limits()

Back to normal with keep_limits(False).

• Inversion of an axis

```
Command: axis_inv('x','y' or 'xy')
```

NB: back to normal with no_axis_inv('x', 'y' or 'xy').

no_axis_inv() without argument corresponds to 'xy'.

• Axis in logarithmic scale

Command: logScale('x','y' or 'xy')

NB: back to normal with no_logScale()

• Modification of an axis label

Command: change_label('axis','label')

• Change the number of minor ticks

Command: set_tickNumber(N)

NB: back to default behaviour (automatic) with set_tickNumber(0)

6.2 Points/curves

• Switch from curves to points

Command: Points(True)

Back to curves with Points(False)

• Choice of line style

Command: set_lineStyle('style')

where style can be:

- cycle_colour so that the line style changes only after a whole colour sequence;
- cycle_all so that each curve is drawn with a different style;
- any of the following: -, -, :, . for solid, dashed, dotted, and dash-dotted lines, respectively

• Choice of point style

Command: set_pointStyle('style')

where 'style' can be:

- cycle_colour so that the points style changes only after a whole colour sequence;
- cycle_all so that each model points are drawn with a different style;
- any of the following: 'o' = \bigcirc , 's' = \bigcirc , 'p' = \bigcirc , 'd' = \diamondsuit , '*' = \diamondsuit , 'v' = \triangledown , '^' = \diamondsuit , '>' = \triangleright , '<' = \triangleleft ;
- a tuple (numsides, style, angle). style can be 0 (polygon), 1 (star-like), or 2 (squeletal). angle can be omitted.

• Choice of point size

Command: set_pointSize(value)

This command accepts either the direct value (set_pointSize(12)), or a factor f=x to multiply the actual size by x: set_pointSize(f=5.)

Called with the argument 'default' (set_pointSize('default')), it recovers the default point size (24).

• Choice of min and max point sizes in Plot_colour() and plotExternal()

Command: set_PSminmax(min,max)

To recover the default size (5,200), call set_PSminmax('default').

• Drawing empty symbols

Command: emptyPoints(True)

Back to filled symbols (default) with emptyPoints(False)

6.3 Colours

• Choice of a given colour for plotting

Command: set_colourFlag('value')

where value can be any of the following:

- a string stating:
 - the shortcut for the main colour names (b, g, r, c, m, y, k, w), or the full colour name as in html palette;
 - · cycle: to recover a colour sequence behaviour;
- a float to set a shade of grey;
- a triplet of floats to code direct RGB values (in a 0 to 1 scale).

• Start the colour sequence n steps further

Command: Plot('y',cshift=n)

• Choice of a colour sequence when plotting more than one model

Command: set_colourSequence('value')

Sets the colour sequence applied to the tracks when more than one is plotted.

value can be one of the following:

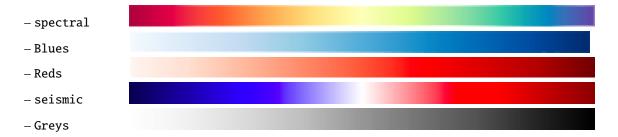
- 'c': for 'contrast', the default sequence (black,red,green,blue,cyan,magenta,orange,olive,pink,brown,grey);
- 'i': for 'iris' (black,blue,cyan,green,yellow,orange,red,magenta,purple,grey);
- 'p': to create a personalised sequence. Following the instructions, you'll have to enter a name for the sequence and then the new list in the format [[colours], [names]];
- any existing sequence name.

• Choice of a colour map for the 3rd variable

Command: set_colourMap('name')

where name can be any known cmap from matplotlib. We particularly recommend the following:





6.4 Windows

• Drawing of 2 or 4 plots on the same window

Command: multiPlot(2 or 4)

NB: back to normal with multiPlot(1)

6.5 Navigation within the graphs

The python window allows to easily zoom on a part of the graph after a clic on . It is also possible to navigate within the graph after clicking on . The original view is recovered with the button.

• Activation of the cursor

Command: cursor()

The coordinates of clicks are written on the terminal. To exit the cursor mode, just hit Enter on the terminal window.

• Measurement of the distance between two points

Command: dist()

The clicks are validated by hitting Enter on the terminal window.

