yt Cheat Sheet

General Info

For everything yt please see http://yt-project.org. Documentation

http://yt-project.org/doc/index.html. Need help? Start here http://yt-project.org/doc/help/ and then try the IRC chat room

http://yt-project.org/irc.html, or the mailing list http://lists.spacepope.org/listinfo.cgi/

yt-users-spacepope.org. Installing yt: The easiest way to install yt is to use the installation script found on the yt homepage or the docs linked above.

Command Line yt

yt, and its convenience functions, are launched from a command line prompt. Many commands have flags to control behavior. Commands can be followed by **--help** (e.g. **yt render --help**) for detailed help for that command including a list of the available flags.

iyt—Load yt and IPython.

yt load dataset — Load a single dataset.

yt help — Print yt help information.

yt stats dataset — Print stats of a dataset.

yt update — Update yt to most recent version.

yt update --all — Update yt and dependencies to most recent version.

yt instinfo — yt installation information.

yt notebook — Run the IPython notebook server. yt serve (*dataset*) — Run yt-specific web GUI (*dataset* is optional).

yt upload_image *image.png* — Upload PNG image to imgur.com.

yt upload_notebook notebook.nb — Upload IPython notebook to hub.yt-project.org.

yt plot dataset — Create a set of images.

yt render dataset — Create a simple volume rendering. yt mapserver dataset — View a plot/projection in a Gmaps-like interface.

yt pastebin *text.out* — Post text to the pastebin at paste, yt-project.org.

yt pastebin.grab *identifier* — Print content of pastebin to STDOUT.

yt hub_register — Register with hub.yt-project.org.

yt hub_submit — Submit hg repo to hub.yt-project.org. yt bootstrap_dev — Bootstrap a yt development environment.

yt bugreport — Report a yt bug.

yt hop dataset — Run hop on a dataset.

yt rpdb — Connect to running rpd session.

yt Imports

In order to use yt, Python must load the relevant yt modules into memory. The import commands are entered in the Python/IPython shell or used as part of a script. from yt.mods import * — Load base yt modules. from yt.config import ytcfg — Used to set yt configuration options. If used, must be called before importing any other module. from yt.analysis_modules.api import * — Load all yt analysis modules.

from yt.analysis_modules.halo_finding.api import * — Load halo finding modules. Other modules are loaded in a similar way by swapping the *emphasized* text. See the **Analysis Modules** section for a listing and short descriptions of each.

Numpy Arrays

Simulation data in yt is returned in Numpy arrays. The Numpy package provides a wealth of built-in functions that operate on Numpy arrays. Here is a very brief list of some useful ones. Please see

http://docs.scipy.org/doc/numpy/reference/ for the full numpy documentation.

v = a.max(), a.min() — Return maximum, minimum of a.

index = a.argmax(), a.argmin() — Return index of
max, min value of a.

v = a[index] — Select a single value from a at location index.

b = a[i:j] — Select the slice of values from a between locations *i* to *j*-1 saved to a new Numpy array b with length *j*-*i*.

sel = (a > const) — Create a new boolean Numpy
array sel, of the same shape as a, that marks which
values of a > const. Other operators (e.g. <, !=, %)
work as well.</pre>

b = **a**[**se1**] — Create a new Numpy array **b** made up of elements from **a** that correspond to elements of **se1** that are *True*. In the above example **b** would be all elements of **a** that are greater than **const**.

a.dump(filename.dat) — Save a to the binary file filename.dat.

a = np.load(filename.dat) — Load the contents of filename.dat into a.

IPython Tips

These tips work if IPython has been loaded, typically either by invoking iyt or yt load on the command line, or using the IPython notebook (yt notebook). Tab complete — IPython will attempt to auto-complete a variable or function name when the Tab key is pressed, e.g. *HaloFi*-Tab would auto-complete to *HaloFinder*. This also works with imports, e.g. *from numpy.random.*-Tab would give you a list of random functions (note the trailing period before hitting Tab). ?, ?? — Appending one or two question marks at the end of any object gives you detailed information about it, e.g. *variable_name*?. Below a few IPython "magics" are listed, which are

IPython-specific shortcut commands. **%paste** — Paste content from the system clipboard into

the IPython shell.

