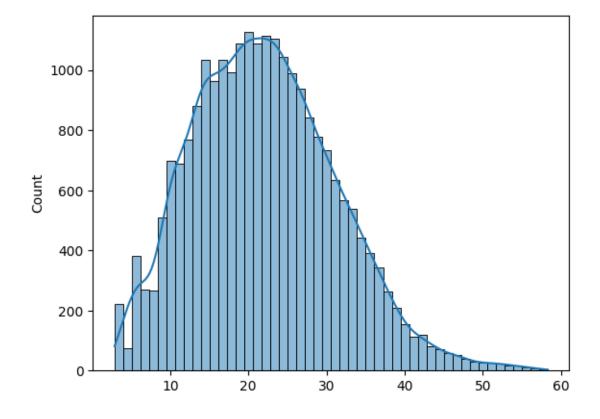
Notebook

April 17, 2023

```
[35]: import scipy as sp
  import numpy as np
  import seaborn as sns
  import matplotlib.pyplot as plt

[36]: data = sp.io.loadmat('coord1PHP.mat')
  protdist = data['protdist'][:,0]
  A = data['A']
  coord = data['coord']
  Nnode = A.shape[0]
  sns.histplot(protdist, kde=True, bins=50)
```

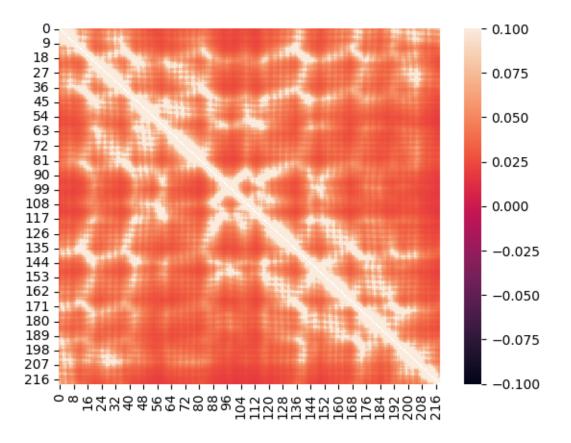
[36]: <Axes: ylabel='Count'>



```
[37]: D = sp.spatial.distance.squareform(protdist) sns.heatmap(1./D)
```

/tmp/ipykernel_450/2899854611.py:2: RuntimeWarning: divide by zero encountered
in divide
 sns.heatmap(1./D)

[37]: <Axes: >



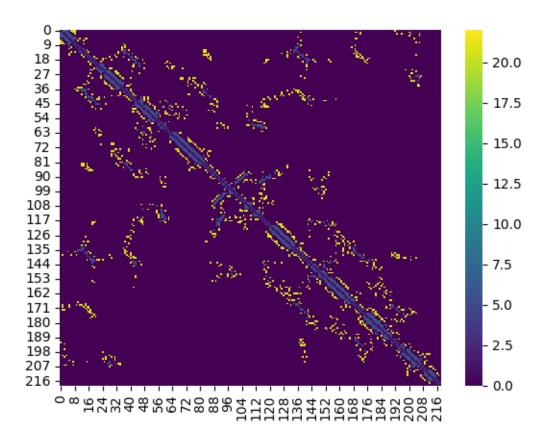
```
[38]: Net6 = (D < 6) * D

Net10 = (D > 10) * (D < 11) * D

N1 = (D < 11) * D

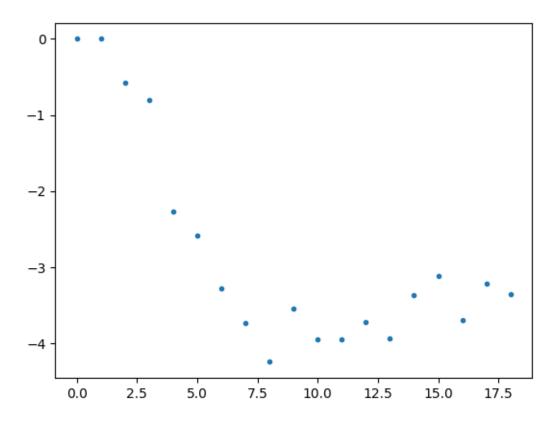
sns.heatmap(Net6 + 2*Net10, cmap='viridis')
```

[38]: <Axes: >



```
[39]: Kdiag = []
for index in range(Nnode):
    Kdiag.append(np.sum(np.diag(A, index))/(Nnode - index))
plt.plot(np.log(Kdiag[1:20]), '.')
```

