Kruskal's algorithm: RT: O(nlogn), Finds the minimum spanning tree in an undirected edge-weighted graph

Basic algo: sort all edges in increasing order by weight, pick smallest edge and check if it forms a cycle with spanning tree formed so far. If no cycle, include it, otherwise discard, repeat until (V-1) edges in spanning tree

Starts with each vertex in its own component and repeatedly merges two components into one by choosing a light edge that connects them, scans the set of edges in monotonically increasing order by weight, uses a disjoint-set data structure to determine whether an edge connects vertices in different components Kosaraju's Algo: RT: O(n), finds strongly connected components in graph Check every node in the graph, if one of the nodes allows you to visit every other node in the graph, then it is a strongly connected component.

1. Call DFS(graph) to compute the finish time for each vertex, 2. Compute G(t), 3. Call DFS(Gt) on vertices in decreasing order of their finish times, 4. Output vertices as separate SCC's to do this. SSC's must have a path that is cyclical

1. Create empty stack 'S', 2. Do DFS traversal of a graph, while after calling recursive DFS for adjacent vertices of a vertex, push the vertex to the stack. 3. Reverse directions of all arcs to obtain the traverse graph. 4. One by one pop a vertex from S while it;s not empty. Let the popped vertex be 'v' Take v as source and do DFS call on V. The DFS staring from v prints strongly connected components of v

Dijkstra algorithm: RT: assume O(n), finds shortest path between nodes Optimal substructure simply that subpath of any shortest path is itself a shortest path, and the shortest path length between some u and v is less than or equal to the shortest path from u to x to v. (triangle inequality)

See other side for algo Dijkstra's is a greedy algorithm

BFS, RT: O(V+E), Shortest Path (Unweighted), Shortest Cycle (Unweighted, Directed), Can find Connected Components (A set of vertices in a graph that are linked to

The proof that vertices are in this order by breadth first search goes by induction on the level number. By the induction hypothesis, BFS lists all vertices at level k-1 before those at level k. Therefore it will place into L all vertices at level k before all those of level k+1, and therefore so list those of level k before those of level k+1.

Prim's algo: RT O(nlogn), Finds the minimum spanning tree in an undirected edge-weighted graph

- 1. Create a set mstSet that keeps track of vertices already included in
- 2. Assign a key value to all vertices in the input graph, initialize all key values as ∞, Assign key value as 0 for first vertex so it's picked first. 3. While mstSet doesn't include all vertices, either A: pick a vertex u which is not in mstSet and has minimum key value, B. Include u to mstSet, C: Update key value of all adjacent vertices of u.To update the key values, iterate through all adjacent vertices. For every adjacent vertex v, if weight of edge u-v is less than the previous key value of v,

update the key value as weight of u-v a greedy algorithm that finds a minimum spanning tree for a connected weighted undirected graph. Builds on tree so A is always a tree, starts from an arbitrary "root" r, and at each step adds a light edge crossing cut (Va, V - Va) to A where Va = vertices that A is incident on. Using different data structures for representing and linearly searching array of weights to find the minimum weight edge: adj. matrix O(|V |^2), bin heap and adj lst O((|V | + |E|)log|V |) = O(|E|log|V|), Fibonacci heap and adj list O(|E| + |V|log|V|).

In the method that uses binary heaps, we can observe that the traversal is executed O(V+E) times (similar to BFS). Each traversal has operation which takes O(LogV) time. So overall time complexity is O(E+V)*O(LogV) which is O((E+V)*LogV) = O(E*LogV) (For a connected graph, V = O(E))

15 Common Recurrence Relations

$$T(n) = 2T(n/2) + n$$
 (10)

$$T(n) = 4[2T(n/8) + n/4] + 2n$$
 (11)

$$T(n) = 2^{k}T(n/(2^{k})) + kn$$
 (12)

$$T(n) = n + nlog_{2}n$$
 (13)

$$T(n) = 2log_{2}nT(1) + (log_{2}n)nT(n) = O(nlogn)$$
 (14)

