EE 232E Project 1

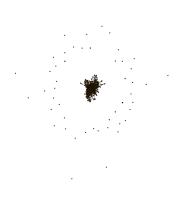
Random Graph and Random Walks

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I. GENERATING RANDOM NETWORKS

A. Create random networks using Erdös-Rényi model

(a) By creating the undirected networks with number of nodes N=1000 and p=0.003, 0.004, 0.01, 0.05, and 0.1, we observe the binomial distribution. The results are shown in Fig 1, 2, 3, 4, 5, 6, 7, 8, 9, 10.



Degree distribution of the network with n = 1000, p=0.003

Fig. 1: Random network G_1 with N = 1000 and p = 0.003.

Fig. 2: Degree distribution of G_1



Fig. 3: Random network G_2 with N=1000 and p=0.004.

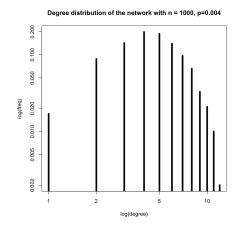


Fig. 4: Degree distribution of G_2

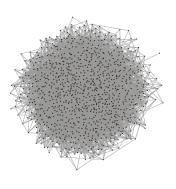


Fig. 5: Random network G_3 with N=1000 and p=0.01.

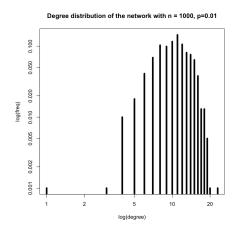
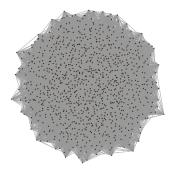


Fig. 6: Degree distribution of G_3



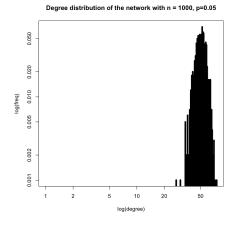
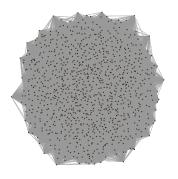


Fig. 7: Random network G_4 with N=1000 and p=0.05.

Fig. 8: Degree distribution of G_4



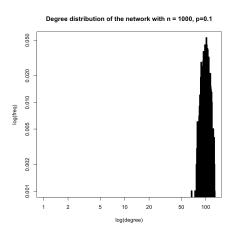


Fig. 9: Random network G_5 with N=1000 and p=0.05.

Fig. 10: Degree distribution of G_5

Below is the comparison between the empirical and theoretical values of mean and variance for the above 5 random networks.

For G_1

$$\tilde{\mathbb{E}}[\deg(G_1)] = 4.122, \quad \text{var}(\deg(G_1)) = 3.305116$$

$$\mathbb{E}[\deg(G_1)] = 2.997, \quad \operatorname{var}(\deg(G_1)) = 2.988009$$

For G_2

$$\tilde{\mathbb{E}}[\deg(G_2)] = 5.028, \quad \text{var}(\deg(G_2)) = 4.461216$$

$$\mathbb{E}[\deg(G_2)] = 3.996, \quad \operatorname{var}(\deg(G_2)) = 3.980016$$

For G_3

$$\tilde{\mathbb{E}}[\deg(G_3)] = 10.796, \quad \text{var}(\deg(G_3)) = 9.142384$$

$$\mathbb{E}[\deg(G_3)] = 9.99, \quad \text{var}(\deg(G_3)) = 9.8901$$

For G_4

$$\tilde{\mathbb{E}}[\deg(G_4)] = 51.222, \quad \text{var}(\deg(G_4)) = 45.54472$$

$$\mathbb{E}[\deg(G_4)] = 49.95, \quad \operatorname{var}(\deg(G_4)) = 47.4525$$

For G_5

$$\tilde{\mathbb{E}}[\deg(G_5)] = 100.772, \quad \text{var}(\deg(G_5)) = 97.11202$$

$$\mathbb{E}[\deg(G_5)] = 99.9, \quad \operatorname{var}(\deg(G_5)) = 89.91$$

- (b) The network G_1 , G_2 and G_3 are disconnected whereas the network G_4 and G_5 are connected. Taking G_1 as an instance, the size of GCC is 941 and the diameter is 13.
- (c) The normalized GCC size versus the probability p in the range of $(0, \frac{\ln n}{n}]$ is given in Fig 11, where n is the number of nodes in the network. The result matches the theoretical values derived in class.

