Tight Bounds for Distributed Selection

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Motivation: Distributed Aggregation

Growing interest in distributed aggregation!

→ Sensor networks, distributed databases...

Aggregation functions?

→ Distributive (max, min, sum, count)
→ Algebraic (plus, minus, average)
→ Holistic (median, kth smallest/largest value) ← Distributed selection

Combinations of these functions enable complex queries!

→ „What is the average of the 10% largest values?“

What cannot be computed using these functions?

Distributed selection

Motivation: Model

How difficult is it to compute these aggregation primitives?

Model:

✓ Connected graph \( G = (V,E) \) of diameter \( D_G \), \( |V| = n \).
✓ Nodes \( v_i \) and \( v_j \) can communicate directly if \( (v_i,v_j) \in E \).
✓ A spanning tree is available (diameter \( D \leq 2 \cdot D_G \))
✓ Asynchronous model of communication.
✓ All nodes hold a single element.
✓ Messages can contain only a constant number of elements.

Simple breadth-first construction!

Can easily be generalized to an arbitrary number of elements!

Motivation: Distributive & Algebraic Functions

How difficult is it to compute these aggregation primitives?

→ We are interested in the time complexity!

→ Distributive (sum, count...) and algebraic (plus, minus...) functions are easy to compute:
Use a simple flooding-echo procedure → convergecast!

Time complexity: \( \Theta(D) \)

What about holistic functions (such as k-selection)???

Is it (really) harder...?

Impossible to perform in-network aggregation?

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Motivation: Holistic Functions

It is widely believed that holistic functions are hard to compute using in-network aggregation. Example: TAG is an aggregation service for ad-hoc sensor networks → It is fast for other aggregates, but not for the MEDIAN aggregate:

"Thus, we have shown that (...) in network aggregation can reduce communication costs by an order of magnitude over centralized approaches, and that, even in the worst case (such as with MEDIAN), it provides performance equal to the centralized approach."

Thus, we have shown that (...).

Motivation: Really so Difficult?

However, there is quite a lot of literature on distributed k-selection:

A straightforward idea: Use the sequential algorithm by Blum et al. also in a distributed setting → Time Complexity: \(O(D \cdot n^{0.914})\).

A simple idea: Use binary search to find the k\textsuperscript{th} smallest value → Time Complexity: \(O(D \cdot \log x_{\text{max}})\), where \(x_{\text{max}}\) is the maximum value.

→ Assuming that \(x_{\text{max}} \in O(n^{O(1)})\), we get \(O(D \cdot \log n)\)...

A better idea: Select values randomly, check how many values are smaller and repeat these two steps! → Time Complexity: \(O(D \cdot \log n)\) in expectation!

Outline

I. Motivation/Model
II. Algorithms
III. Lower Bound
IV. Conclusion

Algorithms: Randomized Algorithm

Choosing elements uniformly at random is a good idea...

How is this done?

→ Assuming that all nodes know the sizes \(n_1, \ldots, n_t\) of the subtrees rooted at their children \(v_1, \ldots, v_t\), the request is forwarded to node \(v_i\) with probability:

\[ p_i := n_i / (1 + \sum_k n_k). \]

With probability \(1 / (1 + \sum_k n_k)\) node \(v\) chooses itself.

Key observation: Choosing an element randomly requires \(O(D)\) time!

Use pipe-lining to select several random elements!
**Algorithms: Randomized Algorithm**

Our algorithm also operates in **phases** → The set of **candidates decreases** in each phase!

A **candidate** is a node whose element is possibly the solution.

A phase of the randomized algorithm:

1. Count the **number of candidates** in all subtrees
2. Pick $O(D)$ elements $x_1, \ldots, x_d$ uniformly at random
3. For all those elements, count the **number of smaller elements**!

Each step can be performed in $O(D)$ time!

Using these counts, the **number of candidates** can be reduced by a **factor of D** in a constant number of phases with **high probability**.

We get the following result:

**Theorem**: The time complexity of the randomized algorithm is $O(D \cdot \log_D n)$ w.h.p.

We further proved a time lower bound of $Ω(D \cdot \log_D n)$.

This simple randomized algorithm is asymptotically **optimal**!

The only remaining question: What can we do **deterministically**???

**Algorithms: Deterministic Algorithm**

Why is it difficult to find a good deterministic algorithm???

→ Hard to find a good selection of elements that provably reduces the set of candidates!

Simple idea: Always propagate the median of all received values!

