

Parallelizing The Data Cube

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Abstract. This paper presents a general methodology for the *efficient parallelization of existing data cube construction algorithms*. We describe two different partitioning strategies, one for top-down and one for bottom-up cube algorithms. Both partitioning strategies assign subcubes to individual processors in such a way that the loads assigned to the processors are balanced. Our methods reduce inter-processor communication overhead by partitioning the load in advance instead of computing each individual group-by in parallel as is done in previous parallel approaches. In fact, after the initial load distribution phase, each processor can compute its assigned subcube without any communication with the other processors. Our methods enable code reuse by permitting the use of existing sequential (external memory) data cube algorithms for the subcube computations on each processor. This supports the transfer of optimized sequential data cube code to a parallel setting.

The bottom-up partitioning strategy balances the number of single attribute external memory sorts made by each processor. The top-down strategy partitions a weighted tree in which weights reflect algorithm specific cost measures like estimated group-by sizes. Both partitioning approaches can be implemented on any *shared disk* type parallel machine composed of p processors connected via an interconnection fabric and with access to a shared parallel disk array. Experimental results presented show that our partitioning strategies generate a close to optimal load balance between processors.

1 Introduction

Data cube queries represent an important class of On-Line Analytical Processing (OLAP) queries in decision support systems. The precomputation of the different group-bys of a data cube (i.e., the forming of aggregates for every combination of GROUP BY attributes) is critical to improving the response time of the queries [16]. Numerous solutions for generating the data cube have been proposed. One of the main differences between the many solutions is whether they are aimed at sparse or dense relations [4, 17, 20, 21, 27]. Solutions within a

category can also differ considerably. For example, top-down data cube computations for dense relations based on sorting have different characteristics from those based on hashing.

To meet the need for improved performance and to effectively handle the increase in data sizes, parallel solutions for generating the data cube are needed. In this paper we present a general framework for the efficient parallelization of existing data cube construction algorithms. We present load balanced and communication efficient partitioning strategies which generate a subcube computation for every processor. Subcube computations are then carried out using existing sequential, external memory data cube algorithms.

Balancing the load assigned to different processors and minimizing the communication overhead are the core problems in achieving high performance on parallel systems. The heart of this paper are two partitioning strategies, one for top-down and one for bottom-up data cube construction algorithms. Good load balancing approaches generally make use of application specific characteristics. Our partitioning strategies assign loads to processors by using metrics known to be crucial to the performance of data cube algorithms [1, 4, 21]. The bottom-up partitioning strategy balances the number of single attribute external sorts made by each processor [4]. The top-down strategy partitions a weighted tree in which weights reflect algorithm specific cost measures such as estimated group-by sizes [1, 21].

The advantages of our load balancing methods compared to the previously published parallel data cube construction methods [13, 14] are:

- Our methods reduce inter-processor communication overhead by partitioning the load in advance instead of computing each individual group-by in parallel (as proposed in [13, 14]). In fact, after our load distribution phase, each processor can compute its assigned subcube without any inter-processor communication.
- Our methods maximize code reuse from existing sequential data cube implementations by using existing sequential (external memory) data cube algorithms for the subcube computations on each processor. This supports the transfer of optimized sequential data cube code to the parallel setting.

Our partitioning approaches are designed for standard, *shared disk* type, parallel machines: p processors connected via an interconnection fabric where the processors have standard-size local memories and access to a shared disk array. We have implemented our top-down partitioning strategy in MPI and tested it on a multiprocessor cluster. We also tested our bottom-up partitioning strategy through a simulation. Our experimental results indicate that our partitioning strategies generate close to optimal load balancing. Our tests on the multiprocessor cluster showed close to optimal (linear) speedup.

The paper is organized as follows. Section 2 describes the parallel machine model underlying our partitioning approaches as well as the input and the output configuration for our algorithms. Section 3 presents our partitioning approach for parallel bottom-up data cube generation and Section 4 outlines our method for parallel top-down data cube generation. In Section 5 we indicate how our

top-down cube parallelization can be easily modified to obtain an efficient parallelization of the ArrayCube method [27]. Section 6 presents the performance analysis of our partitioning approaches. Section 7 concludes the paper and discusses possible extensions of our methods.

