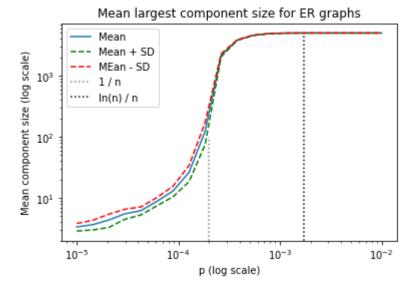
```
import networkx as nx
import numpy as np
import matplotlib.pyplot as plt
from collections import Counter
```

1. ER phase transitions

a) ER graph component sizes

```
In [ ]:
         def analyze er graph components(n, p range, graphs per p=20):
             means, sds = [], []
             for p in p range:
                 print('Analyzing ER graphs for p={}'.format(p))
                 comp sizes = []
                 for i in range(graphs per p):
                     G = nx.generators.fast gnp random graph(n, p, seed=i)
                     comp sizes.append(max([len(comp) for comp in nx.connected components(G)]))
                 means.append(np.mean(comp sizes))
                 sds.append(np.std(comp sizes))
             return np.array(means), np.array(sds)
In [ ]:
         n = 5000
         p range = np.logspace(-5, -2, num=20)
         means, sds = analyze er graph components(n, p range)
        Analyzing ER graphs for p=1e-05
        Analyzing ER graphs for p=1.438449888287663e-05
        Analyzing ER graphs for p=2.06913808111479e-05
        Analyzing ER graphs for p=2.9763514416313192e-05
        Analyzing ER graphs for p=4.281332398719396e-05
        Analyzing ER graphs for p=6.158482110660267e-05
        Analyzing ER graphs for p=8.858667904100833e-05
        Analyzing ER graphs for p=0.00012742749857031334
        Analyzing ER graphs for p=0.00018329807108324357
        Analyzing ER graphs for p=0.00026366508987303583
        Analyzing ER graphs for p=0.000379269019073225
        Analyzing ER graphs for p=0.0005455594781168515
```

```
Analyzing ER graphs for p=0.0007847599703514606
        Analyzing ER graphs for p=0.0011288378916846883
        Analyzing ER graphs for p=0.001623776739188721
        Analyzing ER graphs for p=0.002335721469090121
        Analyzing ER graphs for p=0.003359818286283781
        Analyzing ER graphs for p=0.004832930238571752
        Analyzing ER graphs for p=0.0069519279617756054
        Analyzing ER graphs for p=0.01
In [ ]:
         def plot with sds(x, y, y sds, n=5 000):
             plt.plot(x, y)
             plt.plot(x, y - y sds, c='green', linestyle='dashed')
             plt.plot(x, y + y_sds, c='red', linestyle='dashed')
             # plot vertical lines at 1 / n and ln(n) / n
             plt.vlines(1 / n, 0, 100, linestyle='dotted', colors='gray')
             plt.vlines(np.log(n) / n, 0, 4900, linestyle='dotted', colors='black')
             plt.yscale('log')
             plt.xscale('log')
             plt.title('Mean largest component size for ER graphs')
             plt.xlabel('p (log scale)')
             plt.ylabel('Mean component size (log scale)')
             plt.legend(['Mean', 'Mean + SD', 'MEan - SD', '1 / n', 'ln(n) / n'])
In [ ]:
         plot with sds(p range, means, sds)
```



b) Interesting values of p

From class, we have that when p = lambda / n, lambda > 1 makes the largest component size converge to n, while lambda < 1 makes it converge to ln(n). Here, lambda = 1 represents a point at which the largest component size is transitioning rapidly, as it has the graph's peak slope. Also, l notice that as lambda shrinks less than 1, the size remains low (near $ln(n) \sim = 8$), while as lambda grows above 1, the size approaches n.

Also, we have that when p = lambda * ln(n) / n, lambda < 1 makes the graph's probability of being connected converge to 0, while lambda > 1 makes it converge to 1. Here, to the left of ln(n) / n (i.e. lambda > 1), the mean component size has not yet converged to n, but to the right, the mean is equal to n and the standard deviation is 0, meaning that, as expected, all sampled graphs converge to being connected.

