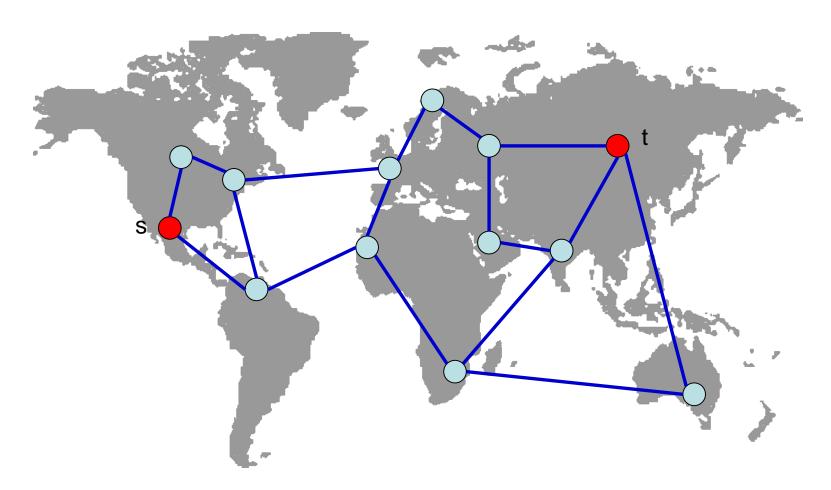
Fundamental Algorithms Chapter 4: Shortest Paths

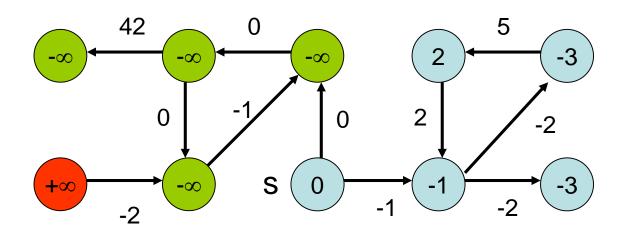
Christian Scheideler WS 2017



Central question: Determine fastest way to get from s to t.

Shortest Path Problem:

- directed/undirected graph G=(V,E)
- edge costs c:E→ℝ
- SSSP (single source shortest path): find shortest paths from a source node to all other nodes
- APSP (all pairs shortest path): find shortest paths between all pairs of nodes



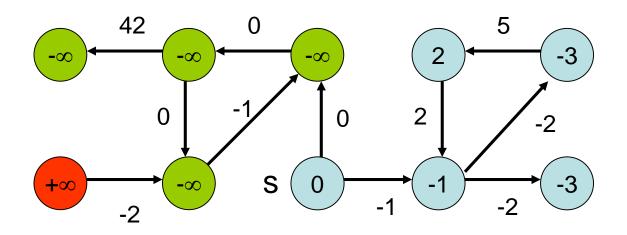
 $\mu(s,v)$: distance between s and v

$$\mu(s,v) = \begin{cases} \infty & \text{no path from s to } v \\ -\infty & \text{path of arbitrarily low cost from s to } v \end{cases}$$

$$\min\{c(p) \mid p \text{ is a path from s to } v\}$$

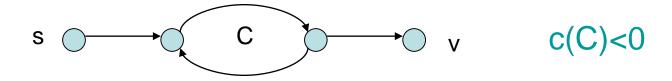
$$\text{Chapter 4}$$

29.11.2017 Chapter 4



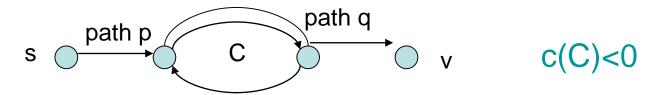
When is the distance $-\infty$?

If there is a negative cycle:



Negative cycle necessary and sufficient for a distance of $-\infty$.

Negative cycle sufficient:



Cost for i-fold traversal of C:

$$c(p) + i \cdot c(C) + c(q)$$

For $i \rightarrow \infty$ this expression approaches $-\infty$.

Negative cycle necessary and sufficient for a distance of -∞.

