

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 pdbs

Model Building

- xmipp3 - extract unit cell
- chimera - rigid fit
- scipion - model from template
- powerfit - powerfit
- ccp4 - coot refinement

Protocol Run: ChimeraProtOperate finished [Cite](#) [Help](#)

Protocol: chimera - chimera operate

Run

Run name chimera - chimera operate **Comment**

Run mode ☒ Continue ☐ Restart **Host** localhost

Use queue? ☐ Yes ☒ No

Wait for

Input **Help**

Input

Input Volume

PDBx/mmCIF file scipion - import atomic structure 5ni

Other PDBx/mmCIF files

Object	Info

Execute **Close** **Save**

Protocols | Data

[Refres](#)

atomic structure 5
finished

- chimera operate
finished

Analyze Results