

Supporting information for:

B(OH)₄⁻ Hydration and Association in Sodium Metaborate Solutions by X-Ray Diffraction and Empirical Potential Structure Refinement

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Fig. S6 A schematic view of distances related to B(OH)_4^- hydration and B(OH)_4^- - Na^+ association in aqueous sodium metaborate solutions.

Table S1 Bond lengths and angles used to describe the molecules B(OH)₄⁻ and H₂O

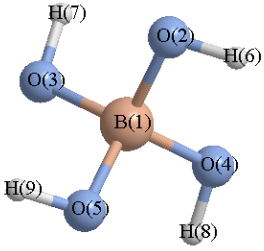

	Bonds/Å			Angle/°				Dihedral angle/°				
	atom 1	atom 2	distance	atom 1	atom 2	atom 3	Angle	atom 1	atom 2	atom 3	atom 4	Angle
 <p>B(OH)₄⁻</p> <p>The structure was based on the crystal structure and optimized with Gaussian at B3LYP/Aug-cc-pvdz</p>	1	2	1.48965	2	1	3	106.93602	3	1	2	6	-162.13789
	1	3	1.48965	2	1	4	106.95973	4	1	2	6	-38.86043
	1	4	1.48889	2	1	5	114.61903	5	1	2	6	79.52378
	1	5	1.48943	3	1	4	114.65141	2	1	3	7	38.9278
	2	6	0.96238	3	1	5	106.9454	4	1	3	7	-79.44763
	3	7	0.96177	4	1	5	106.96913	5	1	3	7	162.15872
	4	8	0.96238	1	2	6	106.74967	2	1	4	8	162.13585
	5	9	0.96268	1	3	7	106.77118	3	1	4	8	-79.50208
				1	4	8	106.76966	5	1	4	8	38.87816
				1	5	9	106.80371	2	1	5	9	79.46024
								3	1	5	9	-38.87276
								4	1	5	9	-162.16083
								3	1	2	6	-162.13789
 <p>H₂O</p>	1	2	0.9650	1	2	3	109.450					
	2	3	0.9650									

Table S2 EPSR simulation boxes setup details.

simulation name	# water molecules	# Na ⁺ cations	# anions	Density /atoms Å ⁻³	Box side length /Å	Iterations times
Water	1000	-	-	0.099987	31.074	10000
B1N1	1000	18	18	0.10315	31.353	11000
B1N2	1000	55	55	0.10735	32.100	12000
B1N3	1000	125	125	0.11128	33.677	13200

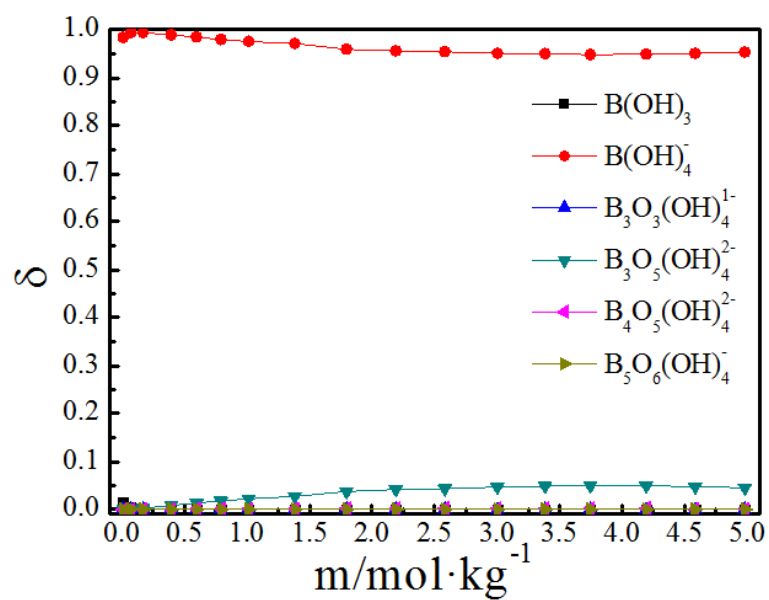


Fig. S1. The distribution of boron species with salt concentration in aqueous NaB(OH)₄ solutions at 298.15 K

■, H₃BO₃; ●, B(OH)₄⁻; ▲, [B₃O₃(OH)₄]⁻; ▼, B₃O₅(OH)₄²⁻; ★, B₄O₅(OH)₄²⁻; ►, B₅O₆(OH)₄⁻

See: Zhou Y. Fang C. Fang Y., Chin. J. Chem. Eng. 2013, 21 (9): 1048-1056.