2 Parallel Computing Model

We use the standard *shared disk* parallel machine model. That is, we assume p processors connected via an interconnection fabric where processors have standard size local memories and concurrent access to a shared disk array. For the purpose of parallel algorithm design, we use the *Coarse Grained Multicomputer* (CGM) model [5, 8, 15, 18, 23]. More precisely, we use the *EM-CGM* model [6, 7, 9] which is a multi-processor version of Vitter’s *Parallel Disk Model* [24–26].

For our parallel data cube construction methods we assume that the d -dimensional input data set R of size N is stored on the shared disk array. The output, i.e. the group-bys comprising the data cube, will be written to the shared disk array. For the choice of output file format, it is important to consider the way in which the data cube will be used in subsequent applications. For example, if we assume that a visualization application will require fast access to individual group-bys then we may want to store each group-by in striped format over the entire disk array.

3 Parallel Bottom-Up Data Cube Construction

In many data cube applications, the underlying data set R is sparse; i.e., N is much smaller than the number of possible values in the given d -dimensional space. Bottom-up data cube construction methods aim at computing the data cube for such cases. Bottom-up methods like *BUC* [4] and *PartitionCube* [part of [20]] calculate the group-bys in an order which emphasizes the reuse of previously computed sort orders and in-memory sorts through data locality. If the data has previously been sorted by attribute A then, creating an AB sort order does not require a complete resorting. A local resorting of *A-blocks* (blocks of consecutive elements that have the same attribute A) can be used instead. The sorting of such A-blocks can often be performed in local memory and, hence, instead of another external memory sort, the AB order can be created in one single scan through the disk. Bottom-up methods [4, 20] attempt to break the problem into a sequence of single attribute sorts which share prefixes of attributes and can be performed in local memory with a single disk scan. As outlined in [4, 20], the total computation time of these methods is dominated by the number of such *single attribute sorts*.

In this section we describe a partitioning of the group-by computations into p independent subproblems. Our goal is to balance the number of single attribute sorts required to solve each subproblem and to ensure that each subproblem has overlapping sort sequences in the same way as for the sequential methods (thereby avoiding additional work).

Let A_1, \dots, A_d be the attributes of the data cube such that $|A_1| \geq |A_2| \geq \dots \geq |A_d|$ where $|A_i|$ is the number of different possible values for attribute A_i . As observed in [20], the set of all groups-bys of the data cube can be partitioned into those that contain A_1 and those that do not contain A_1 . In our partitioning approach, the groups-bys containing A_1 will be sorted by A_1 . We indicate this by saying that they contain A_1 as a *prefix*. The group-bys not containing A_1 (i.e., A_1 is projected out) contain A_1 as a *postfix*. We then recurse with the same scheme on the remaining attributes. We shall utilize this property to partition the computation of all group-bys into independent subproblems computing group-bys. The load between subproblems will be balanced and they will have overlapping sort sequences in the same way as for the sequential methods. In the following we give the details of our partitioning method.

Let x, y, z be sequences of attributes representing sort orders and let A be an arbitrary single attribute. We introduce the following definition of sets of attribute sequences representing sort orders (and their respective group-bys):

$$B_1(x, A, z) = \{x, xA\} \quad (1)$$

$$B_i(x, Ay, z) = B_{i-1}(xA, y, z) \cup B_{i-1}(x, y, Az), 2 \leq i \leq \log p + 1 \quad (2)$$

The entire data cube construction corresponds to the set $B_d(\emptyset, A_1 \dots A_d, \emptyset)$ of sort orders and respective group-bys, where d is the dimension of the the data cube. We refer to i as the *rank* of $B_i(\dots)$. The set $B_d(\emptyset, A_1 \dots A_d, \emptyset)$ is the union of two subsets of rank $d - 1$: $B_{d-1}(A_1, A_2 \dots A_d, \emptyset)$ and $B_{d-1}(\emptyset, A_2 \dots A_d, A_1)$. These, in turn, are the union of four subsets of rank $d - 2$. A complete example for a 4-dimensional data cube with attributes A, B, C, D is shown in Figure 1.

