# **hmf Documentation**

Release 3.0.2

**Steven Murray** 

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# The halo mass function calculator.



*hmf* is a python application that provides a flexible and simple way to calculate the Halo Mass Function for a range of varying parameters. It is also the backend to HMFcalc, the online HMF calculator.

**Warning:** Due to the general trend of moving to Python 3 by important projects such as IPython and astropy, from version 3.0, hmf is compatible with Python 3, and from version 3.1, it will drop (official) support for Python 2.

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Documentation

Read the docs.

CHAPTER	2
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Attribution

Please cite Murray, Power and Robotham (2013) if you find this code useful in your research.

# CHAPTER 3

# **Features**

- Calculate mass functions and related quantities extremely easily.
- Very simple to start using, but wide-ranging flexibility.
- Caching system for optimal parameter updates, for efficient iteration over parameter space.
- Support for all LambdaCDM cosmologies.
- Focus on flexibility in models. Each "Component", such as fitting functions, filter functions, growth factor models and transfer function fits are implemented as generic classes that can easily be altered by the user without touching the source code.
- Focus on simplicity in frameworks. Each "Framework" mixes available "Components" to derive useful quantities all given as attributes of the Framework.
- Comprehensive in terms of output quantities: access differential and cumulative mass functions, mass variance, effective spectral index, growth rate, cosmographic functions and more.
- Comprehensive in terms of implemented Component models
  - 5+ models of transfer functions including directly from CAMB
  - 4 filter functions
  - 20 hmf fitting functions
- Includes models for Warm Dark Matter
- Nonlinear power spectra via HALOFIT
- Functions for sampling the mass function.
- CLI scripts both for producing any quantity included, or fitting any quantity.
- Python 2 and 3 compatible

8 Chapter 3. Features

# CHAPTER 4

# Installation

*hmf* is built on several other packages, most of which will be familiar to the scientific python programmer. All of these dependencies should be automatically installed when installing *hmf*. Explicitly, the dependencies are numpy, scipy, astropy and camb.

You will only need *emcee* if you are going to be using the fitting capabilities of *hmf*.

Finally the *hmf* package needs to be installed: pip install hmf.

To go really bleeding edge, install the develop branch using pip install git+git://github.com/steven-murray/hmf.git@develop.

# CHAPTER 5

Quickstart

Once you have *hmf* installed, you can quickly generate a mass function by opening an interpreter (e.g. IPython) and doing:

```
>>> from hmf import MassFunction
>>> hmf = MassFunction()
>>> mass_func = hmf.dndlnm
```

Note that all parameters have (what I consider reasonable) defaults. In particular, this will return a Tinker (2008) mass function between  $10^{10}-10^{15}M_{\odot}$ , at z=0 for the default PLANCK15 cosmology. Nevertheless, there are several parameters which can be input, either cosmological or otherwise. The best way to see these is to do

```
>>> MassFunction.parameter_info()
```

We can also check which parameters have been set in our "default" instance:

```
>>> hmf.parameter_values
```

To change the parameters (cosmological or otherwise), one should use the *update()* method, if a MassFunction() object already exists. For example

```
>>> hmf = MassFunction()
>>> hmf.update(Ob0 = 0.05, z=10) #update baryon density and redshift
>>> cumulative_mass_func = hmf.ngtm
```

For a more involved introduction to *hmf*, check out the tutorials, which are currently under construction, or the API docs.

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14 Chapter 6. Author

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# Comments, corrections and suggestions

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# CHAPTER 9

Contents

# 9.1 Usage and Tutorials

One way to pick up how to use hmf is to directly consult the API documentation.

Here, however, we have compiled several more high-level resources on how to get started with hmf, and use it efficiently.

### 9.1.1 Dealing with Cosmological Models

hmf uses the robust astropy cosmology framework to deal with cosmological models. This provides a range of cosmographic functionality for free.

Cosmological models are the most basic Framework within hmf. Every other Framework depends on it. So knowing how to specify the models is important (but very simple!).

```
from hmf import cosmo
%matplotlib inline
import matplotlib.pyplot as plt
import numpy as np
```

## **Default Settings**

Like everything in hmf, the Cosmology framework has all parameters specified with defaults. In this case, there are only two parameters – a base cosmological model, and a dictionary of cosmological parameters with which to alter it. By default, the cosmological model is a Flat LambdaCDM model infused with the Planck15 parameters. The dictionary is empty, so we don't modify anything:

```
my_cosmo = cosmo.Cosmology()
```

The intrinsic astropy object is found as the cosmo attribute of the class we just created. Beware, there is also a cosmo\_model attribute, which should only be treated as a parameter, never used in calculations. It has not been supplemented with any custom parameters. We can check out the parameters defined within the model:

