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
    - 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