

Graphs

17/4/19

- Defined by $g = (V, E)$, i.e a set of vertices and edges
- If **directed**, edges are $v_1 \rightarrow v_2$
- If **undirected**, edges are $v_1 \leftrightarrow v_2$
- A path is a sequence of vertices connected by edges: $v_1 \rightarrow v_2 \rightarrow \dots \rightarrow v_k$
or $v_1 \leftrightarrow v_2 \leftrightarrow \dots \leftrightarrow v_k$
- An undirected graph is **connected** if there is a path between any two vertices
 - ↳ a forest is an undirected acyclic graph
 - ↳ a tree is a connected forest.
- Graphs can be represented as:
 - (1) adjacency lists, i.e an array of linked lists containing neighbours
 - (2) adjacency matrix. Requires $O(V^2)$ space.
 ↳ choose depending on sparsity. If E/V^2 is large, use matrix.

Depth-first search

- Visits all vertices reachable from a starting vertex, by recursively searching neighbours (but marking visited to prevent loops).
- Can be implemented with a stack

def dfs(g, s):

for v in g.vertices:
v.visited = False

toexplore = Stack(s)

s.visited = True

while not toexplore.empty():
v = toexplore.pop()

for w in v.neighbours:

if not w.visited:

toexplore.push(w)

w.visited = True

} run once per edge $\therefore O(CE)$

Thus DFS is $O(V+E)$, using aggregate analysis.

- The code for a breadth-first search is identical, except a queue is used rather than a stack.

Dijkstra's Algorithm

- for a graph whose edges have positive weights, Dijkstra's algo allows us to find the shortest ~~min weight~~ path.
- Similar to BFS, except we use a priority queue to store frontier vertices, and greedily choose the nearest one.
 - if we see a vertex that has already been visited, we update its distance and its position in the PQ.
- Once an item is popped, its distance is the minimum distance and it never gets added back to the PQ ^{true}
- So each vertex called popmin(), and we had to push/decreasekey for each edge.
 - using Fibonacci heap: popmin() is $O(\lg n)$, push/decreasekey $O(1)$
 - Dijkstra runtime is $\boxed{O(E + V \log V)}$

Proof of correctness (by contradiction):

- Let v be the first vertex for which after popmin(), $v.\text{distance}$ is not the true shortest distance.
- Let a shortest path from s to v be $s = v_1 \rightarrow \dots \rightarrow v_k = v$
- Let v_i be the first vertex that has not been popped.

$$\text{distance}(s \text{ to } v) < v.\text{distance}$$

$$\leq v_i.\text{distance}$$

$$\leq v_{i-1}.\text{distance} + \text{cost}(v_{i-1} \rightarrow v_i)$$

$$= \text{distance}(s \text{ to } v_{i-1}) + \text{cost}(v_{i-1} \rightarrow v_i)$$

$$\leq \text{distance}(s \text{ to } v).$$

Contradiction $\therefore v.\text{distance}$ is the true shortest distance.

- Thus it cannot be pushed back into the PQ
- And thus the algo must terminate

The Bellman equation

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- Let vertices represent states and edges represent actions. The goal is to find the best sequence of actions.
- Uses different terminology to Dijkstra (more general)
 - cost \rightarrow weight
 - distance \rightarrow minweight
 - shortest path \rightarrow minimal weight path.
- e.g. finding the best FX rate, with weight = $-\ln(\text{exchange rate})$.
- Same as dynamic programming:

$$W_{ij} = \begin{cases} 0, & i = j \\ \text{weight}(i \rightarrow j), & \text{if there is an edge} \\ \infty, & \text{otherwise.} \end{cases}$$

\hookrightarrow minweight action to go from state i to j

$M_{ij}^{(l)}$ is the minimal weight path from i to j in l steps.

$$M_{ij}^{(1)} = W_{ij} \quad M_{ij}^{(l)} = \min_k \{ W_{ik} + M_{kj}^{(l-1)} \}$$

NB: Assumes no negative weight cycles.

\nearrow i.e. make an intermediate jump then choose best path.

- Can be reformulated as 'matrix multiplication':

$$M_{ij}^{(l)} = (W_{ij} + M_{i1}^{(l-1)}) \wedge (W_{i2} + M_{2j}^{(l-1)}) \wedge \dots \wedge (W_{in} + M_{nj}^{(l-1)})$$

where $x \wedge y = \min(x, y)$, $n = |V|$.

\hookrightarrow the minimal weight path must have $\leq n$ edges because we assumed no negative weight cycles

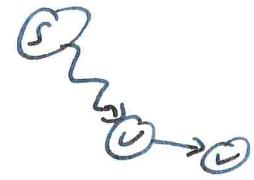
\hookrightarrow requires $\log V$ matrix multiplications, $\therefore O(V^3 \log V)$

- This is a brute-force algo and often cannot be used.

Bellman-Ford

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- If we have a path from s to u and edge $u \rightarrow v$, we can update $v.\text{minweight}$ as follows:

(s) 
if $v.\text{minweight} > u.\text{minweight} + \text{weight}(u \rightarrow v)$:
 $v.\text{minweight} = u.\text{minweight} + \text{weight}(u \rightarrow v)$. } 'relaxing' $u \rightarrow v$

- Bellman-Ford uses this to find the minimal weight path, by relaxing each edge of the graph in $V-1$ rounds

↳ advantage over Dijkstra is that it works for graphs with negative weights, and can detect negative cycles.

negative cycle \Leftrightarrow graph changes in V^{th} relaxation round.