The coordinates are written there, as well as Δx , Δy and the distance.

7 List of the variables

7.1 In EVOL mode

• MODEL:

line:	model num	t_tauH:	$t/ au_{ m H}$
3.6	37.537.3	1	1 (

M: $M[M_{\odot}]$ ageadv: log(time before collapse [yr])

t: t [yr] tauKH: $\tau_{KH} \text{ [yr]}$ t6: t [Myr] rhom: $\rho_{\text{m}} \text{ [g cm}^{3} \text{]}$

Besides, the following global variables are known:

 $\label{eq:file_problem} \textit{FileName}: \qquad \textit{#loaded_file} \qquad \qquad \textit{tau}: \qquad \qquad \textit{lifetimes} \; [\textit{yr}] \; ([\tau_{H}, \tau_{He}, \tau_{C}, \tau_{Ne}, \tau_{O}])$

Mini: $M_{\text{ini}}[M_{\odot}]$ format: #format

• CENTRE:

Mcc : $M_{\rm cc} \, [M_{\odot}]$ rhoc : $\log(\rho_{\rm c} \, [{\rm g \, cm^{-3}}])$ Mccrel : $M_{\rm cc}/M_{\rm tot}$ Tc : $\log(T_{\rm c} \, [K])$

• SURFACE:

Teff: $\log(T_{\rm eff}\,[{\rm K}])$ R: $R\,[R_\odot]$ Teffcorr: $\log(T_{\rm eff}\,[{\rm K}])$ corrected for Rpol: $R_{\rm pol}\,[R_\odot]$

the wind thickness (WR) gsurf: $\log(g_{\text{surf}} [\text{cm s}^{-2}])$

L: $\log(L/L_{\odot})$ gpol: $\log(g_{\rm pol} \ [{\rm cm \, s^{-2}}])$ Mbol: $M_{\rm bol}$ fwg: $\log(g/(T_{\rm eff}/10'000 \ {\rm K})^4)$

sL: $\log(\mathcal{L}/\mathcal{L}_{\odot})$ ZCext: $M_{\text{ZC,ext}}$

 $GammaEdd: \quad \Gamma_{Edd}$

• ROTATION:

 $\Omega_{\rm surf} [\rm s^{-1}]$ Omega surf: oblat: $R_{\rm pol}/R_{\rm eq}$

 $\Omega_{\text{cen}} [s^{-1}]$ Omega_cen: GammaOmega: $\Omega/min(\Omega_{crit,1},\Omega_{crit,2})$

OOc: $\Omega/\Omega_{\rm crit}$ rot corr: F_{Ω}

 $\mathcal{L}_{\text{tot}} [10^{53} \,\mathrm{g \, cm^2 \, s^{-1}}]$ VVc: $V/V_{\rm crit}$ Ltot: $V_{\rm surf}$ [km s⁻¹] $\mathcal{L}_{\text{tot,int}} [10^{53} \,\mathrm{g \, cm^2 \, s^{-1}}]$ Vsurf: Ltotint: $j_{5M_{\odot}}$ [10¹⁶ cm² s⁻¹] $V_{\rm crit,1} \, [{\rm km \, s^{-1}}]$ jspe5: Vcrit1: $j_{3M_{\odot}}$ [10¹⁶ cm² s⁻¹] Vcrit2: $V_{\rm crit,2} \, [{\rm km \, s^{-1}}]$ jspe3: $I [10^{57} \,\mathrm{g\,cm^2}]$ period: P[d] mominert:

• WINDS:

 $V_{\rm esc}$ [km s⁻¹] Mdot: $\log(\dot{M} [M_{\odot} \text{yr}^{-1}])$ Vesc:

Mdot mech: $(\log(\dot{M})_{\rm mech} [M_{\odot} \, \mathrm{yr}^{-1}])$ dlelex: $\Delta \mathcal{L}_{\text{rad+aniso+mech}} [10^{53} \,\text{g cm}^2 \,\text{s}^{-1}]$ Bmin: dMmech:

 $dM_{\rm mech} [M_{\odot}]$ B_{\min} [G] (minimal magn. field $P_{\rm winds}$ [erg s⁻¹] Pwinds: for a wind-surface coupling)

