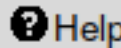




Protocol: phenix - Validation cryoEM viewer



Cite



Help

Volume and models



Summary

MolProbity


Model vs. Data

Data





Model vs. Data

Overall correlation coefficients  

Correlation graphs

Plot CC vs. Chain ID Save chain CC as text 

Chain A

Plot CC vs. Residue number of the selected Chain  Save residue CC of the selected Chain as text  Close