

Efficient Approximate Top- k Query Algorithm Using Cube Index

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Abstract. Exact top- k query processing has attracted much attention recently because of its wide use in many research areas. Since missing the truly best answers is inherent and unavoidable due to the user's subjective judgment, and the cost of processing exact top- k queries is highly expensive for datasets with huge volume, it is intriguing to answer approximate top- k query instead. In this paper, we first define a novel kind of approximate top- k query, called μ -approximate top- k query. Then we introduce an efficient index structure, i.e. *cube index*, based on which, we propose our novel Cube Index Algorithm (CIA). We analyze the complexity of both constructing cube index and CIA algorithm. Moreover, extensive experiments show that CIA performs much better than the well-known approximate TA_θ algorithm [3].

Keywords: Top- k Query Processing, Algorithms, Index.

1 Introduction

Exact top- k query processing has gained more and more attention recently because of its wide use in many fields, such as information retrieval [16][17], multimedia databases [20][21], P2P and sensor networks [18][19], etc. The main reason for such attention is that top- k queries avoid overwhelming the user with a large number of uninteresting answers that are resource-consuming.

However, two main reasons convince us to abandon exact top- k query processing. First, the top- k query concept is heuristic anyway. Hardly any user is interested in all the exact k answers of a top- k query. Instead, they may be only interested in one or several relevant objects in the top- k (e.g. 500 or 2000) answers. So, due to the subjective judgment of the user, missing the truly best answers is inherent and unavoidable. This argument enlightens us to relax exact top- k query to approximate top- k query. Second, the cost of processing exact top- k queries is highly expensive for datasets with huge volume, and the size of datasets in practice is always quite huge. So it's intriguing to answer approximate top- k query instead of exact top- k query.

To answer approximate top- k queries, Fagin et al. propose the TA_θ algorithm in [3], which is based on the TA algorithm. Based on TA_θ , Theobald et al. [6] introduced a scheme to associate probabilistic guarantees with approximate top- k answers. In [8], Amato used a *proximity* measure to decide if a data region should be inspected or not. Only data regions whose *proximity* to the query region is greater than a specified

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threshold are accessed. This method is used to rank the nearest neighbors to some target data object in an approximate manner. Approximate top- k query processing has been also studied in peer-to-peer environments. The KLEE system (Michel et al. [2]) addressed this problem, where distributed aggregation queries are processed based on index lists located at isolated sites. KLEE assumes no random accesses are made to index lists located at each peer.

In this paper, we first define a novel approximate top- k query, called μ -approximate top- k query. Then we introduce an efficient index structure, i.e. *cube index*, based on which, we propose our new Cube Index Algorithm (CIA). We analyze the complexity of both constructing cube index and CIA algorithm. Moreover, extensive experiments show that CIA performs much better than the well-known approximate TA_θ algorithm

The rest of this paper is organized as follows: First, we define the computation model formally and review the TA_θ algorithm in Section 2. In Section 3, we describe our method on setting up the cube index and then analyze its time complexity. Based on these, we show our algorithm CIA and analyze its cost in Section 4. Thereafter, we show the experimental results in Section 5. Finally, in Section 6, we conclude this paper and introduce our future work.

2 Computation Model and TA_θ Algorithm

In this section, we describe the model of top- k problem and review the TA_θ algorithm [3].

Our model of the dataset can be described as follows: assume the database D consists of n objects, which are denoted as $x_1, x_2 \dots x_n$. Each object x is an m -dimensional vector $(s_1(x), s_2(x) \dots s_m(x))$, where $s_i(x)$ is the i th local score of x as a real number in the interval $[0, 1]$. For a given object x , x has a total score of $f(x) = f(s_1(x), s_2(x) \dots s_m(x))$, where the m -dimensional aggregate function f is supposed to be increasingly monotonic:

Definition 2.1 Monotonic Function. An aggregate function f is *increasingly monotonic* if $f(a_1, a_2 \dots a_m) \leq f(a_1', a_2' \dots a_m')$, whenever $a_i \leq a_i'$ for every i .