%hist — Print recent command history.

%quickref — Print IPython quick reference.

 $\ensuremath{\texttt{%pdb}}\xspace -$ Automatically enter the Python debugger at an exception.

%time, %timeit — Find running time of expressions for benchmarking. %lsmagic — List all available IPython magics. Hint: ? works with magics.

Please see http://ipython.org/documentation.html for the full IPython documentation.

Load and Access Data

The first step in using yt is to reference a simulation snapshot. After that, simulation data is generally accessed in yt using *Data Containers* which are Python objects that define a region of simulation space from which data should be selected. pf = load(dataset) - Reference a single snapshot.

dd = pf.h.all_data() — Select the entire volume. a = dd[field_name] — Saves the contents of field into the numpy array a. Similarly for other data containers. pf.h.field_list — A list of available fields in the snapshot.

pf.h.derived_field_list — A list of available derived fields in the snapshot.

val, loc = pf.h.find_max("Density") — Find the value of the maximum of the field Density and its location.

sp = **pf.h.sphere**(*cen, radius*) — Create a spherical data container. *cen* may be a coordinate, or "max" which centers on the max density point. *radius* may be a float in code units or a tuple of (*length, unit*).

re = pf.h.region(*cen*, *left edge*, *right edge*) — Create a rectilinear data container. *cen* is required but not used. *left* and *right edge* are coordinate values that define the region.

di = pf.h.disk(*cen*, *normal*, *radius*, *height*) — Create a cylindrical data container centered at *cen* along the direction set by *normal*, with total length $2 \times height$ and with radius *radius*.

bl = pf.h.boolean(constructor) — Create a boolean data container. constructor is a list of pre-defined non-boolean data containers with nested boolean logic using the "AND", "NOT", or "OR" operators. E.g. constructor= [sp, "NOT", (di, "OR", re)] gives a volume defined by sp minus the patches covered by di and re. pf.h.save_object(sp, "sp_for_later") — Save an object (sp) for later use.

sp = pf.h.load_object("sp_for_later") — Recover a saved object.

Defining New Fields & Quantities

yt expects on-disk fields, fields generated on-demand and in-memory. Quantities reduce a field (e.g. "Density") defined over an object (e.g. "sphere") to get a single value (e.g. "Mass").

def _MetalMassMsun(field, data)

return

data["Metallicity"]*data["CellMassMsun"]
add_field("MetalMassMsun",function=_MetalMassMsun)
Define a new quantity; note the first function operates on
grids and data objects and the second on the results of
the first.

def _TotalMass(data):

baryon_mass = data["CellMassMsun"].sum()

particle_mass = data["ParticleMassMsun"].sum()
return baryon_mass, particle_mass

def _combTotalMass(data, baryon_mass, particle_mass):

Slices and Projections

slc = SlicePlot(pf, axis, field, center=, width=, weight_field=, additional parameters) — Make a slice plot perpendicular to axis of field weighted by weight_field at (code-units) center with width in code units or a (value, unit) tuple. Hint: try SlicePlot? in IPython to see additional parameters.

slc.save(*file_prefix*) — Save the slice to a png with name prefix *file_prefix*. .save() works similarly for the commands below.

prj = OffAxisSlicePlot(pf, normal, fields, center=, width=, depth=, north_vector=, weight_field=) —Make an off-axis slice. Note this takes an array of fields.

prj = OffAxisProjectionPlot(pf, normal, fields, center=, width=, depth=, north_vector=, weight_field=) --Make an off axis projection. Note this takes an array of fields.

Plot Annotations

Plot callbacks are functions itemized in a registry that is attached to every plot object. They can be accessed and then called like prj.annotate.velocity(factor=16, normalize=False). Most callbacks also accept a *plot_args* dict that is fed to matplotlib annotator.

velocity(factor=, scale=, scale_units=, normalize=) — Uses field "x-velocity" to draw quivers

magnetic_field(factor=, scale=, scale_units=,

normalize=) — Uses field "Bx" to draw quivers
quiver(field_x, field_y, factor=, scale=, scale_units=,
normalize=)

contour (field=, ncont=, factor=, clim=, take_log=, additional parameters) —Plots a number of contours ncont to interpolate field optionally using take_log, upper and lower contour limits and factor number of points in the interpolation.

