Bioinformatics - Assignment 1

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Coding Language: Python

To run the assignment:

#To run code

Mutation sequence_length $p(A) p(C) p(G) p(T) no_of_sequences mutation_probability(0:1 interval) outputfilePath$

Split_sequencer inputfilePath min_fragment_length max_fragment_length outputfilePath

Sequence_Assembler inputfilePath s r d outputfilePath s-score for match r-score for replace d- score for delet