

A

Project Help

SCIPION (2018-)

View Protocols SPA

- Classify
- 3D
 - Initial volume
 - Preprocess
 - Classify
 - Refine
 - Analysis
 - Reconstruct
- Tools
 - Sets
 - Calculators
 - xmipp3 - operate particles
 - xmipp3 - operate volumes
 - chimera - chimera operate
 - chimera - chimera restore s
 - phenix - superpose pdbs
- Model Building
 - xmipp3 - extract unit cell
 - chimera - rigid fit
 - scipion - model from template**
 - powerfit - powerfit
 - ccp4 - coot refinement
 - ccp4 - reformat
 - phenix - emringer
 - phenix - real space refine
 - phenix - molprobit

B

Protocol Run: ChimeraModelFromTemplate

Protocol: scipion - model from template

Cite Help

Run

Run name: scipion - model from template

Comment:

Run mode: ☒ Continue ☐ Restart

Host: localhost

Use queue? ☐ Yes ☒ No

Wait for:

Input

PDBx/mmCIF file template:

Chain:

Target sequence:

Additional sequences to align? ☒ Yes ☐ No

Other sequences to align:

Object	Info

Multiple alignment tool: Clustal Omega

Close Save Execute