

View Protocols SPA

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P xmipp3 - operate particles

P xmipp3 - operate volumes

P chimera - chimera operate

P chimera - chimera restore sessi

P phenix - superpose pdbs

▼ Model Building

P xmipp3 - extract unit cell

P chimera - chimera rigid fit

P chimera - model from template

P powerfit_scipion - powerfit

P ccp4 - coot refinement

P ccp4 - refmac

P phenix - emringer

P phenix - real space refine

P phenix - molprobity

P xmipp3 - 3d bionotes



Protocol: ccp4 - coot refinement

Cite Help

Run

Run name ccp4 - coot refinement HBA_H

Comment

Run mode ☒ Continue ☐ Restart

Host localhost

Use queue? ☐ Yes ☒ No

Wait for

Input Help

Input

Input Volume/s

Object

Info

xmipp3 - extract unit cell symC2 offse Volume (101 x 101 x 101)

Normalize

☒ Yes ☐ No

Atomic structure to be refined

chimera - rigid fit HBA_HUMAN.out

Other reference atomic structures

Object

Info

chimera - rigid fit HBA_HUMAN.output PdbFile (pseudoatoms)

Close

Save

Execute