

Cpt S 515 Homework #4

Instructor: Zhe Dang

Student: Sheng Guan

11593929

1. I have k , for some k , water tanks, T_1, \dots, T_k (which are identical in size and shape), whose water levels are respectively denoted by nonnegative real variables x_1, \dots, x_k . Without loss of generality, we assume that x_i equals the amount of water that is currently in T_i . Initially, all the tanks are empty; *i.e.*, $x_i = 0$; $1 \leq i \leq k$. I have m pumps p_1, \dots, p_m , that pump water into tanks. More precisely, a pump instruction, say, PA, c_1, c_2 , where $A \subseteq \{T_1, \dots, T_k\}$, is to pump the same amount of water to each of the tank T_i with $i \in A$ (so water levels on other tanks not in A will not change), where the amount is anywhere between c_1 and c_2 (including c_1 and c_2 , of course we have assumed $0 \leq c_1 \leq c_2$). For instance, $P\{T_2, T_5\}, 1.5, 2.4$ means to pump simultaneously to T_2 and T_5 the same amount of water. However, the amount can be anywhere between 1.5 and 2.4. Suppose that we execute the instruction twice, say:

$P\{T_2, T_5\}, 1.5, 2.4$;

$P\{T_2, T_5\}, 1.5, 2.4$.

The first $P\{T_2, T_5\}, 1.5, 2.4$ can result in 1.8 amount of water pumped into T_2 and T_5 , respectively, and the second $P\{T_2, T_5\}, 1.5, 2.4$ can result in 2.15 amount of water pumped into T_2 and T_5 , respectively. That is, the amount of water can be arbitrarily chosen inside the range specified in the instruction, while the choice is independent between instructions.

Now, let M be a finite state controller which is specified by a directed graph where each edge is labeled with a pump instruction. Different edges may be labeled with the same pump instruction and may also be labeled with different pump instructions. There is an initial node and a final node in M . Consider the following condition $Bad(x_1, \dots, x_k)$:

$$x_1 = x_2 + 1 = x_3 + 2 \wedge x_3 > x_4 + 0.26.$$

A walk in M is a path from the initial to the final. I collect the sequence of pump instructions on the walk. If I carefully assign an amount (of water pumped) for each such pump instruction and, as a result, the water levels x_1, \dots, x_k at the end of the sequence of pump

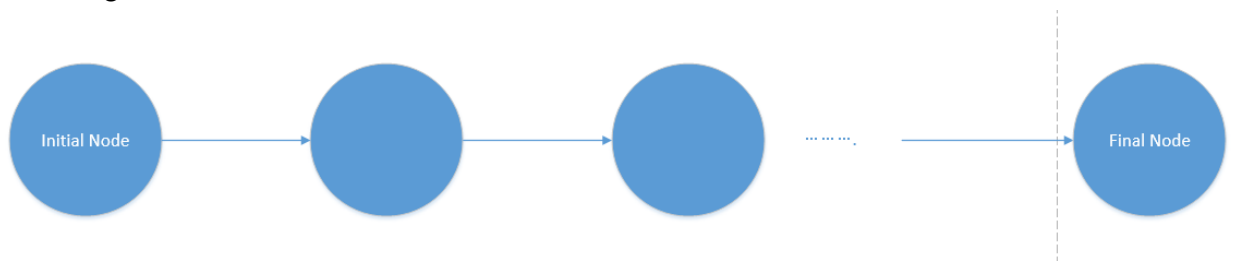
instruction satisfy $Bad(x_1, \dots, x_k)$, then I call the walk is a bad walk. Such a walk intuitively says that there is an undesired execution of M.

Design an algorithm that decides whether M has a bad walk. (Hint: first draw an example M where there is no loop and see what you can get. Then, draw an M that is with a loop and see what you get. Then, draw an M that is with two nested loops and see what you get, and so on.)