Negative cycle necessary:

- I: minimal cost of a simple path from s to v
- suppose there is a non-simple path p from s to v with cost c(r)<l
- p non-simple: continuously remove a cycle C till we are left with a simple path
- since c(p) < I, there must be a cycle C with c(C)<0

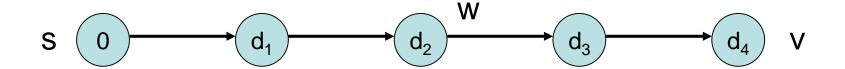
Shortest Paths in Arbitrary Graphs

General Strategy:

- Initially, set d(s):=0 and d(v):=∞ for all other nodes
- Visit nodes in an order that ensures that at least one shortest path from s to every v is visited in the order of its nodes
- For every visited v, update distances to nodes w with (v,w)∈E, i.e., d(w):= min{d(w), d(v)+c(v,w)}

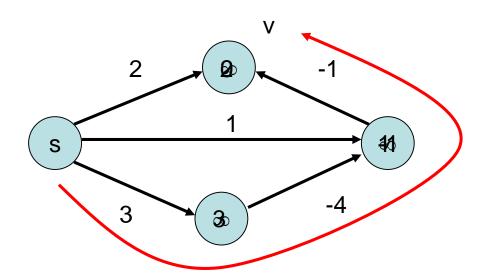
Consider graphs with arbitrary edge costs.

Problem: visit nodes along a shortest path from s to v in the right order



Dijkstra's algorithm cannot be used in this case any more.

Example:



Node v has wrong distance value!

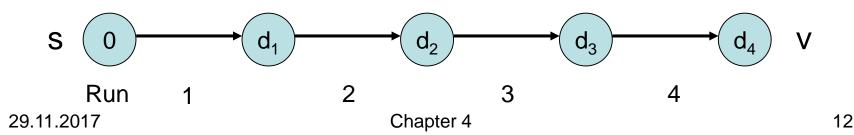
Lemma 4.1: For every node v with $\mu(s,v)>-\infty$ there is a simple path (without cycle!) from s to v of length $\mu(s,v)$.

Proof:

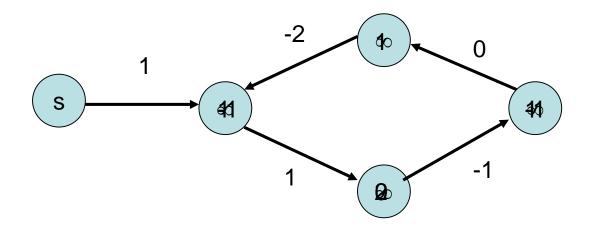
- Path with cycle of length ≥0: removing the cycle does not increase the path length
- Path with cycle of length <0: distance from s is -∞!

Conclusion: (graph with n nodes)
For every node v with $\mu(s,v) > -\infty$ there is a shortest path along <n nodes to v.

Strategy: visit (n-1)-times all nodes in the graph and update distances. Then all shortest paths have been considered.



Problem: detection of negative cycles



Conclusion: in a negative cycle, distance of at least one node keeps decreasing in each round, starting with a round <n

Lemma 4.2:

- No decrease of a distance in a round (i.e., d[v]+c(v,w)≥d[w] for all w):
 Done because d[w]=μ(s,w) for all w
- Decrease of a distance even in n-th round
 (i.e., d[v]+c(v,w)<d[w] for some w):
 There are negative cycles for all of these nodes, so node w has distance μ(s,w)=-∞. If this is true for w, then also for all nodes reachable from w.

Proof: exercise

```
Procedure BellmanFord(s: Nodeld)
  d=<\infty,...,\infty>: NodeArray of \mathbb{R}\cup\{-\infty,\infty\}
  parent=<1,..., \(\perp >: \) NodeArray of NodeId
  d[s]:=0; parent[s]:=s
  for i:=1 to n-1 do // update distances for n-1 rounds
     forall e=(v,w)∈E do
        if d[w] > d[v] + c(e) then // better distance?
           d[w]:=d[v]+c(e); parent[w]:=v
  forall e=(v,w) \in E do // still better in n-th round?
     if d[w] > d[v] + c(e) then infect(w)
Procedure infect(v) // set -∞-distance starting with v
  if d[v] > -\infty then
     d[v]:=-∞
     forall (v,w) \in E do infect(w)
```

Runtime: O(n·m)

Improvements:

- Check in each update round if we still have d[v]+c[v,w]<d[w] for some (v,w)∈E.
 No: done!
- Visit in each round only those nodes w with some edge (v,w)∈E where d[v] has decreased in the previous round.