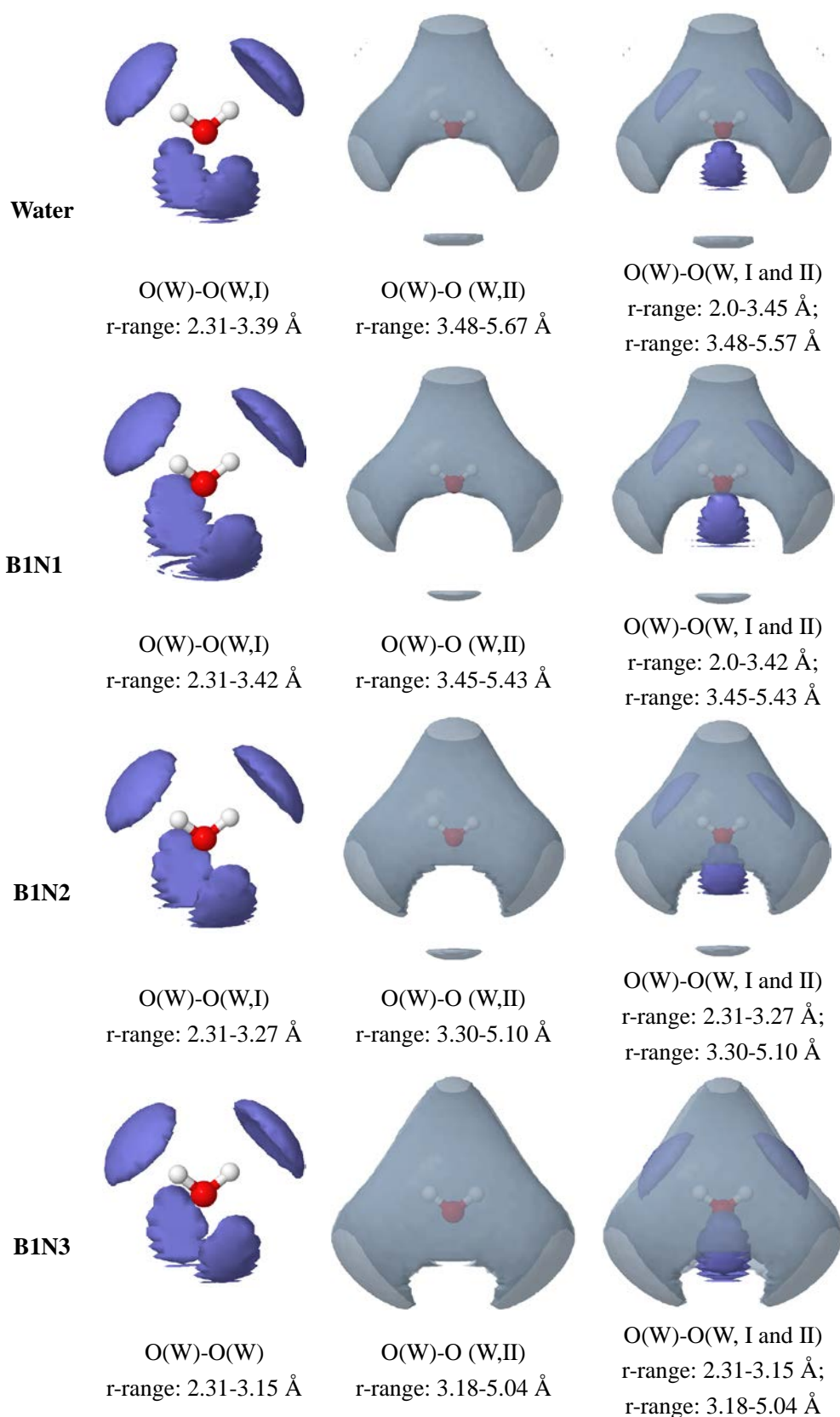


Fig. S2 Spatial density functions of water molecules around a central water molecule in pure water and three aqueous NaB(OH)₄ solutions. The dark blue lobes represent the first sphere and the grayish blue and semitransparent lobes ones represent of the second sphere. The red and white ball in the center represent O and H atom, respectively.