Answer:

(1) We start with the simplest case first that the walk M does not contain any loop:

Assume we have a walk from the initial node to the final node and it looks like the below figure:



The algorithm-1 idea is: first, we use DFS to search such walks from the initial node to the final node and record all walks;

To simplify our analysis, we use the example Bad Walk definition proposed by the question to illustrate our ideas. Now we consider a walk satisfies:

$$x_1 = x_2 + 1 = x_3 + 2 \wedge x_3 > x_4 + 0.26$$

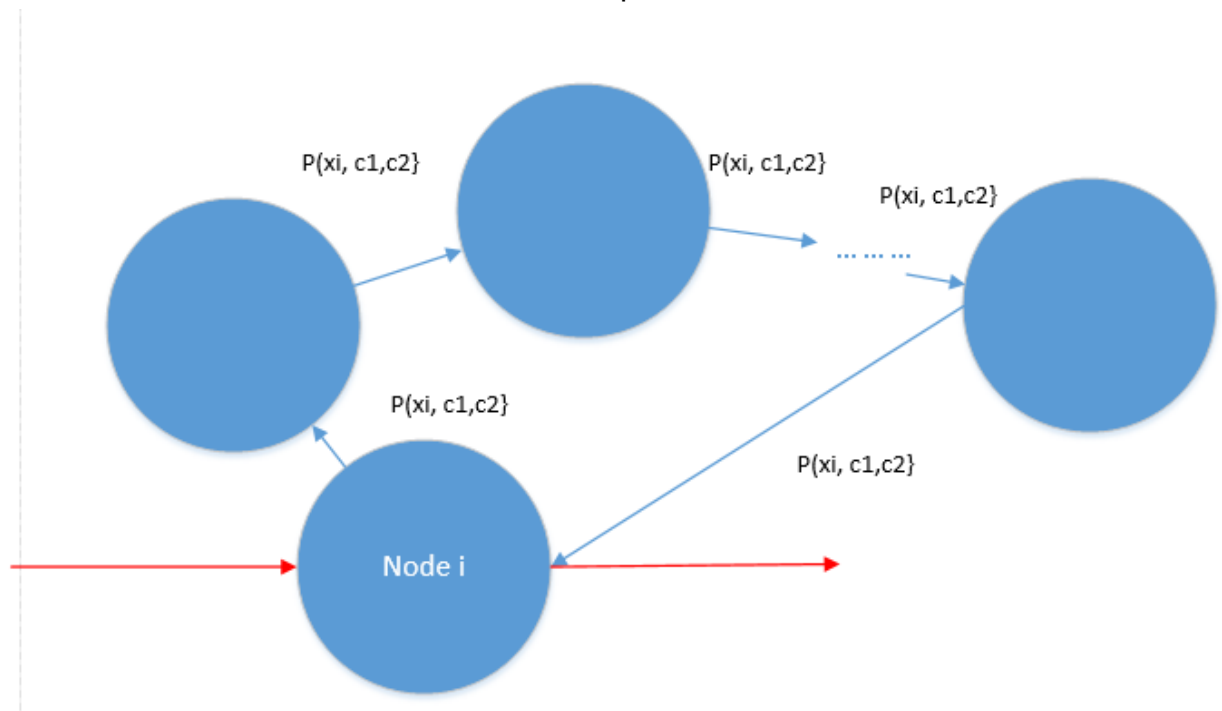
as a bad walk:

Worth to mention that this approach also applies when the bad walk definition is changed and our approach is a generalized one to solve the problem.

Then for each walk, we check all the edges related with T1, T2, T3, T4, and check whether the below linear programming constraints satisfy or not:

$$\begin{aligned} \sum_{i=0}^n x_1 &= 1 + \sum_{i=0}^n x_2 \\ \sum_{i=0}^n x_1 &= \sum_{i=0}^n x_3 + 2 \\ \sum_{i=0}^n x_3 &> 0.26 + \sum_{i=0}^n x_4 \end{aligned}$$

- a) If satisfies, the corresponding walk is a bad walk, the algorithm will decide yes and print out this walk
 - b) Else, the algorithm will say there is no bad walk in M and decide no.
- (2) The directed graph with several nodes that are included in one loop. To illustrate the idea, we need to define one loop first.



At this time, we can consider all other nodes and edges in this loop as One big self-circle edge on the node, and all P instructions can be summed into one instruction $P'\{xi, c'1, c'2\}$.