Recurrence	Algorithm	Big O
$T(n/2) + \Theta(1)$	Binary Search	O(log(n))
$T(n-1)+\Theta(1)$	Sequential Search	O(n)
$2T(n/2) + \Theta(1)$	tree traversal	O(n)
$T(n-1) + \Theta(n)$	Selection Sort (n2 sorts)	$O(n^2)$
$2T(n/2) + \Theta(n)$	Mergesort	O(n log n)
$T(n-1) + T(0) + \Theta(n)$	Quicksort	$O(n^2)$

Merge Sort (Divide and conquer), RT: O(nlogn), Split array into smaller arrays of size 2, sort and combine till back to original size

def mergesort(arr): if len(arr) == 1: return arr m = len(arr) / 2, I = mergesort(arr[:m]), r = mergesort(arr[m:])

if not len(I) or not len(r) return I or r

result = [], i = j = 0,

while (len(result) < len(r)+len(l)): if |[i] < r[j]: result.append(|[i]) i += 1

else: result.append(r[j]) j += 1

if i == len(I) or j == len(r):

result.extend(I[i:] or r[j:]), break

return result

T (n) = aT (n/b) + f (n) where $a \ge 1$ and b > 1 are constants and f (n) is asymptotically positive. Case 1: if f (n) = $(n^{\log_b a - \epsilon})$ for Some ϵ then T (n) = $\Theta(n^{\log_b a})$

Case 2: if $f(n) = \Theta(n^{\log_b a} \log^k n)$ with $k \ge 1$ then $T(n) = \Theta(n^{\log_b a} \log^{k+1} n)^l$ Case 3: if f (n) = $\Omega(n^{\log_b a + \epsilon})$ with $\epsilon > 0$ and f (n) satisfies the regularity condition, then T (n) = $\Theta(f(n))$.

Regularity condition is $af(n/b) \le cf(n)$ for some c < 1 and sufficiently large n. Case on right hand side for examples

Example:

■ The algorithm solves the $T(n) = 3T(n/2) + n^2 = \Theta(n^2)$ 3 (15) $T(n) = 4T(n/2) + n^2 = \Theta(n^2 log(n))$ 2 (16) problem by dividing them $T(n) = T(n/2) + 2^n = \Theta(2^n)$ 3 (17) into four subproblems $T(n) = 2^n T(n/2) + n^n = ...$ a not constant (18) of size n/3 using 4nlog₄(n) $T(n)=16T(n/4)+n=\Theta(n^2)$ 1 (19) time, solving them $T(n) = 2T(n/2) + nlog(n) = \Theta(nlog^2n)$ 2 (20) recursively, and then $T(n) = 2T(n/2) + n/\log(n)$ non poly (21) combining the solutions in $T(n) = 2T(n/4) + n^{0.51} = \Theta(n^{0.51})$ 3 (22) T(n) = .5T(n/2) + 1/na < 1 (23) log 4(n) time. T(n) = 16T(n/4) + n! = O(n!)3 (24)

 \blacksquare a = 4, b = 3, d < (1 + e) $T(n) = \sqrt{2}T(n/2) + log(n) = \Theta(\sqrt{n})$ 1 (25) since nlogn is faster than n $T(n) = 3T(n/3) + \sqrt{n} = \Theta(n)$ 1 (26) but slower than n^(1+e). Thus. $T(n) = 4T(n/2) + cn = \Theta(nlog(n))$ 3 (27) $T(n) = O(n^{\log_3(4)})$ T(n) = T(n/2) + n(2 - cos(n)) regularity violate (28)

Format: $T(n) = a T(n/b) + \Theta(n^k (\log n)^i)$.