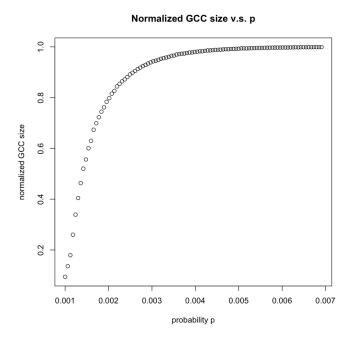


Fig. 11: The normalized GCC size versus the probability p in the range of $(0, \frac{\ln n}{n}]$

(d) Fig 12 shows the expected size of the GCC v.s. the number of nodes n where c=np=0.5.

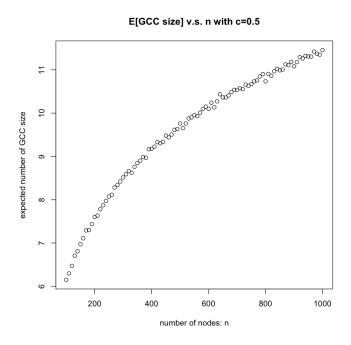


Fig. 12: The expected size of the GCC v.s. the number of nodes n when c = np = 0.5

The result for c = np = 1 is shown in Fig 13.

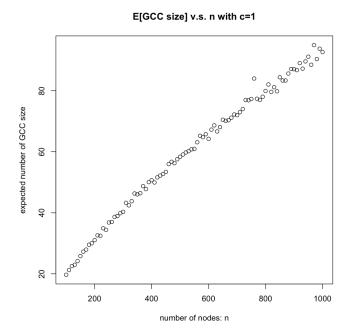


Fig. 13: The expected size of the GCC v.s. the number of nodes n when c = np = 1

The result for c = 1.1, 1.2 and 1.3 is shown in Fig 14.

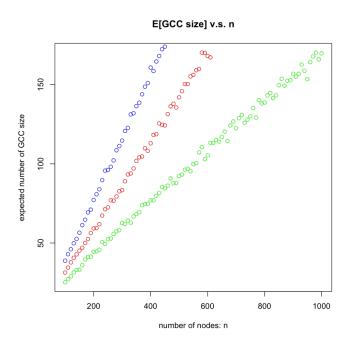


Fig. 14: The expected size of the GCC v.s. the number of nodes n when $c=1.1,\ 1.2$ and 1.3

B. Create networks using preferential attachment model

(a) We created an undirected network under preferential attachment model with n=1000 and m=1, i.e. through the generating process, every step a new node joins the network, we pick only one node proportional to its current degree to connect the new node and there are 1000 steps in total. The network is shown below. We can see that the network we generated is always connected from both practical result and theory analysis.

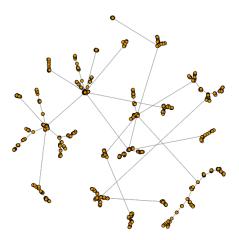


Fig. 15: undirected graph with 1000 nodes based on preferential attachment model

- (b) We used fast greedy method to find the community structure and the modularity of this network equals to 0.932. The Network has a very high modularity which means it has dense connections between the nodes within modules but sparse connections between nodes in different modules.
- (c) This time we tried to generate a larger network with 10000 nodes using the same preferential attachment model. And similarly we applied same algorithm to measure modularity. The modularity of this larger network is 0.978, which is higher in comparison to the smaller network.



Fig. 16: undirected graph with 10000 nodes based on preferential attachment model

- (d) We plotted the degree distribution in a log-log scale for both 1000 nodes and 10000 nodes network. Taking advantage of linear regression, we can estimate the slope of those two plots. The slope of the network plot with 1000 nodes is -2.53 the other plot's slope is -2.91. Theoretically, the ideal log-log degree distribution plot's slope of the network generated under preferential attachment model should be -3.
- (e) We randomly picked a node i, and then randomly picked a neighbor j of that node. The degree distribution plot of nodes j that are picked with this process p in the log-log scale is shown below. This degree distribution is a little bit different from the original degree distribution. The frequency of high degree nodes is higher than the one of original degree distribution.
 - (f) The expected degree of a node that is added at time step i is given by the function below.

$$deg(i) = m(\frac{t}{i})^{\frac{1}{2}} = (\frac{1000}{i})^{\frac{1}{2}}$$

The Plot of relationship between the age of nodes and their expected degree is shown below.

