

Main

July 13, 2018

```
In [1]: import halotools
import numpy as np
import matplotlib.pyplot as plt
from scipy.optimize import curve_fit
%matplotlib inline
```

1 1) Import data

```
In [2]: from halotools import sim_manager
from halotools.sim_manager import CachedHaloCatalog
halocat = CachedHaloCatalog(simname = 'bolplanck', halo_finder = 'rockstar')
```

We begin with the analysis of a subsample of the Bolshoi simulation with Planck cosmology as of 2013 ($h=0.7$, $\Omega_\Lambda=0.69289$, $\Omega_m = 0.30711$, $\Omega_b = 0.048$, $n_s = 0.96$, $\sigma_8 = 0.82$). The simulation started at $z_{ini} = 80$ and followed 2048^3 particles of mass $M_{dm} = 1.55 \times 10^8 M_{sun}/h$ in their evolution in a volume of boxsize $L=250$ Mpc/h.

```
In [3]: #Catalog of halos itself
halos = halocat.halo_table
#To see what properties are available
print(halos.keys())

['halo_vmax_firstacc', 'halo_dmvir_dt_tdyn', 'halo_macc', 'halo_scale_factor', 'halo...
```

2 2) Organize the data into samples

```
In [4]: from halotools.utils import group_member_generator
#Group halos by halo host id
halos.sort('halo_hostid')
grouping_key = 'halo_hostid'
requested_columns = []
group_gen = group_member_generator(halos, grouping_key, requested_columns)

#calculates the number of subalos per host
nsub = np.zeros(len(halos))
for first, last, member_props in group_gen:
```

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        nsub[first:last] = last - first - 1
#Adds a new column to the ith-row of the table which gives the number of subhalos
        halos['num_subhalos'] = nsub

```

```

In [5]: NumberHosthalos=0
        NumberSubhalos=0
        for i in range(0,len(halos)):
            if halos["halo_pid"][i]==-1:
                NumberHosthalos= NumberHosthalos+1
            else:
                NumberSubhalos = NumberSubhalos+1

        print("Number of halos in total, unfiltered bolplank catalog", len(halos))
        print("Number of hosthalos, unfiltered bolplank catalog", NumberHosthalos)
        print("Number of subhalos, unfiltered bolplank catalog", NumberSubhalos)

```

```

Number of halos in total, unfiltered bolplank catalog 1444262
Number of hosthalos, unfiltered bolplank catalog 1078614
Number of subhalos, unfiltered bolplank catalog 365648

```

```

In [9]: #Attribute Subhalos with the Positions of their respective host in order to
        #center of mass frame:
        halos.add_column(halos["halo_x"], name="Host_x")
        halos.add_column(halos["halo_y"], name="Host_y")
        halos.add_column(halos["halo_z"], name="Host_z")
        halos.add_column(halos["halo_vx"], name="Host_vx")
        halos.add_column(halos["halo_vy"], name="Host_vy")
        halos.add_column(halos["halo_vz"], name="Host_vz")

```

The catalog is filtered by the following criteria. We generate two samples :

Sample 1 consists of all host halos with $M_{vir} < 10^{15} M_{sun}$ and with $M_{vir}/M_{Dm,partcl} > 100$ together with the subhalos with $M_{vir}/M_{Dm,partcl} > 100$, $M_{vir}/M_{Vir,Host} > 10^{-3}$ and with $M_{Vir,Host} < 10^{15} M_{sun}$.

Sample 2 is constrained under the same criteria as Sample 1, but one of the conditions on the subhalos, namely $M_{vir}/M_{Vir,Host} > 10^{-3}$, is removed.

