S(L3)

(1. Ch, (1-q). q  $Z = 10^{6} = r = upper$   $Z = 10^{6} + 1 = r = lower bound$ (=0

(in)

C. Willhouse a non-zero asymptote so it is heavy toiled.

i.e. the ratio is non-zero far for b/c the ratio at lags is non-zero. an one end of the scale, it is a very steep core as well. This means heavy tail.

The difference is this ea. is probot a specific word of rooker being chosen, instead of any of length C.

3. a. E[fd] f= c.d~ 1 2 d · Cd = 1 C 2 d

- 1 C 2 d · Cd = - 1 C 2 d

- 1 C 2 d · Cd = - 1 C 2 d 2 - (· (c) = P(C2-1) II) P(DZCA) = E[P]  $| -\rho(0) = ca)^{2} | \frac{ca}{ca}$   $| -\rho(0) = ca)^{2} | \frac{ca}{ca} |$ 1-P[DZCA)Z1-<u>E[D]</u>  $\frac{\partial}{\partial t} \left( \frac{\partial}{\partial t} \left( \frac{\partial$ Soi Lécap(l) gar

1 2 Cd (1-(1-PCL))) m= O(m) arverat O(1) ecn) eci) V) by ii) and iv), we have 2 0 C ( ) 1 /

Cisardu af n.

Sa ve hone

2 (n - t - E) as desired

Plugging into mathematica, muit by l for Ev, we get

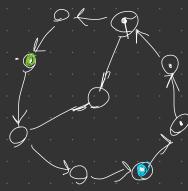
1 = K

b. Expected Avg Shartest path length!

$$\frac{-3+3p+2np-np^2+(1-p)^2(3+(-\lambda+n)p)}{(-1+n)p^2}$$

Scales exponentially w/n.

for no central node, 
$$\frac{1}{N} \cdot \frac{N^{-1}}{N} = \frac{(N-1)(N-1)}{N-1} = \frac{N^{-2}}{N}$$



works in all but this case, in which case it will add 1 to the Shartest path

The algo novid Nark in all but 1 class of cases if B always routes to the clasest node, We always want to toke B, since in most cases, we skip at least I hap af length 1 it we take B. Thus, given a path through B ar not through B, we would want to take B in most cases, The case we wouldn't want to is the one we took, as above going from Blue to gren. In cases like this, the algo takes B, but cames back where it entered and moves on, rather than skip 13. In this case, we add I to tree shartest path, which is pretty close to the real shortest path. An "accidental" traversal at B would only happen are, since if it happened twice, the first traversal would have come out at the Second traversal's paint, a contradiction. Thus, This algo admit) the Shortest path in all but I case, when we have to take B, but the edge clasest to the destination is the one we took, adding I to the resulting path, Which is very close to the real Shortest path.

G. (a. 
$$milt+1)-milt$$
)

 $7 = t+1$ 
 $indiv anty, inc = 1$ 
 $milt=0$ 
 $2t-1$ 

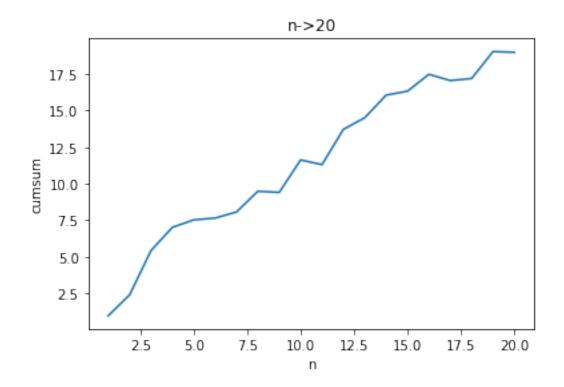
$$m_i(t+i) = m_i(t) + m_i(t)$$

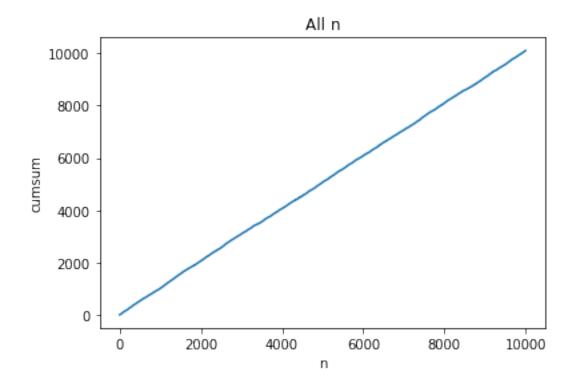
B. Using my approximate contapprox, we conside how The eqn follows from newton's method w/ h=1.

