PDB title: Sars-cov-2-nsp1-40s complex, composite map (2.8Å)

Warning: Valence and nVECSUM parameters should be interpreted with great care due to the presence of multi-nuclear metal clusters around 2:2004 2:2012 2:2031 2:2043 2:2065 2:2071 2:2090 2:2106 2:2145 2:2148

	2:2090 2:2106 2:2145 2:2148											
ID	Res.	Metal	Occupancy	B factor (env.) <sup>1</sup>	Ligands	Valence <sup>2</sup>	nVECSUM³	Geometry <sup>1,4</sup>	gRMSD(°) <sup>1</sup>	Vacancy <sup>1</sup>	Bidentate	Alt. metal
2:2001	MG	Mg	1	N/A	<u>O</u> 2	<u>0.7</u>	0.33	<u>Trigonal</u> <u>Planar</u>	<u>23.8°</u>	<u>33%</u>	0	
2:2002	MG	Mg	1	N/A	<u>0</u> 1	<u>0.09</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2003	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2004	MG	Mg	1	N/A	<u>O</u> 1	<u>0.11</u>	<u>1</u>	Poorly Coordinated	N/A	N/A	0	
2:2005		Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2006	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2007	MG	Mg	1	N/A	<u>0</u> 1	<u>0.2</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2008	MG	Mg	1	N/A	<u>O_1</u>	0.3	0.98	Poorly Coordinated	N/A	N/A	0	
2:2009	MG	Mg	1	N/A	<u>0</u> 1	<u>0.16</u>	1	Coordinated	N/A	N/A	0	
2:2010	MG	Mg	1	N/A	<u>O_2</u>	0.2	0.72	<u>Square</u> <u>Planar</u>	2.3°	<u>50%</u>	0	
2:2011	MG	Mg	1	N/A	<u>0</u> 1	<u>0.2</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2012		Mg	1	N/A	<u>O</u> 2	<u>0.3</u>	0.71	<u>Square</u> <u>Planar</u>	5.2°		0	K
2:2013		Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2014	MG	Mg	1	N/A	<u>O</u> 2	<u>0.3</u>	<u>0.65</u>	<u>Tetrahedral</u>	5.7°	<u>50%</u>	0	K
2:2015	MG	Mg	1	N/A	<u>N</u> 1	<u>0.5</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2016	MG	Mg	1	N/A	$O_3$	<u>0.5</u>	<u>0.67</u>	Octahedral	12.2°	<u>50%</u>	0	
2:2017	MG	Mg	1	N/A	O <sub>2</sub>	<u>0.7</u>	0.078	<u>Linear</u>	5.5°	0	0	
2:2018	MG	Mg	1	N/A	<u>o</u> 1	0.16	1	Poorly Coordinated	N/A	N/A	0	
2:2019	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2020	MG	Mg	1	N/A	<u>O_1</u>	<u>0.4</u>	0.99	Poorly Coordinated	N/A	N/A	0	
2:2021	MG	Mg	1	N/A	O <sub>2</sub>	0.6	<u>0.47</u>	<u>Trigonal</u> <u>Planar</u>	<u>16.9°</u>	<u>33%</u>	0	
2:2022	MG	Mg	1	N/A	<u>o</u> 1	0.3	1	Poorly Coordinated	N/A	N/A	0	
2:2023	MG	Mg	1	N/A	<u>o</u> 1	0.09	1	Poorly Coordinated	N/A	N/A	0	
2:2024		Mg	1	N/A		N/A	N/A	<u>Free</u>		N/A	N/A	
2:2025	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2026	MG	Mg	1	N/A	<u>0</u> 1	<u>0.17</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2027	MG	Mg	1	N/A	<u>o</u> <sub>1</sub>	0.09	1	Poorly Coordinated	N/A	N/A	0	
2:2029	MG	Mg	1	N/A	<u>O</u> 1	0.3	1	Poorly Coordinated	N/A	N/A	0	
2:2030	MG	Mg	1	N/A	<u>o</u> 1	<u>0.3</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2031	MG	Mg	1	N/A	<u>O</u> 2	<u>0.7</u>	0.37	Trigonal Planar	<u>18.5°</u>	<u>33%</u>	0	
2:2032	MG	Mg	1	N/A	<u>o</u> 1	0.09	1	Poorly Coordinated	N/A	N/A	0	
2:2033	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2034	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2035		Mg	1		<u>o</u> 1	0.4	1	Poorly Coordinated			0	
2:2036		Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2037	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	

