

Statistical physics of graphs and networks

Project: study of the configuration model of random graphs

Master in physics of complex systems

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We first import a few packages:

```
In [45]: import numpy as np
import random
import networkx as nx
import matplotlib.pyplot as plt
import scipy.optimize as opt
import copy
```

We will use *networkx* to manage and plot networks.

We also define the parameter for plotting networks:

```
In [46]: options = {
    "with_labels": False,
    "node_size": 3,
    "node_color": "black",
    "edgecolors": "black",
    "linewidths": 2,
    "width": 3,
}

def plot_graph(gr):
    pos = nx.nx_agraph.graphviz_layout(gr, prog="neato")

    C = (gr.subgraph(c) for c in nx.connected_components(gr))
    for g in C:
        c = [np.random.random()] * nx.number_of_nodes(g) # random color...
        nx.draw(g, pos, node_size=40, node_color=c, vmin=0.0, vmax=1.0, with_
```

Problem 1: Generation of instances of the random graph model

First we will implement a few utility functions:

```
In [47]: def choose_and_pop_from_list(items):
    index = random.randrange(len(items))
    return items.pop(index)
```

And then move on to the main asked functions:

```

In [48]: def gen_rgm(N, pi):
    G = nx.Graph()
    for i in range(N):
        G.add_node(i)

    n_d1 = int(np.round(N*(1 - pi)/2)*2) # so that n_d1 + 4*n_d4 is pair
    n_d4 = N - n_d1

    stubs = []
    for i in range(n_d1):
        stubs.append(i)
    for i in range(n_d1, N):
        for j in range(4):
            stubs.append(i)

    while stubs:
        j = choose_and_pop_from_list(stubs)
        k = choose_and_pop_from_list(stubs)
        G.add_edge(j, k)
        G.add_edge(k, j)

    return G

def gen_rgm_biased(N, pi):
    G = nx.Graph()
    for i in range(N):
        G.add_node(i)

    n_d1 = int(np.round(N*(1 - pi)/2)*2) # so that n_d1 + 4*n_d4 is pair

    # use a dictionary of degrees
    degree = {}
    for i in range(N):
        degree[i] = 1 if i < n_d1 else 4

    while len(degree) >= 2:
        unique_stubs = list(degree.keys())

        # select a node and remove it from the temporary list of nodes to sel
        j = choose_and_pop_from_list(unique_stubs)
        while True:
            # try to select another node and remove it from the temporary lis
            k = choose_and_pop_from_list(unique_stubs)
            # only add edge if there isn't already an edge
            if not G.has_edge(j, k):
                G.add_edge(j, k)
                G.add_edge(k, j)

            # delete both index
            degree[j] -= 1
            # and if the degree is 0 remove the entry from the dict
            if degree[j] == 0:
                del degree[j]

            degree[k] -= 1
            if degree[k] == 0:
                del degree[k]

        break

```

```

        # if there is no more node to choose from:
        elif len(unique_stubs) == 0:
            # delete the entry for j because we know that we weren't able
            del degree[j]
            break

    return G

```

Which we then test:

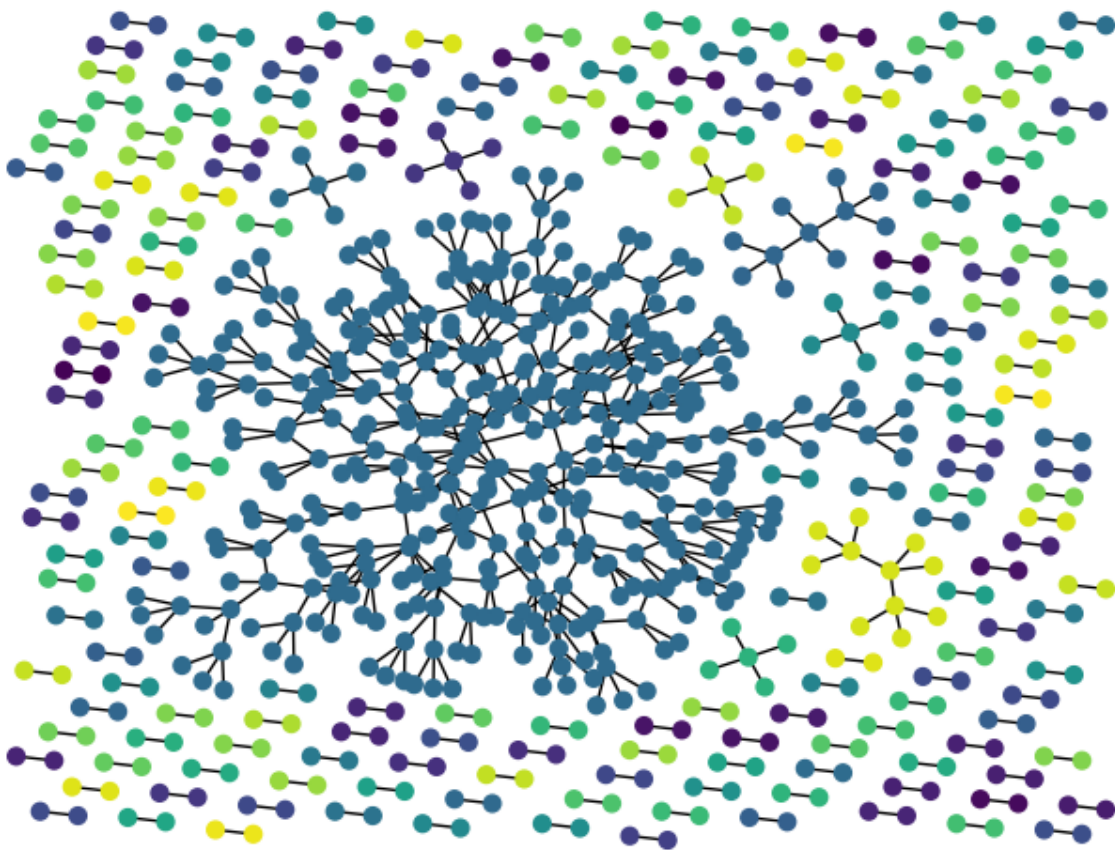
```

In [5]: size, pi = 800, 0.2
graph = gen_rgm_biased(size, pi)

plot_graph(graph)

plt.savefig("./presentation/images/generated_graph.png")

