

Alignment Editor

- Overview
 - Alignment Editor Features
 - Alignment Editor Components
 - Navigation
 - Coloring Schemes
 - Creating Custom Color Scheme
 - Highlighting Alignment
 - Zooming and Fonts
 - Searching for Pattern
 - Consensus
 - Export Consensus
 - Alignment Overview
- Working with Alignment
 - Undo/Redo Framework
 - Selecting Subalignment
 - Moving Subalignment
 - Editing Alignment
 - Removing Selection
 - Filling Selection with Gaps
 - Replacing with Reverse-Complement
 - Replacing with Reverse
 - Replacing with Complement
 - Removing Columns of Gaps
 - Removing All Gaps
 - Saving Alignment
 - Aligning Sequences
 - Pairwise Aligning
 - Working with Sequences List
 - Adding New Sequences
 - Copying Sequences
 - Renaming Sequences
 - Sorting Sequences
 - Shifting Sequences
 - Collapsing Rows
 - Exporting in Alignment
 - Extracting Selected as MSA
 - Exporting Sequence from Alignment
 - Exporting Alignment as Image
- Statistics
 - Distance Matrix
 - Grid Profile
- Advanced Functions
 - Building HMM Profile
- Building Phylogenetic Tree
 - PHYLIP Neighbor-Joining
 - MrBayes
 - PhyML Maximum Likelihood