Protocol Viewer: PhenixProtRunRSRefineViewer	9 📵 😣
Phenix Protocol: phenix - Real Space Refine viewer Cite	⊘ Help
Volume and models MolProbity results Real-space correlation Atomic properties	
MolProbity results	
-Summary MolProbity	
MolProbity Basic Statistics	● 0
Open In Coot	® Ø
Missing atoms	® 0
Basic Geometry: Bond Length Restraints	
Deviations	® Ø
Basic Geometry: Bond Angle Restraints	
Deviations	® 0
Basic Geometry: Dihedral Angle Restraints	
Deviations	® Ø
Outliers	® Ø
Basic Geometry: Chirality Restraints	
Deviations	® 0
Basic Geometry: Planarity Restraints	
Deviations	® Ø
Protein	
Select plot: Ramachandran plot	0
View plot	® Ø
No Ramachandran outliers detected	0
No Rotamer outliers detected	0
No C-beta position outliers detected	0
Recommended Asn/Gin/His sidechain flips:	® 0
No non-trans peptides detected	0
-Clashes	
All atom-contact analysis	⊕ 0

× Close