Chapter 4. (Part 1)

The Greedy Approach

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Greedy Algorithm

- arrives at a solution by making a sequence of choices,
 - each of which simply *looks the best at the moment*.
- That is, each choice is *locally optimal*.
- The hope is that a *globally optimal* solution will be obtained,
 - but this is *not always* the case.
- For a given (greedy) algorithm,
 - we *must determine* whether the (greedy) solution is *always optimal*.





- The problem of *giving change* for a purchase:
 - Our goal is to give the correct change with *as few coins as possible*.
 - A *greedy approach* to the problem:
 - Initially, there are no coins in the change.
 - (selection procedure) Look for the largest coin (in value) you can find.
 - (feasibility check) If the total change does not exceed the amount owed,
 - add the coin to the change
 - (solution check) Check if the change is now equal to the amount owed.
 - If the values are not equal, repeat the process until
 - the value of the change equals the amount owed,
 - or there is no coins left.



• High-level algorithm for the greedy approach:

```
while (there are more coins and the instance is not solved) {
 grab the largest remaining coin; // selection procedure
  if (adding the coin makes the change exceed the amount owed)
    reject the coin;
                                 // feasibility check
  else
    add the coin to the change;
  if (the total value of the change equals the amount owed)
                              // solution check
    the instance is solved;
```



• An example:

- coins = [quarter, dime, dime, nickel, penny, penny] = [25, 10, 10, 5, 1, 1]
- amount owed = 36 cents.
- A greedy algorithm for giving change.
 - change = [25] < 36. Grab.
 - change = [25, 10] < 36. Grab.
 - change = $[25, 10, \frac{10}{10}] > 36$. Reject.
 - change = [25, 10, 5] > 36. Reject.
 - change = [25, 10, 1] = 36. Grab and terminate.





- Does it always result in an optimal solution?
 - Notice here that if we include a 12-cent coin with the U.S. coins,
 - the greedy algorithm *does not always* give an *optimal* solution.
 - coins = [12, 10, 5, 1, 1, 1, 1]
 - amount owed = 16 cents.
 - A greedy algorithm for giving change.
 - change = [12] < 16. Grab.
 - change = $[12, \frac{10}{10}] > 16$. Reject.
 - change = [12, 5] > 16. Reject.
 - change = [12, 1, 1, 1, 1] = 16. Grab and terminate.
 - optimal change = [10, 5, 1]



■ The *Greedy Algorithm*

- starts with an *empty set* and adds items to the set *in sequence*
 - until the set represents a solution to an instance of a problem.
- Each iteration consists of three steps:
 - 1. Selection Procedure:
 - chooses the next item to add to the set.
 - 2. Feasibility Check:
 - determines if the new set if feasible.
 - 3. Solution Check:
 - determines whether the new set constitutes a solution.

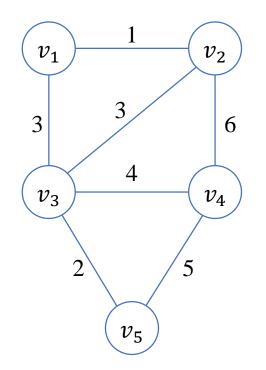


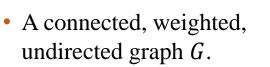
• Minimum Spanning Tree Problem:

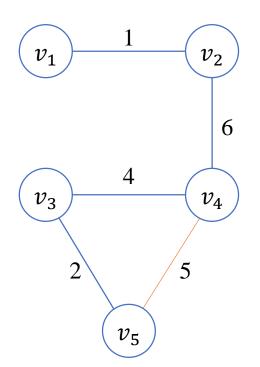
- The problem of removing edges
 - from a connected, weighted, undirected graph G
 - to form a *subgraph* such that *all the vertices* remains *connected*, and
 - the *sum of the weights* on the remaining edges is *as small as possible*.
- A *spanning tree* for *G* is a connected subgraph
 - that contains all the vertices in G and is a tree.
- A minimum spanning tree (MST) is
 - a spanning tree of *minimum weight*.



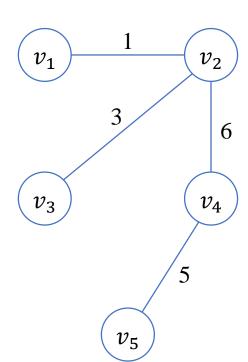




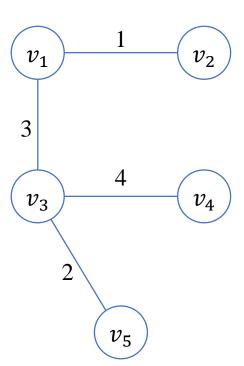




• If (v_4, v_5) were removed, the graph would remain connected.