[39]: [<matplotlib.lines.Line2D at 0x7f42d241c3d0>]

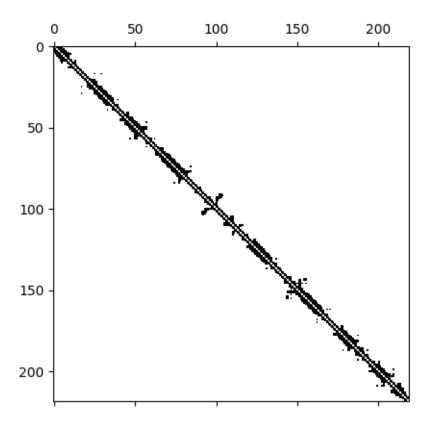


```
[51]: Nkd = 12
Ad = np.zeros((Nnode, Nnode), dtype=int)

for ind in range(-Nkd, Nkd + 1):
    Ad += np.diagflat(A.diagonal(ind), ind)

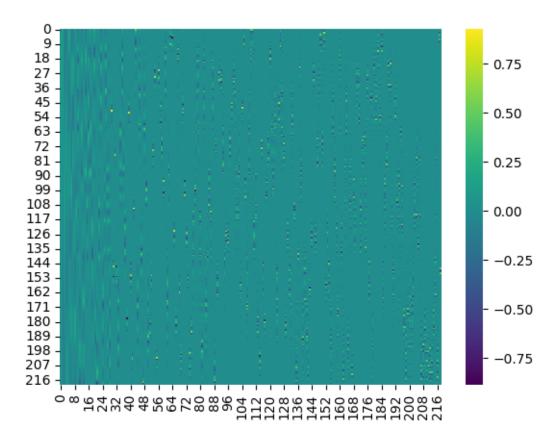
plt.spy(Ad)
```

[51]: <matplotlib.image.AxesImage at 0x7f42d0900670>



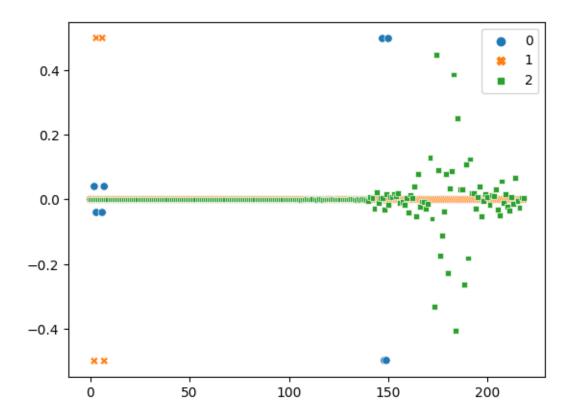
```
[52]: ValAd, VecAd = np.linalg.eig(sp.sparse.csgraph.laplacian(Ad))
sns.heatmap(VecAd, cmap='viridis')
```

[52]: <Axes: >



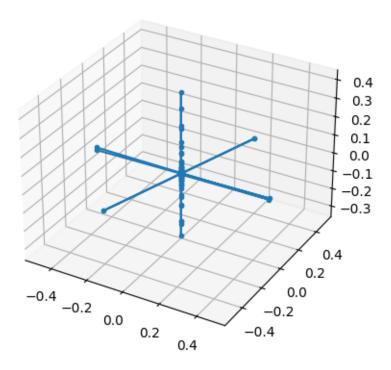
```
[50]: i1 = Nnode - 1
i2 = Nnode - 2
i3 = Nnode - 3
sns.scatterplot(VecAd[:, [i1, i2, i3]])
```

[50]: <Axes: >



```
[68]: fig = plt.figure()
ax = fig.add_subplot(projection='3d')
ax.plot3D(VecAd[:, i1], VecAd[:, i2], VecAd[:, i3], marker='.')
```

[68]: [<mpl_toolkits.mplot3d.art3d.Line3D at 0x7f42cfc2ed10>]



```
[58]: import networkx as nx

D, P = sp.sparse.csgraph.dijkstra(Ad, directed=False, return_predecessors=True)
Cl = 1 / np.mean(D)
K = np.sum(A, axis=0)
BC = nx.betweenness_centrality(nx.from_numpy_array(A))

print(Cl)
print(np.mean(K))
print(BC)
```

