

Supporting Info for:

Formation of water polyhedrons in propofol-water clusters

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KEYWORDS: *electronic spectroscopy, jets, hydrogen bond, solvation, anesthetics, IR spectroscopy.*

Figure S01. Propofol conformers with their relative energies in kJ/mol, as calculated at MP2/6-311++G(d,p) level.

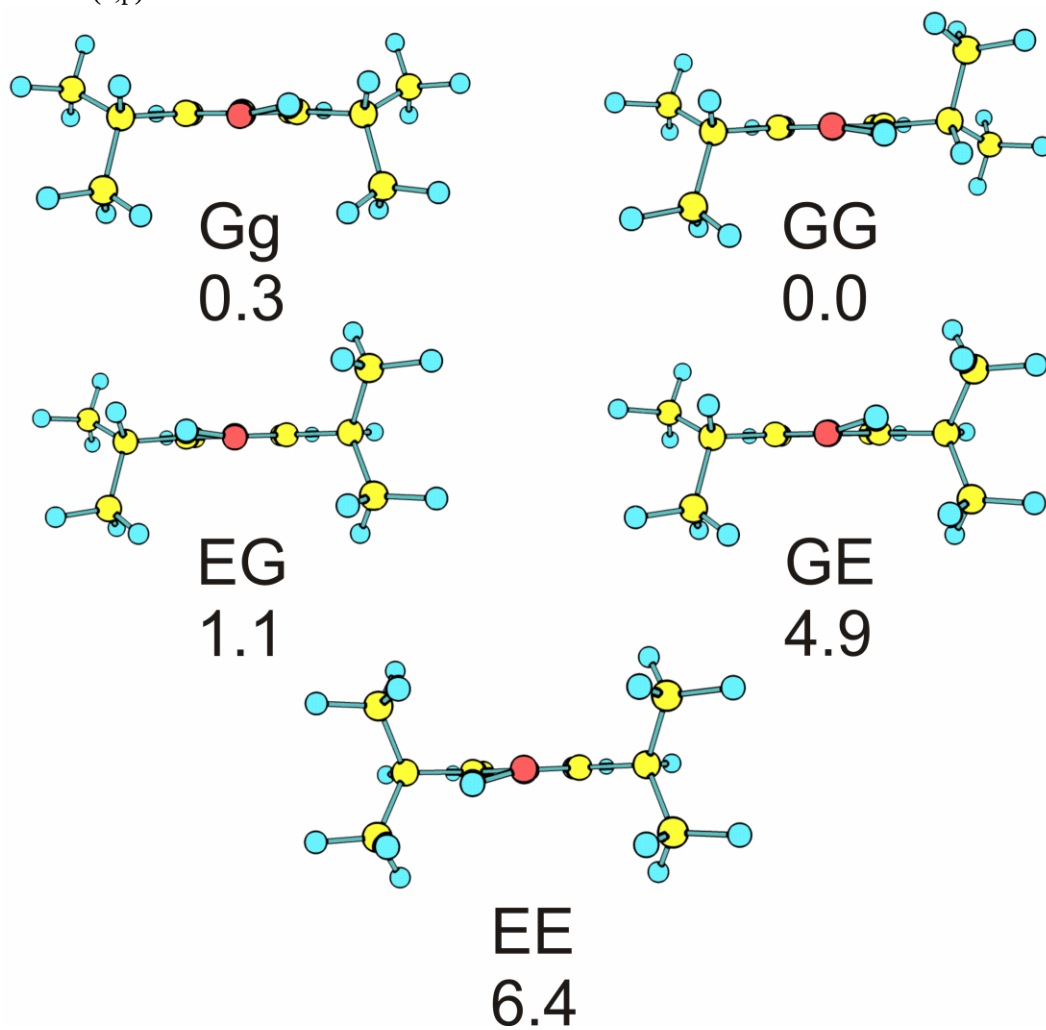


Figure S2. Calculated structures for propofol₁(H₂O)₇ at M06-2X/6-311++G(d,p) level, together with their relative stability in kJ/mol. ZPE correction was applied to all the energy values