```

In [6]: ##We consider two main samples:
        ##Sample 1: Consider all halos consisting of more than 100 dm particles and
        ##Sample 2: Take into account all halos and still exclude hosts with less than
        #The mask for the hosts
        M_dm_particle = 1.55e8
        host_maskSample1 = ((halos['halo_upid'] == -1) & (halos["halo_mvir"]<1e15))
        hostsSample1 = halos[host_maskSample1]

        #mask for the subhalos

        sub_maskSample1 = ((halos["halo_upid"]>0)&((halos["halo_mvir"]/halos["halo_

```

```

subsSample1 = halos[sub_maskSample1]
sub_maskSample2 = ((halos["halo_upid"]>0)&(halos["halo_mvir_host_halo"]< 1e
subsSample2 = halos[sub_maskSample2]

print("Number of hosthalos in Sample 1 = ", len(hostsSample1))
print("The mean concentration of host halos in sample 1 is  ", "r$ \bar{c}$
print("Number of subhalos in Sample 1 = ", len(subsSample1))
print("Number of subhalos in Sample 2 = ", len(subsSample2))

```

C:\Users\ThimoPreis\Anaconda3\lib\site-packages\ipykernel__main__.py:12: RuntimeWarning

```

Number of hosthalos in Sample 1 = 1071702
The mean concentration of host halos in sample 1 is  r$ ar{c}$ = 16.0799
Number of subhalos in Sample 1 = 224818
Number of subhalos in Sample 2 = 289020

```

In []:

3 3) Radial density profile

```

In [7]: from halotools.mock_observables import return_xyz_formatted_array

host_pos_sample1 = return_xyz_formatted_array(hostsSample1['halo_x'], hostsSample1['halo_y'], hostsSample1['halo_z'])

sub_pos_sample1 = return_xyz_formatted_array(subsSample1['halo_x'], subsSample1['halo_y'], subsSample1['halo_z'])
sub_pos_sample2 = return_xyz_formatted_array(subsSample2['halo_x'], subsSample2['halo_y'], subsSample2['halo_z'])

hx1 = hostsSample1["halo_x"]
hy1 = hostsSample1["halo_y"]
hz1 = hostsSample1["halo_z"]
sx1 = subsSample1["halo_x"]
sy1 = subsSample1["halo_y"]
sz1 = subsSample1["halo_z"]
sx2 = subsSample2["halo_x"]
sy2 = subsSample2["halo_y"]
sz2 = subsSample2["halo_z"]

hvx1 = hostsSample1["halo_vx"]
hvy1 = hostsSample1["halo_vy"]
hvz1 = hostsSample1["halo_vz"]
svx1 = subsSample1["halo_vx"]
svy1 = subsSample1["halo_vy"]
svz1 = subsSample1["halo_vz"]
svx2 = subsSample2["halo_vx"]
svy2 = subsSample2["halo_vy"]
svz2 = subsSample2["halo_vz"]

```

```

In [9]: from halotools.mock_observables import radial_profile_3d

rbins_normalized = np.linspace(0.001, 1, 100)
rbins_midpoints = (rbins_normalized[:-1] + rbins_normalized[1:])/2

Density_profileSample1, CountsSample1 = radial_profile_3d(host_pos_sample1,
    rbins_normalized = rbins_normalized,
    normalize_rbins_by = hostsSample1['halo_rvir'],
    period=halocat.Lbox, return_counts = True)

Density_profileSample2, CountsSample2 = radial_profile_3d(host_pos_sample1,
    rbins_normalized = rbins_normalized,
    normalize_rbins_by = hostsSample1['halo_rvir'],
    period=halocat.Lbox, return_counts = True)

In [10]: shell_volumes = 4/3*np.pi*(rbins_normalized[1:]**3-rbins_normalized[:-1]**3)
MeanNumberDensitySample1 = (CountsSample1/float(len(subsSample1)))/shell_volumes
MeanNumberDensitySample2 = (CountsSample2/float(len(subsSample2)))/shell_volumes

In [19]: #Model NFW profil of the hosts
#from halotools import empirical_models
#from halotools.empirical_models import NFWProfile

#nfw = NFWProfile(conc_mass_model='direct_from_halo_catalog', mdef='vir', c=100)
#model_conc = nfw.conc_NFWmodel(table=hostsSample1)

