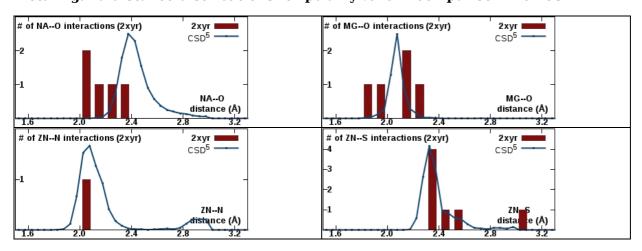
CheckMyMetal(CMM) report for PDB code: 2xyr

PDB title: Crystal structure of the nsp16 nsp10 sars coronavirus complex (2.5Å)

ID	Res.	Metal	Occupancy	B factor (env.) ¹	Ligands	Valence ²	nVECSUM ³	Geometry ^{1,4}	gRMSD(°) ¹	Vacancy ¹	Bidentate	Alt. metal
A:1293	NA	Na	1	<u>42.9</u> (59.5)	O ₅	1.9	0.056	<u>Trigonal</u> <u>Bipyramidal</u>	<u>14.3°</u>	0	0	
A:1294	CL	Cl	1	55.3 (48.6)		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
A:1295	CL	Cl	1	58 (53.6)		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
A:1296	CL	Cl	1	61.2 (42.5)		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
A:1297	CL	Cl	1	<u>55.7</u> (46.6)		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
A:1299	MG	Mg	<u>0.5</u>	39.2 (39.5)	O ₅	1.7	0.14	<u>Trigonal</u> <u>Bipyramidal</u>	13.2°	0	0	Со
B:1132	ZN	Zn	1	51.3 (52.5)	N_1S_3	1.8	0.061	Tetrahedral	4.9°	0	0	
B:1133	ZN	Zn	1	75.4 (76.3)	S_4	<u>1.3</u>	<u>0.17</u>	Tetrahedral	<u>19.8°</u>	0	0	Cu
B:1134	CL	Cl	1	56.6 (63)		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
	Le	gend:	Not applicable Outlier Borderline Acceptable									

Column	Description							
Occupancy	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							
Valence ²	Summation of bond valence values for an ion binding site. <i>Valence</i> accounts for metal-ligand distances							
nVECSUM ³	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							
gRMSD(°) ¹	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							
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 pdb2xyr.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

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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.