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## Contents

1	Introduction	<b>2</b>
	1.1 What is MilkyWay@home?	2
	1.2 What is $mwahpy$ ?	2
	1.3 What can $mwahpy$ do?	3
2	Installing mwahpy	3
3	Core Functionality	5
	3.1 The Timestep Class	5
	3.1.1 Initializing a Timestep	5
	3.1.2 Reading In & Writing Out Data	8
		9
		12
	· · · · · · · · · · · · · · · · · · ·	14
	3.2.2 Functionality of <i>Nbody</i> Objects	15
		15
4	Auxiliary Packages	15
		15
	4.2 Orbit Fitting	15
5	Flags & Settings	16
0		16
	· ·	16
	100	16
_		
6		16
	1	16
	1	16
	1	16
		16
		16
	$6.6  orbit\_fitting\_ac$	16

### 1 Introduction

### 1.1 What is MilkyWay@home?

At best, "MilkyWay@home" is a nebulous term that can refer to several different projects, softwares, and/or groups of people. At its core, MilkyWay@home is a crowd-sourced supercomputer that takes volunteer computing time to perform complex and time-consuming calculations that are designed to improve our understanding of the Milky Way Galaxy. At the moment, this includes both the MilkyWay@home Separation application and the MilkyWay@home N-body application. The MilkyWay@home project was developed at Rensselaer Polytechnic Institute under the direction of Dr. Heidi Jo Newberg, and has utilized countless hours of work from many graduate students and researchers. For more information regarding Milky-Way@home, I direct the user to the MilkyWay forums at https://milkyway.cs.rpi.edu/milkyway/. All of the MilkyWay@home code is publicly available at https://github.com/Milkyway-at-home.

While the Separation application is an important part of MilkyWay@home, the Separation application is difficult to apply to other projects. Additionally, the tools that I have developed for the Separation application are quite situation specific and not designed with user friendliness/adaptability in mind. For this reason, *mwahpy* does not include functionality for the Separation application. For Separation application tools and documentation, see the Separation application github page at https://github.com/Milkyway-at-home/milkywayathome\_client/tree/master/separation.

What I refer to as "MilkyWay@home" in this document is actually the MilkyWay@home N-body software. This software was originally designed to generate model dwarf galaxies in the Milky Way gravitational potential and integrate them forwards in time. This evolved dwarf galaxy would then be compared to observations of stellar streams in the sky in order to determine a quality of fit. By optimizing over the parameters that were used to generate the dwarf galaxy, one could in theory determine the parameters of the progenitor dwarf galaxy. This could even be done for streams with no obvious progenitor (e.g. the Orphan Stream).

It was realized shortly thereafter that the N-body application could be used to integrate the orbits of any bodies in the Milky Way, not just those that were generated as part of a dwarf galaxy. The ability to insert list of bodies to integrate forwards in time was provided. Additionally, the ability to control the size of the timestep of the simulation, the orbit of a dwarf galaxy progenitor, the timescale of the simulation, the dwarf galaxy parameters, the underlying gravitational potential, and other MilkyWay@home functionality made it a versatile and robust tool for N-body integration.

While the MilkyWay@home N-body application is currently used by many graduate and undergraduate students at Rensselaer Polytechnic Institute, the MilkyWay@home team is making an effort to encourage more widespread usage of the MilkyWay@home N-body software. Our goal is to provide a user-friendly, comprehensive experience for N-body integration. This will decrease the time needed to get new students and researchers up to speed with N-body software, which has become commonplace in the dynamical astrophysics community.

We expect that the MilkyWay@home N-body application will become more widely used, and that proper documentation and accompanying tools will quickly become necessary. Additionally, new applications of the N-body software have expanded simulations for upwards of one million bodies. Analysis of these large simulations requires quick, robust software. It is for these reasons that mwahpy was developed.

