

Graph Algorithms: Dynamic Programming

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Dynamic Programming: Algorithms for Graphs

- Most graph properties address optimisation goals, namely
 - a. Shortest paths
 - b. Minimum Spanning Trees
 - c. Minimum Hamiltonian tours (Traveling Salesman)
 - d. Minimum number of colours
- Some of these properties (e.g. **a** and **b**, but not **c** nor **d**), can be computed by polynomial algorithms.
- In most cases, algorithms to compute the optima may follow a methodology, dynamic programming, based on Mathematical Induction on the Integers:
 - Once an optimal solution is obtained with **n** nodes, extend it to **n+1** nodes.
- We will see two examples of this, in the following algorithms
 - Minimum Spanning Tree **Prim's Algorithm**
 - Shortest Paths Floyd-Warshall's Algorithm



- A **spanning tree** is a subset of a connected graph that has the topology of a tree and covers all nodes of the graph.
- It has many applications, namely to provide services to a number of sites (the nodes) that can be interconnected in several ways (by a graph), but using the a minimal number of connections that allow all sites to be reached, i.e. a single path connecting any two nodes.
- Among these spanning trees one is usually interested in **minimum spanning trees** (MST) that minimise the sum of the costs of the arcs selected for the tree.
- There are many polynomial algorithms that may be used to compute these MSTs, the most common ones are the Kruskal's and the Prim's algorithms.
- Given the similarities between the latter and the algorithm to check connectedness of a graph, we will address now the **Prim's Algorithm**.



- The Prim's algorithm is an example of Dynamic Programming that extends a MST with n nodes to n+1 nodes, with an eager selection of the new node (i.e. once the node is selected, the selection is not backtracked for alternatives).
- The algorithm can be understood as a process of increasing the size of a current MST, starting with 1 node and ending with all the nodes, and specified as follows:
 - Maintain two sets of nodes: In and Out, where In is the set of nodes already included in a current MST and Out are those not yet included.
 - 1. Select arbitrarily a node from the tree to initialise the **In** set, and put the others in the **Out** set;
 - 2. While there are nodes in the **Out** set,
 - i. Find which node from the **Out** set has an arc of least cost connecting it to one of the nodes of the **In** set;
 - ii. Transfer the node from the **Out** set to the **In** set and include the least cost arc in the current **MST**.





- Start with an arbitrary node in the In set
- Start with the Out set with all the other nodes
- Initialise the MST to empty





• Chose that with minimum cost







- Move the out node of the arc from the Out to the In set.
- Include the arc in the MST



- Check all arcs between nodes in the In and Out sets
 - Chose that with minimum cost







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- The **Out** set is now empty
- Return the MST.





- Several variants can be used in the implementation of the Prim's algorithm, using appropriate data structures that make it more efficient. Here we present a simple implementation that nonetheless is acceptable for relatively large graphs.
 - 1. Select arbitrarily a node from the tree to initialise the **In** set (here node 0);
 - 2. Put the other nodes in the **Out** set;
 - 3. Initialises the Minimum spanning tree **T** to an empty graph
 - 4. Update and eventually returns the tree **T** as well as the remaining Out nodes

```
def prim(G):
    """ Returns the minimum spanning tree of graph G,
    using the prim algorithm"""
    n = len(G)
    In = [0]
    Out = [i for i in range(1,n)]
    row = [-1 for i in range(n)]
    T = [row.copy() for i in range(n)]
    for i in range(n):
        T[i][i] = 0
    while ...:
    return (T, Out)
```



- The tree is then updated as follows:
 - 1. While there are nodes in the **Out** set,
 - Find the arc of least cost between a node u in the In set and a node v from the Out set (if there is one!);
 - ii. If no arc is selected, that means the graph is not connected and should be returned (together as the remaining Out nodes)
 - iii. Include the least cost arc in the current **MST**.
 - iv. Transfer the node from the **Out** set to the **In** set and



