

Supporting information for:

**On the brink of self-hydration:
the water heptadecamer**

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1 All $(\text{H}_2\text{O})_{17}$ structures of the main paper

In the main paper, four different classes of $(\text{H}_2\text{O})_{17}$ low-energy structures were shown with one representative structure. Here, all twelve structures are displayed. Their atomic coordinates are provided in separate xyz-files. All of them were locally optimized at the B2PYLP-D4/def2-TZVPPD level, with dense grids and tight convergence criteria.

The “prism” class features two fused pentagonal prisms, with one water dimer attached across a square side face of one of these prisms. As displayed in Fig. S1, the structures p1, p2, p3 and p4 have (qualitatively) the same O-atom cage but partly different hydrogen-bond arrangements. p1 is the global minimum candidate at half of the ab-initio/DFT levels employed here, and also for MB-pol2023.

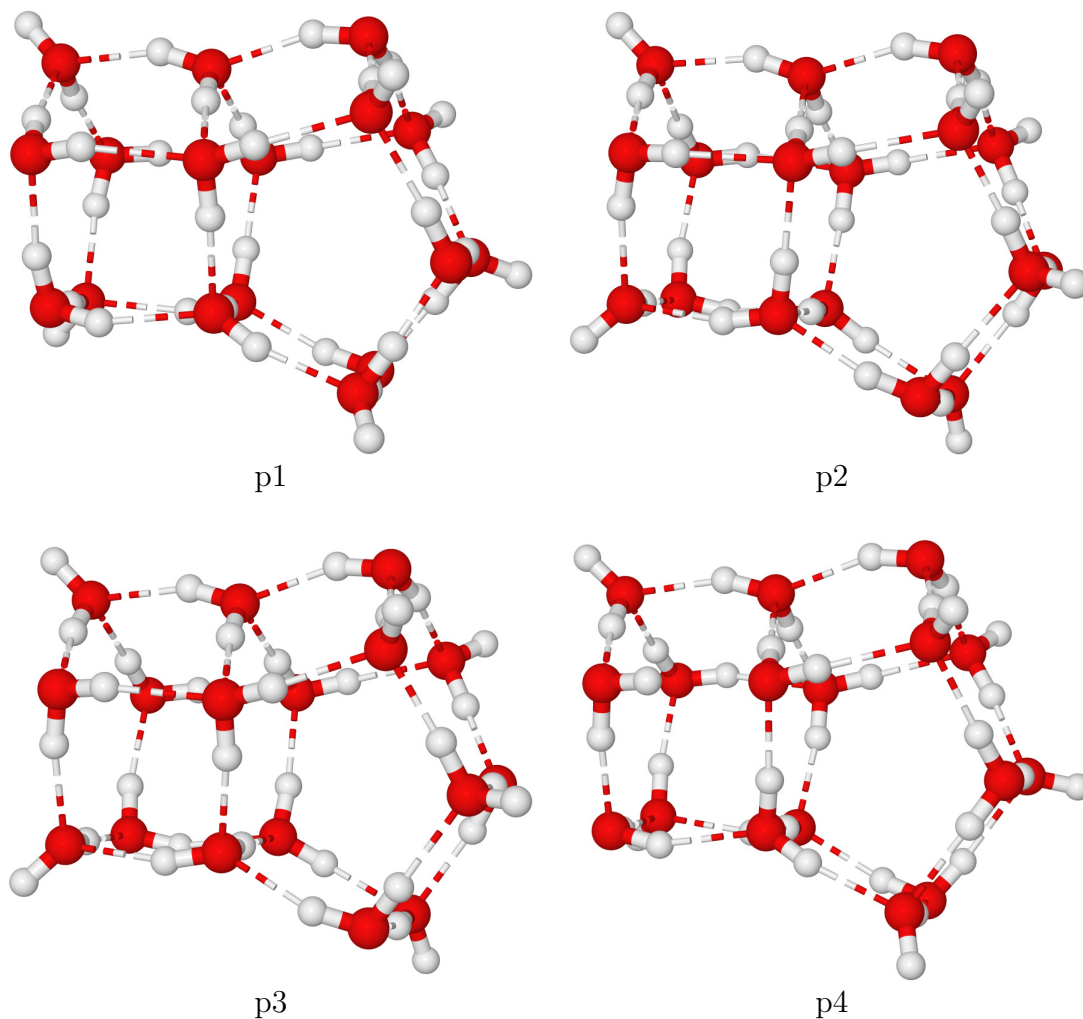


Figure S1: The four p-structures.