$\tt 0.057588259732525154$

10.0

- $\{0: 0.0, 1: 0.01204099367300543, 2: 0.01401334954164753, 3:$
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```

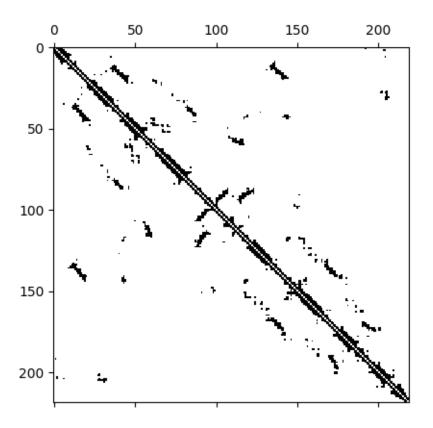
```
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217: 4.2277935145647484e-05, 218: 0.0}
```

1 Permutation effects

```
[20]: import scipy as sp
import numpy as np
from matplotlib import pyplot as plt

[29]: data = sp.io.loadmat('coord1PHP.mat')
A = data['A']
coord = data['coord']
Nnode = A.shape[0]
plt.spy(A)
```

[29]: <matplotlib.image.AxesImage at 0x7fc8e8b04790>



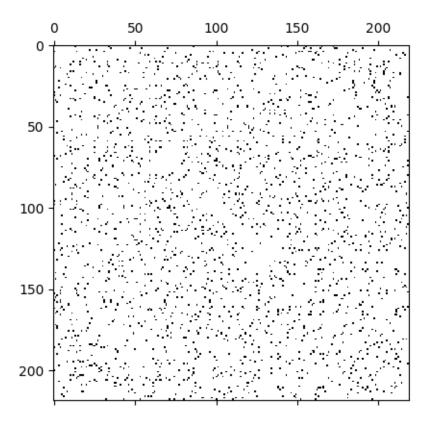
```
[22]: Index = np.arange(Nnode)
PermIndex = np.random.permutation(Nnode)
PermMat = np.zeros((Nnode, Nnode))
```

```
[23]: for ind in Index:
    PermMat[Index[ind], PermIndex[ind]] = 1
```

A faster method could be

```
[25]: A2 = PermMat.dot(A).dot(PermMat.T)
plt.spy(A2)
```

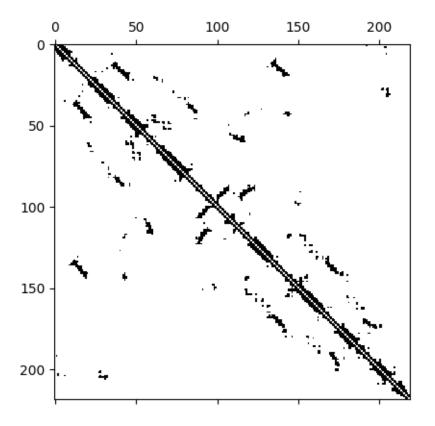
[25]: <matplotlib.image.AxesImage at 0x7fc8e8d9c190>



We can invert the permutation effect

```
[26]: A3 = (PermMat.T).dot(A2).dot(PermMat)
plt.spy(A3)
```

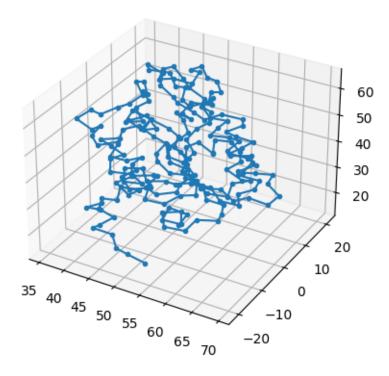
[26]: <matplotlib.image.AxesImage at 0x7fc8e8c082b0>



Let's try to visualize the 3d structure of the protein

```
[38]: fig = plt.figure()
ax = fig.add_subplot(projection='3d')
ax.plot(coord[:,0], coord[:,1], coord[:,2], marker='.')
```

[38]: [<mpl_toolkits.mplot3d.art3d.Line3D at 0x7fc8e19db5b0>]



2 Code from ER class

```
[57]: import scipy as sp
import numpy as np
import sys
import seaborn as sns
import matplotlib.pyplot as plt
import networkx as nx

def rand_canonical_f(Nnode, Plink):
    A = np.triu(np.random.rand(Nnode, Nnode), k=1)
    A = A > 1 - (2*Plink)
    A = A + A.T
    return A
```