2:2038	MC	Mg	1	N/A	$\underline{\mathbf{o}}_{1}$	0.14	1	Poorly	N/A	N/A	0	
								<u>Coordinated</u>				
2:2039	MG	Mg	1	N/A		N/A	N/A	Free Dearly	N/A	N/A	N/A	
2:2040	MG	Mg	1	N/A	<u>O</u> 1	<u>0.18</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2041	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2042	MG	Mg	1	N/A	<u>o</u> 1	0.09	1	Poorly Coordinated	N/A	N/A	0	
2:2043	MG	Mg	1	N/A	<u>O</u> 2	<u>0.3</u>	<u>0.74</u>	<u>Square</u> <u>Planar</u>	1.1°	<u>50%</u>	0	K
2:2044	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2045	MG	Mg	1	N/A	<u>o</u> 1	<u>0.16</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2046	MG	Mg	1	N/A	<u>O</u> 1	<u>0.1</u>	1	<u>Poorly</u> <u>Coordinated</u>	N/A		0	
2:2047		Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2048		Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2049		Mg	1	N/A		N/A	N/A	Free	N/A	N/A	N/A	
2:2050		Mg	1	N/A		N/A	N/A	Free	N/A	N/A	N/A	
2:2051 2:2052	MG	Mg Mg	1	N/A N/A		N/A N/A	N/A N/A	Free Free	N/A N/A	N/A N/A	N/A N/A	
2:2052		Mg	1	N/A		N/A	N/A	Free Free	N/A	N/A	N/A	
2:2053		Mg	1	N/A N/A		N/A	N/A	Free Free	N/A	N/A	N/A	
		Mg	1					Poorly	IN/A			
2:2055	MG	Mg	1		<u>O</u> 1	0.09	1	Coordinated Trigonal	N/A	N/A	0	
2:2056	MG	Mg	1	N/A	O <sub>2</sub>	0.5	0.5	Planar	<u>18°</u>	33%	0	
2:2057		Mg	1		<u>O</u> 1	0.2	1	Poorly Coordinated	N/A	N/A	0	
2:2058	MG	Mg	1	N/A		N/A	N/A	Free	N/A	N/A	N/A	
2:2060	MG	Mg	1	N/A	<u>O</u> 1	0.3	1	Poorly Coordinated	N/A	N/A	0	
2:2061	MG	Mg	1	N/A	<u>O</u> 1	0.13	1	Poorly Coordinated	N/A	N/A	0	
2:2062		Mg	1		<u>O</u> 1	0.11	1	Poorly Coordinated	N/A	N/A	0	
2:2063	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2065	MG	Mg	1	N/A	<u>O</u> 2	<u>0.5</u>	0.84	Square Planar Poorly	13.3°		0	
2:2066		Mg	1	N/A	_	0.3	0.99	Coordinated Square	N/A		0	
2:2067		Mg	1	N/A	_	0.2	0.82	<u>Planar</u>	<u>16.9°</u>		0	
2:2068		Mg	1		<u>O</u> 1	0.2	1	Coordinated Poorly	N/A		0	
2:2069		Mg	1		_	<u>0.5</u>	1	Coordinated Poorly	N/A		0	
2:2070		Mg	1		=	0.11	1	Coordinated Square	N/A		0	
2:2071		Mg	1	N/A	_	0.4	0.8	Planar Poorly	<u>14.6°</u>		0	K
2:2072	MG	Mg	1	N/A	<u>O</u> 1	0.3	1	Coordinated	N/A	N/A	0	
2:2073	MG	Mg	1	N/A	<u>O_2</u>	0.4	0.78	<u>Square</u> <u>Planar</u>	8.8°	<u>50%</u>	0	K
2:2074	MG	Mg	1	N/A	O <sub>3</sub>	<u>0.7</u>	0.37	<u>Square</u> <u>Planar</u>	12.6°	<u>25%</u>	0	Na
2:2075		Mg	1	N/A	<u>O</u> 2	<u>0.4</u>	<u>0.79</u>	<u>Square</u> <u>Planar</u>	13.4°		0	
2:2076	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2077	MG	Mg	1	N/A	<u>o<sub>1</sub></u>	<u>0.12</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2078	MG	Mg	1	N/A	<u>O</u> 2	<u>0.5</u>	0.78	<u>Square</u> <u>Planar</u>	10°	<u>50%</u>	0	
2:2079	MG	Mg	1	N/A	<u>o</u> 1	<u>0.3</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2081	MG	Mg	1	N/A	<u>o</u> 1	0.1	1	Coordinated	N/A	N/A	0	
2 2222	100	3.6	4	37/4	0	0.0		Poorly	27/4	37/4	١	

