Approximation Algorithms

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Approximation algorithms

- Fast. Cheap. Reliable. Choose two.
- NP-hard problems: choose 2 of
 - optimal
 - polynomial time
 - all instances
- Approximation algorithms. Trade-off between time and quality.
- Let A(I) denote the value returned by algorithm A on instance I. Algorithm A is an *a*-approximation algorithm if for any instance I of the optimization problem:
 - A runs in polynomial time
 - A returns a valid solution
 - A(I) $\leq \alpha \cdot OPT$, where $\alpha \geq 1$, for minimization problems
 - $A(I) \ge \alpha \cdot OPT$, where $\alpha \le 1$, for maximization problems

Examples

- **Acyclic Graph** Given a directed graph G=(V,E), pick a maximum cardinality set of edges from E such that the resulting graph is acyclic.
 - Give a 1/2-approximation algorithm for this problem.

Minimum Maximal Matching

- A matching in a graph G=(V,E) is a subset of edges M ⊆ E, such that no two
 edges in M share an endpoint.
- A maximal matching is a matching that cannot be extended, i.e., it is not possible to add an edge from E \ M to M without violating the constraint.
- Design a 2-approximation algorithm for finding a minimum cardinality maximal matching in an undirected graph.

Examples

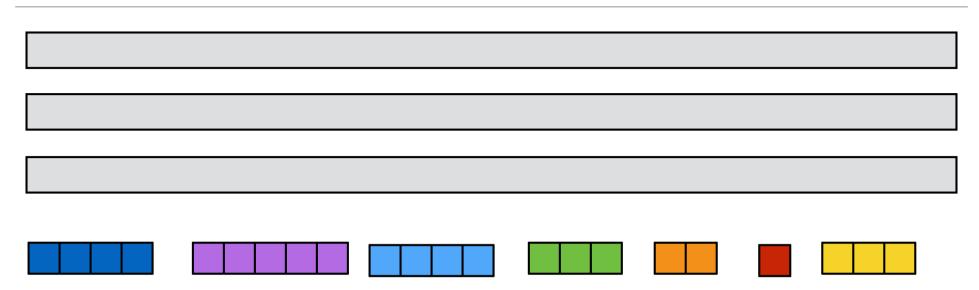
- **Acyclic Graph** Given a directed graph G=(V,E), pick a maximum cardinality set of edges from E such that the resulting graph is acyclic.
 - Give a 1/2-approximation algorithm for this problem.
 - Lower bound what is the best we can hope for?
 - Arbitrarily number the vertices and pick the bigger of the two sets, the forward going edges and the backward going edges.

Minimum Maximal Matching

- A matching in a graph G=(V,E) is a subset of edges M \subseteq E, such that no two edges in M share an endpoint.
- A maximal matching is a matching that cannot be extended, i.e., it is not
 possible to add an edge from E\setminus M to M without violating the constraint.
- Design a 2-approximation algorithm for finding a minimum cardinality maximal matching in an undirected graph.
- Lower bound: Any maximal matching is at least half the maximum maximal matching. Why?