In the “twisted-cubes” class, there are two slightly deformed cubes, not along a common axis and also rotated relative to each other. On one side these cubes are joined directly, on the other side via additional bridging water molecules (cf. Fig. S2). One version of this class, t2, is the TIP4P global minimum discovered by Wales and Hodges [1], which coincides with the “L-shape” structure presented by Xantheas et al. [2]. t2 has a rather high energy at the ab-initio/DFT levels used here. However, t1, a slight structural variation of t2 (not just a different H-bond pattern), has a very low energy

at the ab-initio/DFT levels employed here and is the global minimum candidate for q-AQUA-pol.

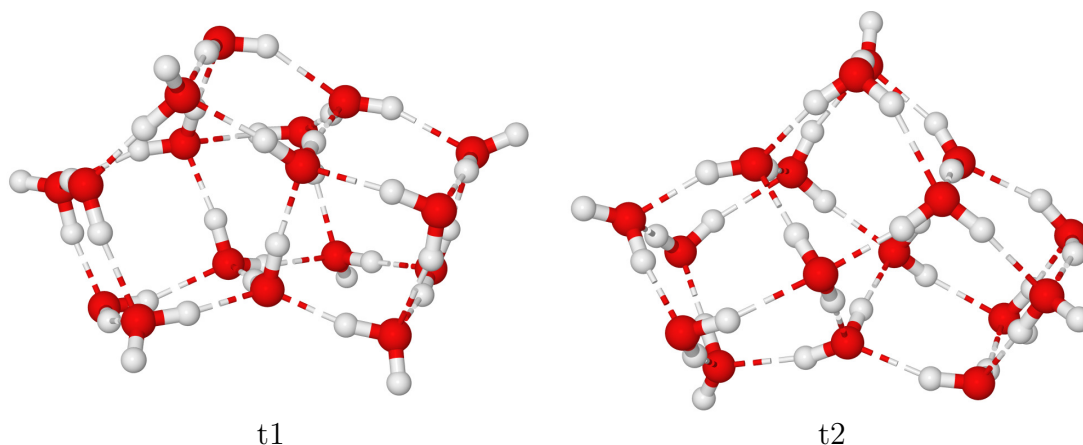


Figure S2: The two t-structures.

The “water-centered” class is the only one with actual water self-hydration, i.e., one water molecule is in or near the cluster center, surrounded by a hydration shell formed by water four-, five- and six-rings. w1 is the global minimum candidate at half of the ab-initio/DFT levels used here, and also for q-AQUA and TTM2-F/TTM3-F. When one of its six-rings is at the bottom (it does not matter which one), the ring sequence around its equator is 5-5-4-4-5-4, capped at the top by 4-5-6. w2, w3 and w4 are different H-bond patterns of a different ring arrangement: When a six-ring is at the bottom again, the ring sequence around the equator is 4-5-4-5-4-5, and the cap consists of three five-rings (cf. Fig. S3).

Finally, in the “cubes” class there are three fused cubes, aligned along a common central axis and not rotated relative to each other. The remaining 17th water molecule can now be added at different locations, as shown in Fig. S4: Adding it to one edge of the middle cube results in c1, which is lower in energy in most cases. Adding it to an end-edge of an end-cube produces c2. Neither c1 nor c2 are among the best structures at the ab-initio/DFT levels employed here, but c1 is the global minimum candidate for MB-pol2016.