In this paper, we assume the aggregate function is weighted summation function,

$f(x) = \sum_{i=1}^m w_i s_i(x)$, where $s_i(x) \in [0, 1]$ and $\sum_{i=1}^m w_i = 1$ ($w_i \neq 0$). We can easily verify that

weighted summation function is increasingly monotonic. Exact top- k query is to find k objects with the highest total scores. For approximate top- k query, Fagin et al. [3] defined a θ -approximation to the top- k answers:

Definition 2.2 θ -Approximation [3]. Let Y be a collection of k objects such that for each y among Y and each z not among Y , there are $\theta f(y) \geq f(z)$, where $\theta > 1$. Then Y is one of the top- k answers with θ -approximation and θ is the *relative approximation coefficient*.

To solve the θ -approximation top- k query, Fagin et al. [3] proposed the TA_θ algorithm, based on the threshold algorithm (i.e. TA). TA_θ is described in Fig. 1.

3 Cube Index

Before proposing our algorithm, we first introduce an efficient indexing structure

Threshold Algorithm with θ -Approximation (TA_θ)

Pre-computing Phase:

For each attribute $i \in \{1, 2 \dots m\}$, get every $s_i(x_j)$ where $j \in \{1, 2 \dots n\}$ and insert them into a sorted list L_i . Sorted list means that objects in each list are sorted in descending order by the $s_i(x_j)$ value.

Computing Phase:

- 1: Do sorted access in parallel to each of the m lists. As an object is seen through sorted access in some list, do random access to the other lists to find all its remaining local scores, and compute its overall score. Maintain a set Y containing the k objects whose overall scores are the highest among all the objects seen so far.
 - 2: For each list L_i , let \underline{s}_i be the last local score seen under sorted access in L_i . Define the *threshold value* τ to be $\tau = f(\underline{s}_1, \underline{s}_2 \dots \underline{s}_m)$.
 - 3: Halt when $\theta M_k \geq \tau$, where $M_k = \min\{f(x) \mid x \in Y\}$.
-

Fig. 1. Threshold Algorithm with θ -Approximation

called cube index to support such μ -approximation top- k query processing.

3.1 Description of Cube Index

We map the database to an m -dimensional hyperspace $[0, 1]^m$; each object x_j with scores $(s_1(x_j), s_2(x_j) \dots s_m(x_j))$ in the database is mapped to an m -dimensional point $p_j = (s_1(x_j), s_2(x_j) \dots s_m(x_j))$ in $[0, 1]^m$. We will not distinguish between object x and its corresponding point p discussed below. Similarly, $s_i(p)$ is the value of p 's i th dimension and $f(p)$ is p 's total score.

Now we define a μ -approximation to the top- k answers.

Definition 3.1 μ -Approximation. Let Y be a collection of k objects such that for each y among Y and each z not among Y , there are $f(y) + \mu \geq f(z)$, where $0 < \mu \leq 1$. Then Y is one of the top- k answers with μ -approximation and μ is the *proportional approximation coefficient*.

Definition 3.2 Dominate [7]. Point p_1 dominates point p_2 if and only if for each $i \in \{1, 2 \dots m\}$, $s_i(p_1) \geq s_i(p_2)$ and there exists at least one member j of $\{1, 2 \dots m\}$ satisfying $s_j(p_1) > s_j(p_2)$.

Observation 3.1. If point p_1 dominates point p_2 , then $f(p_1) > f(p_2)$, where f is an *aggregate monotone function*.

Proof. We can easily get the correctness of Observation 3.1 according to the definitions of *aggregate monotone function* and *dominate*. \square

Definition 3.3 Skyline [7]. The *skyline* of a dataset D is the set of points that are not dominated by any point in D .

Definition 3.4 Bottom Point. The *bottom point* of a hypercube is the vertex whose values of every dimension are all lowest in the hypercube.

For example, the *bottom point* of the 3-dimensional cube $[0.2, 0.3] \times [0.1, 0.2] \times [0.5, 0.6]$ is $(0.2, 0.1, 0.5)$.

Observation 3.2. All other points in a hypercube dominate the *bottom point*.

Proof. We can easily get the correctness of Observation 3.2 according to the definitions of *dominate* and *bottom point*. \square

Now we show the cube partition method on the m -dimensional hyperspace $[0, 1]^m$,

which is described as follows:

Firstly, we set the length of the edge of each hypercube as μ , where $\mu \in [0, 1]$. Then we divide the interval $[0, 1]$ into several μ -segments from 1 to 0 until the rest is shorter than μ . Each dimension is divided in this way so that the m -dimensional hyperspace $[0, 1]^m$ is partitioned into several hypercubes or sub-hyperspaces. Thereafter, we classify all the points in database into several sets: Point p_i belongs to bp_i 's associated point set S_i if and only if p_i is in the hypercube whose *bottom point* is bp_i .