#Write NFW profile function yourself:
from halotools.empirical_models import density_threshold
from astropy import cosmology
from astropy.cosmology import LambdaCDM
cosmo = LambdaCDM(H0=70, Om0=0.3, Ode0=0.7)
m_dm=1.55e8

def NFW(r):
    return density_threshold(cosmo, 0, mdef="vir")*0.3/3*np.mean(hostsSample1)

rhocrit = cosmo.critical_density0
#Another definition, completely equivalent to NFW, not necessary to use
c=np.mean(hostsSample1["halo_nfw_conc"])

def nfwProfile(r):
    return (200*rhocrit*c)/3/(np.log(1+c-c/(1+c))) / (r/np.mean(hostsSample1))

#r_values_hosts=np.linspace(0.001,1,len(hostsSample1))
#rho_nfw = nfw.dimensionless_mass_density(scaled_radius =r_values_hosts, c=c)

```

```

In [25]: plt.rc('text', usetex=True)
         plt.rc('font', family='serif')

         r_values = np.linspace(0.001, 1, 1000)

         # Create a new figure
         fig = plt.figure(figsize=(6, 6), dpi=80)
         plot_profile = fig.add_subplot(111)
         plot_profile.set_xscale('log')
         plot_profile.set_yscale('log')
         plot_profile.grid(True)

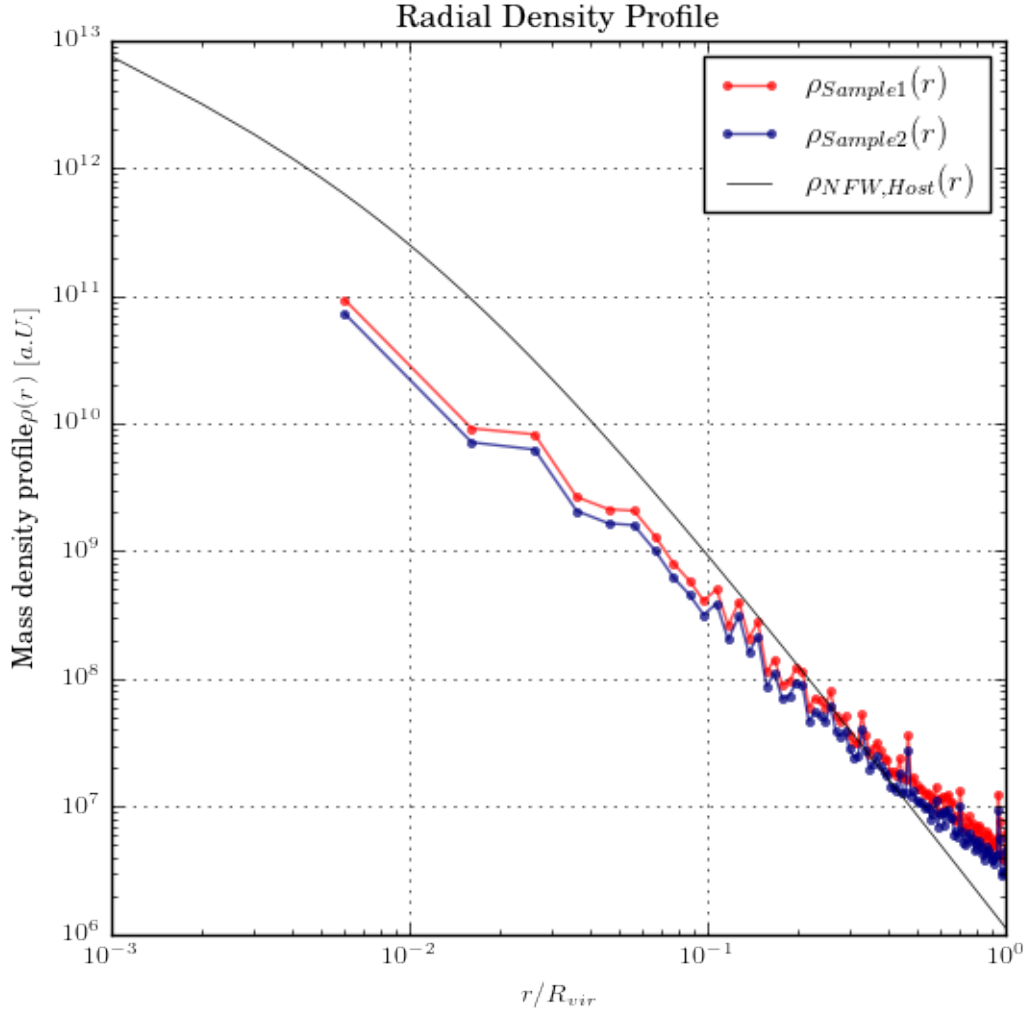
         #plot_ps.set_xlim(1.0e-5, 1.0e5)
         #plot_profile.set_ylim(0.001, 1e3)

