SIGMOD Programming Contest 2018 $Team \, Dataurus$

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Problem

Task: The task is to **evaluate batches of join queries** on a set of pre-defined relations. Each join query specifies a set of relations, (equality) join predicates, and selections (aggregations). The maximum number of relations that a query joins is four. The challenge is to execute the queries as fast as possible without (much) prior indexing. For this reason we only have 1 second for preprocessing.

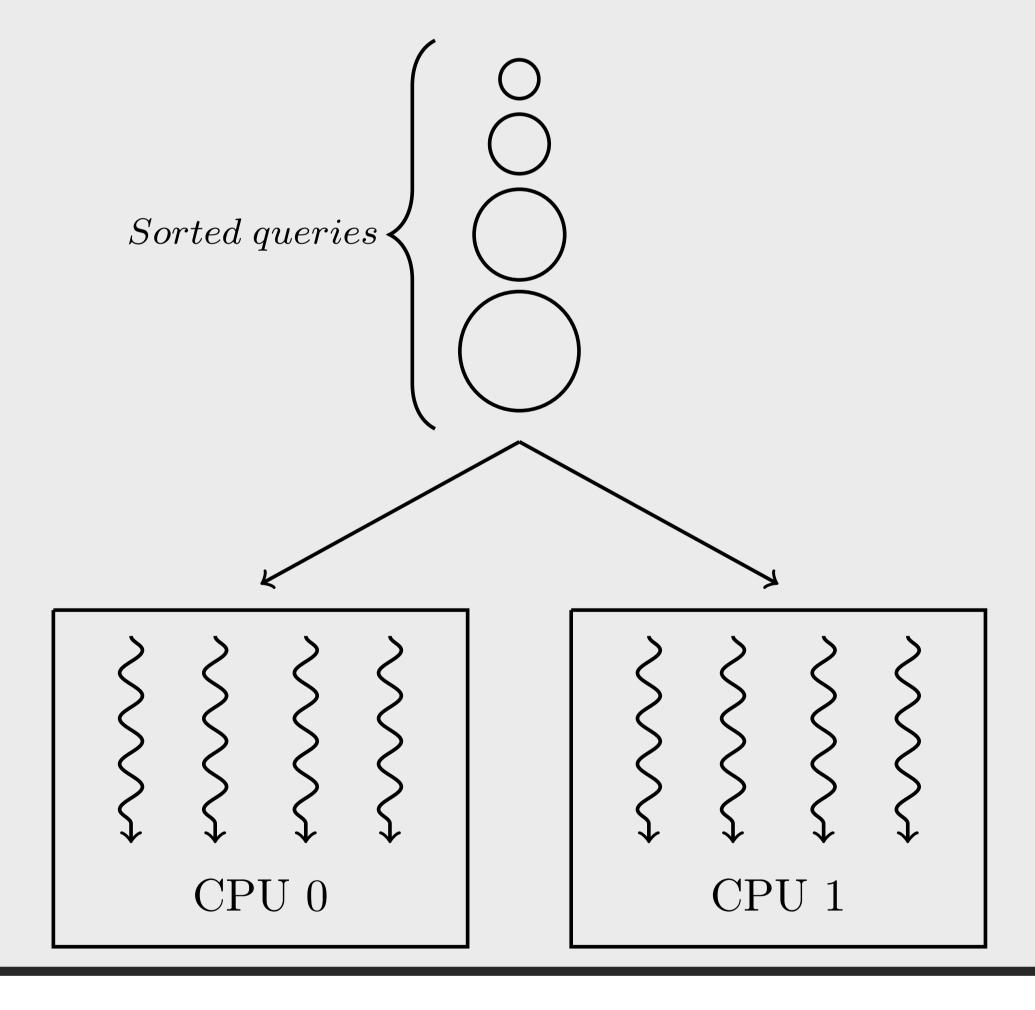
Hardware: 2x Intel Xeon E5-2660 v2 CPU (2.20 GHz, 3.00 GHz turbo) - 20 Cores / 40 Hyperthreads - 256 GB RAM

1. Preprocessing

- Data is stored in **column-store** fashion for cache efficiency.
- Parse the input data in parallel and collect **statistics** per column, such as
 - minimum element value
 - maximum element value
 - number of tuples
 - number of distinct elements
 - spread of values

3. Query Scheduling

- A global queue is used to distribute the queries into two NUMA regions.
- Sort the queries of a batch in descending order (heavy first) based on the estimations (cost) of the optimizer and push to the queue.
- As soon as a cpu finishes the execution of a query, the next one in the queue is assigned to it immediately.
- Clone the dataset on both NUMA regions for better memory bandwidth utilization.



5. Join Operator

- Use Parallel Radix Hash Join^a to compute the join result
 - The two input relations R and S are divided into **partitions**.
 - * Each thread receives a chunk of data and computes a **histogram** over the input data, so that the exact output size is known for each thread and each partition.
 - * Each thread pre-computes the **exclusive location** where it writes its output. Finally, all threads perform their partitioning.
 - Each thread takes a set of R and S partitions. A separate **hash table** is created for each R partition (assuming R is the smaller relation). Each of these hash tables fit into the CPU cache.
 - During the final probe phase, S partitions are scanned and the respective hash table is **probed** for matching tuples.