We call this partition method the μ -cube partition.

Definition 3.5 Sky Point. For a μ -cube partition, the *sky point* is the point whose values in every dimension are all $1 - \mu$, that is, the point $(1 - \mu, 1 - \mu, \dots, 1 - \mu)$.

Apparently, *sky point* is the very *bottom point* which *dominates* all the other *bottom points* and the set $\{\text{sky point}\}$ is the *skyline* of the set of *bottom points*.

Definition 3.6 Neighbor. *Bottom point* bp_1 is a *neighbor* of *bottom point* bp_2 if and only if they satisfy $\sum_{i=1}^m \left\lfloor \frac{|s_i(bp_1) - s_i(bp_2)|}{\mu} \right\rfloor = 1$.

Definition 3.7 Superior. *Bottom point* bp_1 is a *superior* of *bottom point* bp_2 if and only if bp_1 is a *neighbor* of bp_2 and bp_1 *dominates* bp_2 .

Definition 3.8 Inferior. *Bottom point* bp_1 is an *inferior* of *bottom point* bp_2 if and only if bp_1 is a *neighbor* of bp_2 and bp_1 is *dominated* by bp_2 .

Discussions on special cases:

- 1) For the points in the hypercube whose *bottom point* is the *sky point* belong to the 0th set S_0 .
- 2) The points on the intersecting hyperplane of several neighboring hypercubes belong to the hypercube whose *bottom point* *dominates* the others' *bottom point*.
- 3) The points coinciding with bp_i belong to set S_i .
- 4) If $S_i.size = 0$ and $i \neq 0$, then remove bp_i from the set of *bottom points*. Meanwhile, for each *inferior* inf of bp_i , regard all the *superiors* of bp_i as inf 's *superiors* too; for each *superior* sup of bp_i , regard all the *inferiors* of bp_i as sup 's *inferiors* too.

Definition 3.9 μ -Cube Index. For a μ -cube partition, the μ -cube index is an index list or array whose entries are ids of the *bottom points*. Each *bottom point* bp_i has its associated point set S_i as well as its *superiors*' ids and *inferiors*' ids.

3.2 Complexity Analysis of μ -Cube Indexing Method

Now we analyze the time complexity of the method on setting up the cube index, which is done in the pre-computing phase.

According to the description, the most time-consuming calculations in a μ -cube partition are to find the *superiors* and *inferiors* of each *bottom point* and to classify all the points in database into their corresponding sets.

Actually, the *superiors* and *inferiors* of each *bottom point* bp can be determined by the following two simple formulas:

1. For each $i \in \{1, 2 \dots m\}$ and $s_i(bp) \neq 0$, *bottom point* bp' is one *inferior* of bp , satisfying

- a. $s_i(bp') = (s_i(bp) - \mu) \cdot H(s_i(bp) - \mu)$, where $H(x)$ is the Heaviside step function;
 - b. $s_j(bp') = s_j(bp)$ for each $j \in \{1, 2 \dots m\}$ and $j \neq i$.
2. *Bottom point* bp' is one *inferior* of bp if and only if bp is one *superior* of bp' .

There are $\left\lceil \frac{1}{\mu} \right\rceil^m$ *bottom points* in total, so the time complexity to find the *superiors*

and *inferiors* of each *bottom point* is $O\left(\left\lceil \frac{1}{\mu} \right\rceil^m \times m\right)$.

On the other hand, each point p in database belongs to set S_i if and only if set S_i 's corresponding *bottom point* bp_i satisfies that for each $i \in \{1, 2 \dots m\}$,

- a. $s_i(bp) = (1 - \left\lceil \frac{1-s_i(p)}{\mu} \right\rceil \times \mu) \times H(1 - \left\lceil \frac{1-s_i(p)}{\mu} \right\rceil \times \mu)$ if $s_i(p) \neq 1$, where $H(x)$ is the Heaviside step function;
- b. $s_i(bp) = 1 - \mu$ when $s_i(p) = 1$.

Similarly, there are n points in database, so the time complexity to classify all the points in database into their corresponding sets is $O(mn)$.

