Q1- Knapsack problem

The problem involves n objects, b_0 , b_1 ,..., b_{n-1} and a knapsack of capacity C. Each object b_i has a given positive weight or volume w_i and a given positive value v_i (profit, cost, etc) i = 0, ..., n-1. If a fraction f_i of object b_i is placed in the knapsack, then it contributes f_i to the total value of in the knapsack.

The goal is to place objects or fractions of objects in the knapsack without exceeding capacity, so that the total value off the objects in the knapsack is maximized.

Formal Statement:

maximize
$$\sum_{i=0}^{n-1} f_i v_i,$$
subject to the constraints:
$$\sum_{i=0}^{n-1} f_i w_i \le C,$$

$$0 \le f_i \le 1, \quad i = 0, ..., n-1.$$
(6.3.1) p. 248

\

Greedy Strategy: At each iteration, putting objects or fractions of objects into the knapsack according to decreasing rations (densities) v_i/w_i until the knapsack is full yields the optimal solution. Thus, the ratio v_i/w_i is the critical issue. Id.

Exercise 6.5, p. 281

Object Type <i>b_i</i>	Quantity Available <i>w_i</i>	Total Cost for that Quantity <i>V_i</i>
0	30	60
1	100	50
2	10	40
3	10	30
4	8	20
5	8	10
6	1	5
7	1	1

Figure 1.1 - Object *b_i* Inventory

Object Type <i>b_i</i>	Ratio (Density) v _i /w _i	Fraction of Available Quantity Chosen f_i	Quantity Chosen $f_{i}w_{i}$
6	5	1	1
2	4	1	10
3	3	1	10
4	2.5	1	8
0	2	1/30	1
5	1.25	0	0
7	1	0	0
1	0.5	0	0

Figure 1.2 - Solution for capacity C = 30. Objects sorted in decreasing order of ratios v_i/w_i , i = 0,..., n-1

In Figure 1.1, we list the objects and amounts of objects that are available. Figure 1.2 sorts *the* 8 objects, b_0 , b_1 ,..., b_7 in decreasing order of density v_i/w_i , i = 0,..., n-1. The capacity C of the knapsack is 30. Implementing the greedy strategy, we obtain the most valuable (maximized) knapsack (without exceeding capacity) if we first use as much as possible of the object bi whose density is the largest, then as much as possible of the object whose density is second-largest, and so on until we we reach 30.

Here, we place in decreasing order the total quantity available of objects b_6 , b_2 , b_3 , b_4 because they have the highest density, 5, 4, 3, 2.5, respectively. At this point, C = 29, so we have 30-29 = 1 space remaining in the knapsack. The object with the highest density after b_4 is b_2 with a density of 2 and available quantity $w_i = 30$. We only have one space remaining in the knapsack, so the fraction of available quality chosen $f_0 = 1/30$. Since $w_0 = 30$, $f_0*w_0 = 1$. The total value of the knapsack equals (1*5) + (10*4) + (10*3) + (8*2.5) + (1*2) = 5 + 40 + 30 + 20 + 2 = 107, and satisfies capacity constraints (1*1) + (1*10) + (1*10) + (1*8) + [(1/30) *30] = 30.

Q2 - Huffman Codes

The Huffman algorithm is a data compression algorithm based on a greedy strategy of for constructing *optimal binary prefix codes* for a given alphabet of symbols *A*. Binary prefix codes have the property that no binary string in the codes is a prefix of any other string in the code. Optimal binary prefix code are binary prefix codes that minimize the expected length of text generated using the symbols starting from *A*. The problem of finding an optimal binary prefix code is equivalent to finding a 2-tree *T* having minimum *weighted leaf path length WLPT(T)* of *T* defined by,

$$WLPL(T) = \sum_{i=0}^{n-1} \lambda_i f_i.$$
 (6.4.1) p,256

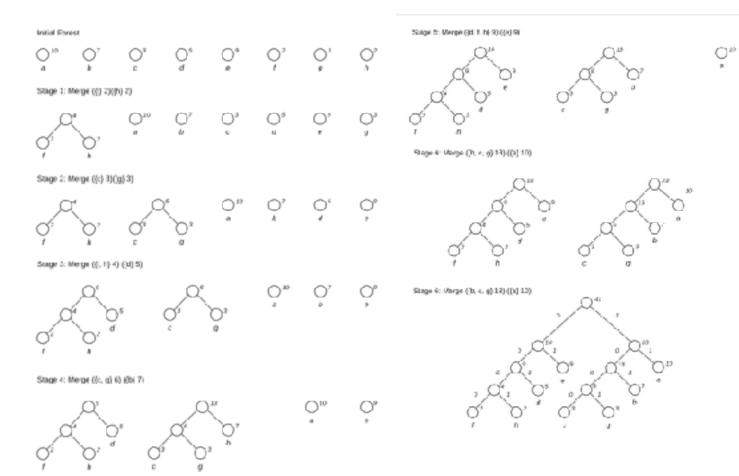
where λ_i denotes the length of the path in T from the root to the leaf node corresponding to ai, i = 0,..., n-1.

