

Datastructure

Data Structures

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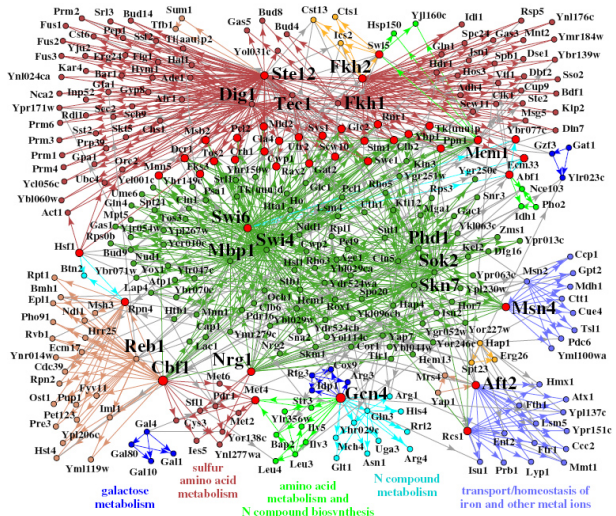
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Contents

- 7 Graphs
 - Representation
 - Graph traversal
 - Disjoint Sets, ADT Union-Find
 - Minimal Spanning Trees
 - Shortest Paths
 - Topological Sort

transcription regulatory interactions



Directed network modules, Palla et al. New Journal of Physics, 2007.

zie ook college SNACS

graph definition

zie FoCS en Algoritmiëk!

Definition

A *graph* is a pair $G = (V, E)$ where:

- V is a set of *vertices*, or *nodes*
- $E \subseteq V \times V$ is a set *edges*, or *arcs*, *lines*

directed / undirected

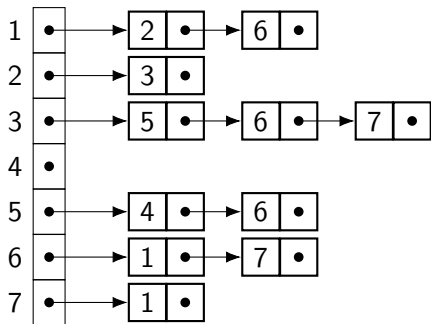
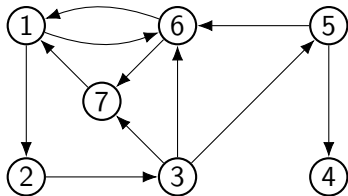
vertices / edges can have labels (string, number)

complexity in $|V|$ and $|E|$ $|E| \leq |V|^2$

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adjacency lists



representation

space matrix $O(|V|^2)$
lists $O(|V| + |E|)$ $|E| \leq |V|^2$

data science / network analysis
huge graphs, few bits per node
sparse graphs

operations 'abstract'

- $(u, v) \in E$
- all outgoing edges
- all incoming edges

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graph traversal

DepthFS pre-order stack

BreadthFS level-order queue

– tree-traversal + marking nodes

DFS nodes can be twice on stack

structure *spanning tree*

tree, forward, back, cross

depth first search

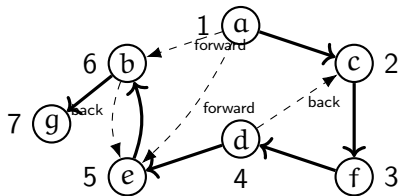
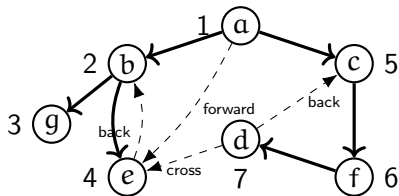
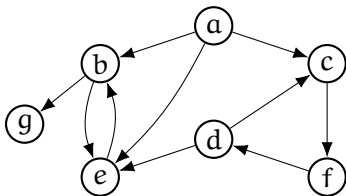
Recursive DFS

```
DFS(v)
  visit(v)
  mark(v)
  for each w adjacent to v
  do if w is not marked
    then DFS(w)
    fi
  od
end // DFS
```