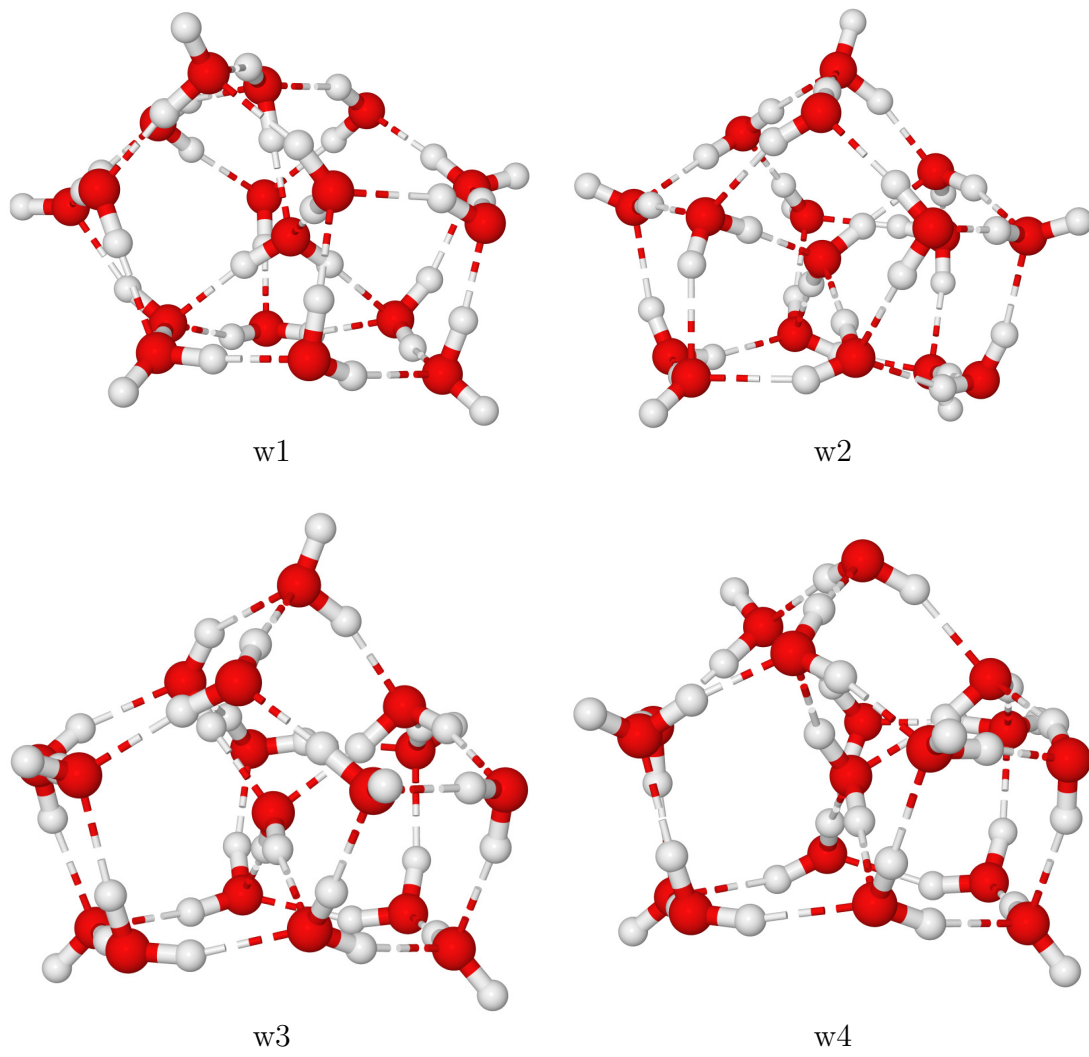


Figure S3: The four w-structures.

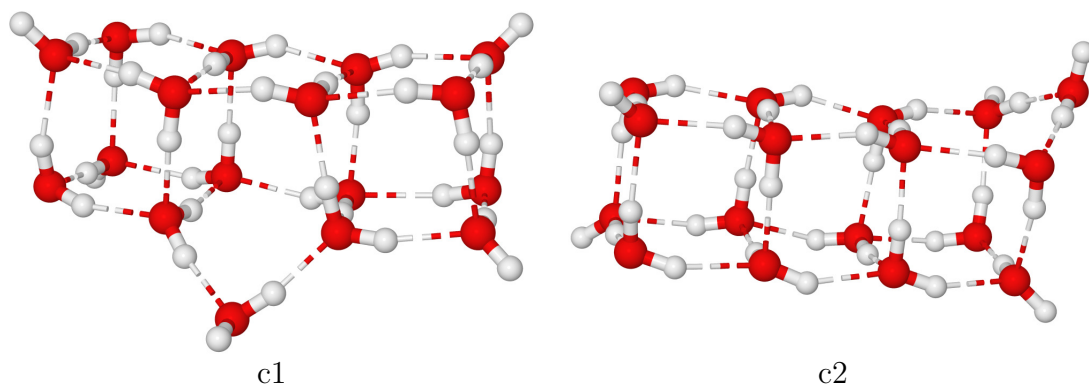


Figure S4: The two c-structures.

