

Gorilla Report

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Results

Our implementation produces the expected results on all pairs of species. Even though the alignments are not *exactly* the same as the out-files, they are still *an* optimal solution.

The output after solving the smallest input file (the Toy-file) is:

INSERT OUTPUT FROM TOY FILE

Implementation details

We have chosen the bottom-up approach by creating a matrix of size $m \times n$. This matrix will contain values for the alignment and takes $O(m * n)$ time to create.