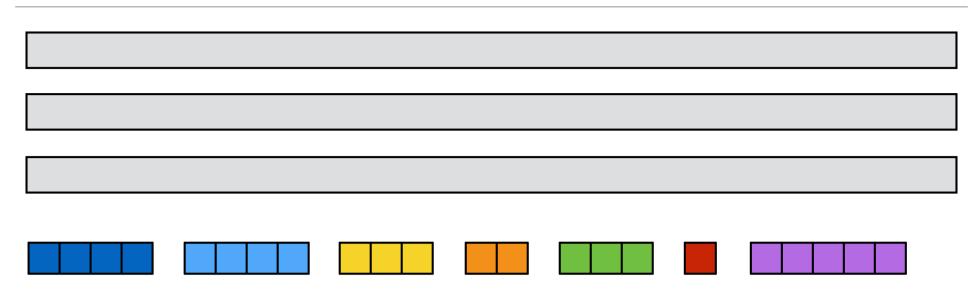
Load balancing

Scheduling on identical parallel machines

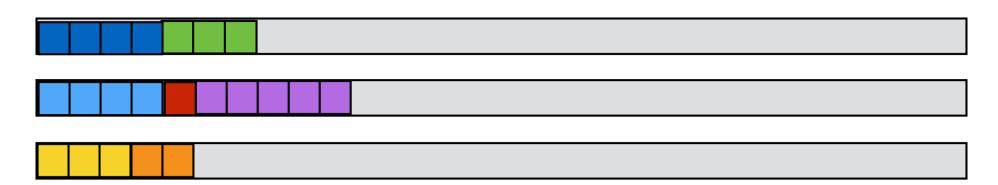


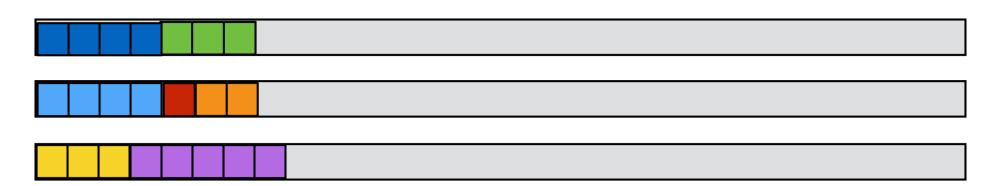
- n jobs to be scheduled on m identical machines.
- Each job has a processing time t_i.
- Once a job has begun processing it must be completed.
- T_{i:} Load of machine j.
- Goal. Schedule all jobs so as to minimize the maximum load (makespan):

minimize
$$T = \max_{i=1...n} T_i$$



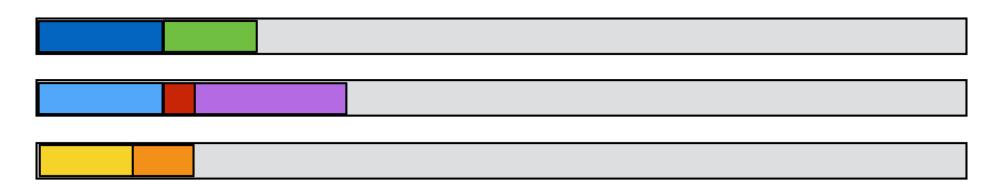
- Simple greedy. Process jobs in any order. Assign next job on list to machine with smallest current load.
- The greedy algorithm above is a 2-approximation algorithm:
 - polynomial time ✓
 - valid solution ✓
 - factor 2







Approximation factor



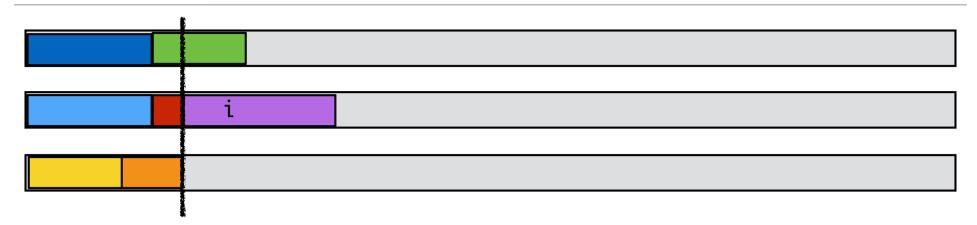
- Lower bounds:
 - Each job must be processed:

$$T^* \ge \max_j t_j$$

• There is a machine that is assigned at least average load:

