Parallel stability-based k-means*

Jason Poulos[†]

Emin Arakelian[‡]

Fadi Kfoury[§]

Abstract

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1. Introduction

Unsupervised learning is a branch of machine learning that infers patterns from data that has no labels. k-means is a popular unsupervised algorithm for finding clusters and cluster centers in data. The goal is to choose k cluster centers to minimize the total squared distance between each data point and its closest center. Given k initial centers chosen uniformly at random from the data points, k-means alternates between two steps until convergence: (assignment step) each point is assigned to the nearest cluster center and; (update step) each center is recomputed as the center of mass of all points assigned to it.

k-means is NP-hard, and can be solved in time $O(n^{dk+1}\log n)$, where n is the number of points in d dimensions. While it's hard to parallelize k-means itself due to its sequential nature, researchers can find shortcuts in the algorithm, or run numerous k-means instances on sub-samples to evaluate the stability of clustering. Our project compares sequential and parallelized versions of the stability-based method.

After briefly reviewing other parallel implementations of k-means in Section 2, we describe the stability-based method and its parallel implementation in Section 3. We then describe experiments and results in Section 4. Finally, we draw conclusions in Section 5.

2. Related work

k-means++ chooses only the first cluster center uniformly at random; subsequent centers are selected from

the data points, weighed by a probability proportional to its contribution to the overall error. This initialization algorithm is shown to obtain a set of initial centers that is close to the optimum solution [1].

k-means|| k-means++ is ill-suited for massive data because it makes k sequential passes over the data points in order to obtain the initial centers. Bahmani et al. [2] propose a method of parallelizing the initialization that reduces the number of passes, which they call k-means||. Instead of sampling a single point in each pass, k-means|| samples O(k) points and repeats for $O(\log n)$ rounds.

3. Stability-based method

Ben-Hur et al. [3] propose an algorithm that uses stability of clustering with respect to perturbations such as sub-sampling as a means of defining meaningful partitions. Computing the stability measure is the bottleneck, so parallelizing the algorithm will involve enhancements to re-use parts of the computations and avoid storing/writing the full matrix multiplication for computing the measure.

3.1. Parallel implementation

4. Experiments

We compare the algorithms using data from a dialect survey, which includes linguistic binary encoded responses [4]. The purpose of the survey is to figure whether a stable number of clusters can be found across the survey participants. We implement the serial and parallel stability-based methods in Python and run all implementations on Edison. Performance is evaluated on three dimensions — clustering cost, running time, and space complexity — for different values of k.

 $^{^\}dagger$ poulos@berkeley.edu

 $^{^\}ddagger$ emin@berkeley.edu

[§]fadi.kfoury@berkeley.edu

^{*}The code used for this project is available on Github: https://github.com/plumSemPy/parallel_kmeans.

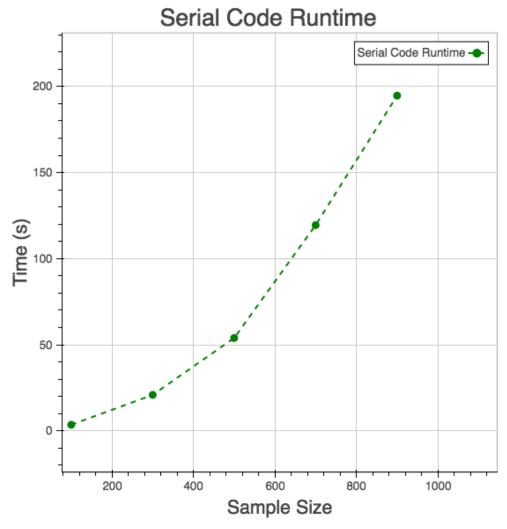


Figure 1. Serial code complexity.

5. Conclusion

References

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