The algorithm Algorithm-2 now is based on Algorithm-1 but is more robust:

- 1) We use DFS to traverse the graph, and from the initial node to the Final node we will get one MAIN PATH(as red edges shown in the above figure) as a simple path M. At the same time we record loop along with this M if any. Continue the same procedure, in the end, we will get a set of M $\{M1, M2, M3, \dots, Mk\}$ with corresponding loops if exist.
- 2) Use Algorithm-1 to search bad walks along with the MAIN PATH of M_i , ($i=1, 2, \dots, k$). If Algorithm-1 decides yes, that means a bad walk existing without considering loop, Algorithm-2 stops, and decides yes; else, go to step 3).

3) we need to check:

for each walk M_i , check if there is one node in the MAIN PATH M_i that is also contained in a loop? Because there's only one loop that contains the node which lies in the MAIN PATH, we can use SCC algorithm to detect such loop. Then in one MAIN PATH M_i , there may contain several such one loops such as P_1' , P_2' , P_3' ... $P_{K'}$.

If no, there is no bad walk in M_i ;

Else, we need to check the new linear programming constraints satisfy or not of M_i :

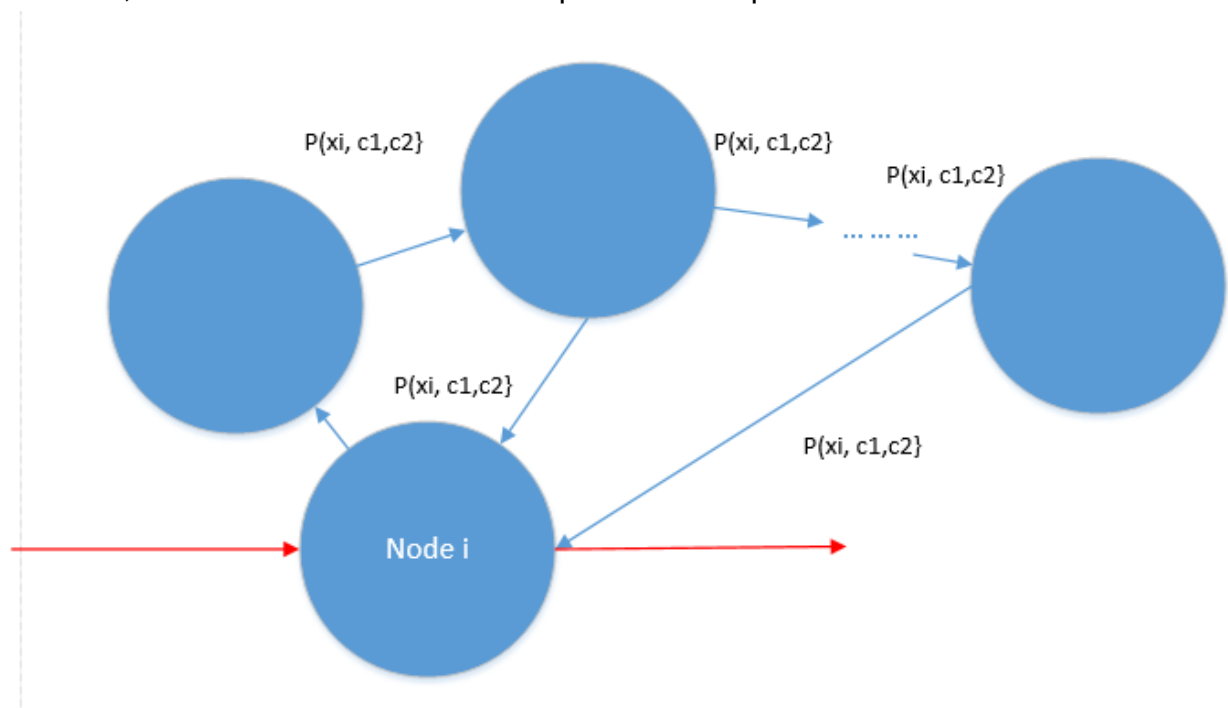
$$\sum_{i=0}^n x_1 \text{ in MAINPATH} + t_1 * (x_1 \text{ in } P_1') + \dots + t_k * (x_1 \text{ in } P_{k'}) = 1 + \sum_{i=0}^n x_2 \text{ in MAINPATH} + t_1 * (x_2 \text{ in } P_1') + \dots + t_k * (x_2 \text{ in } P_{k'})$$

