Gram Matrix Approximation Using Locality Sensitive Hashing on Cluster

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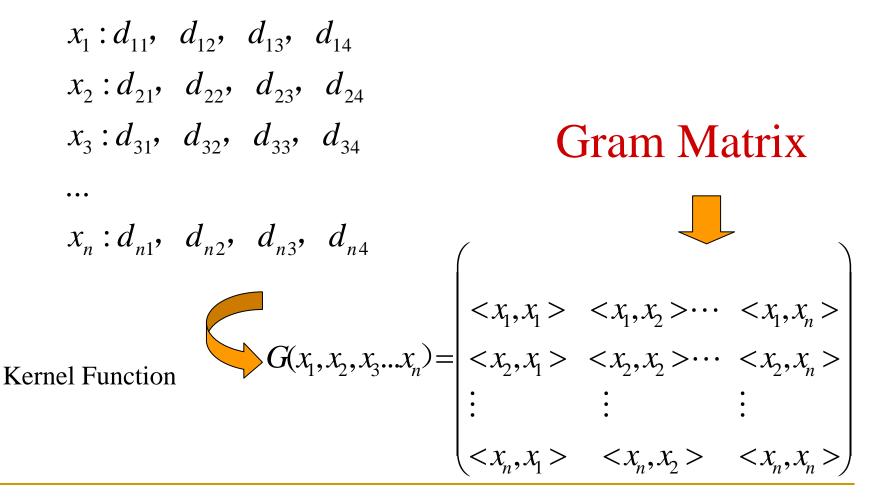


# Outline

 Background and the Problem Gram matrix (G-matrix) The problem we want to solve Approximated G-matrix Scheme How does Hadoop Work? Implementation and Evaluation Conclusion and Future Work

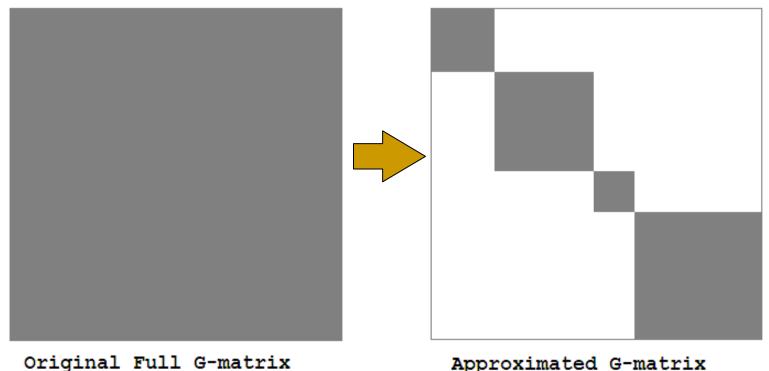
# Background

#### Example



# What if the data set is extremely large?

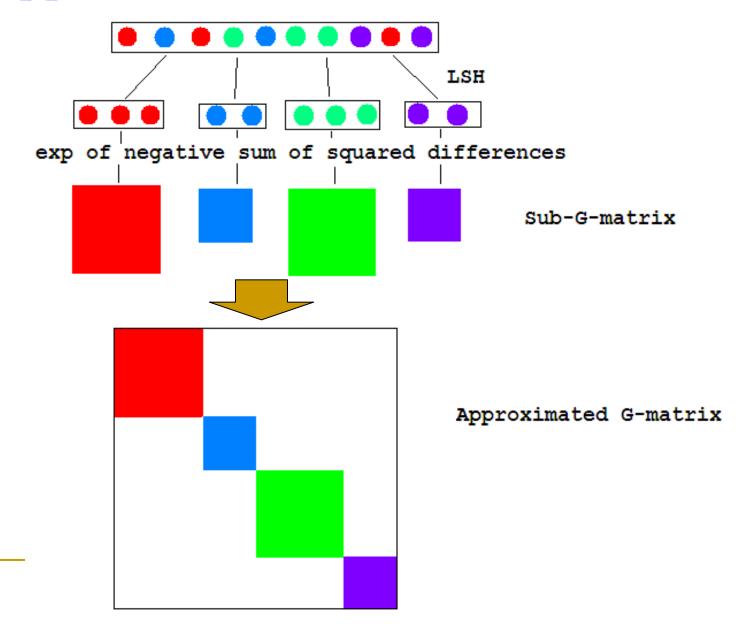
- Partition the dataset into different clusters based on similarity measurement
- Compute the sub-matrix in every cluster