• ENERGETICS:

Vinf:

 V_{∞} [km s⁻¹]

Epot: $E_{\rm pot} [E_{51}]$ Erot: $E_{\rm rot} [E_{51}]$ Egaz:

 $E_{\rm th,gaz}$ [E_{51}] phase: evolutionary phase Erad: $E_{\rm rad}$ [E_{51}]

• ABUNDANCES:

²²Ne (surf., centr.) [mass frac.] H1s,H1c: ¹H (surf., centr.) [mass frac.] Ne22s,Ne22c: ³He (surf., centr.) [mass frac.] ²⁶Al (surf., centr.) [mass frac.] He3s,He3c: Al26s, Al26c:

He4s,He4c: ⁴He (surf., centr.) [mass frac.] Zsurf: Z_{surf} [mass frac.] Be7c: ⁷Be (centr.) [mass frac.] C12C13: $\log(^{12}C/^{13}C \text{ [numb.]})$

⁸B (centr.) [mass frac.] $\log(^{12}C/^{13}C) - \log(^{12}C/^{13}C)_{ini}$ B8c: C12C13rel:

¹²C (surf., centr.) [mass frac.] log(N/H [numb.] + 12)C12s,C12c: NH: ¹³C (surf., centr.) [mass frac.] C13s,C13c: NHrel: $log(N/H) - log(N/H)_{ini}$ ¹⁴N (surf., centr.) [mass frac.] N14s,N14c: NC: log(N/C [numb.])O16s,O16c: ¹⁶O (surf., centr.) [mass frac.] NCrel: $log(N/C) - log(N/C)_{ini}$ ¹⁷O (surf., centr.) [mass frac.] O17s,O17c: NO: log(N/O [numb.]) $log(N/O) - log(N/O)_{ini}$

¹⁸O (surf., centr.) [mass frac.] O18s,O18c: NOrel: ²⁰Ne (surf., centr.) [mass frac.] Ne20s,Ne20c:

In struc mode

• STRUCTURE:

shell: shell number rprev: $r_{\rm prev} [R_{\odot}]$ $g_r \, [\text{cm s}^{-2}]$ Mfrac: $M_r/M_{\rm tot}$ g : $N^2 [s^{-1}]$ N2: Mr: $M_r [M_{\odot}]$ r cm: *r* [cm] Nmu2:

 $N_{\mu}^{2} [s^{-1}]$ $N_{T}^{2} [s^{-1}]$ $r[R_{\odot}]$ NT2: r:

Besides, the following global variables are known:

total shells number M_tot: $M_{\rm tot} [M_{\odot}]$ nshell:

#loaded_file FileName: age: age [yr] Model: model number timestep: δt [s]

• ENERGY:

 $\epsilon(^{16}O(\alpha, \gamma)^{20}Ne) [\text{erg g}^{-1}\text{s}^{-1}]$ L: epsOagNe: $L_r/L_{\rm tot}$

 ϵ (H) [erg g⁻¹s⁻¹] $-\epsilon_{\nu}$ [erg g⁻¹s⁻¹] epsH: epsnu: $\epsilon_{\text{nucl}} + \epsilon_{\nu} [\text{erg g}^{-1} \text{s}^{-1}]$ ϵ (He) [erg g⁻¹s⁻¹] epsHe: eps_reac: ϵ (C) [erg g⁻¹s⁻¹] $\epsilon_{\rm grav}$ [erg g⁻¹s⁻¹] epsC: epsgrav: $\begin{array}{l} \epsilon(3\alpha) \ [\mathrm{erg} \ \mathrm{g}^{-1} \mathrm{s}^{-1}] \\ \epsilon(^{12}C(\alpha,\gamma)^{16}O) \ [\mathrm{erg} \ \mathrm{g}^{-1} \mathrm{s}^{-1}] \end{array}$ eps3a: dEdP: $d \ln E/d \ln P$

epsCagO: dEdT: $d \ln E/d \ln T$

• EOS:

rho: $ho \, [\mathrm{g} \, \mathrm{cm}^3]$ Nabmu: $abla_{\mu}$

mu: μ (mean molecular weight) drhodP: $d \ln \rho / d \ln P$ muprev: μ of previous iteration delta: $\delta = -d \ln \rho / d \ln T$ mufit: μ (smoothed) cs: c_{sound} [cm s⁻¹]

mue : μ_e psi : ψ (indicator of degeneracy)

