RANDOM GRAPH VS PROTEIN GRAPH WITH SAME DEGREE DISTRIBUTION PDB USED = 1H02.PDB

nodes = 158 average degree = 7.72151898734 avg clustering coeff = 0.544386450399 edges = 610 average shortest path = 4.42562283319

degree dist for protein graph =

DEGREE

```
2
            0.0
3
            0.0126582278481
4
            0.0759493670886
5
            0.0506329113924
6
            0.177215189873
7
            0.177215189873
8
            0.151898734177
9
            0.132911392405
            0.0886075949367
10
            0.0886075949367
11
12
            0.0316455696203
13
            0.00632911392405
14
            0.00632911392405
15
            0.0
clustering coeff dist for protein graph =
```

DEGREE DISTR

DEGREE CLUSTERING COEFF DISTR

```
2
            0
3
            1.0
            0.547619047619
4
5
            0.204545454545
6
            0.34666666667
7
            0.192918192918
8
            0.136204481793
9
            0.0853658536585
10
            0.0446066504461
            0.0406983744732
11
12
            0.0130147630148
            0.00269475747183
13
            0.00229517318125
14
15
            0.0
```

degree dist for random graph =

DEGREE DISTR

2 0.0

3 0.0126582278481

```
0.0759493670886
4
5
            0.0506329113924
6
            0.177215189873
7
            0.177215189873
8
            0.151898734177
9
            0.132911392405
10
            0.0886075949367
            0.0886075949367
11
            0.0316455696203
12
            0.00632911392405
13
            0.00632911392405
14
15
            0.0
clustering coeff dist for random graph =
DEGREE
                  CLUSTERING COEFF DISTR
2
            0
3
            0.0
4
            0.0595238095238
5
            0.00454545454545
6
            0.048
7
            0.030525030525
8
            0.0234593837535
9
            0.0178410117435
```

0.01200324412 0.0123419626731

0.0

0.00223387723388

0.000898252490609 0.000765057727083

10

11 12

13

14

15