Design & Analysis of Algorithms

Kruskals algorithm Investigation

Axel Solano

May 6, 2017

Contents

1	Intr	oducti	on	2
2	Analysis of Kruskals algorithm			
	2.1	Theore	ethical Analysis	3
		2.1.1	Heap	3
			Hash	
		2.1.3	Return of the Spanning Tree	4
	2.2	Empir	ical Analysis	5
		2.2.1	Hash	5
		2.2.2	Return of the Spanning Tree	5
	2.3	Summ	ary	12

1 Introduction

In this program, I am going to use the scientific method to demonstrate that Kruskals algorithm runs in $O(m \log(n))$ time, in which m in the number of edges and n is the number of vertices in an undirected graph.

2 Analysis of Kruskals algorithm

The algorithm is going to be divide in three parts. The first part is the construction of the heap with the list of edges. The second part is the construction of a Hash that will contain the id of the vertex as an identifier and each of them will contain a cluster of the respective vertex. The last part consists of building the spanning tree.

```
1
   def mstKruskal(graph)
2
       tree = []
3
       # List of edges to return as spanning tree
4
5
       @h = MinHeap.new
       # Priority Queque that will use the weight of
6
7
       #edges as key
8
9
       mapp = Hash.new
10
       #Hash that will save the cluster for every vertices
11
12
       cluster = Cluster.new
       #Cluster is a data structure for the creation,
13
       #find and union of clusters in vertices
14
15
16
17
       #The first part is creating the heap
       lista = graph.list_edges
18
       @h.buildHeap(lista)
19
20
21
22
       #The second part is saving every cluster for
23
       #each vertex in the Hash
       listvertices = graph.list_vertices
24
25
       listvertices.each do |v|
         mapp[v.getId] = cluster.makeSet(v)
26
27
       end
28
29
       #The third part is to return the spanning tree
       i = 1
30
31
       n = @h.getSize
32
       while ( tree.length!=lista.length-1 &&
         i = i + 1
33
         edge = @h.removeMin(lista)
34
         v1 = edge.getEndVertices.fetch(0)
35
         v2 = edge.getEndVertices.fetch(1)
36
37
         a = cluster.findParent(mapp[v1.getId])
         b = cluster.findParent(mapp[v2.getId])
38
         if a != b
39
40
           tree.push(edge)
           cluster.union(a,b)
41
42
         end
43
       end
44
45
       return tree
46
     end
```

2.1 Theorethical Analysis

2.1.1 Heap

In this part, I used the same implementation from the last assignment but I modify from a max-Heap to min-Heap. The function buildHeap() which construct the heap maintain the same time complexity as O(n) and the function removeMin() also maintain the time complexity of $O(\log(n))$.

buildHeap: This function check from the bottom of the tree to the top and the number of operations in each node is equivalent to the distance between the current level of the tree and the last level. The last level of the tree is founded on the nodes after the division by two of the total number nodes of the tree. In this theoretical analysis, I assume that the operations of each node are in the worst case scenario.

$$\sum_{i=1}^{h} 2^{h-i}i$$

Reducing the expression

$$\sum_{i=1}^{h} 2^{h-i}i = 2^{h} \sum_{i=1}^{h} \frac{i}{2^{i}}$$

$$\sum_{i=1}^{h} \frac{i}{2^{i}} = 1 + \sum_{j=1}^{h-1} \frac{1}{2^{j}} - \frac{h}{2^{h+1}}$$

$$1 + \sum_{j=1}^{h-1} \frac{1}{2^{j}} - 2^{\frac{h}{2^{h+1}}} = 1 + 1 - \frac{1}{2^{h-1}} - \frac{h}{2^{h+1}}$$

$$2^{h} \sum_{i=1}^{h} \frac{i}{2^{i}} = 2^{h} [2 - \frac{1}{2^{h-1}} - \frac{h}{2^{h+1}}]$$

$$\sum_{i=1}^{h} 2^{h-i}i = 2^{h} \sum_{i=1}^{h} \frac{i}{2^{i}} = 2^{h+1} - 2 - \frac{h}{2}$$