• THERMO:

 $P [g cm^{-1} s^{-2}]$ P: Nabrad: ∇_{rad} beta: $\beta = P_{\rm gas}/P_{\rm tot}$ Nabla_int: ∇_{int} $\kappa \, [\mathrm{cm}^2 \, \mathrm{g}^{-1}]$ Hp: H_P [cm] kappa: T[K]dkdP: $d \ln \kappa / d \ln P$ T: K_{ther} [cm² s⁻¹] Kther: dkdT: $d \ln \kappa / d \ln T$ C_P [ergs g⁻¹ K⁻¹] Nabla: Cp: Nabad: $\nabla_{\rm ad}$ V_MLT: $V_{\rm MLT}$ [cm s⁻¹]

• ROTATION:

Omega: $g[s^{-1}]$ $g[s^{-1}]$ $g[s^{-1}]$

Omegaprev: Ω of previous time step jK: j_{Kerr} [cm²s⁻¹] Omegacons: Ω (when only local conservation of angular momentum is applied) jK: j_{Kerr} [cm²s⁻¹] j_{Kerr} [cm²s⁻¹]

 $D_{\rm shear}$ [cm² s⁻¹] Omfit: Ω (smoothed) Dshear: dlodlr: $d \ln \Omega / d \ln r$ Deff: $D_{\rm eff}$ [cm² s⁻¹] $D_{\rm circ}$ [cm² s⁻¹] obla: Dcirc: $r_{\rm pol}/r_{\rm eq}$ $V_r [cm s^{-1}]$ $D_{\rm conv}$ [cm² s⁻¹] Vr: Dconv: Veq: $V_{\rm eq}$ [km s⁻¹] Ur: U_r [cm s⁻¹] \mathcal{L}_r [g cm² s⁻¹] $Ri = N^2/(dV/dz)^2$ Lang: Richardson:

 $jr: j_r [cm^2s^{-1}]$

• MAGNETISM:

Br: $B_r[G]$ etask: η/K $N_{\rm mag}^2 \ [{\rm s}^{-1}]$ Bphi: $B_{\phi}[G]$ N2mag: $D_{\mathrm{mag},\Omega}$ [cm² s⁻¹] qmin: DmagO: q_{\min} $\omega_{\rm Alfven} \, [{\rm s}^{-1}]$ $D_{\text{mag},X} \text{ [cm}^2 \text{ s}^{-1}\text{]}$ alfven: DmagX:

• ABUNDANCES:

²⁸Si [mass frac.] ¹H [mass frac.] H1: Si28,Si28_alu: ³²S [mass frac.] ^{3,4}He [mass frac.] He3,He4: S32: ^{12,13,14}C [mass frac.] ³⁶Ar [mass frac.] Ar36: C12,C13,C14: ⁴⁰Ca [mass frac.] ^{14,15}N [mass frac.] N14,N15: Ca40: ^{16,17,18}O [mass frac.] ⁴⁴Ti [mass frac.] 016,017,018: Ti44: ^{18,19}F [mass frac.] F18.F19: Cr48: ⁴⁸Cr [mass frac.] ^{20,21,22}Ne [mass frac.] ⁵²Fe [mass frac.] Ne20,Ne21,Ne22: Fe52: ²³Na [mass frac.] ⁵⁶Ni [mass frac.] Na23: Ni56: ^{24,25,26}Mg [mass frac.] Mg24,Mg25,Mg26: protons: protons [mass frac.] A126,A127: ^{26,27}Al [mass frac.] neutrons: neutrons [mass frac.]