The goal of the Huffman Algorithm is to generate optimal binary prefix codes with respect to a given set frequencies f_i , i = 0,...,n-1 for each symbol starting from A, thus minimizing the expected length of a coded symbol. The algorithm computes a 2-Tree T minimizing the WLPL(T) by constructing a sequence of forests. At each iteration, the root of each tree in the forest is assigned a frequency. The initial forest F consists of the n single node trees corresponding to the n elements of A. At each stage iteration, the algorithm makes a greedy choice: it finds two trees T_1 and T_2 whose roots R_1 and R_2 have the smallest and second-smallest frequencies over all tree in the current forest. A new internal node R is then added to the forest, together with two edges joining R to R_1 and R to R_2 , so that a new tree is created with R as the root and T_1 and T_2 as the left and right subtrees of R. The frequency off the new root vertex R is taken to be the sum of the frequencies of the old root vertices R_1 and R_2 . The Huffman tree is constructed after n-1 stages, involving the addition of n-1 internal nodes.

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Symbol	Frequency	Encoding
а	10	11
b	7	101
С	3	1000
d	5	001
е	9	01
f	2	0000
g	3	1001
h	2	0001

WLPL(T) = (2)(10) + (3)(7) + (4)(3) + (3)(5) + (2)(9) + (4)(2) + (4)(3) + (4)(2) = 134

Figure 2 - Action of HuffmanCode for the alphabet $A = \{a,b,c,d,e,f,g,h\}$ with frequencies 10,7,3,5,9,2,3,2, respectively.

Q3 - Minimum Spanning Tree

A spanning tree of *G* is a subset of the edges that connects all the vertices and has no cycles. A minimum spanning tree is a spanning tree that has the lowest possible weight, viz, the sum of the weights of the edges must be as low as possible.

6.14 Consider the following weighted graph G

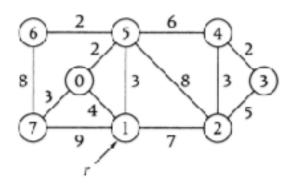


Figure 3.1 - The weighted, connected graph G given in Exercise 6.14, p. 283

a) Trace the action procedure Kruskal for G

Kruskal's assumes that G is connected, and starts with a forest of n isolated trees consisting of a single node. The algorithm uses a greedy strategy to grow a minimum spanning tree by adding to it the lightest edge that connects two trees on each iteration.

The algorithm either sorts all of the edges by weight in advance and process them in order, or uses a min-heap priority queue. Next, the disjoint-set data structure is used to test whether the edge connects two components, rejecting those edges if a cycle is formed. If an edge is found, the *Kruskal's* joins the components using an intermixed sequence of the the disjoint-set ADT's *Union* and *Find* operations.

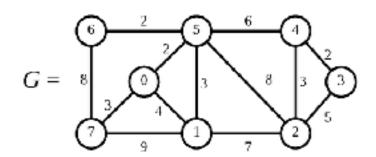
In Figure 3.2, we sort the edges of graph *G* in Figure 3.1 in increasing order of weight or cost. If connecting the two components creates a cycle, the edge is rejected. If a cycle is not formed, the edge is added to the tree. Finally, the weight of the tree is given at each stage or interaction of the process until all vertices are connected in the minimum spanning tree. *Kruskal*'s algorithm is illustrated for the graph in Exercise 6.14 in Figure 3.3.