### 1.2 What is mwahpy?

The MilkyWay@home N-body software has been used by the Galactic dynamics research group at Rensselaer Polytechnic Institute for years. As each new student comes into contact with the software, each one has had to develop their own tools for analyzing the outputs of the software. Each individual software has its own set of bugs, sets the student back a few weeks while developing their own tools, and by not using a standardized set of analysis software we open up the group to problems with sharing data and code. Additionally, MilkyWay@home is written in C, a language which can be daunting to the typical undergraduate student (and is often unecessarily unwieldly if you are just trying to cut data or make a plot). My goal to alleviate these issues was a compilation of the tools that I had built in python and tested over my tenure as an undergraduate (and improved during my time as a graduate student) in the MilkyWay@home dynamics group. This became the mwahpy package.

Before we talk about what mwahpy is, it is good to clarify what mwahpy is not. This package is not able to do N-body integration. Things like generating dwarf galaxies, running simulations, and performing routines on the MilkyWay@home supercomputer are to be delegated to the proper MilkyWay@home software. Additionally, I would like to express that because the Separation application is interacted with by many fewer people than the N-body application, mwahpy will not support Separation application functionality.

There is plenty that mwahpy is meant to be used for, though. At its very base, mwahpy is a python package that is designed to easily and quickly read in N-body data output from the MilkyWay@home N-body software. Once read in, it is fairly simple to cut the data, plot the data, and save the data in a variety of formats. This package is also able to output data in a format that is readable by the MilkyWay@home N-body software. The software will automatically calculate complicated values such as proper motion and energies as needed, instead of computing everything up front. The major benefit of these routines is that new users can be confident that the values that are produced by mwahpy have been tested over several years and are known to be accurate.

There are several associated auxiliary packages in mwahpy as well, such as the coordinate transformation subpackage and the orbit fitting subpackages. These are provided in the package because while working with N-body simulations, I often ran into situations where I used the functionality provided in the auxiliary subpackages. The auxiliary subpackages are less streamlined and complete compared to the main subpackages, but are still fairly well tested and can be trusted.

The code for *mwahpy* is publically available and can be found at https://github.com/thomasdonlon/mwahpy. Not only is collaboration on the code allowed, it is encouraged! If you would like functionality added to *mwahpy* or if you find any bugs in the code, you can either leave a comment on the github page or you can write the code yourself and make a pull request. This code is still being actively maintained, and I plan on eventually getting around to any bug fixes or desired functionality that are brought to my attention.

### 1.3 What can mwahpy do?

The following functionality is provided in mwahpy:

- $\bullet$  Easily & quickly read in data from a Milky Way@home  $N\text{-}\mathrm{body}$  . out file
- MilkyWay@home N-body output only provides 3D Cartesian positions and velocities, as well as mass (in MilkyWay@home structural units). The *mwahpy* package converts this data into many other useful forms, such as R.A. & Dec., Galactic longitude and latitude, angular momenta about the Galactic center, line-of-sight velocity, proper motions, and others.
- Make cuts based on any of *mwahpy*'s supported values, cut the data based on individual components of the data, or subsample the data randomly or symmetrically.
- Quickly plot N-body data in any of the supported values
- A variety of coordinate transformations for typical coordinate systems used in Galactic astronomy
- Two separate orbit fitting routines that have been used in publications

The proper syntax, usage, and details of each of the above capabilities will be outlined in the following sections.

### 2 Installing mwahpy

For a typical user, installing mwahpy is as simple as installing any other python package. For linux machines, go to your terminal and type

Code Block 1: Installing mwahpy

>>> pip3 install mwahpy

Alternatively, you can also install mwahpy using

```
Code Block 2: Installing mwahpy

>>> python3 -m pip install mwahpy
```

It may be necessary to install *mwahpy* only in your user directory due to user access restrictions. It is strongly recommended you do not use sudo to install python packages. Using sudo makes it difficult to track who has the permissions to use what package on your machine, and on machines with multiple users, it can mean you have to install it for each user individually anyways. Misuse of sudo can also result in accidentally using a different version of *mwahpy* than intended, particularly after updates. To install *mwahpy* for only the active user, you can instead type

```
Code Block 3: Installing mwahpy
>>> pip3 install mwahpy --user
```

or

```
Code Block 4: Installing mwahpy

>>> python3 -m pip install mwahpy --user
```

Any of these lines of code will install mwahpy on your machine, as well as install all of the prerequisite packages needed for mwahpy to work.