```



```

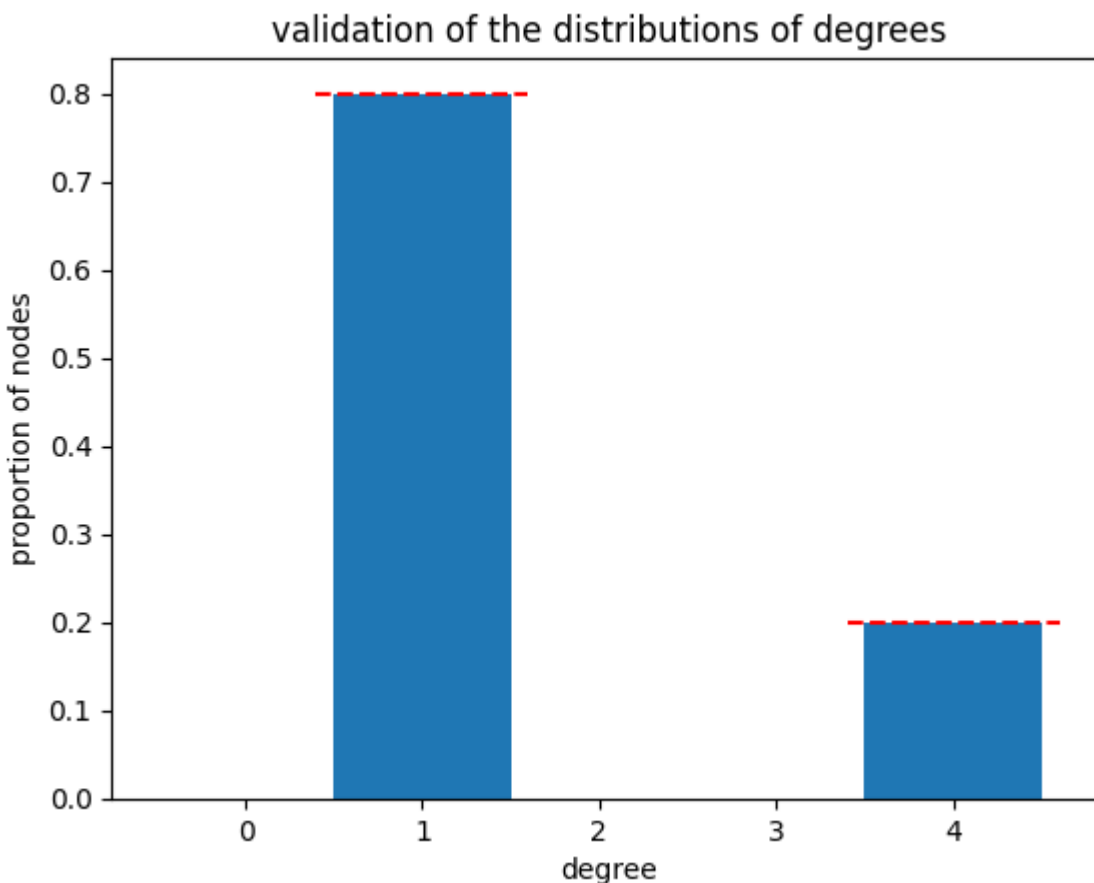
In [6]: degree_list = [graph.degree[i] for i in range(len(graph))]

plt.hist(degree_list, density=True, bins=[-0.5, 0.5, 1.5, 2.5, 3.5, 4.5])
plt.plot([0.4, 1.6], [1-pi, 1-pi], 'r--')
plt.plot([3.4, 4.6], [pi, pi], 'r--')

plt.title("validation of the distributions of degrees")
plt.xlabel("degree")
plt.ylabel("proportion of nodes")

plt.savefig("./presentation/images/degree_distribution.png")
plt.show()

```



Problem 2: The giant component

We first need a function that (recursively) walks through a graph to get the connected component starting from a given node:

```
In [49]: def get_connected_component(graph, node, visited=None):
    if visited is None:
        visited = []
    visited.append(node)

    for neighbor in graph.neighbors(node):
        if neighbor not in visited:
            connected = get_connected_component(graph, neighbor, visited)

            for connected_node in connected:
                if connected_node not in visited:
                    visited.append(connected_node)
    return visited
```

We then use this function to efficiently go through connected components (once per connected components), and return the largest connected components once there is not enough nodes to hope to find another even larger connected component:

```
In [50]: def find_largest_connected_component(graph):
    n_nodes = graph.number_of_nodes()
    not_visited = list(graph.nodes)

    largest_connected_component = []
```

```

while True:
    starting_node = not_visited[0]
    connected_component = get_connected_component(graph, starting_node)

    for node in connected_component:
        not_visited.remove(node)

    if len(connected_component) >= len(largest_connected_component):
        largest_connected_component = connected_component

    if len(not_visited) <= len(largest_connected_component):
        return largest_connected_component

```

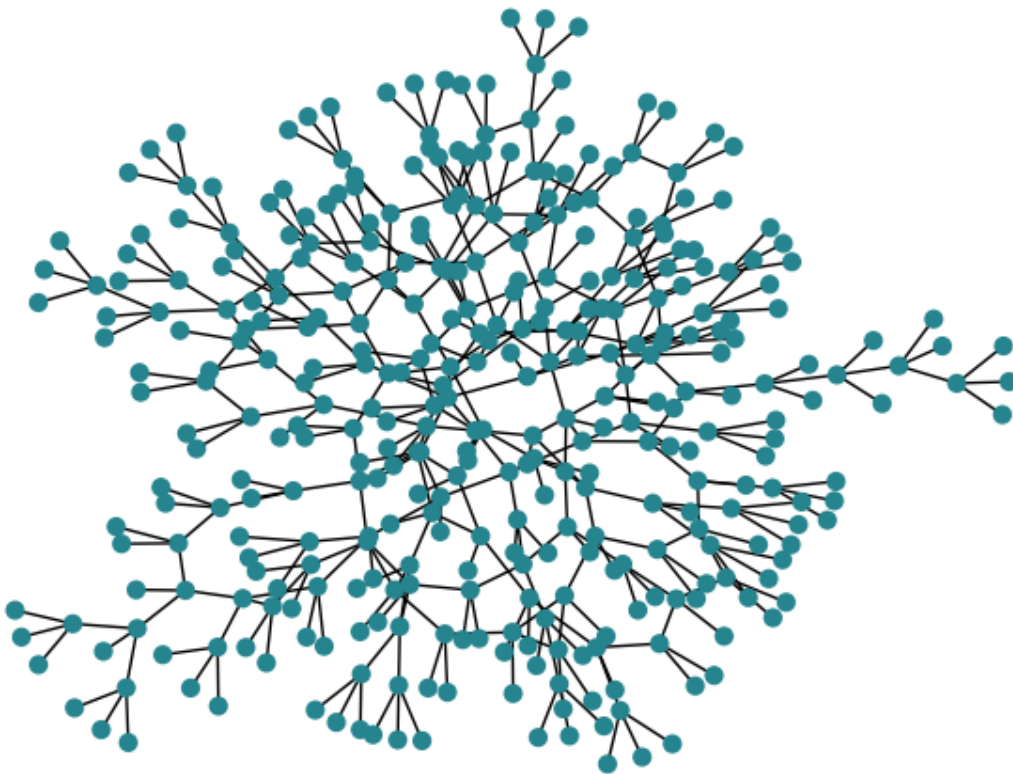
```

In [9]: largest_connected_component = find_largest_connected_component(graph)
largest_connected_subgraph = graph.subgraph(largest_connected_component)

plot_graph(largest_connected_subgraph)

plt.savefig("./presentation/images/generated_graph_giant_connected_component.