$$\sum_{i=0}^n x_1 \text{ in MAINPATH} + t_1 * (x_1 \text{ in } P_1') + \dots + t_k * (x_1 \text{ in } P_{k'}) = \sum_{i=0}^n (x_3 \text{ in MAINPATH}) + 2 + t_1 * (x_3 \text{ in } P_1') + \dots + t_k * (x_3 \text{ in } P_{k'})$$

$$\sum_{i=0}^n x_3 \text{ in MAINPATH} + t_1 * (x_3 \text{ in } P_1') + \dots + t_k * (x_3 \text{ in } P_1') > 0.26 + \sum_{i=0}^n x_4 \text{ in MAINPATH} + t_1 * (x_4 \text{ in } P_1') + \dots + t_k * (x_4 \text{ in } P_1')$$

t_k is the one loop k execution numbers, When the linear programming holds, there is a bad walk in M_i . After checking M_i ($i=1,2,3,\dots,k$), if all M_i don't satisfy the linear programming constraints, the Algorithm-2 decides no.

(3).The directed graph with several nodes that are included in nested loop. To illustrate the idea, here we use a two-nested loop as an example:



this time we use a two-level hashset to make sure that in the second level, the set DFS traversed is unique and every edge is traversed only once. In this way, we will get small cycles respectively.

The algorithm Algorithm-3 now is based on Algorithm-2 but is more robust:

The only difference is we need to detect small cycles in the nested loops and store the information into a data structure. After applying SCC algorithm to allocate the large cycle in the above figure, this time we use a two-level hashset to make sure that in the second level, the set DFS traversed is unique and every edge is traversed only once. In this way, we will get two loops – Loop P, Loop Q separately. Similarly we can generalize this if one MAIN PATH contains k such nested loops, the linear programming constraints will become:

$$\begin{aligned} \sum_{i=0}^n x1 + \sum_{k=1}^k x1 \ tk + \sum_{k=1}^k x1 \ mk &= 1 + \sum_{i=0}^n x2 + \sum_{k=1}^k x2 \ tk + \sum_{k=1}^k x2 \ mk \\ \sum_{i=0}^n x1 + \sum_{k=1}^k x1 \ tk + \sum_{k=1}^k x1 \ mk &= 2 + \sum_{i=0}^n x3 + \sum_{k=1}^k x3 \ tk + \sum_{k=1}^k x3 \ mk \\ \sum_{i=0}^n x3 + \sum_{k=1}^k x3 \ tk + \sum_{k=1}^k x3 \ mk &> 0.26 + \sum_{i=0}^n x4 + \sum_{k=1}^k x4 \ tk + \sum_{k=1}^k x4 \ mk \end{aligned}$$

tk is the one small loop k execution numbers, mk is the other small loop k execution numbers.

This solution can be easily generalized to n-nested loops

When the linear programming holds, there is a bad walk in Mi, the algorithm decides Yes. After checking Mi (i=1,2,3,...,k), if all Mi don't satisfy the linear programming constraints, the Algorithm-3 decides no.

2. The word bit comes from Shannon's work in measuring the randomness in a fair coin. However, such randomness measurement requires a probability distribution of the random variable in consideration. Suppose that a kid tosses a dice for 1000 times and hence he obtains a sequence of 1000 outcomes

$$a1, a2, \dots, a1000$$

where each a_i is one of the six possible outcomes. Notice that a dice may not be fair at all; i.e., the probability of each outcome is not necessarily $\frac{1}{6}$. Based on the sequence

only, can you design an algorithm to decide how “unfair” the dice that the kid tosses is.