Approximated G-matrix Scheme

- Using locality preserving properties of Locality Sensitive Hashing<sup>[1]</sup> to approximate the original gram matrix
- Parallelize the program on cluster using Hadoop Map-reduce framework

# Approximated G-matrix Scheme (cont.)



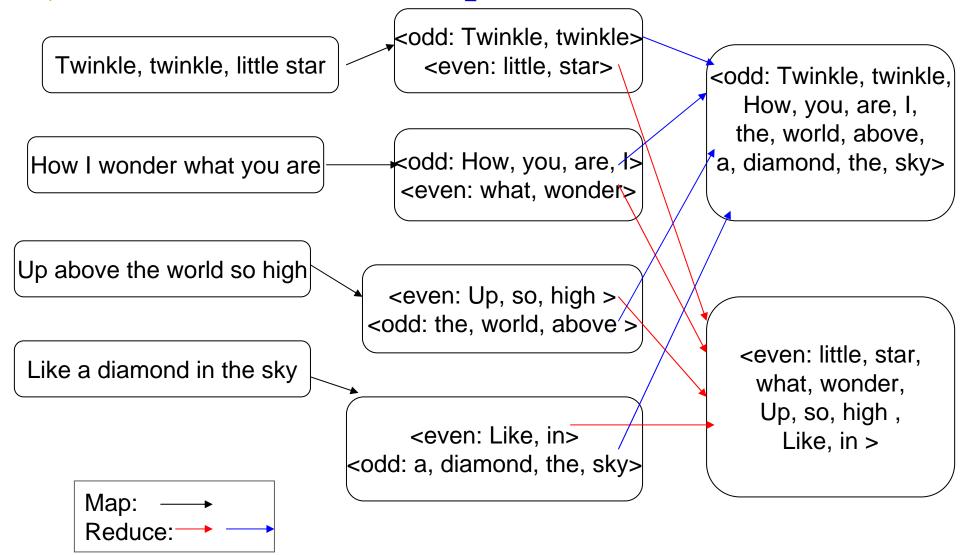
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### How does Map-Reduce Work?

 An example: classify the words in the following nursery rhyme with respect to the parity of word length

Twinkle, twinkle, little star
How I wonder what you are
Up above the world so high
Like a diamond in the sky

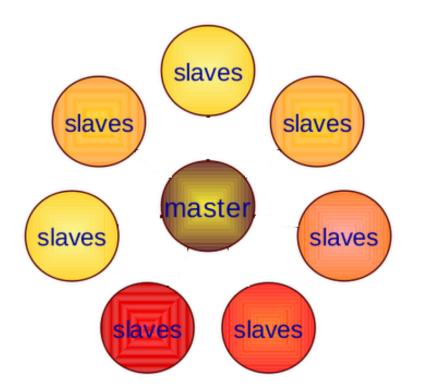
# How does Map-Reduce Work?



