

# A better k-means++ Algorithm via Local Search

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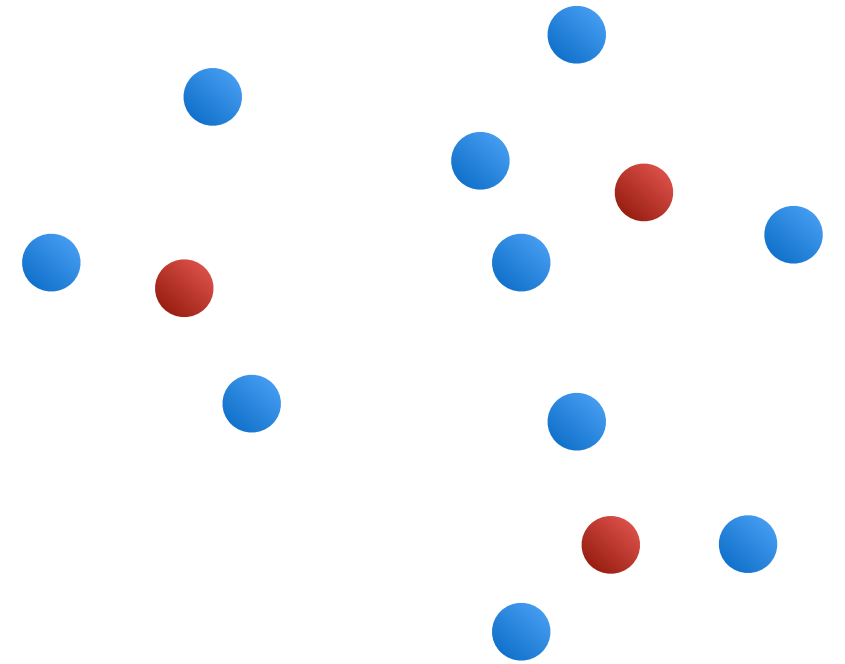
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Google Research

# k-means

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Find a set of k centers

$$\phi(X, C) = \sum_{x \in X} \min_{c \in C} d^2(x, c)$$



Constant approximation algorithms are known.

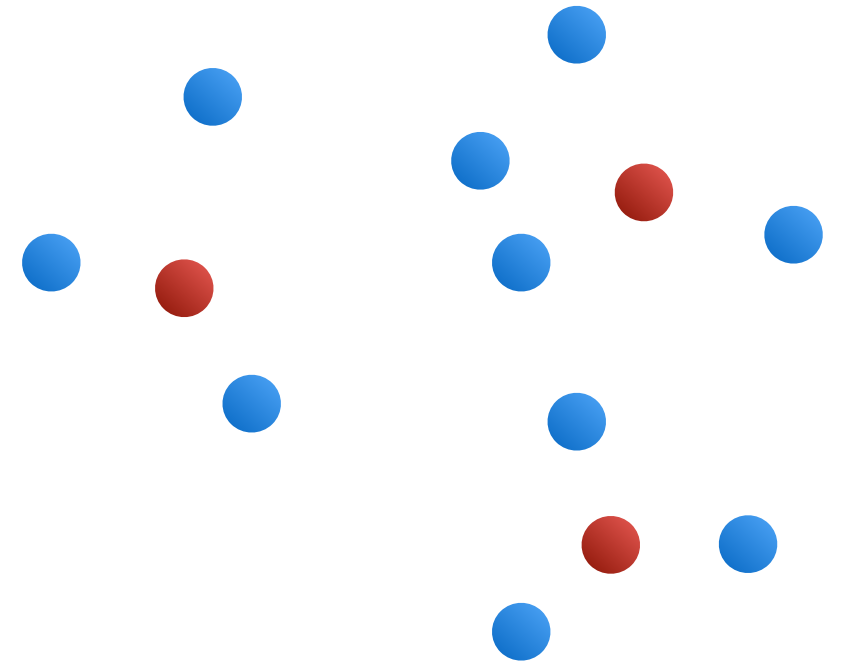
Goal is to design a constant approximation algorithm that is efficient, easy to implement and has good experimental results.

# k-means++ seeding

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Elegant and simple algorithm

```
Uniformly sample  $p \in P$  and set  $C = \{p\}$ .  
for  $i \leftarrow 2, 3, \dots, k$  do  
  Sample  $p \in P$  with probability  $\frac{\text{cost}(\{p\}, C)}{\sum_{q \in P} \text{cost}(\{q\}, C)}$  and  
  add it to  $C$ .  
end for
```



Experimentally gives good results when combined with Lloyd's algorithm.

The solution is a  $O(\log k)$  approximation in expectation.