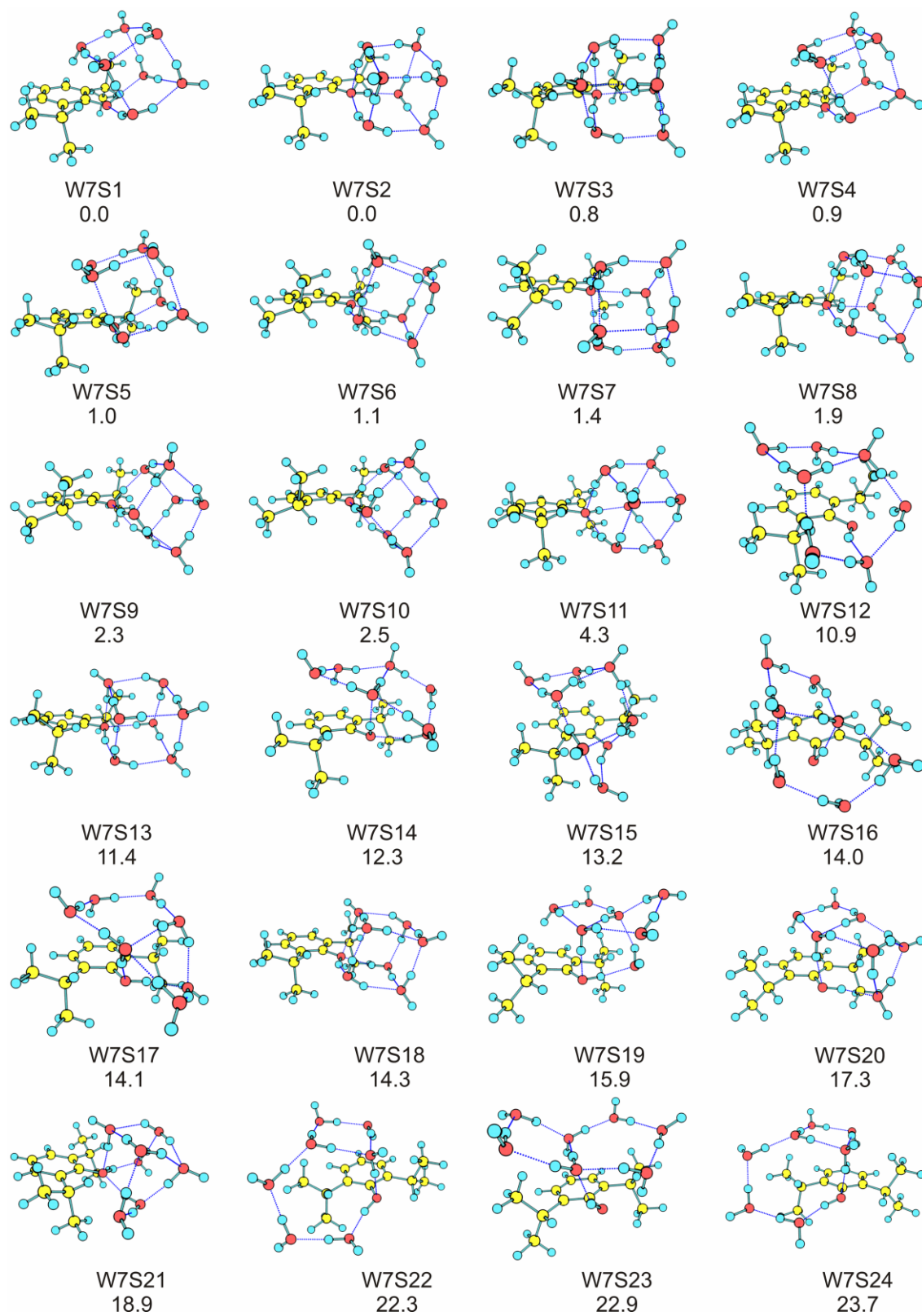


Figure S2.(Cont.)