         plot_profile.set_xlabel(r' $ r/R_{\text{vir}} $ ')
         plot_profile.set_ylabel(r'Mass density profile $\rho(r)$ [a.u.]')

         plot_profile.plot(rbins_midpoints, Density_profileSample1/shell_volumes/fig)
         plot_profile.plot(rbins_midpoints, Density_profileSample2/shell_volumes/fig)
         #plot_profile.plot(r_values_hosts, rho_nfw, color="black", marker=".", label=)
         plot_profile.plot(r_values, NFW(r_values)/4e3, color="black", lw=0.5, label=)

         plot_profile.legend(prop={'size': 12}, loc="best")
         plt.title("Radial Density Profile")
         # Save figure using 80 dots per inch
         plt.savefig("RadialDensityProfile.pdf", dpi=80)

```



```
In [26]: # Create a new figure
fig = plt.figure(figsize=(6,6), dpi=80)
plot_profile = fig.add_subplot(111)
plot_profile.set_xscale('log')
plot_profile.set_yscale('log')
plot_profile.grid(True)

#plot_ps.set_xlim(1.0e-5, 1.0e5)
#plot_profile.set_ylim(0.001, 1e3)

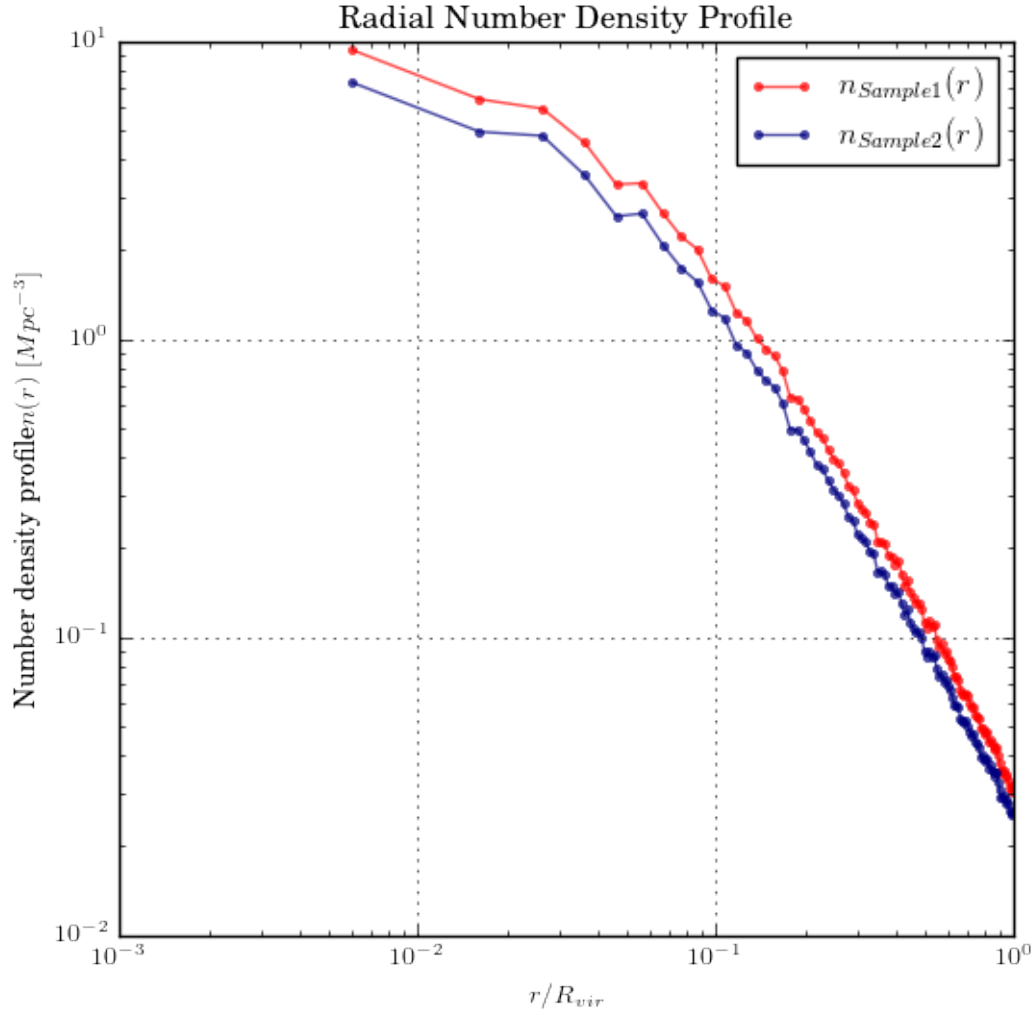
plot_profile.set_xlabel(r' $ r/R_{\text{vir}} $ ')
plot_profile.set_ylabel(r'Number density profile$\rho(r)$ \; [Mpc$^{-3}$]$')