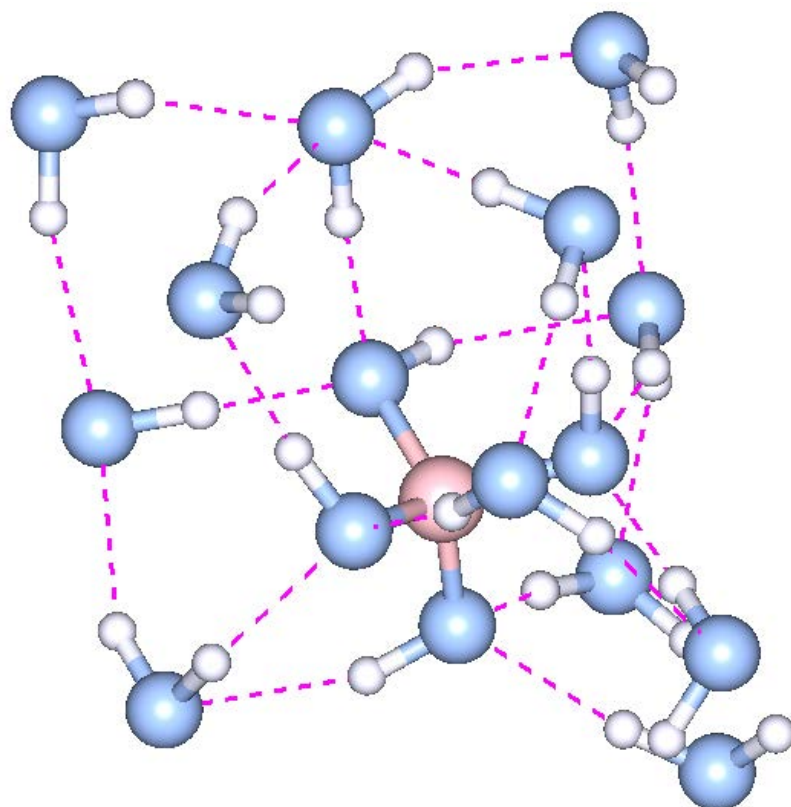


Fig. S3. One of the optimized lowest-energy structures of aqua- B(OH)_4^- [$\text{B(OH)}_4(\text{H}_2\text{O})_{12}]^-$ at B3LYP/aug-cc-pVDZ level.