Assumption: graph with arbitrary edge costs, but no negative cycles

Naive Strategy for a graph with n nodes: run n times Bellman-Ford Algorithm (once for every node as the source)

Runtime: O(n² m)

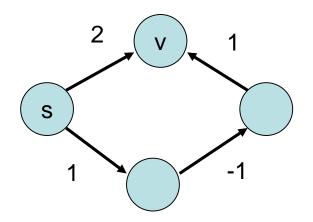
Better Strategy: Reduce n Bellman-Ford applications to n Dijkstra applications

Problem: we need non-negative edge costs

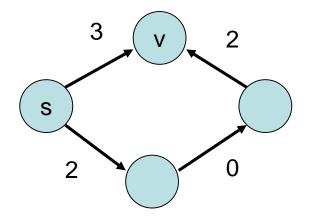
Solution: convert edge costs into nonnegative edge costs without changing the shortest paths (not so easy!)

Counterexample to additive increase by c:





cost +1 everywhere



: shortest path

- The reduced cost of e=(v,w) is:
 r(e) := c(e) + φ(v) φ(w)

Lemma 4.3: Let p and q be paths connecting the same endpoints in G. Then for every potential ϕ : r(p) < r(q) if and only if c(p) < c(q).

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Proof: Let $p=(v_1,...,v_k)$ be an arbitrary path and $e_i=(v_i,v_{i+1})$ for all i. It holds:

$$r(p) = \sum_{i} r(e_{i})$$

$$= \sum_{i} (\phi(v_{i}) + c(e_{i}) - \phi(v_{i+1}))$$

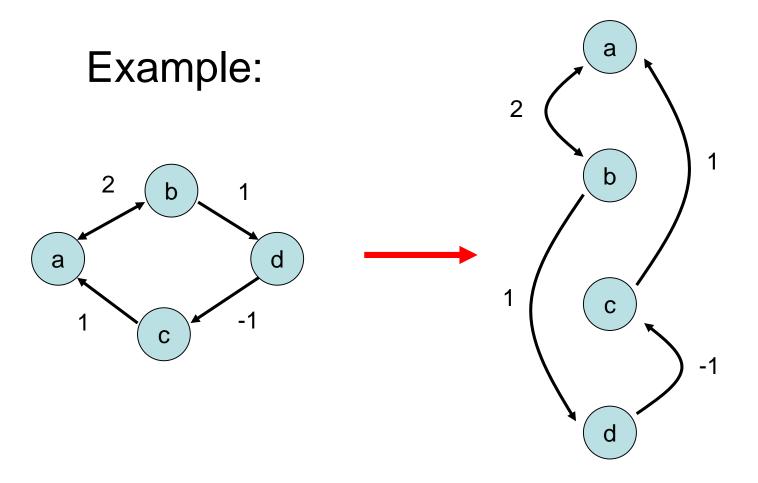
$$= \phi(v_{1}) + c(p) - \phi(v_{k})$$

Lemma 4.4: Suppose that G has no negative cycles and that all nodes can be reached from s. Let $\phi(v)=\mu(s,v)$ for all $v\in V$. With this ϕ , $r(e)\geq 0$ for all e.