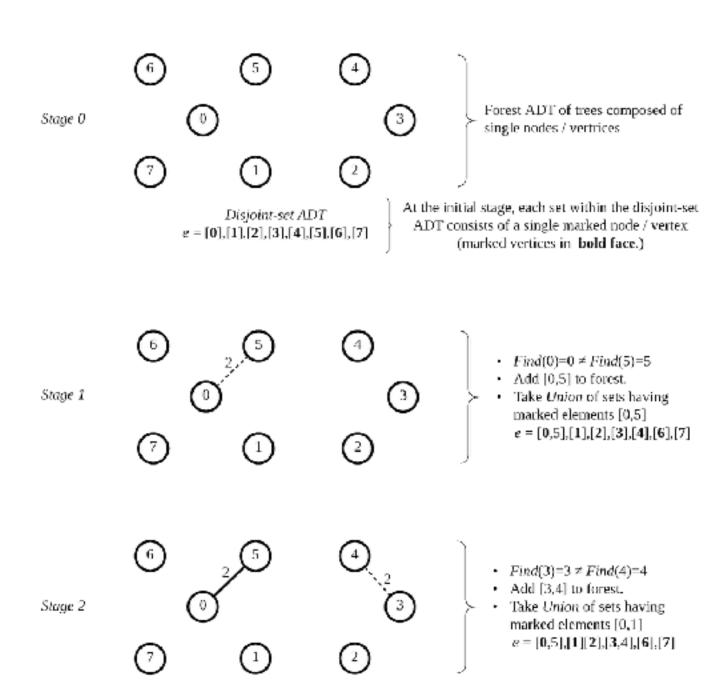
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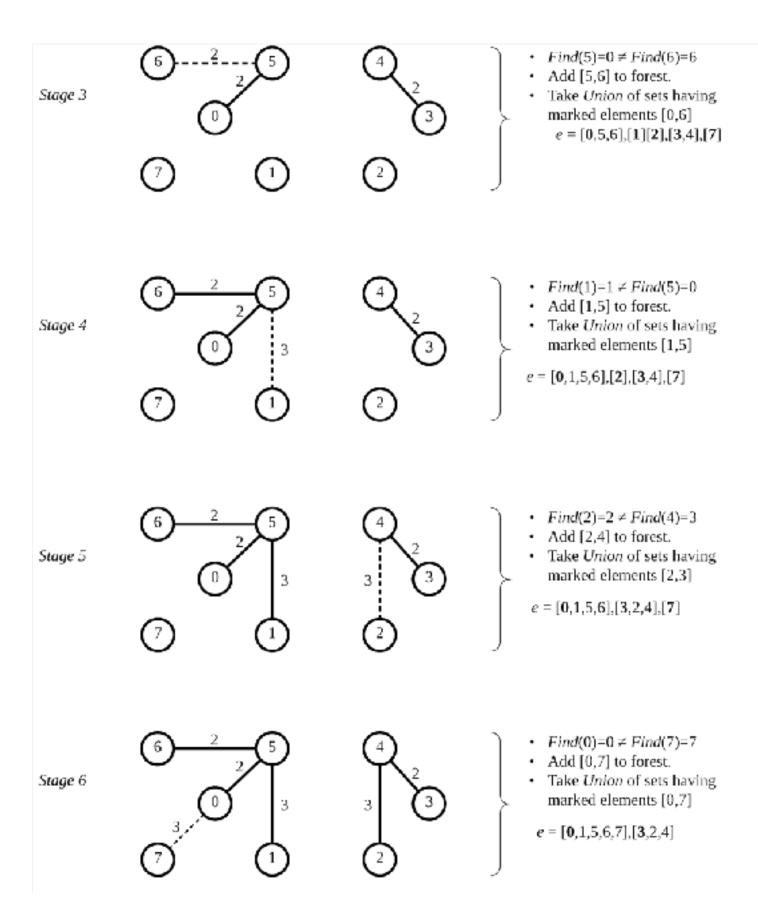
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Stage or Iteration	Edge	Edge Weight	Add or Reject? (cycle formed?)	Tree Weight
Initial	0	0	N/A	0
1	(0,5)	2	Add	2
2	(3,4)	2	Add	4
3	(5,6)	2	Add	6
4	(1,5)	3	Add	9
5	(2,4)	3	Add	12
6	(0,7)	3	Add	15
7	(0,1)	4	Reject	15
8	(2,3)	5	Reject	15
9	(5,4)	6	Add	21

Figure 3.1 - Action of procedure Kruskal for graph G given in Figure 3.1







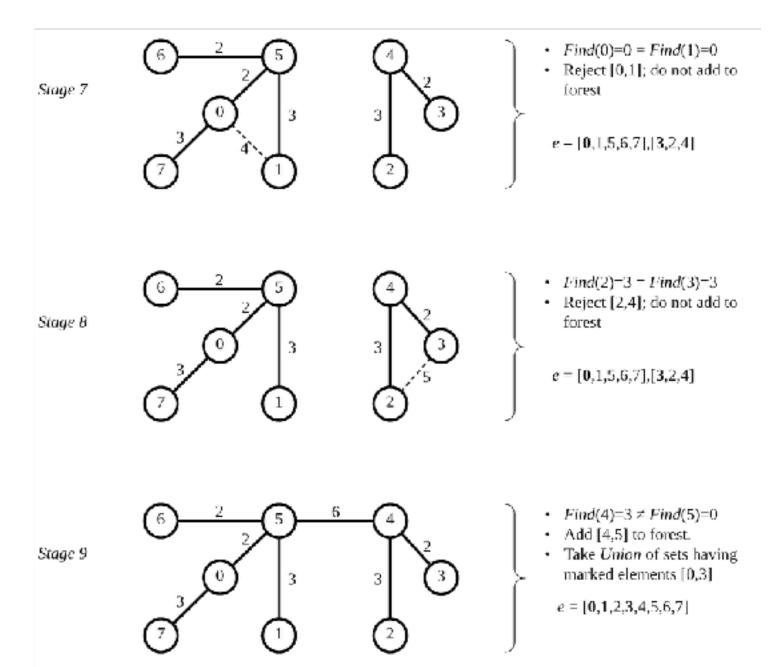


Figure 3.3 - Stages in Kruskal's Algorithm

b) Trace the action of procedure Prim for G with r = 1.