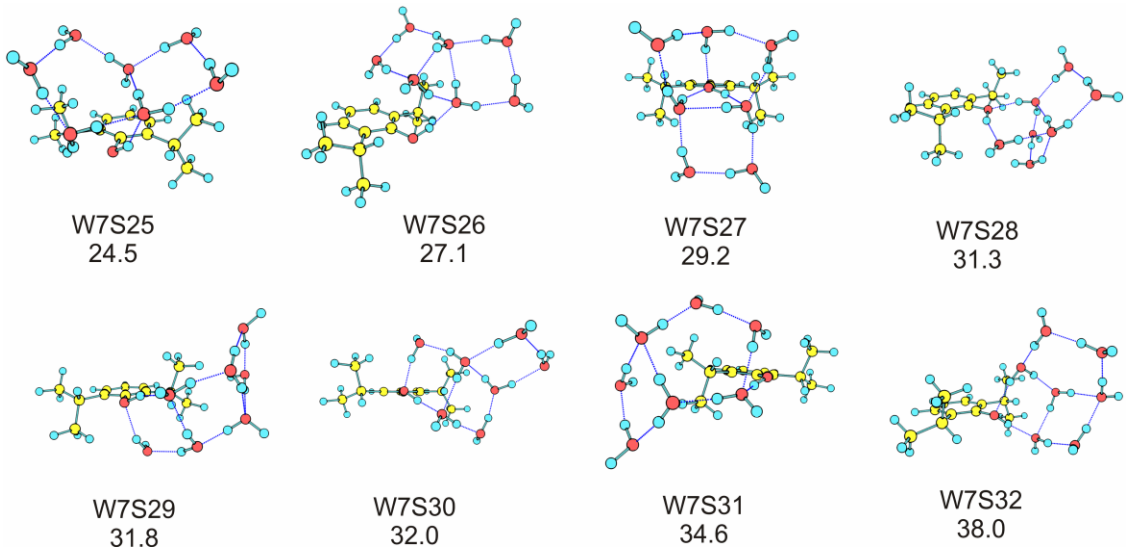


Figure S3. Experimental IDIRS for propofol₁(H₂O)₇ (upper trace) together with the predicted frequencies for each calculated structure. A correction factor of 0.938 was employed.