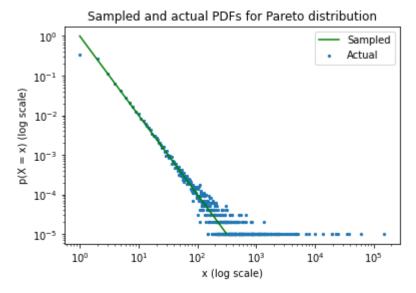
2. Fitting power-law exponents

b) Pareto sampling

```
def sample_pareto(alpha=1, x_0=1, size=100_000):
    pareto_func = np.vectorize(lambda u: x_0 / u**(1 / alpha))
    u_samples = np.random.sample(size=size)
    return pareto_func(u_samples)

def compute_pdf(samples):
```

```
samples_ints = np.round(samples, decimals=0)
             samples int counts = Counter(samples ints)
             x = np.array(sorted(samples int counts.keys()))
             px x = np.array([samples int counts[k] for k in x])
             px x = px x / px x.sum()
             return x, px x
         def compute ccdf(x, px x):
             s = 0
             fbarx x = np.zeros(px x.shape)
             for i in range(len(x) - 1, -1, -1):
                 s += px x[i]
                 fbarx x[i] = s
             return x, fbarx x
         def plot observations(x, y, alpha=1):
             plt.scatter(x, y, s=5)
             plt.xscale('log')
             plt.yscale('log')
             y_expected = 1 / x ** (alpha + 1)
             plt.plot(x[y expected >= 1e-5], y expected[y expected >= 1e-5], c='green')
             plt.title('Sampled and actual PDFs for Pareto distribution')
             plt.legend(['Sampled', 'Actual'])
             plt.xlabel('x (log scale)')
             plt.ylabel('p(X = x) (log scale)')
In [ ]:
         pareto = sample pareto()
         x, px x = compute pdf(pareto)
In [ ]:
         x, fbarx x = compute ccdf(x, px x)
In [ ]:
         plot observations(x, px x)
```



d) Estimating alpha

```
In [ ]:
         def estimate alpha pdf ls(samples):
             x, p_x = compute_pdf(samples)
             x_{\log} = np.\log(x.reshape((x.shape[0], 1)))
             y_{\log} = np.\log(p_x)
             alpha_pred = -np.linalg.lstsq(x_log, y_log, rcond=None)[0] - 1
             return alpha pred
         def estimate alpha ccdf ls(samples):
             x, px_x = compute_pdf(samples)
             x, fbarx x = compute ccdf(x, px x)
             x_log = np.log(x.reshape((x.shape[0], 1)))
             y_log = np.log(fbarx_x)
             # integrating the PDF gives line with slope -alpha
             alpha_pred = -np.linalg.lstsq(x_log, y_log, rcond=None)[0]
             return alpha pred
         def estimate alpha mle(samples, x 0=1):
             n = len(samples)
```

```
# from c), MLE is alpha = n / (sum i(log(x i)) - (n * log(x 0)))
             return n / (np.log(samples).sum() - (n * np.log(x 0)))
         def eval alpha estimate(estimate alpha, name, samples=100):
             preds = []
             for _ in range(samples):
                 samples = sample pareto()
                 preds.append(estimate alpha(samples))
             m, sd = np.mean(preds), np.std(preds)
             print('{}:'.format(name))
             print('Mean: {}, SD: {}'.format(m, sd))
In [ ]:
         eval alpha estimate(estimate alpha pdf ls, 'Least-squares on PDF');
        Least-squares on PDF:
        Mean: 0.7535055686849037, SD: 0.01445416636769601
In [ ]:
         eval alpha estimate(estimate alpha ccdf ls, 'Least-squares on CCDF');
        Least-squares on CCDF:
        Mean: 0.9985720588961311, SD: 0.010272736553571053
In [ ]:
         eval alpha estimate(estimate alpha mle, 'Log-likelihood maximization');
```

Log-likelihood maximization:
Mean: 1.0003310052420762, SD: 0.0032189410484160037

Least-squares regression on the PDF gives a very poor result (25% too small), while least-squares on the CCDF and the log-likelihood MLE both give results very close to the true value. The MLE is closer to the true value than CCDF least-squares by one order of magnitude, and has a smaller standard deviation, so I'd say that the MLE gives the best estimate.