### Problem2

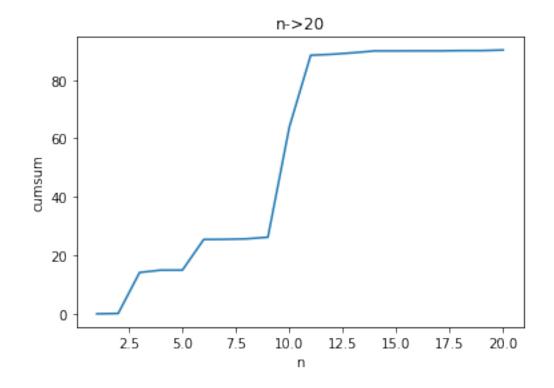
#### February 1, 2023

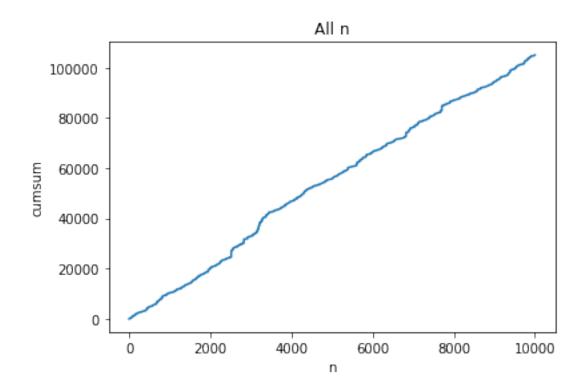
```
In [307]: import numpy as np
          import matplotlib.pyplot as plt
   Part A
1
In [308]: N = 10000
          norm_dist = np.random.normal(1, 1, (N,))
          weib = np.random.weibull(0.3, (N,))
          pareto = (np.random.pareto(0.5, (N,)) + 1) * (1/3)
In [309]: def make_plots_A(dist):
              plt.plot(np.arange(1, 21, 1), np.cumsum(dist)[:20])
              plt.title("n->20")
              plt.xlabel("n")
              plt.ylabel("cumsum")
              plt.show()
              plt.plot(np.arange(N), np.cumsum(dist))
              plt.title("All n")
              plt.xlabel("n")
              plt.ylabel("cumsum")
              plt.show()
In [310]: make_plots_A(norm_dist)
```



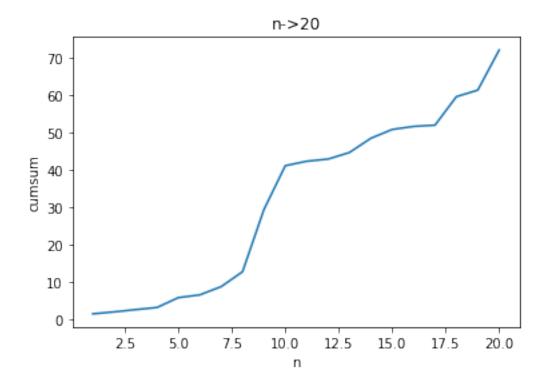


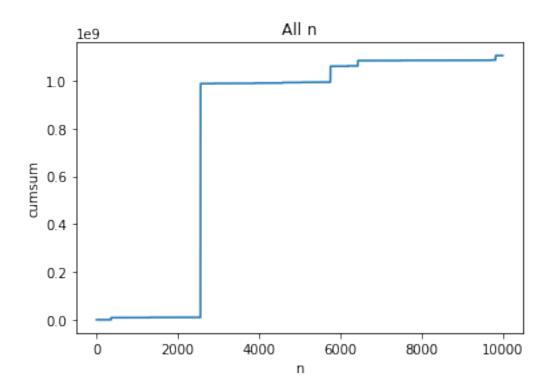
In [311]: make\_plots\_A(weib)



