

# Similarity Group By for Big Data Analytics

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### Motivation

#### The Problem

- Analyzing massive amounts of data is critical for many commercial and scientific applications. However, this task can require processing tens to hundreds of terabytes of data.
- Big Data Systems like Apache Hadoop and Spark and their MapReduce (MR) programming framework enable analyzing very large datasets in a highly parallel and scalable way.
- Grouping operations are among the most useful operators for data processing and analysis.

#### **General Considerations**

- 1. A similarity group is defined as a set of points where each point is within **epsilon** of each other.
- 2. We propose MR-SGB, a MapReduce-based algorithm to efficiently identify similarity groups in large datasets.
- 3. Our algorithm is based on partitioning the data into smaller partitions. Each partitioning round uses a set of special points named pivots. Each data point will be associated with the group corresponding to its closest pivot.
- 4. Even though the algorithm processes the data in parallel over many nodes, it guarantees that each similarity group is generated only once.

## Algorithm for K Pivots

#### 1. Partition the data [Map]

- Duplicate the points in overlapping areas (each base partition is extended by *epsilon*)
- Structure of each record: {RecordID, RecordContent, AssignedPartition, BasePartition}
  - BasePartition: This is the ID of the pivot that is closest to the current record
  - AssignedPartition: This is the ID of the pivot associated to the current partition

#### 2. For each partition $P_i$ , cluster the points in $P_i$ [Reduce]

- For each partition, we know the value of i by looking at the AssignedPartition component of any record
- Structure of each cluster  $C_n$ : {SetOfPoints,  $[f_1, f_2, ..., f_n]$ 
  - Observe that the array has k elements, where k is the number of *pivots*
  - $f_s$  is a binary flag that is 1 if there is at least one record X in the Cluster such that X.BasePartition = s, 0 otherwise
- 3. For each partition  $P_i$ , output the clusters (without duplicating clusters) [Reduce]
  - For each Cluster  $C_n$  in partition  $P_i$ 
    - $minFlag = index of minimum value in C_n.[f_1, f_2,$ ...,  $f_k$ ] that is 1
    - If (i = minFlag) then output  $C_n$ , otherwise don't output it (it will be outputted somewhere else)

### Test Setup

#### Datasets

#### 1. Real dataset

- Source: YearPredictionMSD (UCI ML Repository)
- Data type: numeric vector data (90D)
- Size (Scale Factor 1): 200K records

#### 2. Synthetic dataset

Our generator enables the customization of:

- # of records per group and record repetition
- # of Scale Factors (SF) and # of records per SF
- Epsilon value
- Dimensionality
- Format: Line ID, Aggregation Value, Vector
- Size (Scale Factor 1): 200K records

#### **Experiments**

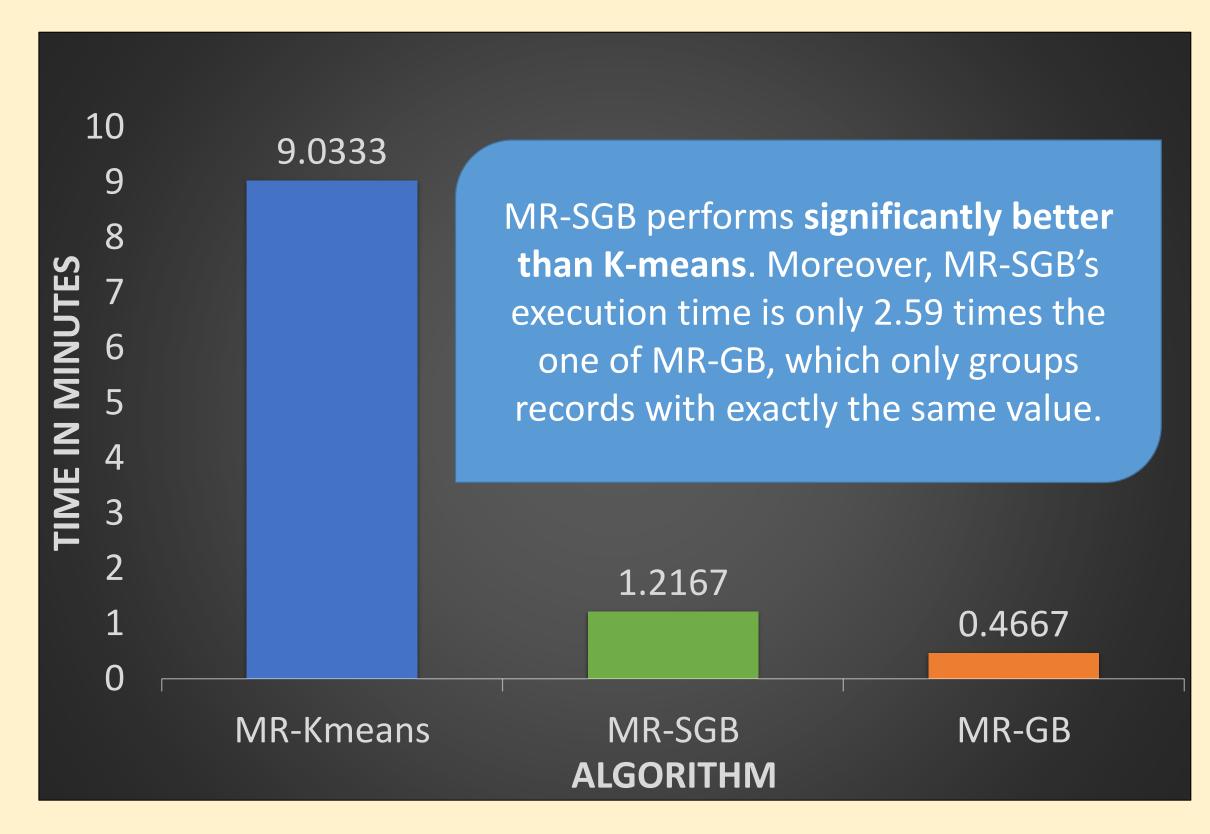
- Execution time varying dataset size (SF1-SF5)
- Execution time varying dimensionality (10D, 25D, 100D, 200D)
- 3. Execution time varying epsilon (1%-5%)