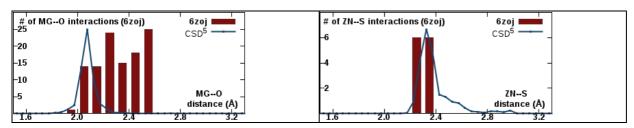
2:2082	MG	Mg	1	N/A	<u>u</u> 1	<u>0.3</u>	1	Coordinated	N/A	N/A	U	
2:2084	MG	Mg	1	N/A	<u>O</u> 1	0.17	1	Poorly	N/A	N/A	0	
2:2085	MG	Mg	1	N/A	-	N/A	N/A	Coordinated Free	N/A	N/A	N/A	
2:2086		J	1		$\underline{\mathbf{o}_1}$	0.2	1	Poorly	N/A	N/A	0	
		3						Coordinated				
2:2087		Mg	1	N/A		N/A	N/A	Free Poorly	N/A	N/A	N/A	
2:2088	MG	Mg	1	N/A	<u>0</u> 1	<u>0.3</u>	<u>1</u>	<b>Coordinated</b>	N/A	N/A	0	
2:2089	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2090	MG	Mg	1	N/A	$O_3$	<u>0.8</u>	0.42	<u>Square</u> <u>Planar</u>	11.7°	<u>25%</u>	0	Na
2:2091	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2092	MG	Mg	1	N/A	<u>0</u> 1	<u>0.18</u>	1	<u>Poorly</u> <u>Coordinated</u>	N/A	N/A	0	
2:2093	MG	Mg	1	N/A	<u>o</u> 1	0.08	1	Poorly Coordinated	N/A	N/A	0	
2:2094	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2095	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2096	MG	Mg	1	N/A	<u>O</u> 1	<u>0.2</u>	<u>1</u>	<u>Poorly</u> <u>Coordinated</u>	N/A	N/A	0	
2:2097	MG	Mg	1	N/A	<u>O</u> 1	0.11	1	Poorly Coordinated	N/A	N/A	0	
2:2098	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2099	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2100	MG	Mg	1	N/A	<u>O</u> 1	<u>0.12</u>	<u>1</u>	<u>Poorly</u> <u>Coordinated</u>	N/A	N/A	0	
2:2101	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2102	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2103	MG	Mg	1	N/A	<u>o<sub>1</sub></u>	0.08	1	Poorly Coordinated	N/A	N/A	0	
2:2105	MG	Mg	1	N/A	<u>O</u> 1	<u>0.1</u>	<u>1</u>	Poorly Coordinated	N/A	N/A	0	
2:2106	MG	Mg	1	N/A	<u>O</u> 1	<u>0.2</u>	0.98	Poorly Coordinated	N/A	N/A	0	
2:2107	MG	Mg	1	N/A	<u>0</u> 1	<u>0.19</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2108	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2109		Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2110		Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2111	MG	Mg	1	N/A		N/A	N/A	Free December	N/A	N/A	N/A	
2:2112	MG	Mg	1	N/A	<u>0</u> 1	<u>0.11</u>	<u>1</u>	Poorly Coordinated	N/A	N/A	0	
2:2113	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2114	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2115	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2116	MG	Mg	1	N/A	<u>O</u> 1	<u>0.4</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2118	MG	Mg	1	N/A	<u>N</u> <sub>1</sub>	0.17	1	Poorly Coordinated	N/A	N/A	0	
2:2119	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2120	MG	Mg	1	N/A	<u>o</u> 1	<u>0.2</u>	<u>1</u>	<u>Poorly</u> <u>Coordinated</u>	N/A	N/A	0	
2:2121		Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2122		Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2123		Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2124		Mg	1	N/A		N/A	N/A	Free	N/A	N/A	N/A	
2:2125		Mg	1	N/A		N/A	N/A	Free Free	N/A	N/A	N/A	
2:2126 2:2127		Mg Mg	1	N/A N/A	<u>O</u> 2	N/A 0.18	N/A 0.75	Free Square	N/A 7°	N/A <b>50%</b>	N/A 0	
2:2128		J	1			0.17	1	Planar Poorly	N/A		0	
			1					<u>Coordinated</u>		•		
2:2129 2:2130		Mg Mg	1	N/A N/A		N/A N/A	N/A N/A	Free Free	N/A N/A	N/A N/A	N/A N/A	
2.2130	1.10	1.19	-	Z V/ZZ		- 1/-T	- 1/1 i	<u> </u>	+ V/ 1 1	11/11	1 1/2 1	

