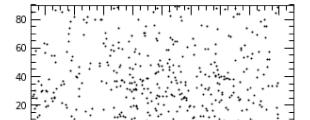
Problem Set 01

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);
```



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

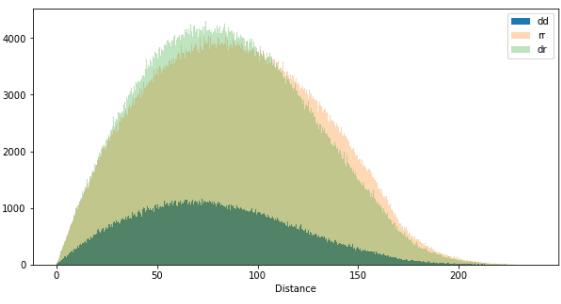
```
In [3]:
         ## Generate Random galaxy positions
         datSize = len(data)
         galMultiplier = 2
         def gen_galaxy_positions(N=datSize*galMultiplier,pkl=False,save=False,timeit=True):
             start_time = time.time() if timeit else None
             Generates N random galaxy 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
```

```
## 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);
```

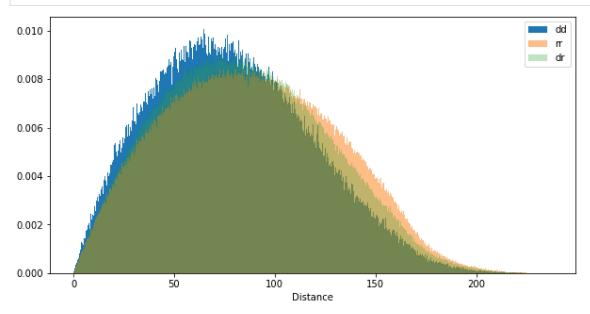
```
## calculate distances to other points
def get_dist(mode='dd',data1=None,data2=None,pkl=False,save=False, timeit=True):
    start_time = time.time() if timeit else None
    if pkl and os.path.isfile(pkl):
        dist = pd.read_pickle(pkl)
    return dist
```

```
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 d1
             if mode == 'dd' or mode == 'rr': ## only take upper triangle of matrix to prevent double coun
                 offset = 1 ## offset to remove self distances in dd and rr (i.e. distance between a point
                 slice idx = np.triu indices from(dist,k=offset) ## get indices for upper triangle of matr
                 #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 ---
In [7]:
         ## plot histograms of distances
         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 pairs
                                                                                          dd
                                                                                          IT
         4000
                                                                                         dr
```

elif data1 is not None and data2 is not None: ##case where I want to explicitly pass in data



```
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()
```



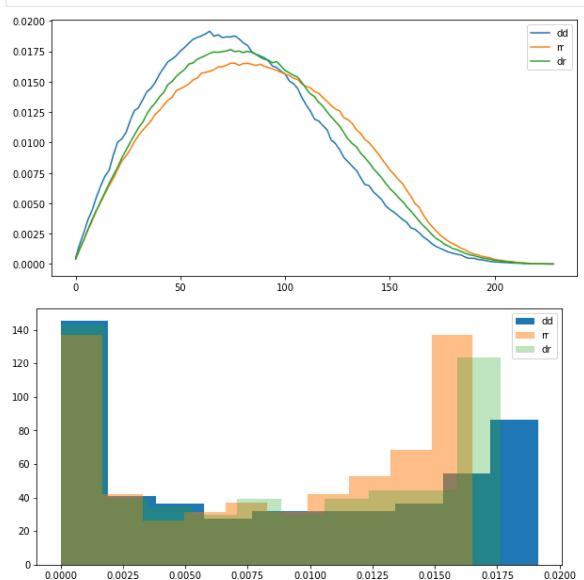
```
In [9]:
         ## develop correlation function
         maxBin = 230
         binWidth = 2
         binArray = np.linspace(0,maxBin,int(maxBin/binWidth)+1) ## the bins to use for the correlation fu
         #print(binArray)
         ## calculate bin averages
         def get_bin_averages(dataset,binWidth=binWidth,double=False,pkl=False,save=False, 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) ## efficien
             binVals = binVals/len(dataset) ## normalize by number of points put in
             # if double: ## accounts for the 2DR case ## 09/20 update: not needed
                   binAvgs = binAvgs*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 w
```