$B_4(\emptyset, ABCD, \emptyset)$	$B_3(\emptyset, BCD, A)$	$B_2(\emptyset, CD, BA)$	$B_1(\emptyset, D, CBA) = \{\emptyset, D\}$
		$B_2(B, CD, A)$	$B_1(C, D, BA) = \{C, CD\}$
			$B_1(B, D, CA) = \{B, BD\}$
	$B_3(A, BCD, \emptyset)$	$B_2(A, CD, B)$	$B_1(BC, D, A) = \{BC, BCD\}$
			$B_1(A, D, CB) = \{A, AD\}$
		$B_2(AB, CD, \emptyset)$	$B_1(AC, D, B) = \{AC, ACD\}$
$B_1(AB, D, C) = \{AB, ABD\}$			
			$B_1(ABC, D, \emptyset) = \{ABC, ABCD\}$

Fig. 1. Partitioning For A 4-Dimensional Data Cube With Attributes A, B, C, D.

For the sake of simplifying the discussion, we assume that p is a power of 2. Consider the $2p$ B -sets of rank $d - \log_2(p) - 1$. Let $\beta = (B^1, B^2, \dots, B^{2p})$ be these $2p$ sets in the order defined by Equation (2). Define

$$\begin{aligned} \text{Shuffle}(\beta) &= \langle B^1 \cup B^{2p}, B^2 \cup B^{2p-1}, B^3 \cup B^{2p-2}, \dots, B^p \cup B^{p+1} \rangle \\ &= \langle \Gamma_1, \dots, \Gamma_p \rangle \end{aligned}$$

We assign set $\Gamma_i = B^i \cup B^{2p-i+1}$ to processor P_i , $1 \leq i \leq p$. Observe that from the construction of all group-bys in each Γ_i it follows that every processor performs the same number of single attribute sorts.

Algorithm 1 Parallel Bottom-Up Cube Construction.

Each processor P_i , $1 \leq i \leq p$, performs the following steps, independently and in parallel:

- (1) Calculate Γ_i as described above.
- (2) Compute all group-bys in Γ_i using a sequential (external-memory) bottom-up cube construction method.

— End of Algorithm —

Algorithm 1 can easily be generalized to values of p which are not powers of 2. We also note that Algorithm 1 requires $p \leq 2^{d-1}$. This is usually the case in practice. However, if a parallel algorithm is needed for larger values of p , the partitioning strategy needs to be augmented. Such an augmentation could, for example, be a partitioning strategy based on the number of data items for a particular attribute. This would be applied after partitioning based on the number of attributes has been done. Since the range $p \in \{2^0 \dots 2^{d-1}\}$ covers current needs with respect to machine and dimension sizes, we do not further discuss such augmentations in this paper.

Algorithm 1 exhibits the following properties:

- (a) *The computation of each group-by is assigned to a unique processor.*
- (b) *The calculation of the group-bys in Γ_i , assigned to processor P_i , requires the same number of single attribute sorts for all $1 \leq i \leq p$.*
- (c) *The sorts performed at processor P_i share prefixes of attributes in the same way as in [4, 20] and can be performed with disk scans in the same manner as in [4, 20].*
- (d) *The algorithm requires no inter-processor communication.*

These four properties are the basis of our argument that our partitioning approach is load balanced and communication efficient. In Section 6, we will also present an experimental analysis of the performance of our method.

4 Parallel Top-Down Data Cube Construction

Top-down approaches for computing the data cube, like the sequential *PipeSort*, *Pipe Hash*, and *Overlap* methods [1, 10, 21], use more detailed group-bys to compute less detailed ones that contain a subset of the attributes of the former. They apply to data sets where the number of data items in a group-by can shrink considerably as the number of attributes decreases (data reduction). A group-by is called a child of some parent group-by if the child can be computed from the parent by aggregating some of its attributes. This induces a partial ordering of the group-bys, called the *lattice*. An example of a 4-dimensional lattice is shown in Figure 2, where A, B, C, and D are the four different attributes. The *PipeSort*, *PipeHash*, and *Overlap* methods select a spanning tree T of the lattice, rooted at the group-by containing all attributes. *PipeSort* considers two cases of parent-child relationships. If the ordered attributes of the child are a