Unlike *Kruskal*'s Algorithm, *Prim*'s Algorithm maintains a tree at each stage instead of a forest. The initial tree T_i may be taken to be any single vertex r of the graph G. At any point, the graph G is composed of a single tree and a bunch of isolated vertices. The algorithm builds a sequence of n trees rooted at r T_0 , T_1 ,..., T_{n-1} , where T_{i+1} is obtained from T_i by adding a single edge e_i+1 , i=0,...,n-2. At every iteration, *Prim*'s makes a greedy choice and adds the smallest possible edge among all edges having exactly one vertex in T_i that connects the tree to an isolated vertex.

The set of all edges in G having exactly one vertex in T_i is denoted by $Cut(T_i)$. At each step, Prim's algorithm greedily chooses the lightest edges among all the edges in $Cut(T_i)$. Thus, we can restrict our attention to the edges in $Cut(T_i)$. Choosing the light edge at each iteration maintains the minimum spanning tree at every step.

In Figure 3.4 and 3.5 we show the action of a straightforward implementation of *Prim's* algorithm for graph G with r = 1 at each iteration. The initial stage shows the current tree, T_0 consisting of the single vertex r = 1 and the set of edges $Cut(T_i)$ available for growing the tree T_i . Subsequent stages also show the edge chosen, individual edge weight and current weight of the tree until all vertices are connected in the minimum spanning tree.

We show the action of procedure for a more efficient implementation of Prim's algorithm based on equation (6.5.1) on pg, 270 in Figure 3.6. Here, we maintain a local array Nearest[i] which keeps track of a vertex u "nearest" to an adjacent vertex v, where the weight of the edge w(u,v) is the smallest of all those in Cut(T), rather than explicitly implementing Cut(T) at each step. The tree T is dynamically maintained using an array Parent[0:n-1]. If vertex v is in tree T, then Parent[v] is the parent of vertex v. If v is not in the tree but adjacent to to at least one vertex of T, then Parent[v] is the vertex nearest to v. Otherwise, Parent[v] is undefined. At each stage, if Parent[v] is defined, then Nearest[v] equals the weight of the edge between v and the vertex u nearest to v.

Stage or Iteration	Edge e _i	Ti	Cut(T _{i)}	Edge Weight	Tree Weight	
0 (r = 1)	Ø	<i>T</i> ₀	[1,0],[1,2],[1,5],[1,7]	0	0	_
1	(1,5)	T_1	[1,0],[1,2],[1,7],[5,0],[5,1],[5,4],[5,6]	3	3	
2	(5,0)	T_2	[0,7],[1,2],[1,7],[5,0],[5,2],[5,4],[5,6]	2	5	
3	(5,6)	<i>T</i> ₃	[0,7],[1,2],[1,7],[5,2],[5,4],[6,7]	2	7	
4	(0,7)	<i>T</i> ₄	[1,2],[5,2],[5,4]	3	10	
5	(5,4)	<i>T</i> ₅	[1,2],[4,2],[4,3],[5,2]	6	16	
6	(4,3)	T_6	[1,2],[4,2],[5,2]	2	18	