Let's compare the memory usage between a sparse and a normal matrix. Moreover, we assume a microcanonical ensemble, fixing the number of nodes and links.

```
[58]: Nnode = int(1e3)
Nlink = int(3e4)
Net1 = sp.sparse.lil_matrix((Nnode,Nnode))
Net2 = np.zeros((Nnode,Nnode))
```

```
print(sys.getsizeof(Net1))
print(sys.getsizeof(Net2))
```

48 8000128

2.1 Microcanonical ensemble

Now, let's try to generate a random graph without using Networkx functions.

```
[59]: for index in range(100):
    # generate a random link
    ii = np.random.randint(Nnode, size=(Nlink, 2))
    # create a sparse matrix using these links
    NetM = sp.sparse.coo_matrix((np.ones(Nlink), (ii[:,0], ii[:,1])),
    shape=(Nnode, Nnode))
    #remove loops
    NetM.setdiag(0)
    # symmetrize
    NetM = NetM + NetM.T
print(NetM)
```

```
(0, 16)
               1.0
(0, 17)
               1.0
(0, 25)
               1.0
(0, 51)
               1.0
(0, 59)
               1.0
(0, 92)
               1.0
(0, 99)
               1.0
(0, 103)
               1.0
(0, 140)
               1.0
(0, 168)
               1.0
(0, 174)
               1.0
(0, 175)
               1.0
(0, 196)
               1.0
(0, 221)
               3.0
(0, 235)
               1.0
(0, 271)
               1.0
(0, 272)
               1.0
(0, 299)
               1.0
(0, 304)
               1.0
(0, 320)
               1.0
(0, 324)
               1.0
(0, 333)
               1.0
(0, 352)
               1.0
(0, 356)
               1.0
(0, 359)
               1.0
```

```
(999, 675)
               1.0
(999, 688)
               1.0
(999, 695)
               1.0
(999, 713)
               1.0
(999, 716)
               1.0
(999, 719)
               1.0
(999, 723)
               1.0
(999, 733)
               1.0
(999, 742)
               1.0
(999, 770)
               1.0
(999, 788)
               1.0
(999, 790)
               1.0
(999, 807)
               1.0
(999, 817)
               1.0
(999, 821)
               1.0
(999, 842)
               1.0
(999, 849)
               1.0
(999, 855)
               1.0
(999, 868)
               1.0
(999, 889)
               1.0
(999, 891)
               1.0
(999, 903)
               2.0
(999, 911)
               1.0
(999, 960)
               1.0
(999, 961)
               1.0
```

NOTE that reciprocal links and repeated links randomly created reduce symmetric links with a probability that scales as $\frac{1}{Nnode^2}$

Let's count the number of links in the matrix.

```
[60]: Nl = int(NetM.count_nonzero()/2)
print('Randomly generated links: {:d}'.format(Nl))
print('Missing links for microcanonical ensemble: {:d}'.format(Nlink-Nl))
```

Randomly generated links: 29035
Missing links for microcanonical ensemble: 965

We notice that we miss some links, then we add one by one missing links until microcanonical constraint fulfilled.

Lastly, let's compute the degree vector, finding minimum and maximum.

```
[62]: K = sum(NetM)
print('Average degree: {:.2f}'.format(np.mean(K)))
print('Minimum degree: {:d}'.format(int(np.min(K))))
print('Maximum degree: {:d}'.format(int(np.max(K))))
```

Average degree: 61.87 Minimum degree: 39 Maximum degree: 87

2.2 Canonical ensemble

Let's now try to build a canonical ensemble.

```
[63]: Nnode = int(1e2)
Plink = 0.3
Nexp = int(1e4)
```

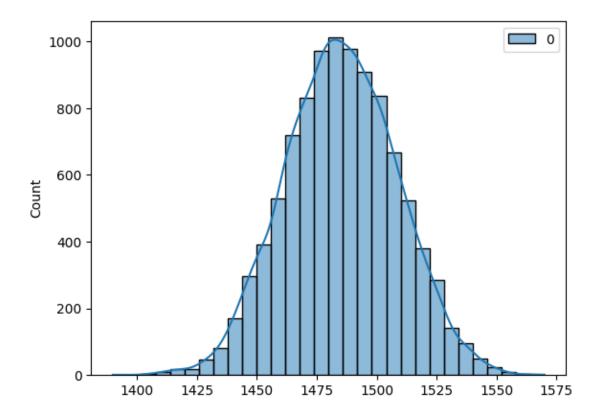
Let's see some stats

Average number of links: 1485.08

Standard deviation: 23.30

Expected number of links: 1485.00 Expected standard deviation: 32.24

[65]: <Axes: ylabel='Count'>



The degree distribution for increasing size - CV decreases. Tends to the delta function - regular graph.

