Grading Criterion

Issues	Highest possible grade
Wrong alignment scores	50% to 70%
Correct scores but wrong alignment (problems with back tracing)	80.00%
Correct program with small bugs on some cases (e.g. out of memory, segmentation fault, etc.)	90.00%
Program did not finish in 20 minutes for some tests	90.00%
Significant restrictions on input file formats (e.g. white space sensitive) or difference from specified output	90.00%
Using (e.g. extending) alignment related classes from biological software packages depending on extent of usage	70% to 90%
Problems with compiling on Glue	90.00%

- Late penalties and speed competition bonuses apply
- Programs with incorrect output were not included in speed competition