(g) Now we repeated all the previous parts for m=2, and m=5. For m=2, the networks generated are shown blow. And the modularity of the network plot with 1000 nodes is 0.521, the larger network's modularity is 0.532.

The regular and randomly walking degree distribution of the networks are shown below.

And the relationship between the age of nodes and expected degree is shown below.

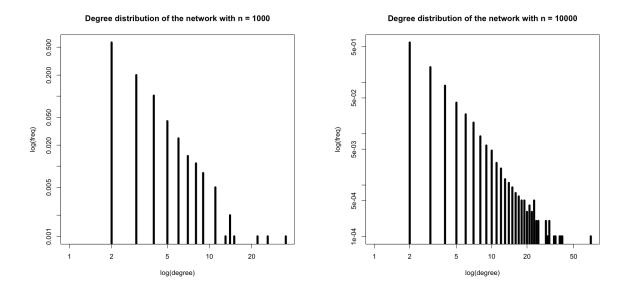


Fig. 17: degree distribution with 1000 nodes Fig. 18: degree distribution with 10000 nodes

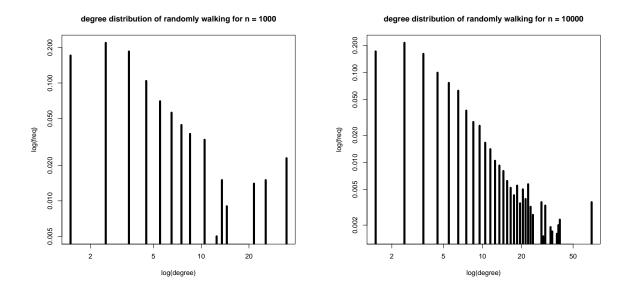


Fig. 19: degree distribution with 1000 nodes Fig. 20: degree distribution with 10000 nodes

Similarly, for m=5, the networks generated are shown blow. And the modularity of the network plot with 1000 nodes is 0.279, the larger network's modularity is 0.272.

The regular and randomly walking degree distribution of the networks are shown below.

And the relationship between the age of nodes and expected degree is shown in Fig 35.

We can see from the comparison that the network with m=1 has the highest modularity.

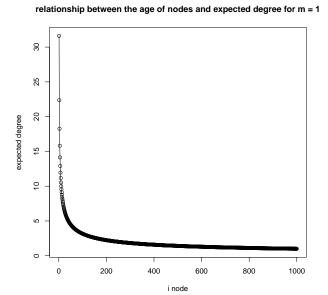


Fig. 21: relationship between the age of nodes and their expected degree

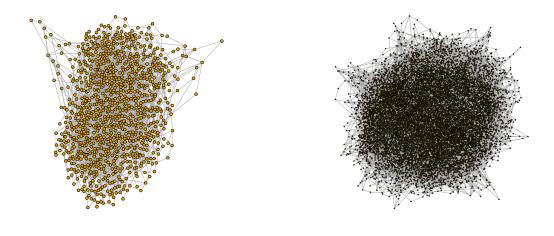
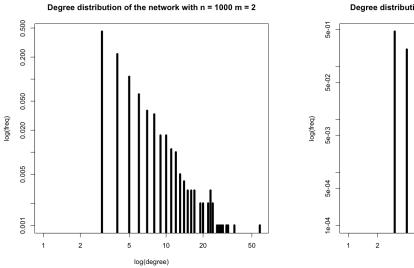


Fig. 22: network with 1000 nodes

Fig. 23: network with 10000 nodes

Since the higher m, the higher the total degree of all the nodes, there is no doubt that the nodes in the network with higher total degree connect to each other much more closely, which means the network becomes a whole instead of several communities.