2 Tabulated energy values

Complementing the energy figures in the main paper, this section collects all numerical energy values depicted there in tabular form. Each theory level has its own table, for all 12 cluster structures, listing both the “absolute” energy value in Hartree (total electronic energy for the ab-initio/DFT methods, and typically with respect to dissociation into water monomers for the water model potentials) and the relative energy differences between the cluster structures in kJ/mol (relative to the lowest absolute energy, at this level of theory).

Structure labels are the same as in the main paper, and as explained and depicted in section 1 of this Supplementary Information.

e34	DLPNO-CCSD(T)/CBS	
structure	absolute energy / Hartree	relative energy / kJ/mol
p1	-1298.683 307	0.00
p2	-1298.682 909	1.04
p3	-1298.682 367	2.46
p4	-1298.682 237	2.80
t1	-1298.683 041	0.69
t2	-1298.681 062	5.89
w1	-1298.682 739	1.49
w2	-1298.682 154	3.02
w3	-1298.681 289	5.29
w4	-1298.681 285	5.31
c1	-1298.682 373	2.45
c2	-1298.681 623	4.42

ccsdT/f12	DLPNO-CCSD(T)/cc-pVQZ-F12	
structure	absolute energy / Hartree	relative energy / kJ/mol
p1	-1298.607 198	0.23
p2	-1298.606 874	1.08
p3	-1298.606 338	2.49
p4	-1298.606 223	2.79
t1	-1298.607 044	0.63
t2	-1298.605 400	4.95
w1	-1298.607 286	0.00
w2	-1298.606 725	1.47
w3	-1298.605 779	3.95
w4	-1298.605 762	4.00
c1	-1298.606 578	1.85
c2	-1298.605 828	3.82

ccsdT/qz		DLPNO-CCSD(T)/def2-QZVPP	
structure	absolute energy / Hartree	relative energy /	kJ/mol
p1	-1298.420 935	0.00	
p2	-1298.420 592	0.90	
p3	-1298.420 053	2.31	
p4	-1298.419 945	2.59	
t1	-1298.420 849	0.22	
t2	-1298.418 925	5.27	
w1	-1298.420 883	0.13	
w2	-1298.420 379	1.46	
w3	-1298.419 453	3.89	
w4	-1298.419 462	3.86	
c1	-1298.419 979	2.51	
c2	-1298.419 128	4.74	

mp2		RI-SCS-MP2/def2-TZVPPD	
structure	absolute energy / Hartree	relative energy /	kJ/mol
p1	-1298.126 007	1.60	
p2	-1298.125 687	2.43	
p3	-1298.125 201	3.71	
p4	-1298.124 957	4.35	
t1	-1298.125 899	1.88	
t2	-1298.123 355	8.56	
w1	-1298.126 616	0.00	
w2	-1298.126 275	0.89	
w3	-1298.125 072	4.05	
w4	-1298.125 237	3.62	
c1	-1298.124 396	5.83	
c2	-1298.123 574	7.98	

b2plyp		B2PLYP-D4/def2-TZVPPD	
structure	absolute energy / Hartree	relative energy /	kJ/mol
p1	-1299.366 933	0.00	
p2	-1299.366 525	1.07	
p3	-1299.366 022	2.39	
p4	-1299.365 961	2.55	
t1	-1299.366 684	0.65	
t2	-1299.364 053	7.56	
w1	-1299.366 612	0.84	
w2	-1299.366 352	1.52	
w3	-1299.365 243	4.43	
w4	-1299.365 154	4.67	
c1	-1299.365 344	4.17	
c2	-1299.364 355	6.76	

wb97		ω B97M-D4/def2-TZVPPD	
structure	absolute energy / Hartree	relative energy /	kJ/mol
p1	-1300.422 829	2.29	
p2	-1300.422 494	3.17	
p3	-1300.421 941	4.62	
p4	-1300.421 832	4.91	
t1	-1300.422 798	2.37	
t2	-1300.421 255	6.42	
w1	-1300.423 703	0.00	
w2	-1300.423 101	1.58	
w3	-1300.422 013	4.43	
w4	-1300.422 232	3.86	
c1	-1300.422 653	2.75	
c2	-1300.421 790	5.02	