Then, make the $h = \log(n/2)$ where n is equal to the number of nodes. Therefore, the build function has the equation of:

$$T(n) = n - \frac{\log(n)}{2} - 3$$

This equation can be reduced in Big-Oh notation as O(m) in which m is the number of edges.

removeMin: This function removes the minimum element in O(1) and then re-arrange the heap for next removal. The re-arragement time takes $O(\log(n))$ because in each iteration the modification and those operations are equivalent to the level of the tree. Also, it is possible to find a relation between the level and the number of nodes of the tree.

$$1 + 2 + 4 + + 2^{h-1} = 2^{h}1$$

$$n \le 2^{h}1$$

$$n + 1 \le 2^{h}$$

Then, I take logarithms of both sides of these two inequalities

$$log(n+1) \le h$$

Finally, we can generalize

$$h = \lceil log(n+1) \rceil$$

$$T(n) = log(n+1) = log(n)$$

This equation can be reduced in Big-Oh notation as $O(\log(m))$ in which m is the number of edges.

2.1.2 Hash

In this part, the theoretical analysis is beyond the scope of the course but based on the book, this data structure present search function in time complexity O(1) in average case and in worst case scenario is in O(n) in which n is the length of the hash. In addition, I will evaluate the time complexity of the search function empirically in the next section to proof that it doesn't have much influence in the performance of the kruskal's algorithm. Also, implementing the cluster with th map will cause a time complexity of O(n) in which n is the number of vertices.

2.1.3 Return of the Spanning Tree

In this part, there is a relevant implementation which is the Cluster class. This class use a Set class in which store the store the element and has linked-connection to itself with the name of parent. The purpose of these classes is to create the cluster of the spanning tree at the same time is creating the spanning tree. On the book, there is an implementation called for Union-Find Structures called Tree-based implementation in which the idea behind is to perform search and insertion in $O(\log(n))$ but with a path compression, this implementation can become faster but it's difficult to measure. The function is represented as $\alpha(n)$ which less than or equal to $\log(n)$ and is called the Ackermann function.

In addition, the time complexity of the loop is represented as O(mlog(m)) in which m is the number of edges. However, we can change the m inside the logarithm function as function of n which is the number of vertices. The total number of edges possible in graph based on number of vertices is:

$$\frac{n*(n-1)}{2} + n = m$$

The first part which is $\frac{n*(n-1)}{2}$ represent the total number of distinct edges that can be formed and the n represent the loops edges of each vertex. Therefore, I am going to show the steps to obtain O(mlog(n)).

$$T(n) = m * log(\frac{n*(n-1)}{2} + n)$$

Elimate the n because loops edges is not allowed to create spanning trees.

$$T(n) = m * log(\frac{n * (n-1)}{2})$$

$$T(n) = m * log(\frac{n*(n-1)}{2})$$

$$T(n) = m*log(n*(n-1)) - m*log(2)$$

$$T(n) = m * log(n) + m * log(n-1) - m$$

This equation can be reduced in Big-Oh notation as O(mlog(n)).

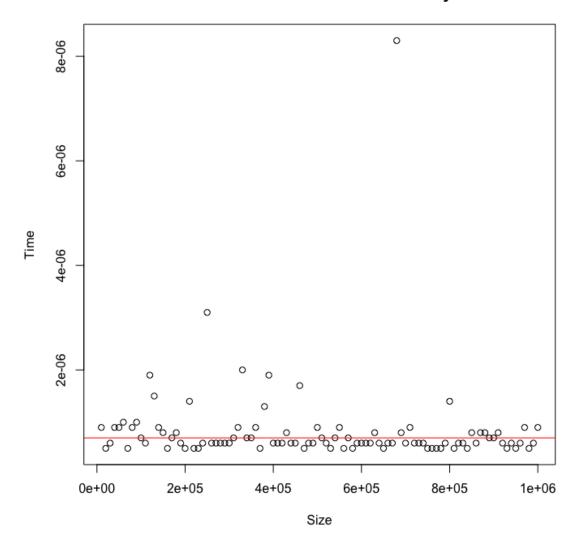
Finally, the summation of the previous time complexities in Big-Oh notation of the used functions is $O(m) + O(n) + O(m\log(n)) = O(m\log(n))$.