• A spanning tree for *G*.



• A minimum spanning tree for *G*.





- Formal Definition of the MST Problem:
 - Given a connected, weighted, undirected graph G = (V, E).
 - A spanning tree T for G has the same vertices V as G,
 - but the set of edges of *T* is a subset *F* of *E*.
 - Denote a spanning tree by T = (V, F).
 - Our problem is to find a subset F of E
 - such that T = (V, F) is a minimum spanning tree for G.





High-level greedy algorithm for the MST problem

```
F = \emptyset;
while (the instance is not solved) {
  select an edge according to some locally optimal consideration;
  if (adding the edge to F does not create a cycle)
     add the edge to the solution;
  else
     discard the edge;
  if (T = (V, F) \text{ is a spanning tree})
     the instance is solved;
```



Prim's Algorithm

- starts with an *empty set* of edges *F*
 - and a *subset of vertices Y* initialized to contain an *arbitrary* vertex (v_1) .
- A vertex *nearest* to Y is a vertex in V-Y
 - that is connected to a vertex in Y by an edge of *minimum weight*.
- The *vertex* that is *nearest* to Y is added to Y
 - and the *edge* is added to *F*. (Ties are broken arbitrarily)
- This process of adding nearest vertices is
 - repeated until Y = V.

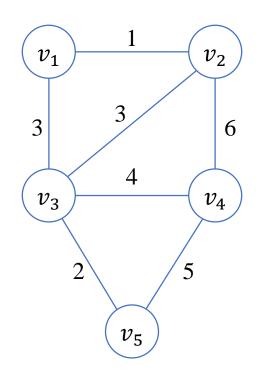


• High-level pseudo-code for the Prim's algorithm

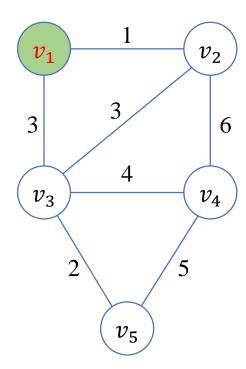
```
F = \emptyset;
Y = \{v_1\};
while (the instance is not solved) {
  select a vertex in V - Y that is nearest to Y;
  add the vertex to Y;
  add the edge to F;
  if (Y = V)
     the instance is solved;
```



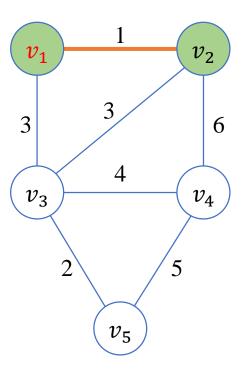
• Determine a minimum spanning tree



1. Vertex v_1 is selected first

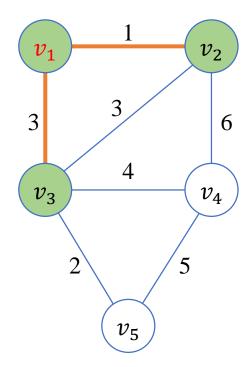


2. Vertex v_2 is selected because it is nearest to $\{v_1\}$

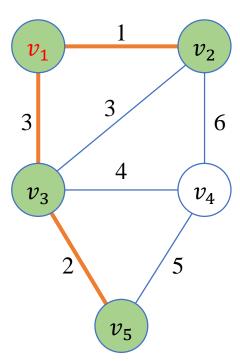




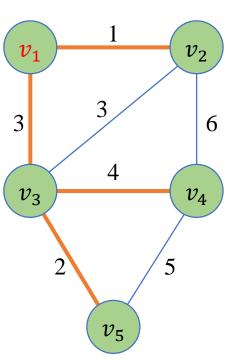
3. Vertex v_3 is selected because it is nearest to $\{v_1, v_2\}$



4. Vertex v_5 is selected because it is nearest to $\{v_1, v_2, v_3\}$



Vertex v_4 is selected because it is nearest to $\{v_1, v_2, v_3, v_5\}$





- Implementing the Prim's algorithm:
 - Represent a weighted graph by its adjacency matrix.