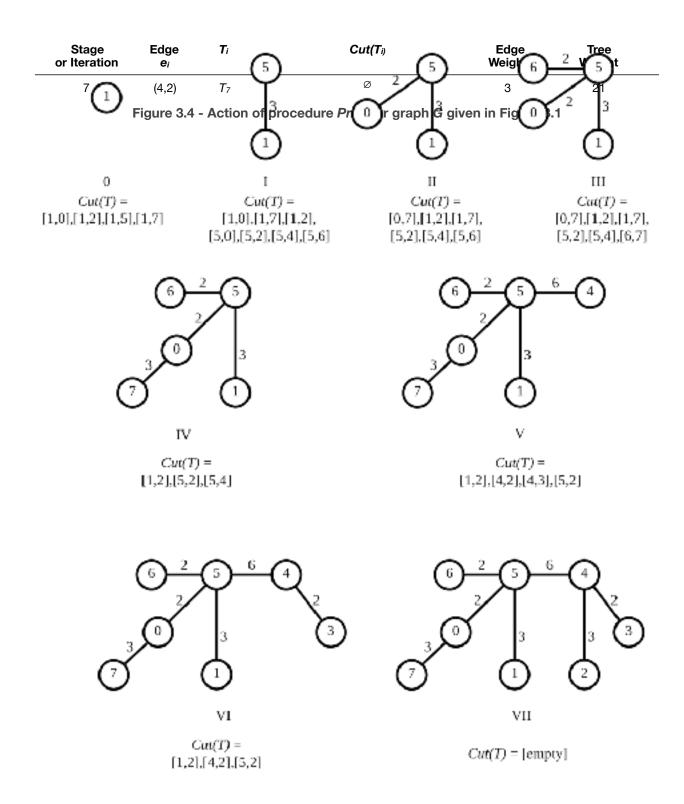
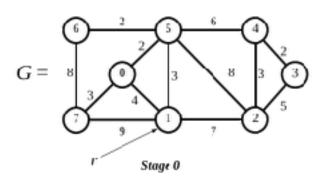
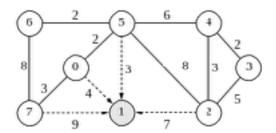


Figure 3.5 - Stages of *Prim's* Algorithm

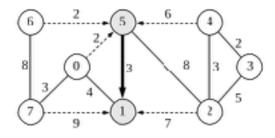




 $i \ 0 \ 1 \ 2 \ 3 \ 4 \ 5 \ 6 \ 7$ $Nearest [i] \ 4 \ 0 \ 7 \ \infty \ \infty \ 3 \ \infty \ 9$ $Parent [i] \ 1 \ -1 \ 1 \ - \ - \ 1 \ - \ 1$ $In The Tree [i] \ F \ \Gamma \ F \ F \ F \ F \ F$

Weight = 0

Stage 1



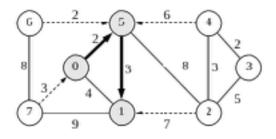
 $I \ 0 \ 1 \ 2 \ 3 \ 4 \ 5 \ 6 \ 7$ Nearest [i] 2 0 8 ∞ 6 3 2 9

Parent [i] 5 ·1 5 - 5 1 5 1

InTheTree [i] F T F F F T F F

Weight = 3

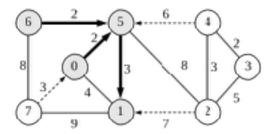
Stage 2



 $i \ 0 \ 1 \ 2 \ 3 \ 4 \ 5 \ 6 \ 7$ $Nearest [i] \ 2 \ 0 \ 8 \ \infty \ 6 \ 3 \ 2 \ 3$ $Parent [i] \ 5 \ -1 \ 5 \ - \ 5 \ 1 \ 5 \ 0$ $In The Tree [i] \ T \ T \ F \ F \ F \ T \ F \ F$

Weight = 5

Stage 3



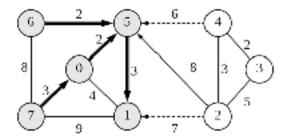
 $i \ 0 \ 1 \ 2 \ 3 \ 4 \ 5 \ 6 \ 7$ Nearest [i] 2 0 7 ∞ 6 3 2 3

Parent [i] 5 -1 1 - 5 1 5 0

In The Tree [i] T T F F F F T T F

Weight = 7

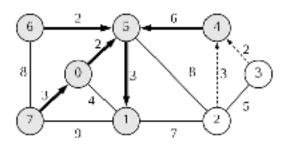
Stage 4



Í	0	1	2	3	4	5	6	7
Neurest [i]	2	0	7	00	6	3	2	3
Parent [i]	5	-1	1	-	5	1	5	0
$In The Tree\ [i]$	T	\mathbf{T}	F	${\rm F}$	F	\mathbf{T}	T	T

Weight = 10

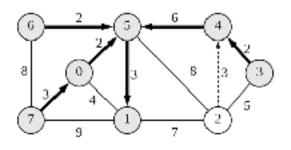
Stage 5



ŕ	D.	1	2	3	4	5	6	7
Nearest [i]	2	0	3	2	6	3	2	3
Parent [i]	5	-1	4	4	5	1	5	0
InTheTree i	Т	Т	F	F	Т	Т	Т	Т

Weight = 16

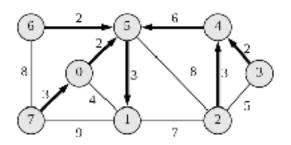
Stage 6



Í	0	1	2	3	4	5	6	7
Nearest [i]	2	0	3	2	6	3	2	3
Parent [i]	5	-1	4	4	5	1	5	0
InTheTree [i]	T	T	F	Т	Т	T	T	Т

Weight = 18

Stage 7



Ī	0	1	2	3	4	5	6	7
Nearest [i]	2	0	3	2	6	3	2	3
Parent [i]	5	-1	4	4	5	1	5	0
InTheTree [i]	Т	Т	Т	Т	Т	Т	Т	Т

Weight = 21

After Stage 6, array *Parent* is now complete, and implements a minimum spanning tree with weight 21.