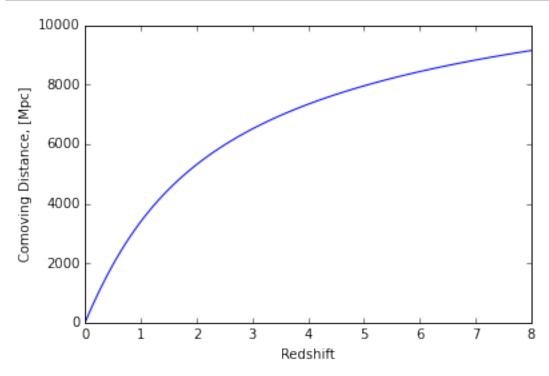
```
print "Matter density: ", my_cosmo.cosmo.Om0
print "Hubble constant: ", my_cosmo.cosmo.H0
print "Dark Energy density: ", my_cosmo.cosmo.Ode0
print "Baryon density: ", my_cosmo.cosmo.Ob0
print "Curvature density: ", my_cosmo.cosmo.Ok0
```

```
Matter density: 0.3075
Hubble constant: 67.74 km / (Mpc s)
Dark Energy density: 0.691009934459
Baryon density: 0.0486
Curvature density: 0.0
```

Or we can check out some cosmographic quantities, like the comoving distance as a function of redshift:

```
z = np.linspace(0,8,100)
plt.plot(z,my_cosmo.cosmo.comoving_distance(z))
plt.ylabel("Comoving Distance, [Mpc]")
plt.xlabel("Redshift")
```





### Passing a cosmological model

The cosmo module contains several pre-made instances of cosmologies which might be useful, which we can input as our default model:

```
my_cosmo = cosmo.Cosmology(cosmo_model=cosmo.WMAP5)
print "WMAP5 baryon density: ", my_cosmo.cosmo.Ob0
```

```
WMAP5 baryon density: 0.0459
```

Alternatively, we can create our own. The astropy package contains the basic tools to do this. To create a standard Flat LambdaCDM cosmology:

```
from astropy.cosmology import FlatLambdaCDM
new_model = FlatLambdaCDM(H0 = 75.0, Om0=0.4, Tcmb0 = 5.0, Ob0 = 0.3)
```

This new model can be used as input to the Cosmology class:

```
my_cosmo = cosmo.Cosmology(cosmo_model = new_model)
print "Crazy cosmology baryon density: ", my_cosmo.cosmo.Ob0
```

```
Crazy cosmology baryon density: 0.3
```

The cosmo\_model needn't be a Flat LambdaCDM. It can be any subclass of FLRW. Thus we could use a non-flat model:

```
from astropy.cosmology import LambdaCDM
new_model = LambdaCDM(H0 = 75.0, Om0=0.4, Tcmb0 = 0.0, Ob0 = 0.3, Ode0=0.4)

my_cosmo = cosmo.Cosmology(cosmo_model = new_model)
print "Crazy cosmology curvature density: ", my_cosmo.cosmo.Ok0
```

```
Crazy cosmology curvature density: 0.2
```

### Passing custom parameters

Instead of passing a pre-made cosmological model, you can pass custom parameters for the default model. This is passed as a dictionary, in which each entry is a valid parameter for the model that has been passed (i.e., if the model is a FlatLambdaCDM, you can't pass <code>Ode0!</code>). This means you can specify the cosmology you want typically in one line, rather than a few. It also means that parameters can be updated in a standard way, so that iterating over parameters, in applications such as fitting models, becomes simple.

When passing the dictionary of parameters, you don't need to specify them all, just whichever ones you want to modify:

```
my_cosmo = cosmo.Cosmology(cosmo_params={"Om0":0.2})
print "Custom cosmology matter density: ", my_cosmo.cosmo.Om0
```

```
Custom cosmology matter density: 0.2
```

New parameters are available for extended cosmological models:

```
my_cosmo = cosmo.Cosmology(new_model, {"Om0":0.2,"Ode0":0.0,"Ob0":0.2})
print "Custom cosmology curvature density: ", my_cosmo.cosmo.Ok0
```

```
Custom cosmology curvature density: 0.8
```

### **Updating parameters**

One of the great things about hmf Frameworks is that any parameter can be updated without re-creating the entire object. This is also true of the Cosmology class.