Therefore, the total time complexity in the pre-computing phase is

$$O\left(m \left\lceil \frac{1}{\mu} \right\rceil^m + mn\right).$$

4 The Cube Index Algorithm

4.1 Description of Cube Index Algorithm

Based on the μ -cube index, we now propose a novel algorithm to answer the μ -approximation top- k query: the Cube Index Algorithm (i.e. CIA), which is described by the pseudo-code in Fig. 2.

Here *Selectively Add* in the pseudo-code is a sub- method to improve the precision of the algorithm qualitatively. It can be to add the points at random, or to add them from the points in *skyline* of S_i or others ways.

4.2 μ -Approximation of Cube Index Algorithm

To proof the μ -approximation of CIA, we first introduce three lemmas and a corollary as follows.

Lemma 4.1. Set T is always the top- $(T.size)$ answers to the set of *bottom points*.

Proof. (By mathematical induction) *Basis:* Set $T = \{sky\ point\}$ is the top-1 answers to the set of *bottom points*. Actually, *sky point* dominates all the other *bottom points* for the formula of μ -cube index and the definition of *sky point*. According to Observation 3.1, the *sky point* is the top-1 in the set of *bottom points*.

Inductive step: Assume that set T is the top- j answers to the set of *bottom points* now, then the *bottom point* bp_i with the highest score in CL is the top- $(j + 1)$ in the set of *bottom points* and is supposed to be moved to set T from CL .

Cube Index Algorithm (CIA)

Pre-computing Phase:

Execute the normalization then set up the μ -cube index on the database.

Computing Phase:

- 1: $Y = \emptyset$, $CL = \emptyset$, $T = \{\text{sky point}\}$, where Y is the result set while CL is the sorted candidate list according to the total scores and T is a temp set.
 - 2: **if** $S_0.size \leq k$ **then**
 - 3: add all points in S_0 into Y
 - 4: **else**
 - 5: *Selectively Add* k points in S_0 into Y .
 - 6: $bp_i = \text{sky point}$.
 - 7: **while** ($Y.size < k$) **do**
 - 8: **for each** *inferior* inf of bp_i **do**
 - 9: **if** inf has not been accessed before and all *superiors* of inf is among T **then**
 - 10: Access inf and insert it into CL
 - 11: **else**
 - 12: Continue.
 - 13: **if** $CL.size > k - Y.size$ **then**
 - 14: Only keep the first $k - Y.size$ points in CL .
 - 15: Let bp_i be the *bottom point* with the highest score in CL and move it into T .
 - 16: **if** $S_i.size \leq k - Y.size$ **then**
 - 17: add all points in S_i into Y
 - 18: **else**
 - 19: *Selectively Add* $k - Y.size$ points in S_i into Y .
 - 20: **Return** Y .
-

Fig. 2. Cube Index Algorithm

Actually, only the points in the CL now have the chance to be the top- $(j + 1)$. Otherwise, for a point bp which is not in CL or set T , either bp has been accessed before or bp has at least one *superior* that is not in set T . In the first case, according to the algorithm, CIA halts if and only if $Y.size = k$, so $Y.size < k$ before the algorithm halts. If bp has been accessed before and be removed from CL , then there exist at least $T.size + (k - Y.size) \geq T.size + 1 = j + 1$ points whose total scores are higher than bp so that bp even has no chance to be one of the top- $(j + 1)$ answers. In the other case, according to the definition of *superior* and Observation 3.1, every *superior* sup of bp satisfies $f(sup) > f(bp)$, so once sup is not in the top- j answers, or set T , bp has no chance to be one of the top- $(j + 1)$ answers. Furthermore, for each point bp in CL , where $bp \neq bp_i$, bp is impossible to be one of the top- $(j + 1)$ answers because even bp_i is not in the top- j answers. Therefore, bp_i is the top- $(j + 1)$ in the set of *bottom points*.

Conclusion: When CIA halts, set T is the top- $(T.size)$ answers to the set of *bottom points*. \square

Corollary 4.1. *Bottom points* are moved into set T in descending order of total score.

Proof. From the proof of Lemma 4.1, we easily conclude that *bottom points* are moved into set T in descending order of total score. \square

Lemma 4.2. When CIA halts, there is at most one *bottom point* bp_j in set T satisfying $S_j \not\subseteq Y$, where bp_j is the one with the lowest score in set T and for each $bp_i \in$

T and $bp_i \neq bp_j$, $S_i \subseteq Y$.