- The tree is then updated as follows:
 - 1. While there are nodes in the **Out** set,
 - Find the arc of least cost between a node u in the In set and a node v from the Out set (if there is one!);
 - ii. If no arc is selected, that means the graph is not connected and should be returned (together as the remaining Out nodes); otherwise
 - iii. Include the least cost arc in the current **MST**.
 - iv. Transfer the node from the Out set to the In set



• The complete algorithm is shown below:

```
def prim(G):
    n = len(G)
    In = [0]
   Out = [i for i in range(1,n)]
    row = [-1 for i in range(n)]
    T = [row.copy() for i in range(n)]
    for i in range(n):
        T[i][i] = 0
    while len(Out) > 0:
        min arc = math.inf
        u = 0
        v = 0
        for i in In:
            for j in Out:
                if G[i][j] > 0 and G[i][j] < min arc:
                    u = i
                    v = j
                    min arc = G[i][j]
        if u == 0 and v == 0:
            return (T, Out)
        T[u][v] = G[u][v]
        T[v][u] = G[v][u]
        In.append(v)
        Out.remove(v)
    return (T, Out)
```



- It is easy to prove, by induction, that the algorithm is correct. If T is an MST with least cost with n nodes, adding to it the least cost arc will make it an MST with least cost with n+1 nodes (adding any other arc would lead to a higher cost spanning tree).
- As to the worst cost complexity of the algorithm, with this implementation, we notice that the while loop is executed **n-1** times (n is the number of nodes of the graph, |V|).

- Finding the minimal cost arc, when k nodes are already in the ln list, requires two nested loops over ranges with k and n-k values, i.e. at most n²/4 (for k = n/2) executions of the body of the loop
- All operations in the loop are "basic", and so the complexity of this implementation of the Prim's algorithm is O(n*n²/4) i.e. O(|V|³) (where |V| = n).
- Note: Implementations with priority queues and other advanced data structures have better complexity, namely O(|E|+Vlog|V|).



- There are many algorithms for finding shortest paths between nodes of weighted graphs. They include algorithms to find one shortest path between two nodes, like the Dijskstra algorithm, or to find all shortest paths between any two nodes of the graph, namely the Floyd-Warshall's (FW) algorithm.
- As the previous one, the **FW** algorithm explores dynamic programming in the following way:
- The initial shortest path between any two nodes, is the direct distance (that can be infinite).
- A list **In** is initialised with all **n** nodes;
- The current shortest distance between two nodes is then updated by checking whether an indirect path exists passing in each of the nodes in list **In**.
- The final result is a matrix with all minimal distances between any two nodes.



- The algorithm can thus be implemented as follows:
 - 1. Initialise a matrix **S** of shortest paths with the adjacency matrix (that is, only direct distances between any two nodes are initially considered).
 - Of course, nodes that are not directly connected by an arc have a distance of -1 at this stage. For convenience, we will assign them to **inf**.
 - 1. Now, for all values **k** in the **In** list (i.e. from 0 to n-1) iterate.
 - In iteration k, update S, by considering all indirect paths between nodes i and j passing through node k.
 - 3. After the last iteration, matrix S contains all shortest paths between any two nodes of G.
- Notice that this algorithm only computes the paths with shortest distance between any two nodes but does not return what these paths are.
 - In fact, a small addition to the algorithm, coming shortly, allows the paths to be obtained.



- The external for loop guarantees that all paths, between nodes i and j, consider, all the paths through nodes k (1, 2, 3, ..., n), previously computed.
- The shortest paths are updated by considering the triangular inequality, with paths passing through the previous values of **k**.

```
def floyd(G):
 """ Returns the minimum distances between any two modes
    of graph G, using the Floyd-Warshall's algorithm."""
    n = len(G)
    S = [ [ math.inf for i in range(n) ] for j in range(n) ]
    for i in range(n):
        for j in range(n):
            if G[i][j] != -1:
                S[i][j]= G[i][j]
    In = [i for i in range(n)]
    for k in In:
      for i in range(n):
         for j in range(n):
            if S[i][k] + S[k][i] < S[i][j]:
               S[i][j] = S[i][k] + S[k][j]
    return S
```