Any parameter passed to the constructor may also be updated:

```
my_cosmo = cosmo.Cosmology(new_model)
my_cosmo.update(cosmo_params={"Om0":0.2,"Ode0":0.0,"Ob0":0.2})
print "Custom cosmology curvature density: ", my_cosmo.cosmo.Ok0
```

```
Custom cosmology curvature density: 0.8
```

The parameter dictionary is persistent, so that updating a different parameter doesn't affect the others:

```
my_cosmo.update(cosmo_params={"H0":10.0})
print "Custom cosmology curvature density: ", my_cosmo.cosmo.Ok0
print "Custom parameters: ", my_cosmo.cosmo_params
```

```
Custom cosmology curvature density: 0.8
Custom parameters: {'H0': 10.0, 'Om0': 0.2, 'Ode0': 0.0, 'Ob0': 0.2}
```

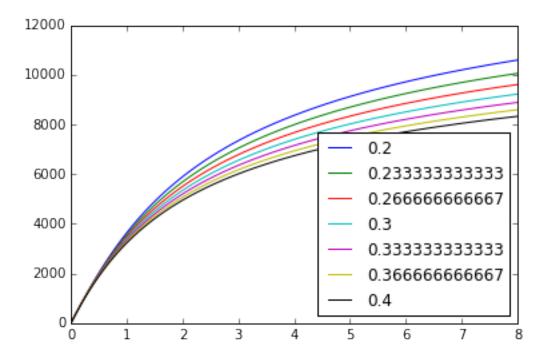
Of course, if we were to update the model to a Flat Lambda CDM model, then the OdeO keyword would give an error. To facilitate this, passing an empty dictionary clears all custom values:

```
my_cosmo.update(cosmo_model=cosmo.Planck13,cosmo_params={})
print "Flat cosmology curvature density: ", my_cosmo.cosmo.Ok0
```

```
Flat cosmology curvature density: 0.0
```

In effect, this gives us an easy way to track changes induced by a cosmological variable:

```
for Om0 in np.linspace(0.2,0.4,7):
    my_cosmo.update(cosmo_params={"Om0":Om0})
    plt.plot(z,my_cosmo.cosmo.comoving_distance(z),label="%s"%Om0)
    = plt.legend(loc=0)
```



### 9.1.2 Mass Definitions

hmf, as of v3.1, provides a simple way of converting between halo mass definitions, which serves two basic purposes:

- 1. the mass of a halo in one definition can be converted to its appropriate mass under another definition
- 2. the halo mass function can be converted between mass definitions.

### Introduction

By "mass definition" we mean the way the extent of a halo is defined. In hmf, we support two main kinds of definition, which themselves can contain different models. In brief, hmf supports Friends-of-Friends (FoF) halos, which are defined via their linking length, b, and spherical-overdensity (SO) halos, which are defined by two criteria: an overdensity  $\Delta_h$ , and an indicator of what that overdensity is with respect to (usually mean background density  $\rho_m$ , or critical density  $\rho_c$ ). In addition to being able to provide a precise overdensity for a SO halo, hmf provides a way to use the so-called "virial" overdensity, as defined in Bryan and Norman (1998).

Converting between SO mass definitions is relatively simple: given a halo profile and concentration for the given halo mass, determine the concentration required to make that profile contain the desired density, and then compute the mass of the halo under such a concentration.

There is no clear way to perform such a conversion for FoF halos. Nevertheless, if one assumes that all linked particles are the same mass, and that halos are spherical and singular isothermal spheres (cf. White (2001)), one can approximate an FoF halo by an SO halo of density  $9\rho_m/(2\pi b^3)$ . hmf will make this approximation if a conversion between mass definitions is desired.

### **Changing Mass Definitions**

Most of the functionality concerning mass definitions is defined in the hmf.halos.mass\_definitions module:

import inspect

Different mass definitions exist inside the mass\_definitions module as Components. All definitions are subclassed from the abstract MassDefinition class:

### **The Mass Definition Component**

To create an instance of any class, optional cosmo and z arguments can be specified. By default, these are the Planck15 cosmology at redshift 0. We'll leave them as default for this example. Let's define two mass definitions, both spherical-overdensity definitions with respect to the mean background density:

```
In [4]: mdef_1 = md.SOMean(overdensity=200)
    mdef_2 = md.SOMean(overdensity=500)
```

Each mass definition has its own model\_parameters, which define the exact overdensity. For both the SOMean and SOCritical definitions, the overdensity can be provided, as above (default is 200). This must be passed as a named argument. For the FOF definition, the linking\_length can be passed (default 0.2). For the SOVirial, no parameters are available. Available parameters and their defaults can be checked the same way as *any* Component within hmf:

```
In [5]: md.SOMean._defaults
Out[5]: {'overdensity': 200}
```

The explicit halo density for a given mass definition can be accessed:

```
In [6]: mdef_1.halo_density, mdef_1.halo_density/mdef_1.mean_density
Out[6]: (17068502575484.857, 200.0)
```

### **Converting Masses**

To convert a mass in one definition to a mass in another, use the following:

The input mass argument can be a list or array of masses also. To convert between masses, the concentration of the input halos, and their density profile, must be known. By default, an NFW profile is assumed, with a concentration-mass relation from Duffy et al. (2008).

One can alternatively pass a concentration directly:

If you have halomod installed, you can also pass any halomod.profiles.Profile instance (which itself includes a concentration-mass relation) as the profile argument.

### **Converting Mass Functions**

All halo mass function fits are measured using halos found using some halo definition. While some fits explicitly include a parameterization for the spherical overdensity of the halo (eg. Tinker08 and Watson), others do not.