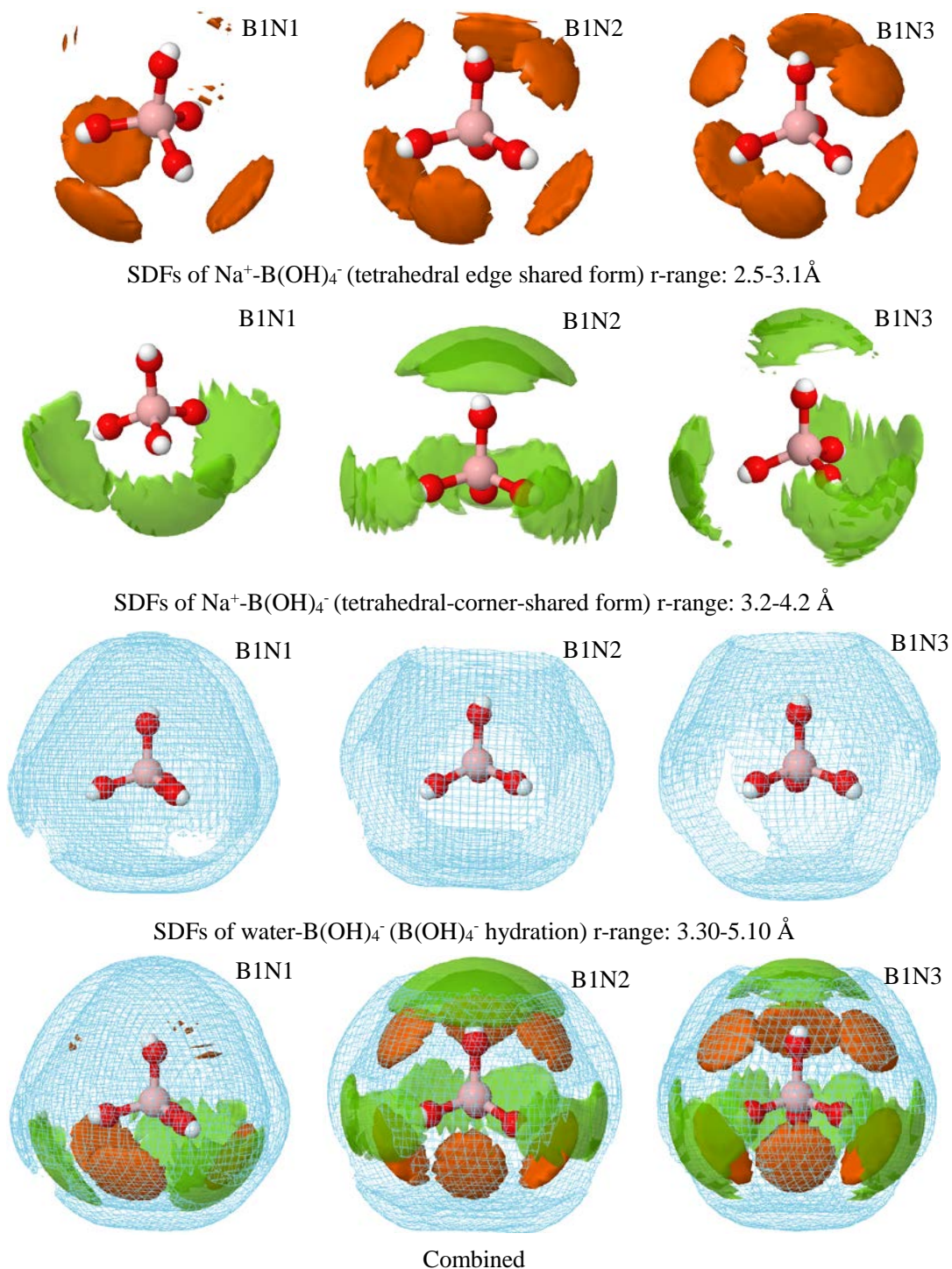


Fig. S4 Spatial density functions for hydrated water molecules and sodium ions to the central B(OH)_4^- . The SDFs surfaces showing the top 25% probability density for correlations of hydrated water molecules to the central B(OH)_4^- within the cutoff distance of 5.0 Å (blue and transparency grid); The top 35% probability density for correlations of Na^+ to the central B(OH)_4^- within the cutoff distance of the local minimal of $g_{\text{Na-B}}(r)$, the tetrahedral-edge-shared bidentate ion pairs (red lobes) and tetrahedral-corner-shared monodentate ion pairs (green and semitransparent lobes). The pink, red

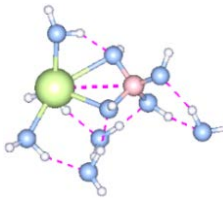
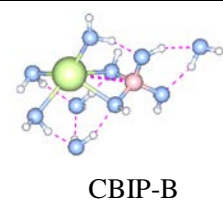
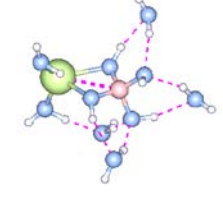
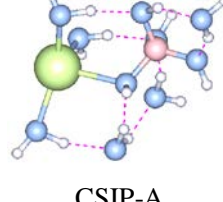
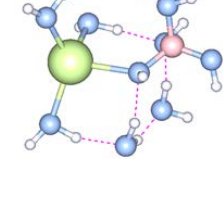
and white balls in the centre represent B, O and H atom of B(OH)_4^- , respectively.