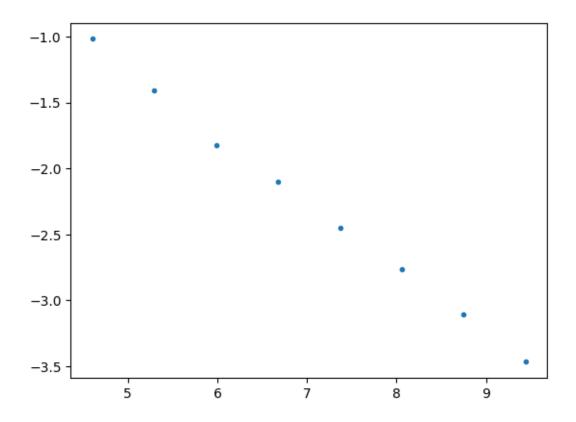
```
[66]: Pow = 4.21 # max network size
Pl = 0.1
Cnt = 0

M = []
S = []
CV = []
Nnodes = []
```

```
for index in np.arange(2, Pow, step=0.3):
    Nnodes.append(int(10**index))
    NetC = sp.sparse.triu(sp.sparse.rand(Nnodes[Cnt], Nnodes[Cnt], density=Pl,_u
    format='csr'), k=1)
    NetC = NetC + NetC.T
    K = sum(NetC).todense()
    M.append(np.mean(K))
    S.append(np.std(K))
    CV.append(np.std(K)/np.mean(K))
```

```
\# sns.histplot(data=K, bins=30, kde=True, label='N=\{:d\}'.
       → format(Nnodes[Cnt]))
          print('N={:d} - Average degree: {:.2f}'.format(Nnodes[Cnt], M[Cnt]))
          print('N={:d} - Standard deviation: {:.2f}'.format(Nnodes[Cnt], S[Cnt]))
          print('N={:d} - CV: {:.2f}'.format(Nnodes[Cnt], CV[Cnt]))
          Cnt += 1
     N=100 - Average degree: 4.50
     N=100 - Standard deviation: 1.62
     N=100 - CV: 0.36
     N=199 - Average degree: 9.76
     N=199 - Standard deviation: 2.39
     N=199 - CV: 0.25
     N=398 - Average degree: 19.72
     N=398 - Standard deviation: 3.19
     N=398 - CV: 0.16
     N=794 - Average degree: 39.65
     N=794 - Standard deviation: 4.83
     N=794 - CV: 0.12
     N=1584 - Average degree: 79.53
     N=1584 - Standard deviation: 6.86
     N=1584 - CV: 0.09
     N=3162 - Average degree: 158.26
     N=3162 - Standard deviation: 9.98
     N=3162 - CV: 0.06
     N=6309 - Average degree: 314.95
     N=6309 - Standard deviation: 14.07
     N=6309 - CV: 0.04
     N=12589 - Average degree: 629.21
     N=12589 - Standard deviation: 19.65
     N=12589 - CV: 0.03
[68]: plt.plot(np.log(Nnodes), np.log(CV), '.')
```

[68]: [<matplotlib.lines.Line2D at 0x7fafb89a6860>]



2.3 Giant component transition

```
[69]: Nnode = int(1e2)

rr = np.triu(np.random.rand(Nnode, Nnode), k=1)
rr = rr + rr.T
```

```
[70]: Pmin = 1 / (20 * Nnode)
Pmax = 6 / Nnode
xP = np.arange(Pmin, Pmax, step=0.00002)
```