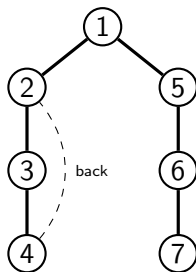
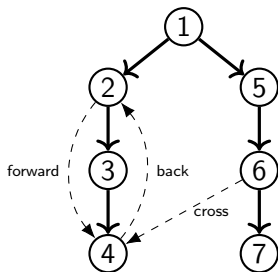
Iterative DFS

```
// start with unmarked nodes
S.push(init)
while S is not empty
do v = S.pop()
  if v is not marked
  then mark v
    for each edge from v to w
    do if w is unmarked
      then S.push(w)
      fi
    od
  fi
od
od
```

dfs tree (directed)



dfs edges



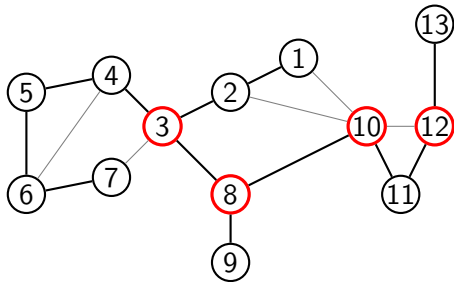
applications of DFS

A DFS traversal itself and the *forest-like representation* of the graph it provides have proved to be extremely helpful for the development of efficient algorithms for checking many important properties of graphs. Note that the DFS yields *two orderings of vertices*: the order in which the vertices are reached for the first time (*pushed onto* the stack) and the order in which the vertices become dead ends (*popped off* the stack). These orders are qualitatively different, and various applications can take advantage of either of them.

[Levitin, Design & Analysis of Algorithms]

- articulation points
- topological sorting

articulation points



articulation points

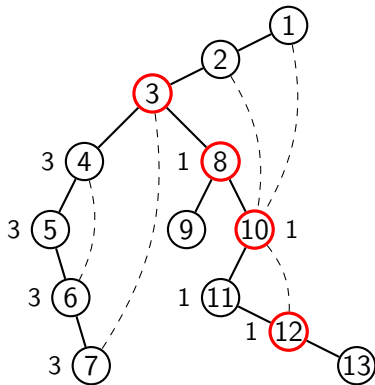
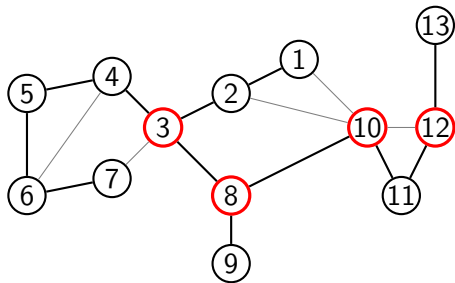
construct DFS tree for graph

Lemma

*A vertex v is an **articulation point** if either*

- *v is the root, and has two or more children, or*
- *v has a (strict) subtree, and no node in the subtree has a back edge that reaches above v .*

dfs and articulation points

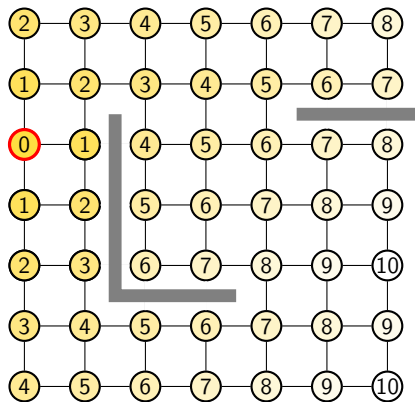


breadth-first search

Iterative BFS

```
// Q is a queue of vertices
// start with unmarked nodes
Q.enqueue(init)
dist[init] = 0
while Q is not empty
do v = Q.dequeue()
  newdist = dist[v] + 1
  for all edges from v to w
  do if w is not marked
    then Q.enqueue(w)
      mark w
      dist[w] = newdist
    fi
  od
od
```

bfs: 'floodfill'



