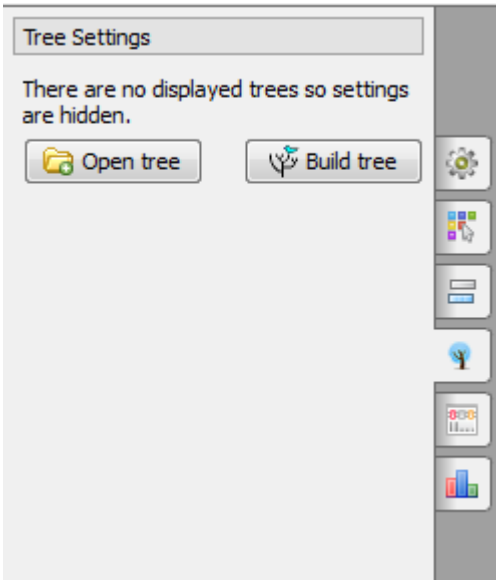
Building Phylogenetic Tree

To build a tree from an alignment either press the *Build Tree* button on the toolbar, select the *Tree Build Tree* item in the alignment context menu or the *Actions Tree Build Tree* item in the main menu.

Also you can use *Tree Settings* tab of the *Options Panel*:



Three methods for building phylogenetic trees are supported:

1. The *PHYLIP Neighbour-Joining* method. The PHYLIP package implementation of the method is used under the hood.
2. The *MrBayes* external tool. Check MrBayes Web Site for more details.
3. PhyML Maximum Likelihood method. Check PhyML Maximum Likelihood Web Site for more details.

- PHYLIP Neighbor-Joining
- MrBayes
- PhyML Maximum Likelihood