In order to use mwahpy after it is installed, simply import the subpackages as you would any other other package in your python script. For example, the following lines are all syntactically acceptable imports:

```
Code Block 5: Importing mwahpy
import mwahpy.output_handler
from mwahpy.data import Data
from mwahpy.coords import *
```

It should be noted that *mwahpy* is only built for and maintained for python v3.2.3 and above. You should get an error when installing if your python installation is not recent enough for *mwahpy*. Depending on your installation, you may instead have to type >>> python -m pip install mwahpy instead of python3, or use pip3 instead of pip. If you are experiencing errors, asking questions on the *mwahpy* github page or trying different permutations of these installation methods are recommended.

If you wish to install <code>mwahpy</code> for development, then you should clone the most recent <code>mwahpy</code> github repository (found at https://github.com/thomasdonlon/mwahpy). Feel free to make your own fork of the master branch if that's what you would prefer. Then, you should download the source code wherever you wish on your machine. If you want to import mwahpy into a python script after doing this, you will have to add a couple extra lines to the import portion of your script:

### Code Block 6: Using mwahpy as a developer

```
import sys
sys.path.insert(1, '/home/path/where/init.py/is/located/')
import mwahpy_subpackage
```

In this case, you would replace the path with the path to your **init.py** file is located (it usually ends with .../mwahpy/mwahpy/), and you would replace import mwahpy\_subpackage with whatever subpackage you wish to use (for example, import output\_handler as oh).

If you then wish to add, remove, or change code in your distribution of *mwahpy*, that can be done easily in the folder where you cloned *mwahpy*. Any changes that you feel are beneficial to the package should be sent as a pull request to the master branch.

### 3 Core Functionality

### WARNING:

MilkyWay@home uses a right-handed Galactic Cartesian coordinate system. This is defined as positive X being in the direction of the Sun towards the Galactic center, positive Y being in the direction of the disk spin at the location of the Sun, and positive Z being in the direction of the right-handed cross product  $X \times Y$  (often called the "Galactic north"). In our coordinate system, the Sun is located at the position (X,Y,Z) = (-8,0,0).

This is in contrast to many other Galactic scientists, who prefer a left-handed coordinate system where the X-axis is flipped and the Sun is located at (X,Y,Z) = (8,0,0). In a left-handed coordinate system the physical interpretation of certain quantities are not always clear (such as the right-handed angular momentum cross product). Many coordinate transformations in mwahpy allow for left-handed coordinates. However, be aware that by default, mwahpy (and MilkyWay@home) output is right-handed.

### 3.1 The *Timestep* Class

The Timestep class is the heart of mwahpy, and it's where the majority of the important and useful calculations in the package are performed. An instance of the Timestep class represents a single timestep of an N-body simulation. In other words, a Timestep instance is the data from a single MilkyWay@home .out file. The code for the Timestep class can be found in mwahpy's timestep.py file.

### 3.1.1 Initializing a Timestep

The most basic usage of Timestep object is a blank instance, to which you can then manually add data:

# import numpy as np from mwahpy.timestep import Timestep t = Timestep() t.id = np.array([0, 1, 2]) t.x = np.array([10, 50, 100]) t.y = np.array([12, 15, 17]) t.z = np.array([1, 2, 3]) t.vx = np.array([13, 183, 102]) t.vy = np.array([0, 50, 180]) t.vz = np.array([23, 69, 12]) t.mass = np.array([1, 1, 1])

These 8 values are all that you need to specify for the Timestep class to do its job. These 8 values (ID, x, y, z, vx, vy, vz, and mass) will be referred to as "provided values". This is in contrast to the "supported values", which are any values that mwahpy can calculate for you (a full list of the supported values is provided in Section 6.1). In the Timestep implementation, all of the particle IDs, x positions, etc. are stored in order as a numpy array of those values. As such, the mwahpy package heavily utilizes features of the numpy package, and some knowledge of numpy can be helpful for those working with mwahpy.