Answer:

The idea to solve this problem is to use the variance to represent the “unfairness”

According to the question, the roll of the dice is not a fair event, which means there is no guarantee that each side of the dice will have the same chance of appearing in 1/6 in 1000 tests. We can use the frequency representation:

$$\mu = \frac{\sum_{i=1}^{1000} a_i}{1000};$$

$$\sigma^2 = \frac{\sum_{i=1}^{1000} (a_i - \mu)^2}{1000}$$

Then in the “fair” case, the outcomes of 1000 tests should be evenly distributed over six possibilities

$$\bar{\mu} = \frac{1 \times \frac{1000}{6} + 2 \times \frac{1000}{6} + 3 \times \frac{1000}{6} + 4 \times \frac{1000}{6} + 5 \times \frac{1000}{6} + 6 \times \frac{1000}{6}}{1000};$$

$$\bar{\sigma}^2 = \frac{\sum_{i=1}^{1000} (a_i - \bar{\mu})^2}{1000}$$

We can use the variance difference ratio to represent the “unfairness” in the end:

$$\theta = \frac{|\sigma^2 - \bar{\sigma}^2|}{\sigma^2}$$

The greater the value of θ , the greater the degree of “unfairness”.

In the algorithm, we can precompute $\bar{\mu}$ and use $a_1, a_2, \dots, a_{1000}$ as input to get the $\sigma^2, \mu, \bar{\sigma}^2$. In the end, we output θ .

However, later I noticed Prof. Zhe Dang gives us the hint to use Kullback-Leibler Divergence to compare two probability distributions.

The ideal distribution of the rolling dice event is a uniform distribution.

And of course we can get our observed distribution of the rolling dice count.

$$H = -\sum_{i=1}^N p(x_i) \cdot \log p(x_i)$$

The information gain simply gives the theoretical lower bound on the number of bits we

need, we have a way to quantify exactly how much information is in our data. Now that we can quantify this, we want to quantify how much information is lost when we substitute our observed distribution for a parameterized approximation.

$$D_{KL}(p||q) = \sum_{i=1}^N p(x_i) \cdot (\log p(x_i) - \log q(x_i))$$

Essentially, what we're looking at with the KL divergence is the expectation of the log difference between the probability of data in the observed distribution with the approximating distribution (uniform distribution in our case). Note that the KL Divergence is not symmetric, here we use:

$$D_{kl}(\text{Observed} || \text{Uniform})$$

to decide how “unfair” the dice that the kid tosses is

3. In below, a sequence is a sequence of event symbols where each symbol is drawn from a known finite alphabet. For a sequence $\alpha = a_1 \cdots a_k$ that is drawn from a known finite set S of sequences, one may think it as a sequence of random variables $x_1 \cdots x_k$ taking values $x_i = a_i$, for each i. We assume that the lengths of the sequences in the set S are the same, say n. In mathematics, the sequence of random variables is called a stochastic process and the process may not be i. i. d at all (independent and identical distribution). Design an algorithm that takes input S and outputs the likelihood on the process being i. i. d.

Answer:

Input: a set S, which contains a finite sequences, and these sequences are with the same length n.

Goal: Output the sequence $\alpha = a_1 \cdots a_k$ drawn from the S and output the likelihood on the process being i.i.d.

According to the question, we start formulating the problem that we have a set of sequences in $S = \{\beta_1, \beta_2, \dots, \beta_m\}$, each β_i is with the same length n. ($i=1,2,3,\dots,m$).