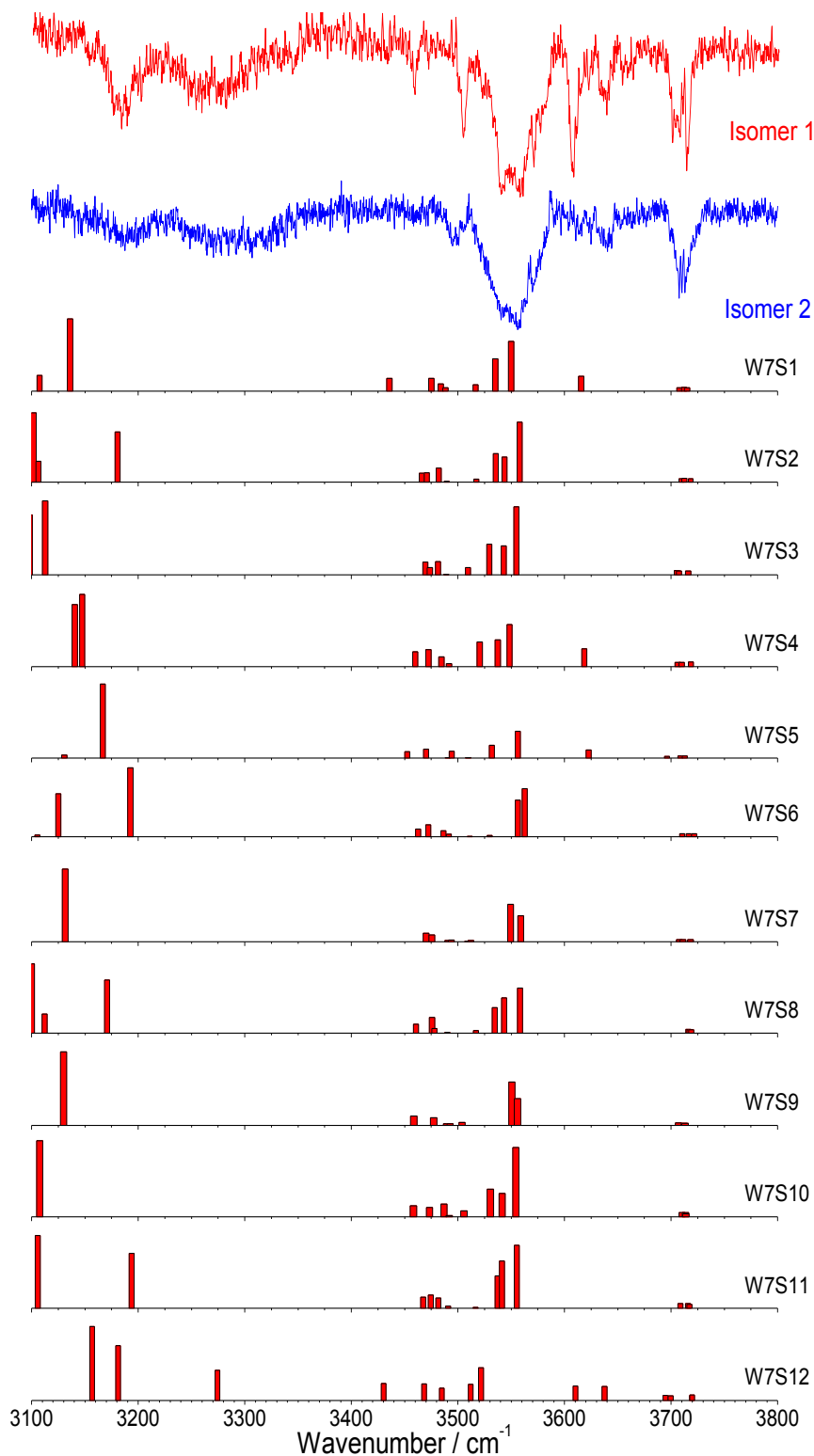


Figure S3.(Cont.)