Figure 3.6 - Action of procedure for *Prim* for graph in Exercise 6.14

Q4 - Shortest Paths

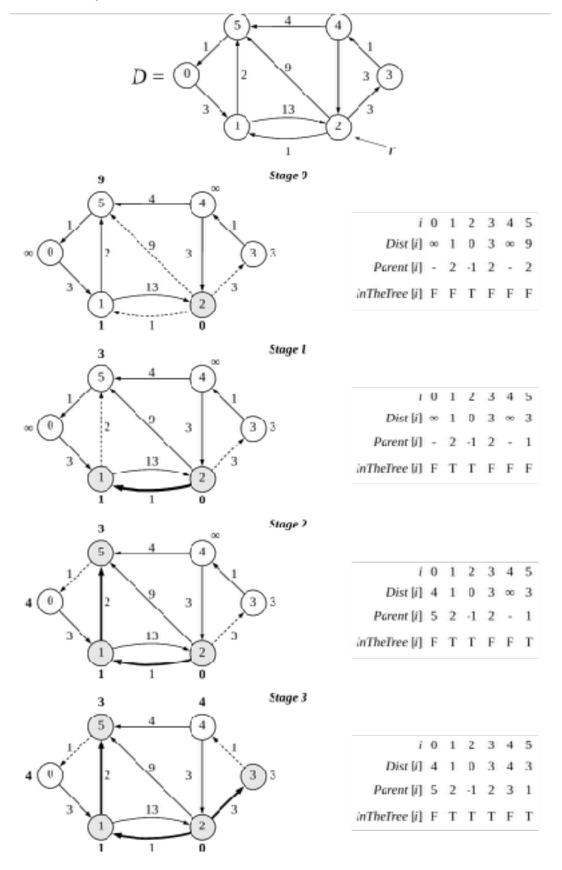
Dijkstra's Algorithm grows a shortest path tree using the greedy method following a similar strategy to Prim's algorithm, differing only in the way the next edge is selected. Like Prim, the tree T is dynamically grown using an array Parent[0:n-1]. A boolean array InTheTree is used to keep track of vertices not in T. At each step, in Dijkstra, instead of selecting an edge of minimum weight in Cut(T), we select an edge uv in Cut(T) where vertex u is in the set of vertices of the tree V(T) such that the path from root r to v through u in the augmented tree T is the shortest path. More precisely, Dist[u] is the weight of the path from r to u in T. We greedily select the edge uv such that Dist[u] + the weight of the edge uv w(uv) is minimized over all edges $uv \in Cut(T)$.

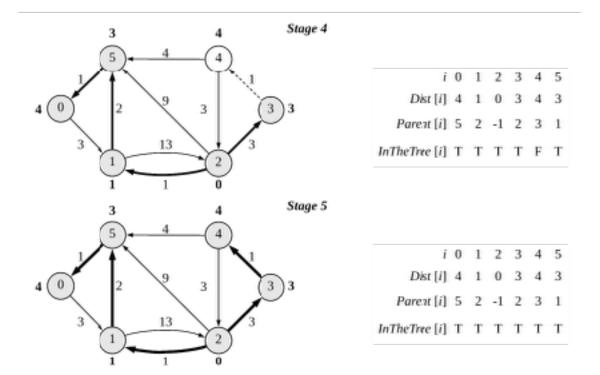
Like Prim, instead of explicitly maintaining the set Cut(Ti), we efficiently select the next edge uv to be added to the tree by maintaining an array Dist[0:n-1], where Dist[v] is the distance from v to r in T and $Dist[v] = \infty$ if v is not in T. Upon completion of Dijsktra, the distance from the root r to all the vertices of G is contained in the array Dist[0:n-1]. Vertices which are candidates for addition to the street T must be : (1) not in the set of vertices in the tree V(T); (2) adjacent to any vertex of T; and (3) Dist[v] is the minimum value of Dist[u] + w(uv) over all edges $uv \in Cut(T)$. Thus, each time a vertex v is added to the tree T, we need only update Parent[v] and Dist[v] for those vertices v that are adjacent to u and do not belong to tree T. If Dist[v] > Dist[u] + w(uv), then we set Parent[v] = u and Dist[v] = Dist[u] + w(uv).