(h) Again, we generated a preferential attachment network with n = 1000, m = 1 and then



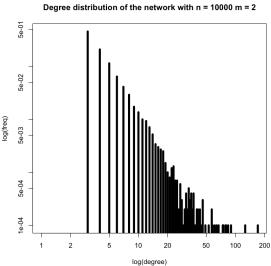
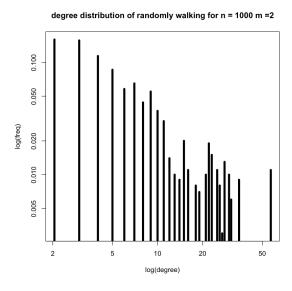
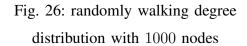


Fig. 24: degree distribution with 1000 nodes Fig. 25: degree distribution with 10000 nodes





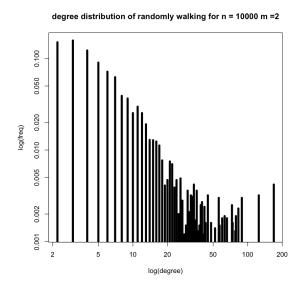


Fig. 27: randomly walking degree distribution with 10000 nodes

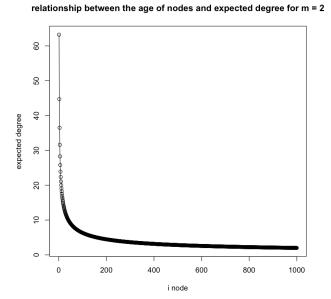


Fig. 28: relationship between the age of nodes and expected degree

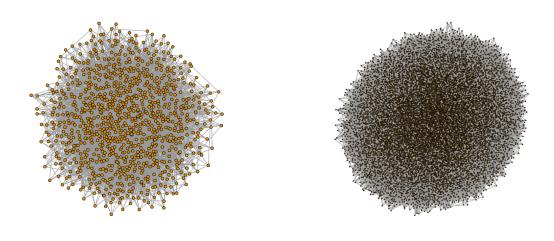


Fig. 29: network with 1000 nodes

Fig. 30: network with 10000 nodes

took its degree sequence and created a new network with the same degree sequence through submatching procedure. Those two networks marked as communities with different color are shown below. The modularity of original network is 0.926, on the other hand, the generated through submatching procedure network's modularity is 0.840.

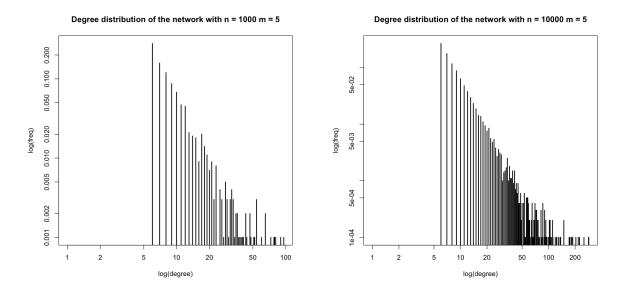


Fig. 31: degree distribution with 1000 nodes Fig. 32: degree distribution with 10000 nodes

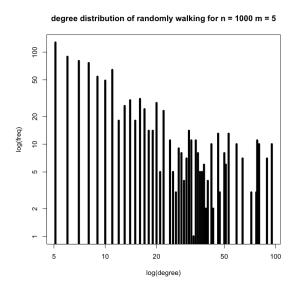


Fig. 33: randomly walking degree distribution with 1000 nodes

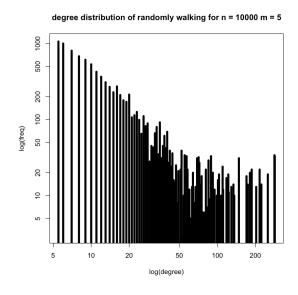


Fig. 34: randomly walking degree distribution with 10000 nodes

relationship between the age of nodes and expected degree for m = 5

Fig. 35: relationship between the age of nodes and expected degree

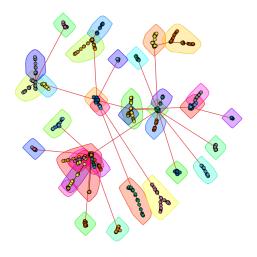


Fig. 36: network marked as communities

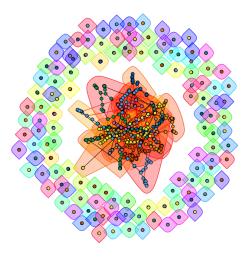


Fig. 37: generated network marked as communities

The submatching method to create random power law network couldn't guarantee that the network is fully connected. And we can induce from the result plot that this method will create a network with high modularity.

