# Mass Sampling Documentation

# **Functions**

# Single Redshifts

#### Main function:

```
A. def mass sampling (mass range, redshift = 0.0, mdef = '200c', model =
   'bocquet16', sample num = 100000):
```

the function to give back a sample of single-redshift cluster mass distribution based on the halo mass function

```
Parameters:
_____
input:
mass range: a tuple of cluster masses, lower limit, and upper limit for
sampling
redshift: a float, 0.0 by default
sample num: an integer of the number of samples, 100000 by default
mdef: The mass definition in which the halo mass M is given
model: the halo mass function model used by colossus
output:
mass chain: a NumPy array of length = sample num.
test func: the likelihood function
```

#### Library:

• Colossus

### step:

- Initiate a NumPy array as cluster masses
- Use the halo mass function given by colossus, give back an array of number density
- ullet Extract power from the mass array using helper 3, from 10^14 M $\odot$  to
- Use helper 2 to calculate the final mass sampling

# **Helpers:**

1. lnpo(mass, min, max, test fun):

likelihood function used by MCMC

#### parameters:

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mass: a float or a 1d NumPy array of cluster mass in 10^n M⊙ unit

2. interpolate\_MCMC(mass\_array\_p, mfunc\_n, mass\_range, sample\_num,
redshift):

interpolate and normalize mfunc\_n, use the result as a likelihood function and perform MCMC method to get the sample.

### parameters:

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input:

mass\_arr\_p: 1d NumPy array of cluster mass power (for example, 10^14 M⊙ represented as 14 in arr)

mfunc\_n: 1d NumPy array of halo number density \* 10^5

mass\_range: a tuple of cluster masses, lower limit, and upper limit for sampling

sample num: an integer of the number of samples

output:

sample\_chain.flatten(): an 1d NumPy array of mass sampling, same unit as
mass\_arr\_p

# Library:

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- scipy(interpolate & integrate)
- Emcee

### step:

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- Interpolate mass\_arr\_p & mfunc\_n, return a function object
- Calculate the integration of function between the mass range, normalize the function using integration, result in test func
- Use emcee package to do MCMC simulation, Helpers 1 (lnpo) used as the likelihood function, return test\_func and mass\_chain.flatten()
- 3. extract power(mass arr):

Function to extract the power of galaxy cluster mass array, switch from  $10^n$  to n

# Multiple Redshifts

#### Main function:

```
B. def mul_redshift_mass_sampling(rs_dist = "skewnorm", rs_range =
(0.0, 1.5), mass_range = (14.0, 16.0), mdef = '200c', model = 'bocquet16',
sample num = 100000, store = True):
```

the function to give back a sample of multi-redshift cluster mass distribution based on halo mass function

```
Parameters:
    input:
    rs_dist: a string, representing the distribution of cluster redshift,
"skewnorm" by default
    rs_range: a tuple of redshift range, (0.0, 1.5) by default
    mass_range: a tuple of cluster masses, lower limit and upper limit for
sampling, [min, max] in 10^min M② unit
    mdef: The mass definition in which the halo mass M is given; see
colossus doc for more info
(https://bdiemer.bitbucket.io/colossus/lss_mass_function.html#lss.mass_function.massFunction)
    model: the halo mass function model used by colossus; see colossus doc
for more info
```

sample\_num: an integer of the number of samples, 100000 by default
store: a boolean, if True store mass array and redshift into a csv file
and return a string of path to file if False returns None

# output:

fin\_cluster: a Pandas dataframe with 2 columns of ["mass\_arr",
"redshift"], NumPy array of cluster mass corresponding to redshift stored
in each row

filepath: str of file path if store=True, else None

# Library:

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• pandas

#### Step:

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- Call helper 2 to obtain the redshift interval (chop) and corresponding cluster number.
- Loop through every redshift in chop, call function A(mass\_sampling) to do single redshift mass sampling, store mass and redshifts in two pandas Series

### **Helpers:**

1. skew\_sample(size = 10000):

the function to give back a sample of redshift based on skew gaussian distribution imitating SPT cluster data:

https://pole.uchicago.edu/public/data/sptsz-clusters/

```
Parameters:
    -----
input:
    size: integer, sample number

output:
    mass_chain: a NumPy array of length = sample_num
    rs_sample: a 1d NumPy array of clusters' redshift sample with length = size
```

#### Library:

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• scipy.stats

#### Step:

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- Use scipy.stats to imitate redshift distribution
- Use skewnorm.rvs to draw a sample from it
- 2. single\_redshift\_num(rs\_range, sample\_num, rs\_dist\_model):

the function to give back redshifts and sample\_num per redshift for multi-redshift sampling

#### Parameters:

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#### input:

 ${\tt rs\_dist\_model:}$  a string, representing the distribution of cluster redshift

rs\_range: a tuple of redshift range, (0.0, 1.5) by default
sample\_num: an integer of the number of samples, 100000 by default

# output:

chop: a NumPy array of redshifts

num\_per\_redshift: a NumPy array of cluster num within the corresponding
redshift interval of the same index number in chop

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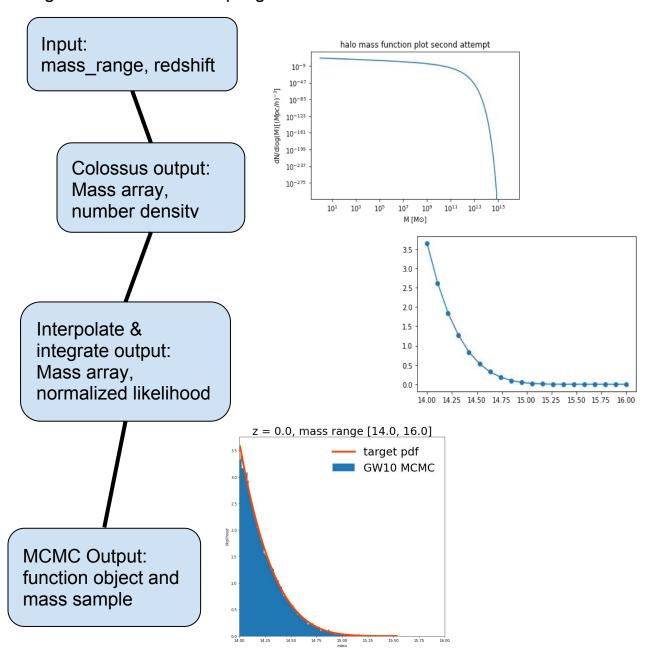
### Step:

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- Call helper 1 to obtain redshift distribution array
- Based on redshift range, divide it to several chops of redshift, each chop used as limit
- Loop through the redshifts array given by helper 1, approximate redshifts between two chops to the lower limit of it.
- Use another NumPy array of the same size to keep track of cluster number in each interval

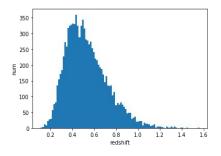
# **Procedure**

Function A [mass\_sampling] Single redshift mass sampling



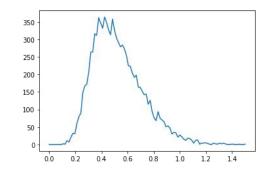
Input: mass\_range, redshift range, sample size

Function B [mul\_redshift\_mass\_sampling] Multiple redshift



[Skew\_sample] output: redshift array based on skewed gaussian distribution imitating

[single\_redshift\_num] output: a chop array of redshift interval, a num\_per\_redshift array of cluster number/redshift interval



Called [mass\_sampling] to sample for each redshift in chop

Output: a pandas dataframe with mass and redshift as two columns

