Databaser och bioinformatik Data structures and Indexing (I)

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IDA / ADIT

Searching and Indexing



- There are 8 Million archives.
- Each archive consists of the personal information such as person number, name, address, age, ...
- The task is given a person number, find the corresponding archive.



Searching and Indexing



- Constraint: the archives are stored in some place, you can go and fetch the archive (each time only one piece), but you can only read the information after you are back in your place.
- How much time do we need to find the archive given a person number n?



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Sort the archives



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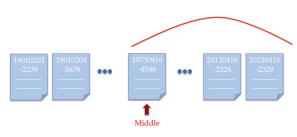
- What is the time cost for finding the archive given a person number n?

· Sort the archives according to the person number

→ Binary search!

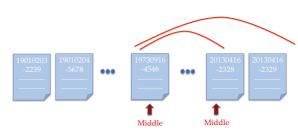


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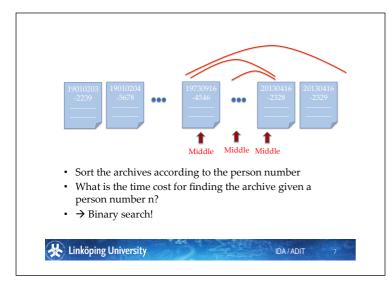


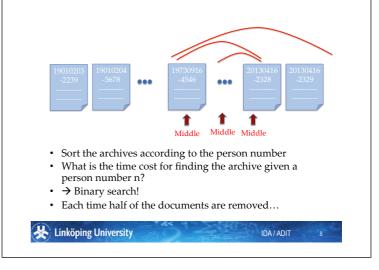


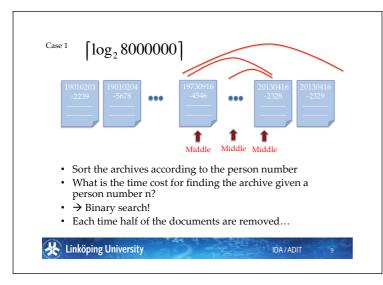
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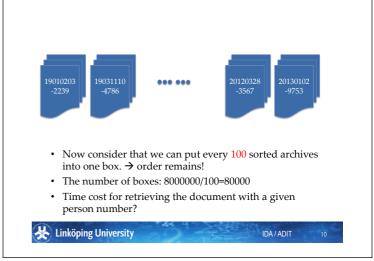


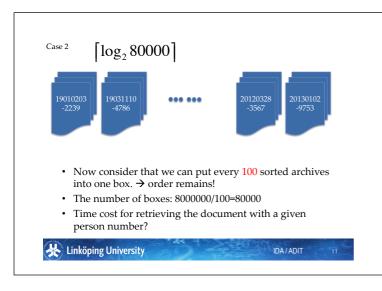
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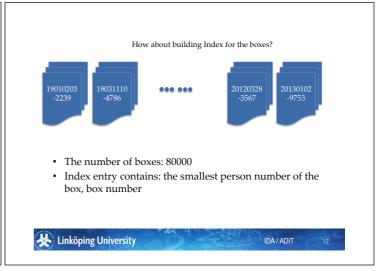