```



(a) We can now plot the average proportion of nodes inside of the giant component versus the value of π :

We here use a depth first search as we recursively search through the graph.

```

In [10]: Ns, sizes = [200, 20, 10], [20, 100, 500]
pis = np.linspace(0, 1, 15)

avg_size = []
for N, size in zip(Ns, sizes):

```

```

avg_size.append([])
for pi in pis:
    avg_size[-1].append(0)
    for n in range(N):
        graph = gen_rgm_biased(size, pi)
        largest_connected_component = find_largest_connected_component(gr
        avg_size[-1][-1] += len(largest_connected_component)
    avg_size[-1][-1] /= N*size

```

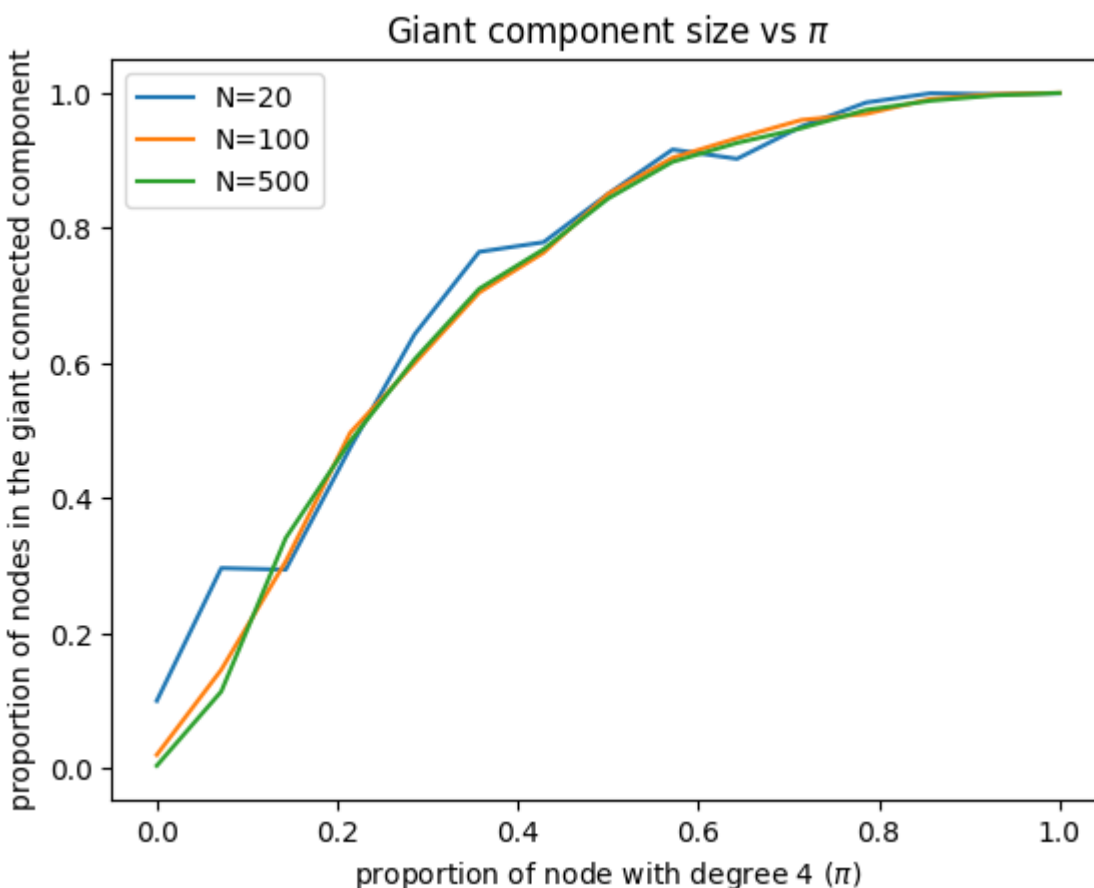
```

In [11]: for size, curve in zip(sizes, avg_size):
        plt.plot(pis, curve, label=f"N={ size }")

plt.title(r"Giant component size vs $\pi$")
plt.xlabel(r"proportion of node with degree 4 ($\pi$)")
plt.ylabel("proportion of nodes in the giant connected component")

plt.legend()
plt.show()

```



We observe that for increasing N (at least for our limited sample range) the curve doesn't evolve too much.

2-b: Theoretical analysis:

In this section, we study the statistics of the giant component of the graph. Let us first discuss the theory for this. Let us note π the probability for a vertex to have degree 4 (as in the exam sheet). Let us suppose I am at node i and I want to know with which probability is it ***not attached to the giant component**. *This is the same as asking the*

probability for one of its neighbours to **not be part of the giant component**. I have to sum the probability that each neighbour is connected to the giant component by one of its neighbours, except i . Let us write σ the probability that a not node is attached to the giant component, knowing it is already connected with a link to another node. We will write q_d the probability for a node to have degree d conditined on the fact that it has already degree 1.

$$\sigma = \sum_{d=0}^{+\infty} q_{d+1} \sigma^d = q_1 + q_4 \sigma^3$$

Now let us write γ th probability for a node to **be in the giant component***.

$$1 - \gamma = \sum_{q=0}^{+\infty} p_d \sigma^d = p_1 + p_4 \sigma^4$$

We have by definition:

$$q_d = \frac{dp_d}{\sum_{q=0}^{+\infty} dp_d}$$

$$q_1 = \frac{1 - \pi}{1 + 3\pi}$$

$$q_4 = \frac{4\pi}{1 + 3\pi}$$

We deduce the equations:

$$\gamma = 1 - \sigma [(1 - \pi) + \pi \sigma^3]$$

$$0 = 1 - \pi - (1 + 3\pi) \sigma + 4\pi \sigma^3$$

$$0 = (\sigma - 1) (4\pi \sigma^2 + 4\pi \sigma - (1 - \pi))$$

We can compute the solutions of the system as:

$$\sigma = 1$$

$$\sigma = \frac{-1 \pm \frac{1}{\sqrt{\pi}}}{2}$$

and

$$\gamma = 0$$

$$\gamma = 1 - \frac{-1 \pm \frac{1}{\sqrt{\pi}}}{2} \left(1 - \pi + \pi \left(\frac{-1 \pm \frac{1}{\sqrt{\pi}}}{2} \right)^3 \right)$$

The solution must be a probability $\gamma, \sigma \in [0, 1]$. So we will only keep the solutions:

$$\sigma = 0$$

$$\sigma = \frac{\frac{1}{\sqrt{\pi}} - 1}{2}$$

We implement the expression for γ in the next cell with a function called `p_giant`.

```
In [51]: def p_giant(pi):
    eps = 1e-18
    if pi == 0:
        return 0

    return max(0,
               1 - (1/np.sqrt(pi) - 1)/2 * (1 - pi + pi*((1/np.sqrt(pi) - 1))
```

(c) We can now compare the theoretical giant component size with our experimental results:

