Problem Set 01

AST 8110, Fall 2022

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```
In [1]: import time
        bigstart = time.time()
        ## Importing packages (some of these may not be useful)
        import numpy as np
        import matplotlib.pyplot as plt
        import pandas as pd
        import seaborn as sns
        import scipy.stats as stats
        #import numba
        import sys
        import os
        # import astropy
        # import astropy.units as u
        # import astropy.constants as const
        # import astropy.coordinates as coord
        import random
        from tqdm import tqdm
        #from numba import jit
        from scipy.stats import binned statistic as binstat
        ## Set Seed
        random.seed(8110)
        ## debug mode for fast testing
        debug = False
        filt = 10 ## factor to reduce the size of the data set by
```

```
In [2]: ## read in data
        data = np.loadtxt('pset1data.txt')
        if debug:
             data = data[::filt]
        ## plot data
        def plot_data(dat=data):
            fig, ax = plt.figure(figsize=(5,5)), plt.gca()
            ax.minorticks on()
            ax.tick_params(axis='both', which='major',direction='in',top=True,right=True,length=10,width
            ax.tick_params(axis='both', which='minor',direction='in',top=True,right=True, length=5,width
             ax.plot(dat[:,0],dat[:,1],'.',markersize=2,color='black');
             plt.xlim(-90,90);
             plt.ylim(-90,90);
             plt.xlabel('x');
             plt.ylabel('y');
             return fig, ax
        plot_data(dat=data);
```

```
80 60 40 20 0 20 40 60 80 x
```

```
In [3]:
       ## Generate Random galaxy positions
        datSize = len(data)
        galMultiplier = 2
        def gen_galaxy_positions(N=datSize*galMultiplier,pkl=False,save=False,timeit=True):
             start_time = time.time() if timeit else None
            Generates N random galaxy positions
             if pkl and os.path.isfile(pkl):
                data = pd.read_pickle(pkl)
                return data
             else:
                x = np.random.uniform(-90,80,N)
                y = np.random.uniform(-80,90,N)
                data = np.column_stack((x,y))
                if save:
                    pd.to_pickle(data, save)
                print("--- %s seconds ---" % (time.time() - start_time)) if timeit else None
                return data
In [4]: ## import random galaxy positions previously generated and plot them
```

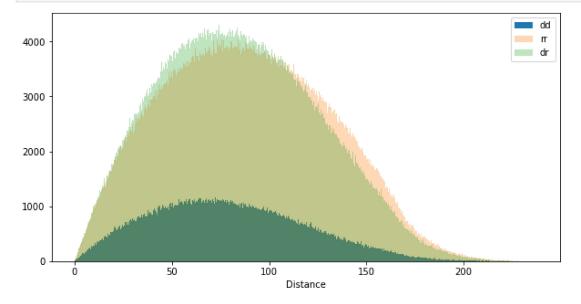
```
In [4]: ## import random galaxy positions previously generated and plot them
  randData = gen_galaxy_positions()
  if debug:
        randData = randData[::filt]
  fig, ax = plot_data(dat=data);
  ax.plot(randData[:,0],randData[:,1],'+',markersize=1,color='red',alpha=0.5);
```

--- 0.0005009174346923828 seconds ---

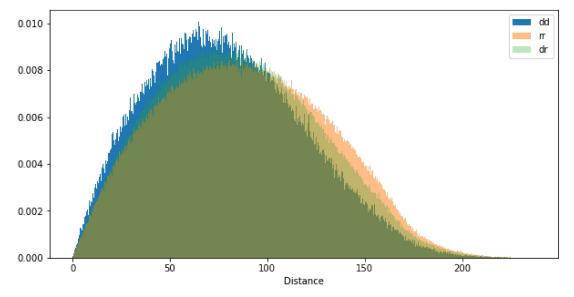
```
80 60 40 20 0 20 40 60 80 x
```

```
In [5]: ## calculate distances to other points
        def get_dist(mode='dd',data1=None,data2=None,pkl=False,save=False, timeit=True):
             start_time = time.time() if timeit else None
             if pkl and os.path.isfile(pkl):
                 dist = pd.read pickle(pkl)
                 return dist
             elif data1 is not None and data2 is not None: ##case where I want to explicitly pass in data
                 d1 = data1
                 d2 = data2
             elif mode == 'dd':
                 d1 = data
                 d2 = data
             elif mode =='rr':
                 d1 = randData
                 d2 = randData
             elif mode == 'dr':
                 d1 = data
                 d2 = randData
             dist = np.linalg.norm(d2[:,None]-d1,axis=-1) ## calculates distances between all points in d
             if mode == 'dd' or mode == 'rr': ## only take upper triangle of matrix to prevent double cou
                 offset = 1 ## offset to remove self distances in dd and rr (i.e. distance between a poin
                 slice_idx = np.triu_indices_from(dist,k=offset) ## get indices for upper triangle of mat
                 #print(slice idx)
                 dist = dist[slice_idx] ## get upper triangle of matrix using indices
             elif mode == 'dr':
                 dist = dist.flatten()
             if save:
                 pd.to_pickle(dist, save)
             print("--- %s seconds ---" % (time.time() - start_time)) if timeit else None
             return dist
In [6]: ## calculate distances
        dd = get_dist('dd')
        rr = get_dist('rr')
        dr = get_dist('dr')
        --- 0.04443359375 seconds ---
         --- 0.1658482551574707 seconds ---
        --- 0.05286979675292969 seconds ---
        ## plot histograms of distances
In [7]:
        fig, ax = plt.figure(figsize=(10,5)), plt.gca()
        plt.hist(dd,bins=1000,density=False,alpha=1,label='dd');
```

