



## Exploring sequence-structure relationships

- **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