Proof. According to the algorithm, the sub-method *Selectively Add* is executed if and only if $S_j.size > k - Y.size$. In this case, we *Selectively Add* $k - Y.size$ points in S_j into Y so that $S_j \not\subseteq Y$. Thus there would be $Y.size = k$ once the *Selectively Add* has been executed, where CIA halts. So the sub-method *Selectively Add* can be executed at most once. For Corollary 4.1, bp_j is the one with the lowest score in set T . However, in the case that $S_i.size \leq k - Y.size$, we add the whole S_i into set Y so that $S_i \subseteq Y$.

Therefore, when CIA halts, there is at most one *bottom point* bp_j in set T satisfying $S_j \not\subseteq Y$, where bp_j is the one with the lowest score in set T and for each $bp_i \in T$ and $bp_i \neq bp_j$, $S_i \subseteq Y$. \square

Lemma 4.3. For point $p_i \in S_i$ and point $p_j \in S_j$, if $f(bp_i) \geq f(bp_j)$, then $f(p_i) + \mu \geq f(p_j)$.

Proof. According to the formula of μ -cube index and the definition of *bottom point*, for each $l \in \{1, 2 \dots m\}$, there is $s_l(bp_j) \leq s_l(p_j) \leq s_l(bp_j) + \mu$. Considering

$$f(x) = \sum_{l=1}^m w_l s_l(x), \text{ where } s_l(x) \in [0, 1] \text{ and } \sum_{l=1}^m w_l = 1, \text{ we have}$$

$$f(bp_j) \leq f(p_j) \leq \sum_{l=1}^m w_l [s_l(p_j) + \mu] = \sum_{l=1}^m w_l s_l(p_j) + \sum_{l=1}^m w_l \mu = f(p_j) + \mu$$

for Observation 3.1 and Observation 3.2. We can also get $f(bp_i) \leq f(p_i)$ in the same way.

Therefore, $f(p_i) + \mu \geq f(bp_i) + \mu \geq f(bp_j) + \mu \geq f(p_j)$. \square

Theorem 4.1. CIA based on μ -cube index finds the top- k answers with μ -approximation.

Proof. According to the algorithm, if $bp_i \notin T$, any member of S_i has no chance to be added into set Y . That is, for each $y \in Y$ and $y \in S_y$, there must be $bp_y \in T$. And from Lemma 4.1, we know that set T is the top- $(T.size)$ answers to the set of *bottom points*. For each point $z \notin Y$ and $z \in S_z$ and for each $y \in Y$ and $y \in S_y$, if $bp_z \notin T$, then $f(bp_y) \geq f(bp_z)$, so $f(y) + \mu \geq f(z)$ for Lemma 4.3. In the other case, if $bp_z \in T$, since $z \notin Y$, meaning $S_z \not\subseteq Y$, bp_z is the one with the lowest score in set T according to Lemma 4.2. So we also have $f(bp_y) \geq f(bp_z)$ and $f(y) + \mu \geq f(z)$.

Therefore, for each y among Y and each z not among Y , there is $f(y) + \mu \geq f(z)$. That is, CIA based on μ -cube index finds the top- k answers with μ -approximation. \square

4.3 Cost Analysis of Cube Index Algorithm

According to Fagin et al. [3], the cost of the top- k query is proportional to the times of accessing or aggregating the objects. For the CIA, the cost is the number of *bottom points* accessed in the query.

First, let \underline{bp} be the last *bottom point* added into set T . Denote $B_1 = \{\text{sky point}\} + \{bp \mid bp \text{ is a bottom point which is accessed in the query}\}$ and $B_2 = T - \{\underline{bp}\}$. According to Lemma 4.1 and Corollary 4.1, B_2 is the top- $(T.size - 1)$ answers to the set of *bottom points*.

Theorem 4.2. The cost of the CIA is $T.size - 2 + \text{skyline}(\bar{B}_2).size$, where \bar{B}_2 is the complementary set of B_2 .

Proof. We only need to show that $B_1 = B_2 + \text{skyline}(\bar{B}_2)$. Actually, it can be proved by apagoge.