C. Create a modified preferential attachment model that penalizes the age of a node

(a) This time, we aimed to penalize the age of a node through creating process under preferential attachment model. That is that each time a new vertex was added, the probability that an old vertex was cited depends on its degree and age. In particular, the probability that a newly added vertex connects to an old vertex was proportional to:

$$P(i) \sim (ck_i^{\alpha} + a)(dl_i^{\beta} + b)$$

where k_i is the degree of vertex i in the current time step, and li is the age of vertex i. We created this network with 1000 nodes and parameters m=1, $\alpha=1$, $\beta=1$, and a=c=d=1, b=0. By linear regression, the practical slope of degree distribution log-log plot equals to -2.745 that is the power law exponent.

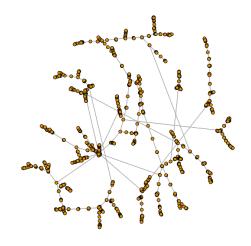


Fig. 38: network plot

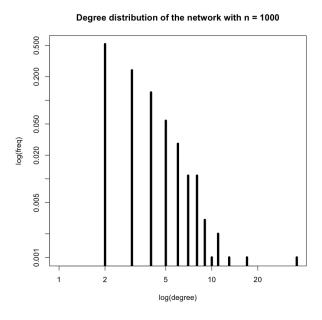


Fig. 39: degree distribution

(b) Again, we applied fast greedy method to find the community structure. The modularity of this network is 0.935.

II. RANDOM WALK ON NETWORKS

A. Random walk on Erdös-Rényi networks

(a) Same as what we have done in the previous part, we generated an undirected random network containing 1000 nodes with the probability of 0.01 to add an edge between any pair of nodes. The graph we got is shown below:

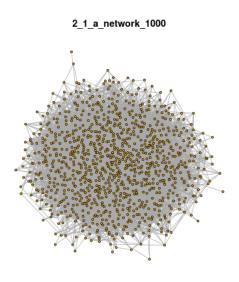


Fig. 40: undirected graph with 1000 nodes based on Erdös-Rényi Model

- (b) After simulating random walk, we got following results:
- (c) The degree distribution of the last reached nodes in N times of the random walk and the degree distribution of the graph are shown below.

The results illustrate that the real graph degree distribution has similar trends with the degree distribution of the ending node by random walk. Both of them follows a Gaussian distribution. This could be proved by doing some simple calculation under the Erdös-Rényi Model. Every time a new node joins the network has the same probability (p) to add an edge to each existing nodes. That is to say, for all the nodes, the degree of a node follows binomial distribution.

$$p_k = \binom{n-1}{k} p^k (1-p)^{n-1-k} \tag{1}$$

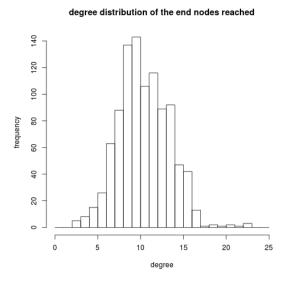
When the number of nodes N is large, Gaussian distribution could be seen the approximation of binomial distribution.

step

standard deviation of distance for 1000 nodes

Fig. 41: s(t) v.s. t with N = 1000

Fig. 42: $\sigma^2(t)$ v.s. t with N=1000



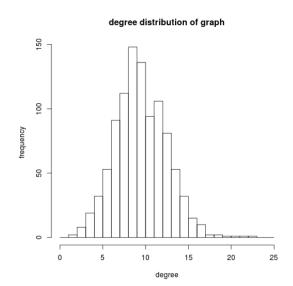
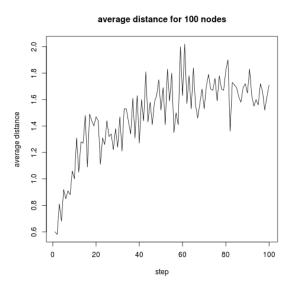


Fig. 43: degree distribution of the ending node with nodes = 1000 in network

Fig. 44: degree distribution of the graph with nodes = 1000 in network

(d) To figure out the difference of the shortest path length with different number of nodes in the network, we repeated what we have done in problem(b) under 100 nodes and 10000 nodes respectively in the network. The following figures show the average distance and standard deviation of distance under 100 nodes and 10000 nodes respectively.