prefix of the ordered attributes of the parent (e.g., $ABCD \rightarrow ABC$) then a simple scan is sufficient to create the child from the parent. Otherwise, a sort is required to create the child. *PipeSort* seeks to minimize the total computation cost by computing minimum cost matchings between successive layers of the lattice. *PipeHash* uses hash tables instead of sorting. *Overlap* attempts to reduce sort time by utilizing the fact that overlapping sort orders do not always require a complete new sort. For example, the ABC group-by has A partitions that can be sorted independently on C to produce the AC sort order. This may allow to perform these independent sorts in memory rather than using external memory sort.

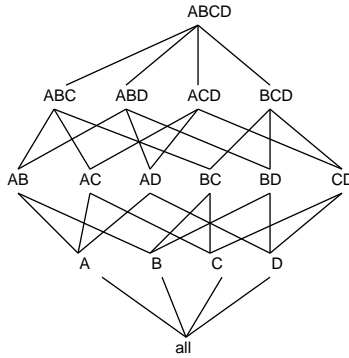


Fig. 2. A 4-Dimensional Lattice.

Next, we outline a partitioning approach which generates p independent subproblems, each of which can be solved by one processor using an existing external-memory top-down cube algorithm. The first step of our algorithm determines a spanning tree T of the lattice by using one of the existing approaches like *PipeSort*, *PipeHash*, and *Overlap*, respectively. To balance the load between the different processors we next perform a storage estimation to determine approximate sizes of the group-bys in T . This can be done, for example, by using methods described in [11] and [22]. We now work with a weighted tree. The most crucial part of our solution is the partitioning of the tree. The partitioning of T into subtrees induces a partitioning of the data cube problem into p subproblems (subsets of group-bys). Determining an optimal partitioning of the weighted tree is easily shown to be an NP-complete problem (by making, for example, a reduction to p processor scheduling). Since the weights of the tree represent estimates, a heuristic approach which generates p subproblems with “some control” over the sizes of the subproblems holds the most promise. While we want the sizes of the p subproblems balanced, we also want to minimize the number of subtrees assigned to a processor. Every subtree may require a scan-

ning of the entire data set R and thus too many subtrees can result in poor IO performance. The solution we develop balances these two considerations.

Our heuristics makes use of a related partitioning problem on trees for which efficient algorithms exist, the *min-max tree k -partitioning problem* [3].

Definition 1. *Min-max tree k -partitioning: Given a tree T with n vertices and a positive weight assigned to each vertex, delete k edges in the tree such that the largest total weight of a resulting subtree is minimized.*

The min-max tree k -partitioning problem has been studied in [3, 12, 19], and an $O(n)$ time algorithm has been presented in [12]. A min-max k -partitioning does not necessarily compute a partitioning of T into subtrees of equal size and it does not address tradeoffs arising from the number of subtrees assigned to a processor. We use tree-partitioning as a preprocessing step for our partitioning. To achieve a better distribution of the load we apply an over partitioning strategy: instead of partitioning the tree T into p subtrees, we partition it into $s \times p$ subtrees, where s is an integer, $s \geq 1$. Then, we use a “packing heuristic” to determine which subtrees belong to which processors, assigning s subtrees to every processor. Our packing heuristic considers the weights of the subtrees and pairs subtrees by weights to control the number of subtrees. It consists of s matching phases in which the p largest subtrees (or groups of subtrees) and the p smallest subtrees (or groups of subtrees) are matched up. Details are described in Step 2b of Algorithm 2.

Algorithm 2 Sequential Tree-partition(T, s, p).

Input: A spanning tree T of the lattice with positive weights assigned to the nodes (representing the cost to build each node from it’s ancestor in T). Integer parameters s (oversampling ratio) and p (number of processors).

Output: A partitioning of T into p subsets $\Sigma_1, \dots, \Sigma_p$ of s subtrees each.