2.2 Empirical Analysis

2.2.1 Hash

Based on the empirical analysis, the hash follow a constant uniform distribution which makes our assumption of constant time complexity correctly.

Time measurement of Hash in Ruby



2.2.2 Return of the Spanning Tree

2.2.3 Testing the algorithm:

In this analysis, I test my knapsack function with the size from 10000 to 1000000 and for every case of the size I run 10 times, and then make an average of the time. In addition, I am going to use statistical analysis using the software R.Also, I create a variable called sum which represent the summation of each weight of the item; I create this variable to use in my test as the input of the maximum weight. In the test, I put the maximum weight as the total weight, therefore, I am

indirectly sorting the items and testing the same time the worst case scenario. I did the previous implementation to prove that my function runs in O(nlogn).

Fitting the points using linear regression

$$TIME = (-1.614e + 01) + (2.350e - 05) * EDGES$$

$$p - value :< 2.2e - 16$$

$$RSE : 13.29$$

$$R^2 : 0.9944$$

$$F - statistic : 1.197e + 04$$

Polynomial function of degree 1

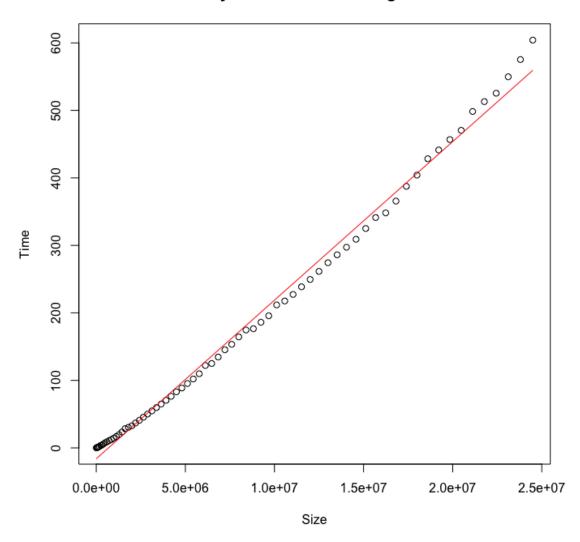


Figure 1: Linear Regression

Fitting the points using Polynomial regression of grade 2

$$TIME = (-1.603) + (1.772e - 05) * EDGES + (2.705e - 13) * EDGES^2$$

$$p - value :< 2.2e - 16$$

$$RSE : 3.297$$

$$R^2 : 0.9997$$

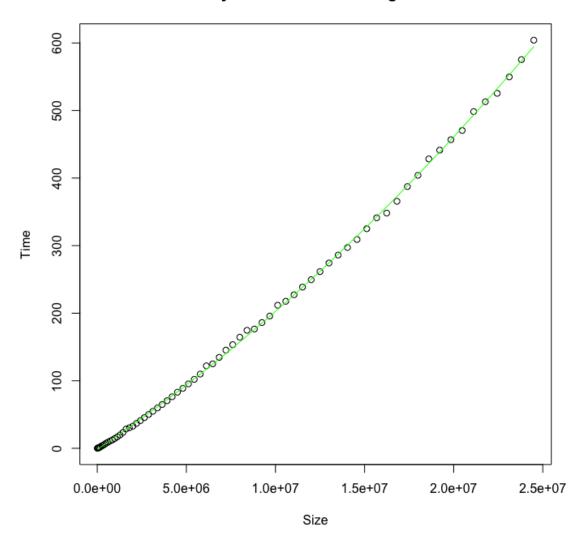


Figure 2: Polynomial regression of grade 2

Fitting the points using Polynomial regression of grade 3

 $TIME = (-2.749) + (1.867e - 05) * EDGES + (1.561e - 13) * EDGES^2 + (3.370e - 2) * EDGES^3$ p - value : < 2.2e - 16 RSE: 3.162 $R^2: 0.9997$