NOTE that $\lambda = xP * Nnode$

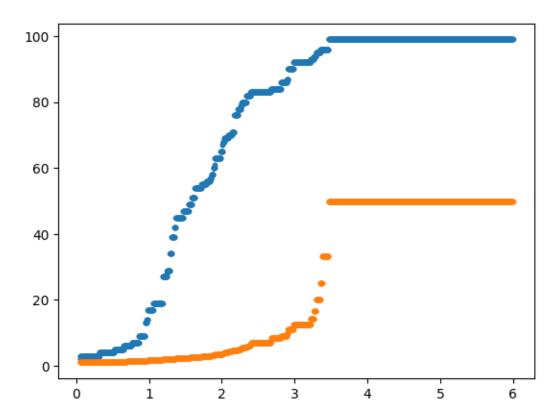
```
[71]: Rc = np.zeros(len(xP))
Rm = np.zeros(len(xP))
Cnt = 0

for index in xP:
    Net = nx.from_numpy_array(rr > 1 - index)
    CompList = nx.connected_components(Net)
    Comps = [len(x) for x in CompList]
    Rc[Cnt] = max(Comps)
```

```
Rm[Cnt] = np.mean(Comps)
Cnt += 1
```

```
[72]: plt.plot(xP*Nnode, Rc, '.')
plt.plot(xP*Nnode, Rm, '.')
```

[72]: [<matplotlib.lines.Line2D at 0x7fafb882d990>]



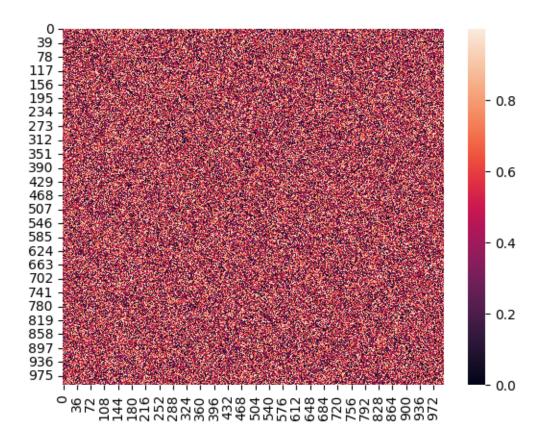
2.4 Semicircle law for ER network spectrum vs RMT (Random Matrix Theory)

```
[73]: Nnode = int(1e3)
Plink = 0.2 # link probability (canonical)

rr = np.triu(np.random.rand(Nnode, Nnode), k=1)
rr = rr + rr.T
np.fill_diagonal(rr, 0) # remove loops

# plot matrix with values
sns.heatmap(rr)
```

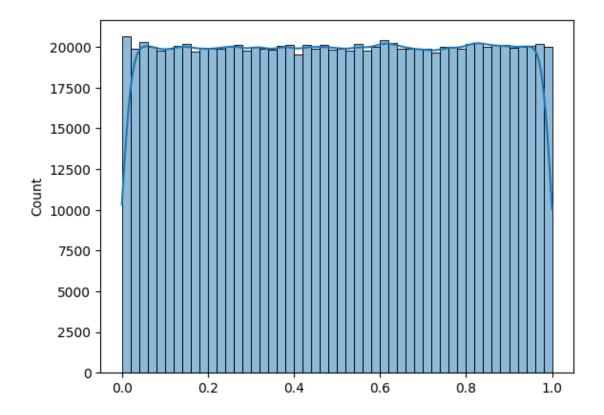
[73]: <Axes: >



Let's see also the histogram of coefficients

```
[74]: sns.histplot(data=rr.ravel(), bins=50, kde=True)
```

[74]: <Axes: ylabel='Count'>

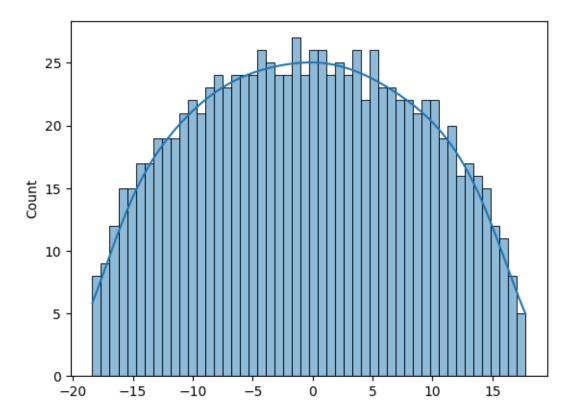


Now the eigenvalues and eigenvectors.