- (1) Compute a min-max tree $s \times p$ -partitioning of T into $s \times p$ subtrees $T_1, \dots, T_{s \times p}$.
- (2) Distribute subtrees $T_1, \dots, T_{s \times p}$ among the p subsets $\Sigma_1, \dots, \Sigma_p$, s subtrees per subset, as follows:
 - (2a) Create $s \times p$ sets of trees named Υ_i , $1 \leq i \leq sp$, where initially $\Upsilon_i = \{T_i\}$. The weight of Υ_i is defined as the total weight of the trees in Υ_i .
 - (2b) For $j = 1$ to $s - 1$
 - Sort the Υ -sets by weight, in increasing order. W.l.o.g., let $\Upsilon_1, \dots, \Upsilon_{sp-(j-1)p}$ be the resulting sequence.
 - Set $\Upsilon_i := \Upsilon_i \cup \Upsilon_{sp-(j-1)p-i+1}$, $1 \leq i \leq p$.
 - Remove $\Upsilon_{sp-(j-1)p-i+1}$, $1 \leq i \leq p$.
 - (2c) Set $\Sigma_i = \Upsilon_i$, $1 \leq i \leq p$.

— End of Algorithm —

The above tree partition algorithm is embedded into our parallel top-down data cube construction algorithm. Our method provides a framework for parallelizing any sequential top-down data cube algorithm. An outline of our approach is given in the following Algorithm 3.

Algorithm 3 Parallel Top-Down Cube Construction.

Each processor P_i , $1 \leq i \leq p$, performs the following steps independently and in parallel:

- (1) Select a sequential top-down cube construction method (e.g., *PipeSort*, *PipeHash*, or *Overlap*) and compute the spanning tree T of the lattice as used by this method.
- (2) Apply the storage estimation method in [22] and [11] to determine the approximate sizes of all group-bys in T . Compute the weight of each node of T ; i.e., the cost to build each node from its ancestor in T .
- (3) Execute Algorithm *Tree-partition*(T, s, p) as shown above, creating p sets $\Sigma_1, \dots, \Sigma_p$. Each set Σ_i contains s subtrees of T .
- (4) Compute all group-bys in subset Σ_i using the sequential top-down cube construction method chosen in Step 1.

— End of Algorithm —

Our performance results described in Section 6 show that an over partitioning with $s = 2$ or 3 achieves very good results with respect to balancing the loads assigned to the processors. This is an important result since a small value of s is crucial for optimizing performance.

5 Parallel Array-Based Data Cube Construction

Our method in Section 4 can be easily modified to obtain an efficient parallelization of the *ArrayCube* method presented in [27]. The *ArrayCube* method is aimed at dense data cubes and structures the raw data set in a d -dimensional array stored on disk as a sequence of “*chunks*”. Chunking is a way to divide the d -dimensional array into small size d -dimensional chunks where each chunk is a portion containing a data set that fits into a disk block. When a fixed sequence of such chunks is stored on disk, the calculation of each group-by requires a certain amount of buffer space [27]. The *ArrayCube* method calculates a minimum memory spanning tree of group-bys, *MMST*, which is a spanning tree of the lattice such that the total amount of buffer space required is minimized. The total number of disk scans required for the computation of all group-bys is the total amount of buffer space required divided by the memory space available. The *ArrayCube* method can now be parallelized by simply applying Algorithm 3 with T being the *MMST*. More details will be given in the full version of this paper.

6 Experimental Performance Analysis

We have implemented and tested our parallel *top-down* data cube construction method presented in Section 4. We implemented sequential pipesort [1] in C++, and our parallel top-down data cube construction method (Section 4) in C++ with MPI [2]. As parallel hardware platform, we use a 9-node cluster. One node is used as the *root node*, to partition the lattice and distribute the work among

the other 8 machines which we refer to as *compute nodes*. The root is an IBM Netfinity server with two 9-G scsi disks, 512 MB of Ram and a 550-MHZ Pentium processor. The compute nodes are 133 MHZ Pentium processors, with 2G IDE hard drives and 32 MB of RAM. The processors run LINUX and are connected via a 100 Mbit Fast Ethernet switch with full wire speed on all ports.