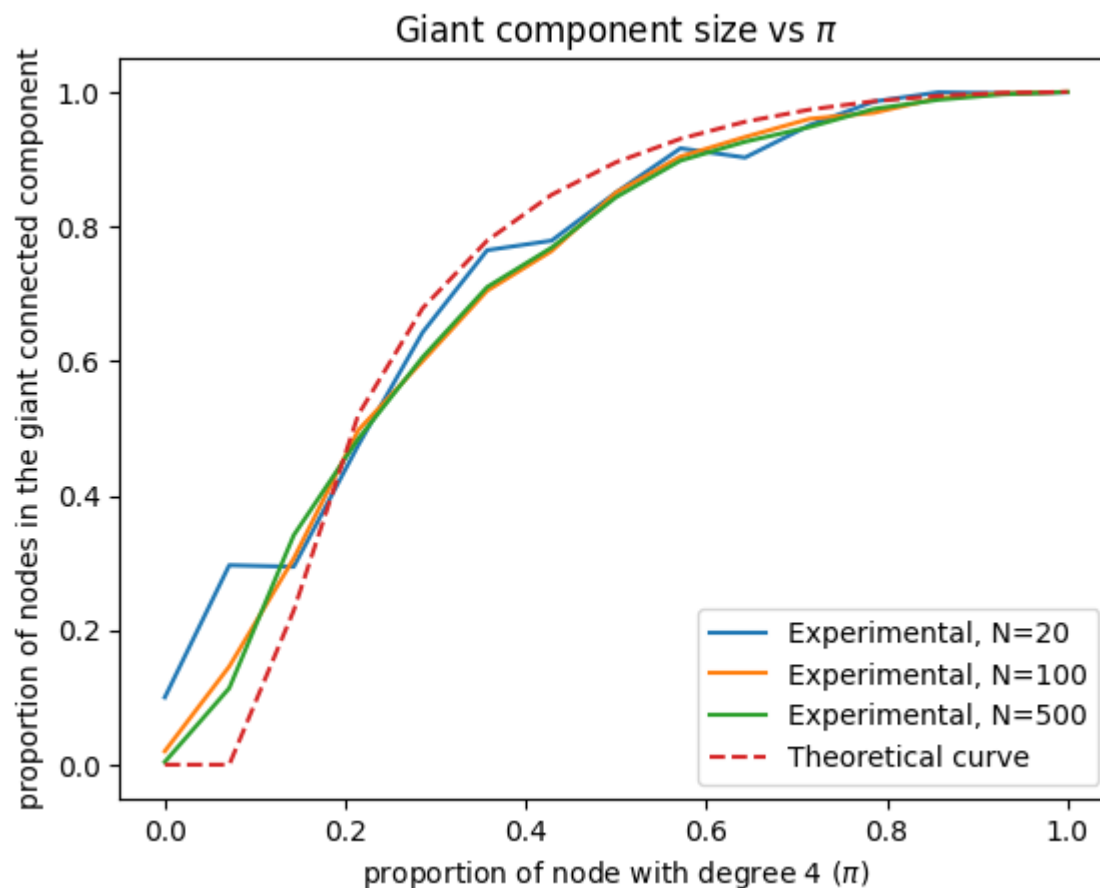
```
In [13]: for size, curve in zip(sizes, avg_size):
    plt.plot(pis, curve, label=f"Experimental, N={ size }")

plt.plot(pis, [p_giant(pi) for pi in pis], "--", label="Theoretical curve")

plt.title(r"Giant component size vs $\pi$")
plt.xlabel(r"proportion of node with degree 4 ($\pi$)")
plt.ylabel("proportion of nodes in the giant connected component")

plt.legend()

plt.savefig("./presentation/images/giant-connected-component.png")
plt.show()
```

We see that the experimental curve is non-zero for small p whereas our theoretically derived curve is zero for $\pi < \pi_0$ with some $\pi_0 > 0$.

We also observe that for all the N with tested, for small π our experimental results gives us bigger average giant connected component, whereas for π closer to 1 we see the opposite with a higher theoretical prediction.

We still observe the same quantization effect.

Problem 3: Emergence of the 3-core

```
In [52]: def find_k_core(graph, k): # pruning algorithm
    G = copy.deepcopy(graph)
    while True:
        to_remove=[]

        for node in G.nodes:
            if G.degree[node] < k:
                to_remove.append(node)

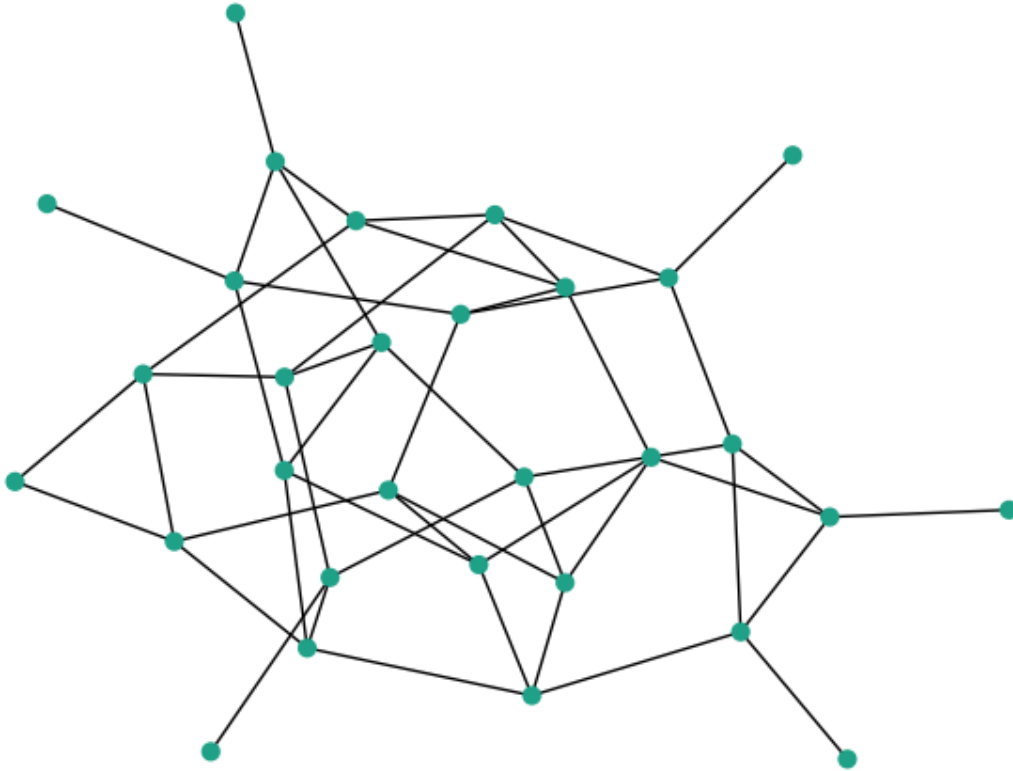
        if len(to_remove) == 0:
            if len(G.nodes) == 0:
                return []

            # the k-core still has to be a single connected graph
            k_core = find_largest_connected_component(G)
            return k_core

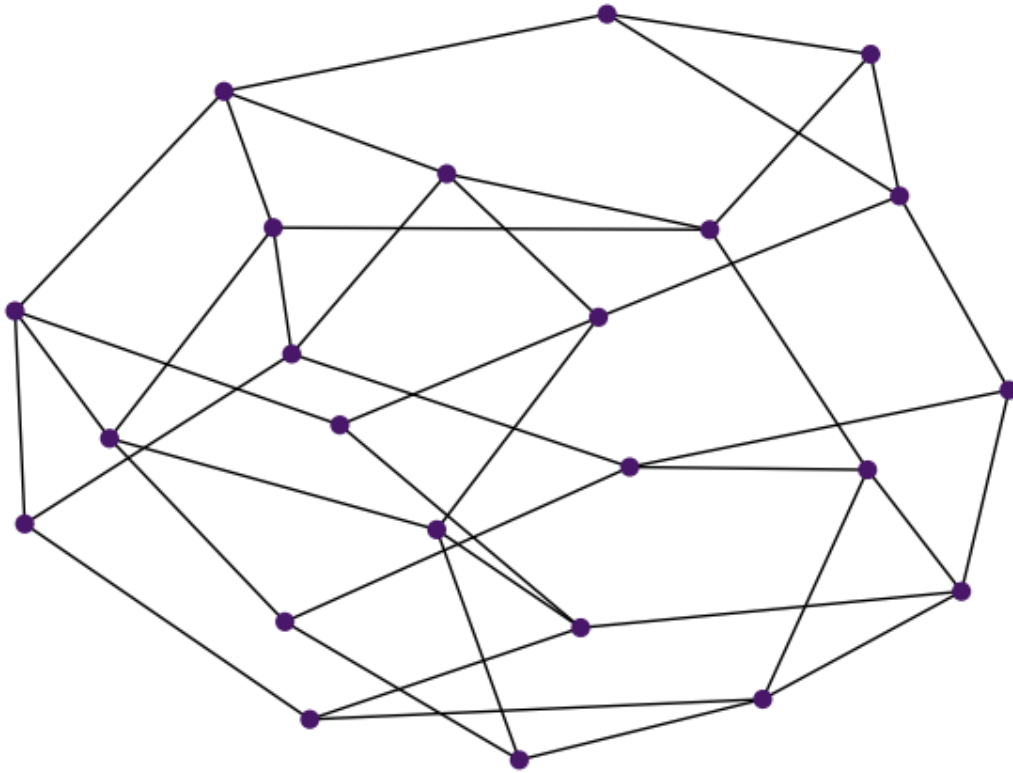
        for node in to_remove:
```

```
G.remove_node(node)
```

```
In [43]: graph = gen_rgm_biased(30, 0.8)
plot_graph(graph)
plt.savefig("./presentation/images/generated_graph2.png")
```



```
In [44]: core3_component = find_k_core(graph, 3)
core3_subgraph = graph.subgraph(core3_component)
plot_graph(core3_subgraph)
plt.savefig("./presentation/images/generated_graph2_3core.png")
```