Case 1: If there exists $bp \in B_2 + \text{skyline}(\bar{B}_2)$ but $bp \notin B_1$, then we know bp is not

the *sky point*.

sub-case 1: If $bp \in B_2$, since $B_2 = T - \{bp\} \Rightarrow B_2 \subset T$, according to the algorithm, bp has no chance to be added into set T if bp has not been accessed. So it will conflict with the algorithm.

sub-case 2: If $bp \in \text{skyline}(\bar{B}_2)$, then all the *superiors* of bp is in B_2 because there is no any point in \bar{B}_2 dominating bp according to the definition of *skyline*. However, in CIA, all points whose all *superiors* are in T must be accessed before the CIA halts. As $B_2 \subset T$, bp must be accessed, which contradicts the assumption that $bp \notin B_1$.

Case 2: If there exists $bp \in B_1$ but $bp \notin B_2 + \text{skyline}(\bar{B}_2)$, then bp belongs to neither B_2 nor $\text{skyline}(\bar{B}_2)$. The fact that $bp \notin B_2$ indicates bp is not in the top- $(T.size - 1)$ answers to the set of *bottom points* so bp is not the *sky point* because $\{\text{sky point}\}$ is the top-1 answers. Therefore, bp has chance to be accessed if and only if all the *superiors* of bp are in B_2 for the algorithm. However, $bp \notin \text{skyline}(\bar{B}_2)$, meaning that bp has at least one *superior* that is not in T so bp cannot be accessed. Thus the assumption has no chance to be true.

Therefore, $B_1 = B_2 + \text{skyline}(\bar{B}_2)$. Besides, since $\text{skyline}(\bar{B}_2) \subseteq \bar{B}_2$, $B_2 \cap \text{skyline}(\bar{B}_2) = \emptyset$. So $B_1.size = B_2.size + \text{skyline}(\bar{B}_2).size$. Moreover, $B_2.size = T.size - 1$ and the cost of the CIA is $B_1.size - 1$, considering that the *sky point* is not accessed in the algorithm.

Therefore, the cost of the CIA is $B_1.size - 1 = T.size - 1 + \text{skyline}(\bar{B}_2).size - 1 = T.size - 2 + \text{skyline}(\bar{B}_2).size$. \square

5 Experiments

In this section, we conduct extensive experiments to evaluate the performance of our algorithm. Our algorithm is implemented in C/C++ language. We perform our experiments on an 8-CPU server with 8GB shared memory and each CPU is 4-core Intel Xeon E5430 2.66GHz.

5.1 Turning μ -Approximation into θ -Approximation

According to the definitions of μ -approximation and θ -approximation, if set Y is the top- k answers with μ -approximation, for each y among Y and each z not among Y , there are $f(y) + \mu \geq f(z)$. So we have $(1 + \frac{\mu}{f(y)})f(y) \geq f(z)$. Let $\underline{f}(y)$ be the k th highest

total score in set Y so that $\frac{\mu}{f(y)} \leq \frac{\mu}{\underline{f}(y)}$. Therefore, the *relative approximation*

coefficient $\theta = \frac{\mu}{\underline{f}(y)}$, or $\mu = \underline{f}(y) \cdot \theta$.

In our experiments, we run the CIA over the databases to find the value of $\underline{f}(y)$ and then the TA_θ runs on θ -approximation of $\theta = \frac{\mu}{\underline{f}(y)}$. We choose the μ -approximation as the criterion of approximation to run our tests.

5.2 Evaluation Metrics

In our tests, the following measures are collected for efficiency comparison [6]:

accesses: the number of items accessed in the query without duplication;

precision: the fraction of top- k results in an approximate result that belongs to the exact top- k result;

recall: the fraction of top- k results in the exact result that were returned by the approximate top- k query;

rank distance: the *footrule distance* [14] between the ranks of the approximate top- k results and their true ranks in the exact top- k result, i.e., $\frac{1}{k} \sum_{i=1}^k |i - \text{truerank}(i)|$,

where $\text{truerank}(i)$ is the i th returned object's true rank in the exact top- k result.

score error: the absolute error between approximate and exact top- k scores, i.e.,

$$\frac{1}{k} \sum_{i=1}^k |\text{totalscore}_i^{(\text{approx})} - \text{totalscore}_i^{(\text{exact})}|,$$

where $\text{score}_i^{(\text{approx})}$ is the total score of the i th object in the approximate top- k result while $\text{score}_i^{(\text{exact})}$ is the total score of the i th object in the exact top- k result.

Because the *precision* and the *recall* have the same denominator k , they have identical values in our setup. We regard the *recall* as a formal measure in our tests, instead of *precision*.