By passing a mass definition to the MassFunction constructor, hmf will attempt to convert the mass function defined by the chosen fitting function to the appropriate mass definition, by solving

$$\int_{\infty}^{m_{\text{old}}} dm \ n(m) = \int_{\infty}^{m_{\text{new}}(m_{\text{old}})} dm \ n'(m)$$

$$(9.1)$$

for n'(m), resulting in

FoF 0.2

$$n'(m) = n(m_{\text{old}}(m_{\text{new}})) \left(\frac{dm_{\text{new}}}{dm_{\text{old}}}\right)^{-1}.$$
(9.2)

By default, the mass definition is None, which turns off all mass conversion and uses whatever mass definition was employed by the chosen fit. This keeps all existing code running as it was previously. Nevertheless, care should be taken here: many of the different fits have different mass definitions, and should not be compared without some form of mass conversion. To see the mass definition intrinsic to the measurement of each fit, use the following:

Here "vir" corresponds to the virial definition of Bryan and Norman (1998), an asterisk indicates that the fit is itself parameterized for SO mass definitions, and 0.2 is the FoF linking length.

Let's convert the mass definition of Tinker08, which has an intrinsic parameterization:

```
In [21]: # Default mass function object
    mf = hmf.MassFunction()

    dndm0 = mf.dndm

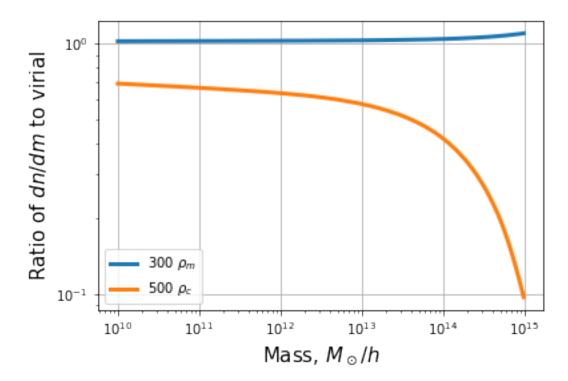
# Change the mass definition to 300rho_mean
    mf.update(
```

```
mdef_model = "SOMean",
             mdef_params = {
                 "overdensity": 300
        )
        plt.plot(mf.m, mf.dndm/dndm0, label=r"300 $\rho_m$", lw=3)
        # Change the mass definition to 500rho_crit
        mf.update(
             mdef_model = "SOCritical",
             mdef_params = {
                 "overdensity": 500
        plt.plot(mf.m, mf.dndm/dndm0, label=r"500 $\rho_c$", lw=3)
        plt.xscale('log')
        plt.yscale('log')
        plt.xlabel(r"Mass, $M_\odot/h$", fontsize=15)
        plt.ylabel(r"Ratio of $dn/dm$ to $200 \rho_m$", fontsize=15)
        plt.grid(True)
        plt.legend();
      10°
Ratio of dn/dm to 200\rho_m
                   300 \rho_{m}
     10^{-1}
                                     1012
                                                  1013
            1010
                         1011
                                                              10^{14}
                                                                           1015
                                     Mass, M_{\odot}/h
```

This did not require any internal conversion. Let's try converting a fit that has no explicit parameterization for overdensity:

```
In [27]: # Default mass function object.
    mf = hmf.MassFunction(hmf_model = "SMT", Mmax=15)
    dndm0 = mf.dndm
# Change the mass definition to 300rho_mean
```

```
mf.update(
             mdef_model = "SOMean",
             mdef_params = {
                 "overdensity": 300
         plt.plot(mf.m, mf.dndm/dndm0, label=r"300 $\rho_m$", lw=3)
         # Change the mass definition to 500rho_crit
         mf.update(
             mdef_model = "SOCritical",
             mdef_params = {
                 "overdensity": 500
         plt.plot(mf.m, mf.dndm/dndm0, label=r"500 $\rho_c$", lw=3)
        plt.xscale('log')
        plt.yscale('log')
        plt.xlabel(r"Mass, $M_\odot/h$", fontsize=15)
        plt.ylabel(r"Ratio of $dn/dm$ to virial", fontsize=15)
        plt.grid(True)
        plt.legend();
332.2445055922226 0.0
300
```