$$- W[i][j] = \begin{cases} weight \ on \ edge \\ \infty \end{cases} \qquad \text{if there is an edge between } v_i \ \text{and } v_j \\ 0 \qquad \qquad \text{if } i = j \end{cases}$$

- We maintain two arrays, *nearest* and *distance*, where, for i = 2, ..., n,
 - $nearest[i] = index of the vertex in Y nearest to <math>v_i$
 - distance[i] = weight on edge between v_i and the vertex indexed by nearest[i]



ALGORITHM 4.1: Prim's Algorithm

```
void prim(int n, matrix_t& W, set_of_edges& F)
    int vnear, min;
    vector<int> nearest(n + 1), distance(n + 1);
    F.clear(); // F = \emptyset;
    for (int i = 2; i <= n; i++) {
        nearest[i] = 1;
        distance[i] = W[1][i];
```



ALGORITHM 4.1: Prim's Algorithm (continued)

```
repeat (n - 1 times) {
    min = \infty;
    for (int i = 2; i <= n; i++)
        if (0 <= distance[i] && distance[i] < min) {</pre>
            min = distance[i];
            vnear = i;
    e = edge connecting vertices indexed by vnear and nearest[vnear];
    add e to F;
    distance[vnear] = -1;
    for (int i = 2; i <= n; i++)
        if (distance[i] > W[i][vnear]) {
            distance[i] = W[i][vnear];
            nearest[i] = vnear;
```



```
#define INF 0xffff
typedef vector<vector<int>> matrix t;
typedef vector<pair<int, int>> set_of_edges;
typedef pair<int, int> edge t;
// e = edge connecting vertices indexed by vnear and nearest[vnear];
// add e to F;
F.push back(make pair(vnear, nearest[vnear]));
set_of_edges F;
prim(n, W, F);
for (edge_t e: F) {
    u = e.first; v = e.second;
    cout << u << " " << v << " " << W[u][v] << endl;
```



							i	2	3	4	5	е
						ini+.	nearest[i]	1	1	1	1	
W	1	2	3	4	5	init:	distance[i]	1	3	∞	∞	
1	0	1	3	∞	∞	step 1:	nearest[i]	1	1	2	1	(2 1 1)
2	1	0	3	6	∞	scep 1.	distance[i]	-1	3	6	∞	(2, 1, 1)
3	3	3	0	4	2	step 2:	nearest[i]	1	1	3	3	(3, 1, 3)
4	∞	6	4	0	5		distance[i]	-1	-1	4	2	
5	∞	∞	2		0	step 3:	nearest[i]	1	1	3	3	(5 2 2)
5		∞	2	J	U		distance[i]	-1	-1	4	-1	(5, 3, 2)
						step 4:	nearest[i]	1	1	3	3	(4 2 4)
						3сер 4.	distance[i]	-1	-1	-1	-1	(4, 3, 4)

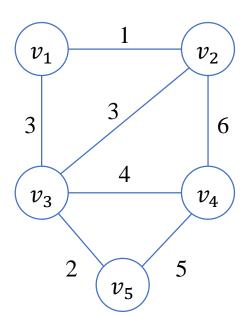




- Time Complexity of Algorithm 4.1:
 - Basic Operation: the *instructions* inside each of two loops.
 - Input Size: *n*, the *number of vertices*.
 - Note that there are two (nested) loops,
 - and the *repeat* loop has n-1 iterations.
 - Therefore,
 - $T(n) = 2(n-1)(n-1) \in \Theta(n^2)$



- Does it always produce an optimal solution?
 - We need to prove that
 - Prim's algorithm *always* produces a minimum spanning tree.
 - Given an undirected graph G = (V, E),
 - A subset *F* and *E* is called *promising*
 - if edges can be added to it so as to *form* a *minimum spanning tree*.



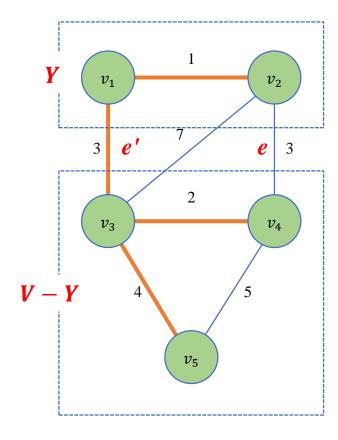
- The subset $\{(v_1, v_2), (v_1, v_3)\}$ is promising.
- The subset $\{(v_2, v_4)\}$ is not promising.