(a) We can now plot the average proportion of nodes inside of the giant component versus the value of π :

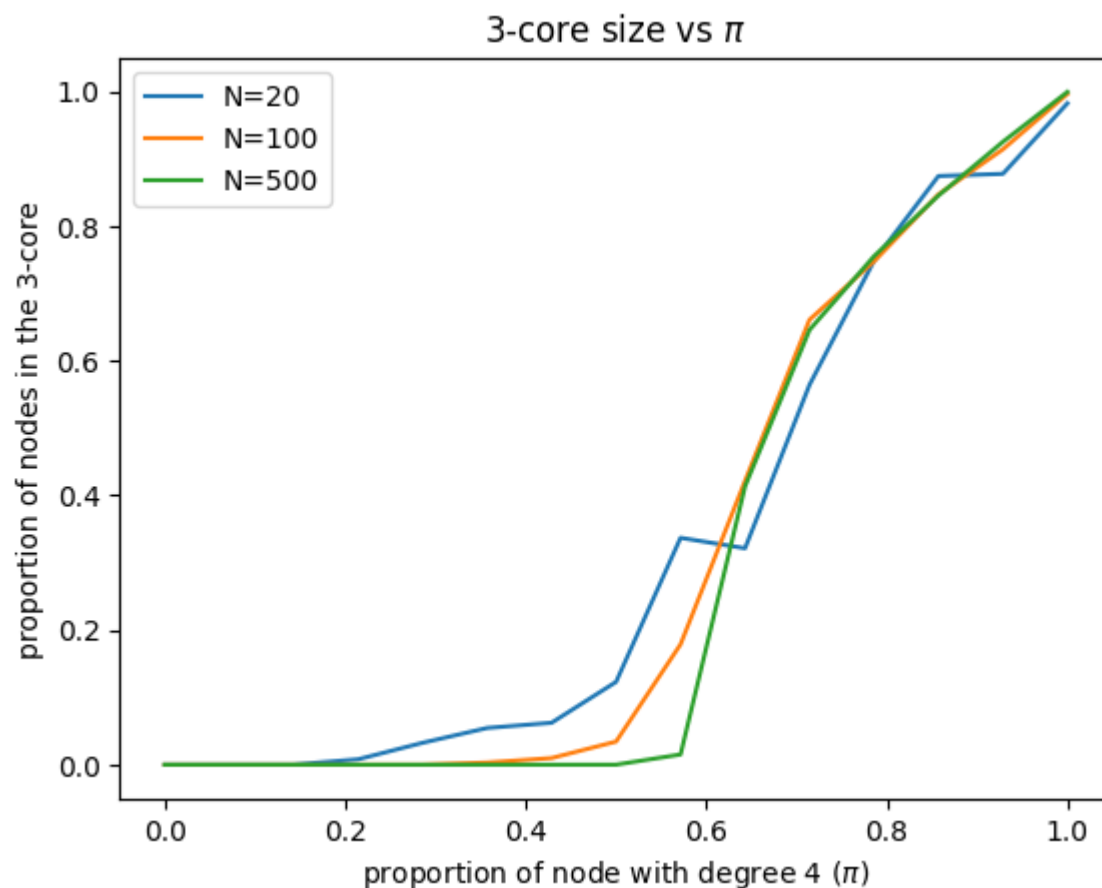
```
In [56]: Ns, sizes = [200, 100, 30], [20, 100, 500]
pis = np.linspace(0, 1, 15)

avg_size_3core = []
for N,size in zip(Ns,sizes):
    avg_size_3core.append([])
    for pi in pis:
        avg_size_3core[-1].append(0)
        for n in range(N):
            graph = gen_rgm_biased(size, pi)
            k_core = find_k_core(graph, 3)
            avg_size_3core[-1][-1] += len(k_core)
        avg_size_3core[-1][-1] /= N*size
```

```
In [57]: for size,curve in zip(sizes, avg_size_3core):
    plt.plot(pis, curve, label=f"N={ size }")

plt.title(r"3-core size vs  $\pi$ ")
plt.xlabel(r"proportion of node with degree 4 ( $\pi$ )")
plt.ylabel("proportion of nodes in the 3-core")

plt.legend()
plt.show()
```



...

