



- **Demo** of selected tools in the Structure Comparison category
  - Multalign Viewer - sequence alignment viewer with many features, including crosstalk to and from associated structures
  - MatchMaker - constructs pairwise sequence alignments and matches structures accordingly (demo: 1jta, 1bn8, 2pec)
  - Match -> Align - constructs pairwise or multiple sequence alignments based on pre-existing structural superpositions
- **Hands-on experience**
  - Sequences and Structures tutorial (the online version can be accessed from the Chimera Help menu)

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## Notes

This image shows a single sheet of white paper with horizontal blue or grey ruling lines. The lines are evenly spaced and run across the width of the page. There are no margins, text, or other markings on the paper.