2:2131	MG	Mg	1	N/A	<u>O_2</u>	<u>0.3</u>	0.82	<u>Square</u> <u>Planar</u>	<u>15.4°</u>	<u>50%</u>	0	K
2:2132	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2134	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2135	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2136	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2137	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2138	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2139	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2140	MG	Mg	1	N/A	<u>o</u> 1	<u>0.1</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2141	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2142	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2143	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2144	MG	Mg	1	N/A	<u>0</u> 1	0.18	1	Poorly Coordinated	N/A	N/A	0	
2:2145	MG	Mg	1	N/A	<u>O</u> 2	0.4	0.87	<u>Square</u> <u>Planar</u>	<u>17.1°</u>	<u>50%</u>	0	
2:2146	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2147	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2148	MG	Mg	1	N/A	<u>0</u> 1	<u>0.4</u>	1	Poorly Coordinated	N/A	N/A	0	
2:2149	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2150	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2151	MG	Mg	1	N/A	<u>0</u> 1	0.08	1	Poorly Coordinated	N/A	N/A	0	
2:2152	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2153	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2154	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2155	MG	Mg	1	N/A	<u>O</u> 2	<u>0.2</u>	<u>0.65</u>	<u>Tetrahedral</u>	8.6°	<u>50%</u>	0	
2:2156	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2157	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2158	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
2:2159	MG	Mg	1	N/A		N/A	N/A	<u>Free</u>	N/A	N/A	N/A	
G:301	MG	Mg	1	N/A	<u>o<sub>1</sub></u>	0.3	1	Poorly Coordinated	N/A	N/A	0	
O:201	MG	Mg	1	N/A	<u>o</u> 1	0.08	1	Poorly Coordinated	N/A	N/A	0	
O:202	MG	Mg	1	N/A	O <sub>3</sub>	<u>0.7</u>	0.7	Tetrahedral	27.7°	<u>25%</u>	0	Na
		Mg	1	N/A		N/A	N/A			N/A	N/A	
S:202	MG		1	N/A	O <sub>3</sub>	0.4	0.37	Square Planar	<u>13.9°</u>	<u>25%</u>	0	K
T:201	MG	Mg	1	N/A	<u>O</u> 2	0.3	0.74	Square Planar	5.2°	<u>50%</u>	0	K
X:2301	MG	Mg	1	N/A		N/A	N/A	Free	N/A	N/A	N/A	
	1	Zn	1	N/A	$S_4$	2.3	0.033			0	0	
d:201	1	Zn	1	N/A	S <sub>4</sub>	2.3	0.081			0	0	
f:500		Zn	1	N/A	S <sub>4</sub>	2.1	0.08			0	0	
1	LIN	Z11	1	11/11	$_{4}$	2.1	0.00	retraileural	1./	U	U	

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							
$I\alpha RMSD(0)1$	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							
	A list of alternative metal(s) is proposed in descending order of confidency, assuming metal							

Alt. metal environment is accurately determined. This feature is still experimental. It requires user discrimination and cannot be blindly accepted

## Metal-ligand distance distributions for pdb6zoj.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.