7.3 In cluster mode

• INITIAL CONDITIONS:

Zini: Z_{ini} Angle: i [$^{\text{o}}$] Mini: M_{ini} [M_{o}] Bin: binary Oini: $\Omega/\Omega_{\text{crit,ini}}$ M1M2: M_1/M_2

Besides, the following global variables are known:

FileName: #loaded_file

• GLOBAL PROPERTIES:

M: $M[M_{\odot}]$ sL: $\log(\mathcal{L}/\mathcal{L}_{\odot})$ R: $R[R_{\odot}]$ Mbol: M_{bol}

Rpol: $R_{\rm pol} [R_{\odot}]$ $\log(g_{\rm surf} [{\rm cm \, s^{-2}}])$ gsurf: L: $\log(g_{\text{pol}} [\text{cm s}^{-2}])$ $\log(L/L_{\odot})$ gpol: L_gd: $\log(g_{\text{mean}} [\text{cm s}^{-2}])$ $\log(L/L_{\odot})_{\text{grav.dark}}$ gmean: $\log(g/(T_{\rm eff}/10'000\,{\rm K})^4)$ L_lgd: $\log(L/L_{\odot})_{limb+grav.dark}$ fwg:

Teff: $\log(T_{\rm eff}\,[{\rm K}])$ rhom: $\rho_{\rm m}\,[{\rm g\,cm^3}]$ Teffcorr: $\log(T_{\rm eff}\,[{\rm K}])$ Gammaedd: $\Gamma_{\rm Edd}$

Teff_gd: $\log(T_{\rm eff}\,[{\rm K}])_{\rm grav.dark}$ Mdot: $\log(\dot{M}\,[M_{\odot}\,{\rm yr}^{-1}])$ Teff_lgd: $\log(T_{\rm eff}\,[{\rm K}])_{\rm limb+grav.dark}$ dMmech: $dM_{\rm mech}\,[M_{\odot}]$

• ROTATION:

 $V_{\text{crit},1} \text{ [km s}^{-1}\text{]}$ OOc: Vcrit1: Ω/Ω_{crit} $\Omega_{\rm surf}$ [s⁻¹] $V_{\text{crit},2} \text{ [km s}^{-1}\text{]}$ Vcrit2: Omega_surf: $V_{\rm surf} \, [{\rm km \, s^{-1}}]$ period: P[d] Vsurf: $V \sin i \left[\text{km s}^{-1} \right]$ Vsini: oblat: $R_{\rm pol}/R_{\rm eq}$

• ABUNDANCES:

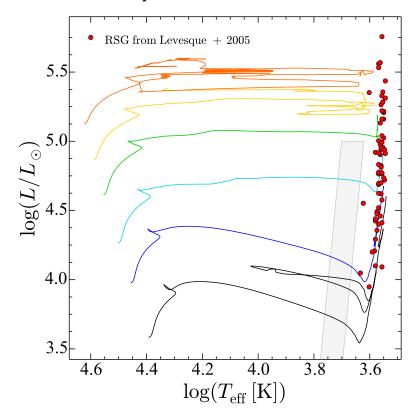
H1s:	¹ H [surf. mass frac.]	Ne22s:	²² Ne [surf. mass frac.]
He4s:	⁴ He [surf. mass frac.]	Al26s:	²⁶ Al [surf. mass frac.]
C12s:	¹² C [surf. mass frac.]	C12C13:	$\log(^{12}C/^{13}C \text{ [numb.]})$
C13s:	¹³ C [surf. mass frac.]	C12C13rel:	$\log(^{12}\text{C}/^{13}\text{C})$ - $\log(^{12}\text{C}/^{13}\text{C})_{\text{ini}}$
N14s:	¹⁴ N [surf. mass frac.]	NH:	log(N/H [numb.]+12)
O16s:	¹⁶ O [surf. mass frac.]	NC:	log(N/C [numb.])
O17s:	¹⁷ O [surf. mass frac.]	NCrel:	log(N/C)- $log(N/C)$ _{ini}
O18s:	¹⁸ O [surf. mass frac.]	NO:	log(N/O [numb.])
Ne20s:	²⁰ Ne [surf. mass frac.]	NOrel:	$log(N/O)$ - $log(N/O)_{ini}$

• COLOURS:

M_V:	$ m M_{ m V}$	U-B:	U-B
M_V_noise:	M _V noised	V-K:	V-K
M_B:	$ m M_{B}$	V-I:	V-I
B-V:	B-V	V-R:	V-R
B-V_noise:	B-V noised	H-K:	H-K
B2_V1:	$B_2 - V_1$	J-K:	J-K