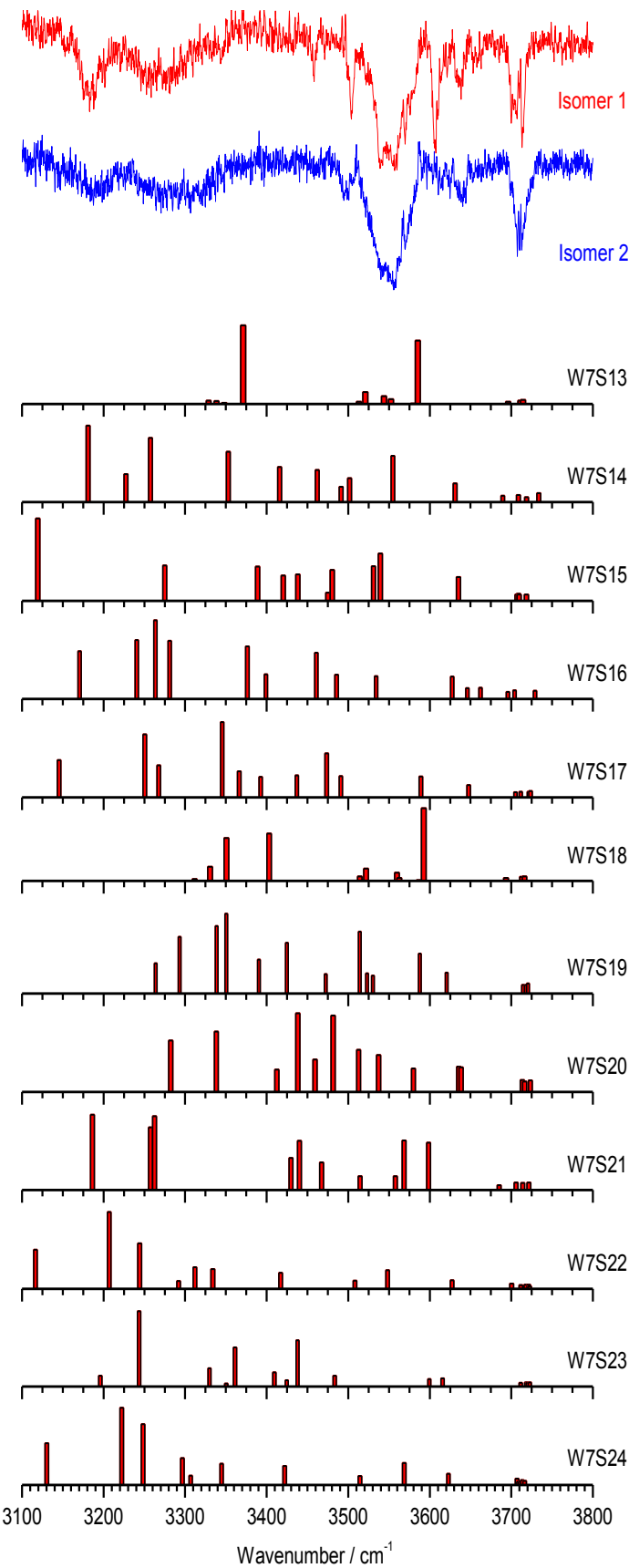
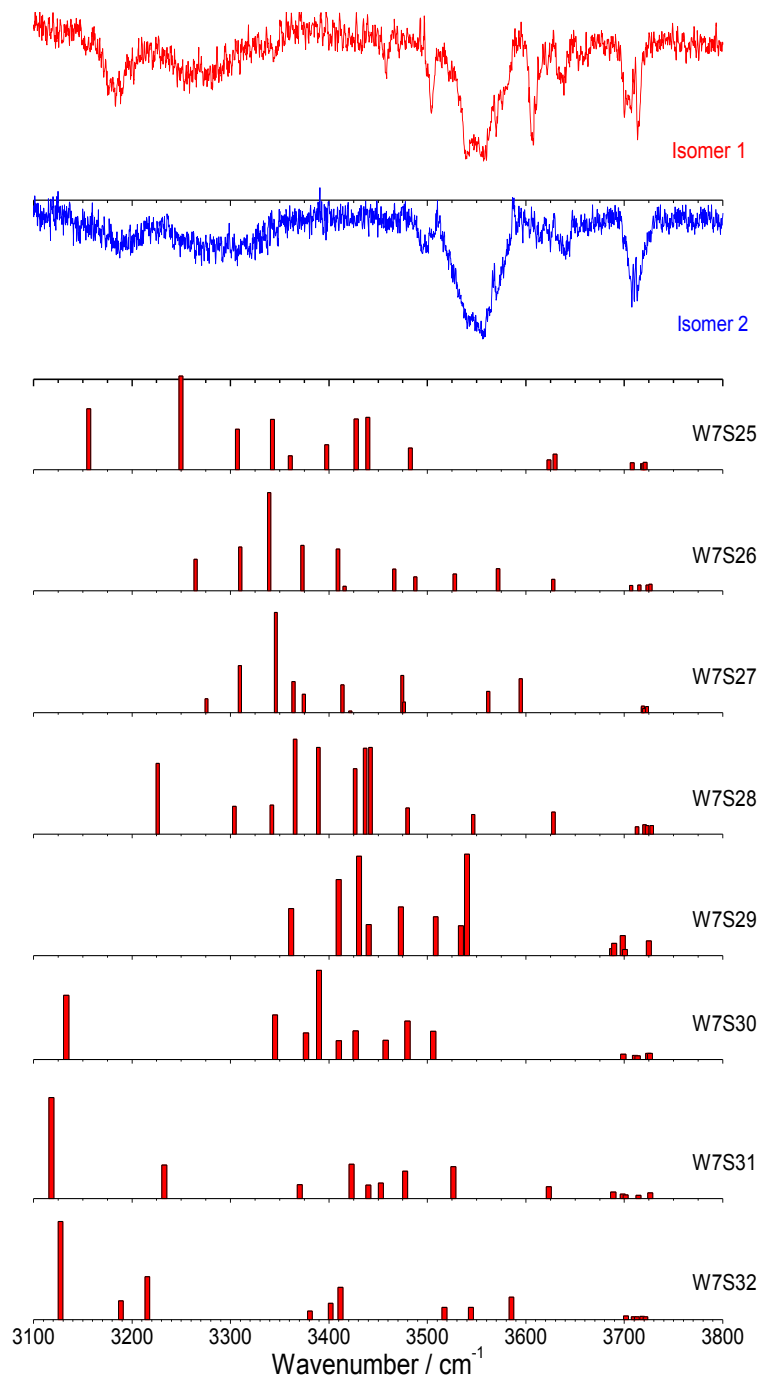


Figure S3 (cont.)



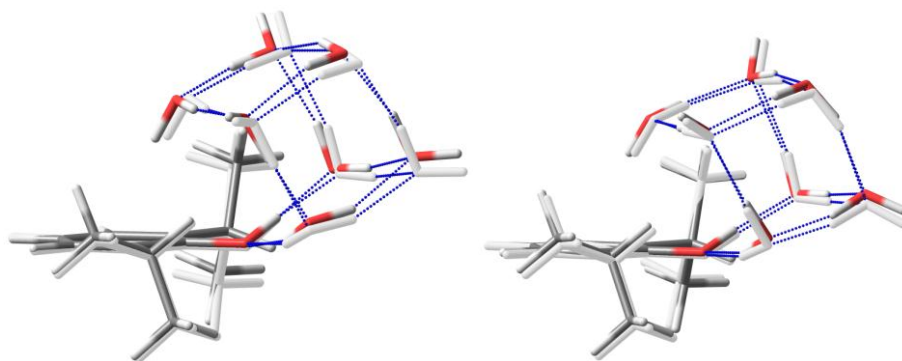


Figure S4. Comparison between the structures of W7S1 (left) and W7S4 (right) calculated at M06-2X/G-311++G(d,p) and at M06-2X/AUG-cc-pVTZ (in white). Small differences between the two structures are found.