F-statistic: 7.096e+04

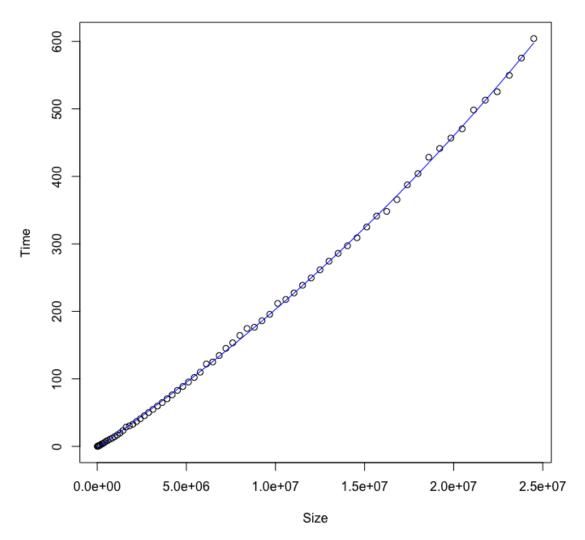


Figure 3: Polynomial regression of grade 3

Fitting the points using Polynomial regression of grade 4

 $TIME = (-1.549e + 00) + (1.698e - 05)*EDGES + (5.299e - 13)*EDGES^2 + (-2.267e - 20)*EDGES^3 + (5.613e - 28)*EDGES^2 + (-2.267e - 20)*EDGES^3 +$

p-value:<2.2e-16

RSE:3.003

 $R^2: 0.9997$

F-statistic: 5.9e+04

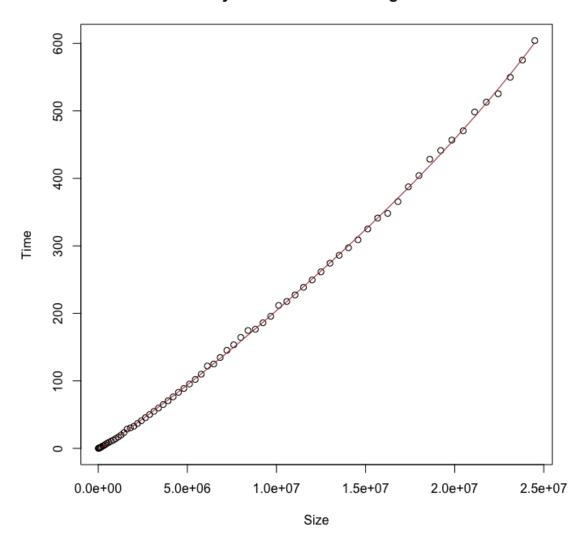


Figure 4: Polynomial regression of grade 4

Fitting the points using Polynomial regression of grade 5

 $TIME = (-2.759e - 01) + (1.424e - 05)*EDGES + (1.480e - 12)*EDGES^2 + (-1.372e - 19)*EDGES^3 + (6.158e - 27)*EDGES^3 + (6.15$

p-value:<2.2e-16

RSE:2.808

 $R^2: 0.9998$

F - statistic : 5.397e + 044

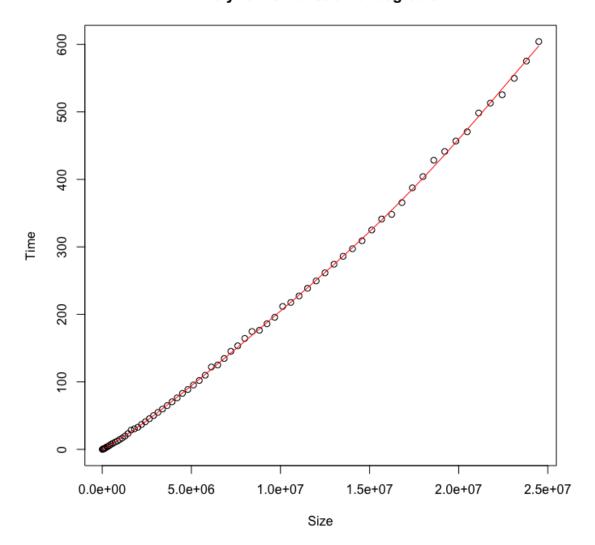


Figure 5: Polynomial regression of grade 5

Fitting the points using n*log(n) curve

TIME = (-9.352) + (9.609e - 07) * EDGES * log(EDGES) p - value :< 2.2e - 16 RSE : 9.608 $R^2 : 0.9971$