#### Look into Hadoop

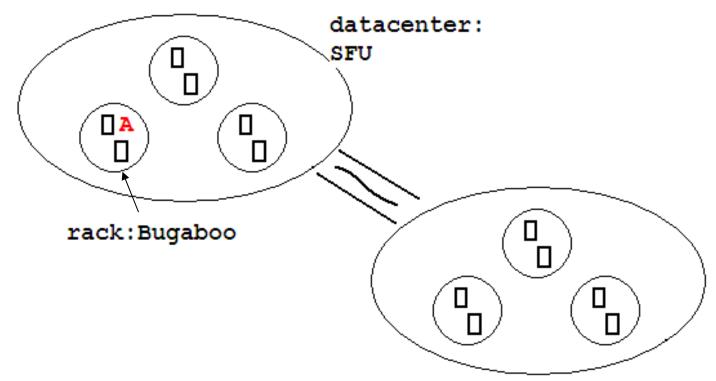


Organization



Master is called JobTracker, slaves called TaskTracker

• Rack, Data centre Determination



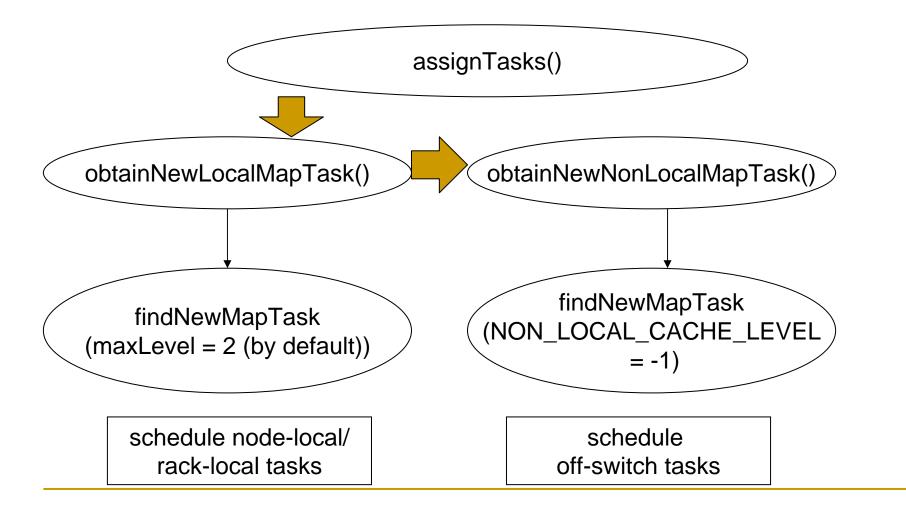
□ The path of machine A: /SFU/Bugaboo/A



- How Input Data is **Split** and **Distributed**?
  - Class InputSplit defines how to do the split of an input file.
  - In most cases Class *FileInputFormat* and its subclass will use 64MB (recommended) to split input file.
  - Function computeReplicationWorkForBlock() in Class FSNamesystem does the replication (default: 3 copies per block).
  - one copy on the local node, one copy on a node in the same rack, and the third copy on a node which is outside the rack.

#### • Task Scheduling

- Task assignment is initiated by a slave (asks for a task), then master decides which task to schedule and responses.
- The Job Tracker knows which node contains the data, and which other machines are nearby. If the work cannot be hosted on the actual node where the data resides, priority is given to nodes in the same rack, then machines in the same datacenter for the nearest located mapping tasks (data split).



### Implementation Using Hadoop

 Step 1: Mapping. For every 64-dimension vector, do LSH in mapping part, the LSH method used is Random Projection from www'07<sup>[2]</sup>. The output pair of this step is <bucketNumber, vectorIndex>.

 $a[i] = \text{vector}[\dim_] \ge \text{threshold}_?1:0, \text{ for every } i \in [0, m-1]$ 

dim\_ and threshold\_ are randomly generated

BucketNumber = 
$$\sum_{i=0}^{m-1} a[i] \cdot 2^i$$

Which is equal to concatenating these binary a[i] bits together. In this way, the maximum number of buckets is  $2^m$ 

# Implementation Using Hadoop

- Step 2: Hadoop merges the vectors which share the same bucket\_number together, into a list.
- Step 3: Reducing. For those vectors hashed into the same bucket, compute the similarity value between every two vectors to form the sub-Gram-matrix. Further, use clustering method.
  - Method of Simlarity Computing

For 64-dimension vector a[] and b[], and pre-set weight[]  $-\sum_{i=0}^{63} weight[i] \cdot (a[i]-b[i])^{2}$ SimValue = e

## Experiment Setup

- Single machine: Intel® Xeon® Processor E5410 (12M Cache, 2.33 GHz, 1333 MHz FSB), 16 GB DRAM. GCC 4.1.3
- Cluster: six machines, each of which is Intel® Core<sup>™</sup>2 Duo Processor E6550 (4M Cache, 2.33 GHz, 1333 MHz FSB), 1 GB DRAM. Hadoop 0.20, JDK 1.6.0\_10
- Dataset property: 64 dimension, the numerical range of every component of the vector is [0,1]

# Evaluation Methods

- Hashing results comparison between cluster and single machine
- Frobenius norm (Fnorm)
  - □ For m-by-n matrix, *Fnorm* =
- Affinity Propagation 3

$$=\sqrt{\sum_{i=1}^{m}\sum_{j=1}^{n}\left|a_{ij}\right|^{2}}$$

- An unsupervised data clustering algorithm
- Basic idea: given a set of data points with unknown cluster structure, the objective is to find a subset to serve as cluster exemplars.
- Approximated G-matrix computing time