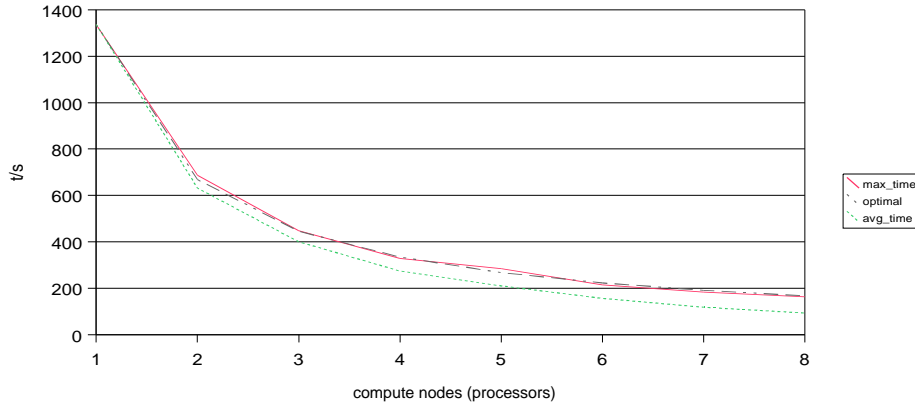


Fig. 3. Running Time (in seconds) As A Function Of The Number of Compute Nodes (Processors)

Figure 3 shows the running time observed (in seconds) as a function of the number of compute nodes used. For the same data set, we measured the sequential time (sequential pipesort [1]) and the parallel time obtained through our parallel top-down data cube construction method (Section 4), using an oversampling ratio of $s = 2$. The data set consisted of 100,000 records with dimension 6. The attribute cardinalities for dimensions 1 to 6 were 5, 15, 500, 20, 1000, and 2, respectively. Our test data values were sparse and uniformly distributed. Figure 3 shows the running times (in seconds) of the algorithm as we increase the number of compute nodes. There are three curves shown. *Max-time* is the time taken by the slowest compute node (i.e. the node that received the largest workload). *Avg-time* is the average time taken by the compute nodes. The time taken by the *root node*, to partition the lattice and distribute the work among the compute nodes, was insignificant. The *optimal* time shown in Figure 3 is the sequential pipesort time divided by the number of compute nodes (processors) used.

We observe that the *max-time* and *optimal* curves are essentially identical. That is, for an oversampling ratio of $s = 2$, the speedup observed is very close to optimal.

Note that, the difference between *max-time* and *avg-time* represents the load imbalance created by our partitioning method. As expected, the difference grows with increasing number of processors. However, we observed that a good part

of this growth can be attributed to the estimation of the cube sizes used in the tree partitioning. We are currently experimenting with improved estimators which appear to improve the result. Interestingly, the *avg-time* curve is below the *optimal* curve, while the *max-time* and *optimal* curves are essentially identical. One would have expected that the *optimal* and *avg-time* curves are similar and that the *max-time* curve is slightly above. We believe that this is caused by another effect which benefits our parallel method: improved I/O. When sequential pipesort is applied to a 10 dimensional data set, the lattice is partitioned into pipes of length up to 10. In order to process a pipe of length 10, pipesort needs to write to 10 open files at the same time. It appears that the number of open files can have a considerable impact on performance. For 100,000 records, writing them to 4 files took 8 seconds on our system. Writing them to 6 files took 23 seconds, not 12, and writing them to 8 files took 48 seconds, not 16. This benefits our parallel method, since we partition the lattice first and then apply pipesort to each part. Therefore, the pipes generated in the parallel method are considerably shorter.

