AIMS Version Update (14/11/2018)

Remaining updates:

- Make selection of evolutionary state generic
- Provide specific entry criteria for this to remove duplicate functions in model.py

AIMS configure.py:

Line 36 - number of steps to be added if MCMC process does not converge.

```
35 nsteps = 200  # number of steps

36 add_steps = 500  # number of steps to add if convergence isn't achieved

37 thin = 10  # thinning parameter (1 out of thin steps will be kept ...)
```

Line 198 - detemines which evolutionary parameter is to be used.

AIMS.py:

Line 64: now import ptemcee to use instead of emcee.PTsampler

```
64 import ptemcee
```

Lines 1786-1800, *load_binary_data* - allows the user to extract information about a specific model based on mass, age and log(Z) value.

```
# for track in grid.tracks:
                      for model in track.models:
    # print model.glb
1787
                             # sys.exit()
# print(model.glb[1]/constants.solar_mass)
1789
                            if (model.glb[1] == 2.5253600000000000E+033) & (model.glb[3] == 0.01) & (model.glb[0] == 4098.5500000000002):
    print model.glb[1]/constants.solar_mass
    print model.glb
1791
1792
1793
                                   print model.gub
print model.get_freq()
model.write_file_simple('test')
print model.find_large_separation()
print model.numax
1794
1795
1796
                                   print model.FeH
1798
                                   model.print_me()
            # sys.exit()
1800
```

Lines 1930-44 (*find_best_model_in_track*): edit to the calculation of the log probability to find the best model along a track. Edit made by Gael therefore for further clarification, please ask him.

```
1930
       for model in grid.tracks[ntrack].models:
1931
          result, rc, rs, rp = prob.evaluate(model)
1932
            reject_classic += rc
1933
           reject_seismic += rs
1934
            reject_prior
           if (config.interp_type == "mHe"):
1935
1936
                istart = max(i-1,0)
                istop = min(i+1,nmodels-1)
1937
               log_slope = math.log(abs((grid.tracks[ntrack].models[istop].string_to_param("Age")
1938
                                        - grid.tracks[ntrack].models[istart].string_to_param("Age")) \
1939
                                        /(grid.tracks[ntrack].models[istop].string_to_param("mHe")
1940
1941
                                        - grid.tracks[ntrack].models[istart].string_to_param("mHe"))))
1942
1943
               log_slope = 0.0
1944
           result += log_slope
```

Run_emcee: have replaced emcee.PTsampler with the purpose specific package ptsampler. Have also included convergence criteria and looping based on whether the threshold set for the Gelman-Rubin statistic is satisfied.

Lines 2174-78 (*find_a_blob*): now provides options depending on the evolutionary parameter used.

Lines 2280-307 (*write_percentiles*): includes additional columns with the sigma values computed. Lenght of percentiles array increased accordingly.

```
2278
             # initialisation
             (m,n) = np.shape(samples)
percentiles = np.empty((n,7), dtype=np.float64)
2279
2280
2281
             for i in range(n):
                   percentiles[i,0] = np.percentile(samples[:,i],100.0*(0.5-two_sigma/2.0))
2282
                   percentiles[i,4] = np.percentile(samples[:,i],100.0*(0.5+two_stgma/2.0))
percentiles[i,1] = np.percentile(samples[:,i],100.0*(0.5+two_stgma/2.0))
percentiles[i,3] = np.percentile(samples[:,i],100.0*(0.5+one_stgma/2.0))
percentiles[i,2] = np.percentile(samples[:,i],50.0)
2283
2284
2285
2286
                    percentiles[i,5] = abs((percentiles[i,0]-percentiles[i,4])/2.)
2287
                    percentiles[i,6] = abs((percentiles[i,1]-percentiles[i,3])/2.)
2288
2289
2290
             # write results to file
             output_file = open(filename,"w")
2291
             output_file.write("Percentiles\n=======\n\n");
output_file.write("Percentiles\n=======\n\n");
output_file.write('{0:25} '.format("Quantity"))
output_file.write('{0:25} '.format("-2sigma"))
output_file.write('{0:25} '.format("Heddian"))
output_file.write('{0:25} '.format("Hisigma"))
output_file.write('{0:25} '.format("+1sigma"))
2292
2293
2294
2295
2296
2297
             output_file.write('{0:25}\n'.format("+2sigma"))
output_file.write('{0:25} '.format("1sig"))
2298
2299
             output_file.write('{0:25}\n'.format("2sig"))
2300
2301
2302
             for i in range(n):
                    output_file.write('{0:25} '.format(labels[i]))
2303
2304
                    for j in range(7):
                          output_file.write('{0:25.15e} '.format(percentiles[i,j]))
2305
                    output_file.write('\n')
2306
             output_file.close()
2307
```

