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Protocol Run: PhenixProtRunMolprobability

Phenix Protocol: phenix - molprobability finished Cite Help

Run

Run name: phenix - molprobability refrac (co) edit Comment: edit

Run mode: ☒ Continue ☐ Restart ? Host: localhost

Parallel Threads: 1 ? Use queue? ☐ Yes ☒ No ?

Wait for: ?

Expert Level: ☒ Normal ☐ Advanced

Input

Input Volume: xmipp3 - extract unit cell sym C2.outputVolume search delete eye ?

Resolution (A): 3.2 ?

Input atomic structure: ccp4 - refrac.outputPdb search delete eye ?

Close Save Execute

SUMMARY

MolProbability statistics:

Ramachandran outliers: 1.43 %	(Goal: < 0.2%)	Ramachandran favored: 97.14 %	(Goal: > 98%)
Rotamer outliers: 7.89 %	(Goal: < 1%)	C-beta outliers: 4	(Goal: 0)
Clashscore: 90.42		Overall score: 3.26	

MolProbity: <http://molprobity.biochem.duke.edu/>

Refre

pl

ra - chimera operate (H
finished

cell) atom structure

Analyze Results

e, volume=False)