For digraphs, uv corresponds to directed edge uv = (u,v) from u to v (as opposed to the undirected edge $uv = \{u,v\}$. Therefore, for digraphs Prim greedily selects edge uv such that Dist[u] + w(uv) is minimized over all directed edges $uv \in Cut(T)$, viz, Dist[v] is the minimum value of Dist[u] + w(uv) over all directed edges $uv \in Cut(T)$.

We show the action of procedure for an efficient implementation of *Dijkstra's* algorithm based on equation (6.5.1) on pg, 270 in Figure 4.1 and 4.2.

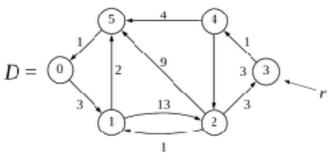
a) Action of procedure Dijkstra for the following digraph D with initial vertex r = 2, exercise 6.23 p. 284

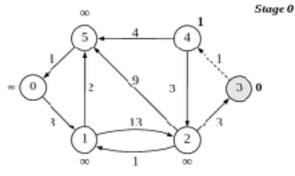


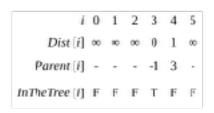


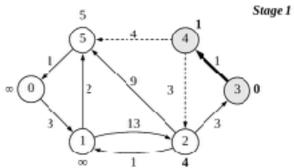
Resulting shortest path spanning-out tree rooted at vertex r=2Figure 4.1 - Action of procedure *Dijkstra* for digraph *D* with initial vertex r=2, exercise 6.23(a), p.284

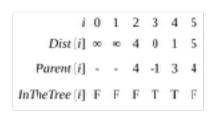
b) Action of procedure Djikstrai for digraph D with initial vertex r=3, exercise 6.23 p. 284

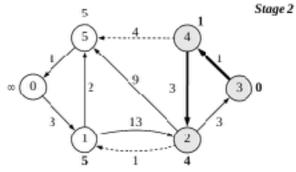




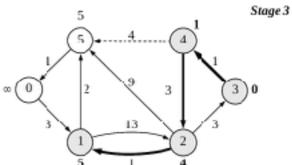




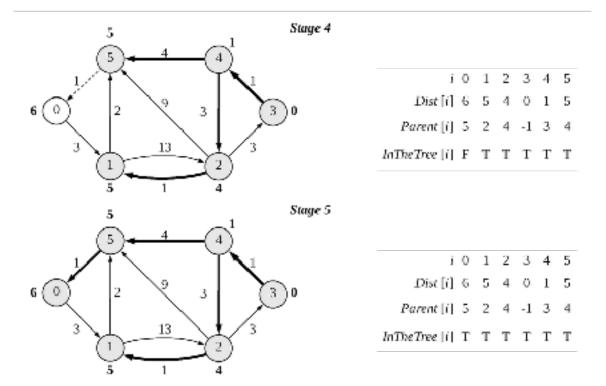




i	0	1	2	3	4	5
Dist[i]	00	5	4	0	1	5
Parent[i]		2	4	-1	3	4
InTheTree [i]	F	F	T	T	T	F



i	0	1	2	3	4	5
Dist[i]	00	5	4	0	1	5
Parent [i]	-	2	4	-1	3	4
$In The Tree \left[i \right]$	F	T	T	T	T	F



Resulting shortest path spanning-out tree rooted at vertex r=3

Figure 4.2 - Action of procedure *Dijkstra* for digraph D with initial vertex r = 3, exercise 6.23(a), p.284

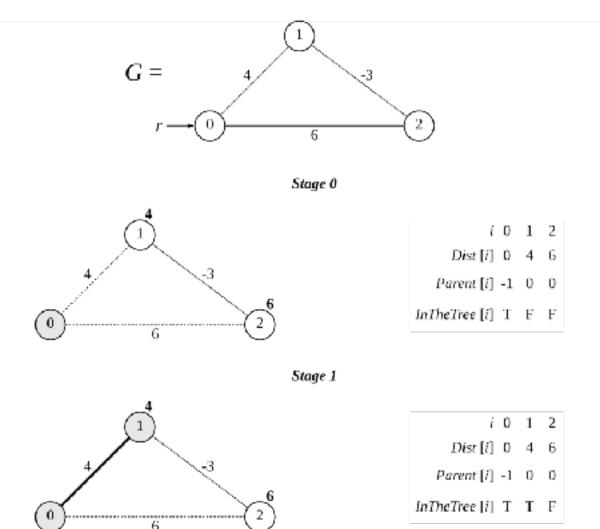
Q5 - Dijkstra's Algorithm Does Not Work on Graphs With Negative Edge Weights and Adding A Sufficiently Large Positive Constant to Each Edge Does Not Always Work Because Dijkstra Is A Greedy Algorithm

As described above, Dijkstra grows a shortest path tree T for a graph G from an initial staring vertex r by greedily choosing an edge uv with the smallest sum of w(u,v) and Dist[u], where Dist[u] is the weight of the path from r to u in T. At each step, the vertex v added to the augmented tree T updates Dist[i] (all the other vertices distance from source r) and Parent[i]. Dijkstra's greedy choice assumes at each step that Dist[I] is the shortest distance from source r, and that **no other vertex can reduce it's** distance, hence why the algorithm adds uv to the tree.