Here, the measured mass function uses "virial" halos, which have an overdensity of  $\sim 330 \rho_m$ , so that converting to  $300 \rho_m$  is a down-conversion of density. Finally, we convert a FoF mass function:

```
In [33]: # Default mass function object.
    mf = hmf.MassFunction(hmf_model = "Jenkins")
    dndm0 = mf.dndm
```

```
# Change the mass definition to 300rho_mean
        mf.update(
            mdef_model = "FOF",
            mdef_params = {
                 "linking_length": 0.3
        plt.plot(mf.m, mf.dndm/dndm0, label=r"FoF $b=0.3$", lw=3)
        # Change the mass definition to 500rho_crit
        mf.update(
            mdef_model = "SOVirial",
            mdef_params = {} # NOTE: we need to pass an empty dict here
                                  # so that the "linking_length" argument is removed.
        plt.plot(mf.m, mf.dndm/dndm0, label=r"SO Virial", lw=3)
        plt.xscale('log')
        plt.yscale('log')
        plt.xlabel(r"Mass, $M_\odot/h$", fontsize=15)
        plt.ylabel(r"Ratio of $dn/dm$ to FoF $b=0.2$", fontsize=15)
        plt.grid(True)
        plt.legend();
Ratio of dn/dm to FoF b = 0.2
                      FoF b = 0.3
                     SO Virial
     2 \times 10^{0}
         10°
    6 \times 10^{-1}
               1010
                           1011
                                        1012
                                                    1013
                                                                 10^{14}
                                                                             1015
```

# 9.2 License

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Mass,  $M_{\odot}/h$ 

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# 9.3 API Summary

hmf.cosmology.cosmo
hmf.cosmology.growth_factor
hmf.density_field.transfer
hmf.density_field.transfer_models
hmf.density_field.halofit
hmf.density_field.filters
hmf.halos.mass_definitions
hmf.mass_function.hmf
hmf.mass_function.fitting_functions
hmf.mass_function.integrate_hmf
hmf.alternatives.wdm
hmf.helpers.sample
hmf.helpers.functional

### 9.4 Releases

### 9.4.1 dev

#### **Features**

- Added new CambGrowth growth factor model, which computes the growth using CAMB. This is useful especially when using w > -1, for which the other growth factor models are inadequate. Solves issue #19 raised by @tijmen.
- Added new module mass\_definitions which more robustly deals with various halo mass definitions, and also includes ability to convert mass functions between different definitions.

### **Enhancement**

• Added *get\_dependencies* method to \_Framework, to enable finding all parameters that a quantity depends on.

### **Bugfixes**

• When using camb for the transfer function, some cosmologies would lead to a segfault (i.e. when Ob0 or Tcmb0 are not set explicitly). This now raises a helpful error.

#### **Internals**

• Removed logging, which was redundant.

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- Moved from nose to pytest
- Significant overhaul of package structure to more modularised form.

### 9.4.2 v3.0.3 [1st Dec 2017]

### **Bugfixes**

- · Fixed usage of deprecated MsolMass in wdm
- Fixed Bhattachrya fitting function (thanks to Benedikt Diemer!)
- Fixed typo in Watson fitting function (thanks to Benedikt Diemer!)
- Update cosmo test to use new Astropy constants.
- Fixed issue with sampling function where zeros in ngtm would yield an error.

# 9.4.3 v3.0.2 [3rd Nov 2017]

### **Bugfixes**

• Changed parameter checks on instantiation to post-conversion.

### 9.4.4 v3.0.1 [31st Oct 2017]

### **Enhancement**

- Normalised all <>\_model properties to be actual classes, rather than either class or string.
- Added consistent checking of dictionaries for <>\_params parameters.

### 9.4.5 v3.0.0 [7th June 2017]

### **Features**

- Now provides compatibility with Python 3.x. Support for 2.x will be removed in hmf v3.1 (whenever that comes).
- Complete overhaul of the caching system. Should be invisible to the user, but streamlines writing of framework code considerably. Removes required manual specification of dependencies between quantities, and adds ability to specify parameter kinds (model, param, res, option, switch).

### **Bugfixes**

- Fixed bug in Caroll1992 GrowthFactor class which affected high-redshift growth factors (thanks to Olmo Piana).
- Fixed astropy dependency to be >= 1.1
- Fixed bug where Takahashi parameters were always passed through regardess of takahashi setting.
- Fixed small bug where the functional get\_label method returned differently ordered parameters because of dicts.
- Note that the fitting subpackage is temporarily unsupported and I discourage its use for the time being.

### Enhancement

• Completely removes dependence on archaic pycamb package. Now supports natively supplied python interface to CAMB. Install camb with pip install --egg camb. This means that much more modern versions of CAMB can be used.

• Many new tests, to bring total coverage up to >80%, and continuous testing on Python 2.7, 3.5 and 3.6

# 9.4.6 v2.0.5 [12th January 2017]

### **Bugfixes**

• Fixed bug in GrowthFactor which gave ripples in functions of z when a coarse grid was used. Thanks to @mirochaj and @thomasguillet!

#### **Enhancments**

- Streamlined the caching framework a bit (removing cruft)
- Totally removed dependency on the Cache (super)class caching indexes now inherent to the called class.
- More robust parameter information based on introspection.

# 9.4.7 v2.0.4 [11th November, 2016]

### **Bugfixes**

• IMPORTANT: Fixed a bug in which updating the cosmology after creation did not update the transfer function.

