

## LSDS Assignment 2

Command for running Problem A : `srun homework_2 1 human.txt human_reads.fa`

Command for running Problem B : `srun homework_2 2 human.txt human_reads.fa`

### A) Unsorted Linear search :

- a. How long did it take you to search for the first 10K, 100K, and 1M 32-character long fragments of the subject dataset within the query dataset?

**Time taken for searching 10k fragments of the subject dataset is : 12190.066 seconds**

**Time taken for searching 100k fragments of the subject dataset is : 117431.09 seconds**

**Time taken for searching 1M fragments of the subject dataset is : 1210354.93 seconds**

- b. How long would it take to search for every possible 32-character long fragment of the subject dataset within the query dataset?

**Time taken for searching 3B fragments of the subject dataset is : 3614457925.939 seconds**

- c. Print the first 10 fragments of the subject dataset that you found within the Query AR object (if any)?

File size of genome data is 3057186663

File size of query dataset is 27772294

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First 10 matching fragments in 10000 fragments are

Matching index in genome data : 0

Matching index in genome data : 1

Matching index in genome data : 2

Matching index in genome data : 3

Matching index in genome data : 4

Matching index in genome data : 5

Matching index in genome data : 6

Matching index in genome data : 7

Matching index in genome data : 8

Matching index in genome data : 9

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First 10 matching fragments in 100000 fragments are

Matching index in genome data : 0

Matching index in genome data : 1

Matching index in genome data : 2

Matching index in genome data : 3

Matching index in genome data : 4

Matching index in genome data : 5

Matching index in genome data : 6

Matching index in genome data : 7

Matching index in genome data : 8

Matching index in genome data : 9

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First 10 matching fragments in 1000000 fragments are

Matching index in genome data : 0
Matching index in genome data : 1
Matching index in genome data : 2
Matching index in genome data : 3
Matching index in genome data : 4
Matching index in genome data : 5
Matching index in genome data : 6
Matching index in genome data : 7
Matching index in genome data : 8
Matching index in genome data : 9
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**B) Sorted Binary search:**

- a. How long did it take you to search for the first 10K, 100K, and 1M 32-character long fragments of the subject dataset within the query dataset?

**Time taken for searching 10k fragments of the subject dataset is : 0.04 seconds**

**Time taken for searching 100k fragments of the subject dataset is : 0.51 seconds**

**Time taken for searching 1M fragments of the subject dataset is : 4.87 seconds**

- b. How long would it take to search for every possible 32-character long fragment of the subject dataset within the query dataset?

**Time taken for searching 3B fragments of the subject dataset is : 14533.358 seconds**

- c. Print the first 10 fragments of the subject dataset that you found within the Query AR object (if any)?

File size of genome data is 3057186663  
File size of query dataset is 27772294  
Sorting and applying binary search

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First 10 matching fragments in 10000 fragments are

Matching index in genome data : 0  
Matching index in genome data : 1  
Matching index in genome data : 2  
Matching index in genome data : 3  
Matching index in genome data : 4  
Matching index in genome data : 5  
Matching index in genome data : 6  
Matching index in genome data : 7  
Matching index in genome data : 8  
Matching index in genome data : 9

Total time taken for program execution after sorting for 10000 fragments are 0.048922 seconds

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Sorting and applying binary search

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First 10 matching fragments in 100000 fragments are

Matching index in genome data : 0  
Matching index in genome data : 1  
Matching index in genome data : 2  
Matching index in genome data : 3  
Matching index in genome data : 4  
Matching index in genome data : 5  
Matching index in genome data : 6  
Matching index in genome data : 7  
Matching index in genome data : 8  
Matching index in genome data : 9

Total time taken for program execution after sorting for 100000 fragments are 0.513192 seconds

Sorting and applying binary search

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First 10 matching fragments in 1000000 fragments are

Matching index in genome data : 0  
Matching index in genome data : 1  
Matching index in genome data : 2  
Matching index in genome data : 3  
Matching index in genome data : 4  
Matching index in genome data : 5  
Matching index in genome data : 6  
Matching index in genome data : 7  
Matching index in genome data : 8  
Matching index in genome data : 9

Total time taken for program execution after sorting for 1000000 fragments are 4.87 seconds

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