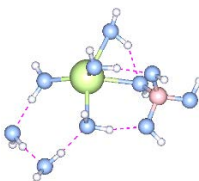
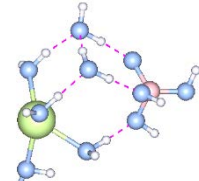
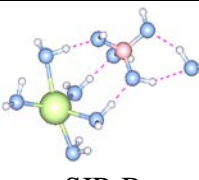
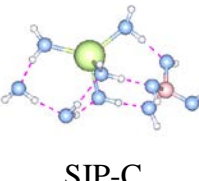
Table S3 Detailed energy parameters of $\text{NaB(OH)}_4(\text{H}_2\text{O})_6$ in the gas and aqueous phase at B3LYP/aug-cc-pVDZ level.

	Clusters	Na^+	B(OH)_4^-	Water	BSSE	Delta	Gscrf	Gss	Gsolv
	Hartree	Hartree	Hartree	Hartree	Hartree	kcal/mol	kcal/mol	kcal/mol	kcal/mol
CBIP-A									
E0	-949.327	-162.092	-328.364	-458.534	0.005	-208.352	-23.750	-0.712	-232.814
E	-949.304	-162.091	-328.358	-458.522	0.005	-205.409	-23.750	-0.712	-229.871
H	-949.303	-162.090	-328.357	-458.521	0.005	-206.628	-23.750	-0.712	-231.090
G	-949.380	-162.107	-328.394	-458.574	0.005	-188.566	-23.750	-0.712	-213.028
CBIP-B									
E0	-949.330	-162.092	-328.363	-458.537	0.005	-208.556	-22.140	-0.712	-231.408
E	-949.306	-162.091	-328.358	-458.524	0.005	-205.705	-22.140	-0.712	-228.558
H	-949.305	-162.090	-328.357	-458.523	0.005	-206.926	-22.140	-0.712	-229.779
G	-949.382	-162.107	-328.392	-458.578	0.005	-188.301	-22.140	-0.712	-211.154
CBIP-C									
E0	-949.318	-162.092	-328.363	-458.529	0.006	-205.201	-25.350	-0.712	-231.263
E	-949.293	-162.091	-328.357	-458.519	0.006	-200.990	-25.350	-0.712	-227.053
H	-949.292	-162.090	-328.356	-458.518	0.006	-202.210	-25.350	-0.712	-228.272
G	-949.373	-162.107	-328.392	-458.570	0.006	-186.681	-25.350	-0.712	-212.743
CSIP-A									
E0	-949.328	-162.092	-328.364	-458.527	0.005	-213.237	-23.770	-0.712	-237.720
E	-949.304	-162.091	-328.357	-458.515	0.005	-210.338	-23.770	-0.712	-234.821
H	-949.303	-162.090	-328.356	-458.515	0.005	-211.558	-23.770	-0.712	-236.040
G	-949.381	-162.107	-328.393	-458.569	0.005	-193.043	-23.770	-0.712	-217.525
CSIP-B									
E0	-949.330	-162.092	-328.363	-458.534	0.005	-210.733	-24.130	-0.712	-235.575
E	-949.306	-162.091	-328.357	-458.522	0.005	-208.272	-24.130	-0.712	-233.114
H	-949.305	-162.090	-328.356	-458.521	0.005	-209.491	-24.130	-0.712	-234.333
G	-949.384	-162.107	-328.393	-458.575	0.005	-190.316	-24.130	-0.712	-215.158
CSIP-C									
E0	-949.327	-162.092	-328.363	-458.544	0.004	-203.099	-24.540	-0.712	-228.351
E	-949.303	-162.091	-328.357	-458.531	0.004	-200.328	-24.540	-0.712	-225.580
H	-949.302	-162.090	-328.356	-458.530	0.004	-201.548	-24.540	-0.712	-226.800
G	-949.382	-162.107	-328.392	-458.585	0.004	-184.097	-24.540	-0.712	-209.349
SIP-A									
E0	-949.323	-162.092	-328.364	-458.520	0.005	-215.029	-24.790	-0.712	-240.532
E	-949.299	-162.091	-328.358	-458.508	0.005	-211.714	-24.790	-0.712	-237.217
H	-949.298	-162.090	-328.357	-458.507	0.005	-212.934	-24.790	-0.712	-238.437
G	-949.379	-162.107	-328.393	-458.562	0.005	-195.160	-24.790	-0.712	-220.662
SIP-B									
E0	-949.322	-162.092	-328.364	-458.520	0.005	-214.034	-24.890	-0.712	-239.636