```
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 ---

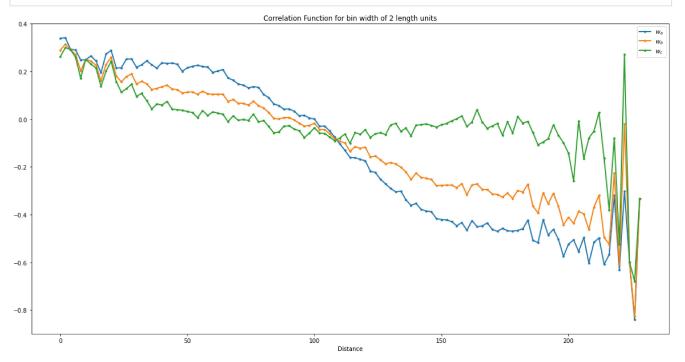
In [11]:
    ## plot bin and vals for dd, rr, and dr
    fig, ax = plt.figure(figsize=(10,5)), plt.gca()
    plt.plot(dd_bin,dd_val,label='dd');
```

```
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?
```

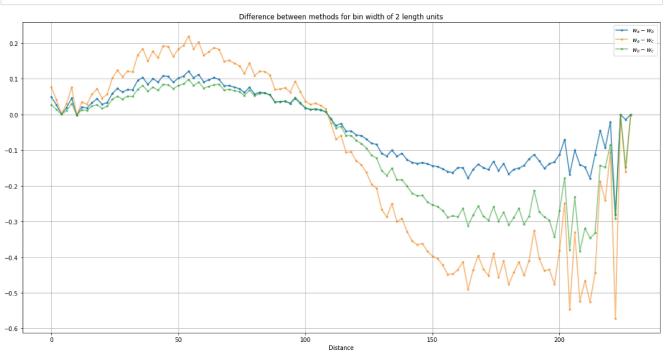


```
In [12]:
          ## calculate correlation function
          # def get_corr_func(dd,rr,dr,binWidth=binWidth,pkl=False,save=False, timeit=True):
                start_time = time.time() if timeit else None
          #
                 if pkl and os.path.isfile(pkl):
          #
                     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)] ## <math>w_c seems
          ## plot correlation function (1 instance)
```

```
fig, ax = plt.figure(figsize=(20,10)), plt.gca()
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));
```



```
## 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
```

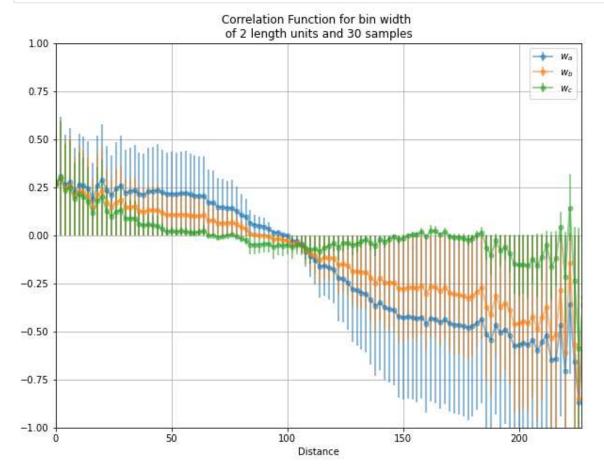


```
def get_rms_error(mode=None,n=10,plot=True,save=False,fun=False):
         start time = time.time()
        randGals = [gen_galaxy_positions(N=10000,timeit=False) for i in range(n)]
        #if mode == 'w_a':
        memDebug = False
        dd = [get_dist('dd',data,data, timeit=False) for i in range(n)] ## 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(dd
        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(dd_i,dr_i)]
        w_c = np.array([[(dd_j - 2*dr_j + rr_j)/(rr_j) for dd_j,dr_j,rr_j in zip(dd_i,dr_i,rr_i)] for
        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 = [np.sqrt(np.mean(item**2)) for item in rms_w_a_unzip]
        \# rms\_w\_b\_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, m
                 plt.errorbar(dr\_bin[0],avg\_w\_b, yerr=rms\_w\_b, label=r'$w\_b$', linestyle='-', linewidth=2, matching and the state of the 
                 plt.errorbar(dr\_bin[0], avg\_w\_c, \ yerr=rms\_w\_c, \ label=r'$w\_c$', linestyle='-', linewidth=2, \ mathematical actions and the second 
                 plt.legend();
                 plt.xlabel('Distance');
                 plt.title('Correlation Function for bin width \n of {binWidth} length units and {n} sampl
                 plt.grid();
                 if save:
                          plt.savefig('correlation_function_n_'+str(n)+'_b_'+str(binWidth)+'.png')
                 plt.show()
```

In [15]:

get_rms_error(n=30, save=True)

Not sure why rms is higher than expected since the expression for rms is correct



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

--- 235.52414774894714 seconds ---