

# Exercises – Phylogenetics

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Dr. Roland Wittler, Prof. Dr. Jens Stoye, M. Sc. Tizian Schulz  
<https://gi.cebitec.uni-bielefeld.de/Teaching/2017winter/Phylogenetik>

## Exercise Sheet 9 — 07.12.2017

Due: 14.12.2017

### Task 1 Neighbor Joining.

(8 points)

Reconstruct a phylogenetic tree from the following matrix. Use

- (a) WPGMA,
- (b) Waterman's algorithm and
- (c) *Neighbor Joining*.

	<i>A</i>	<i>B</i>	<i>C</i>	<i>D</i>	<i>E</i>
<i>A</i> :	0	8	5	8	2
<i>B</i> :		0	8	6	6
<i>C</i> :			0	8	4
<i>D</i> :				0	6
<i>E</i> :					0

Write down all steps.

Hint: When performing Waterman's algorithm, start with the two taxa *A* and *B*, and try to add all taxa to their edge first. As soon as this is not possible anymore, continue with the path between *A* and *C*. You might get negative edge weights.

### Task 2 Minimum Evolution.

(5 points)

Write a linear program (LP) that calculates the *minimum evolution tree* (ME tree) for the distance matrix from Task 2 on exercise sheet 8 and the tree topology resulting for the UPGMA method.

Use an online solver to get a solution for it, for instance:

- <http://www.zweigmedia.com/RealWorld/simplex.html> or
- <http://www.phpsimplex.com/simplex/simplex.htm?l=en>.

Send your LP to your T.A.<sup>1</sup>

What are the minimum total edge length and the individual edge lengths?

In comparison, the resulting topology for UPGMA has a total edge length of 29. Compare these results and the results from Task 2 on exercise sheet 8.

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<sup>1</sup>tischulz[at]cebitec...