```
plt.hist(rr,bins=1000,density=False, alpha=0.3,label='rr');
plt.hist(dr,bins=1000,density=False, alpha=0.3,label='dr');
plt.legend();
plt.xlabel('Distance');
plt.show()
## note here: you need to divide by the number of pairs so that you have the same number of pair
```



```
In [8]: ## do another histogram, but normalized
fig, ax = plt.figure(figsize=(10,5)), plt.gca()
plt.hist(dd,bins=1000,density=True,alpha=1,label='dd');
plt.hist(rr,bins=1000,density=True, alpha=0.5,label='rr');
plt.hist(dr,bins=1000,density=True, alpha=0.3,label='dr');
plt.legend();
plt.xlabel('Distance');
plt.show()
```



```
In [9]: ## develop correlation function
maxBin = 230
binWidth = 2
binArray = np.linspace(0,maxBin,int(maxBin/binWidth)+1) ## the bins to use for the correlation f
#print(binArray)
## calculate bin averages
def get_bin_averages(dataset,binWidth=binWidth,double=False,pkl=False,save=False, timeit=True):
    # startBin = binWidth*round(min(dataset)/binWidth)
    # endBin = binWidth*round(max(dataset)/binWidth)
```

```
# bin = np.arange(startBin,endBin,binWidth)
              #print('Bin Array:',bin)
              start time = time.time() if timeit else None
              bin = binArray
              if pkl and os.path.isfile(pkl):
                 binVals = pd.read pickle(pkl)
                 return bin, binVals
              binVals, sp bin, spc binNum = binstat(dataset,dataset,statistic='count',bins=bin) ## efficie
              binVals = binVals/len(dataset) ## normalize by number of points put in
              # if double: ## accounts for the 2DR case ## 09/20 update: not needed
                   binAvqs = binAvqs*2
             if save:
                 pd.to_pickle(binVals, save)
              print("--- %s seconds ---" % (time.time() - start_time)) if timeit else None
              return bin[:-1], binVals ## remove the max bin value (because binstat returns the bin edges
In [10]:
         dd bin, dd val = get bin averages(dd)
         rr_bin, rr_val = get_bin_averages(rr)
         dr_bin, dr_val = get_bin_averages(dr)
         --- 0.019565343856811523 seconds ---
         --- 0.08364009857177734 seconds ---
         --- 0.07717728614807129 seconds ---
In [11]: ## plot bin and vals for dd, rr, and dr
         fig, ax = plt.figure(figsize=(10,5)), plt.gca()
         plt.plot(dd_bin,dd_val,label='dd');
         plt.plot(rr_bin,rr_val,label='rr');
         plt.plot(dr_bin,dr_val,label='dr');
         plt.legend();
         plt.show()
         ## plot histogram of dd, rr, and dr
         fig, ax = plt.figure(figsize=(10,5)), plt.gca()
         plt.hist(dd_val,density=True,alpha=1,label='dd');
         plt.hist(rr_val,density=True, alpha=0.5,label='rr');
         plt.hist(dr_val,density=True, alpha=0.3,label='dr');
         plt.legend();
         plt.show(); ## does this look weird?
         0.0200
                                                                                            dd
                                                                                            IT
         0.0175
          0.0150
          0.0125
          0.0100
         0.0075
          0.0050
          0.0025
```

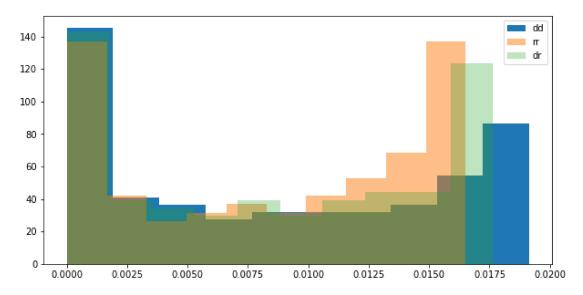
0.0000

50

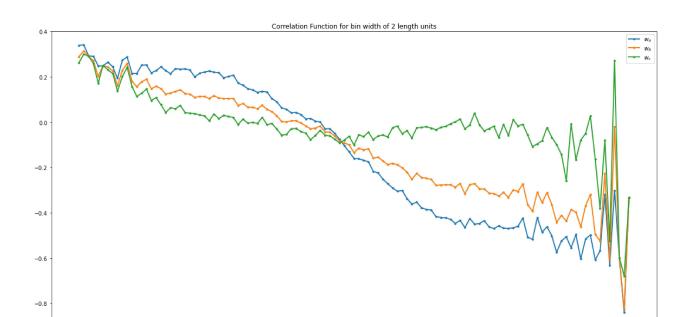
100

150

200