$$T^* \ge \frac{1}{m} \sum_{j} t_j$$

Approximation factor

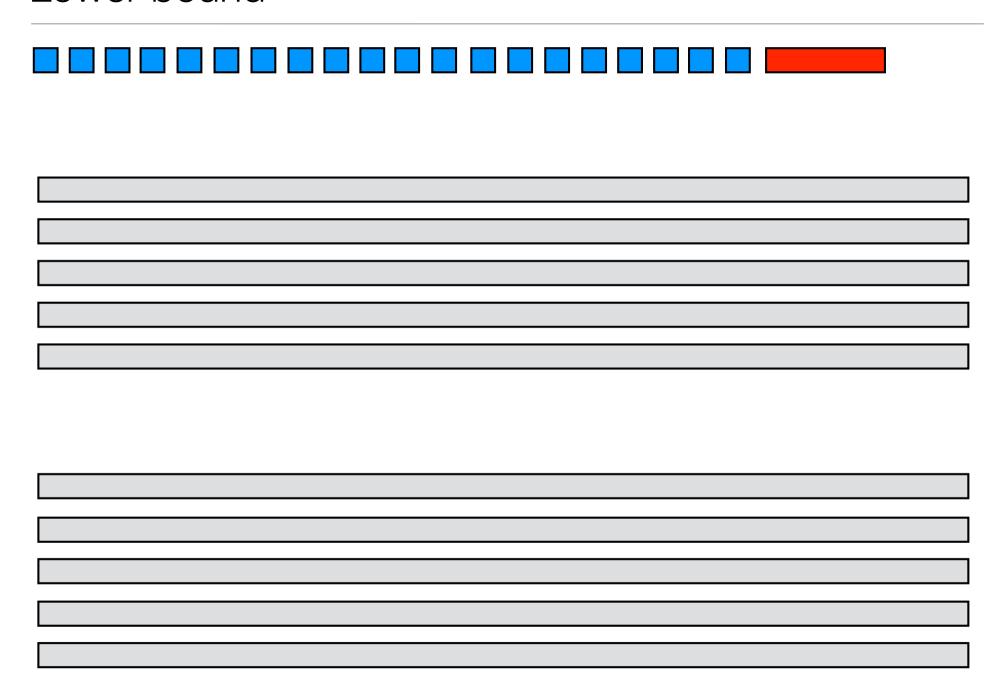


- i: job finishes last.
- All other machines busy until start time s of i. ($s = T_i t_i$)
- Partition schedule into before and after s.
- After $\leq T^*$.
- Before:
 - All machines busy => total amount of work = m·s:

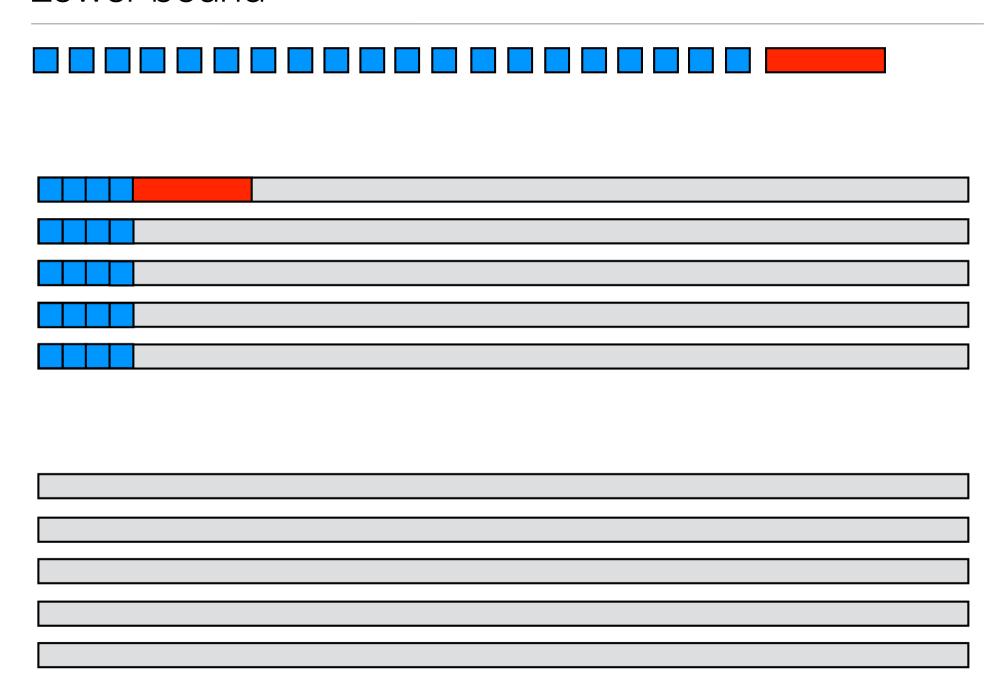
$$m \cdot s \le \sum_{j} t_{j}$$
 $\Rightarrow s \le \frac{1}{m} \sum_{j} t_{j} \le T^{*}$

• Length of schedule = $s + t_i \le T^* + T^* = 2T^*$.