#### Algorithms

- Implemented using Hadoop and MapReduce
- Similarity Group-by (MR-SGB): proposed similarity grouping operator
- 3. K-means (MR-Kmeans): standard clustering algorithm
- 4. Group-by (MR-GB): standard non-similarity-based database grouping operator

### Preliminary Results

- Implemented and compared implemented algorithms using Apache Hadoop
- Compared algorithms: MR-GB, MR-Kmeans, and MR-SGB
- All the algorithms were modified to accept the same input data format



K-means (k=10), SGB (eps=1, pivots=10), GB (count), dataset (2D, small)

### General Algorithm

Key properties of the single-round algorithm:

- We can increase the number of pivots (k) such that all the partitions are small enough to be processed in a single node.
- For the unlikely case that we still have a large partition, we support additional partitioning rounds.

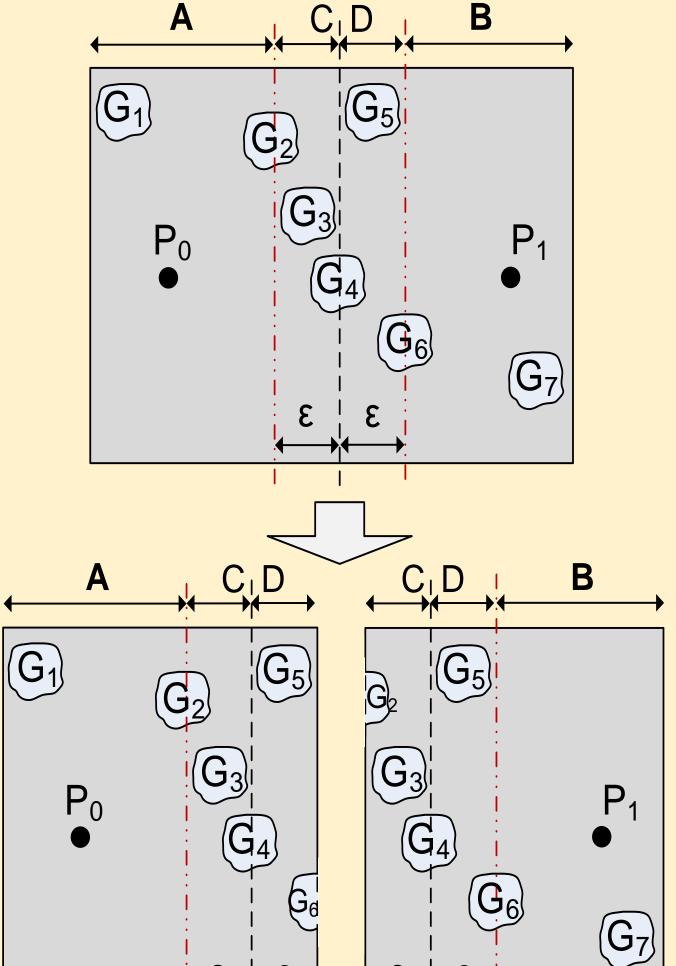
### Overall Algorithm:

- Execute algorithm (round 1)
- For each partition  $P_i$ 
  - If  $P_i$  can be processed in a single node, then we do so
  - Else, we save  $P_i$  in the distributed file system (DFS)
- For each partition  $P_i$  saved into the DFS
  - Execute a new round to re-partition  $P_i$

The algorithm for additional rounds is very similar to the first one, with the difference being that we need to keep track of the history of flags for each partition.

### Example: Case of 2 Pivots

Part1



**Generated Partitions** 

Part0

Initial Dataset (2D space)

Partitioning and Generation of Similarity Groups

### Goals:

- Partition the initial dataset into two partitions such that we can still identify all the similarity groups (G<sub>1</sub>-G<sub>7</sub>)
- Each similarity group should be generated in only one partition

#### **Solution** (using two pivots/partitions):

- Partition the input using two pivots (P<sub>0</sub> and P<sub>1</sub>) such that each point belongs to the partition of its closest pivot
- Additionally, duplicate the points in the ε-windows (C and D). Part0 = A+C+D, Part1 = C+D+B.
- Identify the similarity groups in each partition as follows:

#### In partition Part0: In partition Part1: If group If group Solely in C Solely in A Generate Ignore In A and C In C and D Ignore Generate Solely in C Solely in D Generate Generate Generate In C and D In D and B Generate Solely in B Generate Solely in D Ignore

In the example, similarity groups G<sub>1</sub>, G<sub>2</sub>, G<sub>3</sub>, and G<sub>4</sub> are generated in Part0 while G<sub>5</sub>,G<sub>6</sub>, and G<sub>7</sub> in Part1

### Future Work

- 1. Implement Group-By, K-Means, and Similarity Group-By in Apache Spark to compare the algorithms across the two popular distributed computing frameworks.
- Conduct thorough experimental evaluation using the generated real and synthetic datasets across the two frameworks (Hadoop and Spark).
- Prepare a publication detailing the design, implementation details and performance comparison of Similarity Group-By and alternative algorithms.

### References

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