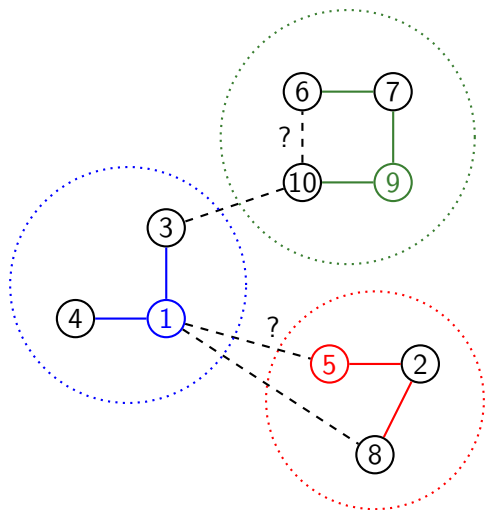
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Algorithms from the Book ☒

124	union-find	Galler and Fischer
116	Knuth-Morris-Pratt	pattern matching
99	Blum,Floyd,Pratt,Rivest,Tarjan	median
94	binary search	
88	Floyd-Warshall	all-pairs shortest path
84	Euclidean algorithm	greatest common divisor (GCD)
75	quicksort	Tony Hoare
63	Huffman coding	data compression
55	Schwartz-Zippel lemma	polynomial identity
54	Miller-Rabin	primality test
48	depth first search	
45	sieve of Eratosthenes	primes
45	Dijkstra	shortest path
44	Gentry	homomorphic encryption
43	Cooley-Tukey	fast Fourier transform

application Union-Find



Union-Find

- **INITIALIZE**: construct the initial partition; each component consists of a singleton set $\{d\}$, with $d \in D$.
- **FIND**: retrieves the name of the component, i.e., $\text{FIND}(u) = \text{FIND}(v)$ iff u and v belong to the same set in the partition.
- **UNION**: given two elements u and v the sets they belong to are merged. Has no effects when u and v already belong to the same set.

Usually it is assumed that u, v are representatives, i.e., names of components, not arbitrary elements.

name array

1	2	3	4	5	6	7	8	9	10		
1	2	3	4	5	6	7	8	9	10		find
1	2	3	4	5	6	7	8	9	10		
1	2	1	4	5	6	6	5	9	9		find

UNION(9, 6)

1	2	3	4	5	6	7	8	9	10		
1	2	1	4	5	9	9	5	9	9		find

lists

1	2	3	4	5	6	7	8	9	10	
1	2	3	4	5	6	7	8	9	10	find
1	2	3	4	5	6	7	8	9	10	next
1	1	1	1	1	1	1	1	1	1	size

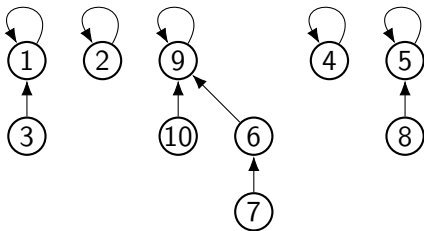
1	2	3	4	5	6	7	8	9	10	
1	2	1	4	5	6	6	5	9	9	find
3	2	1	4	8	7	6	5	10	9	next
2	1	.	1	2	2	.	.	2	.	size

UNION(9, 6)

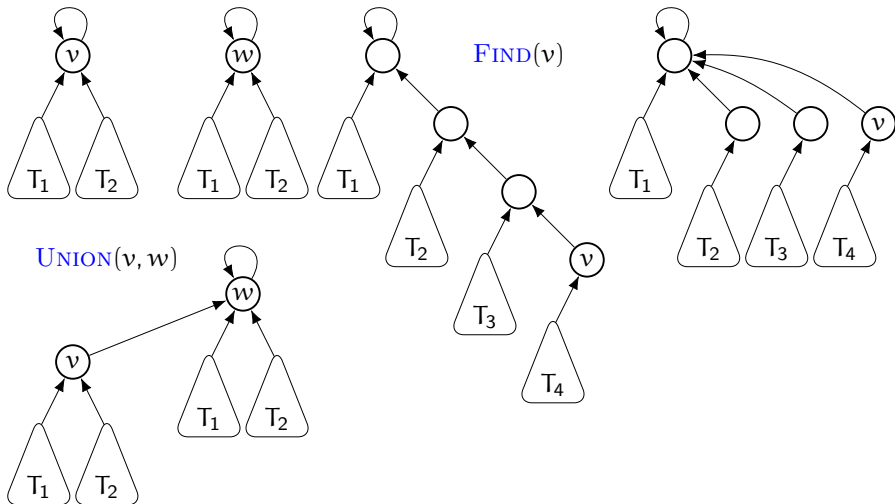
1	2	3	4	5	6	7	8	9	10	
1	2	1	4	5	9	9	5	9	9	find
3	2	1	4	8	10	6	5	7	9	next
2	1	.	1	2	.	.	.	4	.	size