mbpol2016		MB-pol2016	
structure	absolute energy / Hartree	relative energy /	kJ/mol
p1	-0.274 008	0.77	
p2	-0.272 592	4.49	
p3	-0.272 038	5.94	
p4	-0.272 626	4.40	
t1	-0.273 327	2.56	
t2	-0.271 935	6.22	
w1	-0.273 706	1.56	
w2	-0.272 680	4.26	
w3	-0.271 919	6.26	
w4	-0.271 613	7.06	
c1	-0.274 304	0.00	
c2	-0.273 688	1.61	

mbpol2023		MB-pol2023	
structure	absolute energy / Hartree	relative energy /	kJ/mol
p1	-0.277 555	0.00	
p2	-0.275 663	4.96	
p3	-0.275 084	6.48	
p4	-0.275 773	4.67	
t1	-0.276 705	2.23	
t2	-0.274 098	9.07	
w1	-0.276 777	2.04	
w2	-0.275 796	4.61	
w3	-0.275 300	5.92	
w4	-0.273 834	9.76	
c1	-0.276 861	1.82	
c2	-0.276 361	3.13	

qaqua		q-AQUA	
structure	absolute energy / Hartree	relative energy /	kJ/mol
p1	-0.281 220	4.56	
p2	-0.281 651	3.43	
p3	-0.280 649	6.06	
p4	-0.280 528	6.38	
t1	-0.281 699	3.30	
t2	-0.277 311	14.82	
w1	-0.282 959	0.00	
w2	-0.282 638	0.84	
w3	-0.281 140	4.77	
w4	-0.278 712	11.14	
c1	-0.277 228	15.04	
c2	-0.275 978	18.32	

qaquapol		q-AQUA-pol	
structure	absolute energy / Hartree	relative energy /	kJ/mol
p1	-0.275 637	2.52	
p2	-0.275 179	3.72	
p3	-0.275 817	2.04	
p4	-0.276 113	1.27	
t1	-0.276 598	0.00	
t2	-0.273 634	7.78	
w1	-0.274 806	4.70	
w2	-0.275 177	3.73	
w3	-0.275 195	3.68	
w4	-0.271 764	12.69	
c1	-0.274 764	4.81	
c2	-0.273 753	7.46	

ttm3f		TTM3-F	
structure	absolute energy / Hartree	relative energy /	kJ/mol
p1	-0.276 652	6.90	
p2	-0.277 107	5.70	
p3	-0.276 111	8.32	
p4	-0.275 200	10.71	
t1	-0.276 070	8.43	
t2	-0.277 264	5.29	
w1	-0.279 281	0.00	
w2	-0.277 455	4.79	
w3	-0.276 924	6.18	
w4	-0.278 846	1.14	
c1	-0.277 955	3.48	
c2	-0.278 420	2.25	

amoeba+		AMOEBA+	
structure	absolute energy / Hartree	relative energy / kJ/mol	
p1	-0.274 308	3.88	
p2	-0.273 985	4.73	
p3	-0.273 180	6.84	
p4	-0.273 171	6.86	
t1	-0.272 872	7.65	
t2	-0.273 689	5.50	
w1	-0.275 785	0.00	
w2	-0.274 295	3.91	
w3	-0.274 094	4.44	
w4	-0.274 489	3.40	
c1	-0.275 520	0.69	
c2	-0.274 908	2.30	

amoeba14		AMOEBA-2014	
structure	absolute energy / Hartree	relative energy / kJ/mol	
p1	-0.271 896	5.44	
p2	-0.271 985	5.20	
p3	-0.270 965	7.88	
p4	-0.270 730	8.50	
t1	-0.271 025	7.72	
t2	-0.271 928	5.35	
w1	-0.273 967	0.00	
w2	-0.272 673	3.40	
w3	-0.272 309	4.35	
w4	-0.272 800	3.06	
c1	-0.273 125	2.21	
c2	-0.272 895	2.81	