plot_profile.plot(rbins_midpoints, MeanNumberDensitySample1, color="red",
plot_profile.plot(rbins_midpoints, MeanNumberDensitySample2, color="navy",
```

```

plot_profile.legend(prop={'size':12}, loc="best")
plt.title("Radial Number Density Profile")
# Save figure using 80 dots per inch
plt.savefig("RadialNumberDensityProfile.pdf",dpi=80)

```



4 4)Velocity profiles

```

In [ ]: #Replace every position and velocity of the subhalos by the position and velocity
        #frame of their respective host halo:

```

```

for i in range(0, len(hostsSample1)):

```

```

for j in range(0, len(subsSample1)):

    if hostsSample1["halo_id"][i]==subsSample1["halo_hostid"][j]:

        sx1[j] = sx1[j]-hx1[i]
        sy1[j] = sy1[j]-hy1[i]
        sz1[j] = sz1[j]-hz1[i]
        svx1[j] = svx1[j]-hvx1[i]
        svy1[j] = svy1[j]-hvy1[i]
        svz1[j] = svz1[j]-hvz1[i]

    else:

        sx1[j] = 10000
        sy1[j] = 10000
        sz1[j] = 10000
        svx1[j] = 10000
        svy1[j] = 10000
        svz1[j] = 10000

for m in range(0, len(subsSample2)):

    if hostsSample1["halo_id"][i]==subsSample2["halo_hostid"][m]:

        sx2[m] = sx2[m]-hx2[i]
        sy2[m] = sy2[m]-hy2[i]
        sz2[m] = sz2[m]-hz2[i]
        svx2[m] = svx2[m]-hvx2[i]
        svy2[m] = svy2[m]-hvy2[i]
        svz2[m] = svz2[m]-hvz2[i]

    else:

        sx1[m] = 10000
        sy1[m] = 10000
        sz1[m] = 10000
        svx1[m] = 10000
        svy1[m] = 10000
        svz1[m] = 10000

```


5 Put every component to zero, when subhalo does not have a corresponding host halo

```
for i in range(0,len(subsSample1)): if indexReturn(subsSample1["halo_hostid"][i])=="NOT-
FOUND": sx1[i] = 0 sy1[i] = 0 sz1[i] = 0 svx1[i] = 0 svy1[i] = 0 svz1[i] = 0
```

```
else:
```

```
    sx1[i] = sx1[i]- hx1[indexReturn(subsSample1["halo_upid"][i])]
    sy1[i] = sy1[i]- hy1[indexReturn(subsSample1["halo_upid"][i])]
    sz1[i] = sz1[i]- hz1[indexReturn(subsSample1["halo_upid"][i])]
    svx1[i] = svx1[i]- hvx1[indexReturn(subsSample1["halo_upid"][i])]
    svy1[i] = svy1[i]- hvy1[indexReturn(subsSample1["halo_upid"][i])]
    svz1[i] = svz1[i]- hvz1[indexReturn(subsSample1["halo_upid"][i])]
```