union-find with path-compression

1	2	3	4	5	6	7	8	9	10	
1	2	1	4	5	9	6	5	9	9	parent
2	1	.	1	2	.	.	.	3	.	height



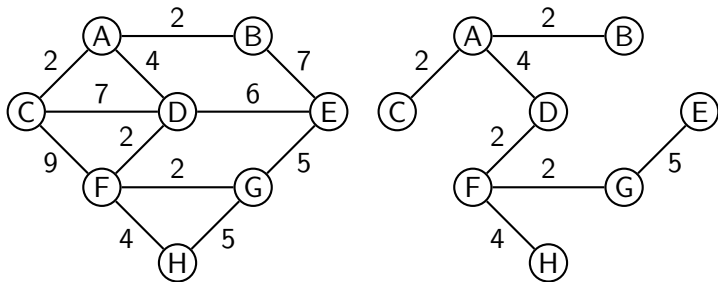
path compression



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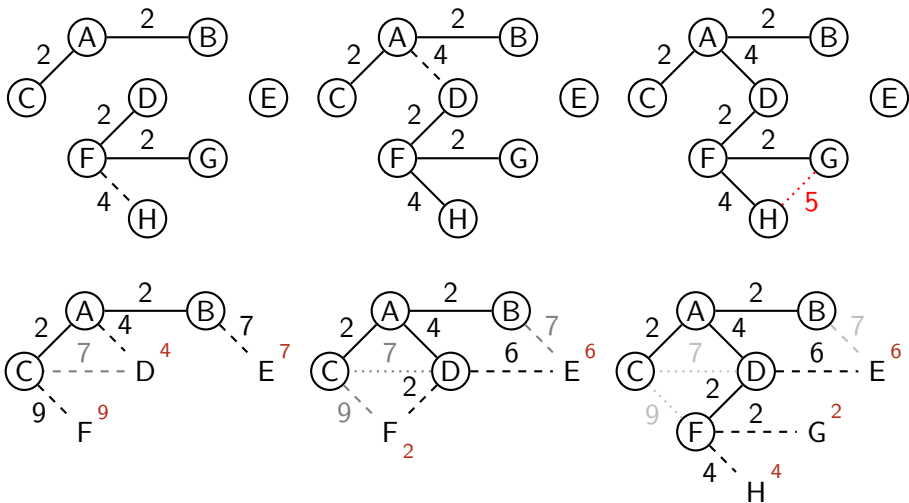
minimal spanning tree



Definition (Minimal spanning tree of weighted graph)

A tree containing all nodes of the graph, with minimal total sum of edge weights

minimal spanning tree Kruskal vs. Prim



minimal spanning tree - Kruskal

Kruskal (high level)

```
repeat
  consider edge with smallest weight
  if it does not yield a cycle
  then add it to the tree
  else discard the edge
fi
until no edges left
```


minimal spanning tree - Prim

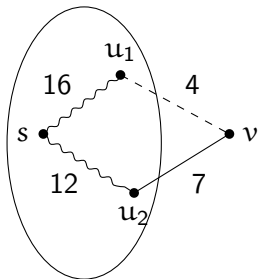
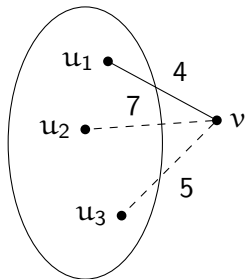
Prim (high level)

```
start with single node
repeat
  consider edge with smallest weight
    that connects node in tree with one outside
  add new node+edge to the tree
until all nodes in tree
```

optimization: for each node outside tree select minimal weight connection to tree

