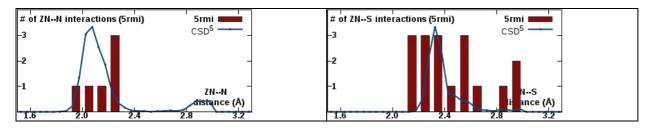
CheckMyMetal(CMM) report for PDB code: 5rmi

PDB title: Pandda analysis group deposition -- crystal structure of sars-cov-2 helicase in complex with z53860899 (2.1Å)

Ю	Res.	Metal	Occupancy	B factor (env.) ¹	Ligands	Valence ²	nVECSUM ³	Geometry ^{1,4}	gRMSD(°) ¹	Vacancy ¹	Bidentate	Alt. metal
A:702	ZN	Zn		74.7 (75.5)	N_1S_3	<u>1.4</u>	0.32	Tetrahedral	4.8°	0	0	Cu, Co
A:703	ZN	Zn		72.6 (77.6)	N_2S_2	1.8	<u>0.22</u>	Tetrahedral	7.1°	0	0	
A:704	ZN	Zn	1	123.4 (106.9)	S_4	1.9	0.32	<u>Trigonal</u> <u>Bipyramidal</u>	7.5°	<u>20%</u>	0	
B:701	ZN	Zn		53.9 (57.2)	N_2S_2	1.9	0.28	Tetrahedral	8.7°	0	0	
B:702	ZN	Zn	1	77.2 (69.7)	S_3	<u>1.6</u>	0.43	Tetrahedral	4.1°	<u>25%</u>	0	Cu, Zn, Co
B:703	ZN	Zn		91.9 (96.2)	N_1S_3	<u>1.4</u>	0.36	Tetrahedral	6.4°	0	0	Cu, Co
	Legend: Not applicable Outlier Borderline Acceptable											

Column	Description							
	Occupancy of ion under consideration							
B factor (env.) ¹	Metal ion B factor, with valence-weighted environmental average B factor in parenthesis							
Ligands	Elemental composition of the coordination sphere							
11/2/Anco	Summation of bond valence values for an ion binding site. <i>Valence</i> accounts for metal-ligand distances							
	Summation of ligand vectors, weighted by bond valence values and normalized by overall valence. Increase when the coordination sphere is not symmetrical due to incompleteness.							
Geometry ^{1,4}	Arrangement of ligands around the ion, as defined by the NEIGHBORHOOD algorithm							
	R.M.S. Deviation of observed geometry angles (L-M-L angles) compared to ideal geometry, in degrees							
Vacancy ¹	Percentage of unoccupied sites in the coordination sphere for the given geometry							
	Number of residues that form a bidentate interaction instead of being considered as multiple ligands							
Alt. metal	A list of alternative metal(s) is proposed in descending order of confidency, assuming metal environment is accurately determined. This feature is still experimental. It requires user discrimination and cannot be blindly accepted							

Metal-ligand distance distributions for pdb5rmi.ent in comparison with CSD



- (1) Zheng H, Chordia MD, Cooper DR, Chruszcz M, Müller P, Sheldrick GM, Minor W (2014) *Nature Protocols*, 9(1), 156-70.
- (2) Brown ID (2009) Chem. Rev., 109, 6858-6919.
- (3) Müller P, Köpke S, Sheldrick GM (2003) Acta Crystallogr. D Biol. Crystallogr., 59, 32-37.
- (4) Kuppuraj G, Dudev M, Lim C (2009) J. Phys. Chem. B, 113, 2952-2960.
- (5) CSD: Cambridge Structural Database
- Maintained by: Heping Zheng <<u>dust@iwonka.med.virginia.edu</u>>

Citing CheckMyMetal (CMM):

Validation of metal-binding sites in macromolecular structures with the CheckMyMetal web server. Zheng, H., Chordia, M.D., Cooper, D.R., Chruszcz, M., Müller, P., Sheldrick, G.M., Minor, W. (2014) Nature Protocols, 9(1), 156-70.