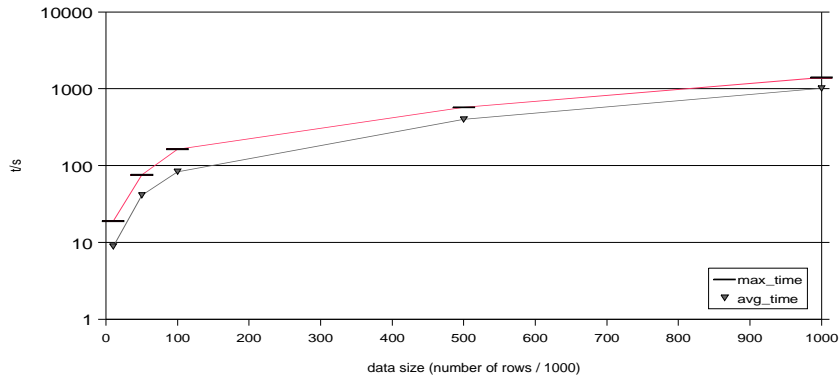


Fig. 4. Running Time (in seconds) As A Function Of The Size Of The Data Set (number of rows / 1000)

Figure 4 shows the running times (in seconds) of our top-down data cube parallelization as we increase the data size from 100,000 to 1,000,000 rows. Note that, the scale is logarithmic. The main observation is that the parallel running time (*max-time*) increases essentially linear with respect to the data size.

Figure 5 shows the running times as a function of the oversampling ratio s . We observe that the parallel running time (i.e., *max-time*) is best for $s = 3$. This is due to the following tradeoff. Clearly, the workload balance improves as s increases. However, as the total number of subtrees, $s \times p$, generated in the tree partitioning algorithm increases, we need to perform more sorts for the root nodes of these subtrees. The optimal tradeoff point for our test case is $s = 3$.

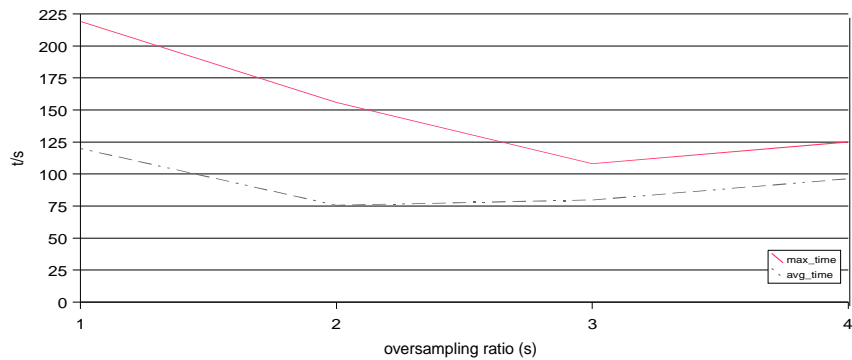


Fig. 5. Running Time (in seconds) As A Function Of The Oversampling Ratio (s)

Figure 6 shows the running times (in seconds) of our top-down data cube parallelization as we increase the dimension of the data set from 2 to 10. Note that, the number of group-bys to be computed grows exponentially with respect to the dimension of the data set. In Figure 6, we observe that the parallel running time grows essentially linear with respect to the output. We also executed our parallel algorithm for a 15-dimensional data set of 10,000 rows, and the resulting data cube was of size more than 1G.

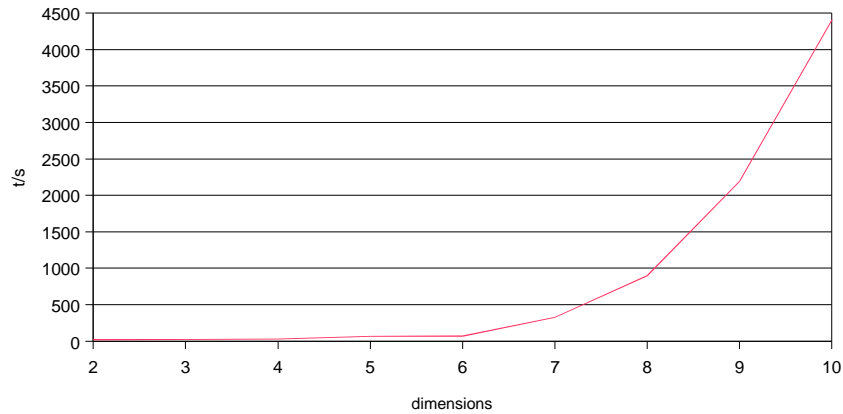


Fig. 6. Running Time (in seconds) As A Function Of The Number Of Dimensions Of The Data Set.