F-statistic: 2.299e+04

m*log2(m) function

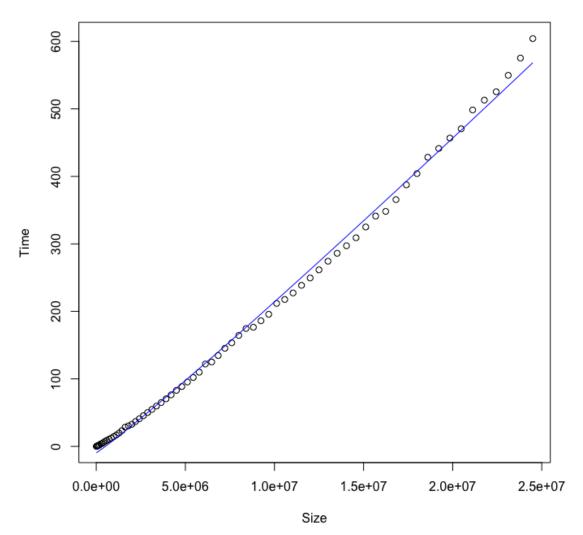


Figure 6: nlogn style

Fitting the points using n*log(n) curve with addition of functions

TIME = (-7.297) + (4.963666e - 07) * (EDGES*log(EDGES) + (-1.170e - 04)EDGES + (2.319e - 02)VERTICES + (2.319e - 02)VERTICES

p-value:<2.2e-16

RSE:4.308

 $R^2: 0.9994$

F-statistic: 3.82e+04

m*log2(m) function type 2

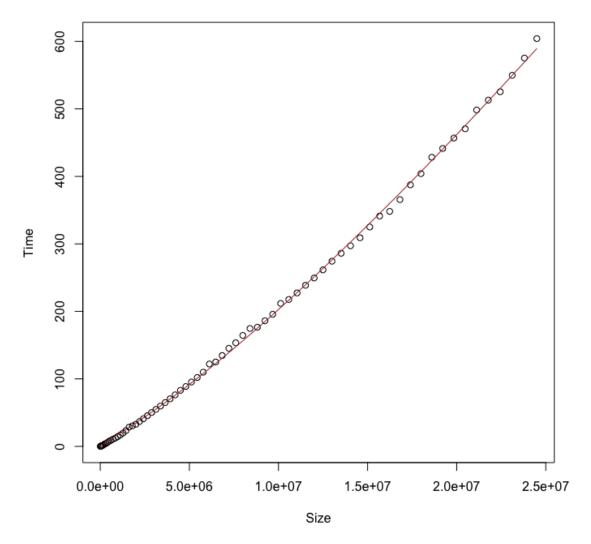


Figure 7: nlogn style

2.3 Summary

In all the cases, the models have presented a R^2 very near to 1 which tell us that is a good model, also the p-value is less than 0.05 which also tella us that the predictors have influence and is good model, and the F-statistic in all the cases show us that models have high influence. The problem is that the model of polynomial regression on degree 2 show the highest F-statistic which tells us the is the best model to predict. Although that model has the best prediction, it does not necessary mean that the last model in which there is $O(m^*log(n))$ is incorrect because this model present all the statistics varibles with a good value and does not indicate as a bad or inaccurate model. In addition, I am going to make an analysis to the coefficients with this R code.

The mean show us a value of 8.458e-07 which is very low and represent as the coefficient to multiplicate to obtain the time. The coefficient in the model 2 is 2.705e-13 which is lower than 8.458e-07 which mean that the function need a very small number to maintain the correct prediction and help us to show that this model is not representing a quadratic growth. Also, the coefficient in the last model is 4.963666e-07 which almost half of the mean and it gives us a hint about how is going to be the growth of the function, therefore coefficient makes sense in our

intuition of choosing the correct model.