• Lemma:

- If *F* is a *promising* subset of *E*
 - then $F \cup \{e\}$ is *promising*,
 - where *e* is an edge of minimum weight that
 - connects a vertex in Y and a vertex in V-Y.
- Proof:
 - Let F' be a set edges in an MST s.t. $F \subseteq F'$.
 - If $e \in F'$, then $F \cup \{e\} \subseteq F'$.
 - If $e \notin F'$, then $F' \cup \{e\}$ must have a cycle.
 - There is an edge $e' \notin F'$ in the cycle
 - Remove e', then the cycle disappears.
 - Hence, $F' \cup \{e\} \{e'\}$ is an MST.
 - Hence, $F \cup \{e\} \subseteq F' \cup \{e\} \{e'\}$.

$$F = \{(v_1, v_2)\}$$



$$F' = \{(v_1, v_2), (v_1, v_3), (v_3, v_4), (v_3, v_5)\}$$

$$F' \cup \{e\} \text{ has a cycle: } [v_1, v_2, v_4, v_3]$$



• Theorem:

- Prim's algorithm *always produces* a minimum spanning tree.
- Proof:
 - Clearly, the *empty set* Ø is *promising*.
 - Assume that, after a given iteration,
 - the selected edges *F* is *promising*.
 - The set $F \cup \{e\}$ is *promising*,
 - where *e* is the edge selected in the next iteration.
 - Because the e is an edge of minimum weight that
 - connects a vertex in Y to a vertex in V Y. (by the Lemma)





Kruskal's Algorithm

- starts by creating *disjoint subsets* of *V*,
 - one for *each vertex* and containing *only that vertex*.
- If then, inspects the edge according to nondecreasing weight
 - ties are broken arbitrarily.
- If an edge *connects* two vertices in *disjoint subsets*,
 - the edge is *added* and the subsets are *merged into one set*.
- This process is repeated
 - until *all the subsets* are *merged into one set*.

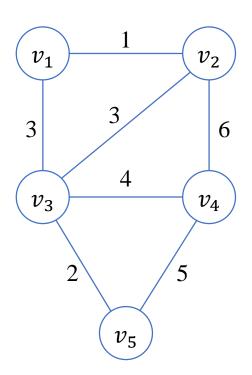


High-level pseudo-code for the Kruskal's algorithm

```
F = \emptyset;
create disjoint subsets of V, one for each vertex and containing only that vertex;
sort the edges in E in nondecreasing order;
while (the instance is not solved) {
  select next edge;
  if (the edge connects two vertices in disjoint subsets) {
    merge the subsets;
    add the edge to F;
  if (all the subsets are merged)
    the instance is solved;
```



• Determine a minimum spanning tree.



1. Edges are sorted by their 2. Disjoint sets are created. weights.

-	
edges	weight
(v_1, v_2)	1
(v_3, v_5)	2
(v_1, v_3)	3
(v_2, v_3)	3
(v_3, v_4)	4
(v_4, v_5)	5
(v_2, v_4)	6

v_1	v_2
v_3	v_4

 v_5



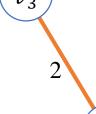
- 3. The first edge (v_1, v_2) is selected
- 4. Next edge (v_3, v_5) is selected
- 5. Next edge (v_1, v_3) is selected