5.3 Description of Datasets

We do experiments on two synthetic datasets. All generated local scores belong to the interval $[0, 1]$. The two synthetic datasets are produced to model different input scenarios. They are UI and NI respectively. UI contains datasets in which objects' local scores are uniformly and independently generated for the different lists. NI contains datasets in which objects' local scores are normally and independently generated for the different lists. For synthetic datasets, our default settings for different parameters are shown in Table 1. As mentioned above, approximate top- k queries are usually applied in the cases that the values of n is fairly large, which could cause considerable cost and delays to return the exact query answers. Therefore, in our tests, the default number of data items in each list is 1,000,000, i.e. $n=1,000,000$. Typically, users are interested in a small number of top answers, thus we set $k = 500$ as the default value of k , which is a tiny value compared with n . We set m as 3 since most previous works evaluate their algorithms on datasets with 3 lists like [4]. Finally, we set 0.05 as the default value of μ .

We run our tests with default precision ($\mu = 0.05$) and high precision ($\mu = 0.005$) over each dataset respectively. Furthermore, we run the algorithms on the datasets

Table 1. Default values of experimental parameters.

| Parameters | Default Values |
|---|----------------------------|
| The number of objects, i.e. n | 1,000,000 |
| The number of lists, i.e. m | 3 |
| The number of results returned, i.e. k | 500 |
| The precision of results returned, i.e. μ | 0.05 |
| Aggregate function | $0.2s_1 + 0.3s_2 + 0.5s_3$ |

with large value of k (2000) to observe the effect of k on the performance.

For real datasets, we choose El Nino dataset¹ and Forest Cover (FC) dataset². El Nino dataset contains 93935 objects and FC dataset contains 581012 objects. El Nino contains oceanographic and surface meteorological readings taken from a series of buoys positioned throughout the equatorial Pacific. The data is expected to aid in the understanding and prediction of El Nino/Southern Oscillation (ENSO) cycles. FC contains 581012 forest land cells (i.e. objects), having four attributes (i.e. lists): horizontal distance to nearest surface water features, vertical distance to nearest surface water features, horizontal distance to nearest roadways, and horizontal distance to nearest wildfire ignition points. For both real datasets, we choose 3 lists and normalize the dataset with the formula: $\frac{s_i(t) - Min}{Max - Min}$, where $s_i(t)$ is t 's i th local score.

5.4 Experimental Results

Fig. 3 illustrates the experimental results where all the parameters are set as default values. Apparently, CIA has significant reduction on the number of accesses over every dataset. Compared with the TA_θ , CIA reduces more than 99% accesses during the query process. Apart from this, CIA is also dominant on other evaluation metrics, namely *recall*, *rank distance* and *score error* over every dataset but FC, where CIA is a little inferior to TA_θ on these aspects.

The experimental results shown in Fig. 4 when $k = 2000$ on each dataset are similar to the results when all the parameters are set as default values. From the results, we can see that CIA also has great reduction on the number of accesses compared with the TA_θ . In terms of the other aspects, CIA performs much better than TA_θ over every dataset except FC.

Fig. 5 shows us the experimental results where the parameters are set as default values except that μ , the precision of results returned is 0.005. Obviously, CIA is more efficient than TA_θ considerably but is transcended in other measures. Therefore, CIA has lower accuracy compared with TA_θ but still keeps its efficiency in the queries with high precision.

| Results for UI | <i>accesses</i> | <i>recall</i> | <i>rank distance</i> | <i>score error</i> |
|----------------|-----------------|---------------|----------------------|--------------------|
| TA_θ | 10527 | 0.50200 | 281.78800 | 0.008390 |
| CIA | 7 | 0.75600 | 88.404000 | 0.002428 |
| Results for NI | <i>accesses</i> | <i>recall</i> | <i>rank distance</i> | <i>score error</i> |
| TA_θ | 10703 | 0.52600 | 242.084000 | 0.007883 |
| CIA | 7 | 0.76800 | 88.180000 | 0.002601 |
| Results for EI | <i>accesses</i> | <i>recall</i> | <i>rank distance</i> | <i>score error</i> |
| TA_θ | 1890 | 0.29200 | 702.208000 | 0.006722 |
| CIA | 2 | 0.66600 | 124.584000 | 0.001354 |
| Results for FC | <i>accesses</i> | <i>recall</i> | <i>rank distance</i> | <i>score error</i> |
| TA_θ | 5031 | 0.99200 | 0.506000 | 0.000017 |
| CIA | 61 | 0.83800 | 35.214000 | 0.001281 |