Proof:

- According to our assumption, $\mu(s,v) \in \mathbb{R}$ for all v
- We know: for every edge e=(v,w), μ(s,v)+c(e)≥μ(s,w) (otherwise, we have a contradiction to the definition of μ!)
- Therefore, $r(e) = \mu(s,v) + c(e) \mu(s,w) \ge 0$

- Create new node s and new edges (s,v) for all v in G with c(s,v)=0 (all nodes reachable!)
- 2. Compute $\mu(s,v)$ using Bellman-Ford and set $\phi(v):=\mu(s,v)$ for all v
- 3. Compute the reduced costs r(e)
- Compute for all nodes v the distances μ(v,w) using Dijkstra with the reduced costs on graph G without node s
- 5. Compute the correct distances $\mu(v,w)$ via $\mu(v,w):=\bar{\mu}(v,w)+\phi(w)-\phi(v)$



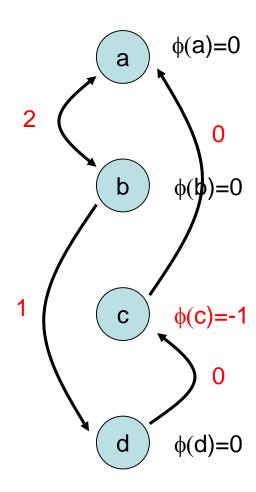
Step 1: create new source s

 $\phi(a)=0$ Step 2: apply Bellman-Ford to s $\phi(\phi)=0$ S $\phi(c)=-1$ $\phi(d)=0$

Step 3: compute r(e)-values

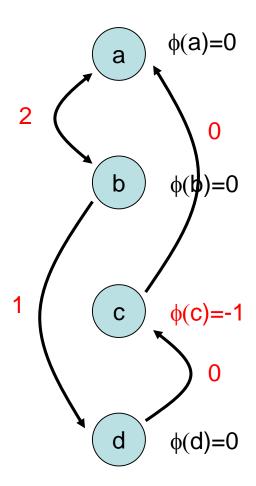
The reduced cost of e=(v,w) is:

$$r(e) := \phi(v) + c(e) - \phi(w)$$



Step 4: compute all distances $\bar{\mu}(v,w)$ via Dijkstra

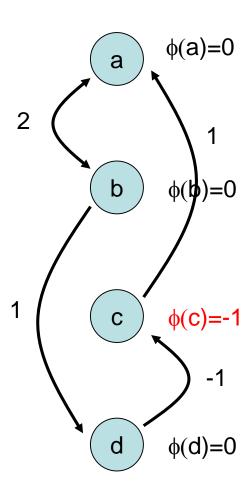
μ	а	b	С	d
a	0	2	3	3
b	1	0	1	1
С	0	2	0	3
d	0	2	0	0



Step 5: compute correct distances via the formula

$$\mu(v,w)=\bar{\mu}(v,w)+\phi(w)-\phi(v)$$

μ	а	b	С	d
а	0	2	2	3
b	1	0	0	1
С	1	3	0	4
d	0	2	-1	0



Runtime of Johnson's Method:

```
O(T_{\text{Bellman-Ford}}(n,m) + n \cdot T_{\text{Dijkstra}}(n,m))
= O(n \cdot m + n(n \log n + m))
= O(n \cdot m + n^2 \log n)
```

when using Fibonacci heaps.

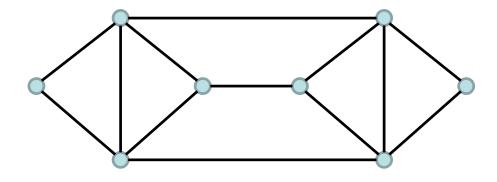
- Problem with the runtime bound: m can be quite large in the worst case (up to ~n²)
- Can we significantly reduce m if we are fine with computing approximate shortest paths?

Definition 4.5: Given an undirected graph G=(V,E) with edge costs $c:E\to\mathbb{R}$, a subgraph $H\subseteq G$ is an (α,β) -spanner of G iff for all $u,v\in V$, $d_H(u,v)\leq \alpha\cdot d_G(u,v)+\beta$

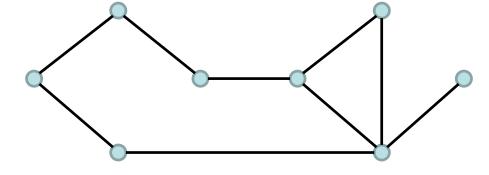
- d_G(u,v): distance of u and v in G
- α: multiplicative stretch
- β: additive stretch

Example: all edge costs are 1

G



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Consider the following Greedy algorithm by Althöfer et al. (Discrete Computational Geometry, 1993):

Theorem 4.6: For any $k \ge 1$, $|E(H)| = O(n^{1+1/k})$ and the graph H constructed by the Greedy algorithm is a (2k-1,0)-spanner.