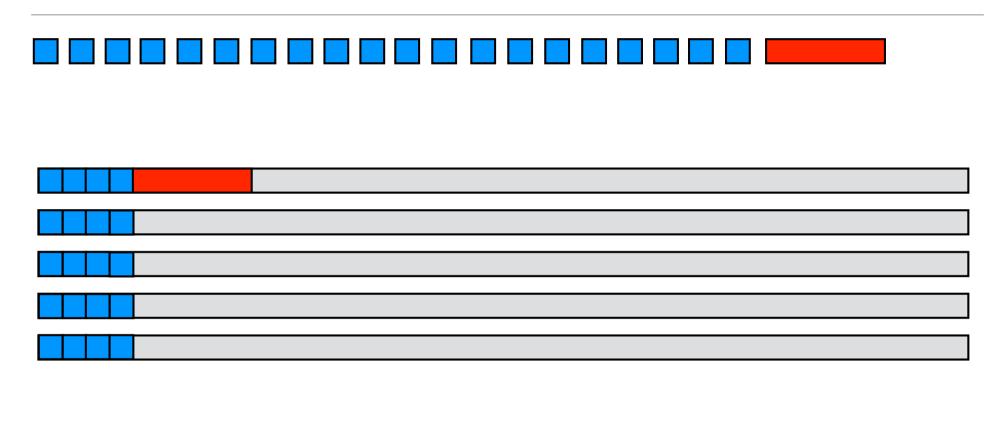
Lower bound



Lower bound

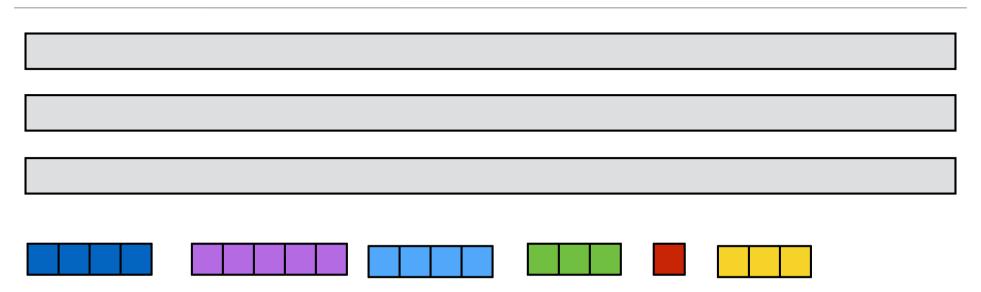


Lower bound



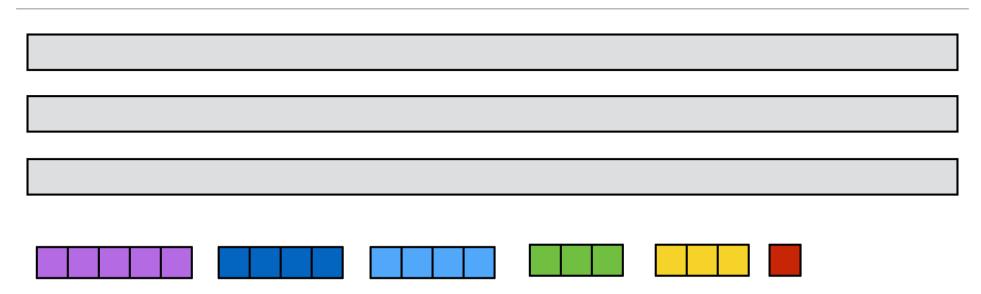


Longest processing time rule



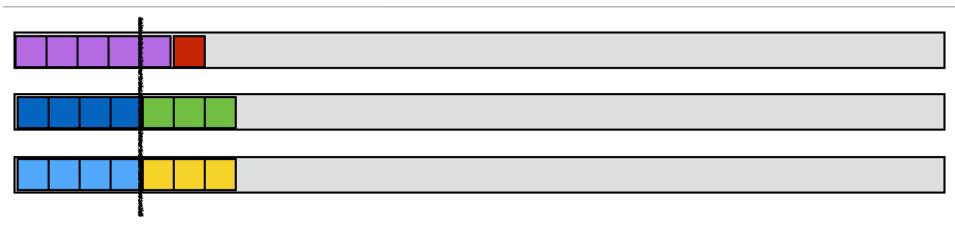
• Longest processing time rule (LPT). Sort jobs in non-increasing order. Assign next job on list to machine as soon as it becomes idle.