```
for k in range(0,len(subsSample2)):
```

```
if indexReturn(subsSample1["halo_hostid"][i])=="NOTFOUND":
```

```
    sx2[k] = 0
    sy2[k] = 0
    sz2[k] = 0
    svx2[k] = 0
    svy2[k] = 0
    svz2[k] = 0
```

```
else:
```

```
    sx2[k] = sx2[k]- hx1[indexReturn(subsSample2["halo_upid"][k])]
    sy2[k] = sy2[k]- hy1[indexReturn(subsSample2["halo_upid"][k])]
    sz2[k] = sz2[k]- hz1[indexReturn(subsSample2["halo_upid"][k])]
    svx2[k] = svx2[k]- hvx1[indexReturn(subsSample2["halo_upid"][k])]
    svy2[k] = svy2[k]- hvy1[indexReturn(subsSample2["halo_upid"][k])]
    svz2[k] = svz2[k]- hvz1[indexReturn(subsSample2["halo_upid"][k])]
```

```
In [12]: #Go into spherical coordinates
```

```
    r_s1 = np.sqrt(sx1**2+sy1**2+sz1**2)
    r_s2 = np.sqrt(sx2**2+sy2**2+sz2**2)
```

```
    vr_s1 = (sx1*svx1+sy1*svy1+sz1*svz1)/r_s1
    vr_s2 = (sx2*svx2+sy2*svy2+sz2*svz2)/r_s2
```

```
    vphi_s1 = (svx1*sy1-sx1*svy1)/(sx1**2+sy1**2)
    vphi_s2 = (svx2*sy2-sx2*svy2)/(sx2**2+sy2**2)
```

```
    vth_s1 = (sz1*(sx1*svx1+sy1*svy1)-svz1*(sx1**2+sy1**2))/np.sqrt(sx1**2+sy1**2)
    vth_s2 = (sz2*(sx2*svx2+sy2*svy2)-svz2*(sx2**2+sy2**2))/np.sqrt(sx2**2+sy2**2)
```

```
In [14]: #Average the radial velocity profiles
```

```

from halotools.mock_observables import radial_profile_3d

rbins_normalized = np.linspace(0.001, 1, 100)
rbins_midpoints = (rbins_normalized[:-1] + rbins_normalized[1:])/2

aver_vr_s1 = radial_profile_3d(host_pos_sample1, sub_pos_sample1, vr_s1,
                               rbins_normalized = rbins_normalized,
                               normalize_rbins_by = hostsSample1['halo_rvir'],
                               period=halocat.Lbox)

aver_vr_s2 = radial_profile_3d(host_pos_sample1, sub_pos_sample2, vr_s2,
                               rbins_normalized = rbins_normalized,
                               normalize_rbins_by = hostsSample1['halo_rvir'],
                               period=halocat.Lbox)

aver_vphi_s1 = radial_profile_3d(host_pos_sample1, sub_pos_sample1, vphi_s1,
                                  rbins_normalized = rbins_normalized,
                                  normalize_rbins_by = hostsSample1['halo_rvir'],
                                  period=halocat.Lbox)

aver_vphi_s2 = radial_profile_3d(host_pos_sample1, sub_pos_sample2, vphi_s2,
                                  rbins_normalized = rbins_normalized,
                                  normalize_rbins_by = hostsSample1['halo_rvir'],
                                  period=halocat.Lbox)

aver_vth_s1 = radial_profile_3d(host_pos_sample1, sub_pos_sample1, vth_s1,
                                 rbins_normalized = rbins_normalized,
                                 normalize_rbins_by = hostsSample1['halo_rvir'],
                                 period=halocat.Lbox)

aver_vth_s2 = radial_profile_3d(host_pos_sample1, sub_pos_sample2, vth_s2,
                                 rbins_normalized = rbins_normalized,
                                 normalize_rbins_by = hostsSample1['halo_rvir'],
                                 period=halocat.Lbox)

```