The diameter of the nodes N=100, N=1000 and N=10000 is about 2.0, 3.5 and 2.5 respectively. This seems not what we have expected, so we ran program several times to generate different networks with certain number of nodes. Not surprisingly, we found that graph with more nodes tend to have smaller diameter and smaller average distance. In addition, we can get another consequence from figures bellow that for the graph with 100 nodes, the average distance converge slowly or even cannot converge if we do not set the start node fixed. On the contrary, the results show less fluctuation with 10000 nodes. In conclusion, the smaller the diameter is, the average distance and standard deviation of the distance converge more quickly and turn out to have smaller fluctuation.



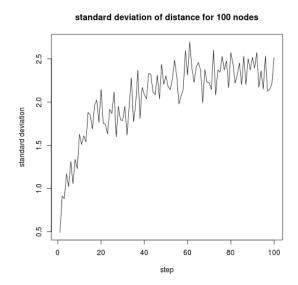
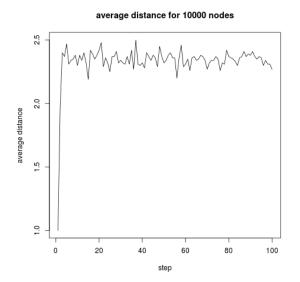


Fig. 45: s(t) v.s. t with nodes = 100

Fig. 46: $\sigma^2(t)$ v.s. t with nodes = 100

B. Random walk on networks with fat-tailed degree distribution

(a) We used function *sample_pa()* to generate a fat-tailed network according to the Barabasi-Albert Model. The graph we got is shown bellow.



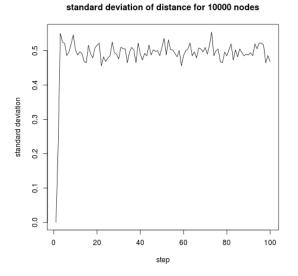


Fig. 47: s(t) v.s. t with nodes = 10000

Fig. 48: $\sigma^2(t)$ v.s. t with nodes = 10000

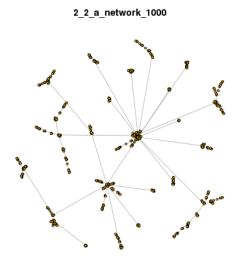
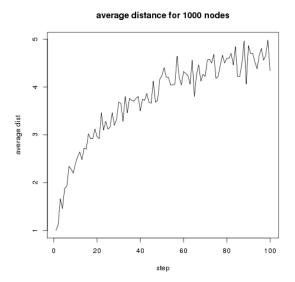


Fig. 49: undirected graph with 1000 nodes based on Barabasi-Albert Model

(b) As what we have dealt with Erdös-Rényi networks, we also simulated random walk on preferential attachment network and Fig 50 and 51 demonstrate the plot s(t) v.s. t and $\sigma^2(t)$ v.s. t respectively. (c) After running several times to get several figures(shown bellow), we could see that after a certain large number of steps of random walk, the results of two kinds of degree



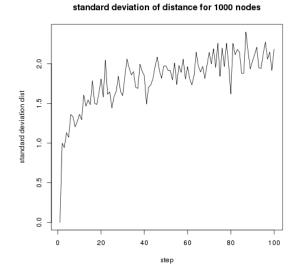
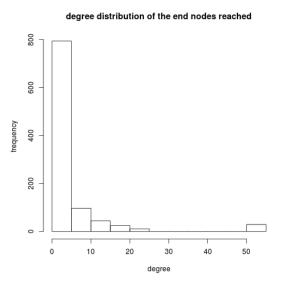


Fig. 50: s(t) v.s. t with nodes = 1000

Fig. 51: $\sigma^2(t)$ v.s. t with nodes = 1000

distribution are similar to each other.