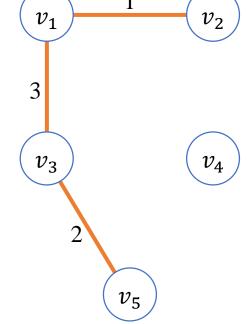








 v_5

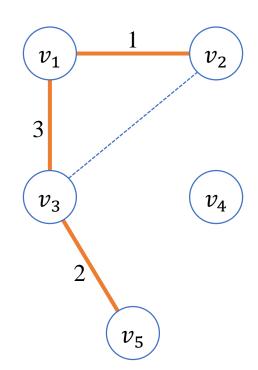


 v_5

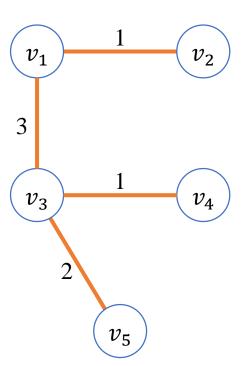
• Ties are broken *arbitrarily*



6. Next edge (v_2, v_3) is *discarded*, 7. Next edge (v_3, v_4) is because it creates a cycle



selected



 (v_4, v_5) and (v_2, v_4) are not considered, because all the subsets are merged



- Abstract Data Type: Disjoint Set
 - To write a formal version of Kruskal's algorithm,
 - we need a *disjoint set* abstract data type: Refer to *Appendix C*.
 - The ADT of the disjoint set defines two data types:
 - index i:
 - set_pointer p, q;
 - Then the routines are defined:
 - initial(n): initialzes n disjoint subsets.
 - p = find(i): makes p point to the set containing index i.
 - merge(p,q): merges the two sets, to which p and q point, into the set.
 - equal(p,q): returns true if p and q both point to the same set.

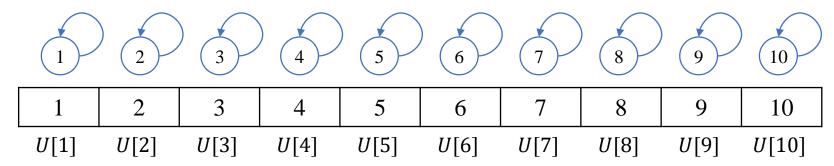


• Let $U = \{A, B, C, D, E\}$ be a universe of elements

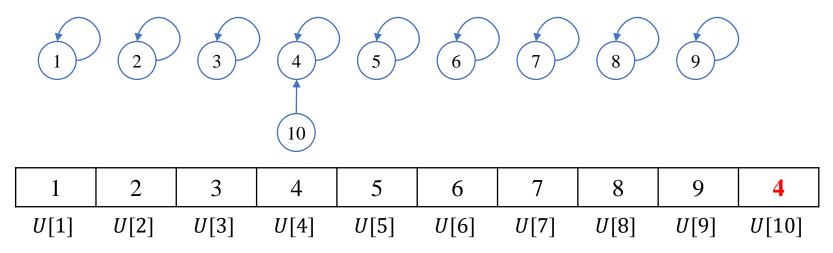
```
for i in U:
                                                             (disjoint sets)
                                    {C}
                                            {D}
                                                     {E}
    initial(i); {A}
                           {B}
   p = find(B);
   q = find(C);
                          {B, C}
                                                     {E}
                                             {D}
   merge(p, q); \{A\}
    p = find(C);
                                                           equal(C, E);
    q = find(E);
                                                              returns false;
                         \{B,C,E\}
   merge(p, q); \{A\}
                                             {D}
   p = find(C);
                                                           equal(C, E);
                              \boldsymbol{q}
   q = find(E);
                                                              returns true;
```



initial(10);



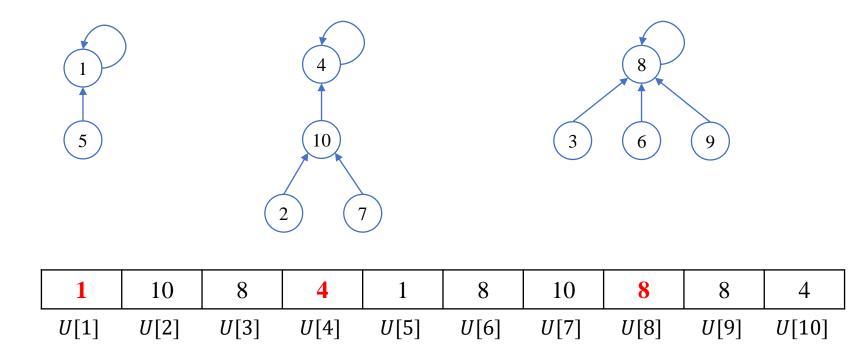
merge(find(4), find(10);







After several union and find:



- Analyze the complexity of union (merge) and find,
 - and improve the efficiency to $\Theta(m \lg m)$,
 - where *m* is the *number of passes* to call the routines(*merge* and *find*).



```
vector<int> dset;
void dset_init(int n) {
    dset.resize(n + 1);
    for (int i = 1; i <= n; i++)
        dset[i] = i;
int dset_find(int i) {
    while (dset[i] != i)
        i = dset[i];
    return i;
void dset_merge(int p, int q) {
    dset[p] = q;
```