Prim vs Dijkstra

growing tree : keep best candidate connection

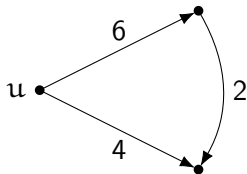


Prim

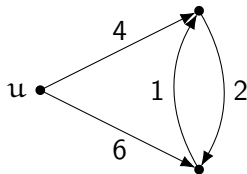
Prim

```
cost[source] = 0           // infinite for other vertices
parent[source] = 0        // code for the root
PQ = V                    // all vertices
while PQ is not empty
do u is vertex in PQ with minimal cost[u]
    remove u from PQ
    for each edge (u,v)
        do if length(u,v) < cost[v]
            then cost[v] = length(u,v)
                parent[v] = u
            fi
    od
od
```

directed graphs not supported



Prim fails



Kruskal fails

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Dijkstra

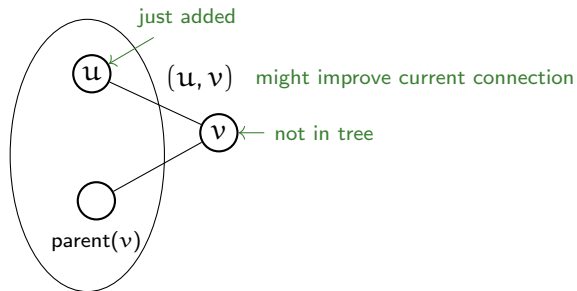
Dijkstra

```
dist[source] = 0    // infinite for other vertices
parent[source] = 0 // code for the root
Q = V              // start with all vertices
while Q is not empty
do u is vertex in Q with minimal dist[u]
  remove u from Q
  for each edge (u,v)
  do if dist[u] + length(u,v) < dist[v]
    then dist[v] = dist[u] + length(u,v)
      parent[v] = u
    fi
  od
od
```

shortest path from *fixed source* node to all other nodes

Prim vs Dijkstra

growing tree



Prim

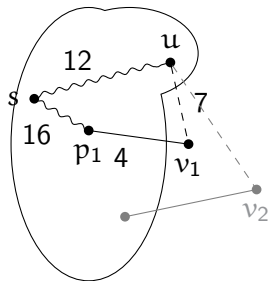
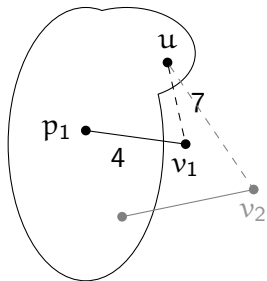
single edge
undirected
negative OK
random start

Dijkstra

length complete path
(un)directed
non-negative
specific source (to all other)

Prim vs Dijkstra update

new node added u : update nodes v_i not in tree



complexity

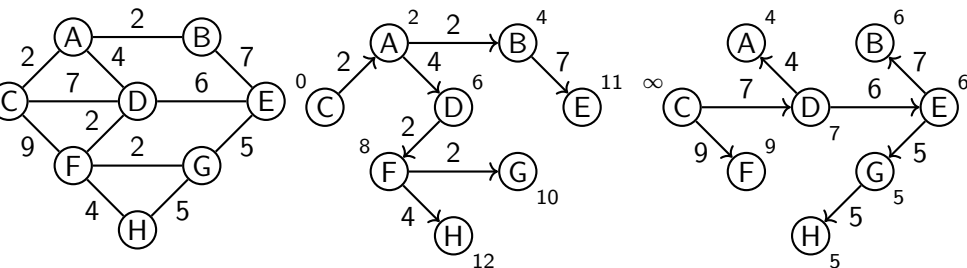
adjacency lists

$$|V| \leq |E| \leq |V|^2$$

		heap	better	
minimal	$ V ^2$	$ V \cdot \lg V $	$ V \cdot \lg V $	findmin
each edge	$ E $	$ E \cdot \lg V $	$ E $	decreasekey
	$ V ^2$	$ E \cdot \lg V $	$ E + V \cdot \lg V $	

heap: better for small number of edges
(or use better priority queue)