If you wish to print the information of a single particle, you can do so with printParticle by specifying the ID of the particle you are interested in:

```
Code Block 8: Printing data for a single particle

>>> t.printParticle(1)
Printing data for Particle 1:
(id:1, x:50, y:15, z:2, vx:183, vy:50, vz:1, mass:1, )
```

### **WARNING:**

The provided values should **always** be numpy arrays of identical length. Attempting to calculate values, plot data, or write out data when the provided values have mismatched length or are not numpy arrays will typically result in an error. It is strongly recommended that the user uses the built-in Timestep methods for cutting or adding data instead of doing it manually.

From this point forward, the user can ask for any *mwahpy* supported value, and it will be calculated for them. For example, if you wanted the line-of-sight velocities of the particles, you could type:

```
Code Block 9: Calculating supported values

>>> t.vlos
array([ 7.894394 , 235.8965911 , 128.90974589])
```

Note that if you call t.printParticle again (this time using the optional dec argument to shorten the output), it now shows more data for the particle!

# Code Block 10: Printing data for a single particle >>>t.printParticle(1, dec=2) Printing data for Particle 1: (id:1, x:50, y:15, z:2, vx:183, vy:50, vz:69, mass:1, msol:222288.47, 1:0.29, b:0.04, ra:266.54, dec:-28.67, dist:59.94, lx:935, ly:3084, lz:-245, lperp:3222.62, ltot:3231.92, r:52.24, R:52.2, vlos:235.9, vgsr:247.14, rad:192.15, rot:-4.69, distFromCOM:52.24, )

This was all calculated in the background when you asked the package to calculate vlos for this Timestep. Due to the overhead on some calculations, the more computationally complex calculations are avoided until the user requests those values. This is explained in more detail in Section ??.

The Timestep class has one very unique property: attributes of a Timestep instance can be accessed via the usual method, or as the key to a dictionary. In fact, comparing the two methods shows that these two actions produce equivalent results.

```
Code Block 11: Accessing Timestep attributes

>>> t.x #accessing data as an attribute
array([ 10, 50, 100])
>>> t['x'] #accessing data as a dict key
array([ 10, 50, 100])
>>> np.all(t.x == t['x'])
True
>>> t.x[0] = 1 #changing the value for Particle 0's x position
>>> t.x[0] #the same value is accessed by both methods
1
>>> t['x'][0]
1
>>> t.x[0] == t['x'][0]
True
```

At first this property may seem confusing and not particularly useful. What is the point of being able to access the same data in two different ways? It turns out that adding this functionality to Timestep allows for some rather powerful behavior. Of the greatest importance to the typical user is the implementation of the iterator for Timestep, which iterates over the *keys* of the class. In this case, that has been implemented as the names of all of the supported values (that have been calculated so far!).

# Code Block 12: Iterating over a Timestep >>> outstr = '' >>> for key in t: ... outstr += (key + ',\_') >>> print(outstr) id, x, y, z, vx, vy, vz, mass, msol, l, b, ra, dec, dist, lx, ly, lz, lperp, ltot, r, R, vlos, vgsr, rad, rot, distFromCOM, >>> outstr = '' >>> for key in t: ... outstr += (str(round(t[key][0],2)) + ',\_') >>> print(outstr) 0, 10, 12, 1, 13, 0, 23, 1, 222288.47, 0.88, 0.06, 266.86, -28.15, 21.66, 276, 217, -156, 351.09, 384.19, 15.65, 15.62, 7.89, 21.43, 9.77, -9.99, 15.65,

This may not seem like a big deal at first. However, if one tries to reproduce this behavior for a class that doesn't have the property c.x == c['x'], then you will quickly run into several problems. I expect that the typical reader would try to iterate over the class' built-in attribute dictionary, which is what I initially tried. What happens if you want to add attributes to the class that cannot be iterated over at the same time as your arrays, such as a single identifying string c.name? I suggest that the reader play around with this idea on their own if they are so inclined.