 ${\bf NOTE}$ A symmetric ER network has a real semicircle spectrum

```
[75]: vv = np.linalg.eigvals(rr)
sns.histplot(data=vv[vv < vv.max()], bins=50, kde=True)
```

[75]: <Axes: ylabel='Count'>

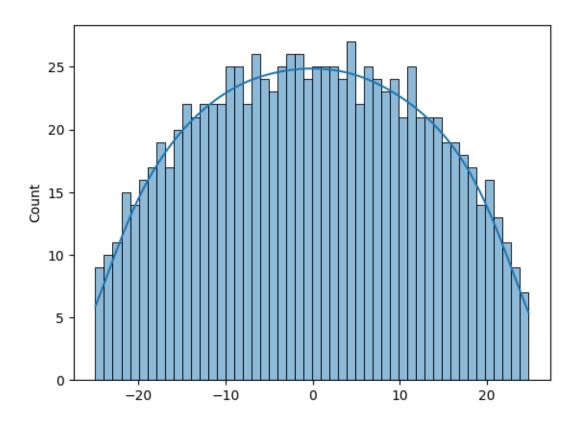


```
[76]: rr = np.triu(np.random.rand(Nnode, Nnode), k=1) > 1 - Plink
rr = rr + rr.T
```

```
[77]: vv, vec = np.linalg.eig(rr)
```

[78]: sns.histplot(data=vv[vv < vv.max()], bins=50, kde=True)

[78]: <Axes: ylabel='Count'>

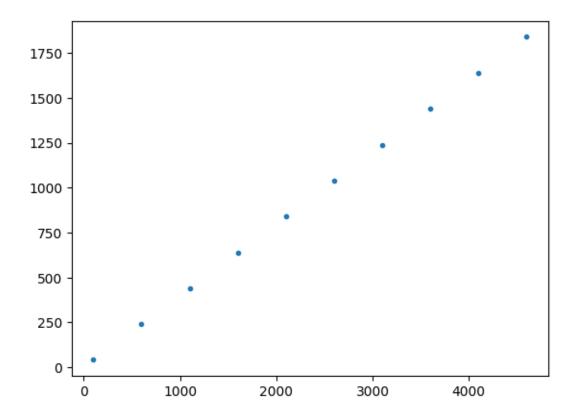


```
[83]: Cnt = 0
    xx = np.arange(100, 5000, step=500)
    vmax = []

for index in xx:
    rr = rand_canonical_f(index, Plink)
    vv = np.linalg.eigvals(rr)
    vmax.append(np.max(vv))
    Cnt += 1

plt.plot(xx, vmax, '.')
```

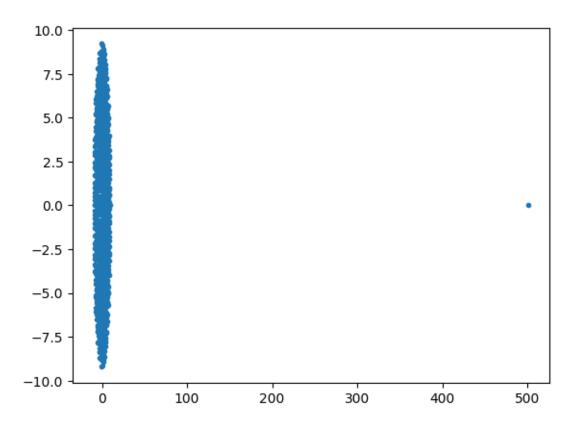
[83]: [<matplotlib.lines.Line2D at 0x7fafb8358f10>]



For a directed random matrix we'll have a complex circular spectrum

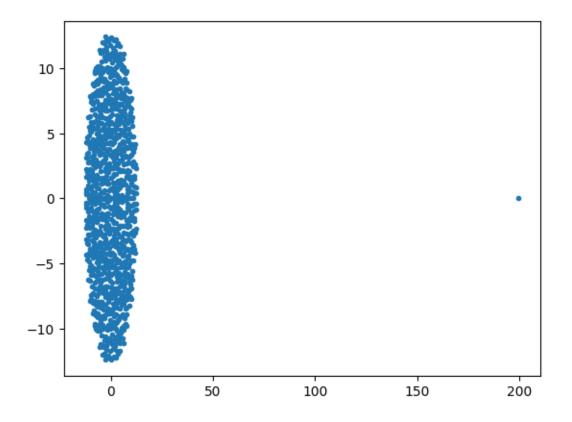
```
[80]: rr = np.random.rand(Nnode, Nnode)
vv = np.linalg.eigvals(rr)
vvr = np.real(vv)
vvi = np.imag(vv)
plt.plot(vvr, vvi, '.')
```