distance vs. bottleneck



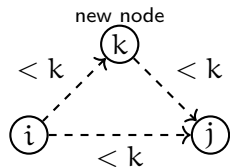
all pairs

$$L^k(i, j) = \min(L^{k-1}(i, j), L^{k-1}(i, k) + L^{k-1}(k, j))$$

nodes $1, 2, \dots, n$

L^k path via nodes $\leq k$

L^0 only single edges \sim adjacency matrix



all pairs shortest distance

$$L^k(i, j) = \min(L^{k-1}(i, j), L^{k-1}(i, k) + L^{k-1}(k, j))$$

Floyd-Warshall

```
// initially dist equals the adjacency matrix
for each edge (i,j)
do next[i,j] = i
od
for k from 1 to n
do for i from 1 to n
do for j from 1 to n
do if dist[i,k] + dist[k,j] < dist[i,j]
then dist[i,j] = dist[i,k] + dist[k,j]
next[i,j] = next[i,k]
fi
od
od
od
```

Floyd example

partial result A^3 ,

$$A^3 = \begin{array}{c} \\ 1 \\ 2 \\ 3 \\ 4 \end{array} \begin{array}{cccc} & 1 & 2 & 3 & 4 \\ \left(\begin{array}{cccc} 0 & 2 & 1 & 6 \\ 3 & 0 & 1 & 4 \\ \boxed{4} & 1 & 0 & 5 \\ -2 & 0 & -1 & . \end{array} \right)$$

and distances via node 4

$$\begin{array}{c} \\ 1 \\ 2 \\ 3 \\ 4 \end{array} \begin{array}{cccc} & 1 & 2 & 3 & 4 \\ \left(\begin{array}{cccc} . & 6+0 & 6-1 & 6 \\ 4-2 & . & 4-1 & 4 \\ \boxed{5-2} & 5+0 & . & \boxed{5} \\ \boxed{-2} & 0 & -1 & . \end{array} \right)$$

path reconstruction

Path-reconstruction

```
Path(u, v)
  if next[u][v] = null
  then return []
  fi
  path = [v]
  while u != v
  do v = next[u][v]
    path.insert_at_end(v)
  od
  return path
```

transitive closure

Warshall

```
// initially conn equals the adjacency matrix
// with additionally 1=true on the diagonal
for k from 1 to n
do for i from 1 to n
  do for j from 1 to n
    do conn[i,j] = conn[i,j] or ( conn[i,k] and conn[k,j] )
    od
  od
od
```

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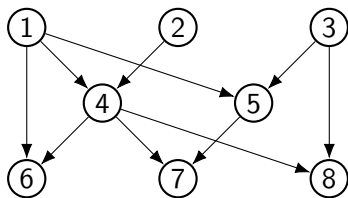
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topological sorting

Let $G = (V, E)$ be a directed graph.

Definition

A *topological ordering* [or *sort*] of G is an ordering (v_1, \dots, v_n) of V , such that if $(v_i, v_j) \in E$ then $i < j$.

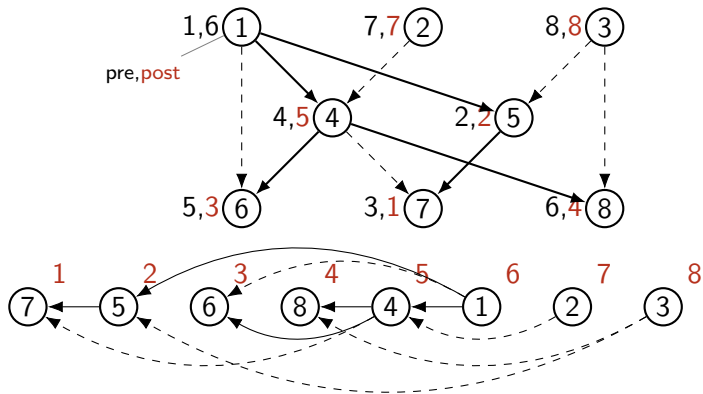


topological sorting

finding a topological sort:

- depth-first search post-order
- source removal Kahn's algorithm
 - 1 pick node without incoming edges
 - 2 remove that node with outgoing edges. go to step 1.

DFS application: topological sort



end.