

# Comparison of Two Way Join Algorithms used in MapReduce Frame work for handling BigData

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**Abstract-Due to the advent of new technologies, devices, and communication means like social networking sites, the amount of data produced by mankind is growing rapidly every year. The amount of data produced by us from the beginning of time till 2003 was 5 billion gigabytes. A vast amount of research going on into how best to handle and process huge amounts of data which is generated by different media and social networks. One such idea for processing enormous quantities of data is Google's Map/Reduce. MapReduce is the heart of Hadoop. It is this programming paradigm that allows for massive scalability across hundreds or thousands of servers in a Hadoop cluster. The MapReduce concept is fairly simple to understand for those who are familiar with clustered scale-out data processing solutions. The MapReduce framework is increasingly being used widely to analyze large volumes of data. One of the techniques that framework is join algorithm. Join algorithms can be divided into two groups: Reduce-side join and Map-side join. The aim of our work is to compare existing join algorithms which are used by the MapReduce framework. We have compared Reducer-side merge join and Map-side replication-join in terms of pre-processing, the number of phases involved, whether it is sensitive to data skew, whether there is need for distributed Cache, memory overflow.**

**Key Words: Join algorithms, MapReduce, optimization.Semi join,Bloom Filter,Hadoop**

## I. INTRODUCTION

Map/Reduce[1] was first introduced by Google engineers - Jeffrey Dean and Sanjay Ghemawat [9]. It was designed for and is still used at Google for processing large amounts of raw data (like crawled documents and web-request logs) to produce various kinds of derived data (like inverted indices, web-page summaries, etc.). It is a simple yet powerful framework for implementing distributed applications without having extensive prior knowledge of the intricacies involved in a distributed system. It is highly scalable and works on a cluster of commodity machines with integrated mechanisms for fault tolerance. Data-intensive applications include large-scale data warehouse systems, cloud computing, data-intensive analysis. These applications have their own specific computational workload. For example, analytic systems produce relatively rare updates but heavy select operation with millions of records to be processed, often with aggregations.

Since its inception at Google, Map/Reduce has found many adopters. Among them, the prominent one is the Apache Software Foundation, which has developed an Open-Source version of the Map/Reduce framework called Hadoop [2]. Hadoop boasts of a number of large web-based corporates like Yahoo, Facebook, Amazon, etc., that use it for various kinds of data-warehousing purposes. Facebook for instance, uses it to store copies of internal logs and uses it as a source for reporting and machine learning. See [4] for other companies that use Hadoop. Applications for large-scale data analysis use such techniques as parallel DBMS, MapReduce (MR) paradigm, and columnar storage. Applications of this type process multiple data sets. This implies need to perform several join operation. It's known join operation is one of the most expensive operations in terms both I /

O and CPU costs. Join algorithms is not directly supported in MapReduce. There are some approaches to solve this problem by using a high-level language PigLatin, HiveQL for SQL queries or implementing algorithms from research papers. The aim of this work is to generalize and compare existing equi-join algorithms with some optimization techniques.

## II. JOIN ALGORITHMS AND OPTIMIZATION TECHNIQUES

In this section we consider various techniques of two-way joins in MapReduce framework. Join algorithms can be divided into two groups: Reduce-side join and Map-side join. The pseudo code presented in Listings, where R – right dataset, L –left dataset, V – line from file, Key – join key, that was parsed from a tuple, in this context tuple is V.

### 2.1 Two-Way Joins

#### 2.1.1 Reduce-Side join

Reduce-side join is an algorithm which performs data pre-processing in Map phase, and direct join is done during the Reduce phase. Join of this type is the most general without any restriction on the data. Reduce-side join is the most time-consuming, because it contains an additional phase and transmits data over the network from one phase to another. In addition, the algorithm has to pass information about source of data through the network. The main objective of the improvement is to reduce the data transmission over the network from the Map task to the Reduce task by filtering the original data through semi-joins. Another disadvantage of this class of algorithms is the sensitivity to the data skew, which can be addressed by replacing the default hash partitioner with a range partitioner.

