

Project 3

Xiao Wei's Problem II

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The story continues...

Xiao Wei's Problem II

- Xiao Wei majors in biology
- He's working on a DNA matching project
- The size of the database is HUGE
- He needs your help, *again*

General Information

- Implement a DNA sequence matching program
- In Java
- Handout: Dec 1, 2014
- Available on FTP **PROJECT** directory

Usage

```
$ java -jar dna_matching.jar db_file query_file output_file
```

Input

- Sequence format
 - String made up of only four characters: A, C, G and T
 - Prefixed by a line starting with a greater than character (>) followed by a description of the sequence
- Database file
 - Consists of numerous *database sequences*
- Query file
 - Consists of numerous *query sequences*

Sample Database File

```
>DB description string 1
```

```
AGCTTTTTCATTCTGACTGCAACGGGCAATATGTCTCTGTGTGGATTAAAAAAAGAGTGTC  
TGATAGCAGCTTCTGAACTGGTTACCTGCCGTGAGTAAATTAAAATTTTATTGACTTAGG  
TCACTAAATACTTTAACCAATATAGGCATAGCGCACAGACAGATAAAAATTACA
```

```
>DB description string 2
```

```
AACGGTGCGGGCTGACGCGTACAGGAAACACAGAAAAAAGCCCGCACCTGACAGTGCGGG  
CTTTTTTTTTTCGACCAAAGGTAACGAGGTAACAACCATGCGAGTGTTGAAGTTCGGCGGT  
ACATCAGTGGCAAATGCAGAACGTTTTCTGCGTGTTGCCGATATTCTGGAAAGCAATGC
```

```
>EOF
```

Sample Query File

[illegible]

Output

- Report the location of an exact match within any given input DNA sequence for each input search query sequence
- If the query sequence matches within multiple database sequences within the database file, report each result
- If the query sequence matches multiple locations within the same database sequence, report the earliest position of match
- Print **NOT FOUND** otherwise

Sample Output File

Query description string 1

[DB description string 1] at offset 7

Query description string 2

[DB description string 1] at offset 47

[DB description string 2] at offset 33

Query description string 3

[DB description string 1] at offset 94

[DB description string 2] at offset 79

Query description string 4

NOT FOUND

How to Solve?

- Algorithms
 - Naïve matching
 - Knuth–Morris–Pratt algorithm
 - Boyer–Moore algorithm
 - Rabin–Karp algorithm
- Techniques
 - Multithreading
 - Hashing
 - MapReduce

Grading

- Correctness: 60%
- Time performance: 20%
 - Prerequisite: pass all correctness tests
- Project development document: 15%
- User manual: 5%
- GUI not required

Submission

- Deadline: Dec. 20 2014, 23:59 (GMT+08:00)
- Package your project
 - Source code
 - Executable jar
 - User manual
 - Development document
- Submit to FTP
- Face-to-face interview

Policy

- No cheating
- No late policy

Thanks!

Q&A

Feel free to contact me via E-mail or WeChat