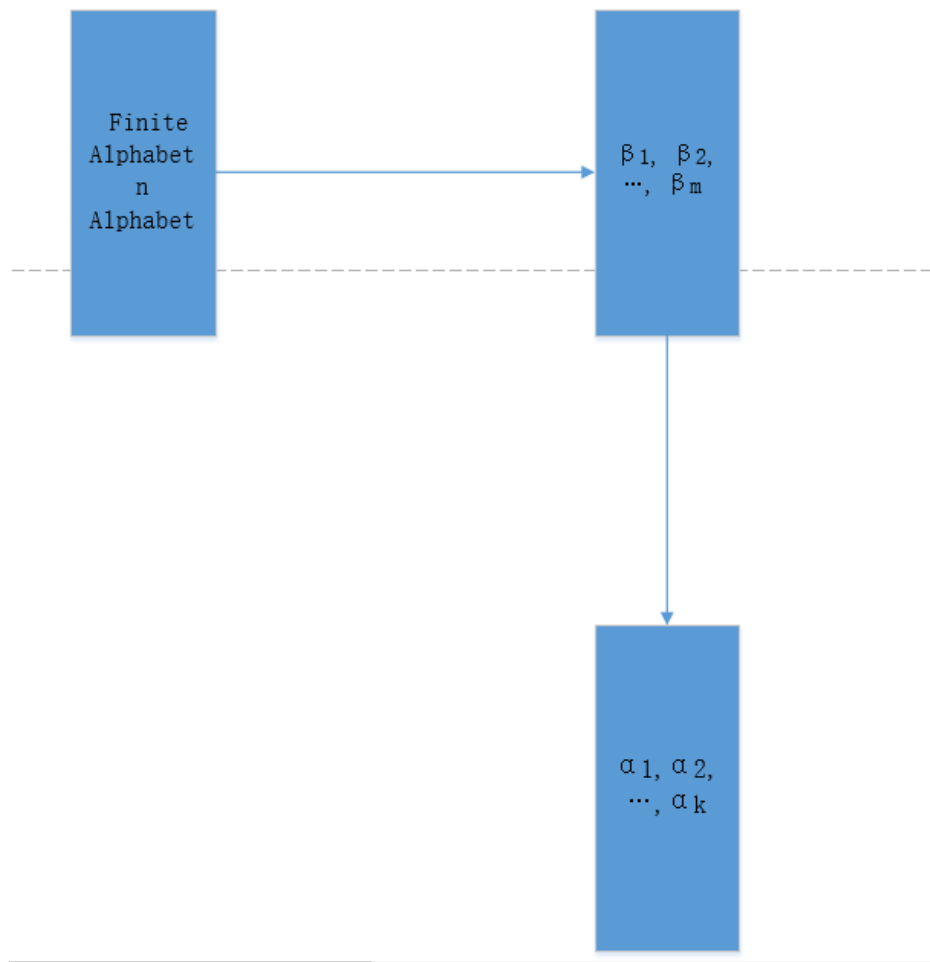
Then to get a sequence $\alpha = a_1 \cdots a_k$ that is drawn from a known finite set S of sequences, from my best understanding, this question could mean both scenarios:

Scenario 1: choose a_i uniformly from β_i , ($i=1,2,\dots,k$)

Scenario 2: choose a_i uniformly from all β_j 's i th element, ($j=1,2,\dots,m$) ($i=1,2,\dots,k$)

I personally tend to select scenario 1 to discuss since in this question we don't have value m in S to discuss. If we choose scenario 2, we have one more parameter that is unknown to consider. However, the analysis is pretty similar for both scenarios.

Because the event symbols in set S are dependent events, we need to show that the likelihood a_i is independently drawn and all a_i obey some kind of identical distribution.



In scenario 1, for each β_i , the n elements is uniformly drawn from the Finite Alphabet, the probability is $\frac{1}{N}$. If we choose a_i uniformly from β_i , then each a_i has the probability to be chosen as

$\frac{1}{n}$. To prove the independence, we need to show that for any $i \neq j$, $a_i \in \alpha$, $a_j \in \alpha$, the

conditional probability equals to the probability. We can certainly compute the $P(a_i = M)$ and $P(a_i=M|a_j=1,\dots,N)$ separately.

If we have $P(a_i=M|a_j=1,\dots,N) = P(a_i=M)$, then any a_i and a_j in α are independent.

Otherwise, they don't satisfy independence property. Similarly, in scenario 2, we still can compute the $P(a_i = M)$ and $P(a_i=M|a_j=1,\dots,N)$ separately, if $P(a_i=M|a_j=1,\dots,N) = P(a_i=M)$, any a_i and a_j in α are independent. Otherwise, they are not independent.

For all a_i satisfied the $P(a_i | a_j)=P(a_i)$ talked above, we add 1 to count. The independent probability of process is count/k .

To prove the identical distribution, we need to collect $\alpha = a_1 \cdot \dots \cdot a_k$ first. Then we can select a bunch of known distribution simulators to fit these data points. We have a total combination of choices N regarding $a_1 \cdot \dots \cdot a_k$ selection since the alphabet is finite. Then for particular α , we can sequentially put α into distribution analyzers and see whether they obey this distribution or not. Then for each α we can get a satisfied count 1 if α obeys one kind of distribution or 0 if α doesn't obey any distribution. Since α is a finite set, in the end we will get

$$\sum_{i=1} 1\{\alpha \text{ obeys certain distribution}\} = M.$$

In the end, the likelihood of identical distribution is $\frac{M}{N}$.