tip4p2005		TIP4P/2005	
structure	absolute energy / Hartree	relative energy / kJ/mol	
p1	-0.302 483	4.37	
p2	-0.302 732	3.71	
p3	-0.302 134	5.28	
p4	-0.302 015	5.59	
t1	-0.302 574	4.13	
t2	-0.304 145	0.00	
w1	-0.303 314	2.18	
w2	-0.302 800	3.53	
w3	-0.302 370	4.66	
w4	-0.303 407	1.94	
c1	-0.302 821	3.48	
c2	-0.303 027	2.94	

tip4p		TIP4P	
structure	absolute energy / Hartree	relative energy /	kJ/mol
p1	-0.274 296	3.64	
p2	-0.274 526	3.04	
p3	-0.273 980	4.47	
p4	-0.273 871	4.76	
t1	-0.274 340	3.52	
t2	-0.275 684	0.00	
w1	-0.274 897	2.06	
w2	-0.274 381	3.42	
w3	-0.273 989	4.45	
w4	-0.274 937	1.96	
c1	-0.274 490	3.13	
c2	-0.274 700	2.58	

gfn2-xtb		GFN2-xTB	
structure	absolute energy / Hartree	relative energy /	kJ/mol
p1	-86.479 330	7.41	
p2	-86.479 382	7.28	
p3	-86.478 902	8.54	
p4	-86.478 527	9.52	
t1	-86.479 299	7.50	
t2	-86.481 660	1.30	
w1	-86.482 156	0.00	
w2	-86.481 114	2.73	
w3	-86.481 518	1.67	
w4	-86.481 499	1.72	
c1	-86.481 428	1.91	
c2	-86.481 625	1.39	

tip3p		TIP3P	
structure	absolute energy / Hartree	relative energy /	kJ/mol
p1	-0.269 678	5.57	
p2	-0.270 697	2.90	
p3	-0.269 411	6.27	
p4	-0.267 847	10.38	
t1	-0.268 016	9.93	
t2	-0.266 757	13.24	
w1	-0.271 802	0.00	
w2	-0.268 872	7.69	
w3	-0.266 597	13.66	
w4	-0.270 062	4.56	
c1	-0.266 512	13.88	
c2	-0.268 489	8.69	

gfn-ff	GFN-FF	
structure	absolute energy / Hartree	relative energy / kJ/mol
p1	-5.836 569	2.30
p2	-5.837 448	0.00
p3	-5.835 457	5.22
p4	-5.833 811	9.54
t1	-5.834 756	7.06
t2	-5.829 560	20.71
w1	-5.836 606	2.21
w2	-5.833 809	9.55
w3	-5.831 423	15.81
w4	-5.833 225	11.08
c1	-5.828 489	23.52
c2	-5.830 776	17.51

3 Deviation measures

To support the $(\text{H}_2\text{O})_{17}$ energy graphs in the main paper, the following table list mean energy differences and standard deviations of these differences, between each method used and a reference. As shown in the main paper as one of the central points, there are small but significant differences between the results of the different ab-initio methods used. However, the differences between those ab-initio results on the one hand and the water model results on the other hand are larger still. Therefore, here it does not matter much which of the ab-initio methods are chosen as reference. Hence, we take as reference “e34”, i.e., DLPNO-CCSD(T) CBS-extrapolated from triple- and quadruple-zeta basis sets. All data are based on the “relative energy” numbers in kJ/mol listed in the tables in the previous section and shown in the graphs in the main paper.

differences to e34		
method	mean / kJ/mol	standard deviation / kJ/mol
ccsd/f12	0.680	0.617
ccsd/qz	0.646	0.611
mp2	1.93	0.837
b2plyp	0.824	0.817
wb97m	1.42	0.686
mbpol2023	1.97	1.55
mbpol2016	1.73	1.13
quaquapol	2.29	1.85
quaqua	5.19	4.38
gfn2-xtb	4.19	2.52
gfn-ff	7.65	6.20
ttn3f	3.77	2.78
amoeba+	2.69	1.92
amoeba14	2.92	2.46
tip4p2005	2.48	1.66
tip4p	2.17	1.58
tip3p	5.54	3.35

In the present case, these statistical measures are less meaningful than one could hope. For example, for q-AQUA-pol, much of the total deviation is due to one single value (w4, easily visible in Fig.4 of the main paper). With this value deleted, the mean value of the differences is somewhat smaller (1.82 kJ/mol) while the standard deviation is cut in half (0.98 kJ/mol, even less than for both MB-pol versions).

