ytree Documentation

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ytree is a tool for working with merger-tree data from multiple sources. ytree is an extension of the yt analysis toolkit and provides a similar interface for merger-tree data that includes universal field names, derived fields, and symbolic units. ytree can create merger-trees from Gadget FoF/Subfind catalogs, either for all halos or for a specific set of halos. ytree is able to load in merger-tree from the following formats:

- Amiga Halo Finder
- Consistent-Trees
- LHaloTree
- Rockstar halo catalogs without consistent-trees
- merger-trees made with ytree

All formats can be saved with a universal format that can be reloaded with ytree. Individual trees for single halos can also be saved.

CHAPTER 1

Table of Contents

1.1 Installation

ytree's main dependency is yt. Once you have installed yt following the instructions here, ytree can be installed using pip.

\$ pip install ytree

If you'd like to install the development version, the repository can be found at https://github.com/brittonsmith/ytree. This can be installed by doing:

```
$ git clone https://github.com/brittonsmith/ytree
$ cd ytree
$ pip install -e .
```

1.2 What version do I have?

To see what version of ytree you are using, do the following:

```
import ytree
print (ytree.__version__)
```

1.3 Sample Data

Sample datasets for every supported data format are available for download from the yt Hub in the ytree data collection. The entire collection (about 358 MB) can be downloaded via the yt Hub's web interface by clicking on "Actions" dropdown menu on the far right and selecting "Download collection." It can also be downloaded through the girder-client interface:

```
$ pip install girder-client
$ girder-cli --api-url https://girder.hub.yt/api/v1 download 59835a1ee2a67400016a2cda_
→ytree_data
```

1.4 Working with Merger-Trees

The *Arbor* class is responsible for loading and providing access to merger-tree data. Below, we demonstrate how to load data and what can be done with it.

1.4.1 Loading Merger-Tree Data

ytree can load merger-tree data from multiple sources using the load command.

```
import ytree
a = ytree.load("consistent_trees/tree_0_0_0.dat")
```

This command will determine the correct format and read in the data accordingly. For examples of loading each format, see below.

Loading Data

Below are instructions for loading all supported datasets.

Amiga Halo Finder

The Amiga Halo Finder format stores data in a series of files, with one each per snapshot. Parameters are stored in ".parameters" and ".log" files, halo information in ".AHF_halos" files, and descendent/ancestor links are stored in ".AHF_mtree" files. Make sure to keep all of these together. To load, provide the name of the first ".parameter" file.

Note: Three important notes about loading AHF data:

- 1. The dimensionless Hubble parameter is not provided in AHF outputs. This should be supplied by hand using the hubble_constant keyword. The default value is 1.0.
- 2. There will be no ".AHF_mtree" file for index 0 as the ".AHF_mtree" files store links between files N-1 and N.
- 3. ytree is able to load data where the graph has been calculated instead of the tree. However, even in this case, only the tree is preserved in ytree. See the Amiga Halo Finder Documentation for a discussion of the difference between graphs and trees.

Consistent-Trees

The consistent-trees format is typically one or a few files with a naming convention like "tree_0_0_0.dat". To load these files, just give the filename

```
import ytree
a = ytree.load("consistent_trees/tree_0_0_0.dat")
```

Rockstar Catalogs

Rockstar catalogs with the naming convention "out_*.list" will contain information on the descendent ID of each halo and can be loaded independently of consistent-trees. This can be useful when your simulation has very few halos, such as in a zoom-in simulation. To load in this format, simply provide the path to one of these files.

```
import ytree
a = ytree.load("rockstar/rockstar_halos/out_0.list")
```

LHaloTree

The LHaloTree format is typically one or more files with a naming convention like "trees_063.0" that contain the trees themselves and a single file with a suffix ".a_list" that contains a list of the scale factors at the time of each simulation snapshot.

In addition to the LHaloTree files, ytree also requires additional information about the simulation from a parameter file (in Gadget format). At minimum, the parameter file should contain the cosmological parameters HubbleParam, Omega0, OmegaLambda, BoxSize, PeriodicBoundariesOn, and ComovingIntegrationOn, and the unit parameters UnitVelocity_in_cm_per_s, UnitLength_in_cm, and UnitMass_in_g. If not specified explicitly (see below), a file with the extension ".param" will be searched for in the directory containing the LHaloTree files.

If all of the required files are in the same directory, an LHaloTree catalog can be loaded from the path to one of the tree files.

```
import ytree
a = ytree.load("lhalotree/trees_063.0")
```

Both the scale factor and parameter files can be specified explicitly through keyword arguments if they do not match the expected pattern or are located in a different directory than the tree files.

The scale factors and/or parameters themselves can also be passed explicitly from python.

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```
0.481261 , 0.502839 , 0.524807 , 0.547136 ,
                                                   0.569789 ,
                                       0.66287 ,
                                                   0.686541 ,
    0.59273 , 0.615919 , 0.639314 ,
                                       0.781371 ,
    0.710278 , 0.734031 , 0.757746 ,
                                                  0.804849,
    0.828124 , 0.851138 , 0.873833 ,
                                       0.896151 ,
                                                   0.918031 ,
    0.939414 , 0.960243 , 0.980457 ,
                                       1.
                                                1
a = ytree.load("lhalotree/trees_063.0",
              parameters=parameters,
              scale_factors=scale_factors)
```

TreeFarm

Merger-trees created with *TreeFarm* (ytree's merger-tree code for Gadget FoF/SUBFIND catalogs) can be loaded in by providing the path to one of the catalogs created during the calculation.

```
import ytree
a = ytree.load("tree_farm/tree_farm_descendents/fof_subhalo_tab_000.0.h5")
```

Saved Arbors

Once merger-tree data has been loaded, it can be saved to a universal format using *save_arbor* or *save_tree*. These can be loaded by providing the path to the primary hdf5 file.

```
import ytree
a = ytree.load("arbor/arbor.h5")
```

Saved Arbors from ytree 1.1

Arbors created with version 1.1 of ytree and earlier can be reloaded by providing the single file created. It is recommended that arbors be re-saved into the newer format as this will significantly improve performance.

```
import ytree
a = ytree.load("arbor.h5")
```

1.4.2 Working with Merger-Tree Data

Very little happens immediately after a dataset has been loaded. All tree construction and data access occurs only on demand. After loading, information such as the simulation box size, cosmological parameters, and the available fields can be accessed.

```
>>> print (a.box_size)
100.0 Mpc/h
>>> print (a.hubble_constant, a.omega_matter, a.omega_lambda)
0.695 0.285 0.715
>>> print (a.field_list)
['scale', 'id', 'desc_scale', 'desc_id', 'num_prog', ...]
```

Similar to yt, ytree supports accessing fields by their native names as well as generalized aliases. For more information on fields in ytree, see *Fields in ytree*.

How many trees are there?

As soon as any information about the collection of trees within the loaded dataset is requested, an array will be constructed containing objects representing the root of each tree, i.e., the last descendent halo. This structure is accessed by querying the loaded Arbor directly. It can also be accessed as a.trees.

```
>>> print (a.size)
Loading tree roots: 100%|| 5105985/5105985 [00:00<00:00, 505656111.95it/s]
327</pre>
```

Root Fields

Field data for all tree roots is accessed by querying the Arbor in a dictionary-like manner.

```
>>> print (a["mass"])
Getting root fields: 100%|| 327/327 [00:00<00:00, 9108.67it/s]
[ 6.57410072e+14 5.28489209e+14 5.18129496e+14 4.88920863e+14, ...,
8.68489209e+11 8.68489209e+11 8.68489209e+11] Msun</pre>
```

ytree uses yt's system for symbolic units, allowing for simple unit conversion.

```
>>> print (a["virial_radius"].to("Mpc/h"))
[ 1.583027 1.471894 1.462154 1.434253 1.354779 1.341322 1.28617, ...,
    0.173696 0.173696 0.173696 0.173696 0.173696] Mpc/h
```

When dealing with cosmological simulations, care must be taken to distinguish between comoving and proper reference frames. Please read *An Important Note on Comoving and Proper Units* before your magical ytree journey begins.

Accessing Individual Trees

Individual trees can be accessed by indexing the Arbor object.

```
>>> print (a[0])
TreeNode[12900]
```

A *TreeNode* is one halo in a merger-tree. The number is the universal identifier associated with halo. It is unique to the whole arbor. Fields can be accessed for any given *TreeNode* in the same dictionary-like fashion.

```
>>> print (a[0]["mass"])
657410071942446.1 Msun
```

The full lineage of the tree can be accessed by querying any TreeNode with the tree keyword.

```
>>> my_tree = a[0]
>>> print (my_tree["tree"])
[TreeNode[12900] TreeNode[12539] TreeNode[12166] TreeNode[11796] ...
TreeNode[591]]
```

Fields can be queried for the tree by including the field name.

```
>>> print (my_tree["tree", "virial_radius"])
[ 2277.73669065 2290.65899281 2301.43165468 2311.47625899 2313.99280576 ...
434.59856115 410.13381295 411.25755396] kpc
```

A halo's ancestors are stored as a list in the ancestors attribute. The descendents are stored in a similar fashion.

```
>>> print (my_tree.ancestors)
[TreeNode[12539]]
>>> print (my_tree.ancestors[0].descendent)
TreeNode[12900]
```

Accessing the Progenitor Lineage of a Tree

Similar to the *tree* keyword, the *prog* keyword can be used to access the line of main progenitors.

```
>>> print (my_tree["prog"])
[TreeNode[12900] TreeNode[12539] TreeNode[12166] TreeNode[11796] ...
TreeNode[62]]
>>> print (my_tree["prog", "mass"])
[ 6.57410072e+14 6.57410072e+14 6.53956835e+14 6.50071942e+14 ...
8.29496403e+13 7.72949640e+13 6.81726619e+13 5.99280576e+13] Msun
```

Customizing the Progenitor Line

By default, the progenitor line is defined as the line of the most massive ancestors. This can be changed by calling the *set_selector*.

```
>>> a.set_selector("max_field_value", "virial_radius")
```

New selector functions can also be supplied. These functions should minimally accept a list of ancestors and return a single TreeNode.

```
>>> def max_value(ancestors, field):
... vals = np.array([a[field] for a in ancestors])
... return ancestors[np.argmax(vals)]
...
>>> ytree.add_tree_node_selector("max_field_value", max_value)
>>>
>>> a.set_selector("max_field_value", "mass")
>>> print (a[0]["prog"])
```

1.4.3 Saving Arbors and Trees

Arbors of any type can be saved to a universal file format with the *save_arbor* function. These can be reloaded with the *load* command. This format is optimized for fast tree-building and field-access and so is recommended for most situations.