```

In [24]: #Normalize by virial velocity
np.max(sx2)

```

```

Out[24]: 0.0

```

```

In [16]: # Create a new figure
fig = plt.figure(figsize=(6,6), dpi=80)
plot_profile = fig.add_subplot(111)
plot_profile.set_xscale('log')
#plot_profile.set_yscale('log')
plot_profile.grid(True)

```

```

#plot_ps.set_xlim(1.0e-5, 1.0e5)
#plot_profile.set_ylim(0.001, 1e3)

plot_profile.set_xlabel(r' $ r/R_{\text{vir}} $')
plot_profile.set_ylabel(r'$\langle v(r) \rangle / V_{\text{vir}}$')

plot_profile.plot(rbins_midpoints, aver_vr_s1, color="red", marker=".", label=
plot_profile.plot(rbins_midpoints, aver_vphi_s1, color="navy", marker=".", label=
#plot_profile.plot(r_values_hosts, rho_nfw, color="black", marker=".", label=
plot_profile.plot(rbins_midpoints, aver_vth_s1, color="black", lw=0.5, label=

plot_profile.legend(prop={'size':12}, loc="best")
plt.title("Radial Average Velocity Profile Sample 1")
# Save figure using 80 dots per inch
plt.savefig("RadialVelocityProfilesSample1.pdf", dpi=80)

```

ValueError Traceback (most recent call last)

```

<ipython-input-16-e09fe14ad7b0> in <module>()
    12 plot_profile.set_ylabel(r'$\langle v(r) \rangle / V_{\text{vir}}$')
    13
--> 14 plot_profile.plot(rbins_midpoints, aver_vr_s1, color="red", marker=".", label=
    15 plot_profile.plot(rbins_midpoints, aver_vphi_s1, color="navy", marker=".", label=
    16 #plot_profile.plot(r_values_hosts, rho_nfw, color="black", marker=".", label=

C:\Users\ThimoPreis\Anaconda3\lib\site-packages\matplotlib\__init__.py in 
1816 warnings.warn(msg % (label_namer, func.__name__),
1817                 RuntimeWarning, stacklevel=2)
-> 1818 return func(ax, *args, **kwargs)
1819 pre_doc = inner.__doc__
1820 if pre_doc is None:

C:\Users\ThimoPreis\Anaconda3\lib\site-packages\matplotlib\axes\_axes.py in 
1384 lines.append(line)
1385
-> 1386 self.autoscale_view(scalex=scalex, scaley=scaley)
1387 return lines
1388

```

C:\Users\ThimoPreis\Anaconda3\lib\site-packages\matplotlib\axes_base.py in

```

2166             x1 += delta
2167         if not _tight:
-> 2168             x0, x1 = xlocator.view_limits(x0, x1)
2169         self.set_xbound(x0, x1)
2170