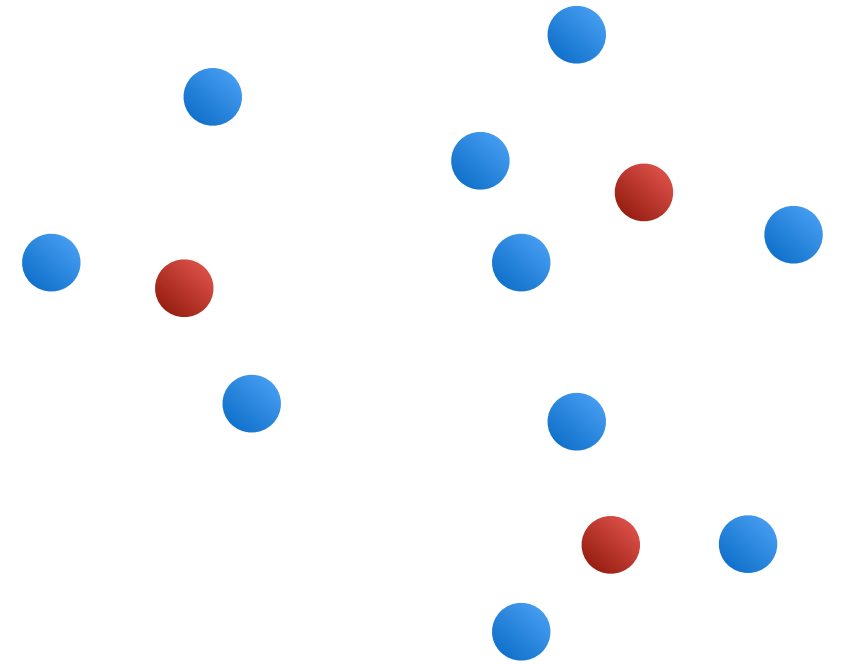
David Arthur, Sergei Vassilvitskii: k-means++: the advantages of careful seeding. SODA 2007: 1027-1035

# Local search

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Elegant and simple algorithm

```
if  $\exists q \in C$  s.t.  $\text{cost}(P, C \setminus \{q\} \cup \{p\}) < \text{cost}(P, C)$   
then  
  Let  $q \in C$  be the  $q$  s.t.  $\text{cost}(P, C \setminus \{q\} \cup \{p\})$  is  
  minimized  
   $C = C \setminus \{q\} \cup \{p\}$   
end if
```



It returns a constant approximation and nice experimental results.

The algorithm is a bit slow.

Tapas Kanungo, David M. Mount, Nathan S. Netanyahu, Christine D. Piatko, Ruth Silverman, Angela Y. Wu:  
A local search approximation algorithm for k-means clustering. *Comput. Geom.* 28(2-3): 89-112 (2004)

# Combining the two algorithms

## Elegant and simple algorithm

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**Algorithm 1**  $k$ -means++ seeding with local search

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**Require:**  $P, k, Z$

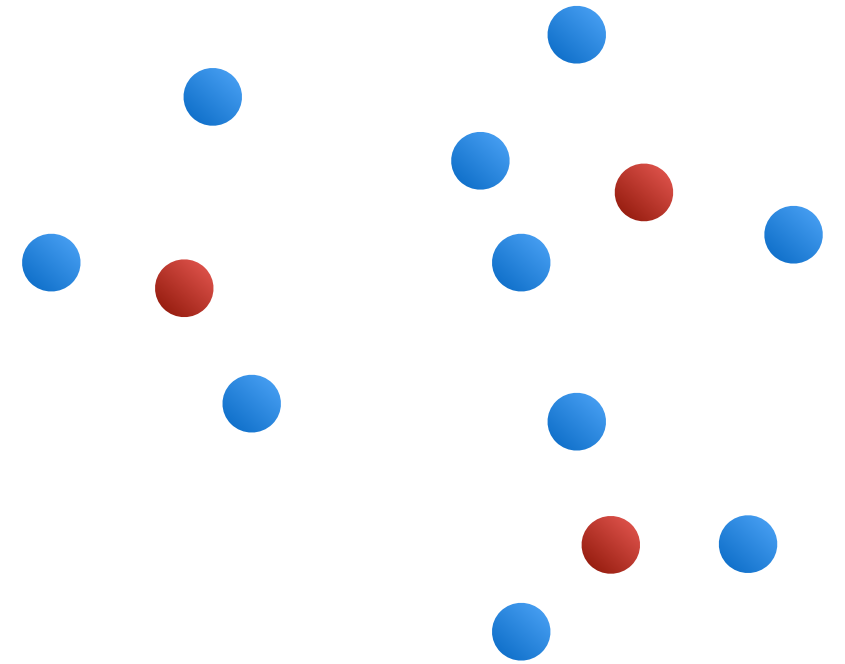
- 1: Uniformly sample  $p \in P$  and set  $C = \{p\}$ .
  - 2: **for**  $i \leftarrow 2, 3, \dots, k$  **do**
  - 3: Sample  $p \in P$  with probability  $\frac{\text{cost}(\{p\}, C)}{\sum_{q \in P} \text{cost}(\{q\}, C)}$  and add it to  $C$ .
  - 4: **end for**
  - 5: **for**  $i \leftarrow 2, 3, \dots, Z$  **do**
  - 6:  $C = \text{LocalSearch++}(P, C)$
  - 7: **end for**
  - 8: **return**  $C$
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**Algorithm 2** LocalSearch++