### 3.1.2 Reading In & Writing Out Data

So far, we know how to initialize a Timestep, how to add data to it manually, and how to access this data. This is all fine and good, but most of the time a user will be interested in using data that has already been generated by the MilkyWay@hoe N-body application. Conveniently, there is a mwahpy function built specifically for this in the mwahpy.output\_handler subpackage:

The oh.readOutput function takes in the path to a MilkyWay@home .out file and outputs a Timestep instance of the data from that file. In the above code block, we read in the test ,out file, which is provided in .../mwahpy/test/. The oh.readOutput function provides you with a progress bar, which is useful for large files. In this case, the progress bar finished almost instantly since the file was small. Typically, the progress bar will reach 100%. The oh.readOutput function also provides you with the number of particles that were read in from the file, and a brief update on when it is converting the data (again, useful for very large files). From this point forward, you can perform any normal Timestep operations on the new Timestep object.

The oh.readOutput function has been heavily optimized, and operates extremely quickly. Even for files with millions of values, reading in the data only takes a few seconds on most machines. If you are working

with N-body output, it is strongly recommended that you use this function to initialize your Timestep instances.

The *mwahpy* package also offers a few different options for printing out Timestep data to a file. Notably, *mwahpy* can print out the data to a .csv file,

```
Code Block 14: Writing out a Timestep

>>> oh.makeCSV(t, '/path/to/my/file.csv')
Writing header...
Printing data...
Timestep output to /path/to/my/file.csv
```

Saving the data to a .csv saves all of the supported values that has been calculated so far. Alternatively, you can write out data from a Timestep to a MilkyWay@home .in file,

```
Code Block 15: Writing out a Timestep

>>> oh.makeNbodyInput(t, '/path/to/my/file.in')
Writing Timestep as N-body input to /path/to/my/file.in...
done
```

Note that a MilkyWay@home .in file will only include the data about the 8 provided values, and not any of the other supported values. This is due to the format that MilkyWay@home requires readable files to be in. Any file that you generate with makeNbodyInput is immediately ready to be used as the manual body input for a MilkyWay@home N-body simulation.

If a user wishes to suspend a Timestep object for later use, I recommend the pickle package (https://docs.python.org/3/library/pickle.html). The pickle package allows for object serialization and saving out/reading in arbitrary objects as binary code.

### 3.1.3 Manipulating Data

After N-body data has been read in, it is useful to be able to only select the data that you are interested in. There are a few different routines for this, namely subsetRect() and subsetCirc(). These methods are n-dimensional cutting routines. For example, say that you took the test data and only wanted particles with a positive X value.

Note that subsetRect() has cut along all axes, not only the axis that we instructed it to cut along. As such, it is easy to see that tcopy.y has the same length as tcopy.x. It is also important to note that

we made a copy of t before using subsetRect(), because the subset methods treat the Timestep object as directly mutable.

While cutting the data, we provided 1000 as an upper value for X. This was because subsetRect() requires both a lower and upper limit for each axis when cutting the data. To get around this, we just chose some arbitrarily large value that would not cut any data off at the high end of X values.

You can also cut on multiple axes with subsetRect(). Say that you wanted to only include particles with positive X and negative Y values:

```
Code Block 17: Cutting data in a Timestep

>>> tcopy = t.copy()
>>> tcopy.subsetRect(['x','y'], [(0,1000),(-1000,0)])
>>> tcopy.x
array([4.53813066, 0.38487162, 0.52268562])
>>> tcopy.y
array([-0.05701174, -0.43899208, -0.34261058])
```

As all methods in Timestep, these routines can be performed on any supported values **after they have been calculated**. Attempting to cut on line-of-sight velocity before it has been calculated, for example, would result in a KeyError.