E	-949.297	-162.091	-328.358	-458.508	0.005	-210.626	-24.890	-0.712	-236.228
H	-949.296	-162.090	-328.357	-458.507	0.005	-211.845	-24.890	-0.712	-237.447
G	-949.376	-162.107	-328.393	-458.562	0.005	-193.737	-24.890	-0.712	-219.340
SIP-C									
E0	-949.329	-162.092	-328.364	-458.539	0.004	-206.652	-25.530	-0.712	-232.895
E	-949.305	-162.091	-328.358	-458.526	0.004	-204.074	-25.530	-0.712	-230.317
H	-949.304	-162.090	-328.357	-458.526	0.004	-205.294	-25.530	-0.712	-231.537
G	-949.382	-162.107	-328.392	-458.581	0.004	-186.490	-25.530	-0.712	-212.732

Table S4 Bond parameters of NaB(OH)₄(H₂O)₆ at B3LYP/ Aug-cc-pVDZ level.

Clusters	Atom pair	Bond length/Å				Averaged/Å
	Na-O(W)	2.297	2.321	2.337		2.318
	Na-O(B)	2.401	2.593		2.497	
	Na-B	3.037				3.037
	B-O(W)	3.572	3.240	3.765		3.526
CBIP-A						
	Na-O(W)	2.375	2.429	2.273		2.359
	Na-O(B)	2.503	2.431		2.467	
	Na-B	2.928				2.928
	B-O(W)	3.335				3.335
CBIP-B						
	Na-O(W)	2.304	2.243		2.273	
	Na-O(B)	2.247	2.427		2.337	
	Na-B	2.847				2.847
	B-O(W)	3.568	3.425	3.317	3.294	3.401
CBIPC						
	Na-O(W)	2.305	2.273	2.269		2.282
	Na-O(B)	2.405				2.405
	Na-B	3.313				3.313
	B-O(W)	3.424	3.844	3.370	3.623	3.905
CSIP-A						
	Na-O(W)	2.257	2.263	2.341		2.287
	Na-O(B)	2.367				2.367
	Na-B	3.448				3.448
	B-O(W)	3.451	3.691	3.807	3.811	3.690
CSIP-B						

	Na-O(W)	2.473	2.385	2.413	2.262		2.383
	Na-O(B)	2.403					2.403
	Na-B	3.349					3.349
	B-O(W)	3.492	3.440	3.878			3.603
CSIP-C							
	Na-O(W)	2.366	2.322	2.251	2.273	2.366	2.303
	Na-O(B)	/					/
	Na-B	4.689					4.689
	B-O(W)	3.5	3.73	3.826			3.685
SIP-A							
	Na-O(W)	2.336	2.453	2.346	2.363	2.425	2.385
	Na-O(B)	/					/
	Na-B	4.14					4.140
	B-O(W)	3.56	3.693	3.321	3.709		3.571
SIP-B							
	Na-O(W)	2.344	2.344	2.273	2.254		2.304
	Na-O(B)	/					/
	Na-B	4.093					4.093
	B-O(W)	3.614	3.663	3.705			3.661
SIP-C							