# 9.4.8 v2.0.3 [22nd September, 2016]

### **Bugfixes**

• SharpK filter integrated over incorrect range of k, now fixed.

#### **Enhancments**

- WDM models now more consistent with MassFunction API.
- Better warning in HALOFIT module when derivatives don't work first time.

### 9.4.9 v2.0.2 [2nd August, 2016]

### **Features**

- Added a bunch of information to each hmf\_model, indicating simulation parameters from which the fit was derived.
- Added FromArray transfer model, allowing an array to be passed programmatically for k and T.
- Added Carroll1992 growth factor approximation model.

#### **Enhancments**

• how\_big now gives the boxsize required to expect at least one halo above m in 95% of boxes.

### **Bugfixes**

- Removed unnecessary multiplication by 1e6 in cosmo.py (thanks @iw381)
- IMPORTANT: normalisation now calculated using convergent limits on k, rather than user-supplied values.
- IMPORTANT: fixed bug in Bhattacharya fit, which was multiplying by an extra delta\_c/sigma.
- fixed issue with nonlinear mass raising exception when mass outside given mass range.

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## 9.4.10 v2.0.1 [2nd May, 2016]

### **Bugfixes**

• Corrects the default sigma\_8 and n (spectral index) parameters to be from Planck15 (previously from Planck13), which corresponds to the default cosmology. **NOTE:** this will change user-code output silently unless sigma\_8 and n are explicitly set.

### 9.4.11 v2.0.0

v2.0.0 is a (long overdue) major release with several backward-incompatible changes. There are several major features still to come in v2.1.0, which may again be backward incompatible. Though this is not ideal (ideally backwards-incompatible changes will be restricted to increase in the major version number), this has been driven by time constraints.

Known issues with this version, to be addressed by the next, are that both scripts (hmf and hmf-fit) are buggy and untested. Don't use these until the next version unless you're crazy.

#### **Features**

- New methods on all frameworks to list all parameters, defaults and current values.
- New general structure for Frameworks and Components makes for simpler delineation and extensibility
- New growth\_factor module which adds extensibility to the growth factor calculation
- New transfer\_models module which separates the transfer models from the general framework
- New Component which can alter dn/dm in WDM via ad-hoc adjustment
- Added a Prior() abstract base class to the fitting routines
- Added a guess() method to fitting routines
- Added ll() method to Prior classes for future extendibility
- New fit from Ishiyama+2015, Manera+2010 and Pillepich+2009

### **Enhancments**

- Removed nz and z2 from MassFunction. They will return in a later version but better.
- Improved structure for FittingFunction Component, with *cutmask* property defining valid mass range. NOTE: the default MassFunction is no longer to mask values outside the valid range. In fact, the parameter cut\_fit no longer exists. One can achieve the effect by accessing a relevant array as dndm[MassFunction.hmf.cutmask]
- Renamed some parameters/quantities for more consistency (esp. M -> m)
- No longer dependent on cosmolopy, but rather uses Astropy (v1.0+)
- mean\_dens now mean\_density0, as per Astropy
- Added exception to catch when dndm has many NaN values in it.
- · Many more tests
- Made the cosmo class pickleable by cutting out a method and using it as a function instead.
- Added normalise() to Transfer class.
- Updated fit.py extensively, and provided new example config files
- Send arbitrary kwargs to downhill solver
- New internal \_utils module provides inheritable docstrings

### **Bugfixes**

- fixed problem with \_gtm method returning nans.
- fixed simple bugs in BBKS and BondEfs transfer models.
- fixes in \_cache module
- simple bug when updating sigma\_8 fixed.
- Made the EnsembleSampler object pickleable by setting \_\_getstate\_\_
- · Major bug fix for EH transfer function without BAO wiggles
- @parameter properties now return docstrings

# 9.4.12 v1.8.0 [February 2, 2015]

#### **Features**

- Better WDM models
- Added SharpK and SharpKEllipsoid filters and overhauled filter system.

#### **Enhancments**

- · Separated WDM models from main class for extendibility
- · Enhanced caching to deal with subclassing

### **Bugfixes**

· Minor bugfixes

# 9.4.13 1.7.1 [January 28, 2015]

### **Enhancments**

• Added warning to docstring of \_dlnsdlnm and n\_eff for non-physical oscillations.

# 9.4.14 1.7.0 [October 28, 2014]

#### **Features**

- Very much updated fitting routines, in class structure
- Made fitting functions more flexible and model-like.