Problem: In one phase, only the $h^n$ smallest element is found if $h$ is the height of the tree...

→ Time complexity: $O(n / h)$

We can do a lot better!!!

Idea: Split the graph into at most $2\sqrt{D}$ groups, each containing at most $\lceil n / \sqrt{D} \rceil$ candidates. Do this recursively!

Each group has a **leader** → "Virtual tree" consisting of **leaders**

Simple bottom-up construction!

Height: $O(\log_D n)$
Algorithms: Deterministic Algorithm

A phase of the algorithm (at leader $\lambda$):

1. Receive $\leq 2\sqrt{D}$ elements from each of $\leq 2\sqrt{D}$ leader children.
2. Count the number of smaller elements for all $\leq 4D$ received elements (in all subtrees).
3. Use those counts to find $\leq 2\sqrt{D}$ elements (locally) that partition all elements into sets of size at most \( \lceil n / \sqrt{D} \rceil \) and report those elements to the next higher leader.

The number of candidates reduces by a factor of $O(\sqrt{D})$ in each phase, thus $O(\log_D n)$ phases are required.

Each phase costs $O(D \cdot \log_D n)$ time.

We get the following result:

**Theorem:** The time complexity of the deterministic algorithm is $O(D \cdot \log_D^2 n)$.

Only a factor $O(\log_D n)$ worse than the randomized algorithm!

In a grid network ($D = \sqrt{n}$), the time complexity is $\Theta(D)$, asymptotically the same complexity as when computing „easy“ aggregates!!

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**Lower Bound**

The proof of the lower bound of $\Omega(D \cdot \log_D n)$ consists of two parts:

**Part I.** Find a lower bound for the case of two nodes $u$ and $v$ with $N$ elements each.

Let $u_0 < u_1 < \ldots < u_{N-1}$ and $v_0 < v_1 < \ldots < v_{N-1}$.

How are the $2N$ elements distributed on $u$ and $v$?
Lower Bound

Assume $N = 2^b$. We use $b$ independent Bernoulli variables $X_0, \ldots, X_{b-1}$ to distribute the elements!

If $X_{b-1} = 0 \rightarrow N/2$ smallest elements go to $u$ and the $N/2$ largest elements go to $v$.

If $X_{b-1} = 1$ it is the other way round.

The remaining $N$ elements are recursively distributed using the other variables $X_0, \ldots, X_{b-2}$.

Ordered list of all $2N$ elements!

Crucial observation: For all $2^b$ possibilities for $X_0, \ldots, X_{b-1}$, the median is a different element.

Determining all $X_i$ is equivalent to finding the median!

We showed that at least $\Omega(\log_2 n)$ rounds are required if $B$ elements can be sent in a single round in this model!

Part II. Find a lower bound for the original model.

Look at the following graph $G$ of diameter $D$:

Theorem: $\Omega(D \cdot \log_D \min\{k, n-k\})$ rounds are needed to find the $k^{th}$ smallest element.

$\Omega(D \cdot \log_D n)$ lower bound to find the median!

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Conclusion

- Simple randomized algorithm with time complexity $O(D \cdot \log_2 n)$ w.h.p.
  - Easy to understand, easy to implement...
  - Even asymptotically optimal! Our lower bound shows that no algorithm can be significantly faster!
- Deterministic algorithm with time complexity $O(D \cdot \log^2 n)$.
  - If $\exists c \leq 1: D = n^c \Rightarrow k$-selection can be solved efficiently in $\Theta(D)$ time even deterministically!

Recall the 50x50 grid used to test out TAG!

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Questions and Comments?

Thank you for your attention!

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Additional Slide: Deterministic Algorithm

A phase of the deterministic algorithm step by step:

1. **A. Count the number of candidates** in all subtrees starting at the leaves.

2. **B. Build groups at the same time** → Link children together as long as each group contains at most $\lceil n / \sqrt{D} \rceil$ candidates. One node in each group becomes its leader.

3. **The leaders** split their group recursively into at most $t \leq 2 \sqrt{D}$ groups.

4. **Groups of size at most $2 \sqrt{D}$** report all values $S_i$ immediately.

5. **Once all $2 \sqrt{D} \cdot 2 \sqrt{D} = 4D$ values from all groups have arrived**, count the elements in each interval and send a selection $S$ of at most $\approx 2 \sqrt{D}$ values to the next higher leader.

Each final interval contains at most $n / \sqrt{D}$ values!