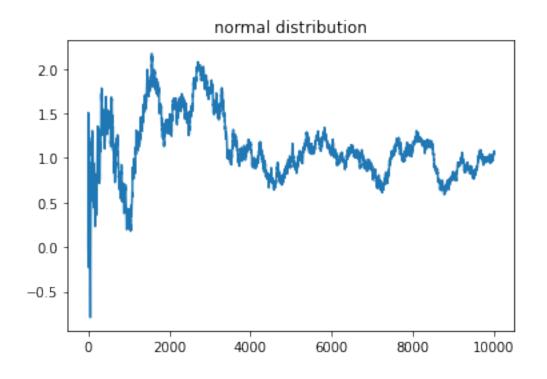


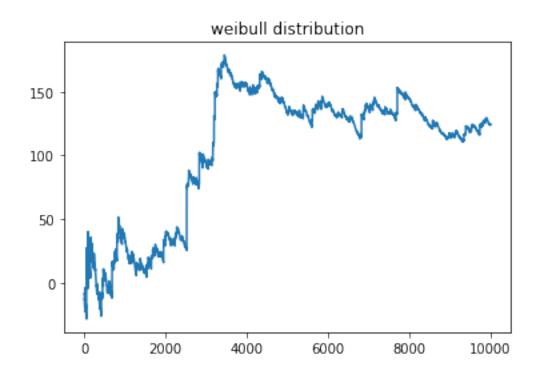
In [312]: make\_plots\_A(pareto)





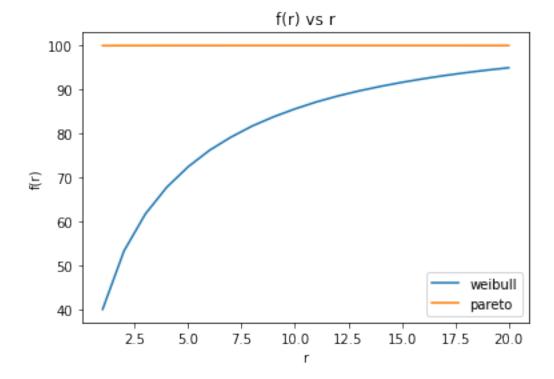
#### 2 Part B





#### 3 Part C

Out[317]: <matplotlib.legend.Legend at 0x7f3836ceb670>



#### 4 Part D

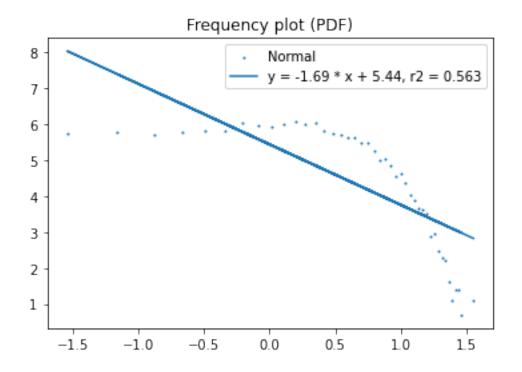
```
In [318]: from statsmodels.distributions.empirical_distribution import ECDF
          from sklearn.linear_model import LinearRegression
          from sklearn.metrics import r2_score
In [319]: # Note: Feel free to modify this template as you wish or build your own from scratch
          # This is a tricky problem to implement so we hope we are helping by providing a tem
          def pdf(data, dx=1):
              '''Takes an array with random samples from a distribution,
              and creates an approximate PDF of points, to use when frequency
              Returns a tuple of two vectors x, y where
              y_i = P(x_i - dx/2 \le data \le x_i + dx/2)'''
              low, high = np.min(data), np.max(data)
              x, y = [], []
              for d in data:
                  i = int((d + dx / 2 - low) / dx)
                  xi = low + i * dx
                  if xi in x:
                      y[x.index(xi)] += 1
                  else:
                      x.append(xi)
                      y.append(1)
              return np.array(x), np.array(y)
          def ccdf(data):
              '''Takes an array with random samples from a
              distribution, and creates an approximate CCDF
              (complementary CDF) of points. Returns a tuple of
              two vectors x, y where y_i = P(data > x_i)'''
              # TODO: Complete this function
              # Use this when creating the rank plot
              # HINT: To generate an approximate CDF (not CCDF), one would sort the random sam
              \# return those as the x values, and then create a range from 1 to n
              # (scaled by 1/n) for the y values. How does this change when creating a CCDF?
              x = 1-np.sort(data)
              y = [i / N \text{ for } i \text{ in } range(1, N+1, 1)]
              return x, y
          def keep_positive(data):
              '''Takes an array with random samples from
              a distribution, and filters our negative and
              zero entries (in both x and y) in data'''
```