```
>>> fn = a.save_arbor()
Setting up trees: 100%|| 327/327 [00:00<00:00, 483787.45it/s]
Getting fields [1/1]: 100%|| 327/327 [00:00<00:00, 36704.51it/s]
Creating field arrays [1/1]: 100%|| 613895/613895 [00:00<00:00, 7931878.47it/s]
>>> a2 = ytree.load(fn)
```

By default, all trees and all fields will be saved, but this can be customized with the trees and fields keywords.

For convenience, individual trees can also be saved by calling *save_tree*.

```
>>> fn = a[0].save_tree()
Creating field arrays [1/1]: 100%|| 4897/4897 [00:00<00:00, 13711286.17it/s]
>>> a2 = ytree.load(fn)
```

1.5 An Important Note on Comoving and Proper Units

Users of yt are likely familiar with conversion from proper to comoving reference frames by adding "cm" to a unit. For example, proper "Mpc" becomes comoving with "Mpccm". This conversion relies on all the data being associated with a single redshift. This is not possible here because the dataset has values for multiple redshifts. To account for this, the proper and comoving unit systems are set to be equal to each other.

```
>>> print (a.box_size)
100.0 Mpc/h
>>> print (a.box_size.to("Mpccm/h"))
100.0 Mpccm/h
```

Data should be assumed to be in the reference frame in which it was saved. For length scales, this is typically the comoving frame. When in doubt, the safest unit to use for lengths is "unitary", which a system normalized to the box size.

```
>>> print (a.box_size.to("unitary"))
1.0 unitary
```

1.6 Fields in ytree

ytree supports multiple types of fields, each representing numerical values associated with each halo in the Arbor. These include the *native fields* stored on disk, *alias fields*, *derived fields*, and *analysis fields*.

1.6.1 The Field Info Container

Each *Arbor* contains a dictionary, called field_info, with relevant information for each available field. This information can include the units, type of field, any dependencies or aliases, and things relevant to reading the data from disk.

```
>>> import ytree
>>> a = ytree.load("tree_0_0_0.dat")
>>> print (a.field_info["Rvir"])
{'description': 'Halo radius (kpc/h comoving).', 'units': 'kpc/h ', 'column': 11,
    'aliases': ['virial_radius']}
>>> print (a.field_info["mass"])
{'type': 'alias', 'units': 'Msun', 'dependencies': ['Mvir']}
```

1.6.2 Fields on Disk

Every field stored in the dataset's files should be available within the Arbor. The field_list contains a list of all fields on disk with their native names.

```
>>> print (a.field_list)
['scale', 'id', 'desc_scale', 'desc_id', 'num_prog', ...]
```

1.6.3 Alias Fields

Because the various dataset formats use different naming conventions for similar fields, ytree allows fields to be referred to by aliases. This allows for a universal set of names for the most common fields. Many are added by default, including "mass", "virial_radius", "position_<xyz>", and "velocity_<xyz>". The list of available alias and derived fields can be found in the derived_field_list.

```
print (a.derived_field_list)
['uid', 'desc_uid', 'scale_factor', 'mass', 'virial_mass', ...]
```

Additional aliases can be added with add_alias_field.

```
>>> a.add_alias_field("amount_of_stuff", "mass", units="kg")
>>> print (a["amount_of_stuff"])
[ 1.30720461e+45, 1.05085632e+45, 1.03025691e+45, ...
1.72691772e+42, 1.72691772e+42, 1.72691772e+42]) kg
```

1.6.4 Derived Fields

Derived fields are functions of existing fields, including other derived and alias fields. New derived fields are created by providing a defining function and calling add_derived_field.

Field functions should only take a single argument, representing the entity for which the field is defined. This argument will also have access to the parent Arbor.

1.6.5 Analysis Fields

Analysis fields provide a means for saving the results of complicated analysis for any halo in the Arbor. This would be operations beyond derived fields, for example, things that might require loading the original simulation snapshots. New analysis fields are created with add_analysis_field and are initialized to zero.

Analysis fields will be automatically saved when the Arbor is saved with save_arbor.

1.7 Making Merger-trees from Gadget FoF/Subfind

The ytree TreeFarm can compute merger-trees either for all halos, starting at the beginning of the simulation, or for specific halos, starting at the final output and moving backward. These two use-cases are covered separately. Halo catalogs must be in the form created by the Gadget FoF halo finder or Subfind substructure finder.

1.7.1 Computing a Full Merger-tree

TreeFarm accepts a yt time-series object over which the merger-tree will be computed.

```
import yt
import ytree
ts = yt.DatasetSeries("data/groups_*/*.0.hdf5")
my_tree = ytree.TreeFarm(ts)
my_tree.trace_descendents("Group", filename="all_halos/")
```

The first argument to trace_descendents specifies the type of halo object to use. This will typically be either "Group" for FoF groups or Subhalo for Subfind groups. This process will create a new halo catalogs with the additional field representing the descendent ID for each halo. These can be loaded using yt like any other catalogs. Once complete, the final merger-tree can be *loaded into ytree*.

1.7.2 Computing a Targeted Merger-tree

Computing a full merger-tree can be extremely expensive when the simulation is large. Instead, merger-trees can be created for specific halos in the final dataset, then working backward. Below is an example of computing the merger-tree for only the most massive halo.

```
import yt
import yt
import ytree

ds = yt.load("fof_subfind/groups_025/fof_subhalo_tab_025.0.hdf5")
i_max = np.argmax(ds.r["Group", "particle_mass"])
my_id = ds.r["particle_identifier"][i_max]

ts = yt.DatasetSeries("data/groups_*/*.0.hdf5")
my_tree = ytree.TreeFarm(ts)
my_tree.trace_ancestors("Group", my_id, filename="my_halo/")
```

Just as above, the resulting catalogs can then be loaded into a *TreeFarm Arbor*.

1.7.3 Optimizing Merger-tree Creation

Computing merger-trees can often be an expensive task. Below are some tips for speeding up the process.

Running in Parallel

ytree uses the parallel capabilities of yt to divide up the halo ancestor/descendent search over multiple processors. In order to do this, yt must be set up to run in parallel. See here for instructions. Once this is done, a call to yt.enable_parallelism() must be added to your script.

```
import yt
yt.enable_parallelism()
import ytree
ts = yt.DatasetSeries("data/groups_*/*.0.hdf5")
my_tree = ytree.TreeFarm(ts)
my_tree.trace_descendents("Group", filename="all_halos/")
```

That script must then be run with mpirun.

```
mpirun -np 4 python my_script.py
```

Optimizing Halo Candidate Selection

Halo ancestors and descendents are typically found by comparing particle IDs between two halos. The method of selecting which halos should be compared can greatly affect performance. By default, TreeFarm will compare a halo against all halos in the next dataset. This is both the most robust and slowest method of matching ancestors and descendents. A smarter method is to select candidate matches from only a region around the target halo. For example, TreeFarm can be configured to select halos from a sphere centered on the current halo.

```
my_tree = ytree.TreeFarm(ts)
my_tree.set_selector("sphere", "virial_radius", factor=5)
my_tree.trace_descendents("Group", filename="all_halos/")
```

In the above example, candidate halos will be selected from a sphere that is five times the value of the "virial_radius" field. While this will speed up the calculation, a match will not be found if the ancestor/descendent is outside of this region. Some experimentation is recommended to find the optimal balance between speed and robustness.

Currently, the "sphere" selector is the only other selection method implemented, although others can be created easily. For an example, see sphere_selector.

Searching for Fewer Ancestors

When computing a merger-tree for specific halos (*Computing a Targeted Merger-tree*), you only be interested in the most massive or the few most massive progenitors. If this is the case, TreeFarm can be configured to end the ancestor search when these have been found, rather than searching for all possible progenitors.

The set_ancestry_filter function places a filter on which ancestors of any given halo will be returned and followed in successive rounds of the merger-tree process. The "most_massive" filter instructs the TreeFarm to only keep the most massive ancestor. This will greatly reduce the number of halos included in the merger-tree and, therefore, speed up the calculation considerably. For an example of how to create a new filter, see most_massive.

The filtering will only occur after all candidates have been checked for ancestry. An additional operation an be added to end the ancestry search after certain criteria have been met. In the call to set_ancestry_short below, the ancestry search will end as soon as an ancestor with at least 50% of the mass of the target halo has been found. For an example of how to create a new function of this type, see most_massive.

