

CS321: Numerical Methods in Comp Mol Bio

Homework 11

Due: Thursday, Dec 1 2005 at the beginning of the section

Problem 1

Write a Matlab function that accepts two sequences, a similarity matrix and a gap penalty score and returns an optimal local alignment of the two sequences along with its score.

Use Blosun50 and a gap penalty of -5 to find the optimal local alignment for all the pairs of sequences seq1 to seq4.

- a) report your alignments and their scores.
- b) comment on which of the sequences you believe are related to one another.

Problem 2

On the same sequences as problem 1, find the percentage of global identity between all pairs.

- a) report your alignments and their scores.
- b) comment on which of the sequences you believe are related to one another.

Problem 3

For extra credit:

Write a Matlab function that accepts two sequences, a similarity matrix a gap open penalty score and a gap extension penalty score and returns an optimal local alignment of the two sequences along with its score.

Use Blosun50 a gap open penalty of -10 and a gap extension penalty of -3 to find the optimal local alignment for all the pairs of the sequences:

- a) report your alignments and their scores.
- b) comment on which of the sequences you believe are related to one another.