Quick Sort, RT: O(nlogn) Worst Case: O(n^2), Choose pivot and sort around the pivot This is a divide and conquer algorithm, first divide, conquer, then combine def quickSort(alist):

quickSortHelper(alist,0,len(alist)-1)

def quickSortHelper(alist,first,last):

if first<last: splitpoint = partition(alist,first,last)

quickSortHelper(alist,first,splitpoint-1),quickSortHelper(alist,splitpoint+1,last

def partition(alist,first,last):

pivotvalue = alist[first],leftmark = first+1, rightmark = last, done = False while not done:

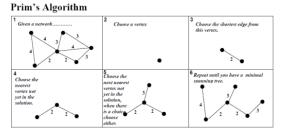
> while leftmark <= rightmark and \ alist[leftmark] <= pivotvalue: leftmark = leftmark + 1

while alist[rightmark] >= pivotvalue and \ rightmark >= leftmark: rightmark = rightmark -1

if rightmark < leftmark: done = True

Else: temp = alist[leftmark], alist[leftmark] = alist[rightmark] alist[rightmark] = temp

temp = alist[first], alist[first] = alist[rightmark], alist[rightmark] = temp return rightmark



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Random Lingo
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• NP-Complete problems to reduce from:

Independent Set: A set of k vertices that don't have any edges connecting them, This algorithm finds maximum vertices you can select that aren't connected to each other. You can also make a yes/no algorithm by asking "are there more than n vertices that are independent".

Vertex Cover: A set of vertices that cover all the edges. This algorithm finds the minimum vertices that cover all the edges. You can make this a yes/no algorithm by asking "are there fewer than n vertices that are independent". Vertex Cover and Independent Set are closely related. (With v being the number of vertices) If you have an n size independent set, you'd have a v-n size vertex cover. And it'd just be all other vertices. And if you have an n size vertex cover, you'd have a v-n size independent set.

Clique: Literally just "is there a complete subgraph of size k?" Set of vertices all connected

Rudrata Path (directed or undirected): A path that visits every vertex once

Rudrata Cycle (directed or undirected): A cycle that visits every vertex once, Find median in linear time

K-selection: Find x-th element in linear time

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Topological Sort :RT: O(n), Organizes a DAG with all edges pointing from left to right 1. Call DFS to compute finishing times for all vertices, as each vergtex finishes, insert it onto the front of a linked list, and return the linked list of vertices, this is \Theta(V+E). Proof: show that if (u, v) \in E then f[v] < f[u]: when we explore (u, v) what are the colors of u and v, if us is grey then v cannot be grey because then v would be ancestor of u therefore (u, v) is a black edge, which is a contradiction of DAG, if v is white than becomes descendant of u and parenthesis theorem work
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Dijsktras algo
def minDistance(self, dist, sptSet):
     min = sys.maxint
     for u in range(self.V):
        if dist[u] < min and sptSet[u] == False:
          min = dist[u]
          min index = u
     return min index
def dijkstra(self, src):
     dist = [sys.maxint] * self.V , dist[src] = 0, sptSet = [False] * self.V
     for cout in range(self.V):
        x = self.minDistance(dist, sptSet), sptSet[x] = True
        for y in range(self.V):
          if self.graph[x][y] > 0 and sptSet[y] == False and \
          dist[y] > dist[x] + self.graph[x][y]:
                dist[y] = dist[x] + self.graph[x][y]
     Return dist
```

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Kruskal algo
 def KruskalMST(self):
     result = [] # This will store the resultant MST
     i = 0
     self.graph = sorted(self.graph, key=lambda item: item[2]) #sort edges in non dec
     parent = [],
                   rank = ∏
     for node in range(self.V): parent.append(node),
                                                             rank.append(0)
     while e < self.V - 1:
       u, v, w = self.graph[i], i = i + 1 ,x = self.find(parent, u), y = self.find(parent, v)
       if x != y: e = e + 1, result.append([u, v, w]), self.union(parent, rank, x, y)
     minimumCost = 0
     for u, v, weight in result:
       minimumCost += weight
     Return minimumCost
```

def dfs(graph, start):
 visited, stack = set(), [start]
 while stack:
 vertex = stack.pop()
 if vertex not in visited:
 visited.add(vertex)
 stack.extend(graph[vertex] - visited)
 return visited

Greedy algorithms- algorithms that make the optimal choice at each step as it tries to find the overall optimal way to solve the problem. This can backfire when the solution is hidden behind a less optimal path

If both of the properties below are true, a greedy algorithm can be used

to solve the problem.