However, this focus on one big deviation is not true for the other methods. In fact, deviation distributions turn out to be highly method/model-dependent, which tends to be hidden by the integrative statistical indicators shown in the above table. Hence, it is more informative to inspect the 12 actual values for each method, as shown in the main paper.

4 Further water model versions

To avoid cluttering the figures in the main paper, only one version of AMOEBA and TIP4P, respectively, is shown there. Fig. S5 demonstrates that for the $(\text{H}_2\text{O})_{17}$ cluster structures under study, TIP4P [3] and TIP4P/2005 [4] data are very similar, as well as the AMOEBA2014 [5] and AMOEBA+ [6,7] results, respectively.

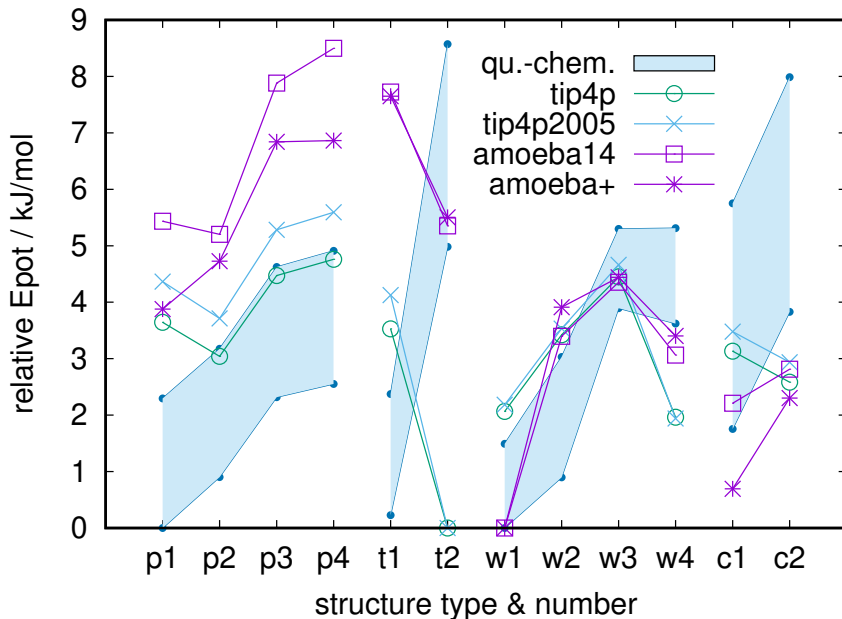


Figure S5: Relative potential energies for the full set of 12 structures, comparing TIP4P, TIP4P2005, AMOEBA2014 and AMOEBA+. As in the main paper, the shaded regions represent the quantum-chemistry results.

5 Water hexamer

To help gauging the DFT and ab-initio approaches used in the main paper, the following table lists total electronic energies of the well-known low-energy $(\text{H}_2\text{O})_6$ structures “cage”, “book” and “6-ring”, relative to the “prism” and in direct comparison to literature values [8–13]. Since there is some spread in these literature values, due to differences in basis sets, CBS-extrapolations and other methodical issues, they are summarized in

value intervals. However, MP2 and CCSD(T) from the literature are shown separately. All values are in kJ/mol; the method abbreviations from the main paper are used. As in the main paper for $(\text{H}_2\text{O})_{17}$, local structure optimizations were performed for SCS-MP2 and B2PLYP, while all CCSD(T) results are single-point calculations for the B2PLYP-optimized $(\text{H}_2\text{O})_6$ structures.

structure	b2plyp	mp2	ccsdt/qz	ccsdt/f12	e34	lit.mp2	lit.ccsdt
prism	0.0	0.0	0.0	0.0	0.0	0.0	0.0
cage	0.22	0.42	1.49	1.31	1.73	0.38 – 0.39	0.84 – 1.26
book	0.32	0.62	2.86	2.93	2.54	1.13 – 1.54	2.64 – 5.02
6ring	3.78	3.23	6.74	7.15	5.84	4.27 – 5.31	6.44 – 9.12

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