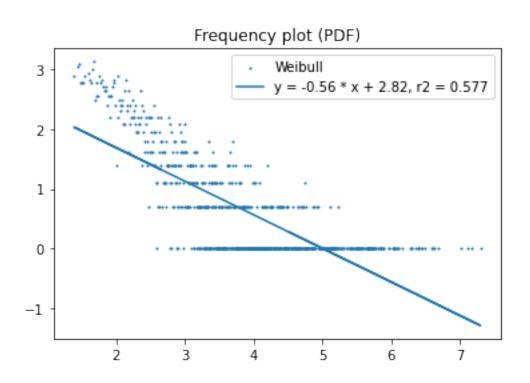
```
# TODO: Complete this function
    for dist in range(len(data)):
        x, y = data[dist]
        x = np.array(x)
        y = np.array(y)
        x_pos = x > 0
        y_pos = y > 0
        x = x[x_pos \& y_pos]
        y = y[y pos \& x pos]
        data[dist] = (x, y)
    return data
def non_outliers(x, m):
    '''Takes an array x of data and an integer m,
    and returns a list z of boolean values such
    that z_i indicates whether the mean-centered
    value x_i is within m std devs of the values in x'''
    # TODO: Complete this function
    x_{mean} = np.mean(x)
    x std = np.std(x)
    x centered = x - x mean
    z = np.abs(x centered) <= m * x std</pre>
    return z
def reject_outliers(data, m):
    '''Takes an array of data (data here is a
    tuple (x,y), where x and y are arrays:
    in the form resulting from pdf() or ccdf(), for example)
    and an integer m, and removes outliers from the data'''
    # TODO: Complete this function
    # Note that since we are plotting on log-log scale
    # Removing outliers should happen AFTER we move to log-log scale
    # HINT: This function returns the (x, y) tuples in data that
    # are not outliers (non_outliers() is helpful here)
    # HINT: You must set m to a default value that defines how many
    # standard deviations away an outlier is. We are testing that
    # these distributions are heavy-tailed, so too tight of a range
    # won't be helpful.
    x, y = data
    m = 2
    x_non = non_outliers(x, m)
    y_non = non_outliers(y, m)
    x = x[x_non \& y_non]
    y = y[x_non & y_non]
    return (x, y)
```

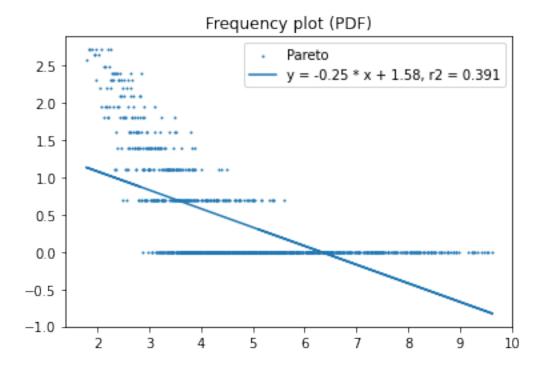
```
'''Takes (one-dimensional) training data X
              and labels y and fits the simple linear regression
              model y = mX + b, and computes the coefficient
              of determination (r^2). Returns a tuple (m, b, r^2)'''
              # TODO: Complete this function
              reg = LinearRegression()
              reg.fit(X.reshape(-1, 1), y)
              ypred = reg.predict(X.reshape(-1, 1))
              return (reg.coef_, reg.intercept_, r2_score(y, ypred))
          def make_graphs_d(data, title, labels, ylabel='', xlabel=''):
              Build respective PDF/CCDF for all three distributions
              Attempts to create a best-fit line
              And plots this line alongside the points
              data: List of pdf() outputs from all 3 distributions
              title: Either
                  Frequency plot (PDF)
                  Rank plot (CCDF)
              labels: List of distribution names
                  i.e. ["Normal, Weibull, Pareto"]
              # General plotting function for points, as well as best-fit line
              for (X, y), label in zip(data, labels):
                  m, b, r2 = linear_regression(X, y)
                  plt.scatter(X, y, label=label, s=1)
                  plt.plot(X, b + m * X, label='y = %.2f * x + %.2f, r2 = %.3f' % (m, b, r2))
                  plt.title(title)
                  plt.xlabel(xlabel)
                  plt.ylabel(ylabel)
                  plt.legend()
                  plt.show()
In [320]: # Concatenate data into one list to reduce code repetition
          Xi = [norm_dist, weib, pareto]
          names = ["Normal", "Weibull", "Pareto"]
          # Build the PDF and turn it into a log-log scale,
          # first removing all negative values and corresponding indices
          # TODO: When (if at all) do we remove outliers here?
          # Before or after we convert to log-scale?
          data = [pdf(Xi[i], dx=0.1) for i in range(3)]
          data = keep_positive(data)
          data = [(np.log(X), np.log(y)) for (X, y) in data]
          data = [reject_outliers(data[i], 4) for i in range(len(data))]
```

def linear\_regression(X, y):