Thorup and Zwick have shown that for any graph G with non-negative edge costs a structure related to H can be built in expected time $O(k \cdot m \cdot n^{1/k})$, which implies that we can then solve the (2k-1)-approximate APSP in time $O(k \cdot m \cdot n^{1/k} + n^{2+1/k})$.

We will get back to that when we talk about distance oracles.

Proof of Theorem 4.6:

Lemma 4.7: H is a (2k-1,0)-spanner of G.

Proof:

- Consider any edge {u,v}∈E(G)\E(H).
- Since {u,v} was rejected by the algorithm, d_H(u,v)≤(2k-1)·c(u,v).
- Consider now any shortest path p from node a to b in G.
- For every edge {u,v} in p, there is a path of length at most (2k-1)·c(u,v) in H.
- Replacing each edge {u,v} in p by this path results in a path from a to b in H of length at most (2k-1)·c(p).

Proof of Theorem 4.6:

Lemma 4.8: Let C be any cycle in H. Then |C|>2k. Proof:

- Assume that there is a cycle C of length at most 2k in H.
- Let {u,v} be the last edge in C that was added by the algorithm.
- Clearly, {u,v} has the largest cost of all edges in C.
- Also, when {u,v} was considered, (2k-1)·c(u,v)<d_H(u,v) as otherwise {u,v} would not have been added to H.
- However, since C\{u,v\} results in a path of length at most (2k-1)·c(u,v) from u to v, d_H(u,v)≤(2k-1)·c(u,v), leading to a contradiction.
- Hence, the lemma is true.

Proof of Theorem 4.6:

Lemma 4.8 implies that H has a girth (defined as the minimum cycle length in H) of more than 2k.

Lemma 4.9: Let H be a graph of size n with girth >2k. Then $|E(H)|=O(n^{1+1/k})$. Proof:

- Let H be be any graph with girth >2k and at least n+2n^{1+1/k} edges.
- Repeatedly remove any node from H of degree at most node any edges incident to that node, until no such node exists.
- The total number of edges removed in this way is at most $n \cdot (n^{1/k}+1)$.
- Hence, we obtain a subgraph H´ of H of minimum degree more than n¹/k with at least n¹+¹/k edges connecting at most n nodes.
- Exercise: show that there cannot be a graph G of size n with girth >2k and minimum degree more than $\lceil n^{1/k} \rceil$.
- Thus, H' must have a girth of at most 2k, and therefore also the original graph H. This, however, is a contradiction.

If we restrict ourselves to unweighted graphs (i.e., all edges have a cost of 1), we can also construct good additive spanners.

Theorem 4.10: Any n-node graph G has a (1,2)-spanner with O(n^{3/2} log n) edges.

Proof:

We first need the notion of hitting sets.

Definition 4.11: Given a collection M of subsets of V, a subset S⊆V is a hitting set of M if it intersects every set in M.

Lemma 4.12: Let $M=(S_1,...,S_n)$ be a collection of subsets of $V=\{1,...,n\}$ with $|S_i| \ge R$ for all i. There is an algorithm running in $O(nR \log n + (n/R)\log^2 n)$ time that finds a hitting set S of M with $|S| \le (n/R) \ln n$.

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• Assume w.l.o.g. that $|S_i|=R$ for all i. Run the following greedy algorithm:

```
 \begin{split} |S| := & \varnothing \\ \text{for each } 1 \leq j \leq n, \text{ keep a counter } \mathbf{c}(j) = |\{S_i \in M \colon j \in S_i\}| \\ \text{while } M \neq \varnothing \text{ do} \\ \text{k} := & \operatorname{argmax}_j \mathbf{c}(j) \\ S := & S \cup \{k\} \\ \text{remove any subsets from M containing k and } \\ \text{update the counters } \mathbf{c}(j) \text{ accordingly} \\ \end{split}
```