Lines 2505-12 (*write_combinations*): Addition of other evolutionary parameter to perform model interpolation with.

```
output_file.write("{0:s} {1:e}\n".format(grid.prefix,constants.G))
for params in samples:
    results = model.find_combination(grid,params[0:ndims-nsurf])
    if config.interp_type == "Age":
        my_model = model.interpolate_model(grid,params[0:ndims-nsurf],grid.tessellation,grid.ndx)
    if config.interp_type == "mHe":
        my_model, slope = model.interpolate_model_mHe(grid,params[0:ndims-nsurf],grid.tessellation,grid.ndx)
```

Line 3342: changed "Age" to config.interp_type to access the relevant evolutionary parameter functions.

```
# load grid and associated quantities
grid = load_binary_data(config.binary_grid)
plot_frequencies(grid) # this will stop the program
grid_params_MCMC = grid.grid_params + (config.interp_type,)
grid_params_MCMC_with_surf = grid_params_MCMC \
```

Lines 3464-67: functions selected based on the evolutionary parameter chosen to find best MCMC model.

```
# find best MCMC model

a460 ndx_max = lnprob.argmax()

best_MCMC_result = lnprob[ndx_max,0]

best_MCMC_params = samples[ndx_max,1:]

best_MCMC_params.reshape(ndims)

a461 if config.interp_type == "Age":

best_MCMC_model = model.interpolate_model(grid,best_MCMC_params[0:ndims-nsurf],grid.tessellation,grid.ndx)

elif config.interp_type == "mHe":

best_MCMC_model, slope = model.interpolate_model_mHe(grid,best_MCMC_params[0:ndims-nsurf],grid.tessellation,grid.ndx)
```

Lines 3479-82: functions selected based on the evolutionary parameter chosen to find best statistical model.

```
# find statistical model
statistical_params = samples[:,1:].sum(axis=0, dtype=np.float64)/(1.0*config.nwalkers*config.nsteps)
statistical_params.reshape(ndims)
statistical_result = prob(statistical_params)
if config.interp_type == "Age":
    statistical_model = model.interpolate_model(grid,statistical_params[0:ndims-nsurf],grid.tessellation,grid.ndx)
elif config.interp_type == "mHe":
    statistical_model, slope = model.interpolate_model_mHe(grid,statistical_params[0:ndims-nsurf],grid.tessellation,grid.ndx)
```

model.py:

Many of the changes in this section are replications of current functions for when age
is used as the evolutionary parameter. In these cases, just the name of the function is
included as the core code remains the same, only age is changed for mHe

Series of changes to global parameter numbers. These all depend upon whether the core helium mass is being included in as a grid parameter. grid_SUN_DIFF2 does not include this, hence the '-1' included. When He-core mass is included, the '-1' disappears. imHe is included as a global parameter for all other grids used in the AIMS paper.

```
94 nglb
                   = 9 + len(config.user_params) - 1
  95 """ total number of global quantities in a model (see :py:data:`Model.glb`)."""
124 # imHe
                  = 5
125 """ index of the parameter corresponding to He core mass in the :py:data:`Model.glb` array """
127 ifreq_ref
                = 6 + len(config.user params) - 1
128
129 index of the parameter corresponding to the reference frequency
130 (used to non-dimensionalise the pulsation frequencies of the model)
131 in the :py:data: `Model.glb` array
132 "
133
134 iradius
                = 7 + len(config.user_params) - 1
135 """ index of the parameter corresponding to radius in the :py:data: `Model.glb` array """
136
137 iluminosity = 8 + len(config.user_params) - 1
138 """ index of the parameter corresponding to luminosity in the :py:data:`Model.glb` array """
```

string to latexl: Inclusion of mHe.

```
166 if (string == "mHe"): return r'Mass of He core, $%sM_{He-core}/M_{\odot}%s$'%(prefix,postfix)
```

Model:

string to param: Inclusion of mHe.

```
237 if (string == "mHe"): return self.glb[imHe]/constants.solar_mass
```

Lines 284-5: Assertion of mHe to be non-zero and positive if used as the evolutionary parameter. Value can be zero if an MS grid is used that includes the He-core mass.