### **Enhancments**

- Modified get\_hmf to be more general
- Moved get\_hmf and related functions to "functional.py"

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# 9.4.15 1.6.2 [September 16, 2014]

#### **Features**

• New HALOFIT option for original co-oefficients from Smith+03

### **Enhancments**

- Better Singleton labelling in get\_hmf
- Much cleaning of mass function integrations. New separate module for it.
- IMPORTANT: Removal of nltm routine altogether, as it is inconsistent.
- IMPORTANT: mltm now called rho ltm, and mgtm called rho gtm
- IMPORTANT: Definition of rho\_ltm now assumes all mass is in halos.
- Behroozi-specific modifications moved to Behroozi class
- New property hmf which is the actual class for mf\_fit

### **Bugfixes**

- Fixed bug in Behroozi fit which caused an infinite recursion
- Tests fixed for new cumulants.

## 9.4.16 1.6.1 [September 8, 2014]

### **Enhancments**

· Better get hmf function

### **Bugfixes**

- Fixed "transfer" property
- Updates fixed for transfer\_fit
- · Updates fixed for nu
- Fixed cache bug where unexecuted branches caused some properties to be misinterpreted
- Fixed bug in CAMB transfer options, where defaults would overwrite user-given values (introduced in 1.6.0)
- Fixed dependence on transfer\_options
- Fixed typo in Tinker10 fit at z = 0

# 9.4.17 1.6.0 [August 19, 2014]

### **Features**

• New Tinker10 fit (Tinker renamed Tinker08, but Tinker still available)

### **Enhancments**

- Completely re-worked caching module to be easier to code and faster.
- Better Cosmology class more input combinations available.

### **Bugfixes**

· Fixed all tests.

### 9.4.18 1.5.0 [May 08, 2014]

### **Features**

• Introduced \_cache module: Extracts all caching logic to a separate module which defines decorators – much simpler coding!

# 9.4.19 1.4.5 [January 24, 2014]

#### **Features**

- Added get\_hmf function to tools.py easy iteration over models!
- Added hmf script which provides cmd-line access to most functionality.

#### **Enhancments**

- · Added Behroozi alias to fits
- Changed kmax and k\_per\_logint back to have transfer\_\_ prefix.

### **Bugfixes**

- Fixed a bug on updating delta\_c
- Changed default kmax and k\_per\_logint values a little higher for accuracy.

# 9.4.20 1.4.4 [January 23, 2014]

### **Features**

• Added ability to change the default cosmology parameters

### **Enhancments**

• Made updating Cosmology simpler.

### **Bugfixes**

- Fixed a bug in the Tinker function (log was meant to be log10): thanks to Sebastian Bocquet for pointing this out!
- Fixed a bug in updating n and sigma\_8 on their own (introduced in 1.4.0)
- Fixed a bug when using a file for the transfer function.

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# 9.4.21 1.4.3 [January 10, 2014]

### **Bugfixes**

· Changed license in setup

# 9.4.22 1.4.2 [January 10, 2014]

### **Enhancments**

- · Mocked imports of cosmolopy for setup
- · Cleaner imports of cosmolopy

# 9.4.23 1.4.1 [January 10,2014]

### **Enhancments**

• Updated setup requirements and fixed a few tests

# 9.4.24 1.4.0 [January 10, 2014]

### **Enhancments**

- Upgraded API once more: Now Perturbations -> MassFunction
- Added transfer.py which handles all k-based quantities
- Upgraded docs significantly.

# 9.4.25 1.3.1 [January 06, 2014]

### **Bugfixes**

• Fixed bug in transfer read-in introduced in 1.3.0

# 9.4.26 1.3.0 [January 03, 2014]

#### **Enhancments**

- A few more documentation updates (especially tools.py)
- Removed new\_k\_bounds function from tools.py
- Added w parameter to cosmolopy dictionary in cosmo.py
- · Changed cosmography significantly to use cosmolopy in general
- Generally tidied up some of the update mechanisms.
- API CHANGE: cosmography.py no longer exists I've chosen to utilise cosmolopy more heavily here.
- Added Travis CI usage

### **Bugfixes**

- Fixed a pretty bad bug where updating h/H0 would crash the program if only one of omegab/omegac was updated alongside it
- Fixed a compatibility issue with older versions of numpy in cumulative functions

# 9.4.27 1.2.2 [December 10, 2013]

### **Bugfixes**

• Bug in "EH" transfer function call

# 9.4.28 1.2.1 [December 6, 2013]

### **Bugfixes**

• Small bugfixes to update() method

### 9.4.29 1.2.0 [December 5, 2013]

### **Features**

• Addition of cosmo module, which deals with the cosmological parameters in a cleaner way

#### **Enhancments**

- Major documentation overhaul most docstrings are now in Sphinx/numpydoc format
- Some tidying up of several functions.

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## 9.4.30 1.1.10 [October 29, 2013]

**Enhancements** - Better updating – checks if update value is actually different. - Now performs a check to see if mass range is inside fit range.