• To obtain the runtime, we store the counts c(j) in a data structure that can support the following operations in $O(\log n)$ time: insert an element, return element j with maximum c(j), decrement a given c(j).

```
 \begin{split} |S| &:= \varnothing \\ \text{for each } 1 \leq j \leq n, \text{ keep a counter } c(j) = |\{S_i \in M: j \in S_i\}| \\ \text{while } M \neq \varnothing \text{ do} \\ & k := \underset{j \in S}{\operatorname{argmax}_j c(j)} \\ & S := S \cup \{k\} \\ \text{remove any subsets from M containing k and update the counters } c(j) \text{ accordingly} \end{split}
```

- total number of inserts: n because of n counters
 → runtime O(n log n)
- total number of decrements: nR because each of the n sets contains just R elements and each of them can only cause one decrement
 → runtime O(nR log n)
- total number of argmax calls: depends on number of iterations of while loop

Proof of Lemma 4.12 (continued):

- We still need an upper bound on |S| (which gives an upper bound on while loop)
- Let m_j be the number of sets remaining in M after j passes of the while loop. Then $m_0=n$.
- Let k_i be the j-th element added to S, so $m_i=m_{i-1}-c(k_i)$.
- Just before we add k_j , the sum of c(j) over all $j \in V \setminus \{k_1, \dots, k_{j-1}\}$ must be $m_{j-1}R$, so $c(k_j)$ must be at least the average count, which is $m_{j-1}R/(n-j+1)$.
- Therefore,

```
\begin{split} m_j &\leq (1-R/(n-j+1)) \cdot m_{j-1} \leq n \ \Pi_{l=0}^{j-1} \ (1-R/(n-l)) \\ &< n \cdot (1-R/n)^j \leq n \cdot e^{-Rj/n} \ \ \text{(using the fact that } 1-x \leq e^{-x} \text{ for all } x \in [0,1]) \end{split}
```

- Taking j=(n/R) In n gives $m_j<1$, and therefore $m_j=0$.
- Hence, |S|≤(n/R) In n.
- Thus, the total runtime over all argmax calls is O((n/R)log² n).

Proof of Theorem 4.10 (continued):

- Let S be a hitting set of minimal size for M={ N(v) | deg(v) ≥ √n }.
- From Lemma 4.12 ($R = \sqrt{n}$) we know that $|S| = O(\sqrt{n} \log n)$.
- Do a BFS search from each s∈S and add the resulting n edges of the BFS tree to E(H).
- For every u∈V with deg(u)<√n (the low-degree nodes), add all edges incident to u to E(H).
- By construction, $|E(H)| = |S| \cdot n + n \sqrt{n} = O(n^{3/2} \log n)$.
- Consider any pair u,v∈V with shortest path p in G. We have two cases:
- (a): p contains only low-degree nodes. Then p is also contained in H, so d_H(u,v) = d(u,v).
- (b): p contains a high-degree node x. Let s∈S be a node adjacent to x. Then
 we append the shortest paths from u to s and s to v in H to obtain a path
 from u to v in H. It holds:

$$d_H(u,v) \le d_H(u,s)+d_H(s,v) = d(u,s)+d(v,s)$$

 $\le (d(u,x)+1)+(d(v,x)+1) = d(u,v)+2$

Hence, H is indeed a (1,2)-spanner.

Runtime of the algorithm for (1,2)-spanner:

- O(n^{3/2} log n): construction of hitting set S
- O(\int \log n (n+m)): BFS for all nodes in S
- O(n^{3/2}): adding all edges of low-degree nodes to H

Total runtime: $O(\sqrt{n} (m+n) \log n)$.

Runtime of approximate APSP algorithm for an unweighted graph G based on (1,2)-spanner H:

```
O(\sqrt{n} (m+n) \log n)) + O(n \cdot n^{3/2} \log n + n^2 \log n)
= O(n^{5/2} \log n)
```

With a more complex approach the runtime can be reduced to $O(n^{7/3} \log n)$. For the details see:

D. Dor, S. Halperin, and U. Zwick. All-pairs almost shortest paths. SIAM Journal of Computing, 29(5): 1740-1759, 2000.

Interestingly, the following two results are known:

Theorem 4.13: Any n-node graph G has a (1,6)-spanner with $O(n^{4/3})$ edges.

