

View Model building

- Model building
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 - import sequence
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Protocol Run: ChimeraModelFromTemplate

Protocol: chimera - model from template

Run

Run name: Comment:

Run mode: ☒ Continue ☐ Restart Host:

Use queue? ☐ Yes ☒ No

Wait for:

Input

Atomic structure used as template:

Chain:

Target sequence:

Options to improve the alignment:

Other sequences to align

Object	Info
pwem - import sequence HBA_TURTLE.out	Sequence (name = HBA_TURTLE.out)
pwem - import sequence HBA_TURKEY.out	Sequence (name = HBA_TURKEY.out)
pwem - import sequence HBA_RABBIT.out	Sequence (name = HBA_RABBIT.out)
pwem - import sequence HBA_HORSE.out	Sequence (name = HBA_HORSE.out)

Multiple alignment tool:

Additional target sequence to include? ☐ Yes ☒ No

Analyze Results

Execute