ALGORITHM 4.2: Kruskal's Algorithm

```
void kruskal(int n, int m, set of edges& E, set of edges& F) {
    int p, q;
    edge t e;
    PriorityQueue PQ;
    sort the m edges in E by weight in nondecreasing order;
    F.clear(); // F = \emptyset;
    dset_init(n);
    while (number of edges in F is less than n - 1) {
         e = PQ.top(); PQ.pop(); // edge with least weight not yet considered;
        p = dset find(e.u);
        q = dset_find(e.v);
        if (p != q) {
            dset merge(p, q);
            F.push back(e); // add e to F
```



4.1 Minimum Spanning Trees

```
typedef struct edge {
    int u, v, w;
} edge t;
struct edge_compare {
    bool operator()(edge_t e1, edge_t e2) {
        if (e1.w > e2.w) return true;
        else return false;
};
typedef vector<edge t> set of edges;
typedef priority_queue<edge_t, vector<edge_t>, edge_compare> PriorityQueue;
// sort the m edges in E by weight in nondecreasing order;
for (edge_t e: E)
    PQ.push(e);
```





4.1 Minimum Spanning Trees

- Time Complexity of Algorithm 4.2:
 - Basic Operation: a *comparison* instruction.
 - Input Size: *n*, the *number of vertices*, and *m*, the *number of edges*.
 - Three considerations in this algorithm:
 - 1. The time to sort the edges: $\Theta(m \lg m)$
 - 2. The time to initialize n disjoint sets: $\Theta(n)$.
 - 3. The time in the while loop: $\Theta(m \lg m)$
 - The time complexity of *Union-Find* (Appendix C)
 - Since $m \ge n 1$, $W(m, n) \in \Theta(m \lg m)$
 - In worst-case, the number of edges is m = n(n-1)/2
 - $w(m,n) \in \Theta(n^2 \lg n^2) = \Theta(n^2 \lg n)$



4.1 Minimum Spanning Trees

- Comparing Prim's Algorithm with Kruskal's Algorithm:
 - The time complexity of two algorithms:
 - Prim's: $T(n) \in \Theta(n^2)$
 - Kruskal's: $W(m, n) \in \Theta(m \lg m) = \Theta(n^2 \lg n)$
 - We can show that $n-1 \le m \le \frac{n(n-1)}{2}$.
 - For a *sparse graph*,
 - whose number of edges m is near the low end of these limits,
 - Kruskal's algorithm is $\Theta(n \lg n)$, which is *faster than Prim's*.
 - For a dense graph,
 - whose number of edges m is near the high end of those limits,
 - Kruskal's algorithm is $\Theta(n^2 \lg n)$, which is *slower than Prim's*.



- The Problem of *Single-Source-Shortest-Paths*
 - Find a shortest path from *one particular vertex* to *all the others*.
 - *Dijkstra's Algorithm* uses the *greedy approach*
 - to develop a $\Theta(n^2)$ algorithm for this problem.



Dijkstra's Algorithm

- initializes a set Y to contain only the source vertex v_1 ,
 - and initializes a set *F* of edges to being empty.
- First, choose a vertex v that is *nearest* to v_1 ,
 - add it to Y, and add the edge $\langle v_1, v \rangle$ (a *shortest edge*) to F.
- Next, check the paths from v_1 to the vertices in V-Y
 - that allow only vertices in *Y* as intermediate vertices.
- The vertex at the end of such a path is added to *Y*,
 - and the edge (on the path) that *touches* that vertex is added to *F*.
- Continue this procedure, until *Y* equals *V*.
 - At this point, *F* contains the edges in *shortest paths*.



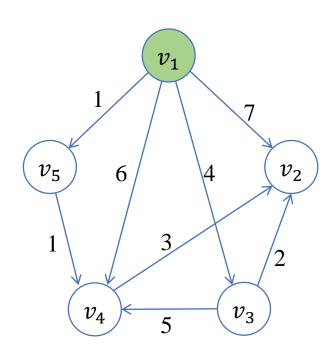


• High-level pseudo-code for the Dijkstra's algorithm:

```
Y = \{v_1\};
F = \emptyset;
while (the instance is not solved) {
  select a vertex v from V-Y that has a shortest path from v_1,
      using only vertices in Y as intermediates;
  add the new vertex v to Y;
  add the edge (on the shortest path) that touches v to F;
  if (Y = V)
     the instance is solved;
```

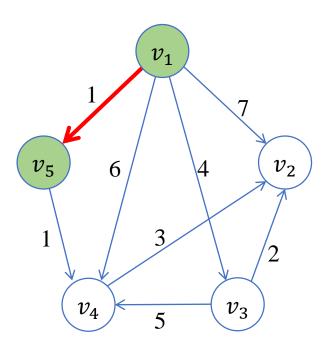


• Compute shortest paths from v_1 .