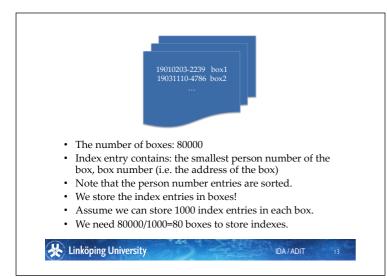


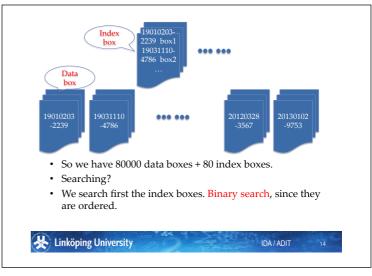


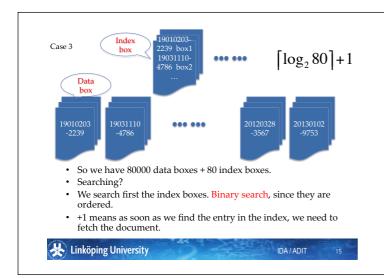


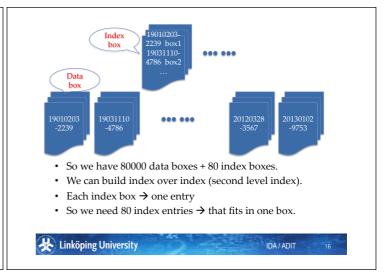


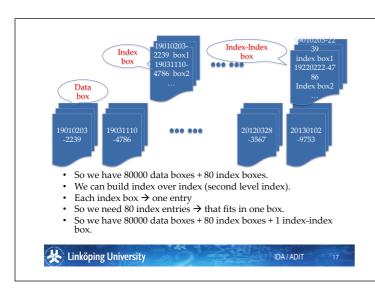


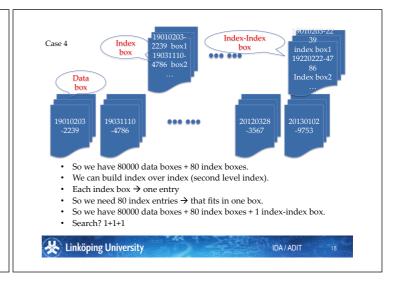


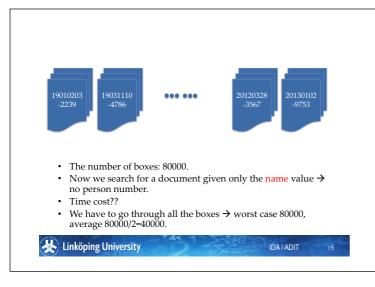


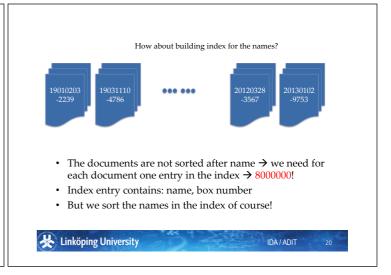


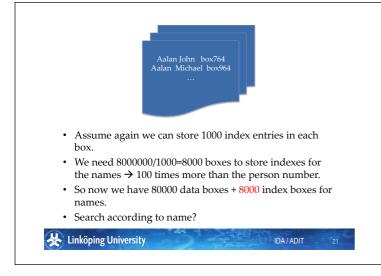


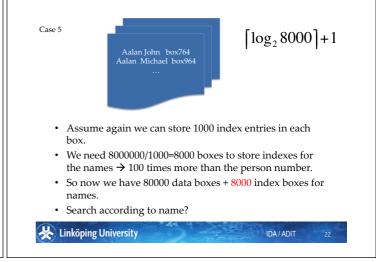


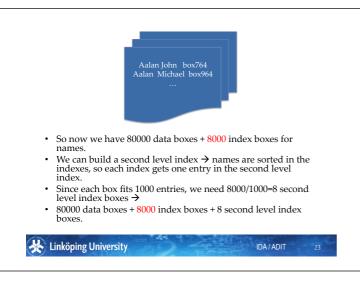


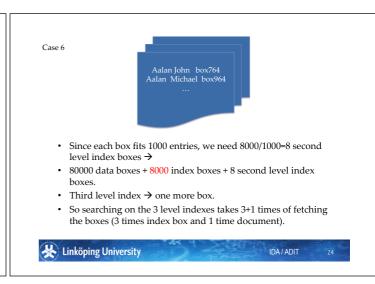


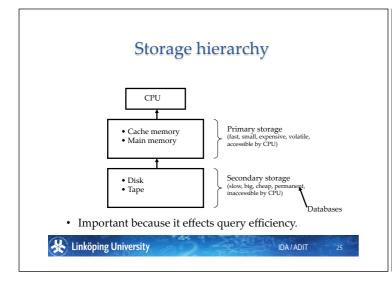


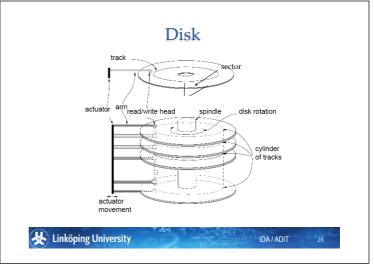


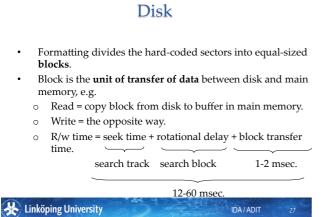


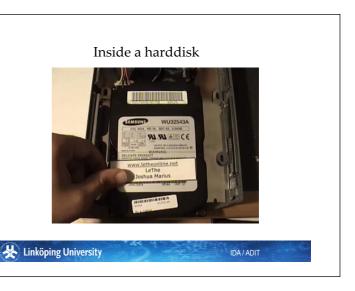












Files and records

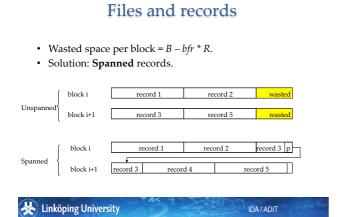
- Data stored in files.
- File is a **sequence** of **records**.
- Record is a set of field values.
- For instance, file = relation, record = entity, and field = attribute.
- Records are allocated to file **blocks**.

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Files and records

- Let us assume
 - \circ *B* is the size in bytes of the block.
 - \circ *R* is the size in bytes of the record.
 - o *r* is the number of records in the file.
- Blocking factor: $bfr = \left| \frac{B}{R} \right|$
- Blocks needed: $b = \left[\frac{r}{bfr} \right]$
- What is the space wasted per block?





From file blocks to disk blocks

- Contiguous allocation: cheap sequential access but expensive record addition. Why ?
- Linked allocation: expensive sequential access but cheap record addition. Why ?
- Linked clusters allocation.
- Indexed allocation.



File organization

- Heap files.
- · Sorted files.
- · Hash files.
- File organization != access method, though related in terms of efficiency.