3-b: Theoretical analysis:

We know from the course that the probability that a vertex is in the q-core is:

$$F_q(\beta_f) = 1 - \sum_{d=0}^{q-1} e^{-\beta_f} \frac{\beta_f^d}{d!}$$

With β_f is the fixed point of the series

$$\frac{\beta_{i+1}}{c} = F_{q-1}(\beta_i) \text{ And } c \text{ is the average degree of the vertexes: } c = \langle d \rangle = \sum_{d=0}^{\infty} d p_d$$

In our case, we obtain:

- $F_3 = 1 - e^{-\beta_f} \left(1 + \beta_f + \frac{\beta_f^2}{2} \right)$
- $\frac{\beta_f}{1+3\pi} = 1 - e^{-\beta_f} (1 + \beta_f)$
- $c = 1 - \pi + 4\pi = 1 + 3\pi$

We thus implement a function that returns the fraction of nodes in the 3-core, and the parameter β_f :

```
In [54]: def Three_Core_Theory(pi):
def F2(beta):
    return 1 - np.exp(-beta)*(1 + beta)
def F3(beta):
```

```

    return 1 - np.exp(-beta)*(1 + beta + beta**2*0.5)
def beta_func(beta, pi):
    return beta/(1 + 3*pi)

# function and its derivatives for
func      = lambda beta,pi: beta_func(beta, pi) - F2(beta)
dfdbeta   = lambda beta,pi: 1/(1 + 3*pi) - np.exp(-beta)*beta
df2dbeta2 = lambda beta,pi: np.exp(-beta)*(beta - 1)

try:
    Beta = opt.newton(func, 4., fprime=dfdbeta, fprime2=df2dbeta2, args=(
except: # If error return the (0,0) solution
    return 0., 0.

return F3(Beta), Beta

```

(c) We can now compare the theoretical giant component size with our experimental results:

```

In [58]: for size, curve in zip(sizes, avg_size_3core):
    plt.plot(pis, curve, label=f"Experimental N={ size }")

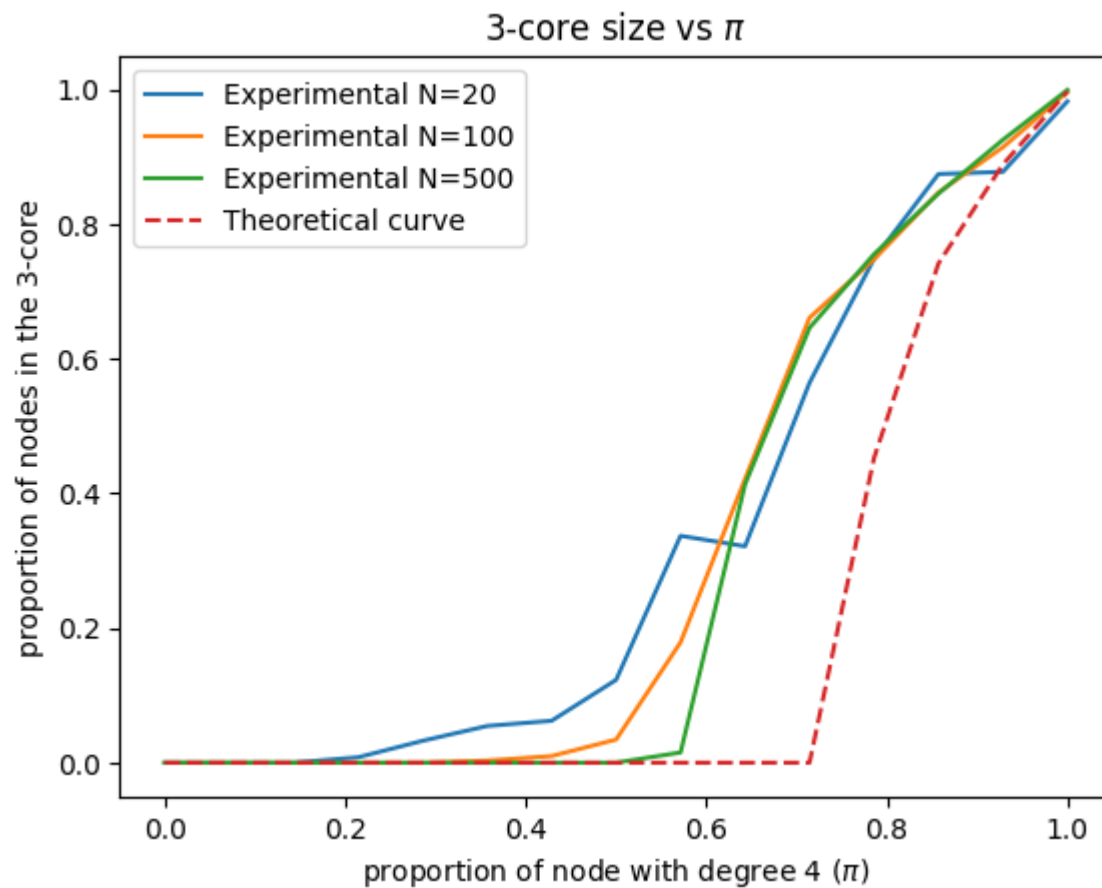
plt.plot(pis, [Three_Core_Theory(pi)[0]*3/2 for pi in pis], "--", label="Theo

plt.title(r"3-core size vs $\pi$")
plt.xlabel(r"proportion of node with degree 4 ($\pi$)")
plt.ylabel("proportion of nodes in the 3-core")

plt.legend()

plt.savefig("./presentation/images/3-core.png")
plt.show()

```



We see that for higher N the π_0 at which the 3-core emerges (the average size of the 3 core becomes non-zero) is pushed toward the high- π , and thus toward our theoretically-derived curve.

Problem 4: The ferromagnetic Ising model

```
In [21]: def gen_random_state(N_node, p=0.5):
    random_state = np.random.rand(N_node)
    state = 2*(random_state < p) - 1
    return state
```

```
def get_state_magnetisation(state):
    return np.mean(state)
```

```
In [22]: def monte_carlo_simulate(graph, state, T, N_it_per_node):
    n_nodes = graph.number_of_nodes()

    for i in range(n_nodes*N_it_per_node):
        node = np.random.randint(0, n_nodes)

        total_surrounding_spin = 0
        for neighbor in graph.neighbors(node):
            total_surrounding_spin += state[neighbor]

        delta_E = 2*state[node]*total_surrounding_spin
        if delta_E < 0:
            state[node] = -state[node]
        else:
            P_flip = np.exp(-delta_E/T)
```

```

        if np.random.rand() < P_flip:
            state[node] = -state[node]

    return state

```

```

In [23]: graph = gen_rgm_biased(100, 0.8)
        state = gen_random_state(100)

        print(get_state_magnetisation(monte_carlo_simulate(graph, state, 10, 100)))
        print(get_state_magnetisation(monte_carlo_simulate(graph, state, 0.5, 100)))

-0.18
-0.98

(1) TODO