- The correction of the algorithm can be proved by induction on the number of nodes considered in indirect paths (left as exercise).
- As to the complexity, it is easy to see that the algorithm requires 3 nested loops of size **n**, with a basic operation in the body,

```
for k in In: # In = [0..n-1]
for i in range(n):
    for j in range(n):
```

- The complexity of the algorithm is thus $O(|V|^3)$.
- Notice that algorithms to compute shortest paths between 2 nodes, like the Dijkstra algorithm, have complexity O(|V|²), but they do not consider the distance between all the nodes.



- The previous algorithm does not provide the shortest paths between any two nodes, but rather the shortest distances of any path between the nodes.
- Nevertheless, these paths may be easily reconstructed if a matrix is computed during the FW algorithm, to it possible to later compute the shortest path from some node **i** to another node **j**.
- Matrix Next plays this role. In such matrix, Next[i][j] = k means that to follow the shortest path to node j, the path should start by the arc i → k.
- All that is needed is to compute matrix Next during the FW algorithm. To do so, all values Next[i][j] should be initialised with j (in the beginning, i.e. before exploring the graph, the best path is the direct path).
- In fact, if there is no connection between nodes i and j, Next[i][j] should be initialised to inf, to account for that non-connection.
- Then, if a better path is found through node k, Next[i][j] must be updated to Next[i][k], i.e. to go from i to j, one should start in the best arc to go from i to k.
- Better paths are found in the inner loop of the FW algorithm, so one needs simply to add some extra instructions to function floyd just developed.

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Dynamic Programming: Graph Algorithms



• The initialisation of matrix Next (i.e. **Next[i][j] = j**) can be implemented as:

Next = [[j for j in range(n)] for i in range(n)]

• If there is no connection, this should be accounted for in Next

```
if G[i][j] != -1:
    ...
else:
    Next[i][j] = math.inf
```

- The update of the elements of Next may be done in the inner loop of the floyd function, taking into account that
 - If a better path is found, through node k, Next[i][j] must be updated to Next[i][j], i.e. to go from i to j, one should start in the best arc to go from i to k.

```
for k in In:
    for i in range(n):
        for j in range(n):
            if S[i][k] + S[k][i] < S[i][j]:
                S[i][j] = S[i][k] + S[k][j]
                Next[i][j] = Next[i][k]</pre>
```



• The completed Floyd function is thus shown below (changes in bold).

```
def floyd(G):
    ....
   n = len(G)
   S = [ [ math.inf for i in range(n) ] for j in range(n) ]
    Next = [ [ j for j in range(n) ] for i in range(n) ]
    for i in range(n):
        for j in range(n):
            if G[i][i] != -1:
                S[i][i]= G[i][i]
            else:
                Next[i][j] = math.inf
    In = [i for i in range(n)]
    for k in In:
      for i in range(n):
         for j in range(n):
            if S[i][k] + S[k][i] < S[i][j]:
               S[i][j] = S[i][k] + S[k][j]
               Next[i][j] = Next[i][k]
    return (S, Next)
```



• Once the matrix Next is returned the path between any two nodes, u and v, can be obtained by following the trail indicated by this matrix, as shown below

```
def path(u,v,Next):
    """ Returns the shortest path between nodes u and v,
    according to matrix Next computed with function floyd."""
    if Next[u][v] == math.inf:
        return []
    P = [u]
    while u != v:
        u = Next[u][v]
        P.append(u)
    return P
```

- The first test checks whether there is any path between nodes **u** and **v**. If not it returns an empty path.
- Otherwise the path is "reconstructed", starting in node u ...
- ... and continuing until reaching node v
- With this reconstruction technique, the complexity of the FW algorithm is not changed, and the paths are only computed when needed.