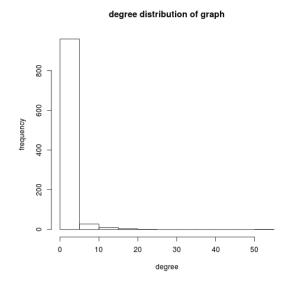


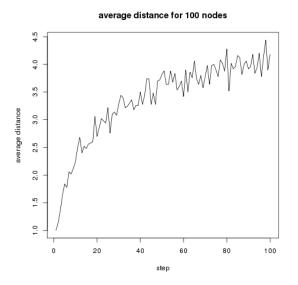
Fig. 52: degree distribution of the ending node with nodes = 1000 in network

Fig. 53: degree distribution of the graph with nodes = 1000 in network

(d) Repeated the process in (b), we got the diameters of graphs with 100 and 10000 nodes are 4.5 and 5 respectively. Though we got this result, according to the theory, we knew that this seemed incorrect. Therefore, we generate network several times and calculate the average of the

diameter. Then, we got the conclusion that in general, the graph with more nodes has smaller diameter, which means it also has smaller average distance.

The results also illustrate some differences between two different graph generation models. Considering the graph with 1000 nodes, in previous model(Erdös-Rényi Model), we can see that the average distance increases dramatically within not more than 10 steps, and then the value keeps fluctuating along the step t and gradually converges. However in Barabasi-Albert Model, the average distance shows a slight increase along the step t.



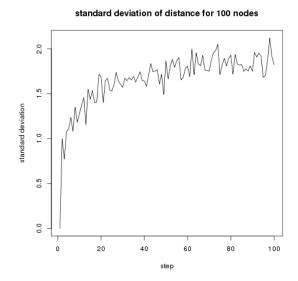


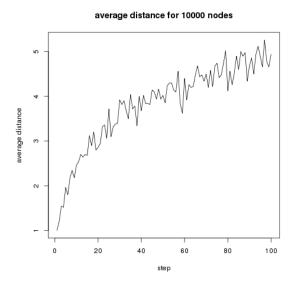
Fig. 54: s(t) v.s. t with nodes = 100

Fig. 55: $\sigma^2(t)$ v.s. t with nodes = 100

C. PageRank

PageRank is a famous algorithm designed by Larry Page, one of the founder of Google. Although there are a few limitations of this algorithm, many other ranking algorithms are invented under its inspiration. In this section, we applied random walk based on PageRank and got some interesting results which illustrate the principle of PageRank.

(a) The graph we generated based on preferential attachment is shown bellow.



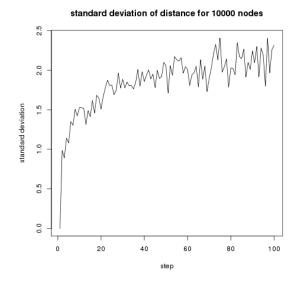


Fig. 56: s(t) v.s. t with nodes = 10000

Fig. 57: $\sigma^{2}(t)$ v.s. t with nodes = 10000

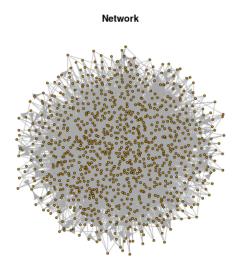
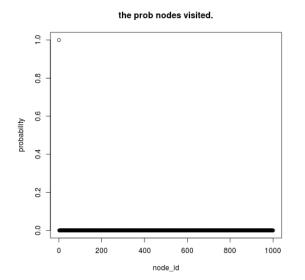


Fig. 58: directed graph with 1000 nodes using preferential attachment

From the results shown bellow, it could be concluded that the visit probability is proportional to the node degree. The higher the node degree is, the more possible the walker would visit the node. (b) Considering the teleportation probability of $\alpha=0.15$, we get slightly different results. It demonstrates that the difference of visiting probability along degree of the node is getting