```

```

In [24]: N, size, n_it = 30, 50, 100
        Ts = np.linspace(0.5, 8, 30)

        graph = gen_rgm_biased(size, 0.8)

        magnetisations = []
        for T in Ts:
            magnetisations.append([])
            for i in range(N):
                state = gen_random_state(100)
                final_state = monte_carlo_simulate(graph, state, T, n_it)
                magnetisation = get_state_magnetisation(final_state)

                magnetisations[-1].append(magnetisation)

```

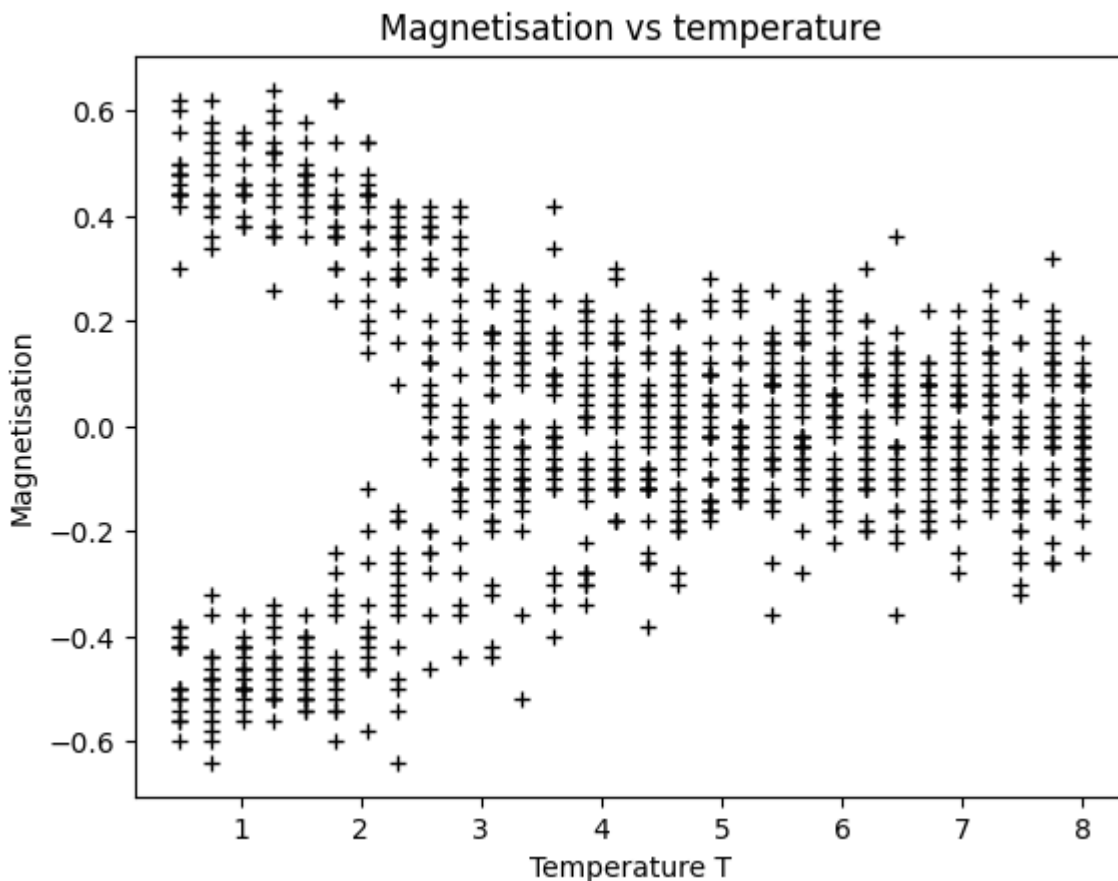
```

In [25]: for T, mag in zip(Ts, magnetisations):
        plt.plot([T]*N, mag, "k+")

        plt.title("Magnetisation vs temperature")
        plt.xlabel(r"Temperature T")
        plt.ylabel("Magnetisation")

        plt.savefig("./presentation/images/magnetisation.png")
        plt.show()

```



We here can see a phase transition from $m = \pm m_{eq}$ to $m = 0$ at $T \approx 3$.

```
In [26]: N, size, n_it = 10, 50, 100
pis = np.linspace(0, 1, 20)
Ts = np.linspace(0.5, 5, 20)

PIS, TS = np.meshgrid(pis, Ts)

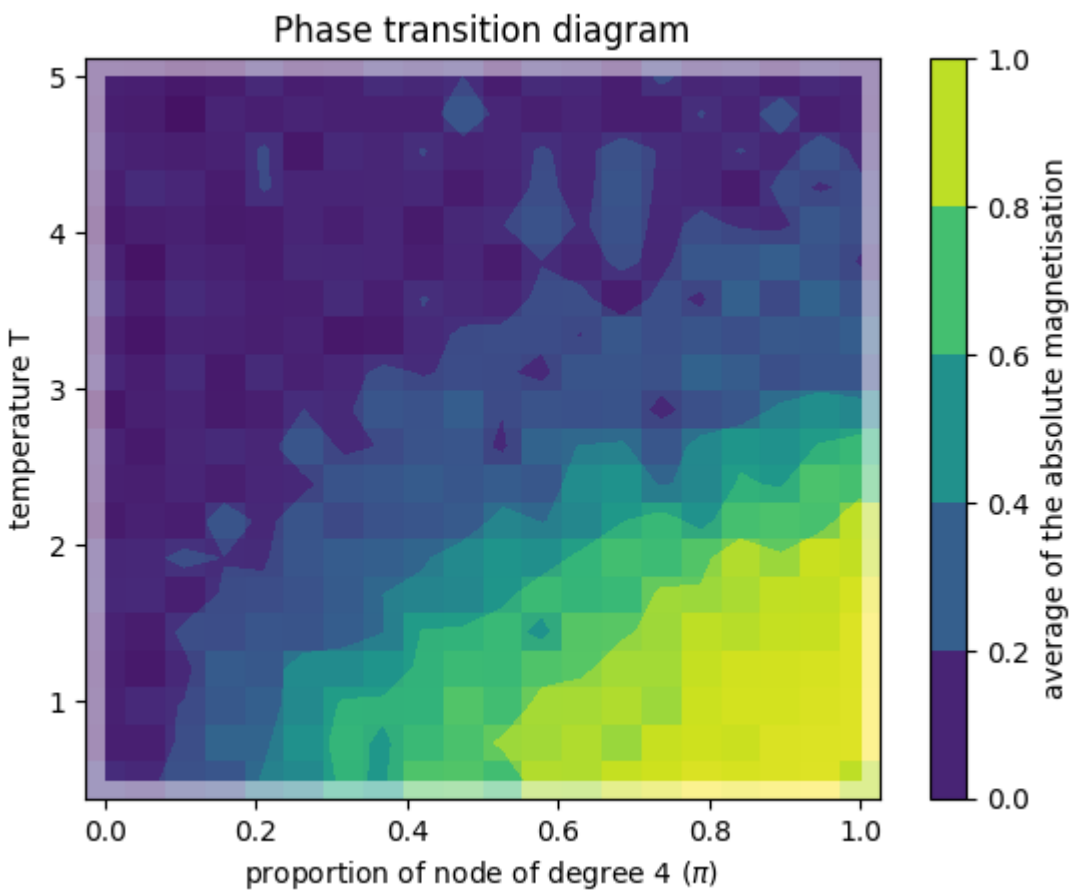
avg_m = np.zeros_like(PIS)
for i in range(PIS.shape[0]):
    for j in range(PIS.shape[1]):
        for n in range(N):
            graph = gen_rgm_biased(size, PIS[i, j])
            state = gen_random_state(size)

            equilibrium_state = monte_carlo_simulate(graph, state, TS[i, j],
            avg_m[i, j] += abs(get_state_magnetisation(equilibrium_state))
avg_m /= N
```

```
In [27]: plt.contourf(PIS, TS, avg_m, 5)
plt.colorbar(label="average of the absolute magnetisation")
plt.pcolormesh(PIS, TS, avg_m, alpha=0.5)

plt.title("Phase transition diagram")
plt.xlabel(r"proportion of node of degree 4 ( $\pi$ )")
plt.ylabel("temperature T")

plt.savefig("./presentation/images/phase_transition_diagram.png")
plt.show()
```