Theorem 4.14: In general, there is no additive spanner with $O(n^{4/3-\epsilon})$ edges for n-node graphs for any $\epsilon>0$.

For more information on that see:

Amir Abboud and Greg Bodwin. The 4/3 additive spanner exponent is tight. Proc. of the 48th ACM Symposium on Theory of Computing (STOC), 2016.

How to quickly answer distance requests?

Naive approach:

Run an APSP algorithm and store all answers in a matrix

Problems:

- High runtime (O(nm + n² log n))
- High storage space (⊕(n²))

Alternative approach, if approximate answers are sufficient:

- Compute additive or multiplicative spanner, and run an APSP algorithm on that spaner.
 - → lower runtime
- But storage space is still high

Better solutions concerning the storage space have been investigated under the concept of distance oracles.

Definition 4.15: An α -approximate distance oracle is defined by two algorithms:

- a preprocessing algorithm that takes as its input a graph G=(V,E) and returns a summary of G, and
- a query algorithm based on the summary of G that takes as its input two vertices u,v∈V and returns an estimate D(u,v) such that d(u,v) ≤ D(u,v) ≤ α·d(u,v).

The quality of an α -approximate distance oracle is defined by its query time q(n), preprocessing time p(m,n), and storage space s(n). The goal is to minimize all of these quantities.

Thorup and Zwick (STOC 2001) have shown the following result for graphs of non-negative edge costs:

Theorem 4.16: For all $k \ge 1$ there exists a (2k-1)-approximate distance oracle using $O(k \cdot n^{1+1/k})$ space and $O(m \cdot n^{1/k})$ time for preprocessing that can answer queries in O(k) time (where we hide logarithmic factors in the O-notation).

Theorem 4.16: For all $k \ge 1$ there exists a (2k-1)-approximate distance oracle using $O(k \cdot n^{1+1/k})$ space and $O(m \cdot n^{1/k})$ time for preprocessing that can answer queries in O(k) time (where we hide logarithmic factors in the O-notation).

```
k=1: trivial. Just run our APSP algorithm.
```

k=2: Consider the following preprocessing algorithm:

```
A:= random subset S \subseteq V of size O(\sqrt{n} \log n) for each a \in A run Dijkstra to compute d(a,v) for all v \in V for each v \in V \setminus A do p_A(v) := argmin_{y \in A} d(v,y) run Dijkstra to compute A(v) := \{x \in V \mid d(v,x) < d(v,p_A(v))\} B(v) := A \cup A(v) store p_A(v) under v for all x \in B(v), store d(v,x) under key (v,x) in a hash table
```

It is easy to show that with high probability A is a hitting set of
 M={ N_{√n}(v) | v∈V}, where N_{√n}(v) is the set containing the closest √n nodes
 to v.

```
Lemma 4.17: |B(v)| \le O(\sqrt{n} \log n). Proof:
```

- It suffices to show that |A(v)| ≤ \(\bar{n} \) since by construction |A|=O(\(\bar{n} \) log n).
- Because A hits the closest \n nodes to v with high probability, some node a∈A is also in N_{√n} (v).
- Thus, all nodes closer to v than a are also in N_{In} (v).
- By the definition of A(v), this implies that $A(v) \subseteq N_{\Gamma \Gamma}(v)$, and therefore, $|A(v)| \le \sqrt{\Gamma}$.

Proof of Theorem 4.16 (continued):

Now we can bound the quality of the distance oracle.

- Lemma 4.17 implies that the storage space needed by our distance oracle is O(n·√n log n).
- A query (u,v) is processed as follows:

```
if d(u,v) is stored in the hash table then
  return d(u,v)
else
  return d(u,p<sub>A</sub>(u))+d(v,p<sub>A</sub>(u))
```

 This obviously takes constant time. We also need to show that d(u,v)≤D(u,v)≤3d(u,v).

```
Claim: d(u,v) \leq D(u,v) \leq 3d(u,v).
```

- Suppose that v∉B(u), as otherwise D(u,v)=d(u,v).
- By the triangle inequality we have

```
d(u,v) \leq d(u,p_A(u))+d(v,p_A(u)) = D(u,v)
```

 In order to show that D(u,v)≤3d(u,v), we use the triangle inequality again to obtain

```
\begin{split} D(u,v) &= d(u,p_A(u)) + d(p_A(u),v) \\ &\leq d(u,p_A(u)) + (d(p_A(u),u) + d(u,v)) \\ &= 2d(u,p_A(u)) + d(u,v) \\ &\leq 3d(u,v) \end{split}
```

Proof of Theorem 4.16 (continued):

It remains to determine the runtime for the preprocessing.