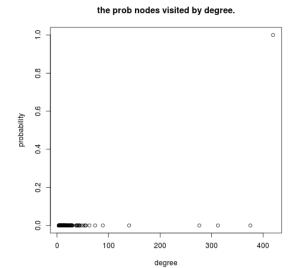
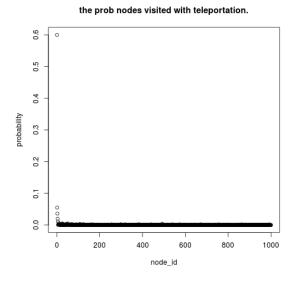


Fig. 59: probability that the walker visits each node

Fig. 60: visit probability v.s. degree



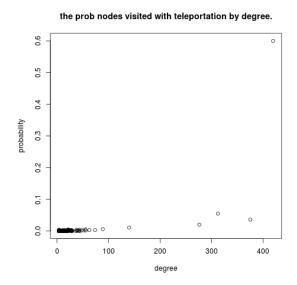


Fig. 61: probability that the walker visits each node with teleportation probability $\alpha=0.15$

Fig. 62: visit probability v.s. degree with teleportation probability $\alpha=0.15$

larger.

D. Personalized PageRank

(a) In Personalized PageRank, we set the teleportation probability proportionally to PageRank of each node. The results are shown bellow. Comparing to 3(a), the results show that Personalized

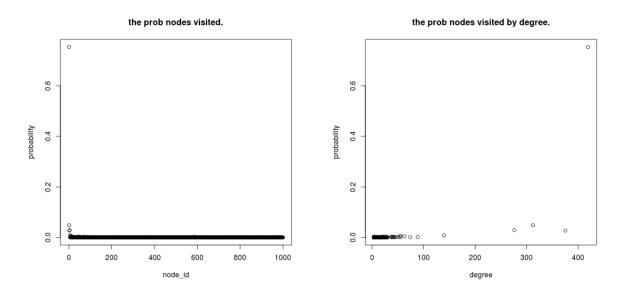
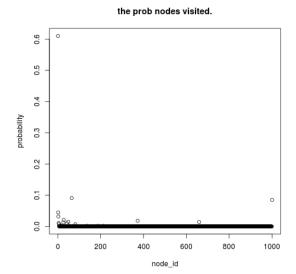


Fig. 63: probability that the walker visits each node under Personalized PageRank

Fig. 64: visiting probability v.s. degree under Personalized PageRank

PageRank can make degree of the node influence more to the visiting probability, which is in fact similar to what we get in 3(b). The higher degree the node is, the more possible it will be visited. In other words, since it is a preferential attachment model, the longer time a node stays in the network, the more possible it will be visited.

(b) The results are shown bellow. Comparing to all the results we get from PageRank or Personalized PageRank, under this circumstance, the visiting probability varies more significantly along the degree of a node. Specifically, some nodes with less than 100 degree have high probability to be visited than some other nodes with degree around less than 50. Nevertheless, this rarely happens in previous model, where nodes with degree of 300 or higher may be more possible to be visited than other nodes, while nearly most of nodes within the degree of 100 have similar visiting probability of around 0. Though it appears some irregular results within the degree of 100 (the visiting probability does not follow the rule exactly), the general rule that



the prob nodes visited by degree.

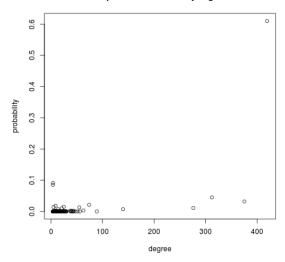


Fig. 65: probability that the walker visits each node under fixed teleportation

Fig. 66: visiting probability v.s. degree under fixed teleportation

nodes with higher degree have more chance to be visited does not change. (c) Since at this time, nodes can only teleport to trusted nodes, which is similar to the previous problem that only two nodes are allowed to teleport to. Here, only the nodes belong to trusted node set are allowed to teleport to. Thus, the equation would be like this:

$$r = \alpha \cdot T \cdot r + (1 - \alpha) \cdot d$$

where T is the transition matrix and vector d can be used to assign a non-zero score to the set of trusted pages only. In this way, trusted pages will have higher probability to be visited, and what these trusted pages point to can also be regarded as trustworthy pages, and their visiting probability will increase since we make restriction to teleport to trusted pages only.