### **Bugfixes**

• Fixed bug in mltm property

# 9.4.31 1.1.9 [October 4, 2013]

### **Bugfixes**

• Fixed some issues with n(<m) and M(<m) causing them to give NaN's

# 9.4.32 1.1.85 [October 2, 2013]

### **Enhancments**

• The normalization of the power spectrum now saved as an attribute

# 9.4.33 1.1.8 [September 19, 2013]

### **Bugfixes**

· Fixed small bug in SMT function which made it crash

### 9.4.34 1.1.7 [September 19, 2013]

#### **Enhancments**

- Updated "ST" fit to "SMT" fit to avoid confusion. "ST" is still available for now.
- Now uses trapezoid rule for integration as it is faster.

# 9.4.35 1.1.6 [September 05, 2013]

### **Enhancments**

• Included an option to use delta\_halo as compared to critical rather than mean density (thanks to A. Vikhlinin and anonymous referree)

### **Bugfixes**

- Couple of bugfixes for fitting\_functions.py
- Fixed mass range of Tinker (thanks to J. Tinker and anonymous referee for this)

# 9.4.36 1.1.5 [September 03, 2013]

### **Enhancments**

-Added a whole suite of tests against genmf that actually work

### **Bugfixes**

- Fixed bug in mgtm (thanks to J. Mirocha)
- Fixed an embarrassing error in Reed07 fitting function
- Fixed a bug in which dndlnm and its dependents (ngtm, etc..) were calculated wrong if dndlog10m was called first.
- Fixed error in which for some choices of M, the whole extension in ngtm would be NAN and give error

## 9.4.37 1.1.4 [August 27, 2013]

### **Features**

• Added ability to change resolution in CAMB from hmf interface (This requires a re-install of pycamb to the newest version on the fork)

### 9.4.38 1.1.3 [August 7, 2013]

### **Features**

• Added Behroozi Fit (thanks to P. Behroozi)

### 9.4.39 1.1.2 [July 02, 2013]

### **Features**

• Ability to calculate fitting functions to whatever mass you want (BEWARE!!)

# 9.4.40 1.1.1 [July 02, 2013]

### **Features**

• Added Eisenstein-Hu fit to the transfer function

### **Enhancments**

• Improved docstring for Perturbations class

# **Bugfixes**

- Corrections to Watson fitting function from latest update on arXiv (thanks to W. Watson)
- IMPORTANT: Fixed units for k and transfer function (Thanks to A. Knebe)

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# 9.4.41 1.1.0 [June 27, 2013]

#### **Enhancments**

 Massive overhaul of structure: Now dependencies are tracked throughout the program, making updates even faster

# 9.4.42 1.0.10 [June 24, 2013]

#### **Enhancments**

• Added dependence on Delta\_vir to Tinker

# 9.4.43 1.0.9 [June 19, 2013]

### **Bugfixes**

• Fixed an error with an extra ln(10) in the mass function (quoted as dn/dlnM but actually outputting dn/dlog10M)

# 9.4.44 1.0.8 [June 19, 2013]

### **Enhancments**

- Took out log10 from cumulative mass functions
- Better cumulative mass function logic

# 9.4.45 1.0.6 [June 19, 2013]

### **Bugfixes**

• Fixed cumulative mass functions (extra factor of M was in there)

# 9.4.46 1.0.4 [June 6, 2013]

### **Features**

· Added Bhattacharya fitting function

### **Bugfixes**

· Fixed concatenation of list and dict issue

# 9.4.47 1.0.2 [May 21, 2013]

### **Bugfixes**

• Fixed some warnings for non-updated variables passed to update()

# 9.4.48 1.0.1 [May 20, 2013]

#### **Enhancments**

- Added better warnings for non-updated variables passed to update()
- Made default cosmology WMAP7

# 9.4.49 0.9.99 [May 10, 2013]

### **Enhancments**

• Added warning for k\*R limits

# **Bugfixes**

- Couple of minor bugfixes
- Important Angulo fitting function corrected (arXiv version had a typo).

# 9.4.50 0.9.97 [April 15, 2013]

### **Bugfixes**

• Urgent Bugfix for updating cosmology (for transfer functions)

# 9.4.51 0.9.96 [April 11, 2013]

### **Bugfixes**

· Few bugfixes

### 9.4.52 0.9.95 [April 09, 2013]

### **Features**

- Added cascading variable changes for optimization
- Added the README
- Added update() function to simply change parameters using cascading approach

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# 9.4.53 0.9.9 [April 08, 2013]

#### **Features**

- First version in its own package
- Added pycamb integration

### **Enhancments**

- Removed fitting function from being a class variable
- Removed overdensity form being a class variable

# 9.4.54 0.9.7 [March 18, 2013]

### **Enhancments**

- Modified set\_z() so it only does calculations necessary when z changes
- Made calculation of dlnsdlnM in init since it is same for all z
- Removed mean density redshift dependence

# 9.4.55 0.9.5 [March 10, 2013]

### **Features**

• The class has been in the works for almost a year now, but it currently will calculate a mass function based on any of several fitting functions.

# 9.5 Indices and tables

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- · modindex
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