Parallel K-means

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Abstract

This project aims to implement and compare sequential and parallel versions of the K-means algorithm using both OpenMP and CUDA. The objective is to analyze performance and measure the speedup achieved by the parallel version compared to the sequential one.

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1. System Specification

CPU: Intel i3-4005U (4 Threads) 1.600GHz **GPU**: NVIDIA GeForce 920M

2. Introduction to K-means

K-means clustering is an unsupervised machine learning algorithm used for partitioning a dataset into a predetermined number of clusters. The objective of K-means is to group similar data points together and discover underlying patterns or structures within the data.

At its core, K-means operates by iteratively assigning each data point to the nearest cluster centroid and then recalculating the centroids based on the mean of the data points assigned to each cluster. This process continues until the centroids no longer change significantly, or a specified number of iterations is reached.

One of the key parameters in K-means is 'k', which represents the number of clusters the algorithm should identify.

One iteration of the algorithms works in this way

- 1. for every point in the dataset, the point is assigned to the nearest centroid
- 2. the centroid are updated and convergence is checked

The algorithm stops when

- 1. the maximum number of iteration is reached
- 2. convergence is reached. This happens when two consecutive iteration of the algorithm don't change the centroid position (given a certain tollerance).

3. Sequential implementation

I implemented a KMeans algorithm in C++ with a KMeans class. I used a array of structure approach (AoS) over a structure of arrays (SoA) because the SoA implementation will have drawbacks. In fact the K-means algorithm needs to access, for each point, all the coordinates of the point, for this reason the SoA approach will have too many cache misses.

In my implementation the public method:

void fit(std::vector<Point>& dataPoints,int
maxIteration, bool useStopCondition);

takes a vector of data points, the maximum iteration number and if the stop condition should be used or not. For test purpose data points are generated by a python script ($create_dataset.py$) which generates a csv file that is imported as a Point vector in c++ using the function loadCsv (defined in utils.cpp).

At each iteration of the algorithm the centroids vector attribute of the class is updated, and at the end it is possible to export the result of the *fit*

method using the exportCsv function (defined in utils.cpp).

An example of the result of the algorithm can be seen in Figure 1.

3.1. centroids initialization

The function *fit* of the class K-means expect the vector *centroids* to be already initialized. Centroids can be initialized in many ways, for example with an algorithm known as K-means++. For the sake of semplicity i decided to choose random K random points in the dataset and making them the initial centroids.

4. parallel implementation with openMP

It is clear that the part which takes more time in every iteration of the algorithm is the first one, because it has to cycle through every point of the dataset. Luckily all operaion done for each point is indipendent one from another. This make this problem an embarassingly parallel one. Using openmp we can make the code parallel in this way: we put this openmp directive before the first step of k-means

```
#pragma omp parallel for schedule(static) default
(none) shared(dataPoints,newCentroids)
num_threads(4)
```

The schedule is chosen to be static because every thread has the same amount of work to do.

For the second step of kmeans another openmp directive is used

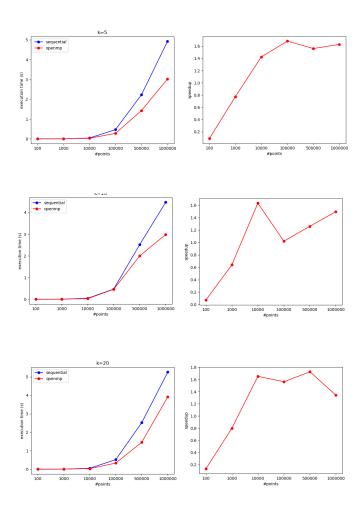
```
#pragma omp parallel for default(none) shared(
newCentroids,converged,useStopCondition)
num_threads(4)
```

because each update of the centroids is independent of each other.

5. speedup analysis

For the analysis i have set the use of stop condition to be false, because i noticed that many times

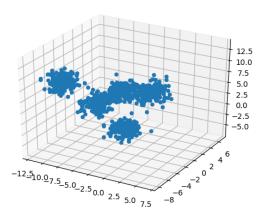
the number of iteration is less than the thread available, so the speedup is not noticeable. For this analysis the same centroids are used for both the sequential and parallel run. Dataset are generated with increasing number of points and the times of the sequential and parallel program are compared.



We can see that the maximum speedup we get is about 1.8.

6. CUDA

CUDA version of kmeans can be implemented as the sequential one, the only thing to take into account is that doing an atomicAdd for each point (and dimension) of the dataset can lead to too much time waste. The strategy i adopted is a sort of reduction, implemented in the kernel centroidUpdate.



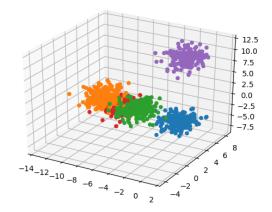


Figure 1. on the left a 1000 points dataset with 3 dimension, on the right the result of K-means with K=5

The other kernel i implemented is centroidAssign, it's purpose is to assign a cluster to each point. This operation is an embarassingly parallel one and doesn't require any coordination between threads.

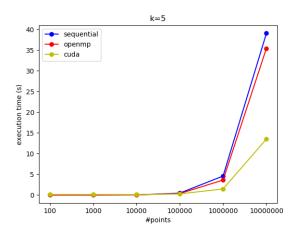
For data storage i decided to follow the sequential implementation and go for 3 dimension point dataset. Every point is stored with each coordinate next to each other and than there is another component that rapresent the cluster label which the point is assigned to during the algorithm. Same thing happens for centroids, but every 4th cell the centroid cardinality is stored.

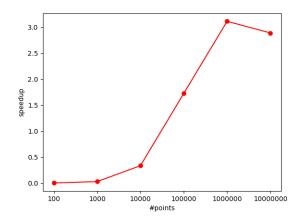
So after the execution of the kernel centroidAssign, every 4th*i (i=1,..,N) cell of the data points array is labeled with the corresponding closest centroid label. Similarly, after the execution of centroidUpdate kernel, every 4th*i (i=1,..,K) cell of the centroids points is labeled with the corresponding cardinality and the points component are updated with the sum of associated points.

The kernel *centroidUpdate* uses shared memory and performs a reduction in order to reduce the number of atomicAdd performed. Every first thread of a block calculates the sum of the dimension of the points associated to a cluster and then performs the atomicAdd.

7. Speedup analysis

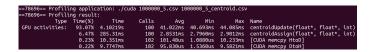
Here all 3 versions are compared, with datasets up to 10^7 points.

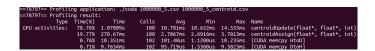




We can see that the speedup for the CUDA ver-

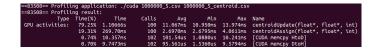
sion is more than 3. Blocks of 512 threads were chosen for the parallel version because analyzing performance with nvprof with higher thread counts leads to performance degradation. For example here we can see first the profiled output with thread blocks of dimension 1024 than 512.





We can notice how the bottleneck is the centroidUpdate kernel as we expected and that with smaller block sizes the performance is better.

Of course if we choose block sized too small, perfomance might be worse too.



Here block sizes of 256 are tested.