- For each a∈A run Dijkstra to compute d(a,v) for all v∈V: runtime O((m+n log n). √n log n).
- Determine $p_A(v)$ for each $v \in V \setminus A$: total runtime $O(n \cdot \sqrt{n \log n})$
- Run Dijkstra to compute A(v):={x∈V | d(v,x)<d(v,p_A(v))}: We need to modify Dijkstra so that we start with a Fibonacci heap only containing v and we only include a node y into the heap (and therefore process it) if d(x)+c(x,y)<d(v,p_A(v)) for some processed x. Then the runtime for each v∈V\A is O(|E(A(v))|+|A(v)| log n), where E(A(v)) is the set of all edges containing vertices of A(v).
- Let C(v)={ w∈V | v∈A(w) }. One can show (similarly to A(v)) that the expected size of C(v) ist at most 2\n. Hence,

```
\begin{split} \Sigma_{\mathsf{w}\in\mathsf{V}} \left| \mathsf{E}(\mathsf{A}(\mathsf{w})) \right| &\leq \Sigma_{\mathsf{w}\in\mathsf{V},\;\mathsf{v}\in\mathsf{A}(\mathsf{w})} \left| \mathsf{E}(\mathsf{v}) \right| = \Sigma_{\mathsf{v}\in\mathsf{V},\;\mathsf{w}\in\mathsf{C}(\mathsf{v})} \left| \mathsf{E}(\mathsf{v}) \right| \\ &= \Sigma_{\mathsf{v}\in\mathsf{V}} \left( \left| \mathsf{C}(\mathsf{v}) \right| \cdot \left| \mathsf{E}(\mathsf{v}) \right| \right) = \mathsf{O}(\mathsf{m} \cdot \sqrt{\mathsf{n}}) \end{split}
```

- Thus, the overall runtime for the A(v)'s is $O(m \cdot \sqrt{n} + n \sqrt{n} \log n)$.
- Therefore, the preprocessing needs $O((m + n \log n) \cdot \sqrt{n \log n})$ time.

Proof of Theorem 4.16 (continued):

The algorithm for general k proceeds by taking many related samples $A_0,...,A_k$ instead of just a single sample A. Concretely, it does the following:

- Let $A_0 = V$ and $A_k = \emptyset$. For each $1 \le i \le k-1$, choose a random $A_i \subseteq A_{i-1}$ of size $(|A_{i-1}|/n^{1/k}) \log n = O(n^{1-i/k} \log n)$.
- Let $p_i(v)$ be the closest node to v in A_i . If $d(v,p_i(v))=d(v,p_{i+1}(v))$ then let $p_i(v)=p_{i+1}(v)$.
- For all v∈V and i<k-1, define

```
\begin{array}{l} A_i(v) = \{ \; x {\in} A_i \; | \; d(v, x) < d(v, p_{i+1}(v)) \; \} \\ B(v) = A_{k-1} {\cup} (U_{i=0}^{k-2} \; A_i(v)) \end{array}
```

• For all $v \in V$ and all $x \in B(v)$, store d(v,x) in a hash table. Also store for each $v \in V$ and each $i \le k-1$, $p_i(v)$.

A query for (u,v) then works as follows:

```
w:=p_0=v for i=1 to k do if w\in B(u) then return d(u,w)+d(w,v) else w:=p_i(u); swap u and v
```

For more information see:

Mikkel Thorup and Uri Zwick. Approximate distance oracles. In Proc. of the 33rd ACM Symposium on Theory of Computing (STOC), 2001.

Next Chapter

Matching algorithms...