Algorithmic Data Science - Exercise Series 2

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Exercise 1

(a)

We have that

$$h_{a,b}(x) = h_{a,b}(y)$$
$$\Longleftrightarrow ax + b \equiv ay + b \mod m$$
$$\Longleftrightarrow a(x - y) \equiv 0 \mod m$$

which in the case of $x = m, \ y = 0$ is true $\forall a, b$ therefore

$$P(h_{a,b}(x) = h_{a,b}(y)) = 1 > \frac{1}{m}$$

meaning that the family is not universal.

(b)

This exercise is *Theorem 11.5* in [1].

Let $x, y \in \mathbb{Z}_p : x \neq y$.

Define

$$u := ax + b \mod p$$
$$v := ay + b \mod p$$

We have that $u \neq v$ since $u - v \equiv a(x - y) \not\equiv 0 \mod p$ because $a \neq 0$, $x - y \not\equiv 0 \mod p$, and $\mathbb{Z}_p$ is a field. Note that $x - y \not\equiv 0 \mod p$ holds because by hypothesis $x \neq y$ and $x, y < p$.

Therefore, there are no collisions when we apply $x \mapsto ax + b \mod p$.

We proceed to show that $(a, b) \mapsto (ax+b \mod p, \ ax+b \mod p)$ is a bijection between the pairs $(a, b) \in \mathbb{Z}_p^* \times \mathbb{Z}_p$ and the pairs $(u, v) \in \mathbb{Z}_p \times \mathbb{Z}_p : u \neq v$.

We can solve for a, b and get a unique solution

$$a = \frac{u - v}{x - y} \mod p$$
$$b = r - ak \mod p$$

Where $\frac{1}{t}$ is the inverse of t in $\mathbb{Z}_p$

Therefore the mapping is one to one. Since we also have that both the domain and the co-domain have $p(p-1)$ elements, the mapping is a bijection. Thus, if (a, b) is uniformly distributed, so is (u, v).

Therefore, the probability that $x, y \in \mathbb{Z}_p : x \neq y$ collide is equal to the probability that $u \equiv v \mod m$ collide when $(u, v) \in \mathbb{Z}_p \times \mathbb{Z}_p : u \neq v$ are chosen uniformly randomly. We proceed to calculate that probability.

Given u, of the $p - 1$ possible remaining values for v we have that at most $\lceil \frac{p}{m} \rceil - 1 \leq \frac{p-1}{m}$ can collide with u.

Therefore the probability of collision is $\leq \frac{1}{m}$, meaning that the hash function family is universal.

(c)

The proof in is still valid, since $x \in U \implies x < p$ is still valid. Therefore the hash function family remains universal.

Exercise 2

(a)

From what I understand, the expected number of probes for a successful search is not equal to the expected number of probes for an insertion. What [1] says in the proof of *Corollary 11.7* is that the expected number of probes for an *Unsuccessful* search is equal to the expected number of probes for an insertion, and that number is bounded above by $\frac{1}{1-a}$ where a is the load factor.

The expected number of probes a for a successful search on the other hand is bounded above by $\frac{1}{a} \ln \frac{1}{1-a}$, as shown in *Theorem 11.8*.

If the expected values were equal, wouldn't the book bound them by the same number?

(b)

First we need to show that the expected number of probes in an unsuccessful search is bounded by $\frac{1}{1-a}$.

Let X be a random variable describing the number of probes in an unsuccessful search.

We have

$$\mathrm{E}[X] = \sum_{i=0}^{\infty} i \Pr(X = i)$$

$$= \sum_{i=1}^{\infty} \Pr(X \geq i)$$

Now,

$$\Pr(X \geq i) = \frac{n}{m} \cdot \frac{n-1}{m-1} \cdots \frac{n-i+2}{m-i+2}$$

$$\leq \left(\frac{n}{m}\right)^{i-1}$$

$$= a^{i-1}$$

Therefore,

$$\mathrm{E}[X] \leq \sum_{i=1}^{\infty} a^{i-1} = \frac{1}{1-a}$$

Since an element is inserted only if there is room in the table (thus $a < 1$), inserting a key requires an unsuccessful search followed by placing the key into

the first empty slot found. Thus, the expected number of probes for an insertion is at most $\frac{1}{1-a}$.

We now proceed to prove the result about the successful search.

A search for a key reproduces the same probe sequence as when the element with key was inserted. By the above result, if the element was the $(i+1)$st element inserted, then the expected number of probes made in a search for it is at most $\frac{1}{1-1/m} = \frac{m}{m-i}$.

Therefore the expected number of probes for a successful search is

$$\frac{1}{n} \sum_{i=0}^{n-1} \frac{m}{m-i} = \frac{1}{a} \sum_{i=0}^{n-1} \frac{1}{m-i}$$

$$= \frac{1}{a} \sum_{k=m-n+1}^{m} \frac{1}{k}$$

$$\leq \frac{1}{a} \int_{m-n}^{m} \frac{1}{x} dx$$

$$= \frac{1}{a} \ln \frac{m}{m-n}$$

$$= \frac{1}{a} \ln \frac{1}{1-a}$$

Exercise 3

(a)

We proceed to calculate the time complexity of the Girvan-Newman algorithm, shown in Algorithm 1. We iterate over $|V|$ nodes, so everything will be multiplied by $|V|$. The BFS step takes $O(|V|+|E|)$ time. The accumulation with DFS also takes $O(|V|+|E|)$ time. The betweenness calculation takes $O(|V|+2|E|) = O(|V|+|E|)$ time as well. Therefore, we end up with $O(|V|(|V|+|E|))$ time.