Similar results can be obtained from subsetCirc. If we wanted all particles with X values within 1 kpc of the Galactic center:

Or if we wanted all particles with X and Y values within 1 kpc of the Galactic center:

This routine is a bit of a misnomer, as it can be extended to any n-dimensional spheroidal volume of your phase space. For example, say that we wanted particles located within a spheroid defined with an semiaxis of length 1 on the X-axis, a semiaxis of length 2 on the Y-axis, and a semiaxis of length 5 on the Z-axis, centered at (X, Y, Z) = (0, 1, 3). This can be done with

```
Code Block 20: Cutting data in a Timestep

>>> tcopy = t.copy()
>>> tcopy.subsetCirc(['x', 'y', 'z'],[1,2,5],[0,1,3])
>>> tcopy.x
array([-0.59395581, -0.00613003, 0.11393187, 0.35803147, 0.3565866])
>>> tcopy.y
array([0.87634781, 0.43348783, 0.39540971, 0.14577695, 0.0082146])
>>> tcopy.z
array([ 0.55739028, -0.18463063, 0.32116868, 0.17315727, -0.09600029])
```

Unlike subsetRect(), this routine becomes somewhat unclear when you begin mixing values with different units. In that case, it is suggested that you either make multiple circular cuts where the units of each value match, or make rectangular cuts instead.

Other data manipulation routines include sampling and splitting Timesteps. There are three types of sampling that *mwahpy* supports: manual, incremental, and random sampling, shown below.

```
Code Block 21: Sampling a Timestep
>>> #manual sampling
>>> tcopy = t.copy()
\Rightarrow tcopy.take([0,2,3,6,9]) #take particles at indices 0, 2, 3, 6, and 9
>>> tcopy.x
array([ 4.53813066, -1.41000385, -0.00613003, 0.35803147, 3.57179938])
>>> #incremental sampling
>>> tcopy = t.copy()
>>> tcopy.subsample(2) #take every 2nd particle, starting at 0
>>> tcopy.x
array([ 4.53813066, -1.41000385, 0.11393187, 0.35803147, 0.52268562])
>>> #random sampling
>>> tcopy = t.copy()
>>> tcopy.randSample(5) #randomly take 5 particles from the data
>>> tcopy.x
array([ 4.53813066, -0.59395581, 0.38487162, -0.00613003, 0.35803147])
>>> #run again to demonstrate randomness
>>> tcopy = t.copy()
>>> tcopy.randSample(5)
>>> tcopy.x
array([ 0.52268562, -0.59395581, 0.3565866, -0.00613003, 0.11393187])
```

It is easy to split a Timestep by providing the index at which you want to split:

```
Code Block 22: Splitting a Timestep

>>> tcopy, tcopy2 = t.split(4) #split at index 4
>>> tcopy.x
array([ 4.53813066, -0.59395581, -1.41000385, -0.00613003])
>>> tcopy2.x
array([ 0.11393187,  0.38487162,  0.35803147,  0.3565866,  0.52268562,
3.57179938])
```

Finally, if you would like to add two Timestep instances together, use the appendTimestep method:

You should now have a fundamental undertsanding of how to use the Timestep class. The rest of the methods that were not covered in this section can be found in Section 6.1. Continue reading for information about handling entire simulations with multiple timesteps and plotting timesteps and simulations.

### 3.2 The *Nbody* Class

The Nbody class is one rung above the Timestep class in the *mwahpy* hierarchy; one Nbody instance is composed of many individual Timestep instances. This is useful when you need to compare many timesteps from a single MilkyWay@home simulation. The simplest example of a Nbody instance is the default instance:

```
Code Block 24: Initializing an Nbody

>>> from mwahpy.nbody import Nbody
>>> n = Nbody()
```

From here, you can add a Timestep to the dictionary in the Nbody instance:

Note that the Nbody instance itself does not carry the Timestep data, but just points to it.

```
Code Block 26: An Nbody does not carry simulation data

>>> n.x
Traceback (most recent call last):
   File "<stdin>", line 1, in <module>
AttributeError: 'Nbody' object has no attribute 'x'
```

In this case, the Timestep t still exists on its own, but the Nbody instance forms a nice functional container for it. This can be seen by altering t, as the change is also present when accessing the data through the Nbody. Thus, the Nbody structure does not copy the Timestep data, it just references it.