Longest processing time rule



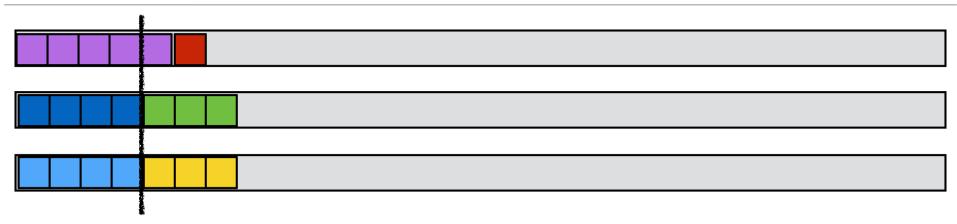
- Longest processing time rule (LPT). Sort jobs in non-increasing order. Assign next job on list to machine as soon as it becomes idle.
- LPT is a is a 3/2-approximation algorithm:
 - polynomial time ✓
 - valid solution
 - factor 3/2

Longest processing time rule: factor 3/2



- Longest processing time rule (LPT). Sort jobs in non-increasing order. Assign next job on list to machine as soon as it becomes idle.
- Assume $t_1 \ge \ge t_n$.
- If $n \le m$ then optimal.
- Lower bound: If n > m then $T^* \ge 2t_{m+1}$.
- Factor 3/2:
 - Before $s \le T^*$
 - After: i job that finishes last.
 - $t_i \le t_{m+1} \le T^*/2$.
 - $T \le T^* + T^*/2 \le 3/2 T^*$.
- Tight?

Longest processing time rule: factor 4/3



- Longest processing time rule (LPT). Sort jobs in non-increasing order. Assign next job on list to machine as soon as it becomes idle.
- Assume $t_1 \ge \ge t_n$.
- Assume wlog that smallest job finishes last.
- If $t_n \le T^*/3$ then $T \le 4/3 T^*$.
- If t_n > T*/3 then each machine can process at most 2 jobs in OPT.
- Lemma. For any input where the processing time of each job is more than a third of the optimal makespan, LPT computes an optimal schedule.
- Theorem. LPT is a 4/3-approximation algorithm.

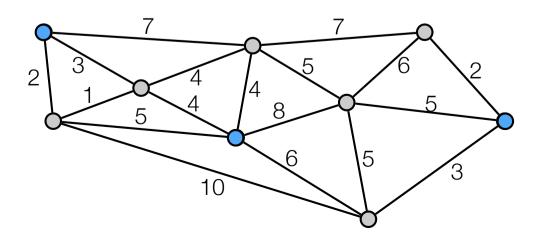
k-center

The k-center problem

- Input. An integer k and a set of sites S with distance d(i,j) between each pair of sites i,j ∈ S.
- · d is a metric:
 - dist(i,i) = 0
 - dist(i,j) = dist(j,i)
 - dist(i,l) ≤ dist(i,j) + dist(j,l)
- Goal. Choose a set $C \subseteq S$, |C| = k, of k centers so as to minimize the maximum distance of a site to its closest center.

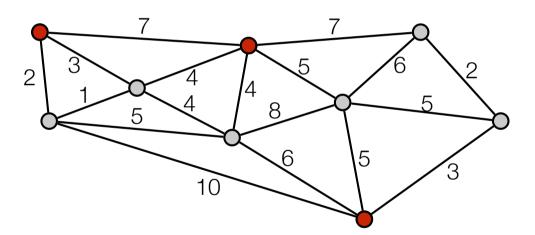
$$C = \operatorname{argmin}_{C \subseteq V, |C| = k} \operatorname{max}_{i \in V} \operatorname{dist}(i, C)$$

Covering radius. Maximum distance of a site to its closest center.



k-center: Greedy algorithm

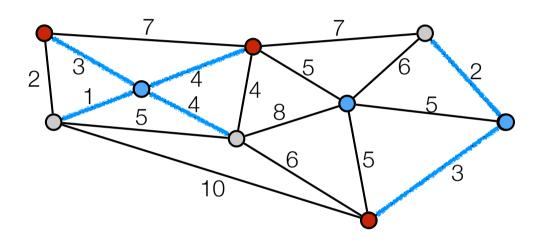
- · Greedy algorithm.
 - Pick arbitrary i in S.
 - Set $C = \{i\}$
 - while |C| < k do
 - Find site j farthest away from any cluster center in C
 - Add j to C
 - Return C



- Greedy is a 2-approximation algorithm:
 - polynomial time ✓
 - valid solution ✓
 - factor 2

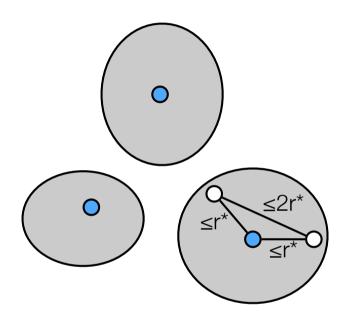
k-center analysis: optimal clusters

• Optimal clusters: each site assigned to its closest optimal center.

