

View Protocols SPA

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▼ Model Building

▶ xmipp3 - extract unit cell

▶ chimera - rigid fit

▶ scipion - model from template

▶ powerfit - powerfit

▶ ccp4 - coot refinement

▶ ccp4 - reflat

▶ phenix - emringer

▶ phenix - real space refine

▶ phenix - molprobt

Edit Copy

ccp4 - coot

phenix - real

Summary Me

▼ Input

inputVolume (from xmipp3 - extract unit cell symC2 offset -45 -> outputVolume Volume (101 x 101 x 101, 1.05 Å/px)

SUMMARY

Protocol Run: PhenixProtRunRSRefine

Phenix

Protocol: phenix - real space refine

Cite Help

Run

Run name phenix - real space refine HBA

Comment

Run mode

☒ Continue

☐ Restart

Host

localhost

Use queue?

☐ Yes

☒ No

Wait for

Expert Level

☒ Normal

☐ Advanced

Input

Input

Input Volume xmipp3 - extract unit cell symC2 off

Resolution (Å): 3.2

Input atomic structure chimera - rigid fit HBA_HUMAN.out

Output format

pdb

Close

Save

Execute

Protocols | D

Ref

scipion - import sequen
finished

HBA_HUMAN

scipio

Analyze Results