Greedy choice property: A global (overall) optimal solution can be

reached by choosing the optimal choice at each step.

Optimal substructure: A problem has an optimal substructure if an optimal solution to the entire problem contains the optimal solutions to the sub-problems.

This can be made by identifying an optimal substructure or subproblem

Memoization (Top Down): memoized program for a problem is like the recursive version with a small modification that looks into a lookup table before computing solutions. We initialize a lookup array with all initial values as NIL. Whenever we need the solution to a subproblem, we first look into the lookup table. If the precomputed value is there then we return that value, otherwise, we calculate the value and put the result in the lookup table so that it can be reused later.

Dynamic Programming- solving a problem by recursively breaking it down into simpler subproblems as the optimal solution to the overall problem depends on the optimal solution to the individual subproblems. Each subproblem is done once and the solution is stored, this makes things less complex and is good for optimization. This fails when there isn't an overlapping sub problem, or an optimal sub-structure, for example shortest path can be solved using DP but longest path can't solved, because longest path doesn't hold optimal sub-structure property. Dynamic Programming is not useful when there are no common (overlapping) subproblems because there is no point storing the solutions if they are not needed again. For example, Binary Search doesn't have common subproblems Not going to give an instance of a problem and then slave to solve it (building a table, bottom up)

Tabulation (Bottom Up): The tabulated program for a given problem builds a table in bottom-up fashion and returns the last entry from the table. For example, for the same Fibonacci number, we first calculate fib(0) then fib(1) then fib(2) then fib(3), and so on. So literally, we are building the solutions of subproblems bottom-up.

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\begin{array}{c|c} \text{log}_{a} = \frac{\log_{a} n}{\log_{a} c} \\ \text{log}_{a} = \frac{\log_{a} n}{\log_{a} c} \\ \text{log}_{a} = 1 \\ \text{log}_{a} =
```



Checklist

- ◆ Dijktra's Algo
- → Runtime: Assume O(n2)
- Use: Find the shortest distance to each node in a graph
- Kosaraju's Algo
- → Runtime: O(n)
- Topological Sort
- → Runtime: O(n)
- Use: Organizes a DAG with all edges pointing from left to right
- Kruskal's Algorithm
- → Runtime: O(nlogn)
- Use: Finds the minimum spanning tree in an undirected edge-weighted graph
- Prim's Algorithm
- · Runtime: O(nlogn)
- Use: Finds the minimum spanning tree in an undirected edge-weighted graph
- BFS
- → Runtime: O(V+E)
- Shortest Path (Unweighted)
- Shortest Cycle (Unweighted, Directed)
- Can find Connected Components (A set of vertices in a graph that are linked to each other by paths)
- DFS
- → Runtime: O(V+E)
- Shortest Path (Unweighted)
- NP-Complete problems to reduce from:
- → Independent Set
- A set of k vertices that don't have any edges connecting them
- This algorithm finds maximum vertices you can select that aren't connected to each other. You can also make a yes/no algorithm by asking "are there more than n vertices that are independent".
- Vertex Cover
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- Vertex Cover and Independent Set are closely related. (With v being the number of vertices) If you have an n size independent set, you'd have a v-n size vertex cover. And it'd just be all other vertices. And if you have an n size vertex cover, you'd have a v-n size independent set.
- Clique
- Literally just "is there a complete subgraph of size k?"
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- K-selection
- → Find x-th element in linear time
- Merge Sort (Divide and conquer)
- → O(nlogn)
- Split array into smaller arrays of size 2, sort and combine till back to original size
- Quick Sort
- → O(nlogn)
- → Worst Case: O(n^2)
- · Choose pivot and sort around the pivot
- Masters Theorem
- https://www.nayuki.io/page/master-theorem-solver-javascript
- The algorithm solves the problem by dividing them into four subproblems of size n/3 using 4nlog_4(n) time, solving them recursively, and then combining the solutions in log_4(n) time.
- a = 4, b = 3, d < 1 +\e since nlogn is faster than n but slower than n^{1+} . Thus, $T(n)=O(n^{{\log_3(4)}})$