

SCIPION v2.0 (2019-0)

View: Model building

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Protocol Run: ChimeraProtRigidFit

Protocol: chimera - chimera rigid fit finished [Cite](#) [Help](#)

Run

Run name: chimera - chimera rigid fit [Edit](#) Comment: [Edit](#)

Run mode: ☒ Continue ☐ Restart [?](#) Host: localhost [?](#)

Use queue? ☐ Yes ☒ No [?](#)

Wait for: [?](#)

Input [Help](#)

Input

Input Volume: xmipp3 - extract unit cell sym C2.outputVolume [Search](#) [Delete](#) [View](#) [Help](#)

Atomic structure to be fitted: powerfit_scipion - powerfit.outputPDBs.2 [Search](#) [Delete](#) [View](#) [Help](#)

Other reference atomic structures:

Object	Info
powerfit_scipion - powerfit.outputPDBs.1	AtomStruct(ps)

Close **Save** **Execute**