# # Plot the frequencies make\_graphs\_d(data, 'Frequency plot (PDF)', names)

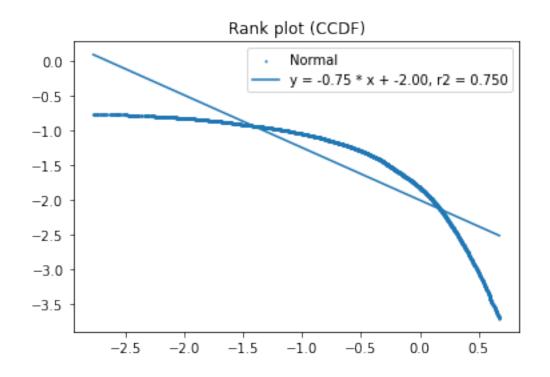


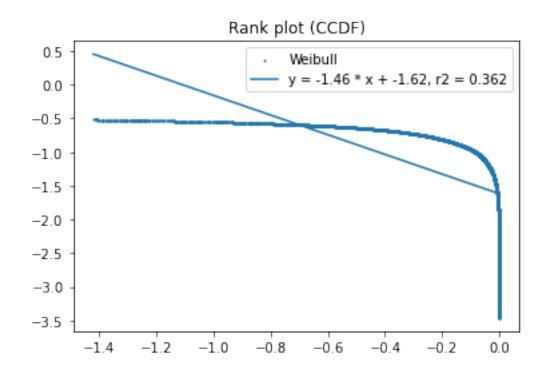


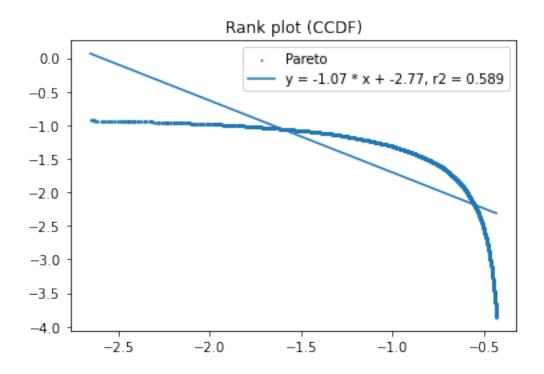


```
In [321]: data = [ccdf(Xi[i]) for i in range(3)]

# Build the CCDF and turn it into a log-log scale,
# first removing all negative values and corresponding indices
# TODO: Do we remove outliers here?
data = keep_positive(data)
data = [(np.log(X), np.log(y)) for (X, y) in data]
data = [reject_outliers(data[i], 4) for i in range(len(data))]
# Plot the ranks
make_graphs_d(data, 'Rank plot (CCDF)', names)
```







For the plots,

Filtering out outliers makes sense because on a log scale, big outliers can mess with the graph ranges. Small outliers become big outliers in log, while big outliers become even bigger.

We see that the normal distribution is not heavy-tailed, but the other 2 are. It's easier to see this because best fits have better linearity for normal, but they are worse for weibull and pareto, and we see a sharp drop off in the points in the emperical ccdf.