```
## calculate correlation function
In [12]:
         # def get_corr_func(dd,rr,dr,binWidth=binWidth,pkl=False,save=False, timeit=True):
               start time = time.time() if timeit else None
         #
               if pkl and os.path.isfile(pkl):
         #
                   corrFunc = pd.read_pickle(pkl)
         #
                   return corrFunc
         #
               corrFunc = None
         #
               if save:
         #
                   pd.to_pickle(corrFunc, save)
         #
               print("--- %s seconds ---" % (time.time() - start_time)) if timeit else None
               return corrFunc
         w_a = [dd_i/rr_i -1 for dd_i,rr_i in zip(dd_val,rr_val)]
         w_b = [dd_i/dr_i -1 for dd_i,dr_i in zip(dd_val,dr_val)]
         w_c = [(dd_i - 2*dr_i +rr_i)/(rr_i) for dd_i,dr_i,rr_i in zip(dd_val,dr_val,rr_val)] ## w_c seem
         ## plot correlation function (1 instance)
         fig, ax = plt.figure(figsize=(20,10)), plt.gca()
         \verb|plt.plot(|dd_bin,w_a,label=r'$w_a$',linestyle='-',linewidth=2, marker='o',markersize=3);|
         plt.plot(dr_bin,w_b,label=r'$w_b$',linestyle='-',linewidth=2, marker='o',markersize=3,alpha=1);
         plt.plot(dr_bin,np.array(w_c),label=r'$w_c$',linestyle='-',linewidth=2, marker='o',markersize=3,
         plt.legend();
         plt.xlabel('Distance');
         plt.title('Correlation Function for bin width of %s length units'%str(binWidth));
```

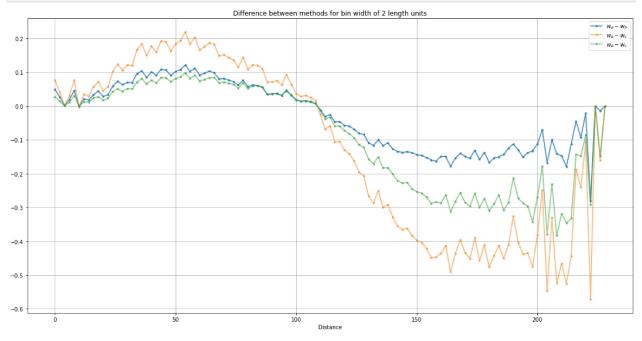


Distance

150

200

```
In [13]: ## plot differences between methods
    fig, ax = plt.figure(figsize=(20,10)), plt.gca()
    plt.plot(dd_bin,np.array(w_a)-np.array(w_b),label=r'$w_a-w_b$',linestyle='-',linewidth=2, marker
    plt.plot(dd_bin,np.array(w_a)-np.array(w_c),label=r'$w_a-w_c$',linestyle='-',linewidth=2, marker
    plt.plot(dd_bin,np.array(w_b)-np.array(w_c),label=r'$w_b-w_c$',linestyle='-',linewidth=2, marker
    plt.legend();
    plt.xlabel('Distance');
    plt.title('Difference between methods for bin width of %s length units'%str(binWidth));
    plt.grid()
    ## could probably normalize this somehow to get a better idea of the differences
```

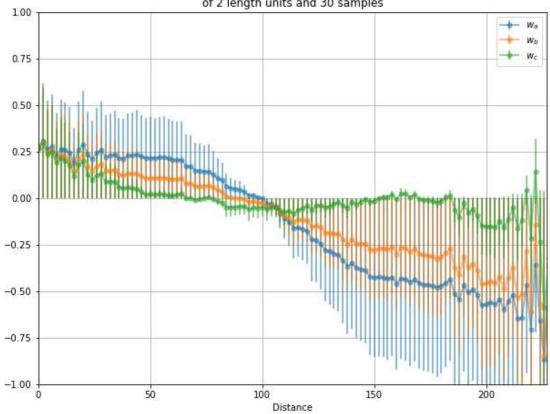