There are three algorithms in this group:

- General reducer-side join
- Optimized reducer-side join
- Hybrid Hadoop join

General reducer-side join is the simplest one. The same algorithms are called Standard Repartition Join in [6]. The abbreviation is GRSJ and pseudo code is presented below:

General reducer-side join:

```

-----
Map (K: null, V from R or L)
    Tag = bit from name of R or L;
    emit (Key, pair(V,Tag));
Reduce (K': join key, LV: list of V with key K')
create buffers B r and B l for R and L;
for t in LV do
    add t.v to B r or B l by t.Tag;
for r in B r do
for l in B l do
    emit (null, tuple(r.V,l.V));
  
```

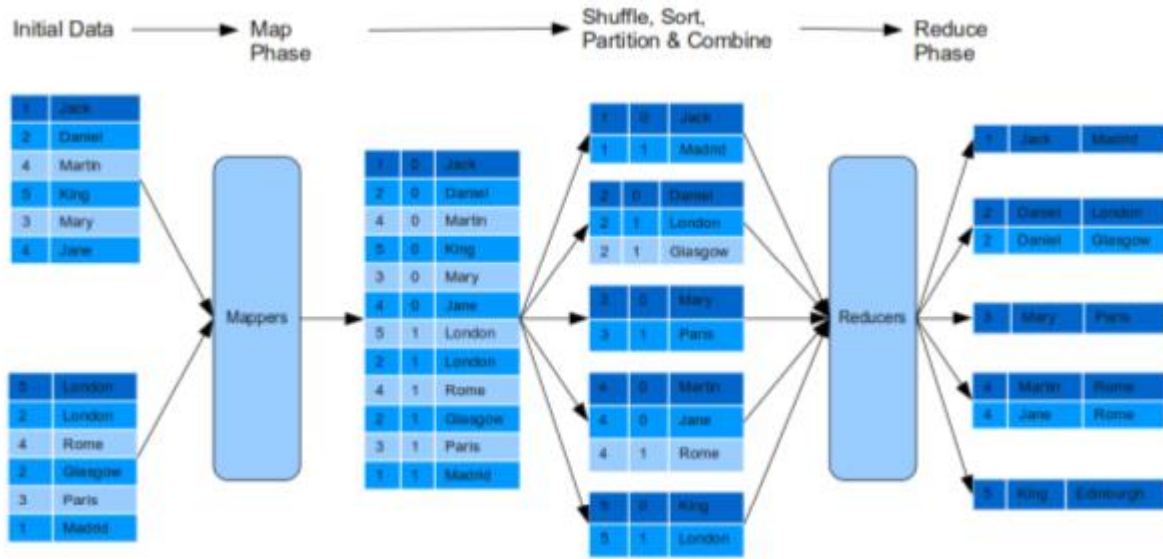


Figure-1. Reduce Side Join - Two Way

This algorithm has both Map and Reduce phases. In the Map phase, data are read from two sources and tags are attached to the value to identify the source of a key/value pair. As the key is not effected by this tagging, so we can use the standard hash partitioner. In Reduce phase, data with the same key and different tags are joined with nested-loop algorithm. The problems of this approach are that the reducer should have sufficient memory for all records with a same key; and the algorithm sensitivity to the data skew.

### 2.1.2 Map-Side join

Map-side join is an algorithm without Reduce phase. This kind of join can be divided into two groups. First of them is partition join, when data previously partitioned into the same number of parts with the same partitioner. The relevant parts will be joined during the Map phase. This map-side join is sensitive to the data skew. The second is in memory join, when the smaller dataset send whole to all mappers and bigger dataset is partitioned over the mappers. The problem with this type of join occurs when the smaller of the sets cannot fit in memory.

There are three methods to avoid this problem:

- JDBM-based map join,
- Multi-phase map join,
- Reversed map join

Limitation of Map-Side Join:

Limitation	Why
All datasets must be sorted using the same comparator	The sort ordering of the data in each dataset must be identical for datasets to be joined.
All datasets must be partitioned using the same partitioner.	A given key has to be in the same partition in each dataset so that all partitions that can hold a key are joined together.
The number of partitions in datasets must be identical	A given key has to be in the same partition in each dataset so that all partitions that can hold a key are joined together.

### 2.1.3 Broadcast Join

Broadcast Join was studied by Blanas et. al. in [6]. If one of the datasets is very small, such that it can fit in memory, then there is an optimization that can be exploited to avoid the data transfer overhead involved in transferring values from Mappers to Reducers. This kind of scenario is often seen in real-world applications. For

instance, a small users database may need to be joined with a large log. This small dataset can be simply replicated on all the machines. This can be achieved by simply using `-files` or `-archive` directive to send the file to each machine while invoking the Hadoop job. Broadcast join is a Map-only algorithm.

#### *Map Phase*

The Mapper loads the small dataset into memory and calls the map function for each tuple from the bigger dataset. For each (key, value), the map function probes the in- memory dataset and finds matches. This process can be further optimized by loading the small dataset into a Hashtable. It then writes out the joined tuples. The below shown configure function is part of the Mapper class and is called once for every Mapper. The below code reads the 'broadcasted' file and loads the same into an in- memory hash table.

#### *Loading the small dataset into a Hash table:*

```

-----
public void configure ( JobConf conf ) {
//Read the broadcasted file
T1 = new File ( conf . get (" broadcast . file "));
//Hashtable to store the tuples
ht = new HashMap < String , ArrayList < String > >();
BufferedReader br = null;
String line = null;
try{
br = new BufferedReader (new FileReader ( T1 ));
while(( line = br . readLine () ) !=null)
{
String record [] = line . split ("\t" , 2);
if( record . length == 2)
{
//Insert into Hashtable
if( ht . containsKey ( record [0]) )
{
ht . get ( record [0]) . add ( record [1]);
}
else
{
ArrayList < String > value = new ArrayList <
String >();
value . add ( record [1]);
ht . put ( record [0] , value );
}
}
}
}
catch( Exception e)
{
e . printStackTrace ();
}
}
}
-----

```

The next piece of code is the actual map function that receives the records from the HDFS and probes the HashTable containing the tuples from the broadcasted file. Notice that it takes care of duplicate keys as well.

#### *Broadcast Join Map Function:*

```

-----
public void map( LongWritable lineNumber , Text value ,
OutputCollector < Text , Text > output , Reporter
reporter )
throws IOException {
String [] rightRecord = value . toString () . split ("\t" ,2);
if( rightRecord . length == 2)
{
for( String leftRecord : ht . get ( rightRecord [0]) )

```

```

{
    output . collect (new Text ( rightRecord [0]) , new Text
    ( leftRecord + "\t" + rightRecord [1] ) );
}
}
}

```

-----

Broadcast Join benefits from the fact that it uses a small ‘local’ storage on the individual nodes instead of the HDFS. This makes it possible to load the entire dataset into an in-memory hash table, access to which is very fast. But on the down side, it will run into problems when both the datasets are large and neither of them can be stored locally on the individual nodes.

### III. SEMIJOIN

Sometimes a large portion of the data set does not take part in the join. Deleting of tuples that will not be used in join significantly reduces the amount of data transferred over the network and the size of the dataset for the join. This preprocessing can be carried out using semi-joins by selection or by a bitwise filter.

However, these filtering techniques introduce some cost (an additional MR job), so the semi-join can improve the performance of the system only if the join key has low selectivity. There are three ways to implement the semi-join operation:

- a semi-join using bloom-filter,
- semi-join using selection,
- an adaptive semi-join.

Bloom-filter is a bit array that defines a membership of element in the set. False positive answers are possible, but there are no false-negative responses in the solution of the containment problem. The accuracy of the containment problem solution depends on the size of the bitmap and on the number of elements in the set. These parameters are set by the user. It is known that for a bitmap of fixed size  $m$  and for the data set of  $n$  tuples, the optimal number of hash functions is  $k=0.6931*m/n$ . In the context of MapReduce, the semi-join is performed in two jobs. The first job consists of the Map phase, in which keys from one set are selected and added to the Bloom-filter. The Reduce phase combines several Bloom-filters from first phase into one. The second job consists only of the Map phase, which filters the second data set with a Bloom-filter constructed in previous job.

The accuracy of this approach can be improved by increasing the size of the bitmap. However in this case, a larger bitmap consumes more amounts of memory. The advantage of this method is its the compactness. The performance of the semi-join using Bloom-filter highly depends on the balance between the Bloom-filter size, which increases the time needed for its reconstruction of the filter in the second job, and the number of false positive responses in the containment solution. The large size of the data set can seriously degrade the performance of the join.

*Semi-join using Bloom Filter*

-----

**Job 1:** construct Bloom filter

```

Map (K:null, V from L)
  Add Key to BloomFilter B1
close() //for Map phase
emit(null, B1);
Reduce (K?: key, LV) //only 1 Reducer
  for l in LV do
    union filters by operation Or
close() // for Reduce phase
  write resulting filter into file;

```

**Job 2:** filter dataset

```

init() //for Map phase
  read filter from file in B1
Map (K:null, V from R)
  if (Key in B1) then
emit (null, V);

```

**Job 3:** do join with L dataset and filtered dataset from Job 2.

---

### Semi join with selection:

---

Job 1: find unique keys  
 Map (K:null, V from L)  
 Create HashMap H;  
 if (not Key in H) then  
   add Key to H;  
 emit (Key, null);  
 Reduce (K': key, LV) //only one Reducer  
 emit (null,key);

Job 2: filter dataset  
 init() //for Map phase  
 add to HashMap H unique keys from job 1;  
 Map (K:null, V from R)  
 if (Key in H) then  
   emit (null,V);

Job 3: do join with L dataset and filtered dataset from Job 2.

---

Semi-join with selection extracts unique keys and constructs a hash table. The second set is filtered by the hash table constructed in the previous step. In the context of MapReduce, the semi-join is performed in two jobs. Unique keys are selected during the Map phase of the first job and then they are combined into one file during the Map phase. The second job consists of only the Map phase, which filters out the second set. The semi-join using selection has some limitations. Hash table in memory, based on records of unique keys, can be very large, and depends on the key size and the number of different keys.

The Adaptive semijoin is performed in one job, but filters the original data on the flight during the join. Similar to the Reduce-side join at the Map phase the keys from two data sets are read and values are set equal to tags which identify the source of the keys. At the Reduce phase keys with different tags are selected. The disadvantage of this approach is that additional information about the source of data is transmitted over the network.

### *Adaptive Semijoin*

---

Job 1: find keys which are present in two datasets  
 Map (K:null, V from R or L)  
 Tag = bit from name of R or L;  
 emit (Key,Tag);  
 Reduce (K': join key,  
 LV: list of V with key K')  
 Val = first value from LV;  
 for t in LV do  
 if (not Val==Val2) then  
   emit (null, K');

Job 2: before joining it is necessary to filter the smaller dataset  
 dataset by keys from the Job 1 that will be loaded into hash map.  
 Then the bigger dataset is joined with filtered one

---

## VI. EXPERIMENTS AND ANALYSIS

### *Dataset*

Data are the set of tuples, which attributes are separated by a comma. Tuple is split into a pair of a key and a value, where value is the remaining attributes. Generation of synthetic data was done as in [4]. Join keys are distributed randomly except experiment with the data skew.

### *Cluster configuration*

HYB 1 data 2 - yes - Part size is large Hash

MSPJ 2 data 1 - yes - Part size is large Hash

Cluster consists of three virtual machines, where one of them is master and slave at the same time, the remaining two are the slaves. Host configuration consists of 1 processor, 512 mb of memory for nodes, 5 gb is the disk size. Hadoop 20.203.0 runs on Ubuntu 10.10.

*The General Case:*

The base idea of this experiment is to compare executions time of different phases of various algorithms. Some parameters are fixed: the number of Map and Reduce tasks is 3, the input size is  $10^4 \times 10^5$  and  $10^6 \times 10^6$  tuples.

*Performance of two-way joins on increasing input data size*

A series of experiments were conducted to compare the runtime and scalability of the various algorithms. The first of these involved running the three two-way algorithms mentioned earlier on increasing data sizes .

Both datasets involved in the joins had the following characteristics -

- Same number of tuples
- Same key space
- #keys = 1/10 #tuples

We tested on two kinds of key distributions as shown in Figure 2 and Figure 3. In the case of skewed data, one of the datasets had a key that was repeated 50% of the times. Which means that given n rows, n / 2 of them were the same key. The other dataset had uniform key distribution

Concentrating first on the Uniform Key Distribution graph (Figure 2), it can be seen that all the three algorithms gave comparable numbers at lower input data sizes. But Map-side join always performed the best, even with large input datasets. Broadcast join deteriorated the fastest as the data size increased. This is because a larger file had to be replicated across all the machines in the cluster as the input size increased. Figure 3 shows something interesting. Map-side join gave a worse time than the other two algorithms for the input with 5 million rows when then input dataset had skewed key distribution. This can be attributed to the unequal partitioning that will occur due to the skew.

Apart from testing for runtimes, we also checked for the amount of data transferred over the network from the machines running the Reduce tasks. A simple number like the number of bytes transferred might not be very meaningful. Hence we took the ratio as

*Bytes transferred over the network*

*Initial Data Size*

We call this the Data Shuffle ratio. This ratio is interesting because it gives an idea of how well Hadoop achieves data locality and whether it improves with bigger data sizes. Table 1 and Table 2 show the actual run-times of the different algorithms shown in Figure 2 and Figure 3. Also given are the number of bytes shuffled across machines from the Map phase to the Reduce phase in the Reduce-Side Join. Note that Map-Side Join and Broadcast Join do not have a reduce side and hence no data is shuffled across machines between the Map phase and the Reduce phase. The sixth column shows the time taken for pre-processing in the case of the Map-Side Join algorithm.

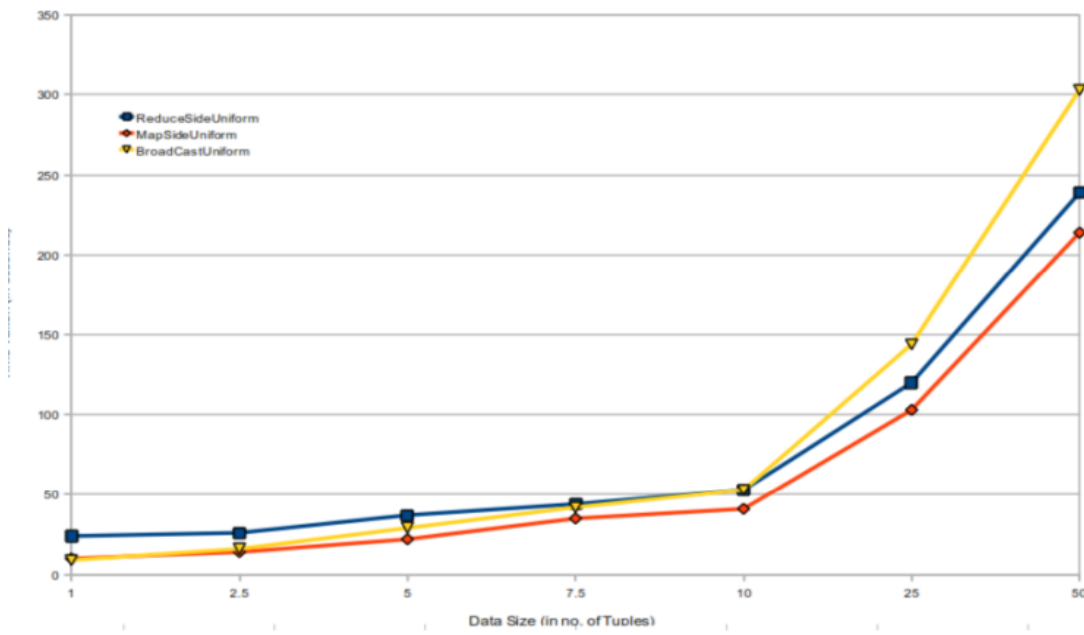


Figure-2. Uniform Key Distribution

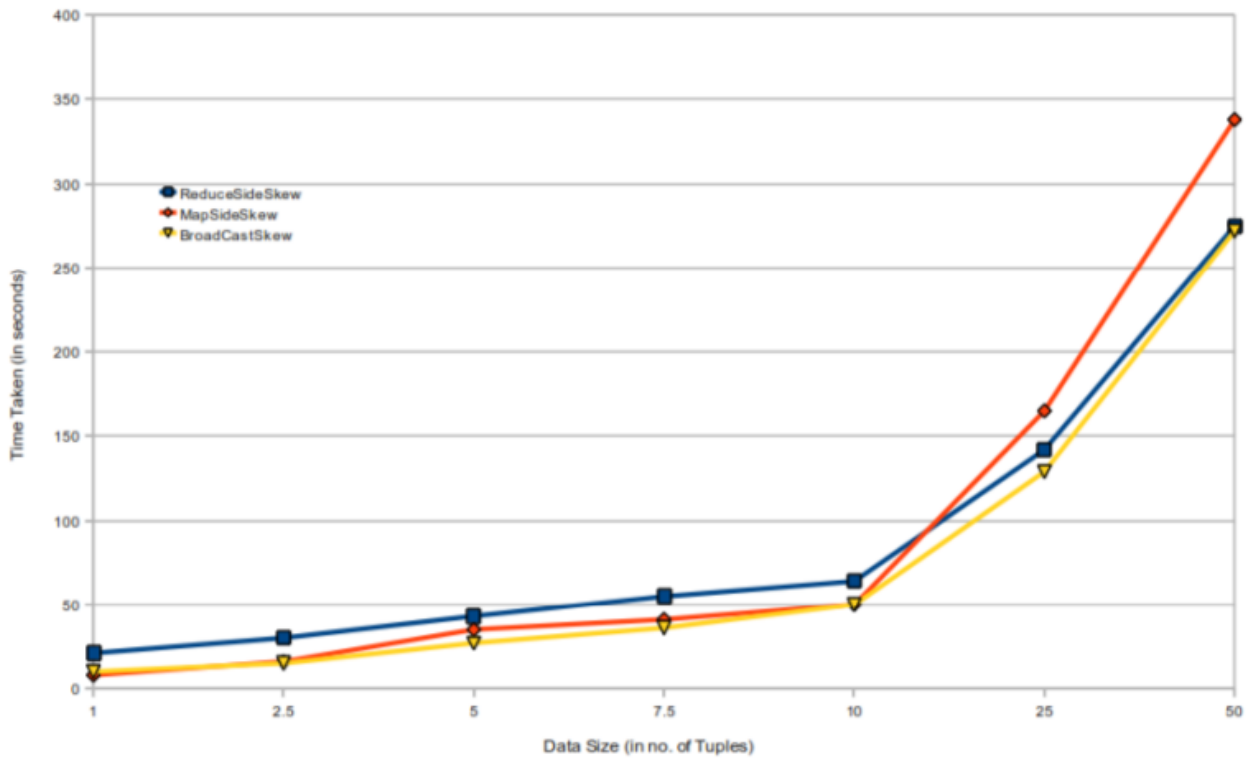


Figure-3. Skewed Key Distribution



Table -1 Uniform Key Distribution

Tuples (x100000)	Reduce-Side Join	Data Shuffled (bytes)	Data Shuffle Ratio	Map-Side Join	Pre-Processing	Broadcast Join
1	24s	9178000	1.150	9s	38s	10s
2.5	26s	23278000	1.148	14s	45s	16s
5	37s	46778000	1.147	22s	46s	29s
7.5	44s	70278000	1.147	35s	47s	42s
10	53s	93778000	1.147	41s	49s	53s
25	120s	237778000	1.144	103s	56s	144s
50	239s	477778000	1.144	214s	71s	303s

Table -2 Skewed Data Distribution

Tuples (x100000)	Reduce-Side Join	Data Shuffled (bytes)	Data Shuffle Ratio	Map-Side Join	Pre-Processing	Broadcast Join
1	21s	8194793	1.127	8s	41s	10s
2.5	30s	23243259	1.148	16s	42s	15s
5	43s	46805826	1.147	35s	42s	27s
7.5	55s	70118029	1.147	41s	46s	36s
10	64s	93806021	1.147	50s	46s	50s
25	142s	238055988	1.144	165s	54s	129s
50	275s	478056409	1.144	338s	76s	272s

There are two main things worth noticing in Table 1 and Table 2. The first is that the Data Shuffle ratio remains around 1.14 for both uniform and skewed key distribution. This is because as the data size increases, Hadoop is able to better manage data distribution and data locality. The other thing is that though Map-Side performed best most of the times, it took an average time of 50.29 seconds (uniform key distribution) and 49.57 seconds (skewed key distribution) for the pre-partitioning phase. This may not seem a lot, but is likely to be quite high if the input datasets are much larger than the ones used in this experiment.

*Semi-Join Comparisons:*

The main idea of this experiment is to compare different semi-join algorithms. These parameters are fixed: the number of Map and Reduce tasks is 3, the bitmap size of Bloom-filter is  $25 \times 10^5$ , the number of hash-functions in Bloom-filter is 173, built-in Jenkins hash algorithm is used in Bloom-filter. Adaptive semi-join (ASGRSJ) does not finish because of memory overflow. The abbreviation of Bloom-filter semi-join for GRSJ is BGRSJ. The abbreviation of semi-join with selection for GRSJ is SGRSJ respectively.

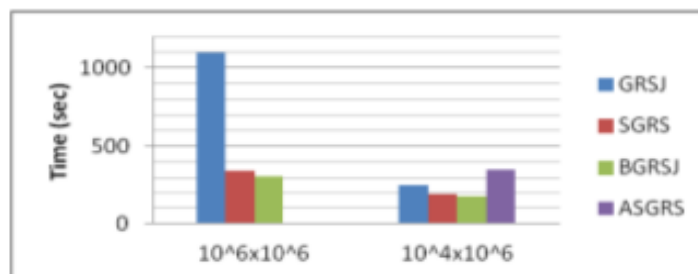


Figure-4. Comparison of different semi-join implementations

## V. CONCLUSION

We explored the practicality of using Map/Reduce for joining datasets and found it quite easy to use and implement over different sizes of cluster. In this work we found some quite interesting results from our experiments on Join algorithms using Map/Reduce. Even though our clusters were of relatively small sizes, we could notice from our experiments how the Hadoop framework exploited the cluster architecture when more nodes were added (as was seen with the Data Shuffle ratio that remained the same throughout our experiments). This gave us a first-hand feel of why Hadoop seems to be enjoying such a high popularity among data warehousing specialists. We found that Map-Side joins performs the best when the datasets have relatively uniform key distribution. But this comes at the cost of a pre-processing step. Broadcast Join is a better option when one of the datasets is much smaller compared to the other and the dataset is small enough to be easily replicated across all the machines in the cluster. It deteriorated the fastest when the input data size was increased. Reduce-Side Join gave almost the same performance for both uniform and skewed key distributions whereas Map-Side Join seemed to give slightly unpredictable results with skewed data.

In this work we describe the state of the art in the area of massive parallel processing, presented our comparative study of these algorithms reduced side join, map side join, broad cast join and semi-join. In future work multi way join can be tested and analysed using more number of Hadoop clusters.

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