Global Sequence Alignment lab

Name: _____

_____ 50 Code
_____ 10 recursiveEditDistance
_____ 10 editDistance
_____ 20 printEditDistance
_____ 10 timeRecursiveEditDistance, timeEditDistance, randomDNAString
_____ 20 Comments and Style
_____ 5 Adequate comments
_____ 10 Single copy of code used by editDistance and printEditDistance
_____ 5 Choice of names, indentation, other style issues
_____ 30 Questions in readme.txt
_____ 10 running time of recursiveEditDistance
_____ 5 distance and timing for the various E. coli
_____ 5 doubling hypotheis

____ TOTAL

____ 10 running time estimation

MCS-178