Dijkstra's Algorithm Is Not Applicable on Graphs With Negative Edge Weights Because Dijkstra's Greedy Choice Will Not Always Choose the Shortest Path

This assumption fails when edge weights are negative because the greedy choice that at each step that Dist[v] is the shortest distance from source r, and that no other vertex can reduce it's distance no longer holds. If edge weights are negative, as new vertices v not in T become adjacent to those vertices $u \in V(T)$, newly available edges could expose edge weights which would reveal paths to vertices $v \in V(T)$ shorter than those already added to the shortest-path tree T. In other words, we can no longer choose a vertex v at stage i based on locally minimizing Dist[v] = Dist[u] + w(uv) over all edges $uv \in Cut(T)$ because a new edge with negative weight discovered at later stage could reveal paths shorter than those selected at an earlier stage. Thus, Dijkstra may not always choose the shortest path if weights are negative and the algorithm is not applicable to graphs which contain negative edge weights.

Figure 5.1 illustrates the problem for a simple graph with initial vertex r = 0. Since *Dijkstra* restricts itself with vertices v belonging to $uv \in Cut(T)$, the algorithm ignores all other edges in G. The algorithm makes the greedy choice and selects edge [0,1] because Dist[1] = 4 < Dist[6] = 6. Here, the greedy choice fails because [0,1] is not the shortest path from vertex 0 to vertex 1, but rather edge [0,2,1] whose distance is 6-3=3.



Since *Dijkstra*'s greedy strategy only selects the edge in Cut(T) with the shortest corresponding path length from r to v, w[0,1] = 4 the algorithm ignores [1,2] and cannot forsee that later, a negative edge weight w[2,1] = -3 can bring the weight of the path from 0 to 1 below 4, w[0,2,1] = 3

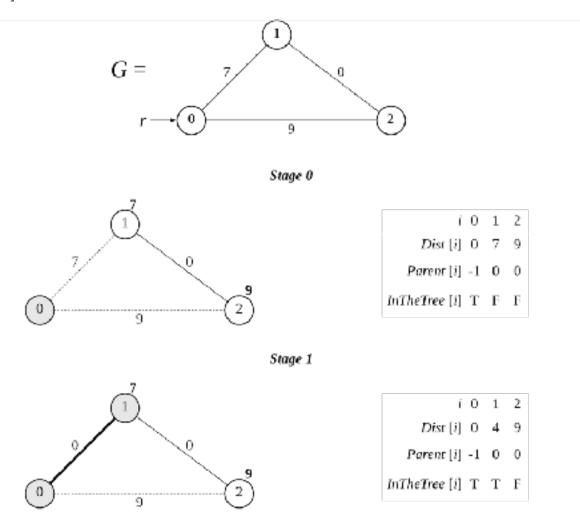
Figure 5.1 - *Dijkstra*'s algorithm fails to find the shortest path in sample graph *G* because greedy strategy cannot foresee negative edge weight

Adding A Sufficiently Large Positive Constant to Each Edge Does Not Always Work Because it Changes the Shortest Path and Distorts the Graph

Adding a sufficiently large, positive constant c to all the edges in the graph G could only work if it does not actually distort the structure graph. Unfortunately, altering the graph is exactly what happens because it can change the shortest path. Since

distance is dependent on the number of edges, adding c to each edge/path segment "penalizes" paths using more than one segment. In other words, if a path uv contains n segments, updating G by adding c to each segment means adding c to path uv n times. In the case where the shortest path is made up of more than one edge / segment, adding c to each edge in G overcompensates for the negative weight of one or more of its edges and can change which edge or edges constitute the shortest path. Thus, adding c to each edge in G changes the shortest path and alters the structure of the graph.

Figure 5.2 illustrates the problem using the same sample graph G in Figure 5.1, with c = 3. Now, Dist[1] = 4 + 3 = 7 and Dist[2] = 6 + 3 + 0 = 9. Previously in Figure 5.1, the weight of shortest path w[0,2,1] = 3. After adding c to each edge in each path, the new shortest path is [0,1] with w[0,1] = 7, and the weight of the old shortest path w[0,1,2] = 9.



Adding constant c = 3 to each edge in G distorts the graph because it changes the shortest path. Dijkstra now returns [0,1] as the shortest path when it should still be [0,2,1] as in Figure 5.1

Figure 5.1 - adding constant C changes the shortest path and alters G