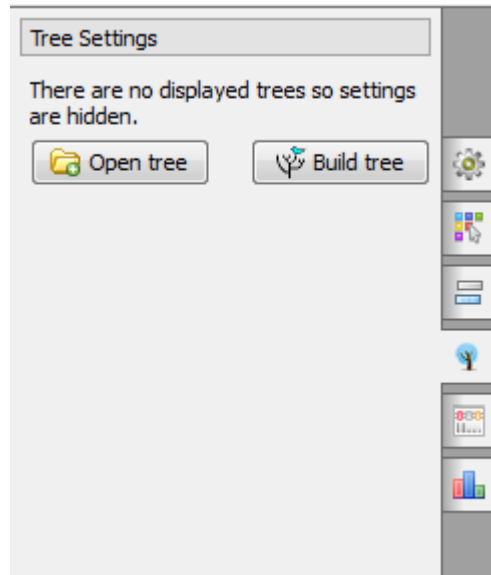


Building Phylogenetic Tree

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 *PHYMLIP Neighbour-Joining* method. The [PHYMLIP](#) 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](#)