```
ts = yt.DatasetSeries("data/groups_*/*.0.hdf5")
my_tree = ytree.TreeFarm(ts)
my_tree.trace_ancestors("Group", my_id, filename="my_halo/")
my_tree.set_ancestry_filter("most_massive")
my_tree.set_ancestry_short("above_mass_fraction", 0.5)
```

1.8 Community Code of Conduct

ytree is a project by members of the yt community. As such, we stand by the yt Community Code of Conduct.

Below is the ytree version of this code.

ytree Community Code of Conduct

The community of participants in open source Scientific projects is made up of members from around the globe with a diverse set of skills, personalities, and experiences. It is through these differences that our community experiences success and continued growth. We expect everyone in our community to follow these guidelines when interacting with others both inside and outside of our community. Our goal is to keep ours a positive, inclusive, successful, and growing community.

As members of the community,

- We pledge to treat all people with respect and provide a harassment- and bullying-free environment, regardless of sex, sexual orientation and/or gender identity, disability, physical appearance, body size, race, nationality, ethnicity, and religion. In particular, sexual language and imagery, sexist, racist, or otherwise exclusionary jokes are not appropriate.
- We pledge to respect the work of others by recognizing acknowledgment/citation requests of original authors. As authors, we pledge to be explicit about how we want our own work to be cited or acknowledged.
- We pledge to welcome those interested in joining the community, and realize that including people with a variety of opinions and backgrounds will only serve to enrich our community. In particular, discussions relating to pros/cons of various technologies, programming languages, and so on are welcome, but these should be done with respect, taking proactive measure to ensure that all participants are heard and feel confident that they can freely express their opinions.
- We pledge to welcome questions and answer them respectfully, paying particular attention to those new to the community. We pledge to provide respectful criticisms and feedback in forums, especially in discussion threads resulting from code contributions.
- We pledge to be conscientious of the perceptions of the wider community and to respond to criticism respectfully. We will strive to model behaviors that encourage productive debate and disagreement, both within our community and where we are criticized. We will treat those outside our community with the same respect as people within our community.

We pledge to help the entire community follow the code of conduct, and to not remain silent when we see violations of the code of conduct. We will take action when members of our community violate this code such as contacting the project manager, Britton Smith (brittonsmith@gmail.com). All emails will be treated with the strictest confidence or talking privately with the person.

This code of conduct applies to all community situations online and offline, including mailing lists, forums, social media, conferences, meetings, associated social events, and one-to-one interactions.

This Community Code of Conduct comes the yt Community Code of Conduct, which was adapted from the Astropy Community Code of Conduct, which was partially inspired by the PSF code of conduct.

1.9 Contributing to ytree

ytree is a community project and it will be better with your contribution.

Contributions are welcome in the form of code, documentation, or just about anything. If you're interested in getting involved, please do!

ytree is developed using the same conventions as yt. The yt Developer Guide is a good reference for code style, communication with other developers, working with git, and issuing pull requests. For information specific to ytree, such as testing and adding support for new file formats, see the ytree Developer Guide.

If you'd like to know more, contact Britton Smith (brittonsmith@gmail.com).

You can also find help on the yt developers list.

1.10 Developer Guide

ytree is developed using the same conventions as yt. The yt Developer Guide is a good reference for code style, communication with other developers, working with git, and issuing pull requests. Below is a brief guide of aspects that are specific to ytree.

1.10.1 Contributing in a Nutshell

Step zero, get out of that nutshell!

After that, the process for making contributions to ytree is roughly as follows:

- 1. Fork the main ytree repository.
- 2. Create a new branch.
- 3. Make changes.
- 4. Run tests. Return to step 3, if needed.
- 5. Issue pull request.

The yt Developer Guide and github documentation will help with the mechanics of git and pull requests.

1.10.2 Testing

The ytree source comes with a series of tests that can be run to ensure nothing unexpected happens after changes have been made. These tests will automatically run when a pull request is issued or updated, but they can also be run locally very easily. At present, the suite of tests for ytree takes about three minutes to run.

Testing Data

The first order of business is to obtain the sample datasets. See *Sample Data* for how to do so. Next, ytree must be configure to know the location of this data. This is done by creating a configuration file in your home directory at the location ~/.config/ytree/ytreerc.

```
$ mkdir -p ~/.config/ytree
$ echo [ytree] > ~/.config/ytree/ytreerc
$ echo test_data_dir = /Users/britton/ytree_data >> ~/.config/ytree/ytreerc
$ cat ~/.config/ytree/ytreerc
[ytree]
test_data_dir = /Users/britton/ytree_data
```

This path should point to the outer directory containing all the sample datasets.

Run the Tests

Before running the tests, you will the pytest and flake8 packages. These can be installed with pip.

```
$ pip install pytest flake8
```

Once installed, the tests are run from the top level of the ytree source.

1.10.3 Adding Support for a New Format

The *Arbor* class is reasonably generalized such that adding support for a new file format should be relatively straightforward. The existing frontends also provide guidance for what must be done. Below is a brief guide for how to proceed. If you are interested in doing this, we will be more than happy to help!

Where do the files go?

As in yt, the code specific to one file format is referred to as a "frontend". Within the ytree source, each frontend is located in its own directory within ytree/arbor/frontends. Name your directory using lowercase and underscores and put it in there.

To allow your frontend to be directly importable at run-time, add the name to the _frontends list in ytree/ arbor/frontends/api.py.

Building Your Frontend

To build a new frontend, you will need to make frontend-specific subclasses for a few components. The easiest way to do this is to start with a blank Arbor subclass first. Create a sample script that loads your data with load, prints the number of trees, and queries some fields. Within the base classes, the necessary functions will raise a NotImplementedError if you have not added them yet. Keep running your script and implementing the function raising this error and before you know it, you'll be done.

The components and the files in which they belong are:

- 1. The Arbor itself (arbor.py).
- 2. The file i/o (io.py).
- 3. Recognizing frontend-specific fields (fields.py).

In addition to this, you will need to add a file called __init__.py, which will allow your code to be imported. This file should minimally import the frontend-specific Arbor class. For example, the consistent-trees __init__.py looks like this:

```
from ytree.arbor.frontends.consistent_trees.arbor import \
    ConsistentTreesArbor
```

Two Types of Arbors

There are generally two types of merger-tree data that ytree ingests:

1. all merger-tree data (full trees, halos, etc.) contained within a single file. An example of this is the consistent-trees frontend.

2. halos in files grouped by redshift (halo catalogs) that contain the halo id for the descendent halo which lives in the next catalog. An example of this is the rockstar frontend.

Depending on your case, different base classes should be subclassed. This is discussed below.

The _is_valid Function

Within every Arbor subclass should appear a function called <u>is</u>_valid. This function is used by load to determine if the provide file is the correct type. This function can examine the file's naming convention and/or open it and inspect its contents, whatever is required to uniquely identify your frontend. Have a look at the various examples.

Merger-Tree Data in One File (or a few)

If this is your case, then the consistent-trees and "ytree" frontends are the best examples to follow.

In arbor.py, your subclass of Arbor should implement two functions, _parse_parameter_file and _plant_trees.

_parse_parameter_file: This is the first thing called when your dataset is loaded. It is responsible for determining things like box size, cosmological parameters, and the list of fields.

_plant_trees: This function is responsible for constructing the array containing the roots of all trees in the Arbor. This should not fully build the trees, but just create *TreeNode* instances for each root and put them in the array.

In io.py, you will implement the machinery responsible for reading field data from disk. You must create a subclass of the *TreeFieldIO* class and implement the _read_fields function. This function accepts a single root node (a TreeNode that is the root of a tree) and a list of fields and should return a dictionary with NumPy arrays for each field.

Halo Catalog-style Data

If this is your case, then the rockstar and tree_farm frontends are the best examples to follow.

For this type of data, you will subclass the *CatalogArbor* class, which is itself a subclass of *Arbor* designed for this type of data.

In arbor.py, your subclass should implement two functions, _parse_parameter_file and _get_data_files. The purpose of _parse_parameter_file is described above.

_get_data_files: This type of data is usually loaded by providing one of the set of files. This function needs to figure out how many other files there are and their names and construct a list to be saved.

In io.py, you will create a subclass of *CatalogDataFile* and implement two functions: _parse_header and _read_fields.

_parse_header: This function reads any metadata specific to this halo catalog. For exmaple, you might get the current redshift here.

_read_fields: This function is responsible for reading field data from disk. This should minimally take a list of fields and return a dictionary with NumPy arrays for each field for all halos contained in the file. It should also, optionally, take a list of *TreeNode* instances and return fields only for them.

Field Units and Aliases (fields.py)

The *FieldInfoContainer* class holds information about field names and units. Your subclass can define two tuples, known_fields and alias_fields. The known_fields tuple is used to set units for fields on disk. This is useful especially if there is no way to get this information from the file. The convention for each entry is (name on disk, units).

By creating aliases to standardized names, scripts can be run on multiple types of data with little or no alteration for frontend-specific field names. This is done with the alias_fields tuple. The convention for each entry is (alias name, name on disk, field units).

You made it!

That's all there is to it! Now you too can do whatever it is people do with merger-trees. There are probably important things that were left out of this document. If you find any, please consider making an addition or opening an issue. If you're stuck anywhere, don't hesitate to ask for help. If you've gotten this far, we really want to see you make it to the finish!

Everyone Loves Samples

It would be especially great if you could provide a small sample dataset with your new frontend, something less than a few hundred MB if possible. This will ensure that your new frontend never gets broken and will also help new users get started. Once you have some data, make an addition to the arbor tests by following the example in tests/test_arbors.py. Then, contact Britton Smith to arrange for your sample data to be added to the ytree data collection on the yt Hub.

Ok, now you're totally done. Take the rest of the afternoon off.

1.11 Help

If you encounter problems, we want to help and there are lots of places to get help. As an extension of the yt project, we are members of the yt community. Any questions regarding ytree can be posted to the yt users list. You will also find interactive help on the yt slack channel.

Bugs and feature requests can also be posted on the ytree issues page.

See you out there!

1.12 Citing ytree

If you use ytree in your work, please cite the following:

Britton Smith, & Meagan Lang. (2018, February 16). ytree: merger-tree toolkit. Zenodo. http://doi.org/10.5281/ zenodo.1174374

For BibTeX users:

If possible, please also add a footnote pointing to http://ytree.readthedocs.io.

1.13 API Reference

1.13.1 Working with Merger-Trees

The *load* can load all supported merger-tree formats. Once loaded, the *save_arbor* and *save_tree* functions can be used to save the entire arbor or individual trees.

load(filename[, method])	Load an Arbor, determine the type automatically.
Arbor(filename)	Base class for all Arbor classes.
<pre>save_arbor([filename, fields, trees,])</pre>	Save the arbor to a file.
<pre>save_tree([filename, fields])</pre>	Save the tree to a file.
<pre>set_selector(selector, *args, **kwargs)</pre>	Sets the tree node selector to be used.
<pre>TreeNodeSelector(function[, args, kwargs])</pre>	The TreeNodeSelector is responsible for choosing
	which one of a halo's ancestors to return when query-
	ing the line of main progenitors for a halo.
<pre>add_tree_node_selector(name, function)</pre>	Add a TreeNodeSelector to the registry of known selec-
	tors, so they can be chosen with set_selector.
<pre>max_field_value(ancestors, field)</pre>	Return the TreeNode with the maximum value of the
	given field.
	Continued on next next

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Table 1 – continued from previous page		
<pre>min_field_value(ancestors, field)</pre>	Return the TreeNode with the minimum value of the given field.	

ytree.arbor.arbor.load

ytree.arbor.arbor.load (*filename*, *method=None*, ***kwargs*) Load an Arbor, determine the type automatically.

filename [string] Input filename.

method [optional, string] The type of Arbor to be loaded. Existing types are: ConsistentTrees, Rockstar, TreeFarm, YTree. If not given, the type will be determined based on characteristics of the input file.

****kwargs** [optional, dict] Additional keyword arguments are passed to _is_valid and the determined type.

Arbor

```
>>> import ytree
>>> # saved Arbor
>>> a = ytree.load("arbor/arbor.h5")
>>> # consistent-trees output
>>> a = ytree.load("rockstar_halos/trees/tree_0_0_0.dat")
>>> # Rockstar catalogs
>>> a = ytree.load("rockstar_halos/out_0.list")
>>> # TreeFarm catalogs
>>> a = ytree.load("my_halos/fof_subhalo_tab_025.0.h5")
>>> # LHaloTree catalogs
>>> a = ytree.load("my_halos/trees_063.0")
>>> # Amiga Halo Finder
>>> a = ytree.load("ahf_halos/snap_N64L16_000.parameter",
... hubble_constant=0.7)
```

ytree.arbor.arbor.Arbor