↳ runtime $O(VE)$

Proof of correctness (induction):

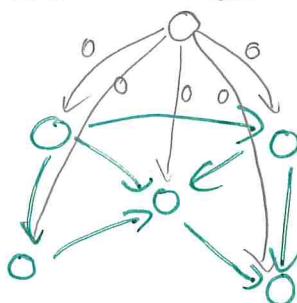
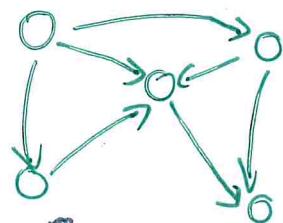
- Consider the minimal-weight path from s to some v :

$$s = v_0 \rightarrow v_1 \rightarrow \dots \rightarrow v_k = v$$

- Initially, $v_0.\text{minweight} = 0$
- After one relaxation round, $v_1.\text{minweight}$ is correct because we are on a minimal-weight path. Proceed by induction.
- At most $|V|-1$ edges, hence at most $|V|-1$ iterations.
- If the graph has a negative-weight cycle, an exception will be thrown.

Johnson's algorithm

- Used to find all shortest paths between all pairs of vertices
 - ↳ needed to calculate **betweenness centrality** - number of ~~not~~ shortest paths that use a given edge.
- We could run Dijkstra for each vertex, b in $O(VE + V^2 \log V)$, but this fails for negative weights.
- Using Bellman-Ford for each vertex would be $O(V^2 E)$.
- Johnson's algo uses BF then Dijkstra
 - ↳ build a **helper graph** with a new node s , with $d_v = w(s \rightarrow v)$



(d_{v_i}) ?
min weight from $s \rightarrow v_i$

- ↳ ~~BF~~ run BF on this graph to look for neg. weight cycles.
- ↳ recreate the original graph with weights: $w'(u \rightarrow v) = d_u + w(u \rightarrow v) - d_v$
 - ↳ $w'(u \rightarrow v) \geq 0$ by relaxation
- ↳ run Dijkstra on every vertex.
- ↳ total runtime $O(VE + V^2 \log V)$

Proof of correctness:

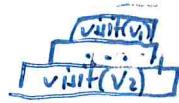
- The minweight path using w' instead of w is the same. Consider some path $v_0 \rightarrow \dots \rightarrow v_k$
 - original weights: $w'(v_0 \rightarrow v_1) + w'(v_1 \rightarrow v_2) + \dots + w'(v_{k-1} \rightarrow v_k)$
 - new weights: $= d_{v_0} + w(v_0 \rightarrow v_1) - d_{v_1} + d_{v_1} + w(v_1 \rightarrow v_2) - d_{v_2} + \dots + w(v_{k-1} \rightarrow v_k) - d_{v_k}$

i.e telescoping sum

$$\therefore \text{weight of path using } w' = \text{weight of path using old } w - d_{v_0} + d_{v_k}$$

Topological sort

- A directed acyclic graph (DAG) can always be used to represent a total ordering, such that if $v_1 \rightarrow v_2$, then v_1 is before v_2 in the total order.
- The algorithm is based on a recursive DFS, adding v to the total order when $\text{visit}(v)$ returns.
- We can use a breakpoint proof of correctness:
 - nodes are initially white
 - grey once visited
 - black once it has been added to the total order.
- Consider an edge $v_1 \rightarrow v_2$, and we have just entered $\text{visit}(v_1)$, i.e. v_1 is grey.
 - if v_2 is black, it is already in the list, so prepending v_1 will be correct
 - if v_2 is white, $\text{visit}(v_2)$ will be called, so v_2 will be prepended before v_1 . Thus v_1 will be before v_2
 - if v_2 is grey, it means we are inside $\text{visit}(v_2)$ when we called $\text{visit}(v_1)$, implying $v_2 \rightarrow v_1$. But we are in a DAG, so this contradicts $v_1 \rightarrow v_2$.
 $\hookrightarrow v_2$ cannot be grey.



Minimum Spanning trees

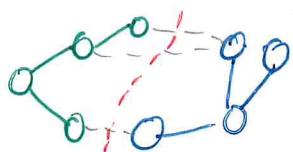
- Given a connected weighted undirected graph, the MST connects all vertices and has minimal weight.

Prim's algorithm

- Greedily builds an MST by choosing the lowest weight connector to a frontier vertex
- Very similar to Dijkstra, except:
 - need to keep track of the tree
 - we want 'distance from tree' instead of 'distance from start'
- Runtime is the same as Dijkstra, i.e. $O(E + V \log V)$.

Kruskal's algorithm

- Builds an MST by agglomerating smaller subtrees greedily.
- Edges need to be sorted by weight: runtime is $O(E \log E)$
- Despite worse runtime than Prim's, Kruskal has useful intermediate states and can be used to build clusters.
- Can be proved by considering two minimal spanning subtrees



Kruskal picks the min weight across the **cut**, so if t_1 and t_2 are MSTs within their partition, the result will also be an MST.