

import volumes import atomic structure

import sequence

mxmipp3 - create 3d mask

xmipp3 - local MonoRes

mxmipp3 - extract unit cell

rigid fit

phenix - real space refine ccp4 - coot refinement ccp4 - refmac Validation

phenix - emringer

phenix - molprobity

chimera - contacts

phenix - superpose pdbs atomstructutils - operator

chimera - chimera restore session

EMRinger Score: 1.41

Preprocess map

Initial model

Rigid fitting

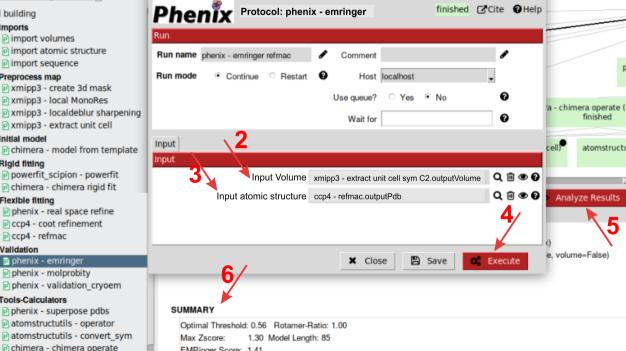
Flexible fitting

Model building

Demo Modeling

Protocols | Da

A CRefre



Protocol Run: PhenixProtRunEMRinger