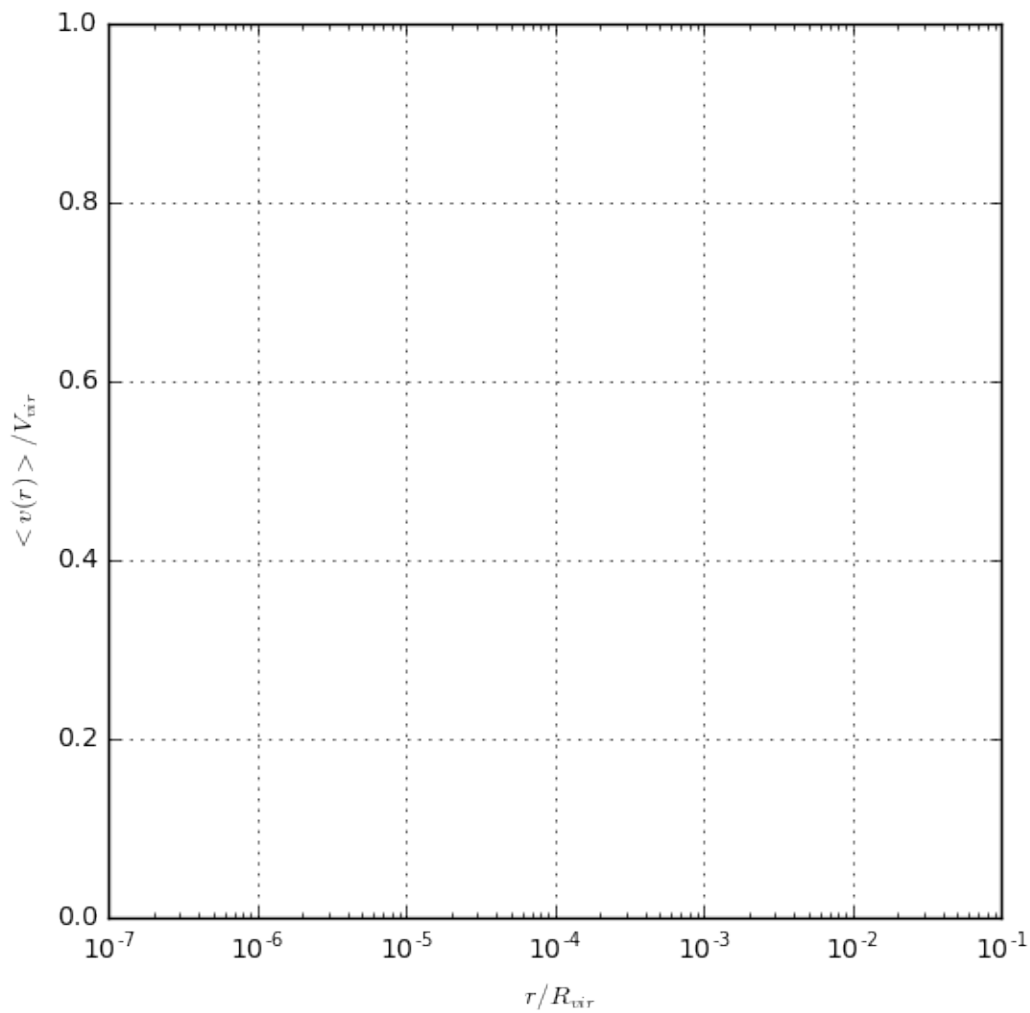
```

```

C:\Users\ThimoPreis\Anaconda3\lib\site-packages\matplotlib\ticker.py in view
1615         if minpos <= 0 or not np.isfinite(minpos):
1616             raise ValueError(
-> 1617                 "Data has no positive values, and therefore can not be
1618                 "log-scaled.")
1619

```

ValueError: Data has no positive values, and therefore can not be log-scaled



```

In [ ]: # Create a new figure
fig = plt.figure(figsize=(6,6), dpi=80)
plot_profile = fig.add_subplot(111)
plot_profile.set_xscale('log')
plot_profile.set_yscale('log')
plot_profile.grid(True)

#plot_ps.set_xlim(1.0e-5, 1.0e5)
#plot_profile.set_ylim(0.001, 1e3)

plot_profile.set_xlabel(r' $ r/R_{\text{vir}} $ ')
plot_profile.set_ylabel(r' $ \langle v(r) \rangle / V_{\text{vir}} $ ')

plot_profile.plot(rbins_midpoints, aver_vr_s1, color="red", marker=".", label=)
plot_profile.plot(rbins_midpoints, aver_vphi_s1, color="navy", marker=".", label=)
#plot_profile.plot(r_values_hosts, rho_nfw, color="black", marker=".", label=)
plot_profile.plot(rbins_midpoints, aver_vth_s1, color="black", lw=0.5, label=)

plot_profile.legend(prop={'size':12}, loc="best")
plt.title("Radial Average Velocity Profile Sample 2")
# Save figure using 80 dots per inch
plt.savefig("RadialVelocityProfilesSample2.pdf",dpi=80)

```

In []:

In [52]:

```

2821926828
1071307
243.125
243.142
0.0174408

```

In []:

In []:

In []:

In []:

In []:

In []: