## Efficient Processing of k Nearest Neighbor Joins using MapReduce

## I. SUMMARY

The paper illustrates a efficient processing of kNN Joins using MapReduce

In the preliminaries, the authors shows the basic concepts of kNN Join, MapReduce Framework, and particularly the Voronoi Diagram-based Partitioning, which is the mathematic fundamental for partitioning the input set R and S.

A native and straightforward idea of performing kNN join in MapReduce is similar to the hash join algorithm. R is split into disjoint subsets, each subset  $R_i$  is distributed to a reducer, and without any pruning rule, the entire set S has to be sent to each reducer to be joined with  $R_i$ .

In order to reduce the shuffling cost, a better strategy is that R is partitioned into N disjoint subsets and for each subset  $R_i$ , find a subset of  $S_i$  that  $R_i \ltimes S = R_i \ltimes S_i$  and  $R \ltimes S = \bigcup_{1 \leq i \leq N} R_i \ltimes S_j$ . So  $S_i$  is sent to the reducer that  $R_i$  belongs to and the KNN join is performed between  $R_i$  and  $S_i$  only.

For the purpose of reducing the size of  $S_i$ , we need to derive a distance bound based on the partitionning of R. Using basic geometry knowledge, the upper bound distance from  $s \in P_j^S$  to  $r \in P_i^R$ , denoted as  $ub(s, P_i^R)$ , can be calculated easily. Then a bound(denoted as  $\theta_i$ ) of the KNN distance for all objects in  $P_i^R$ , can be touched by continually adding  $ub(s, P_i^R)$ , which  $s \in KNN(p_j, P_j^S)$ , to a priority queue for each  $P_j^S$ , until the size grows to k and there is no more  $ub(s, P_i^R)$  smaller than the top of the queue.

The corresponding lower bound can be derived in a similar way, and after that, we get the necessary condition s is assigned to  $S_i$ :

$$|s, p_i| \ge LB(p_i^S, P_i^R) \tag{1}$$

So, given the input set R and S, the partition of  $R_i$  and  $S_i$  should be calculated first in the map procedure, and then each  $R_i, S_i$  pair will be sent to the same reducer to execute KNN join respectively. The details are shown below, and it takes three steps to complete the kNN Join.

- First, the master node invokes a preprocessing step, which takes the original R and S as input and finds out a set of pivot objects based on the input dataset R. The pivots can be got by Random Selection, Farthest Selection or kmeans selection, and the pivots is used to create a Voronoi diagram, which can help partition objects in R effectively while preserving their proximity.
- Second, the first MapReduce job consists of a single Map phase, which takes the selected pivots and datasets R and S as the input. The output of the mapping phase is a partitioning on R, based on the Voronoi diagram of

the pivots. Meanwhile, the mappers also collects some statistics about each partition  $R_i$ .

The partitioning on R can be represented as a table, which including each object o along with its partition id, original dataset name (R or S), distance to the closest pivot.

The statistics are kept in two summary table  $T_R$  and  $T_S$ .  $T_R$  maintains the following information for every partition of R: the partition id, the number of objects in the partition, the minimum distance  $\mathrm{L}(P_i^R)$  and maximum distance  $\mathrm{U}(P_i^R)$  from an object in partition  $P_i^R$  to the pivot. Moreover,  $T_S$  also maintains the distances between objects in  $\mathrm{KNN}(p_i, P_i^S)$  and  $p_i$ 

• Third, taking T<sub>R</sub>, T<sub>S</sub> and statistics from steps 2 as input, S<sub>i</sub> will be built with the constraint of equation 1. The R<sub>i</sub> and S<sub>i</sub> will be sent to the same reducer to execute R<sub>i</sub> κ S<sub>i</sub>. In the reducer, each value pair will be parsed to derive the partition P<sub>i</sub><sup>R</sup> and subset S<sub>i</sub> that consists of P<sub>j1</sub><sup>S</sup>,..., P<sub>jM</sub><sup>S</sup>, then, a similar pruning process as the second step does, can be applied to S<sub>i</sub> to get LB(P<sub>j</sub><sup>S</sup>, P<sub>i</sub><sup>R</sup>) for every KNN distance, θ<sub>i</sub>, for all objects of P<sub>i</sub><sup>R</sup>. Hence, we can issue a range search with query and threshold θ<sub>i</sub> over dataset S<sub>i</sub>. After checking all partitions of S<sub>i</sub> with pruning and updating KNN(r,S), the reducer outputs KNN(r,S).

The way  $S_i$  derived will bring replications of S. To minimize the number of replicas of objects in S, a intuitive way is to increase the number of pivots. However, this requires a large number of reducers, which may not be practical. A natural idea is to divide partitions of R into disjoint groups. There are two strategies for grouping.

- Geometric Grouping first select N  $p_i$  as the basic element of each group  $G_i$ , which is faraway from each other, and assign the rest  $p_j$  to the nearest group  $G_i$ .
- Greedy Grouping tries to minimize the increasing of  $RP(S,G_i)$  when assigning a new partition  $P_j^R$  to  $G_i$ . But for the sake of reducing the computation cost, once  $\exists s \in P l^S$  satisfying the necessity of being added to  $S_i$ , we add all objects of partition  $P_i^S$  to  $RP(S,G_i)$ .

The draw back of the algorithm is that, improper pivot selection will cause large difference in partition size, which degrades performance due to unbalanced workload. And this paper doesn't present a pivot selection algorithm that will guarantee the uniformity of partitioning.