(b)

In the case of a tree, since a tree is by definition an undirected acyclic graph, it can also be viewed as a DAG, meaning that we don't need to perform the BFS step. Also, in the second DFS part, since tree nodes have only one parent, we don't need to compute the u.npaths/s factor because it's always equal to 1. The complexity doesn't change in the end, since the DFS still takes $O(|V|+|E|)$ time. Finally in a tree we have that $|E| = |V| - 1$. Thus the complexity can now be written as $O(|V|^2)$.

Algorithm 1 Girvan-Newman

1: **for** each e in E **do**
2:     e.betweenness ← 0
3: **end for**
4:
5: **for** each r in V **do**
6:     G ← DAG by performing BFS from r
7:     **for** each v in G **do**
8:         v.npaths ← 1
9:     **end for**
10:     Cumulatively sum node npaths from leaves to root (post-order DFS)
11:     **for** each v in G **do**
12:         v.credit ← 1
13:     **end for**
14:     **for** v in post-order DFS(r) **do**
15:         s ← 0
16:         **for** neighbor u of v incoming from above **do**
17:             s ← s + u.npaths
18:         **end for**
19:         **for** each edge e incoming to v from the above node u **do**
20:             e.tempbetweenness ← v.credit * u.npaths / s
21:             u.credit ← u.credit + e.tempbetweenness
22:         **end for**
23:     **end for**
24:     **for** each e in E **do**
25:         e.betweenness ← e.betweenness + e.tempbetweenness/2
26:     **end for**
27: **end for**

Exercise 4

Exercise 10.4.1 of [2]

The adjacency matrix is

$$A = \begin{pmatrix} 0 & 1 & 1 & 0 & 0 & 0 & 0 & 0 & 0 \\ 1 & 0 & 1 & 0 & 0 & 0 & 0 & 1 & 0 \\ 1 & 1 & 0 & 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 1 & 0 & 1 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 1 & 0 & 1 & 1 & 0 & 0 \\ 0 & 0 & 0 & 1 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 & 0 & 1 & 1 \\ 0 & 1 & 0 & 0 & 0 & 0 & 1 & 0 & 1 \\ 0 & 0 & 0 & 0 & 0 & 0 & 1 & 1 & 0 \end{pmatrix}$$

The degree matrix is

$$D = \mathrm{diag}(2, 3, 3, 3, 3, 2, 3, 3, 2)$$

The Laplacian matrix is

$$L = \begin{pmatrix}
2 & -1 & -1 & 0 & 0 & 0 & 0 & 0 & 0 \\
-1 & 3 & -1 & 0 & 0 & 0 & 0 & -1 & 0 \\
-1 & -1 & 3 & -1 & 0 & 0 & 0 & 0 & 0 \\
0 & 0 & -1 & 3 & -1 & -1 & 0 & 0 & 0 \\
0 & 0 & 0 & -1 & 3 & -1 & -1 & 0 & 0 \\
0 & 0 & 0 & -1 & -1 & 2 & 0 & 0 & 0 \\
0 & 0 & 0 & 0 & -1 & 0 & 3 & -1 & -1 \\
0 & -1 & 0 & 0 & 0 & 0 & -1 & 3 & -1 \\
0 & 0 & 0 & 0 & 0 & 0 & -1 & -1 & 2
\end{pmatrix}$$

Exercise 10.4.2 of [2]

Using Wolfram Mathematica's function `Eigensystem` we find out that second smallest eigenvalue of L after 0 is $\frac{5-\sqrt{13}}{2}$ with corresponding eigenvectors

$$v_1 = \begin{pmatrix}
\frac{5+\sqrt{13}}{2} \\
\frac{3+\sqrt{13}}{2} \\
\frac{1+\sqrt{13}}{2} \\
-\frac{7+\sqrt{13}}{1+\sqrt{13}} \\
\frac{-3-\sqrt{13}}{2} \\
-\frac{3(3+\sqrt{13})}{1+\sqrt{13}} \\
-1 \\
1 \\
0
\end{pmatrix}, \quad v_2 = \begin{pmatrix}
-\frac{2(4+\sqrt{13})}{1+\sqrt{13}} \\
\frac{-1-\sqrt{13}}{2} \\
-2 \\
1 \\
2 \\
\frac{7+\sqrt{13}}{1+\sqrt{13}} \\
\frac{\sqrt{13}-1}{2} \\
0 \\
1
\end{pmatrix}$$

The partition is based on the signs of the coordinates of the eigenvector. Eigenvector v_1 partitions the graph into $\{A, B, C, H, I\}$ and $\{D, E, F, G\}$, while eigenvector v_2 partitions the graph into $\{A, B, C\}$ and $\{D, E, F, G, H, I\}$.

Exercise 5

(a)

References

[1] T. H. Cormen, C. E. Leiserson, R. L. Rivest, and C. Stein, *Introduction to Algorithms*, 3rd ed. MIT Press, 2009.

[2] J. Leskovec, A. Rajaraman, and J. Ullman, *Mining of Massive Datasets*, 3rd ed. Cambridge University Press, 2019.