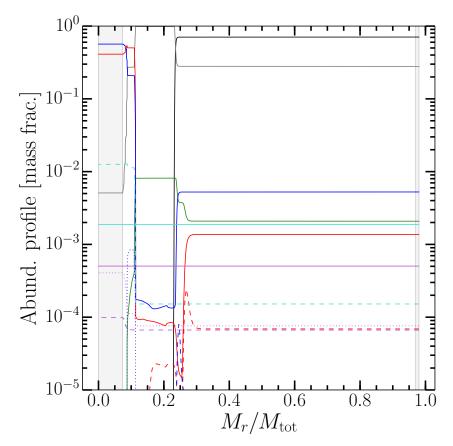
8 Examples

8.1 HR diagram with observational data points



```
Loading of the package:
      from origin_tools.Origin_Tools import *
Loading of the evolution files:
      loadE('/Path/to/file/P009z14S0.dat',1)
      loadE('/Path/to/file/P012z14S0.dat',2)
      loadE('/Path/to/file/P015z14S0.dat',3)
      loadE('/Path/to/file/P020z14S0.dat',4)
      loadE('/Path/to/file/P025z14S0.dat',5)
      loadE('/Path/to/file/M032Z14V0.dat',6)
Choice of the colour sequence 'iris':
      set_colourSequence('i')
Change of the upper limit in y:
      Limits(ymax=5.9)
Drawing of the HR diagram (predefined command):
      HRD_plot()
Passing to 'points' mode:
      Points(True)
Drawing of external data, read in a file:
      plotExternal('RSG_Levesque2005.dat',0,2,log='x',style='ro')
Drawing the legend of the external data:
      \label{lem:condition} $\operatorname{dotxy}(4.6,5.75,\text{style='ro',label='}\mathbb{R}SG\ from\ Levesque+2005}\)$', fontsize=16)$
Saving the figure:
      MyFig('HRDrainbow_RSG')
```

8.2 Abundances profiles with the convective zones shaded



Loading of a structure file:

loadS('/Path/to/file/P007z14S0.v0022721',1)

Labels written with LATEX:

iLatex(True)

Enhancement of the axes ticks:

set_tickSize(length=12,width=2)

Modification of the lower limit in *y*:

Limits(ymin=1.e-5)

y-axis in logarithmic scale:

logScale('y')

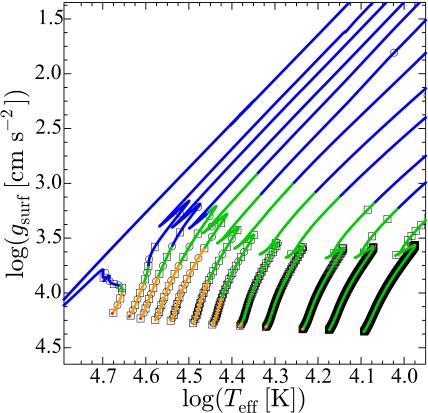
Drawing of the abundances profiles (predefined command):

Abund('p')

Addition of the convective zones as shaded areas:

convZones(1)

