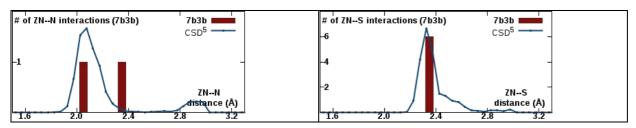
## **CheckMyMetal(CMM)** report for PDB code: 7b3b

PDB title: Structure of elongating sars-cov-2 rna-dependent rna polymerase with remdesivir at position -3 (structure 1) (3.1Å)

ID	Res.	Metal	Occupancy	B factor (env.) <sup>1</sup>	Ligands	Valence <sup>2</sup>	nVECSUM <sup>3</sup>	Geometry <sup>1,4</sup>	gRMSD(°) <sup>1</sup>	Vacancy <sup>1</sup>	Bidentate	Alt. metal
A:2000	ZN	Zn	1	N/A	$N_1S_3$	2	0.11	Tetrahedral	8.9°	0	0	
A:2001	ZN	Zn	1	N/A	$N_1S_3$	1.8	0.17	Tetrahedral	<u>21.1°</u>	0	0	
	Le	gend:	Not applicable <b>Outlier <i>Borderline</i></b> Acceptable									

Column	Description				
Occupancy	Occupancy of ion under consideration				
B factor (env.) <sup>1</sup>	Metal ion B factor, with valence-weighted environmental average B factor in parenthesis				
Ligands	Elemental composition of the coordination sphere				
Valence <sup>2</sup>	Summation of bond valence values for an ion binding site. <i>Valence</i> accounts for metal-ligand distances				
nVECSUM <sup>3</sup>	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 <sup>1,4</sup>	Arrangement of ligands around the ion, as defined by the NEIGHBORHOOD algorithm				
gRMSD(°) <sup>1</sup>	R.M.S. Deviation of observed geometry angles (L-M-L angles) compared to ideal geometry, in degrees				
Vacancy <sup>1</sup>	Percentage of unoccupied sites in the coordination sphere for the given geometry				
Bidentate	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 pdb7b3b.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 < dust@iwonka.med.virginia.edu >

## 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.