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**Require:**  $P, C$

- 1: Sample  $p \in P$  with probability  $\frac{\text{cost}(\{p\}, C)}{\sum_{q \in P} \text{cost}(\{q\}, C)}$
  - 2: **if**  $\exists q \in C$  s.t.  $\text{cost}(P, C \setminus \{q\} \cup \{p\}) < \text{cost}(P, C)$  **then**
  - 3: Let  $q \in C$  be the  $q$  s.t.  $\text{cost}(P, C \setminus \{q\} \cup \{p\})$  is minimized
  - 4:  $C = C \setminus \{q\} \cup \{p\}$
  - 5: **end if**
  - 6: **return**  $C$
- 



It returns a constant approximation, it is slightly slower than  $k$ -means++ and has better experimental results.

# Main theoretical result

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**Theorem 1.** *Let  $P \subseteq \mathbb{R}^d$  be a set of points and  $C$  be the output of Algorithm 1 with  $Z \geq 100000k \log \log k$  then we have  $E[\text{cost}(P, C)] \in O(\text{cost}(P, C^*))$ , where  $C^*$  is the set of optimum centers. The running time of the algorithm is  $O(dnk^2 \log \log k)$ .*

Main idea is to adapt local search analysis to show that in every step with constant probability we reduce the cost of the solution by a multiplicative  $\left(1 - \frac{1}{100k}\right)$  factor

# Experimental results

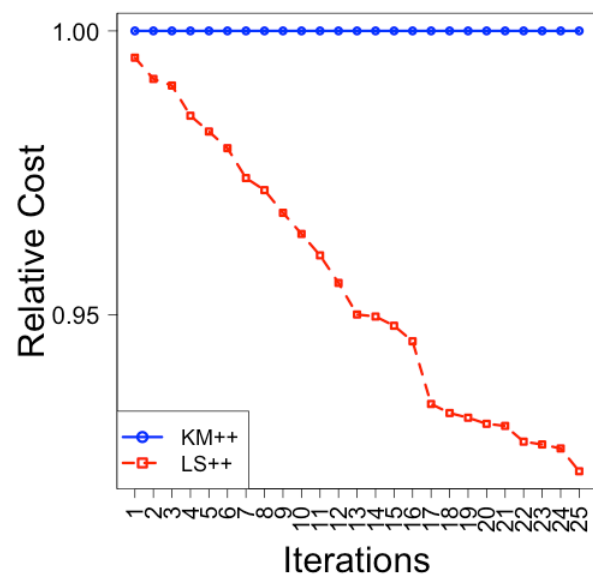
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## Datasets:

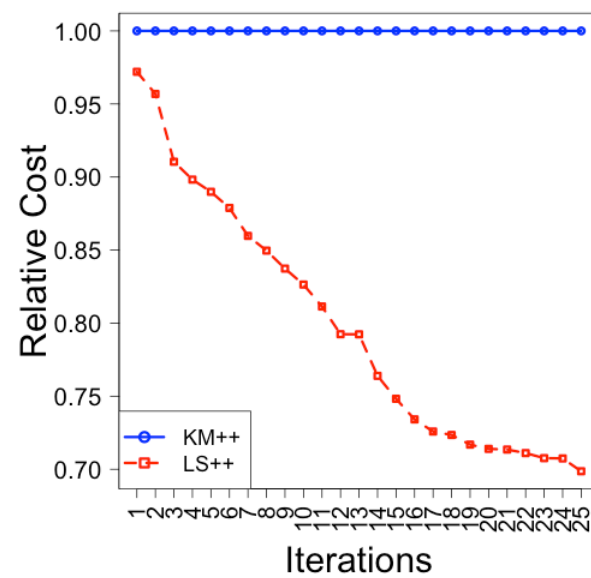
- **RNA**: 8 features from 488565 RNA input sequence pairs (Uzilov et al., 2006)
- **KDD-BIO**: 145751 samples with 74 features measuring the match between a protein and a native sequence (KDD)
- **KDD-PHY**: 100000 samples with 78 features representing a quantum physic task (KDD)