```
class ytree.arbor.arbor.Arbor (filename)
Base class for all Arbor classes.
```

Loads a merger-tree output file or a series of halo catalogs and create trees, stored in an array in trees. Arbors can be saved in a universal format with *save_arbor*. Also, provide some convenience functions for creating YTArrays and YTQuantities and a cosmology calculator.

__init__ (*filename*) Initialize an Arbor given an input file.

Methods

			Initialize an Arbor given an input file.
add_alias_field(alias,	field[,	units,	Add a field as an alias to another field.
force_add])			
add_analysis_field(name, units)			Add an empty field to be filled by analysis opera-
			tions.
			Continued on poyt page

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add_derived_field(name, function[, units,	Add a field that is a function of other fields.
])	
is_grown(tree_node)	Return True if a tree has been fully assembled, i.e.,
	the hierarchy of ancestor tree nodes has been built.
is_setup(tree_node)	Return True if arrays of uids and descendent uids
	have been read in.
query(key)	If given a string, return an array of field values for
	the roots of all trees.
<pre>save_arbor([filename, fields, trees,])</pre>	Save the arbor to a file.
<pre>select_halos(criteria[, trees, select_from,])</pre>	Select halos from the arbor based on a set of criteria
	given as a string.
<pre>set_selector(selector, *args, **kwargs)</pre>	Sets the tree node selector to be used.

Table 2 – continued from previous page

Attributes

arr	Create a YTArray using the Arbor's unit registry.
box_size	The simulation box size.
field_info	A dictionary containing information for each avail-
	able field.
hubble_constant	Value of the Hubble parameter.
quan	Create a YTQuantity using the Arbor's unit registry.
size	Return length of tree list.
trees	Array containing all trees in the arbor.
unit_registry	Unit system registry.

ytree.arbor.arbor.Arbor.save_arbor

```
Arbor.save_arbor (filename='arbor', fields=None, trees=None, max_file_size=524288)
Save the arbor to a file.
```

The saved arbor can be re-loaded as an arbor.

filename [optional, string] Output file keyword. If filename ends in ".h5", the main header file will be just that. If not, filename will be <filename>/<basename>.h5. Default: "arbor".

fields [optional, list of strings] The fields to be saved. If not given, all fields will be saved.

header_filename [string] The filename of the saved arbor.

```
>>> import ytree
>>> a = ytree.load("rockstar_halos/trees/tree_0_0_0.dat")
>>> fn = a.save_arbor()
>>> # reload it
>>> a2 = ytree.load(fn)
```

ytree.arbor.tree_node.TreeNode.save_tree

```
TreeNode.save_tree (filename=None, fields=None)
Save the tree to a file.
```

The saved tree can be re-loaded as an arbor.

filename [optional, string] Output file keyword. Main header file will be named <filename>/<filename>.h5. Default: "tree_<uid>".

fields [optional, list of strings] The fields to be saved. If not given, all fields will be saved.

filename [string] The filename of the saved arbor.

```
>>> import ytree
>>> a = ytree.load("rockstar_halos/trees/tree_0_0_0.dat")
>>> # save the first tree
>>> fn = a[0].save_tree()
>>> # reload it
>>> a2 = ytree.load(fn)
```

ytree.arbor.arbor.Arbor.set_selector

```
Arbor.set_selector(selector, *args, **kwargs)
```

Sets the tree node selector to be used.

This sets the manner in which halo progenitors are chosen from a list of ancestors. The most obvious example is to select the most massive ancestor.

selector [string] Name of the selector to be used.

Any additional arguments and keywords to be provided to the selector function should follow.

```
>>> import ytree
>>> a = ytree.load("rockstar_halos/trees/tree_0_0_0.dat")
>>> a.set_selector("max_field_value", "mass")
```

ytree.arbor.tree_node_selector.TreeNodeSelector

```
class ytree.arbor.tree_node_selector.TreeNodeSelector(function, args=None,
```

kwargs=None)

The TreeNodeSelector is responsible for choosing which one of a halo's ancestors to return when querying the line of main progenitors for a halo.

ancestors [list of TreeNode objects] List of TreeNode objects from which to select.

The function should return a single TreeNode.

```
>>> import ytree
>>> def max_value(ancestors, field):
... vals = np.array([a[field] for a in ancestors])
... return ancestors[np.argmax(vals)]
>>> ytree.add_tree_node_selector("max_field_value", max_value)
>>> a = ytree.load("tree_0_0_0.dat")
>>> a.set_selector("max_field_value", "mass")
>>> print (a[0]["prog"])
```

__init___(function, args=None, kwargs=None)
Initialize self. See help(type(self)) for accurate signature.

Methods

ytree.arbor.tree_node_selector.add_tree_node_selector

ytree.arbor.tree_node_selector.add_tree_node_selector (name, function)
Add a TreeNodeSelector to the registry of known selectors, so they can be chosen with set_selector.

name [string] Name of the selector.

function [callable] The associated function.

```
>>> import ytree
>>> def max_value(ancestors, field):
... vals = np.array([a[field] for a in ancestors])
... return ancestors[np.argmax(vals)]
>>> ytree.add_tree_node_selector("max_field_value", max_value)
>>> a = ytree.load("tree_0_0_0.dat")
>>> a.set_selector("max_field_value", "mass")
>>> print (a[0]["prog"])
```

ytree.arbor.tree_node_selector.max_field_value

ytree.arbor.tree_node_selector.max_field_value(ancestors, field)

Return the TreeNode with the maximum value of the given field.

ancestors [list of TreeNode objects] List of TreeNode objects from which to select.

field [string] Field to be used for selection.

TreeNode object

ytree.arbor.tree_node_selector.min_field_value

ancestors [list of TreeNode objects] List of TreeNode objects from which to select.

field [string] Field to be used for selection.

TreeNode object

1.13.2 Making Merger-Trees

TreeFarm(time_series[, setup_function])	TreeFarm is the merger-tree creator for Gadget FoF and
	Subfind halo catalogs.
<pre>trace_ancestors(halo_type, root_ids[,])</pre>	Trace the ancestry of a given set of halos.
<pre>trace_descendents(halo_type[, fields, filename])</pre>	Trace the descendents of all halos.
<pre>set_selector(selector, *args, **kwargs)</pre>	Set the method for selecting candidate halos for tracing
	halo ancestry.
<pre>set_ancestry_checker(ancestry_checker,)</pre>	Set the method for determing if a halo is the ancestor of
	another halo.

Continued on next page

	nom previous page
<pre>set_ancestry_filter(ancestry_filter, *args,)</pre>	Select a method for determining which ancestors are
	kept.
<pre>set_ancestry_short(ancestry_short, *args,)</pre>	Select a method for cutting short the ancestry search.
AncestryChecker(function[, args, kwargs])	An AncestryCheck is a function that is responsible for
	determining whether one halo is an ancestor of another.
add_ancestry_checker(name, function)	Add an ancestry checking function to the registry.
<pre>common_ids(descendent_ids, ancestor_ids[,])</pre>	Determine if at least a given fraction of ancestor's mem-
	ber particles are in the descendent.
AncestryFilter(function[, args, kwargs])	An AncestryFilter takes a halo and a list of ancestors
	and returns a filtered list of filtered list of ancestors.
<pre>add_ancestry_filter(name, function)</pre>	Add an ancestry filter function to the registry.
<pre>most_massive(halo, ancestors)</pre>	Return only the most massive ancestor.
AncestryShort(function[, args, kwargs])	An AncestryShort takes a halo and an ancestor halo and
	determines if the ancestry search should come to an end.
add_ancestry_short(name, function)	Add an ancestry short-out function to the registry.
above_mass_fraction(halo, ancestor, fraction)	Return only the most massive ancestor.
HaloSelector(function[, args, kwargs])	A HaloSelector is a function that is responsible for cre-
	ating a list of ids of halos that are potentially ancestors
	of a given halo.
add_halo_selector(name, function)	Add a HaloSelector to the registry of known selectors,
	so they can be chosen with set_selector.
<pre>sphere_selector(hc, ds2, radius_field[,])</pre>	Select halos within a sphere around the target halo.
all_selector(hc, ds2)	Return all halos from the ancestor dataset.

Table 5 – continued from previous page

ytree.tree_farm.tree_farm.TreeFarm

class ytree.tree_farm.tree_farm.**TreeFarm**(*time_series*, *setup_function=None*) TreeFarm is the merger-tree creator for Gadget FoF and Subfind halo catalogs.

TreeFarm can be used to create a merger-tree for the full set of halos, starting from the first catalog, or can be used to trace the ancestry of specific halos, starting from the last catalog. The merger-tree process will create a new set of halo catalogs, containing necessary fields (positions, velocities, masses), user-requested fields, and descendent IDs for each halo. These halo catalogs can be loaded at yt datasets.

- **time_series** [yt DatasetSeries object] A yt time-series object containing the datasets over which the merger-tree will be calculated.
- **setup_function** [optional, callable] A function that accepts a yt Dataset object and performs any setup, such as adding derived fields.

To create a full merger tree:

```
>>> import nummpy as np
>>> import yt
>>> import ytree
>>> from ytree.tree_farm import TreeFarm
>>> ts = yt.DatasetSeries("data/groups_*/fof_subhalo_tab*.0.hdf5")
>>> my_tree = TreeFarm(ts)
>>> my_tree.trace_descendents("Group", filename="all_halos/")
>>> a = ytree.load("all_halos/fof_subhalo_tab_000.0.h5")
>>> m = a["particle_mass"]
>>> i = np.argmax(m)
>>> print (a.trees[i]["prog", "particle_mass").to("Msun/h"))
```

To create a merger tree for a specific halo or set of halos:

