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Protocol Run: ChimeraModelFromTemplate

**Protocol: chimera - model from template** finished Cite Help

**Run**

Run name: chimera - model from template Comment

Run mode: ☒ Continue ☐ Restart ? Host: localhost

Use queue? ☐ Yes ☒ No ?

Wait for: ?

**Input** Help

Atomic structure used as template: scipion - import atomic structure 1PBX.output Q 🗑 👁 ?

Chain: [{"model": 0, "chain": "A", "residues": 142}] 🔧 ?

Target sequence: scipion - import sequence HBA\_HUMAN.outputSequence Q 🗑 👁 ?

Additional sequences to align? ☒ Yes ☐ No ?

Other sequences to align

Object	Info
scipion - import sequence ALDGI.outputSequence	Sequence (name = HBA_ALDGI_P8313
scipion - import sequence MELGA.outputSequence	Sequence (name = HBA_MELGA_P810
scipion - import sequence RABIT.outputSequence	Sequence (name = HBA_RABIT_P0194
scipion - import sequence HORSE.outputSequence	Sequence (name = HBA_HORSE_P019

Multiple alignment tool: Clustal Omega Q 🗑 👁 ?

Close Save Execute