HOMEWORK ASSIGNMNENT- I

LARGE SCALE DATA STRUCTURES & ORG - INF 503



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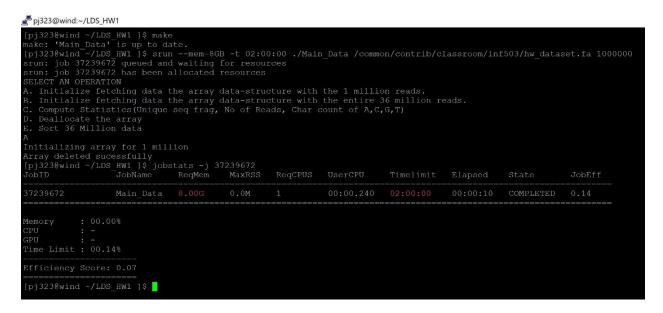
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Question A

Using the first 1 million reads, estimate, and report the total CPU time and RAM it will take to initialize (fill up) the array data-structure with the entire 36 million reads. Note that this may mean creating custom constructor to read first X reads rather than to the End-Of-File.

Initialized Job ID = 37239672

We re fetching 1 Million Reads or data and computing the system usage as asked in the Question above. Attaching a Screen shot of the same below.



Here we can see that the JOB was completed in **10 Seconds**. This shows nothing but 10 Seconds of the CPU usage time.

We can see that allocated RAM Size was **8GB**, but the used RAM was 0.0.

Manual Calculation

```
If 1000000 (1M) Data = 10 Seconds
```

Then for 36000000 (36M) Data = ?

On Manual calculation we know the general formula RAM = O(n)

So we can use general formula and we find that it takes 3 Minutes for initializing 36 Million Data.

we can find the actual time it takes to do the same by initialising to fetch 36 Million data by running the Problem statement B.

Question B

Test your prediction using the entire 36 million read set – report actual RAM and CPU time used. Refer to Monsoon workshop notes for help in estimating actual runtime and RAM usage of your run. Were you accurate? If not, explain what you think caused the discrepancy.

Here we initialize to find the actual CPU time for initializing 36 Million Data and compare the statistics with the manual calculation.

```
[pj323@wind ~/LDS_HW1 ] $ srun --mem=8GB -t 02:00:00 ./Main_Data /common/contrib/classroom/inf503/hw_dataset.fa 36000000 srun: job 37239040 queued and waiting for resources srun: job 37239040 has been allocated resources
SELECT AN OPERATION
A. Initialize fetching data the array data-structure with the 1 million reads.
B. Initialize fetching data the array data-structure with the entire 36 million reads.
C. Compute Statistics(Unique seq frag, No of Reads, Char count of A,C,G,T)
D. Deallocate the array
E. Sort 36 Million data
B
Initializing array for 36 million
Array deleted successfully
[pj323@wind ~/LDS_HW1 ] $ jobstats -j 37239840
Job1D JobName ReqMem MaxRSS ReqCPUS UserCPU Timelimit Elapsed State JobEff

37239840 Main_Data 8.00G 0.0M 1 00:07.365 02:00:00 00:00:30 COMPLETED 0.42

Memory : 00.00%
CPU :-
Time Limit : 00.42%

Efficiency Score: 0.21

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```

To initialize 30M data it just took 30 Seconds

The Average CPU time used here is just = 30 Seconds on a comparison it is faster and processing the more amount of data it just took 30 seconds.

Question C

Compute the following statistics for your read set

- Total number of unique sequence fragments (here, safe to assume this is the total number of sequence fragments in the file).
- Total number of reads for each 'data set' separately (recall there are 14 data sets in our example here). You will need 14 different totals.
- Number of A, C, G, and T characters in the dataset.

This is the Output when we tried to compute the data of **36000000** (**36M**) **reads**. Was taking longer time for computation.

The Count

A = 493102372

C = 406639890

G = 408544523

T = 489180159

Computing for **100,000 Data** and checking the unique Sequence so we are calculating all the reads which are unique.

It copares with every read and chech for the data

Time Taken = 28:01 min

Ram Used = **13.8 GB**

No of Unique Sequence = 100000

Complete computation was done and the Output was given.

Question D

Implement a destructor for your class to delete / deallocate your array data structure. How long did it take? Does this make sense to you?

Basically, we are trying to Deallocate the Array.

~Main_Data()- destructor deallocates the memory of the array as soon as control reaches the end of the class.

When the function is run on Monsoon it takes around **22 Seconds** to Deallocate the Memory that is present in the Array.

Question E

Implement a function that would sort the genomic sequences (fragments not characters within a fragment) in your array in alphabetic order.

- What is the 'big O' notation of your approach (linear / quadratic / cubic / etc)? Please note that depending on the efficiency of your algorithm, you may not be able to alphabetically sort the entire 36 million reads in a reasonable amount of time (24-36 CPU hours). If this happens, reduce the problem size (by using a smaller subset of the reads) and estimate the final run time.
- Print the first 20 lines of the sorted output.

```
|pj323@wind ~/LDS_INI | $ srun --mem=866 -t 02:00:00 ./Main_Data /common/contrib/classroom/inf503/hw_dataset.ta 36000000 srun: job 37239867 queued and waiting for resources SULDCT AN OPENATION been allocated resources SULDCT AN OPENATION.

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```

Time complexity of Merge sort is **O**(nLogn)-which is same in worst case.

It is a logarithmic approach. Space complexity of Merge sort is O(n).

It took RAM = 4.96GB to sort

Time= 1 minute 53 seconds.