

A

Project Help



View Protocols SPA

3D

- ▶ Initial volume
- ▶ Preprocess
- ▶ Classify
- ▶ Refine
- ▶ Analysis
- ▶ Reconstruct

Tools

- ▶ Sets
- ▶ Calculators
 - ▶ xmipp3 - operate particles
 - ▶ xmipp3 - operate volumes
 - ▶ chimera - chimera operate
 - ▶ chimera - chimera restore session
 - ▶ phenix - superpose pdb

Model Building

- ▶ xmipp3 - extract unit cell
- ▶ chimera - rigid fit
- ▶ scipion - model from template
- ▶ powerfit - powerfit
- ▶ ccp4 - coot refinement
- ▶ ccp4 - reftmac
- ▶ phenix - emringer
- ▶ phenix - real space refine
- ▶ phenix - molprobability

B

Protocol Run: ChimeraProtRigidFit



Protocol: chimera - rigid fit

Cite Help

Run

Run name chimera - rigid fit (5)

Comment

Run mode

☒ Continue
 ☐ Restart

Host

localhost

Use queue?

☐ Yes
 ☒ No

Wait for

Input

Help

Input

Input Volume

PDBx/mmCIF file to be refined

Other reference PDBx/mmCIF files

Object

Info

Close

Save

Execute