(2) We can now study the

In []:

(3) TODO

In []:

Problem 5: Inverse Ising model

In []:

(1) TODO

```
In [28]: size, T, pi = 50, 4, 0.4
         N, n_it = 500, 100

         graph = gen_rgm_biased(size, pi)
```

```
In [29]: states = np.zeros((N, size))
         for n in range(N):
             state = gen_random_state(size)
             equilibrium_state = monte_carlo_simulate(graph, state, T, n_it)
             states[n, :] = equilibrium_state
```

```
In [30]: correlations = np.zeros((size, size))
         avgs = np.mean(states, axis=0)
```

```

for i in range(size):
    for j in range(i): # range(i + 1), force corr[i, i] = 0 so self edges are
        for n in range(N):
            correlations[i, j] += states[n, i]*states[n, j]
        correlations[i, j] /= N

    correlations[i, j] -= avgs[i]*avgs[j]
    correlations[j, i] = correlations[i, j]

pairs = []
for i in range(size):
    for j in range(i):
        pairs.append((i, j))
cor_pairs = zip([correlations[pair] for pair in pairs], pairs)
cor_pairs_sorted = reversed(sorted(cor_pairs))
pairs_sorted = [pair for cor, pair in cor_pairs_sorted]

```

```

In [31]: n_predictions = np.arange(1, int(graph.number_of_edges()*2))
prediction_acc = np.zeros_like(n_predictions, dtype=float)

for i in range(len(n_predictions)):
    for pair in pairs_sorted[:n_predictions[i]]:
        prediction_acc[i] += float(graph.has_edge(pair[0], pair[1]))
    prediction_acc[i] = float(prediction_acc[i])/n_predictions[i]

```

```

In [32]: plt.plot(n_predictions, prediction_acc)

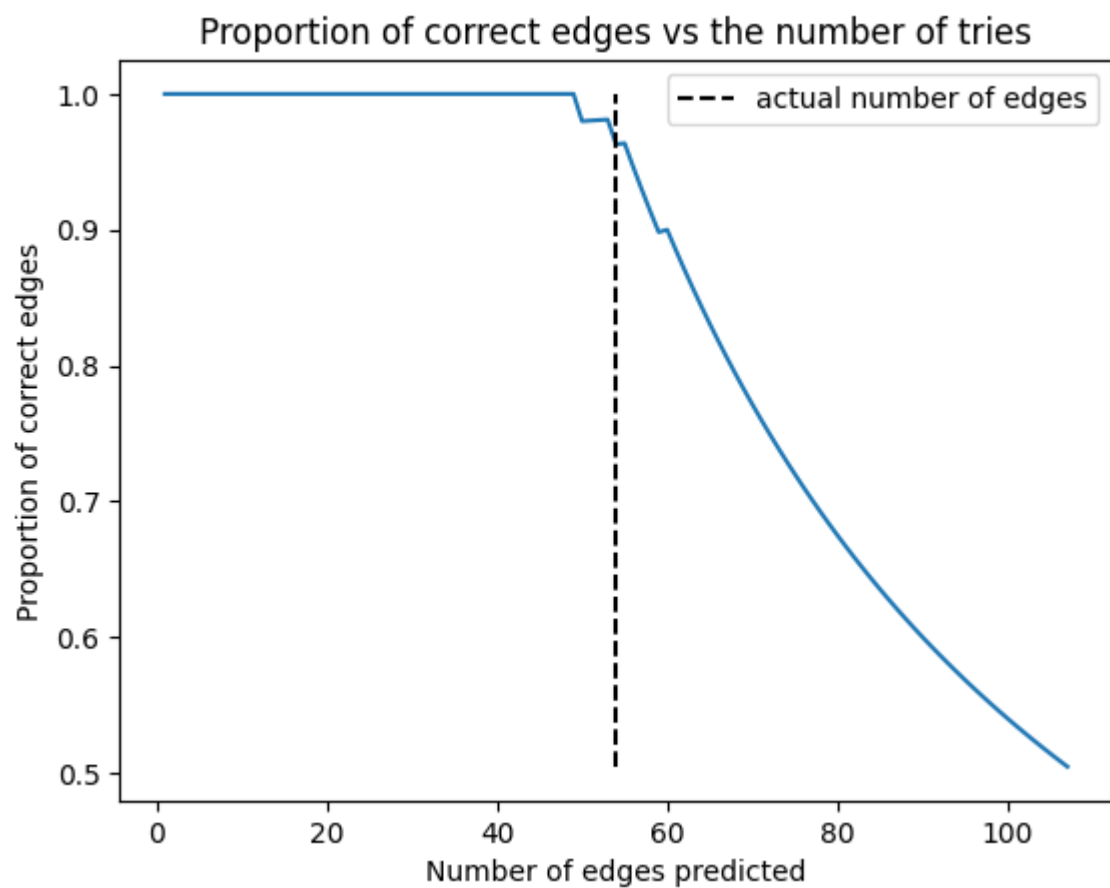
plt.plot([graph.number_of_edges(), graph.number_of_edges()],
         [min(prediction_acc), max(prediction_acc)],
         "k--", label="actual number of edges")

plt.title("Proportion of correct edges vs the number of tries")
plt.xlabel("Number of edges predicted")
plt.ylabel("Proportion of correct edges")

plt.legend()

plt.savefig("./presentation/images/reconstruction_accuracy_plot.png")
plt.show()

```



(2) TODO

```
In [33]: Jij = np.linalg.inv(correlations)
```

```
In [ ]:
```

(3) TODO

```
In [ ]:
```