grids(alpha=, draw_ids=, periodic=, min_level=, max_level=) —Add grid boundaries.

streamlines(field_x, field_y, factor=, density=)

clumps(clumplist) — Generate clumplist using the clump finder and plot.

arrow(pos, code_size) Add an arrow at a position. point(pos, text) — Add text at a position. marker(pos, marker=) — Add a matplotlib-defined marker at a position.

sphere(center, radius, text=) — Draw a circle and append text.

hop_circles(hop_output, max_number=, annotate=, min_size=, max_size=, font_size=, print_halo_size=, fixed_radius=, min_mass=, print_halo_mass=, width=) — Draw a halo, printing it's ID, mass, clipping halos depending on number of particles (size) and optionally fixing the drawn circle radius to be constant for all halos. hop_particles(hop_output, max_number=, p_size=, min_size, alpha=) — Draw particle positions for member halos with a certain number of pixels per particle. particles(width, p_size=, col=, marker=, stride=, ptype=, stars_only=, dm_only=, minimum_mass=, alpha=) — Draw particles of p_size pixels in a slab of width with color using a matplotlib marker plotting only every stride number of particles. title(text)

The \sim /.yt/ Directory

yt will automatically check for configuration files in a special directory (\$HOME/.yt/) in the user's home directory.

The config file — Settings that control runtime behavior. The my_plugins.py file — Add functions, derived fields, constants, or other commonly-used Python code to yt.

Analysis Modules

The import name for each module is listed at the end of each description (see **vt Imports**).

Absorption Spectrum — (absorption_spectrum).

Clump Finder — Find clumps defined by density

thresholds (level_sets).

Coordinate Transformation -

(coordinate_transformation).

Halo Finding — Locate halos of dark matter particles (halo_finding).

Halo Mass Function — Find halo mass functions from data and from theory (halo_mass_function).

Halo Profiling — Profile and project multiple halos (halo_profiler).

Àalo Merger Tree — Create a database of halo mergers (halo_merger_tree).

Light Cone Generator — Stitch datasets together to perform analysis over cosmological volumes.

Light Ray Generator — Analyze the path of light rays. Radial Column Density — Calculate column densities around a point (radial_column_density).

Rockstar Halo Finding — Locate halos of dark matter using the Rockstar halo finder (halo_finding.rockstar).

Star Particle Analysis — Analyze star formation
history and assemble spectra (star_analysis).
Sunrise Exporter — Export data to the sunrise
visualization format (sunrise_export).
Two Point Functions — Two point correlations
(two_point_functions).

Parallel Analysis

Nearly all of yt is parallelized using MPI. The *mpi4py* package must be installed for parallelism in yt. To install *pip install mpi4py* on the command line usually works. Execute python in parallel similar to this: *mpirun -n 12 python script.py -parallel* This command may differ for each system on which you use yt; please consult the system documentation for details on how to run parallel applications. from yt.pmods import * — Load yt faster when in parallel. This replaces the usual from yt.mods import *. parallel_objects() — A way to parallelize analysis over objects (such as halos or clumps).

Pre-Installed Versions

yt is pre-installed on several supercomputer systems. NICS Kraken — $module \ load \ yt$

Mercurial

Please see http://mercurial.selenic.com/ for the full Mercurial documentation.

hg clone https://bitbucket.org/yt_analysis/yt — Clone a copy of yt.

hg status — Files changed in working directory.

hg diff — Print diff of all changed files in working directory.

hg diff -rRevX - rRevY — Print diff of all changes between revision RevX and RevY.

hg log — History of changes.

hg cat -rRevX file — Print the contents of file from revision RevX.

hg heads — Print all the current heads.

hg revert -rRevX file — Revert file to revision RevX.

On-disk changed version is moved to *file.orig*.

hg commit — Commit changes to repository.

hg push — Push changes to default remote repository.

hg pull — Pull changes from default remote repository. hg serve — Launch a webserver on the local machine to

examine the repository in a web browser.

FAQ

pf.field_info['field'].take_log = False — When plotting field, do not take log. Must enter pf.h before this command.