

CS639: Data Management for Data Science

Lecture 8: Reasoning about Scale & The MapReduce Abstraction

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Logistics/Announcements

• Submission template for PA2

• Bonus problem for PA2

• Other questions on PA2?

Today's Lecture

- 1. Scalability and Algorithmic Complexity
- 2. Data-Parallel Algorithms
- 3. The MapReduce Abstraction

1. Scalability and Algorithmic Complexity

What does scalable mean?

- Operationally:
 - Works even if the data does not fit in main memory
 - Use all available resources (cores/memory) on a single node (aka scale up)
 - Can make use of 1000s of cheap computers (cloud) elastic (aka scale out)
- Algorithmically:
 - If you have N data items you should not perform more than N^m operations (polynomial complexity)
 - In many cases it should be N*log(N) operations (streaming or too large data)
 - If you have N data items, you must do no more than N^m/k operations for some large k (k = number of cores/threads)

A sketch of algorithmic complexity

- Example: Find matching string sequences
- Given a set of string sequences
- Find all sequences equal to "GATTACGA"





Time = 0: TACCTGCC ? GATTACGA



Time = 0:TACCTGCC ? GATTACGANo move cursor to next data entry



Time = 1:EFTAAGCA ? GATTACGANo move cursor to next data entry



Time = 2:XXXXXX ? GATTACGANo move cursor to next data entry



Time = n:GATTACGA ? GATTACGAYes! Output matching sequence



If we have 40 records we need to perform 40 comparisons



For N records we perform N comparisons The algorithmic complexity is order N: O(N)

What if we knew the sequences are sorted





Time = 0: Start at 50% mark CTGTACA < GATTACGA



Time = 1: Start at 50% mark CTGTACA < GATTACGA Skip to 75% mark (you know your sequence is in the second half)



Time = 2: We are at the 75% mark TTGTCCA > GATTACGA Skip to 62.5% mark Match: GATTACGA = GATTACGA We find our sequence in three steps. Now we can scan entries



How many comparisons?

For N records we did log(N) comparisons

The algorithm has complexity O(log(N)) — much better scalability

2. Data-Parallel Algorithms

New task: Trim string sequences

- Given a set of string sequences
- Trim the final *n* characters of each sequence
- Generate a new dataset



Time = 0: TACCTGCC -> TACCTG



Time = 1: GATTCTGC -> GATTC



Time = 2: CCCGAAT -> CCCG

Can we use a data structure to speed this operation?



Time = 2: CCCGAAT -> CCCG

Can we use a data structure to speed this operation? No. We have to touch every record! The task is O(N).







Time = 1: Process first element of each group



Time = 2: Process second element of each group



Time = 3: Process third element of each group Etc.. How much time does this take?



We only need O(N/k) operations where k is the number of groups (workers)

Schematic of Parallel Algorithms





- Convert TIFF images to PNG
- Run thousands of simulations for different model parameters
- Find the most common word in each document
- Compute the word frequency of every word in a single document
- Etc....

- Convert TIFF images to PNG
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- Etc....
- There is a common pattern in all these applications

- A function that *maps* a string to a trimmed string
- A function that *maps* a TIFF images to a PNG image
- A function that *maps* a set of parameters to simulation results
- A function that *maps* a document to its most common word
- A function that *maps* a document to a histogram of word frequencies

• What if we want to compute the word frequency across *all* documents?



3. The MapReduce Abstraction

Compute the word frequency **across** 5M documents

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Challenge: in this task

 How can we make sure that a single computer has access to every occurrence of a given word regardless of which document it appeared in?

• Ideas?

Compute the word frequency across 5M documents









A hash function is any function that can be used to map data of arbitrary size to a data of a fixed size



The Map Reduce Abstraction for Distributed Algorithms **Distributed Data Storage** Map (Shuffle) Reduce