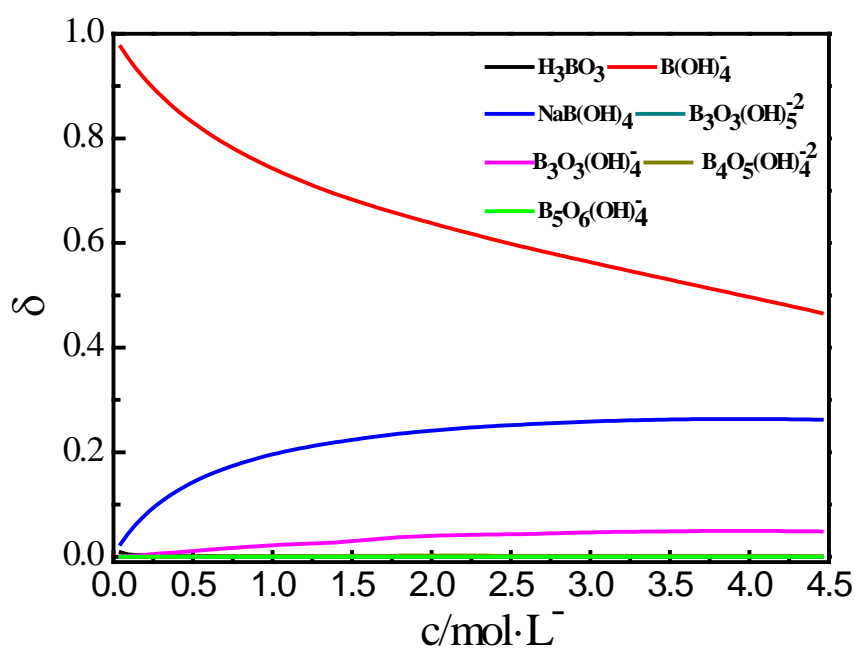


Fig. S5 Speciation of polyborates and ion association (in the apparent form of NaB(OH)_4) vs concentration in aqueous NaB(OH)_4 solutions at 298.15K and 1atm.

(using the ion association constant suggested by Rowe *et al. J. Solution Chem.*, **19**, 149(1990).)

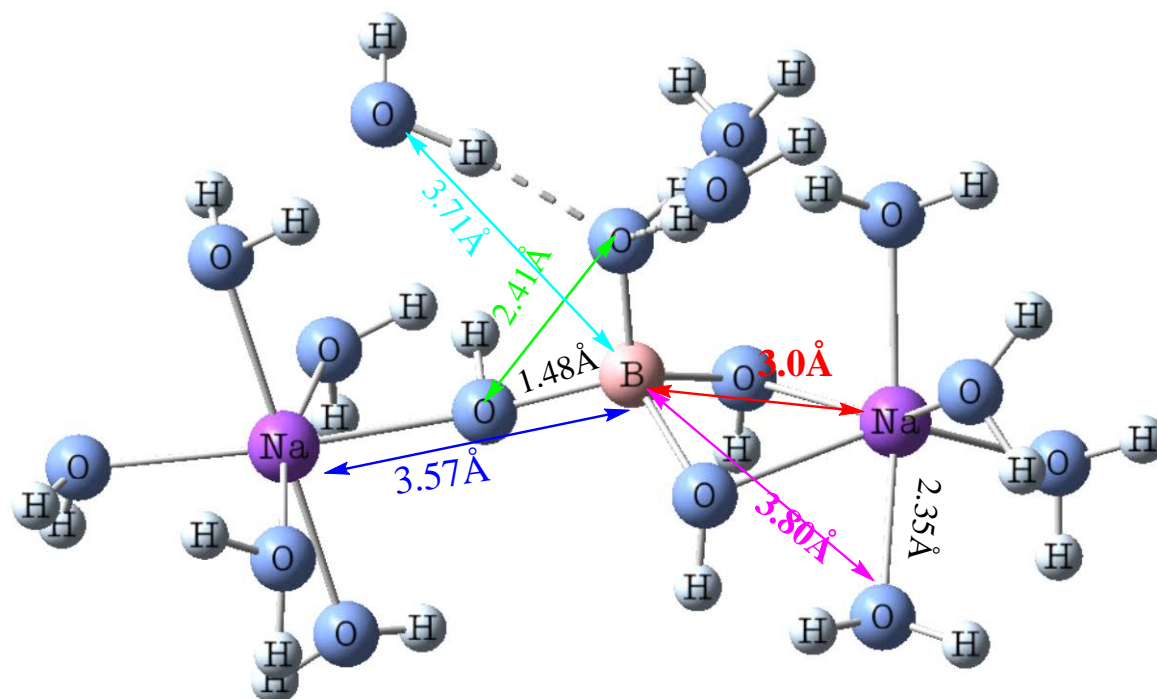


Fig. S6 A schematic view of distances related to B(OH)_4^- hydration and B(OH)_4^- - Na^+ association in aqueous sodium metaborate solutions