Lines 511-20: function to return core helium mass.

```
511 def get_mHe(self):
```

Line 897 (*print_me*): Allows for direct inclusion of core helium mass in model properties printed to the screen.

```
# print "Mass He core (in M_Sun): %.4f" % self.glb[imHe]
```

Track:

Line 1214: Beginning of the core helium mass functions in Track class. What follows are the function names unless there are any major changes to the included code.

```
1214
            ''' Mass of He core versions '''
        def sort mHe(self):
1215
1220
        def is_sorted_mHe(self):
1231
        def duplicate_mHe(self):
        def interpolate_model_mHe(self, mHe):
1253
        def find_combination_mHe(self, mHe, coef):
1291
        def find modes mHe(self, ntarget, ltarget):
1326
        def test_interpolation_mHe(self, nincr):
1353
```

Model_grid:

Lines 1494-1496 (*read_model_list*): hard coded inclusion of imHe to global parameter list and extension of the array length for the addition of any other parameters.

```
1494  # glb[imHe] = utilities.to_float(columns[8])
1495
1496  i = 9 - 1
```

Lines 1527-32: Sort tracks depending upon the evolutionary parameter selected.

```
# sort tracks:
for track in self.tracks: track.sort()
for track in self.tracks: track.sort
for track in self.tracks: track.sort_age()
for track in self.tracks: track.sort_mHe()
for track in self.tracks: track.sort_mHe()
```

 Test_interpolation (line 1668) contains multiple changes to allow original model and interpolated model properties to be saved out. These are currently commented out in the code, but can be reinstated accordingly.

```
# initialisation
1685
                           results = []
1686
1687
                          ndim = self.ndim+1
1688
                           # print ndim
                          # output_folder = '/home/buldgen/AIMS-master_New/AIMS_BEN/'
1689
1690
                         # filename = os.path.join(output_folder,"combinations_Delaunay.txt")
                        # f2 = os.path.join(output_folder,"Delaunay_MS_Models_mHe.txt")
1691
                        # f3 = os.path.join(output_folder,"Delaunay_MS_Mod_vals_mHe.txt")
1692
1693
                        # output_file = open(filename,"w")
                         # out2 = open(f2,"w")
# out3 = open(f3,"w")
1694
1695
1606
1702
                         for i in range(nmodels):
                                aModel1 = self.tracks[j].models[i]
if config.interp_type == "Age":
1703
1704
                                       pt[-1] = aModel1.glb[iage]
                                        aModel2 = interpolate_model(self,pt,tessellation,ndx2) #,aModel1.name,aModel1,out2,out3)
1706
1707
                                elif config.interp_type == "mHe":
                                       pt[-1] = aModel1.glb[imHe]
aModel2, slope = interpolate_model_mHe(self,pt,tessellation,ndx2)
1708
1709
1710
                                aResult[i,0:ndim] = pt
1711
                              if (res is None): continue # filter out combinations outside the grid
# output_file.write("{0:d} {1:.2f} {2:.3f} {3:.3f} {4:.4f} {5:.3f} {6:.3f} {7:.2f}\n".format( \
# len(res), aModel1.glb[imass]/constants.solar_mass, aModel1.glb[iradius]/constants.solar_radius, \
# aModel1.glb[iuminosity]/constants.solar_luminosity, aModel1.glb[iz0], \
# aModel1.glb[ix0], aModel1.glb[iage], \
# aModel1.glb[ix0], aModel1.glb[imHe]/constants.solar_mass))
1722
1723
1724
1725
1726
1727
                  # aModel2.glb[iuminosity]/constants.solar_mass, aModel2.glb[iradius],
# aModel2.glb[iuminosity]/constants.solar_luminosity,aModel2.glb[iz0], \
# aModel2.glb[ix0],aModel2.glb[iaga], \
# aModel2.glb[itemperature], aModel1.glb[imHe]/constants.solar_mass))
# for (coef,model_name) in res:
# output_file.write("{0:.15f} {1:s}\n".format(coef, model_name))
# output_file.write("\n")
results.append(aResult)
                               " output_file.write("\{0:d\} \{1:.2f\} \{2:.3f\} \{3:.3f\} \{4:.4f\} \{5:.3f\} \{6:.3f\} \{7:.2f\}\n".format( \ # len(res), aModel2.glb[imass]/constants.solar_mass, aModel2.glb[iradius]/constants.solar_radius, \
1730
1731
1732
1733
1734
               # output_file.close()
# out2.close()
# out3.close()
1737
```

(sorry it kept getting smaller)

Final interpolation copy-cats for mHe:

```
2262 def find_mHes(coefs, tracks, mHe):
2263
        Find mHes to which each track needs to be interpolated for a specified
2264
2265
        mHe. Follows the same structure as find_ages() and uses the same scaled and
2266
        fixed parameter for determining the method used to calculate the values.
2267
2293 def interpolate_model_mHe(grid,pt,tessellation,ndx):
2294
          Interpolate model in grid using provided parameters.
2295
2365 def find combination mHe(grid,pt):
2366
         Find linear combination of models which corresponds to interpolating
2367
2368
         the model based on the provided parameters.
```