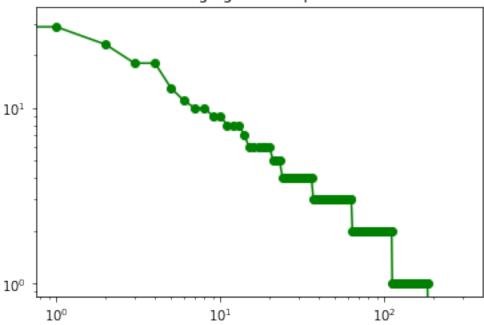
#### Problem4

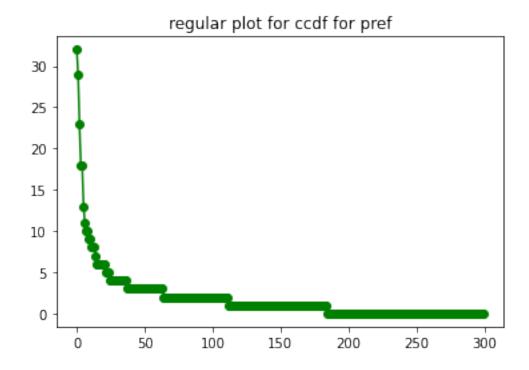
#### February 1, 2023

```
In [57]: import numpy as np
         import networkx as nx
         import matplotlib.pyplot as plt
In [66]: def get_deg_sum(G):
             nodes = list(G.nodes())
             tot = 0
             for node in nodes:
                 tot += G.degree(node)
             return tot
         # A
         def gen_pref(T):
             G = nx.Graph()
             G.add_edge(0, 1)
             for newcomer in range(2, T, 1):
                 G.add_node(newcomer)
                 deg_sum = get_deg_sum(G)
                 for existing in range(newcomer):
                     p = G.degree(existing) / deg_sum
                     if np.random.random() < p:</pre>
                         G.add_edge(newcomer, existing)
             degrees = [d for n, d in G.degree()]
             return G, degrees
         # B
         def configuration_graph(degrees):
             v = []
             G = nx.Graph()
             for i, k in enumerate(degrees):
                 v.extend([i for x in range(k)])
             v = np.random.permutation(v)
             for i in range(0, len(v), 2):
                 G.add_edge(v[i], v[i+1])
             return G
```

#### 1 A

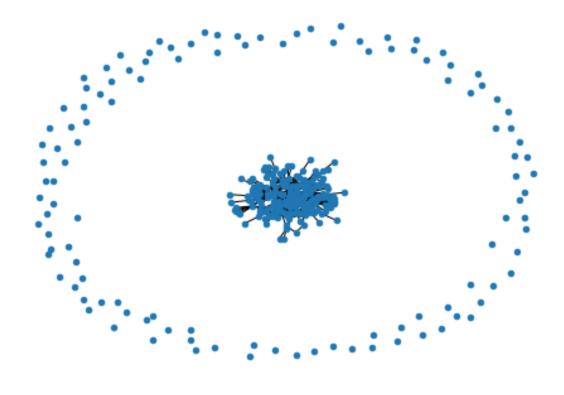
## loglog ccdf for pref



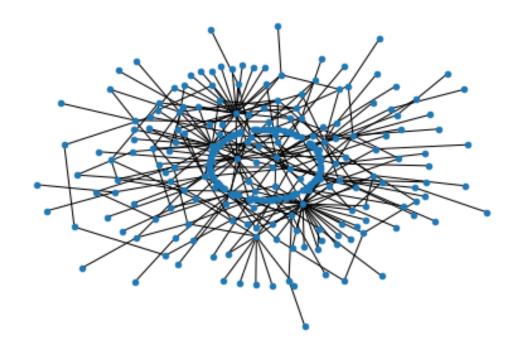


We definitely see a heavy-tail here given these plots.

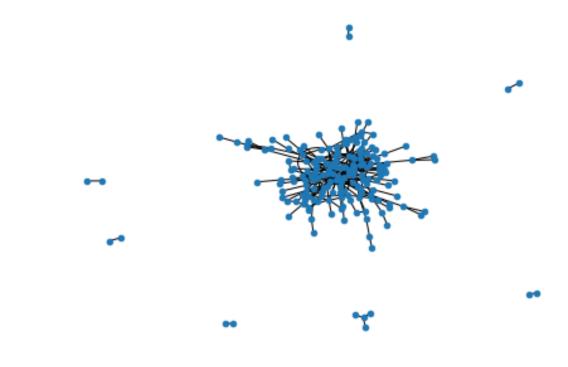
#### 2 B



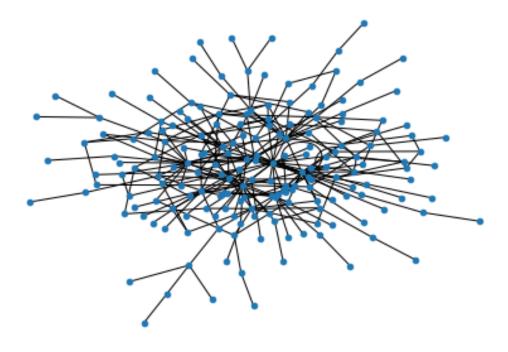
In [79]: nx.draw\_kamada\_kawai(G\_pref, node\_size=20)



In [81]: nx.draw(G\_config, node\_size=20)



In [80]: nx.draw\_kamada\_kawai(G\_config, node\_size=20)



#### 3 C

The differences I see are that the pref generated graph has a more uniform structure of disconnected components, while the config graph has smaller, disconnected clusters all around a main cluster in the middle. The pref graph has a bit more structure to it as well, forming a "social circle" of sorts with the highly connected nodes.