The Timestep does recognize that it belongs to an Nbody class however, and it remembers its designated time and the Nbody object which it belongs to:

```
Code Block 28: Accessing an Nbody

>>> t.time
1
>>> t.nbody
<mwahpy.nbody.Nbody object at 0x[...]>
```

### WARNING:

Currently, Timestep instances only weakly retain information about parent Nbody instances. If a Timestep is added to a second Nbody, it will replace its self.time and self.nbody from the first Nbody instance with the new information from the second Nbody. This is made more confusing by the fact that multiple Nbody instances can point to the same Timestep, or a single Nbody can point to the same Timestep more than once. In these cases, the data can be properly accessed through the Nbody objects, but is not guaranteed to refer back to the desired information when accessed from the Timestep object.

### 3.2.1 Reading In & Writing Out Data

As with the Timestep class, the Nbody class is typically not created manually, but is initialized by reading in data. This is also done through the mwahpy.output\_handler subpackage by providing a folder with the oh.readFolder() function.

```
Code Block 29: Reading data into an Nbody
>>> n = oh.readFolder('<...path/to/mwahpy>/mwahpy/test/nbody_test/')
Reading in data from directory ../test/nbody_test/...
Reading in data from ../test/nbody_test/1...
[-----
                    ] 73%
10 objects read in
Converting data...done
Reading in data from ../test/nbody_test/2...
[---->
                    ] 73%
10 objects read in
Converting data...done
Reading in data from ../test/nbody_test/3...
[---->
                    ] 73%
10 objects read in
Converting data...done
>>> n[1].x
array([ 4.53813066, -0.59395581, -1.41000385, -0.00613003, 0.11393187,
       0.38487162, 0.35803147, 0.3565866, 0.52268562, 3.57179938])
>>> n[2].x
array([21.51019042, 17.97537383, 15.99136357, 12.30622142, 10.80896895,
      10.56226125, 12.94243578, 11.16187291, 11.17781449, 8.04042707])
```

This data has been read in from the .../mwahpy/test/nbody\_test/ folder that is included in mwahpy.

### WARNING:

When reading in a Nbody from simulation data, the **only** files in the provided folder must be MilkyWay@home .out files, and **must** be named only their time **without the** .out **extension**. This is the default output scheme of MilkyWay@home N-body.

Currently, there is not support for writing out Nbody data. This can be done fairly simply by iterating over the Nbody (see the next subsection for details) and writing out each Timestep. It is also not recommended that a user writes out Nbody data using the pickle package.

### 3.2.2 Functionality of Nbody Objects

Nbody objects have a few nice capabilities that allow for robust treatment of simulation data. To begin, iterating over an Nbody object returns its component Timestep objects sorted by time.

Being able to read in and iterate over large groups of Timestep instances from the same simulation is much easier than going through each one individually, not to mention it is often faster.

The times of each timestep can be scaled in order to reflect the actual physical simulation time of each timestep using the scaleTimes() method. For example, if one timestep is equal to 0.5 Myr in physical time, then we can scale the corresponding timesteps. We see that the original information is retained, although renamed:

### 3.3 Plotting

Text

### 4 Auxiliary Packages

Text

### 4.1 Coordinate Transformations

Text

### 4.2 Orbit Fitting

The subpackages mwahpy.orbit\_fitting and mwahpy.orbit\_fitting\_gc are both still under development. Once the code has been refactored and brought to an acceptable level of quality, the documentation for these subpackages will be provided.

### 5 Flags & Settings

Text

5.1 flags

Text

 $5.2 \quad mwahpy\_glob$ 

Text

5.3 pot

Text

### 6 Functions & Methods

Text

6.1 timestep

Text

 $6.2 \quad output\_handler$ 

Text

6.3 plot

Text

6.4 coords

Text

### 6.5 orbit\_fitting

The subpackage mwahpy.orbit\_fitting is still under development. Once the code has been refactored and brought to an acceptable level of quality, the documentation for this subpackage will be provided.

### $6.6 \quad orbit\_fitting\_gc$

The subpackage mwahpy.orbit\_fitting\_gc is still under development. Once the code has been refactored and brought to an acceptable level of quality, the documentation for this subpackage will be provided.