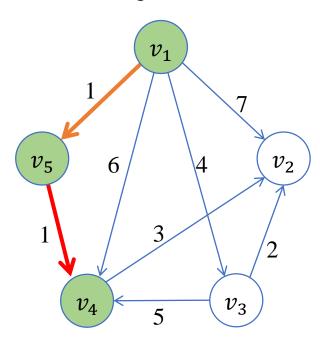




1. Vertex v_5 is selected because it is nearest to v_1 .



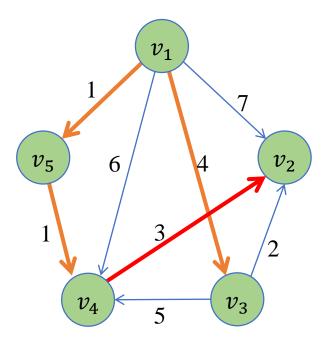
2. Vertex v_4 is selected because it has the shortest path from v_1 using only vertices in $\{v_5\}$ as intermediates.





- 3. Vertex v_3 is selected because it has the shortest path from v_1 using only vertices in $\{v_4, v_5\}$ as intermediates.
 - v_5 v_2 6 v_4 v_3

The shortest path from v_1 to v_2 is $[v_1, v_5, v_4, v_2].$





- Implementation of Dijkstra's Algorithm
 - It is very *similar* to *Prim's* Algorithm.
 - The difference is that, instead of the arrays *nearest* and *distance*,
 - we have arrays *touch* and *length*, where for $i = 2, \dots, n$.
 - Let us define:
 - touch[i] = index of $vertex\ v$ in Y such that the edge $\langle v, v_i \rangle$ is the $last\ edge$ on the $current\ shortest\ path$ from v_1 to v_i using $only\ vertices$ in Y as intermediates.
 - length[i] = length of the current shortest path from v_1 to v_i using only vertices in Y as intermediates.



ALGORITHM 4.3: Dijkstra's Algorithm

```
void dijkstra(int n, matrix_t& W, set_of_edges& F)
    int vnear, min;
    vector<int> touch(n + 1), length(n + 1);
    F.clear();
    for (int i = 2; i <= n; i++) {
        touch[i] = 1;
        length[i] = W[1][i];
```



ALGORITHM 4.3: Dijkstra's Algorithm (continued)

```
repeat (n - 1 times) {
    min = INF;
    for (int i = 2; i <= n; i++)
        if (0 <= length[i] && length[i] < min) {</pre>
            min = length[i];
            vnear = i;
    e = edge from vertex indexed by touch[vnear];
    add e to F;
    for (int i = 2; i <= n; i++)
        if (length[i] > length[vnear] + W[vnear][i]) {
            length[i] = length[vnear] + W[vnear][i];
            touch[i] = vnear;
    length[vnear] = -1;
```



init

step 1

step 2

step 3

step 4

	ı				
W	1	2	3	4	5
1		7	4	6	1
2	∞	0	∞	∞	∞
3	∞	2	0	5	∞
4	∞	3	∞	0	∞
5	∞	∞	∞	1	0

	i	2	3	4	5	е	
: -	touch[i]	1	1	1	1		
	length[i]	7	4	6	1		
:	touch[i]	1	1	5	1	(1 5 1)	
	length[i]	7	4	2	-1	(1, 5, 1)	
:	touch[i]	4	1	5	1	(5, 4, 1)	
	length[i]	5	4	-1	-1		
:	touch[i]	4	1	5	1	(1, 3, 4)	
	length[i]	5	-1	-1	-1		
·: _	touch[i]	4	1	5	1	(4, 2, 3)	
	length[i]	-1	-1	-1	-1		



- The Lengths of Shortest Paths:
 - Algorithm 4.3 determines only the edges in the shortest paths.
 - It does not produce the lengths of those paths.
 - These lengths could be obtained from the edges.
 - Alternatively, they can be computed and stored in an array as well.

- Time Complexity of Algorithm 4.3
 - is the same with that of Algorithm 4.1 (Prim's Algorithm)
 - $T(n) = 2(n-1)^2 \in \Theta(n^2)$

Any Questions?