# Experimental results

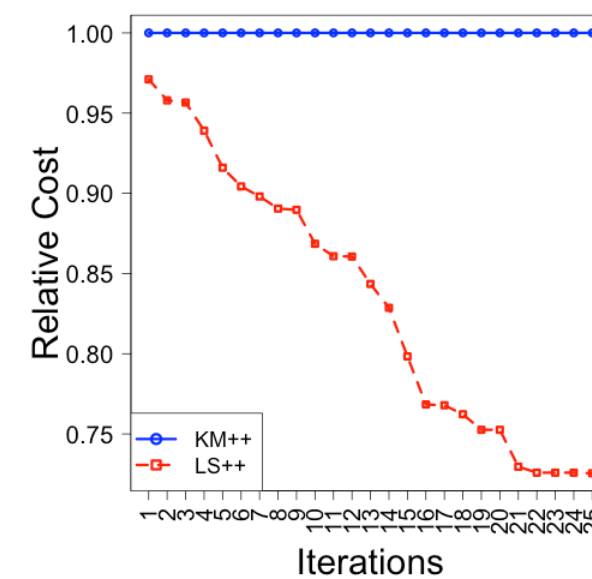
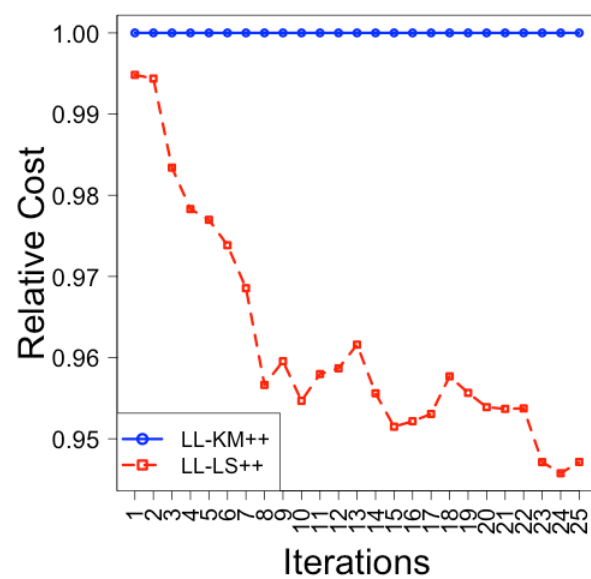
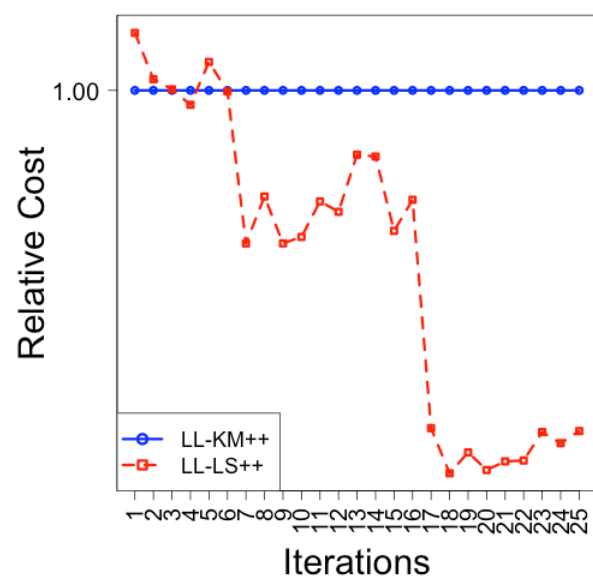
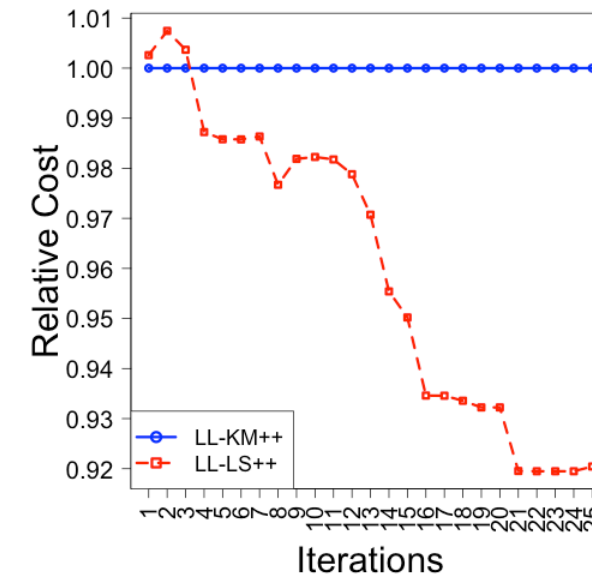
## KDD-BIO



## RNA



## KDD-PHY





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# Thanks