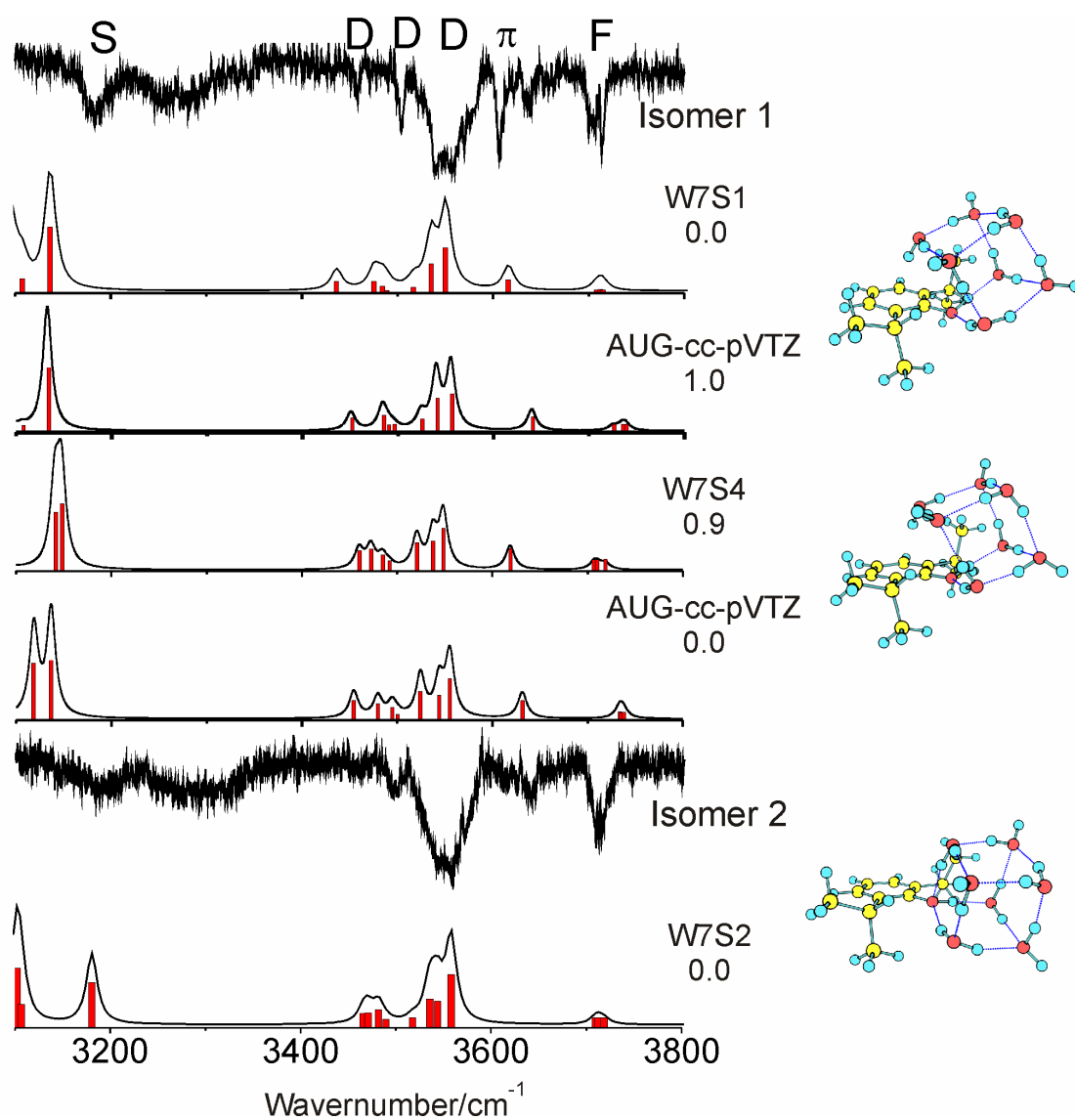


Figure S5. Comparison between the experimental IDIRS of the two propofol₁(H₂O)₇ isomers and the spectra predicted for W7S1 and W7S4 using two basis sets: 6-311++G(d,p) and AUG-cc-pVTZ. A correction factor of 0.950 was used for the second basis set.

