

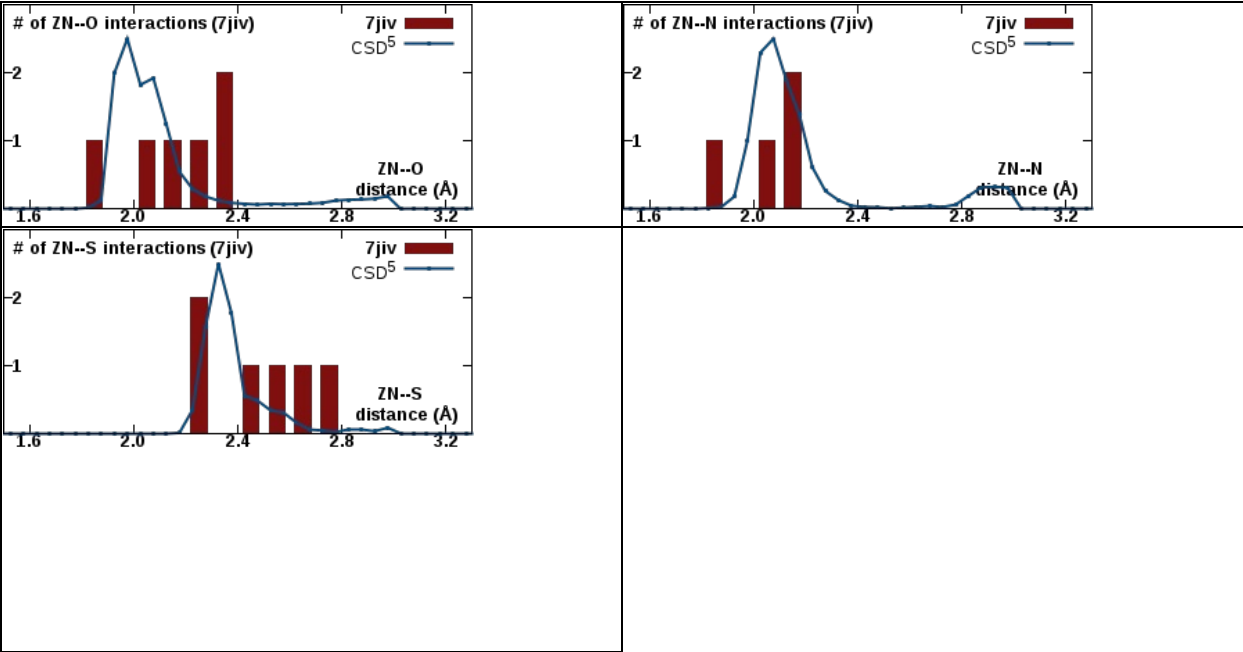
CheckMyMetal(CMM) report for PDB code: 7jiv

PDB title: The crystal structure of papain-like protease of sars cov-2 , c111s mutant, in complex with plp\_snyder530 (2.0Å)

Warning: Valence and nVECSUM parameters should be interpreted with great care due to the presence of multi-nuclear metal clusters around A:504												
Warning: Coordinating ligands by symmetry operation are labeled with prefix 'sym-'												
Warning: Partial occupancy of the metal is not adjusted upon symmetry operation												
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:502	ZN	Zn	1	100.2 (97.2)	S <sub>4</sub>	1.4	0.33	Tetrahedral	16.7°	0	0	Cu, Co
A:503	ZN	Zn	1	104.7 (85.2)	O <sub>1</sub> N <sub>1</sub>	1	0.67	Linear	33°	0	0	
A:504	ZN	Zn	0.5	39.8 (49.7)	O <sub>2</sub> N <sub>1</sub> S <sub>1</sub>	1.3	0.092	Tetrahedral	12.7°	0	0	
A:504	ZN	Zn	0.5	53.6 (52.8)	O <sub>2</sub> N <sub>1</sub> S <sub>1</sub>	1.4	0.29	Tetrahedral	11.5°	0	0	Cu
A:505	ZN	Zn	1	55.3 (56)	O <sub>1</sub> N <sub>1</sub>	1.9	0.11	Trigonal Planar	0.9°	33%	0	Cu
A:507	CL	Cl	0.7	77.9 (66.7)		N/A	N/A	Free	N/A	N/A	N/A	
A:508	CL	Cl	1	57 (70.4)		N/A	N/A	Free	N/A	N/A	N/A	
A:509	CL	Cl	1	59.1 (59)		N/A	N/A	Free	N/A	N/A	N/A	
Legend:			Not applicable	Outlier	Borderline	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. Valence 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 pdb7jiv.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.