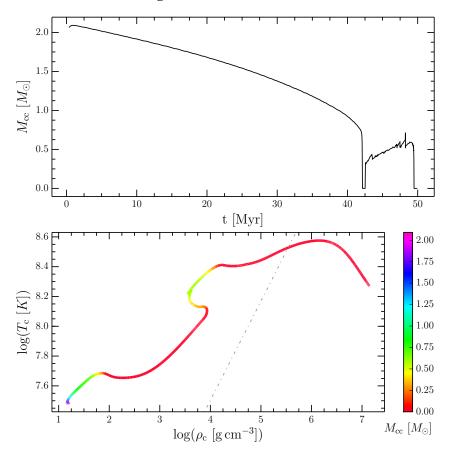
8.3 g_{surf} vs T_{eff} with time steps and various colours for velocities zones



```
Loading of a set of evolution files listed in a text file:
                                                                    Drawing g_{\text{surf}}:
       loadEFromList('loadModels.txt')
                                                                           Plot('gsurf')
Setting the line width to zero (no curves drawn):
                                                                    Switching off the markers for time steps:
       set_lineStyle('-',width=0)
                                                                           timesteps(False)
Switching on the markers for time steps:
                                                                    Setting the line width to 2:
       timesteps(True)
                                                                           set_lineStyle('-',width=2)
Setting the timesteps marker to squares:
                                                                    Selection of all models again:
       timestep_marker('s')
                                                                           select_model([1,2,3,4,5,6,7,8,9,10,11,12])
Definition of the x axis:
                                                                    Changing the colour to blue:
       defX('Teffncorr')
                                                                           set_colourFlag('b')
Changing the natural limits (can be done in one command):
                                                                    Drawing g_{\text{surf}} only if V_{\text{eq}} < 100 \,\text{km s}^{-1}:
                                                                           Plot('gsurf',plotif=['Vsurf','<100.'])</pre>
       Limits(xmin=3.95, xmax=4.79)
      Limits(ymin=1.35,ymax=4.65)
                                                                    Changing the colour to customed green:
Inversion the y axis:
                                                                           set\_colourFlag((0.,0.8,0.))
                                                                    Drawing g_{\text{surf}} only if 100 < V_{\text{eq}} \le 200 \,\text{km s}^{-1}:
       axis_inv('y')
                                                                           Plot('gsurf',plotif=[['Vsurf','Vsurf'],
Choice of a unique colour for all curves (black):
       set_colourFlag('k')
                                                                                                           [['>100.','<=200']])
                                                                    Changing the colour to orange:
Drawing g_{\text{surf}}:
       Plot('gsurf')
                                                                           set_colourFlag((1.,0.6,0.))
                                                                    Drawing g_{\text{surf}} only if 200 < V_{\text{eq}} \le 300 \,\text{km s}^{-1}:
Keeping the window for further plotting:
                                                                           Plot('gsurf',plotif=[['Vsurf','Vsurf'],
       plot2var()
Selection of a subset of models:
                                                                                                           ['>200.','<=300']])
       select_model([4,5,6,7,8,9,10,11,12])
                                                                    Changing the colour to red:
Changing the time step for those models:
                                                                           set_colourFlag('r')
                                                                    Drawing g_{\text{surf}} only if V_{\text{eq}} > 300 \,\text{km s}^{-1}:
       set_deltat(5.e5)
                                                                           Plot('gsurf',plotif=['Vsurf','>300.'])
Setting the timestep markers to circles:
```

timestep_marker('o')

8.4 Double figure with one of them having a 3rd variable in colour code



Loading of an evolution file:

loadE('/Path/to/file/P007z14S0.dat',1)

Labels written with LATEX:

iLatex(True)

Passing to two figures in a window mode:

multiPlot(2)

Reduction of the labels size:

set_fontSize(12)

Drawing of the first figure (default *x*-axis):

Plot('Mcc')

Choice of the *x*-axis for the 2nd figure:

defX('rhoc')

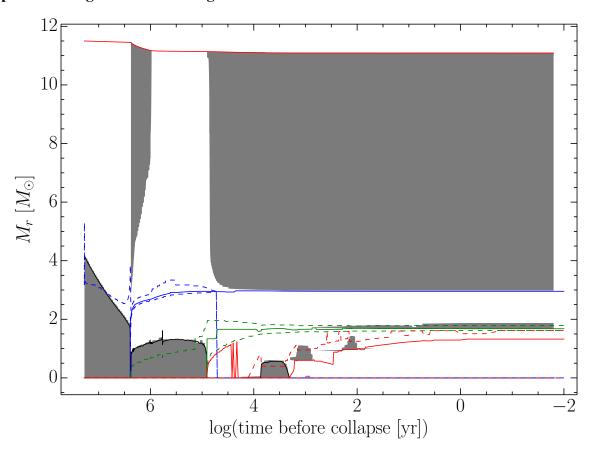
Drawing with a 3rd variable (Mcc) in colour code:

Plot_colour('Tc','Mcc')

Addition of the gas degeneracy line on the plot:

degenerate_line()

8.5 Kippenhahn diagram with burning zones



Loading of the evolution file:

loadE('/Path/to/file/P011p5z14S0.dat',1)

Labels written with LATEX:

iLatex(True)

