

SI

Substitution of Ca^{2+} and changes in the H-bond network near the oxygen-evolving complex of photosystem II

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Table S1. Functional/basis set and calculated distances in the open-cubane S₂ conformation in Å. For efficient analysis, a smaller QM region was adopted (i.e., the Mn₄CaO₅ cluster; the ligand side-chains of D1-Asp170, D1-Glu189, D1-His332, D1-Glu333, D1-Asp342, and CP43-Glu354; ligand carboxy-terminal group of D1-Ala344; and ligand water molecules, W1–W4; and CP43-Arg357): thus, the calculated distances may not be consistent with those calculated using the large QM region (e.g., Figure 1b).

functional	B3LYP						M05
basis set	LACVP*	LACV3P	LACV3P*	LACV3P**	LACV3P***	LACV3P****	LACVP*
Mn1-Mn2	2.77	2.84	2.80	2.80	2.80	2.80	2.77
Mn1-Mn3	3.18	3.19	3.18	3.18	3.18	3.18	3.18
Mn1-Mn4	4.67	4.68	4.66	4.65	4.66	4.66	4.66
Mn2-Mn3	2.74	2.80	2.76	2.76	2.77	2.77	2.74
Mn3-Mn4	2.72	2.80	2.75	2.75	2.75	2.75	2.72
Mn1-O1	1.83	1.86	1.84	1.84	1.85	1.85	1.82
Mn1-O3	1.90	1.92	1.91	1.91	1.91	1.91	1.90
Mn1-O5	2.71	2.67	2.67	2.68	2.68	2.68	2.71
Mn1-N _{D1-His332}	2.01	1.98	2.00	2.00	2.00	2.00	2.01
Mn1-O _{D1-Asp342}	2.18	2.20	2.20	2.20	2.20	2.20	2.18
Mn1-O _{D1-Glu189}	1.90	1.90	1.91	1.91	1.91	1.91	1.91
Mn2-O1	1.81	1.84	1.83	1.83	1.83	1.83	1.82
Mn2-O2	1.81	1.84	1.82	1.82	1.82	1.82	1.81
Mn2-O3	1.84	1.89	1.87	1.87	1.87	1.87	1.84
Mn2-O _{D1-Asp342}	2.01	1.97	1.99	1.99	1.99	1.99	2.01
Mn2-O _{D1-Ala344}	1.90	1.88	1.90	1.90	1.90	1.90	1.91
Mn2-O _{CP43-Glu354}	2.00	1.98	2.00	1.99	2.00	2.00	2.00
Mn3-O2	1.84	1.88	1.86	1.86	1.86	1.86	1.84
Mn3-O3	1.91	1.92	1.92	1.92	1.92	1.92	1.91
Mn3-O4	1.80	1.82	1.81	1.81	1.81	1.81	1.80
Mn3-O5	1.82	1.87	1.84	1.84	1.84	1.84	1.83
Mn3-O _{CP43-Glu354}	2.06	2.03	2.05	2.05	2.05	2.05	2.08
Mn3-O _{D1-Glu333}	1.99	1.97	1.99	1.99	1.99	1.99	1.99
Mn4-O4	1.77	1.81	1.79	1.79	1.79	1.79	1.77
Mn4-O5	1.97	2.03	1.99	1.99	1.99	1.99	1.97
Mn4-W1	2.17	2.11	2.15	2.17	2.18	2.18	2.19
Mn4-W2	2.05	2.00	2.03	2.04	2.04	2.04	2.06
Mn4-O _{D1-Asp170}	2.16	2.12	2.15	2.14	2.15	2.15	2.17

Mn4-O _{D1-Glu333}	2.08	2.10	2.10	2.09	2.09	2.09	2.08
Ca-O5	2.50	2.53	2.52	2.51	2.52	2.52	2.51
Ca-W3	2.42	2.38	2.41	2.43	2.44	2.44	2.42
Ca-W4	2.42	2.38	2.41	2.42	2.45	2.45	2.43
Ca-O _{D1-Glu189}	3.13	3.17	3.17	3.18	3.17	3.17	3.13

Table S2. Distances in the open-cubane Mn_4MO_5 clusters in S_2 ($\text{M} = \text{Mg}^{2+}$, Ca^{2+} , Sr^{2+} , and Ba^{2+}) in Å using M06 functional (i.e., dispersion correction).

	Mg²⁺	Ca²⁺	Sr²⁺	Ba²⁺	Ba²⁺ + water
ionic radius ^a	0.66	0.99	1.12	1.34	1.34
surface area ^b	0.44	1	1.28	1.83	1.83
W3... M	2.08	2.41	2.56	2.66	2.71
W4... M	2.12	2.39	2.55	2.69	2.75
O1... M	2.16	2.38	2.50	2.65	2.66
O2... M	2.25	2.56	2.68	2.81	2.77
O5... M	2.89	2.59	2.67	2.78	2.80
O5...Mn1	2.87	3.03	3.06	3.11	3.11
O5...Mn4	1.82	1.83	1.82	1.81	1.82

Table S3. Contributions of the Mn_4MO_5 components to the HOMO of S_2 (%). The component that has the largest contribution to the HOMO is in bold. To specifically focus on the difference in the electronic state of the Mn_4MO_5 cluster, the QM region was defined as Mn_4MO_5 ; side chains of D1-Asp170, D1-Glu189, D1-His332, D1-Glu333, D1-Asp342, and CP43-Glu354; carboxyl terminal group of D1-Ala344; and water molecules, W1–W4.

HOMO	contribution from								
	Mn1 ^a	Mn2	Mn3	Mn4	O1	O2	O3	O4	O5
S_2									
Mg ²⁺ -PSII	86	0	0	0	3	2	3	0	0
Ca ²⁺ -PSII	83	0	0	0	4	2	4	0	0
Sr ²⁺ -PSII	83	1	1	0	4	2	5	0	0
Ba ²⁺ -PSII	82	1	1	0	4	2	5	0	0

^a Including ligand residues.