[80]: [<matplotlib.lines.Line2D at 0x7fafb83ea710>]



```
[81]: rr = np.random.rand(Nnode, Nnode) > 1 - Plink
    np.fill_diagonal(rr, 0)
    vv = np.linalg.eigvals(rr)
    vvr = np.real(vv)
    vvi = np.imag(vv)
    plt.plot(vvr, vvi, '.')
```

[81]: [<matplotlib.lines.Line2D at 0x7fafb84633d0>]

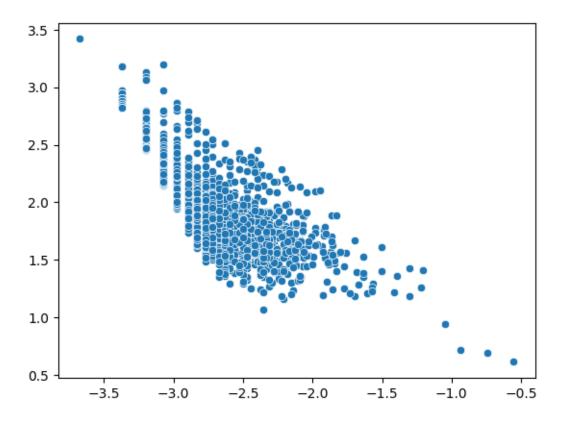


```
[9]: import scipy as sp
     import numpy as np
     import networkx as nx
     from tqdm import tqdm # necessary because shuffling takes a while
     def shuffle_net_sym_f(Net, Nsh):
         # Nsh = number of shuffles (* number of links)
         Shuf = Net
         I = list(nx.edges(Shuf))
         Nl = len(I)
         for ind in tqdm(range(Nl * Nsh)):
             L = np.random.randint(N1, size = 2)
             while L[0] == L[1]:
                  L[1] = np.random.randint(N1)
             if Shuf.has_edge(I[L[0]][0], I[L[0]][1]) and Shuf.has_edge(I[L[1]][0], __
      \BoxI[L[1]][1]) and (Shuf.has_edge(I[L[0]][0], I[L[1]][1]) == False) and (Shuf.
      \neghas_edge(I[L[1]][0], I[L[0]][1]) == False) and (I[L[1]][0] != I[L[0]][1])_u
      \hookrightarrow and (I[L[0]][0] != I[L[1]][1]) and (I[L[0]][0] != I[L[0]][1]) and
      \hookrightarrow (I[L[1]][0] != I[L[1]][1]):
                  # delete links
                  Shuf.remove_edge(I[L[0]][0], I[L[0]][1])
```

```
Shuf.remove_edge(I[L[1]][0], I[L[1]][1])
                  if Shuf.has_edge(I[L[0]][1], I[L[0]][0]):
                      Shuf.remove_edge(I[L[0]][1], I[L[0]][0])
                  if Shuf.has_edge(I[L[1]][1], I[L[1]][0]):
                      Shuf.remove_edge(I[L[1]][1], I[L[1]][0])
                  # shuffle links
                  Shuf.add_edge(I[L[0]][0], I[L[1]][1])
                  Shuf.add_edge(I[L[1]][1], I[L[0]][0])
                  Shuf.add_edge(I[L[1]][0], I[L[0]][1])
                  Shuf.add_edge(I[L[0]][1], I[L[1]][0])
                  # remap links
                  I = list(nx.edges(Shuf))
                  N1 = len(I)
          return Shuf
[10]: data = sp.io.loadmat('Yeast_DIP.mat')
      YSTnet = data['YSTnet']
     Let's process the network, which is not symmetrized with loops
[11]: Ndir = np.sum(YSTnet - YSTnet.T)
      Nloop = YSTnet.trace()
      print(Ndir)
      print(Nloop)
      YSTnet.setdiag(0)
      YSTnetSym = YSTnet + YSTnet.T
      GraphYST = nx.from_scipy_sparse_array(YSTnetSym)
     0.0
     4713.0
[12]: Kyst = nx.degree centrality(GraphYST)
      Kyst = np.array(list(Kyst.values()))
      Knn = np.divide(YSTnetSym * Kyst, Kyst)
[13]: import seaborn as sns
```

[13]: <Axes: >

sns.scatterplot(x=np.log10(Kyst), y=np.log10(Knn))



Let's shuffle links - microcanonical ensemble. **NOTE**: microcanonical shuffle does not work with few iterations.

[15]: <Axes: >