```
In [14]: ## run 10-30 times to calculate the rms error of the correlation function for each bin

def get_rms_error(mode=None,n=10,plot=True,save=False,fun=False):
    start_time = time.time()
    randGals = [gen_galaxy_positions(N=10000,timeit=False) for i in range(n)]
    #if mode == 'w_a':
    memDebug = False
    dd = [get_dist('dd',data,data, timeit=False) for i in range(n)] ## I know this is redundant,
    rr = [get_dist('rr',r1, r2,timeit=False) for r1,r2 in zip(randGals,randGals)]
```

```
dr = [get dist('dr',data,r2, timeit=False) for r2 in randGals]
print('1') if memDebug else None
dd get bins = [get bin averages(d i, timeit=False) for d i in dd]
dd bin, dd avg = zip(*dd get bins)
rr_get_bins = [get_bin_averages(r_i, timeit=False) for r_i in rr]
rr_bin, rr_avg = zip(*rr_get_bins)
dr_get_bins = [get_bin_averages(d_i, timeit=False) for d_i in dr]
dr_bin, dr_avg = zip(*dr_get_bins)
print('2') if memDebug else None
## commented out was causing a memory error
\# w_a, w_b, w_c = [], [], []
# for dd_i, rr_i, dr_i in zip(dd_avg, rr_avg, dr_avg):
         #print(i)
          w_a.append([dd_j/rr_j -1 for dd_j,rr_j in zip(dd_i,rr_i)])
          w_b.append([dd_j/dr_j -1 for dd_j,dr_j in zip(dd_i,dr_i)])
          w_c.append([(dd_j - 2*dr_j + rr_j)/(rr_j) for dd_j,dr_j,rr_j in zip(dd_i,dr_i,rr_i)])
w_a = np.array([[dd_j/rr_j -1 for dd_j,rr_j in zip(dd_i,rr_i)] for dd_i, rr_i, dr_i in zip(d
w_b = np.array([[dd_j/dr_j -1 for dd_j,dr_j in zip(dd_i,dr_i)] for dd_i, rr_i, dr_i in zip(d
w_c = np_array([[(dd_j - 2*dr_j + rr_j)/(rr_j) \text{ for } dd_j, dr_j, rr_j \text{ in } zip(dd_i, dr_i, rr_i)] \text{ for } dd_j, dr_j, dr_j,
print('3') if memDebug else None
\# w_a = np.array(w_a)
\# w b = np.array(w b)
\# w_c = np.array(w_c)
## actually, you want to average the value of each bin when calculating <
#print(w a)
## could do this across axis 0 instead I think
# print(w_a.shape)
# print(w_b.shape)
# print(w_c.shape)
avg_w_a = np.mean(w_a,axis=0);
avg_w_b = np.mean(w_b,axis=0);
avg_w_c = np.mean(w_c,axis=0);
print('4') if memDebug else None
rms_w_a = np.sqrt(np.mean((w_a)**2,axis=0));
rms_w_b = np.sqrt(np.mean((w_b)**2,axis=0));
rms_w_c = np.sqrt(np.mean((w_c)**2,axis=0));
print('5') if memDebug else None
\# rms\_w\_a\_unzip = np.array([[item[i] for item in w\_a] for i in range(len(w\_a[0]))]) ## worka
# rms_w_a = [np.sqrt(np.mean(item**2)) for item in rms_w_a_unzip]
\# rms_wb_unzip = np.array([[item[i] for item in w_b] for i in range(len(w_b[0]))])
# rms_w_b = [np.sqrt(np.mean(item**2)) for item in rms_w_b_unzip]
# rms_w_c_unzip = np.array([[item[i] for item in w_c] for i in range(len(w_c[0]))])
# rms_w_c = [np.sqrt(np.mean(item**2)) for item in rms_w_c_unzip]
if plot: ## just going to plot the first sample with the rms error for ease of code
       fig, ax = plt.figure(figsize=(10,7.5),facecolor='white'), plt.gca()
       plt.xlim(right=227)
       plt.ylim((-1,1))
       plt.errorbar(dd_bin[0],avg_w_a, yerr=rms_w_a, label=r'$w_a$',linestyle='-',linewidth=2,
       plt.errorbar(dr_bin[0],avg_w_b, yerr=rms_w_b, label=r'$w_b$',linestyle='-',linewidth=2,
       plt.errorbar(dr_bin[0],avg_w_c, yerr=rms_w_c, label=r'$w_c$',linestyle='-',linewidth=2,
       plt.legend();
       plt.xlabel('Distance');
       plt.title('Correlation Function for bin width \n of {binWidth} length units and {n} samp
       plt.grid();
              plt.savefig('correlation function n '+str(n)+' b '+str(binWidth)+'.png')
       plt.show()
```

Correlation Function for bin width of 2 length units and 30 samples



In [16]: print("--- %s seconds ---" % (time.time() - bigstart))

--- 235.52414774894714 seconds ---