```
>>> import nummpy as np
>>> import yt
>>> import ytree
>>> from ytree.tree_farm import TreeFarm
>>> ts = yt.DatasetSeries("data/groups_*/fof_subhalo_tab*.0.hdf5")
>>> ds = yt[-1]
>>> i = np.argmax(ds.r["Group", "particle_mass"].d)
>>> my_ids = ds.r["Group", "particle_identifier"][i_max]
>>> my_tree = TreeFarm(ts)
>>> my_tree.set_ancestry_filter("most_massive")
>>> my_tree.set_ancestry_short("above_mass_fraction", 0.5)
>>> my_tree.trace_ancestors("Group", my_ids, filename="my_halos/")
>>> a = ytree.load("my_halos/fof_subhalo_tab_025.0.h5")
>>> print (a[0]["prog", "particle_mass").to("Msun/h"))
```

___init___(*time_series*, *setup_function=None*) Initialize self. See help(type(self)) for accurate signature.

Methods

	Initialize self.
<pre>set_ancestry_checker(ancestry_checker,</pre>	Set the method for determing if a halo is the ancestor
)	of another halo.
<pre>set_ancestry_filter(ancestry_filter, *args,</pre>	Select a method for determining which ancestors are
)	kept.
<pre>set_ancestry_short(ancestry_short, *args,</pre>	Select a method for cutting short the ancestry search.
)	
<pre>set_selector(selector, *args, **kwargs)</pre>	Set the method for selecting candidate halos for trac-
	ing halo ancestry.
<pre>trace_ancestors(halo_type, root_ids[,])</pre>	Trace the ancestry of a given set of halos.
trace_descendents(halo_type[, fields, file-	Trace the descendents of all halos.
name])	

ytree.tree_farm.tree_farm.TreeFarm.trace_ancestors

TreeFarm.trace_ancestors (halo_type, root_ids, fields=None, filename=None)
Trace the ancestry of a given set of halos.

A merger-tree for a specific set of halos will be created, starting with the last halo catalog and moving backward.

halo_type [string] The type of halo, typically "FOF" for FoF groups or "Subfind" for subhalos.

root_ids [integer or array of integers] The halo IDs from the last halo catalog for the targeted halos.

fields [optional, list of strings] List of additional fields to be saved to halo catalogs.

filename [optional, string] Directory in which merger-tree catalogs will be saved.

ytree.tree_farm.tree_farm.TreeFarm.trace_descendents

```
TreeFarm.trace_descendents(halo_type, fields=None, filename=None)
```

Trace the descendents of all halos.

A merger-tree for all halos will be created, starting with the first halo catalog and moving forward.

halo_type [string] The type of halo, typically "FOF" for FoF groups or "Subfind" for subhalos.

fields [optional, list of strings] List of additional fields to be saved to halo catalogs.

filename [optional, string] Directory in which merger-tree catalogs will be saved.

ytree.tree_farm.tree_farm.TreeFarm.set_selector

TreeFarm.set_selector(selector, *args, **kwargs)

Set the method for selecting candidate halos for tracing halo ancestry.

The default selector is "all", i.e., check every halo for a possible match. This can be slow. The "sphere" selector can be used to specify that only halos within some sphere be checked.

selector [string] Name of selector.

ytree.tree_farm.tree_farm.TreeFarm.set_ancestry_checker

TreeFarm.**set_ancestry_checker** (*ancestry_checker*, **args*, ***kwargs*) Set the method for determing if a halo is the ancestor of another halo.

The default method defines an ancestor as a halo where at least 50% of its particles are found in the descendent.

ancestry_checker [string] Name of checking method.

ytree.tree_farm.tree_farm.TreeFarm.set_ancestry_filter

TreeFarm.set_ancestry_filter(ancestry_filter, *args, **kwargs)

Select a method for determining which ancestors are kept. The kept ancestors will have their ancestries tracked. This can be used to speed up merger-trees for targeted halos by specifying that only the most massive ancestor be kept.

ancestry_filter [string] Name of filter method.

ytree.tree_farm.tree_farm.TreeFarm.set_ancestry_short

TreeFarm.set_ancestry_short (ancestry_short, *args, **kwargs)
Select a method for cutting short the ancestry search.

This can be used to speed up merger-trees for targeted halos by specifying that the search come to an end when an ancestor with greater than 50% of the halo's mass has been found, thereby ensuring that the most massive halo has already been found.

ancestry_short [string] Name of short-out method.

ytree.tree_farm.ancestry_checker.AncestryChecker

class ytree.tree_farm.ancestry_checker.**AncestryChecker**(*function*, *args=None*,

kwargs=None)

An AncestryCheck is a function that is responsible for determining whether one halo is an ancestor of another.

descendent_ids [list of ints] Member ids for first halo.

ancestor_ids [list of int] Member ids for second halo.

The function should return True or False.

__init___(function, args=None, kwargs=None) Initialize self. See help(type(self)) for accurate signature.

Methods

ytree.tree_farm.ancestry_checker.add_ancestry_checker

ytree.tree_farm.ancestry_checker.add_ancestry_checker(name, function)
Add an ancestry checking function to the registry.

ytree.tree_farm.ancestry_checker.common_ids

ytree.tree_farm.ancestry_checker.common_ids (descendent_ids, ancestor_ids, threshold=0.5)

Determine if at least a given fraction of ancestor's member particles are in the descendent.

descendent_ids [list of ints] Member ids for first halo.

ancestor_ids [list of int] Member ids for second halo.

threshold [float, optional] Critical fraction of ancestor's particles ending up in the descendent to be considered a true ancestor. Default: 0.5.

True or False

ytree.tree_farm.ancestry_filter.AncestryFilter

class ytree.tree_farm.ancestry_filter.AncestryFilter(function, args=None,

kwargs=None)

An AncestryFilter takes a halo and a list of ancestors and returns a filtered list of filtered list of ancestors. For example, a filter could return only the most massive ancestor.

halo: halo data container Data container of the descendent halo.

ancestors [list of halo data containers] List of data containers for the ancestor halos.

The function should return a list of data containers.

___init___(function, args=None, kwargs=None) Initialize self. See help(type(self)) for accurate signature.

Methods

______(function[, args, kwargs]) Initialize self.

ytree.tree_farm.ancestry_filter.add_ancestry_filter

ytree.tree_farm.ancestry_filter.most_massive

halo: halo data container Data container of the descendent halo.

ancestors [list of halo data containers] List of data containers for the ancestor halos.

filtered_ancestors [list of halo data containers] List containing data container of most massive halo.

ytree.tree_farm.ancestry_short.AncestryShort

class ytree.tree_farm.ancestry_short.**AncestryShort** (*function*, args:

args=None,

kwargs=None) An AncestryShort takes a halo and an ancestor halo and determines if the ancestry search should come to an end.

halo: halo data container Data container of the descendent halo.

ancestor [halo data container] Data container for the ancestor halo.

The function should return True or False.

___init___(function, args=None, kwargs=None) Initialize self. See help(type(self)) for accurate signature.

Methods

ytree.tree_farm.ancestry_short.add_ancestry_short

ytree.tree_farm.ancestry_short.above_mass_fraction

halo: halo data container Data container of the descendent halo.

ancestor [halo data container] Data containers for the ancestor halo.

True or False

ytree.tree_farm.halo_selector.HaloSelector

class ytree.tree_farm.halo_selector.**HaloSelector** (*function*, *args=None*, *kwargs=None*) A HaloSelector is a function that is responsible for creating a list of ids of halos that are potentially ancestors of a given halo.

hc [halo container object] Halo container associated with the target halo.

ds2 [halo catalog-type dataset] The dataset of the ancestor halos.

The function should return a list of integers representing the ids of potential halos to check for ancestry.

___init___(function, args=None, kwargs=None) Initialize self. See help(type(self)) for accurate signature.

Methods

ytree.tree_farm.halo_selector.add_halo_selector

ytree.tree_farm.halo_selector.add_halo_selector(name, function)

Add a HaloSelector to the registry of known selectors, so they can be chosen with set_selector.

name [string] Name of the selector.

function [callable] The associated function.

ytree.tree_farm.halo_selector.sphere_selector

Select halos within a sphere around the target halo.

hc [halo container object] Halo container associated with the target halo.

ds2 [halo catalog-type dataset] The dataset of the ancestor halos.

radius_field [str] Name of the field to be used to get the halo radius.

factor [float, optional] Multiplicative factor of the halo radius in which potential halos will be gathered. Default: 1.

min_radius [YTQuantity or tuple of (value, unit)] An absolute minimum radius for the sphere.

my_ids [list of ints] List of ids of potential halos.

ytree.tree_farm.halo_selector.all_selector

ytree.tree_farm.halo_selector.**all_selector**(*hc*, *ds2*) Return all halos from the ancestor dataset.

hc [halo container object] Halo container associated with the target halo.

ds2 [halo catalog-type dataset] The dataset of the ancestor halos.

my_ids [list of ints] List of ids of potential halos.

1.13.3 Internal Classes

Arbor(filename)

Base class for all Arbor classes.

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Table 11 – continued from previous page			
CatalogArbor(filename)	Base class for Arbors created from a series of halo cata-		
	log files where the descendent ID for each halo has been		
	pre-determined.		
FieldInfoContainer(arbor)	A container for information about fields.		
FieldContainer(arbor)	A container for field data.		
FakeFieldContainer(arbor[, name])	A fake field data container used to calculate dependen-		
	cies.		
FieldIO(arbor)	Base class for FieldIO classes.		
TreeFieldIO(arbor)	IO class for getting fields for a tree.		
RootFieldIO(arbor)	IO class for getting fields for the roots of all trees.		
FallbackRootFieldIO(arbor)	Class for getting root fields from arbors that have no		
	specialized storage for root fields.		
DataFile(filename)	Base class for data files.		
CatalogDataFile(filename, arbor)	Base class for halo catalog files.		
TreeNode(uid[, arbor, root])	Class for objects stored in Arbors.		
<pre>TreeNodeSelector(function[, args, kwargs])</pre>	The TreeNodeSelector is responsible for choosing		
	which one of a halo's ancestors to return when query-		
	ing the line of main progenitors for a halo.		
AHFArbor(filename[, hubble_constant])	Arbor for Amiga Halo Finder data.		
AHFFieldInfo(arbor)			
AHFDataFile(filename, arbor)			
ArborArbor(filename)	Class for Arbors created with ytree version 1.1.0 or ear-		
	lier.		
ArborArborFieldInfo(arbor)			
ArborArborTreeFieldIO(arbor)			
ArborArborRootFieldIO(arbor)			
ConsistentTreesArbor(filename)	Arbors from consistent-trees output files.		
ConsistentTreesFieldInfo(arbor)			
ConsistentTreesDataFile(filename)			
ConsistentTreesTreeFieldIO(arbor)			
LHaloTreeArbor(*args, **kwargs)	Arbors for LHaloTree data.		
LHaloTreeFieldInfo(arbor)			
LHaloTreeTreeFieldIO(arbor)			
LHaloTreeRootFieldIO(arbor)			
RockstarArbor(filename)	Class for Arbors created from Rockstar out_*.list files.		
RockstarFieldInfo(arbor)			
RockstarDataFile(filename, arbor)			
TreeFarmArbor(filename)	Class for Arbors created with TreeFarm.		
TreeFarmFieldInfo(arbor)			
TreeFarmDataFile(filename, arbor)			
TreeFarmTreeFieldIO(arbor)			
YTreeArbor(filename)	Class for Arbors created from the <i>save_arbor</i> or <i>save_tree</i> functions.		
YTreeDataFile(filename)	—		
YTreeTreeFieldIO(arbor)			
YTreeRootFieldIO(arbor)			
/			

Table 11 – continued from previous page

ytree.arbor.arbor.CatalogArbor

class ytree.arbor.arbor.CatalogArbor(filename)

Base class for Arbors created from a series of halo catalog files where the descendent ID for each halo has been pre-determined.

___init___(*filename*) Initialize an Arbor given an input file.

Methods

init(filename)	Initialize an Arbor given an input file.
add_alias_field(alias, field[, units,	Add a field as an alias to another field.
force_add])	
add_analysis_field(name, units)	Add an empty field to be filled by analysis opera-
	tions.
add_derived_field(name, function[, units,	Add a field that is a function of other fields.
])	
is_grown(tree_node)	Return True if a tree has been fully assembled, i.e.,
	the hierarchy of ancestor tree nodes has been built.
is_setup(tree_node)	Return True if arrays of uids and descendent uids
	have been read in.
query(key)	If given a string, return an array of field values for
	the roots of all trees.
<pre>save_arbor([filename, fields, trees,])</pre>	Save the arbor to a file.
<pre>select_halos(criteria[, trees, select_from,])</pre>	Select halos from the arbor based on a set of criteria
	given as a string.
<pre>set_selector(selector, *args, **kwargs)</pre>	Sets the tree node selector to be used.

Attributes

arr	Create a YTArray using the Arbor's unit registry.
box_size	The simulation box size.
field_info	A dictionary containing information for each avail-
	able field.
hubble_constant	Value of the Hubble parameter.
quan	Create a YTQuantity using the Arbor's unit registry.
size	Return length of tree list.
trees	Array containing all trees in the arbor.
unit_registry	Unit system registry.

ytree.arbor.fields.FieldInfoContainer

class ytree.arbor.fields.FieldInfoContainer(arbor)

A container for information about fields.

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.	
clear()		
copy()		
		Continued on next page

fromkeys	Returns a new dict with keys from iterable and values
2	equal to value.
get(k[,d])	
items()	
keys()	
pop(k[,d])	If key is not found, d is returned if given, otherwise
	KeyError is raised
popitem()	2-tuple; but raise KeyError if D is empty.
resolve_field_dependencies(fields[,	Divide fields into those to be read and those to gen-
fcache])	erate.
<pre>setdefault(k[,d])</pre>	
<pre>setup_aliases()</pre>	Add aliases defined in the alias_fields tuple for each
	frontend.
<pre>setup_derived_fields()</pre>	Add stock derived fields.
<pre>setup_known_fields()</pre>	Add units for fields on disk as defined in the
	known_fields tuple.
update([E,]**F)	If E is present and has a .keys() method, then does:
	for k in E: $D[k] = E[k]$ If E is present and lacks a
	.keys() method, then does: for k, v in E: $D[k] = v$ In
	either case, this is followed by: for k in F: $D[k] =$
	F[k]

Table 14 – continued from previous page

values()

Attributes

alias_fields		
known_fields		

ytree.arbor.fields.FieldContainer

```
class ytree.arbor.fields.FieldContainer (arbor) A container for field data.
```

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
clear()	
copy()	
fromkeys	Returns a new dict with keys from iterable and values
	equal to value.
get(k[,d])	
items()	
keys()	
pop(k[,d])	If key is not found, d is returned if given, otherwise
	KeyError is raised
	Continued on post name

Continued on next page

	continued norm previous page
popitem()	2-tuple; but raise KeyError if D is empty.
<pre>setdefault(k[,d])</pre>	
update([E,]**F)	If E is present and has a .keys() method, then does: for k in E: D[k] = E[k] If E is present and lacks a .keys() method, then does: for k, v in E: D[k] = v In either case, this is followed by: for k in F: D[k] = F[k]
values()	

Table 16 – continued from previous page	6 – continued from previous pa	age
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ytree.arbor.fields.FakeFieldContainer

class ytree.arbor.fields.FakeFieldContainer(arbor, name=None)

A fake field data container used to calculate dependencies.

__init___(arbor, name=None)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor[, name])	Initialize self.
clear()	
сору()	
fromkeys	Returns a new dict with keys from iterable and values
	equal to value.
get(k[,d])	
items()	
keys()	
pop(k[,d])	If key is not found, d is returned if given, otherwise
	KeyError is raised
popitem()	2-tuple; but raise KeyError if D is empty.
<pre>setdefault(k[,d])</pre>	
update([E,]**F)	If E is present and has a .keys() method, then does:
	for k in E: $D[k] = E[k]$ If E is present and lacks a
	.keys() method, then does: for k, v in E: $D[k] = v$ In
	either case, this is followed by: for k in F: $D[k] =$
	F[k]
values()	

Attributes

default factory	Factory for default value called by missing ().
deradre_raccory	ractory for default value caned by <u></u> missing().

ytree.arbor.io.FieldIO

class ytree.arbor.io.FieldIO(arbor)

Base class for FieldIO classes.

This object is resposible for field i/o for an Arbor.

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
<pre>get_fields(data_object[, fields])</pre>	Load field data for a data object into storage struc-
	tures.

ytree.arbor.io.TreeFieldIO

class ytree.arbor.io.TreeFieldIO(arbor)

IO class for getting fields for a tree.

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
<pre>get_fields(data_object[, fields])</pre>	Load field data for a data object into storage struc-
	tures.

ytree.arbor.io.RootFieldIO

class ytree.arbor.io.**RootFieldIO**(*arbor*) IO class for getting fields for the roots of all trees.

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
<pre>get_fields(data_object[, fields])</pre>	Load field data for a data object into storage struc-
	tures.

ytree.arbor.io.FallbackRootFieldIO

class ytree.arbor.io.FallbackRootFieldIO(arbor)

Class for getting root fields from arbors that have no specialized storage for root fields.

__init__ (*arbor*) Initialize self. See help(type(self)) for accurate signature.

init(arbor)	Initialize self.
<pre>get_fields(data_object[, fields])</pre>	Load field data for a data object into storage struc-
	tures.

ytree.arbor.io.DataFile

class ytree.arbor.io.**DataFile**(*filename*) Base class for data files.

This class allows us keep files open during i/o heavy operations and to keep things like caches of fields.

___init___(filename)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(filename)	Initialize self.
close()	
open()	

ytree.arbor.io.CatalogDataFile

class ytree.arbor.io.**CatalogDataFile** (*filename*, *arbor*) Base class for halo catalog files.

___init___(filename, arbor) Initialize self. See help(type(self)) for accurate signature.

Methods

	Initialize self.	
close()		
open()		

ytree.arbor.tree_node.TreeNode

class ytree.arbor.tree_node.**TreeNode** (*uid*, *arbor=None*, *root=False*) Class for objects stored in Arbors.

Each TreeNode represents a halo in a tree. A TreeNode knows its halo ID, the level in the tree, and its global ID in the Arbor that holds it. It also has a list of its ancestors. Fields can be queried for it, its progenitor list, and the tree beneath.

```
___init___(uid, arbor=None, root=False)
Initialize a TreeNode with at least its halo catalog ID and its level in the tree.
```

init(uid[, arbor, root])	Initialize a TreeNode with at least its halo catalog ID
	and its level in the tree.
add_ancestor(ancestor)	Add another TreeNode to the list of ancestors.
<pre>clear_fields()</pre>	If a root node, delete field data.
pwalk()	An iterator over all TreeNodes in the progenitor list,
	starting with this TreeNode.
query(key)	Return field values for this TreeNode, progenitor list,
	or tree.
reset()	Reset all data structures.
<pre>save_tree([filename, fields])</pre>	Save the tree to a file.
twalk()	An iterator over all TreeNodes in the tree beneath,
	starting with this TreeNode.

ancestors	
descids	
is_root	
nodes	
tree_size	
uids	

ytree.arbor.frontends.ahf.arbor.AHFArbor

class ytree.arbor.frontends.ahf.arbor.**AHFArbor** (*filename*, *hubble_constant=1.0*) Arbor for Amiga Halo Finder data.

____init___ (*filename*, *hubble_constant=1.0*) Initialize an Arbor given an input file.

	Initialize an Arbor given an input file.	
add_alias_field(alias, field[, units,	Add a field as an alias to another field.	
force_add])		
add_analysis_field(name, units)	Add an empty field to be filled by analysis opera-	
	tions.	
<pre>add_derived_field(name, function[, units,</pre>	Add a field that is a function of other fields.	
])		
is_grown(tree_node)	Return True if a tree has been fully assembled, i.e.,	
	the hierarchy of ancestor tree nodes has been built.	
is_setup(tree_node)	Return True if arrays of uids and descendent uids	
	have been read in.	
query(key)	If given a string, return an array of field values for	
	the roots of all trees.	
<pre>save_arbor([filename, fields, trees,])</pre>	Save the arbor to a file.	
<pre>select_halos(criteria[, trees, select_from,])</pre>	Select halos from the arbor based on a set of criteria	
	given as a string.	
<pre>set_selector(selector, *args, **kwargs)</pre>	Sets the tree node selector to be used.	

arr	Create a YTArray using the Arbor's unit registry.
box_size	The simulation box size.
field_info	A dictionary containing information for each avail-
	able field.
hubble_constant	Value of the Hubble parameter.
quan	Create a YTQuantity using the Arbor's unit registry.
size	Return length of tree list.
trees	Array containing all trees in the arbor.
unit_registry	Unit system registry.

ytree.arbor.frontends.ahf.fields.AHFFieldInfo

class ytree.arbor.frontends.ahf.fields.AHFFieldInfo(arbor)

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

init(arbor)	Initialize self.
clear()	
copy()	
fromkeys	Returns a new dict with keys from iterable and values equal to value.
get(k[,d])	
items()	
keys()	
pop(k[,d])	If key is not found, d is returned if given, otherwise KeyError is raised
popitem()	2-tuple; but raise KeyError if D is empty.
resolve_field_dependencies(fields[,	Divide fields into those to be read and those to gen-
fcache])	erate.
<pre>setdefault(k[,d])</pre>	
<pre>setup_aliases()</pre>	Add aliases defined in the alias_fields tuple for each frontend.
<pre>setup_derived_fields()</pre>	Add stock derived fields.
<pre>setup_known_fields()</pre>	Add units for fields on disk as defined in the known_fields tuple.
update([E,]**F)	If E is present and has a .keys() method, then does: for k in E: $D[k] = E[k]$ If E is present and lacks a
	.keys() method, then does: for k, v in E: $D[k] = v$ In either case, this is followed by: for k in F: $D[k] = F[k]$
values()	

alias_fields	 	 	
known_fields			

ytree.arbor.frontends.ahf.io.AHFDataFile

class ytree.arbor.frontends.ahf.io.AHFDataFile(filename, arbor)

___init___(*filename*, *arbor*) Initialize self. See help(type(self)) for accurate signature.

Methods

	Initialize self.
close()	
open()	

Attributes

links

ytree.arbor.frontends.arborarbor.arbor.ArborArbor

class ytree.arbor.frontends.arborarbor.arbor.**ArborArbor**(*filename*) Class for Arbors created with ytree version 1.1.0 or earlier.

___init___(*filename*) Initialize an Arbor given an input file.

			Initialize an Arbor given an input file.
add_alias_field(alias,	field[,	units,	Add a field as an alias to another field.
force_add])			
add_analysis_field(name,	, units)		Add an empty field to be filled by analysis opera-
			tions.
add_derived_field(name,	function[,	units,	Add a field that is a function of other fields.
])			
is_grown(tree_node)			Return True if a tree has been fully assembled, i.e.,
			the hierarchy of ancestor tree nodes has been built.
is_setup(tree_node)			Return True if arrays of uids and descendent uids
			have been read in.
query(key)			If given a string, return an array of field values for
			the roots of all trees.
save_arbor([filename, fields, t	trees,])		Save the arbor to a file.
			Continued on next page

Table 35 – Continued nom previous page		
<pre>select_halos(criteria[, trees, select_from,])</pre>	Select halos from the arbor based on a set of criteria	
	given as a string.	
<pre>set_selector(selector, *args, **kwargs)</pre>	Sets the tree node selector to be used.	

Table 33 – continued from previous page

Attributes

arr	Create a YTArray using the Arbor's unit registry.
box_size	The simulation box size.
field_info	A dictionary containing information for each avail-
	able field.
hubble_constant	Value of the Hubble parameter.
quan	Create a YTQuantity using the Arbor's unit registry.
size	Return length of tree list.
trees	Array containing all trees in the arbor.
unit_registry	Unit system registry.

ytree.arbor.frontends.arborarbor.fields.ArborArborFieldInfo

class ytree.arbor.frontends.arborarbor.fields.ArborArborFieldInfo(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
clear()	
сору()	
fromkeys	Returns a new dict with keys from iterable and values
	equal to value.
get(k[,d])	
items()	
keys()	
pop(k[,d])	If key is not found, d is returned if given, otherwise
	KeyError is raised
popitem()	2-tuple; but raise KeyError if D is empty.
resolve_field_dependencies(fields[,	Divide fields into those to be read and those to gen-
fcache])	erate.
<pre>setdefault(k[,d])</pre>	
<pre>setup_aliases()</pre>	Add aliases defined in the alias_fields tuple for each
	frontend.
<pre>setup_derived_fields()</pre>	Add stock derived fields.
<pre>setup_known_fields()</pre>	Add units for fields on disk as defined in the
	known_fields tuple.

Continued on next page

__init__(arbor)

Table	35 – continued from previous page
update([E,]**F)	If E is present and has a .keys() method, then does: for k in E: D[k] = E[k] If E is present and lacks a .keys() method, then does: for k, v in E: D[k] = v In either case, this is followed by: for k in F: D[k] = F[k]

values()

Attributes

alias_fields			
known_fields			

ytree.arbor.frontends.arborarbor.io.ArborArborTreeFieldIO

class ytree.arbor.frontends.arborarbor.io.ArborArborTreeFieldIO(arbor)

___init___(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
<pre>get_fields(data_object[, fields])</pre>	Load field data for a data object into storage struc-
	tures.

ytree.arbor.frontends.arborarbor.io.ArborArborRootFieldIO

class ytree.arbor.frontends.arborarbor.io.ArborArborRootFieldIO(arbor)

___init___(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
<pre>get_fields(data_object[, fields])</pre>	Load field data for a data object into storage struc-
	tures.

ytree.arbor.frontends.consistent_trees.arbor.ConsistentTreesArbor

class ytree.arbor.frontends.consistent_trees.arbor.ConsistentTreesArbor(filename)
 Arbors from consistent-trees output files.

```
___init___(filename)
```

Initialize an Arbor given an input file.

Methods

init(filename)	Initialize an Arbor given an input file.
add_alias_field(alias, field[, units,	Add a field as an alias to another field.
force_add])	
add_analysis_field(name, units)	Add an empty field to be filled by analysis opera-
	tions.
add_derived_field(name, function[, units,	Add a field that is a function of other fields.
])	
is_grown(tree_node)	Return True if a tree has been fully assembled, i.e.,
	the hierarchy of ancestor tree nodes has been built.
is_setup(tree_node)	Return True if arrays of uids and descendent uids
	have been read in.
query(key)	If given a string, return an array of field values for
	the roots of all trees.
<pre>save_arbor([filename, fields, trees,])</pre>	Save the arbor to a file.
<pre>select_halos(criteria[, trees, select_from,])</pre>	Select halos from the arbor based on a set of criteria
	given as a string.
<pre>set_selector(selector, *args, **kwargs)</pre>	Sets the tree node selector to be used.

Attributes

arr	Create a YTArray using the Arbor's unit registry.
box_size	The simulation box size.
field_info	A dictionary containing information for each avail-
	able field.
hubble_constant	Value of the Hubble parameter.
quan	Create a YTQuantity using the Arbor's unit registry.
size	Return length of tree list.
trees	Array containing all trees in the arbor.
unit_registry	Unit system registry.

ytree.arbor.frontends.consistent_trees.fields.ConsistentTreesFieldInfo

class ytree.arbor.frontends.consistent_trees.fields.ConsistentTreesFieldInfo(arbor)

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
clear()	
copy()	
fromkeys	Returns a new dict with keys from iterable and values
	equal to value.
get(k[,d])	

Continued on next page

items()	
keys()	
pop(k[,d])	If key is not found, d is returned if given, otherwise
	KeyError is raised
popitem()	2-tuple; but raise KeyError if D is empty.
resolve_field_dependencies(fields[,	Divide fields into those to be read and those to gen-
fcache])	erate.
<pre>setdefault(k[,d])</pre>	
<pre>setup_aliases()</pre>	Add aliases defined in the alias_fields tuple for each
	frontend.
<pre>setup_derived_fields()</pre>	Add stock derived fields.
<pre>setup_known_fields()</pre>	Add units for fields on disk as defined in the
	known_fields tuple.
update([E,]**F)	If E is present and has a .keys() method, then does:
	for k in E: $D[k] = E[k]$ If E is present and lacks a
	.keys() method, then does: for k, v in E: $D[k] = v$ In
	either case, this is followed by: for k in F: $D[k] =$
	F[k]
values()	

Table 41 – continued from previous page

Attributes

alias_fields			
known_fields			

ytree.arbor.frontends.consistent_trees.io.ConsistentTreesDataFile

class ytree.arbor.frontends.consistent_trees.io.ConsistentTreesDataFile(filename)

___init___(filename)

Initialize self. See help(type(self)) for accurate signature.

Methods

	Initialize self.
close()	
open()	

ytree.arbor.frontends.consistent_trees.io.ConsistentTreesTreeFieldIO

class ytree.arbor.frontends.consistent_trees.io.ConsistentTreesTreeFieldIO(arbor)

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
<pre>get_fields(data_object[, fields])</pre>	Load field data for a data object into storage struc-
	tures.

ytree.arbor.frontends.lhalotree.arbor.LHaloTreeArbor

__init__(*args, **kwargs)

Added reader class to allow fast access of header info.

Methods

i n i + (* orga * * laworga)			Added reader class to allow fast access of header
(*args, **kwargs)			
			info.
add_alias_field(alias,	field[,	units,	Add a field as an alias to another field.
force_add])			
add_analysis_field(name,	, units)		Add an empty field to be filled by analysis opera-
			tions.
add_derived_field(name,	function[,	units,	Add a field that is a function of other fields.
])			
is_grown(tree_node)			Return True if a tree has been fully assembled, i.e.,
			the hierarchy of ancestor tree nodes has been built.
is_setup(tree_node)			Return True if arrays of uids and descendent uids
			have been read in.
query(key)			If given a string, return an array of field values for
			the roots of all trees.
save_arbor([filename, fields, t	trees,])		Save the arbor to a file.
select_halos(criteria[, trees,	select_from	l,])	Select halos from the arbor based on a set of criteria
			given as a string.
<pre>set_selector(selector, *args,</pre>	**kwargs)		Sets the tree node selector to be used.
<pre>]) is_grown(tree_node) is_setup(tree_node) query(key) save_arbor([filename, fields, t select_halos(criteria[, trees,</pre>	trees,]) select_from		Add a field that is a function of other fields. Return True if a tree has been fully assembled, it the hierarchy of ancestor tree nodes has been buil Return True if arrays of uids and descendent us have been read in. If given a string, return an array of field values the roots of all trees. Save the arbor to a file. Select halos from the arbor based on a set of crite given as a string.

Attributes

The simulation box size.
A dictionary containing information for each avail-
able field.
Value of the Hubble parameter.
Create a YTQuantity using the Arbor's unit registry.
Return length of tree list.
Array containing all trees in the arbor.
Unit system registry.

class ytree.arbor.frontends.lhalotree.arbor.**LHaloTreeArbor**(*args, **kwargs) Arbors for LHaloTree data.

ytree.arbor.frontends.lhalotree.fields.LHaloTreeFieldInfo

class ytree.arbor.frontends.lhalotree.fields.LHaloTreeFieldInfo(arbor)

___init___(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
clear()	
copy()	
fromkeys	Returns a new dict with keys from iterable and values
	equal to value.
get(k[,d])	
items()	
keys()	
pop(k[,d])	If key is not found, d is returned if given, otherwise
	KeyError is raised
popitem()	2-tuple; but raise KeyError if D is empty.
resolve_field_dependencies(fields[,	Divide fields into those to be read and those to gen-
fcache])	erate.
<pre>setdefault(k[,d])</pre>	
<pre>setup_aliases()</pre>	Add aliases defined in the alias_fields tuple for each
	frontend.
<pre>setup_derived_fields()</pre>	Add stock derived fields.
<pre>setup_known_fields()</pre>	Add units for fields on disk as defined in the
	known_fields tuple.
update([E,]**F)	If E is present and has a .keys() method, then does:
	for k in E: $D[k] = E[k]$ If E is present and lacks a
	.keys() method, then does: for k, v in E: $D[k] = v$ In
	either case, this is followed by: for k in F: $D[k] =$
	F[k]
values()	

values()

Attributes

alias_fields		
known_fields		

ytree.arbor.frontends.lhalotree.io.LHaloTreeTreeFieldIO

class ytree.arbor.frontends.lhalotree.io.LHaloTreeFieldIO(arbor)

___init___(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
<pre>get_fields(data_object[, fields])</pre>	Load field data for a data object into storage struc-
	tures.

ytree.arbor.frontends.lhalotree.io.LHaloTreeRootFieldIO

class ytree.arbor.frontends.lhalotree.io.LHaloTreeRootFieldIO(arbor)

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
<pre>get_fields(data_object[, fields])</pre>	Load field data for a data object into storage struc-
	tures.

ytree.arbor.frontends.rockstar.arbor.RockstarArbor

class ytree.arbor.frontends.rockstar.arbor.RockstarArbor(filename)

Class for Arbors created from Rockstar out_*.list files. Use only descendent IDs to determine tree relationship.

__init__ (*filename*) Initialize an Arbor given an input file.

init(filename)	Initialize an Arbor given an input file.
add_alias_field(alias, field[, units	Add a field as an alias to another field.
force_add])	
add_analysis_field(name, units)	Add an empty field to be filled by analysis opera-
	tions.
add_derived_field(name, function[, units	Add a field that is a function of other fields.
])	
is_grown(tree_node)	Return True if a tree has been fully assembled, i.e.,
	the hierarchy of ancestor tree nodes has been built.
is_setup(tree_node)	Return True if arrays of uids and descendent uids
	have been read in.
query(key)	If given a string, return an array of field values for
	the roots of all trees.
<pre>save_arbor([filename, fields, trees,])</pre>	Save the arbor to a file.
<pre>select_halos(criteria[, trees, select_from,])</pre>	Select halos from the arbor based on a set of criteria
	given as a string.
<pre>set_selector(selector, *args, **kwargs)</pre>	Sets the tree node selector to be used.

arr	Create a YTArray using the Arbor's unit registry.
box_size	The simulation box size.
field_info	A dictionary containing information for each avail-
	able field.
hubble_constant	Value of the Hubble parameter.
quan	Create a YTQuantity using the Arbor's unit registry.
size	Return length of tree list.
trees	Array containing all trees in the arbor.
unit_registry	Unit system registry.

y tree. arbor. frontends. rock star. fields. Rock star Field Info

class ytree.arbor.frontends.rockstar.fields.RockstarFieldInfo(arbor)

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

init(arbor)	Initialize self.
clear()	
сору()	
fromkeys	Returns a new dict with keys from iterable and values
	equal to value.
get(k[,d])	
items()	
keys()	
pop(k[,d])	If key is not found, d is returned if given, otherwise
	KeyError is raised
popitem()	2-tuple; but raise KeyError if D is empty.
resolve_field_dependencies(fields[,	Divide fields into those to be read and those to gen-
fcache])	erate.
setdefault(k[,d])	
setup_aliases()	Add aliases defined in the alias_fields tuple for each
	frontend.
_setup_derived_fields()	Add stock derived fields.
<pre>setup_known_fields()</pre>	Add units for fields on disk as defined in the
	known_fields tuple.
update([E,]**F)	If E is present and has a .keys() method, then does:
	for k in E: $D[k] = E[k]$ If E is present and lacks a
	.keys() method, then does: for k, v in E: $D[k] = v$ In
	either case, this is followed by: for k in F: $D[k] =$
	F[k]
values()	

alias_fields			
known_fields			

ytree.arbor.frontends.rockstar.io.RockstarDataFile

class ytree.arbor.frontends.rockstar.io.RockstarDataFile(filename, arbor)

___init___(*filename*, *arbor*) Initialize self. See help(type(self)) for accurate signature.

Methods

	Initialize self.
close()	
open()	

ytree.arbor.frontends.tree_farm.arbor.TreeFarmArbor

class ytree.arbor.frontends.tree_farm.arbor.**TreeFarmArbor**(*filename*) Class for Arbors created with TreeFarm.

___init___(*filename*) Initialize an Arbor given an input file.

init(filename)	Initialize an Arbor given an input file.
add_alias_field(alias, field[, units,	Add a field as an alias to another field.
force_add])	
add_analysis_field(name, units)	Add an empty field to be filled by analysis opera-
	tions.
<pre>add_derived_field(name, function[, units,</pre>	Add a field that is a function of other fields.
])	
is_grown(tree_node)	Return True if a tree has been fully assembled, i.e.,
	the hierarchy of ancestor tree nodes has been built.
is_setup(tree_node)	Return True if arrays of uids and descendent uids
	have been read in.
query(key)	If given a string, return an array of field values for
	the roots of all trees.
<pre>save_arbor([filename, fields, trees,])</pre>	Save the arbor to a file.
<pre>select_halos(criteria[, trees, select_from,])</pre>	Select halos from the arbor based on a set of criteria
	given as a string.
<pre>set_selector(selector, *args, **kwargs)</pre>	Sets the tree node selector to be used.

arr	Create a YTArray using the Arbor's unit registry.
box_size	The simulation box size.
field_info	A dictionary containing information for each avail-
	able field.
hubble_constant	Value of the Hubble parameter.
quan	Create a YTQuantity using the Arbor's unit registry.
size	Return length of tree list.
trees	Array containing all trees in the arbor.
unit_registry	Unit system registry.

ytree.arbor.frontends.tree_farm.fields.TreeFarmFieldInfo

class ytree.arbor.frontends.tree_farm.fields.TreeFarmFieldInfo(arbor)

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

init(arbor)	Initialize self.
clear()	
сору()	
fromkeys	Returns a new dict with keys from iterable and values equal to value.
get(k[,d])	
items()	
keys()	
pop(k[,d])	If key is not found, d is returned if given, otherwise KeyError is raised
popitem()	2-tuple; but raise KeyError if D is empty.
resolve_field_dependencies(fields[,	Divide fields into those to be read and those to gen-
fcache])	erate.
<pre>setdefault(k[,d])</pre>	
<pre>setup_aliases()</pre>	Add aliases defined in the alias_fields tuple for each frontend.
<pre>setup_derived_fields()</pre>	Add stock derived fields.
<pre>setup_known_fields()</pre>	Add units for fields on disk as defined in the known_fields tuple.
update([E,]**F)	If E is present and has a .keys() method, then does:
	for k in E: $D[k] = E[k]$ If E is present and lacks a
	.keys() method, then does: for k, v in E: $D[k] = v$ In
	either case, this is followed by: for k in F: $D[k] =$
	F[k]
values()	

alias_fields			
known_fields			

ytree.arbor.frontends.tree_farm.io.TreeFarmDataFile

class ytree.arbor.frontends.tree_farm.io.TreeFarmDataFile(filename, arbor)

```
___init___(filename, arbor)
Initialize self. See help(type(self)) for accurate signature.
```

Methods

	Initialize self.	
close()		
open()		

ytree.arbor.frontends.tree_farm.io.TreeFarmTreeFieldIO

class ytree.arbor.frontends.tree_farm.io.TreeFarmTreeFieldIO(arbor)

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
<pre>get_fields(data_object[, fields])</pre>	Load field data for a data object into storage struc-
	tures.

ytree.arbor.frontends.ytree.arbor.YTreeArbor

class ytree.arbor.frontends.ytree.arbor.**YTreeArbor**(*filename*) Class for Arbors created from the *save_arbor* or *save_tree* functions.

___init___(*filename*) Initialize an Arbor given an input file.

Methods

			Initialize an Arbor given an input file.
add_alias_field(alias,	field[,	units,	Add a field as an alias to another field.
force_add])			

Continued on next page

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add_analysis_field(name, units)	Add an empty field to be filled by analysis opera-	
	tions.	
add_derived_field(name, function[, units,	Add a field that is a function of other fields.	
])		
is_grown(tree_node)	Return True if a tree has been fully assembled, i.e.,	
	the hierarchy of ancestor tree nodes has been built.	
is_setup(tree_node)	Return True if arrays of uids and descendent uids	
	have been read in.	
query(key)	If given a string, return an array of field values for	
	the roots of all trees.	
save_arbor([filename, fields, trees,])	Save the arbor to a file.	
<pre>select_halos(criteria[, trees, select_from,])</pre>	Select halos from the arbor based on a set of criteria	
	given as a string.	
<pre>set_selector(selector, *args, **kwargs)</pre>	Sets the tree node selector to be used.	

Table 62 - continued from previous page

Attributes

arr	Create a YTArray using the Arbor's unit registry.
box_size	The simulation box size.
field_info	A dictionary containing information for each avail-
	able field.
hubble_constant	Value of the Hubble parameter.
quan	Create a YTQuantity using the Arbor's unit registry.
size	Return length of tree list.
trees	Array containing all trees in the arbor.
unit_registry	Unit system registry.

ytree.arbor.frontends.ytree.io.YTreeDataFile

class ytree.arbor.frontends.ytree.io.YTreeDataFile(filename)

___init___(filename)

Initialize self. See help(type(self)) for accurate signature.

Methods

	Initialize self.
close()	
open()	

ytree.arbor.frontends.ytree.io.YTreeTreeFieldIO

class ytree.arbor.frontends.ytree.io.YTreeTreeFieldIO(arbor)

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

Methods

init(arbor)	Initialize self.
<pre>get_fields(data_object[, fields])</pre>	Load field data for a data object into storage struc-
	tures.

ytree.arbor.frontends.ytree.io.YTreeRootFieldIO

class ytree.arbor.frontends.ytree.io.YTreeRootFieldIO(arbor)

__init__(arbor)

Initialize self. See help(type(self)) for accurate signature.

init(arbor)	Initialize self.
<pre>get_fields(data_object[, fields])</pre>	Load field data for a data object into storage struc-
	tures.

CHAPTER 2

Citing ytree

If you use ytree in your work, please cite the following:

Britton Smith, & Meagan Lang. (2018, February 16). ytree: merger-tree toolkit. Zenodo. http://doi.org/10.5281/ zenodo.1174374

For BibTeX users:

If possible, please also add a footnote pointing to http://ytree.readthedocs.io.

CHAPTER $\mathbf{3}$

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