Definition of the *x* variable:

defX('ageadv')

Drawing of the Kippenhahn diagram (predefined command):

Kippen(1,burn=True)

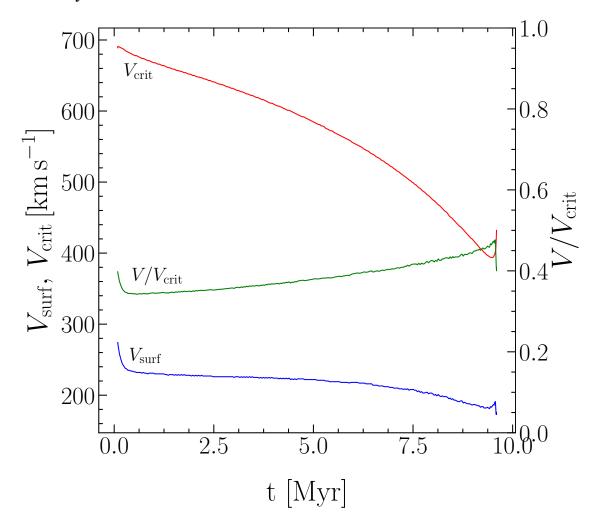
NB: the burning zones (plotted with the optional argument burn=True) are read in a #starname.burn file that must exist somewhere. This .burn file is automatically generated when the star's directory has been cleaned with the cleanfiles routine with the option -b. It can be generated afterwards on the .v file of a model with the script BurningZonesCalc.py. This script can be requested to sylvia.ekstrom@unige.ch.

8.6 Two different y axes

Loading of the evolution file:

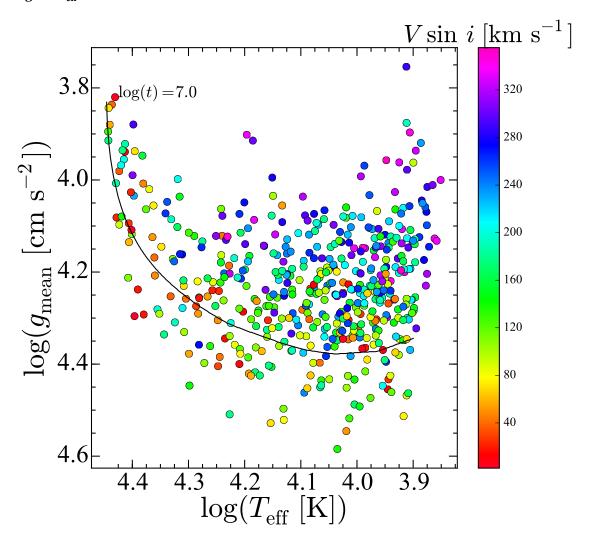
Setting the colour to blue:

loadE('/Path/to/file/P018z14S4.dat')



```
set_colourFlag('b')
Plotting the first curve, for the MS only:
      Plot('Vsurf',plotif=['H1c','>0.'])
Adding the label:
      add_label(0.37,240.,'$V_\mathrm{surf}$',fontsize=18)
Keeping the figure open for further plotting:
      keep_plot(True)
Setting the colour to red:
      set_colourFlag('r')
Plotting the second curve:
      Plot('Vcrit1',plotif=['H1c','>0.'])
Adding the label:
      add_label(0.23,640.,'$V_\mathrm{crit}$',fontsize=18)
Changing the y label:
      change\_label('y','$V\_\mathtt{mathrm}\{surf\},\,V\_\mathtt{mathrm}\{crit\}\,[\mathtt{mathrm}\{km\,s\}^{-1}]$')
Preparing for plotting on the second y axis:
      plot2var('double')
Setting the limits on this axis:
      Limits(ymin=0.,ymax=1.)
Setting the colour to green:
      set_colourFlag('g')
Plotting the last curve:
      Plot('VVc',plotif=['H1c','>0.'])
Adding the label:
      add_label(0.43,0.36,'$V/V_\mathbf{crit}$',fontsize=18)
```

8.7 Noised g vs T_{eff} with $V \sin i$ in colours



add_label(4.425,3.82,'\$\log(t)=7.0\$',fontsize=16)