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Exercise 1

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Exercise 1 a)

Describe the Cheung formula.

Compute the Cheung formula for the values  $N = 1000$ ,  $m = 100$ ,  $k = 15$

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Exercise 2

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Exercise 2 a)

Read the chapter on *Genetic Algorithms* in the *Probabilistic Algorithms* section in the script.

Exercise 2 b)

Implement the genetic algorithm for the join ordering problem. It is sufficient to consider only left deep trees.

Note, on a high level, it is easy to implement the algorithm. However you will need a significant number of helper functions.

Feel free to take the solution code and only implement the high level level algorithm (Run function).

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Exercise 3

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Queries with multiple predicates may require the evaluation of multiple indices. In what order should you process the predicates?

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Exercise 4

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Instead of using B-Tree one can also use hashing based data structures. For instance, one can build on the idea of extendible hashing

[https://en.wikipedia.org/wiki/Extendible\\_hashing](https://en.wikipedia.org/wiki/Extendible_hashing)

For queries with what type of predicates can you apply such data structures?

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Exercise 5

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You are given a relation with 6 tuples. These tuples are equally distributed over 3 pages. Compute the average number of page accesses for reading 2 tuples. Assume that all tuples have the same probability to be read.

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Exercise 6

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*BONUS*

Write a program that allows you to measure some physical properties of your disk.