Simulation results for our *bottom-up* data cube parallelization in Section 3 are shown in Figure 7. For this method we have so far measured its load balancing characteristics through simulation only. As indicated in [4, 20], the main indicator for the load generated by a bottom-up data cube computation is the number

of single attribute sorts. Our partitioning method in Section 3 for bottom-up data cube parallelization does in fact *guarantee* that the subcube computations assigned to the individual processor do all require *exactly* the same number of single attribute sorts. There are no heuristics (like oversampling) involved. Therefore, what we have measured in our simulation is whether the *output sizes* of the subcube computations assigned to the processors are balanced as well. The results are shown in Figure 7. The x-axis represents the number of processors $p \in \{2, \dots, 64\}$ and the y-axis represents the largest output size as a percentage of the total data cube size. The two curves shown are the largest output size measured for a processor and the optimal value (total data cube size / number of processors). Five experiments were used to generate each data point. We observe that the actual values are very close to the optimal values. The main result is that our partitioning method in Section 3 not only balances the number of single attribute sorts but also the sizes of the subcubes generated on each processor.

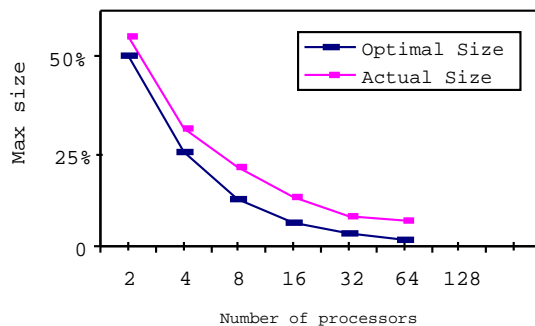


Fig. 7. Bottom-Up Cube. Maximum Output Size For One Processor As Percentage Of Total Data Cube Size.

7 Conclusion

We presented two different, partitioning based, data cube parallelizations for standard shared disk type parallel machines. Our partitioning strategies for bottom-up and top-down data cube parallelization balance the loads assigned to the individual processors, where the loads are measured as defined by the original proponents of the respective sequential methods. Subcube computations are carried out using existing sequential data cube algorithms. Our top-down partitioning strategy can also be easily extended to parallelize the ArrayCube method.

Experimental results indicate that our partitioning methods produce well balanced data cube parallelizations. Compared to existing parallel data cube methods, our parallelization approach brings a significant reduction in inter-processor communication and has the important practical benefit of enabling the re-use of existing sequential data cube code.

A possible *extension* of our data cube parallelization methods is to consider a *shared nothing* parallel machine model. If it is possible to store a duplicate of the input data set R on each processor’s disk, then our method can be easily adapted for such an architecture. This is clearly not always possible. It does solve most of those cases where the total output size is considerably larger than the input data set; for example *sparse* data cube computations. In fact, we applied this strategy for our implementation presented in Section 6. As reported in [20], the data cube can be several hundred times as large as R . Sufficient total disk space is necessary to store the output (as one single copy distributed over the different disks) and a p times duplication of R may be smaller than the output. Our data cube parallelization method would then partition the problem in the same way as described in Sections 3 and 4, and subcube computations would be assigned to processors in the same way as well. When computing its subcube, each processor would read R from its local disk. For the output, there are two alternatives. Since the output data sizes are well balanced, each processor could simply write the subcubes generated to its local disk. This could, however, create a bottleneck if there is, for example, a visualization application following the data cube construction which needs to read a single group-by. In such a case, each group-by should be distributed over all disks, for example in striped format. To obtain such a data distribution, all processors would not write their subcubes directly to their local disks but buffer their output. Whenever the buffers are full, they would be permuted over the network. In summary we observe that, while our approach is aimed at shared disk parallel machines, its applicability to shared nothing parallel machines depends mainly on the distribution and availability of the input data set R . We are currently considering the problem of identifying the “ideal” distribution of input R among the p processors when a fixed amount of replication of the input data is allowed (i.e., R can be copied r times, $1 \leq r < p$).

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