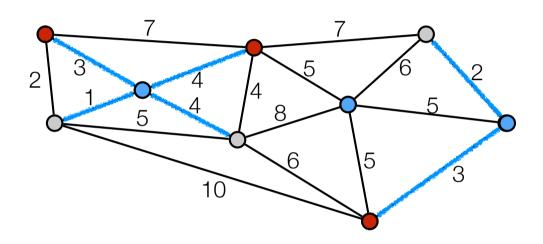


k-center analysis

- r* optimal radius.
- Claim: Two sites in same optimal cluster has distance at most 2r* to each other.

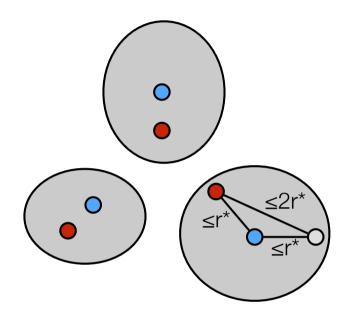


k-center

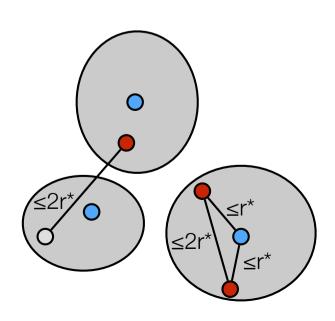


k-center: analysis greedy algorithm

- r* optimal radius.
- Show all sites within distance 2r* from a center.
- Consider optimal clusters. 2 cases.
 - Algorithm picked one center in each optimal cluster
 - distance from any sites to its closest center ≤ 2r*.



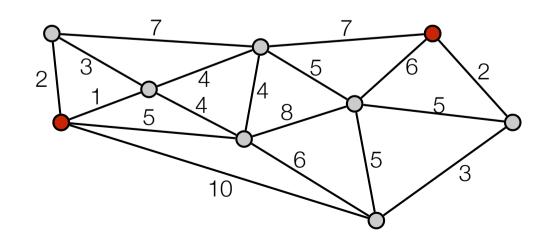
- 2. Some optimal cluster does not have a center.
 - Some cluster have more than one center.
 - Distance between these two centers ≤ 2r*.
 - When second center in same cluster picked it was the site farthest away from any center.
 - Distance from any site to its closest center at most 2r*.



Bottleneck algorithm

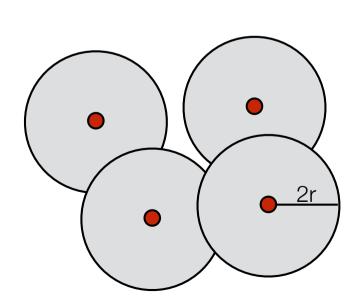
- · Assume we know the optimum covering radius r.
- Bottleneck algorithm.
 - Set R := S and $C := \emptyset$.
 - while $R \neq \emptyset$ do
 - Pick arbitrary j in R.
 - Add j to C
 - Remove all sites with $d(j,v) \le 2r$ from R.
 - Return C

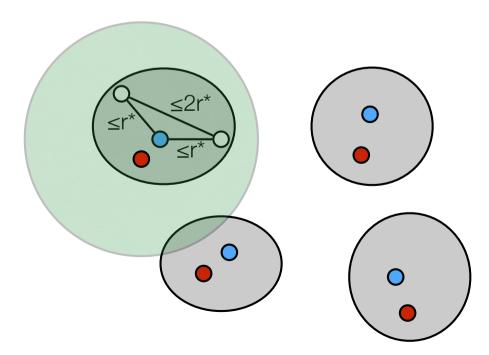
• Example: k = 3. r = 4.



Analysis bottleneck algorithm

- r* optimal radius.
- Covering radius is at most 2r = 2r*.
- Show that we cannot pick more than k centers:
 - We can pick at most one in each optimal cluster:
 - Distance between two nodes in same optimal cluster ≤ 2r.*
 - When we pick a center in a optimal cluster all nodes in same optimal cluster is removed.





Analysis bottleneck algorithm

- r* optimal radius.
- Can use algorithm to "guess" r* (at most n² values).
- If algorithm picked more than k centers then r* > r.
 - If algorithm picked more than k centers then it picked more than one in some optimal cluster.
 - Distance between two nodes in same optimal cluster ≤ 2r.*
 - If more than one in some optimal cluster then 2r < 2r*.