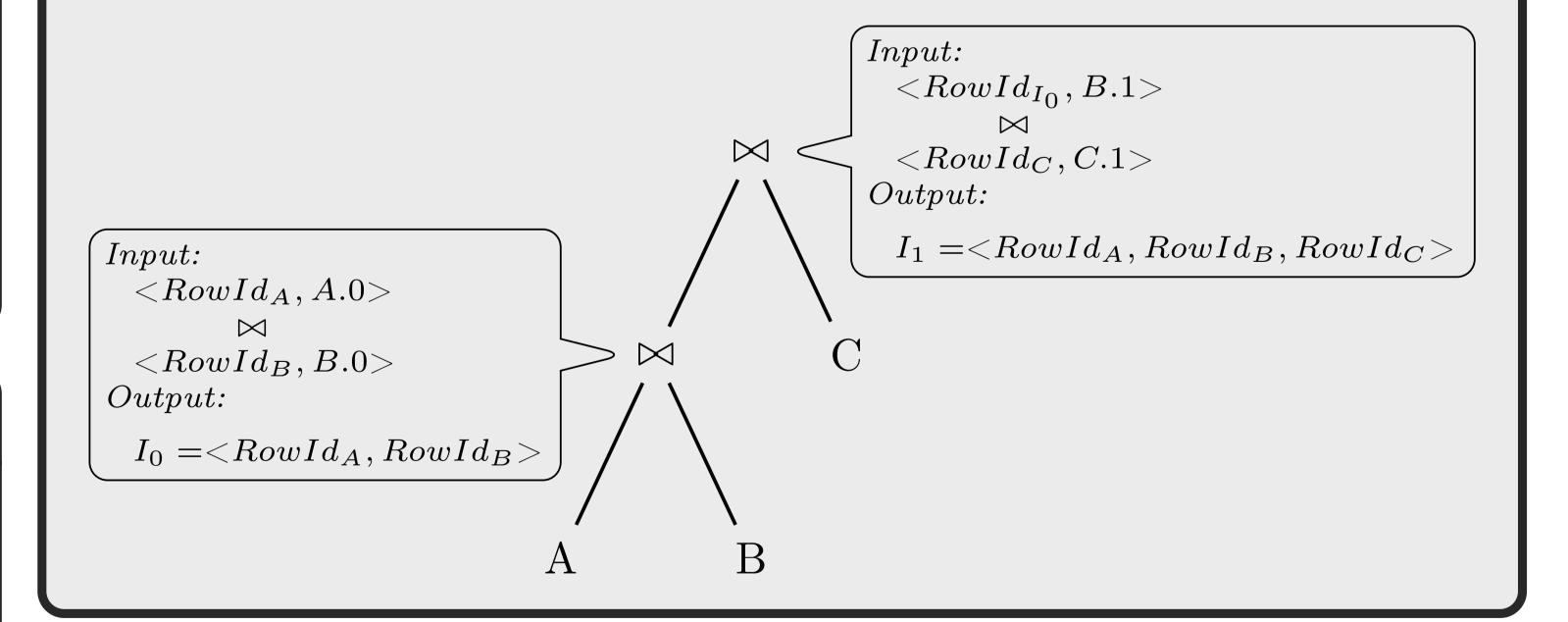
 $\overline{}^a$ source: ETHZ Systems Group's work on Parallel & Distributed Joins at https://www.systems.ethz.ch/node/334

2. Query Optimization

- Assumptions
 - All attribute values are uniformly distributed.
 - The values of all attributes are drawn **independently**.
- Join re-ordering
 - Join tree enumeration using a bottom up dynamic algorithm approach.
 - Consider only **left deep** binary join trees.
 - Cardinality estimation for both sides of the join predicate with statistical formulas taking into consideration foreign key relationships between the two columns.
 - Use as cost function the **sum** of sizes of intermediate results.

4. Plan Execution

- Intermediate results are materialized into RAM to evaluate next-level operations.
- **Filters** are performed first in the tree hierarchy to eliminate as many tuples as possible.
- Each column (suppose from relations A and B) on both sides of the join predicate is represented as a vector of **<RowId**, **Value>** pairs. These vectors are then passed as input to the join operator.
- The join operation results in a vector of pairs $< Row Id_A, Row Id_B>$, where A and B are the joined relations.
- This result is then transformed into a vector of **values** that correspond to the **row ids** of all relations whose join predicates have already been calculated.
- To calculate the final aggregate functions we scan the vector of row ids and probe the initial tables.
- Let's consider the query A.0 = B.0 & B.1 = C.1



6. Caching of indexes

- The Radix-Join algorithm requires a preprocessing phase that divides a relation into partitions (build phase). This phase is the most time consuming in comparison with the other radix hash join phases.
- Every time we apply the build phase to a column of an initial table, we **store** the result for subsequent use.

Conclusion

- Memory's bandwidth seems to be the bottleneck for in memory database systems.
- Occam's Razor: The simplest solution is always the best.