# Hashing Results Comparison

#### • Procedure

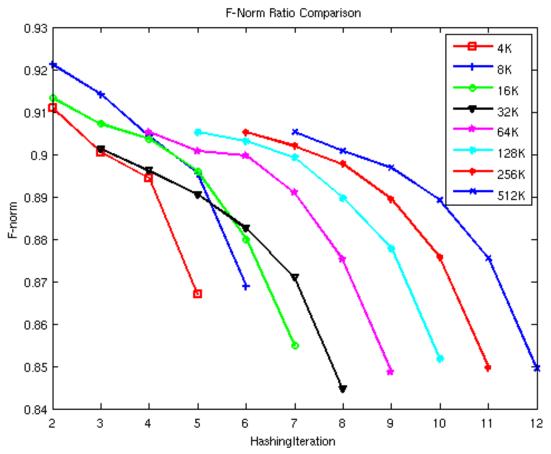
- Write the hashing results into files
- Use the same hashing parameter (hashing iteration 3, 6) and dataset (1K, 8K, 64K, 512K) to do hashing in cluster and single machine code.
- Sort the vector indices in every bucket (for efficiency)
- Count the number of same vectors (N\_same) in the corresponding bucket in cluster and single machine results.
- take the total of the two bucket size, and then subtract by 2\* N\_same, therefore, we get how many different vectors are in the same bucket

#### Result

The two implementations get exactly the same results, as expected.

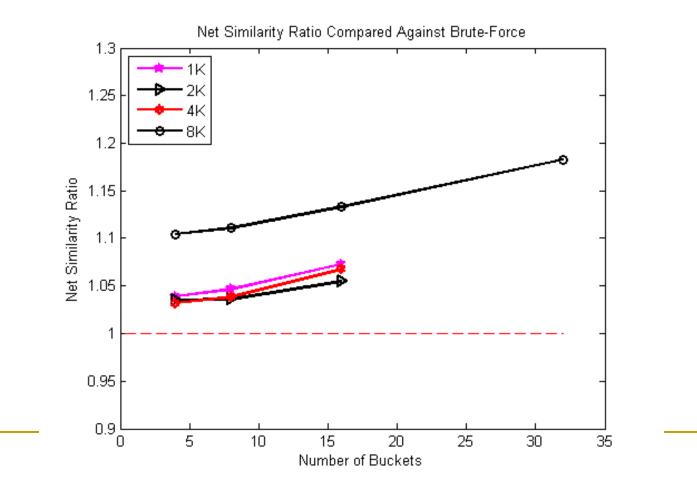
### Frobenius Norm Comparison

 Comparison between Approximated G-matrix and Original matrix



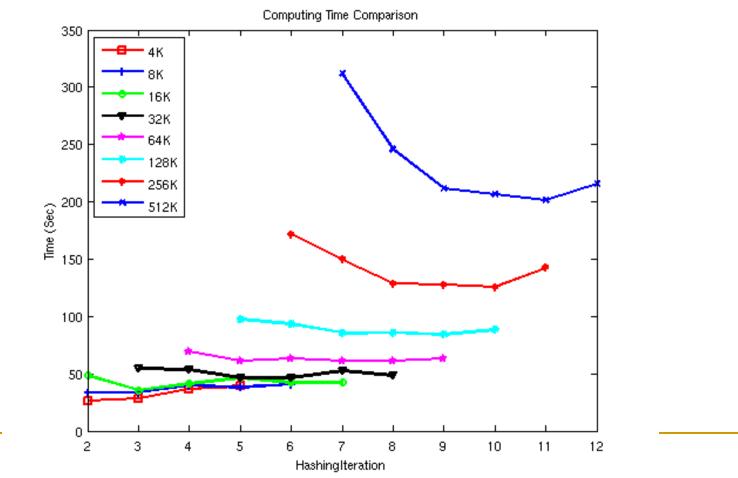
### Affinity Propagation Comparison

• The difference between Approximated and original matrix is small (5%-12% difference when 16 buckets)



#### Approximated G-matrix computing time

• Time comsumed grows sublinearly with the growth of data set size.



### Conclusion and Future Work

- High precision (6%-12%difference) can be achieved in Gram Matrix approximation using Locality Sensitive Hashing (LSH)
  - Frobenius norm (from matrix property side)
  - Clustering error (Net Similarity, from application side)
- LSH and cluster enables the parallelization of Gram matrix approximation and reduction of computing time
- Experiment on large scale cluster (Westgrid) to do optimization, further time reduction expected with guaranteed precision

# Reference

- Gionis, A.; Indyk, P., Motwani, R. (1999)., "Similarity Search in High Dimensions via Hashing". Proceedings of the 25th Very Large Database (VLDB) Conference.
- 2. Gurmeet S. Manku, Arvind Jain, Anish D. Sarma: Detecting nearduplicates for web crawling, In WWW '07: Proceedings of the 16th international conference on World Wide Web (2007), pp. 141-150.

Thank you

Questions and Suggestions ( on deployment of cluster with 2304 cores)