The total likelihood is $\frac{\text{count}}{k} \times \frac{M}{N}$

Another measurement inspired by paper -- *Lossiness of communication channels modeled by transducers*, we can try to compute the information rate of event symbols in S . Here we use $L(S_i)$ denote the number of event symbols in a sequence β_i ($i=1,2,\dots,m$). The information rate $\gamma(S_i) = \lim_{n \rightarrow \infty} \frac{\log L(S_i)}{n}$, since alphabet is finite, we always can have an upper bound of this $\gamma(S_i)$.

Then the total likelihood in this measurement representation is $\frac{\text{count}}{k} \times \frac{1}{m} \sum_{i=1}^m \gamma(S_i)$

4. Let G_1 and G_2 be two directed graphs and v_1, u_1 be two nodes in G_1 and v_2, u_2 be two nodes in G_2 . Suppose that from v_1 to u_1 , there are infinitely many paths in G_1 and that from v_2 to u_2 , there are infinitely many paths in G_2 as well. Design an algorithm

deciding that the number of paths from v_1 to u_1 in G_1 is “more than” the number of paths from v_2 to u_2 in G_2 , even though both numbers are infinite (but countable).

Answer:

We can use adjacency matrix to represent G_1 and G_2 . We first locate v_1 and u_1 in adjacency matrix to prune this matrix and only reserves related information to get A_1 .

Similarly, we can get A_2 .

A_1 is diagonalizable if there is a basis v_1, v_2, \dots, v_n such that $D[A_1]$ - is diagonal. $\{v_1, v_2, \dots, v_n\}$ is a basis of Eigenvectors. Similarly for A_2 .

There is a positive real number r , called the Perron–Frobenius eigenvalue (Perron root), such that r is an eigenvalue of A_1 and any other eigenvalue λ (possibly, complex) is strictly smaller than r in absolute value, $|\lambda| < r$. Thus, the spectral radius $\rho(A_1)$ is equal to r .

Similarly, we have $\rho(A_2) = r_2$.

In the paper “AN ALGORITHM FOR COMPUTING THE PERRON ROOT OF A NONNEGATIVE IRREDUCIBLE MATRIX” by PRAKASH CHANCHANA gives the algorithm to find the Perron–Frobenius eigenvalue.

The algorithm detail can be found in the below picture:

1. Compute the LU factorization of

$$(\lambda^{(i)}I - A) = L^{(i)}U^{(i)},$$

and solve for $\tilde{x}^{(i)}$

$$L^{(i)}U^{(i)}\tilde{x}^{(i)} = x^{(i)}$$

2. Use the same LU factors solve for $x^{(i+1)}$

$$L^{(i)}U^{(i)}x^{(i+1)} = \tilde{x}^{(i)}.$$

3. Compute

$$\bar{\lambda}^{(i)} = \lambda^{(i)} - \min_j \left(\frac{\tilde{x}_j^{(i)}}{(B^{(i)}\tilde{x}^{(i)})_j} \right)$$

and

$$\underline{\lambda}^{(i)} = \lambda^{(i)} - \max_j \left(\frac{\tilde{x}_j^{(i)}}{(B^{(i)}\tilde{x}^{(i)})_j} \right)$$

where $1 \leq j \leq n$; Note that the quantity $(B^{(i)}\tilde{x}^{(i)})_j$ is $x_j^{(i+1)}$.

4. Set

$$\lambda^{(i+1)} = \bar{\lambda}^{(i)}.$$

5. Compute

$$error^{(i)} = (\bar{\lambda}^{(i)} - \underline{\lambda}^{(i)})/\bar{\lambda}^{(i)}.$$

6. If $error > tol$ go back to step 1; otherwise, the Perron root of A is $\lambda^{(i+1)}$.

Then we can compare $p(A1)$ and $p(A2)$ to decide whether that the number of paths from $v1$ to $u1$ in $G1$ is "more than" the number of paths from $v2$ to $u2$