Fig. 3. Performance of CIA vs. TA_θ when $k = 500$ and $\mu = 0.05$

¹From UCI KDD. http://kdd.ics.uci.edu/databases/el_nino/el_nino.html

²From UCI KDD. <http://kdd.ics.uci.edu/databases/coverttype/coverttype.html>

| Results for UI | <i>accesses</i> | <i>recall</i> | <i>rank distance</i> | <i>score error</i> |
|----------------|-----------------|---------------|----------------------|--------------------|
| TA_{θ} | 28778 | 0.77900 | 232.320500 | 0.003214 |
| CIA | 24 | 0.83700 | 158.745000 | 0.001843 |
| Results for NI | <i>accesses</i> | <i>recall</i> | <i>rank distance</i> | <i>score error</i> |
| TA_{θ} | 29375 | 0.80200 | 194.234000 | 0.002834 |
| CIA | 26 | 0.85100 | 136.511000 | 0.001665 |
| Results for EI | <i>accesses</i> | <i>recall</i> | <i>rank distance</i> | <i>score error</i> |
| TA_{θ} | 4519 | 0.70300 | 463.897000 | 0.006876 |
| CIA | 4 | 0.94750 | 17.667000 | 0.000258 |
| Results for FC | <i>accesses</i> | <i>recall</i> | <i>rank distance</i> | <i>score error</i> |
| TA_{θ} | 10084 | 0.94150 | 23.150500 | 0.000418 |
| CIA | 138 | 0.89650 | 75.043000 | 0.001175 |

Fig. 4. Performance of CIA vs. TA_{θ} when $k = 2000$ and $\mu = 0.05$

| Results for UI | <i>accesses</i> | <i>recall</i> | <i>rank distance</i> | <i>score error</i> |
|----------------|-----------------|---------------|----------------------|--------------------|
| TA_{θ} | 40683 | 0.99800 | 0.030000 | 0.000001 |
| CIA | 532 | 0.97400 | 1.112000 | 0.000031 |
| Results for NI | <i>accesses</i> | <i>recall</i> | <i>rank distance</i> | <i>score error</i> |
| TA_{θ} | 40371 | 0.99999 | 0.000001 | 0.000001 |
| CIA | 539 | 0.97800 | 0.678000 | 0.000023 |
| Results for EI | <i>accesses</i> | <i>recall</i> | <i>rank distance</i> | <i>score error</i> |
| TA_{θ} | 8941 | 0.99999 | 0.000001 | 0.000001 |
| CIA | 22 | 0.96200 | 2.200000 | 0.000023 |
| Results for FC | <i>accesses</i> | <i>recall</i> | <i>rank distance</i> | <i>score error</i> |
| TA_{θ} | 10482 | 0.99999 | 0.000001 | 0.000001 |
| CIA | 552 | 0.97400 | 0.840000 | 0.000027 |

Fig. 5. Performance of CIA vs. TA_{θ} when $k = 500$ and $\mu = 0.005$

Summary: From all the experimental results, we know that CIA improves significantly not only on the number of accesses, but also on other evaluation metrics in the queries with default precision. In addition, we can also learn the fact that CIA still keeps its efficiency and accuracy when the value of k is considerable large. However, CIA is not dominant on all the evaluation metrics over some datasets, like FC in our tests. Finally, in the queries with high precision, our algorithm is considerably superior to TA_{θ} on the number of accesses but have little advantage on other respects.

6. Conclusions and Future Work

In this paper, we analyzed the model of the top- k queries and gave some observations. To measure the approximation of the top- k answers, we defined a novel approximation, μ -approximation to the top- k answers. Then we introduce an efficient indexing structure called μ -cube index to support this kind of approximate query. Based on the μ -cube index on the dataset, we proposed our algorithm, the Cube Index Algorithm to answer the μ -approximation top- k queries. The main advantage of CIA is that we choose the *bottom point* of a hypercube to approximately represent the points in the hypercube and run the algorithm to find the top- $T.size$ in the set of

bottom points so that the number of accesses can be reduced significantly. Extensive experimental results on both generated and real-world datasets show that our algorithm owns higher accuracy with less cost, compared with TA_{θ} .

In the future work, we plan to turn our algorithm into exact algorithm based on the cube index ideas.

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