Figure S6. Calculated structures for propofol₁(H₂O)₈ at M06-2X/6-311++G(d,p) level, together with their relative stability in kJ/mol. ZPE correction was applied to all the energy values

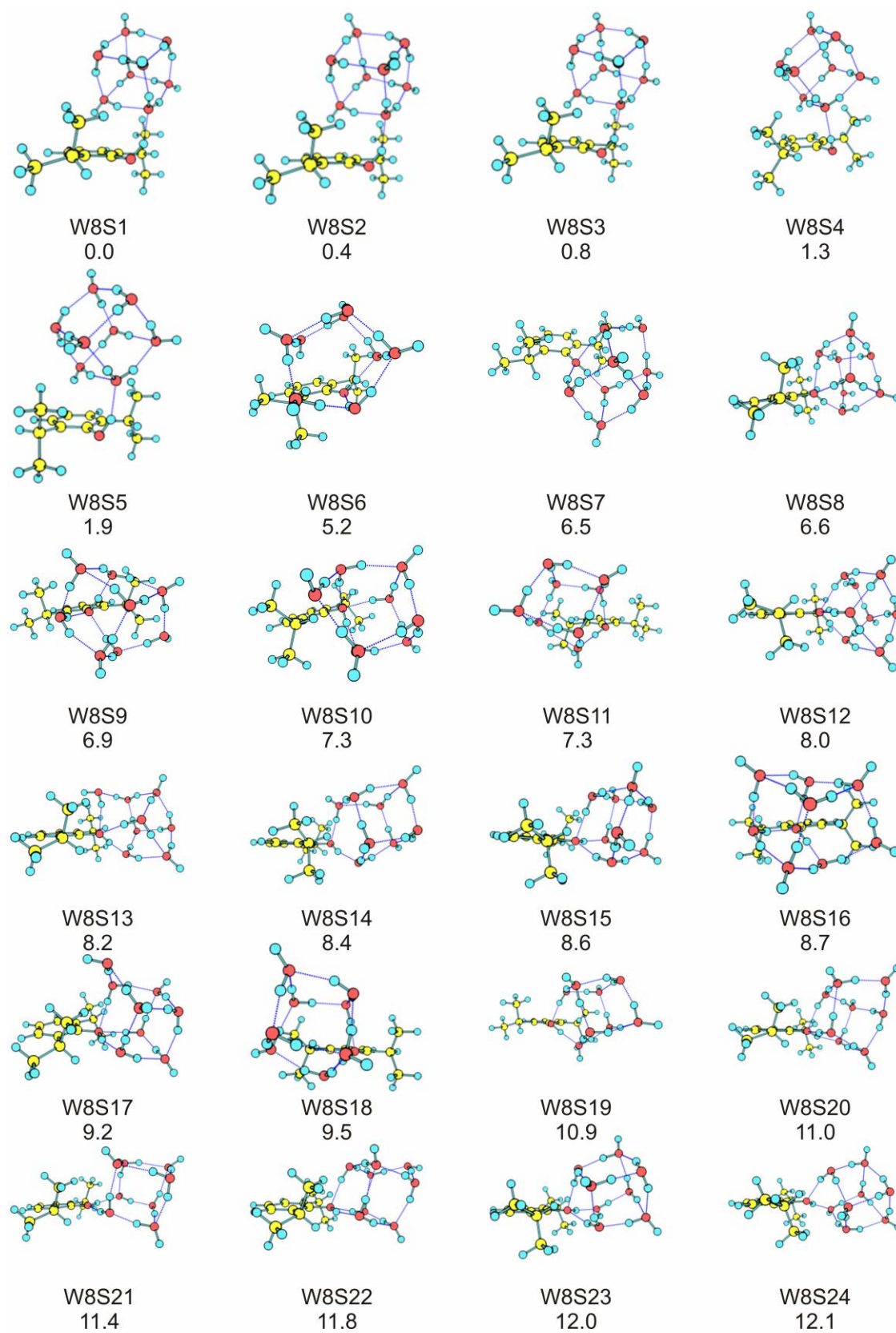


Figure S6. (Cont.)

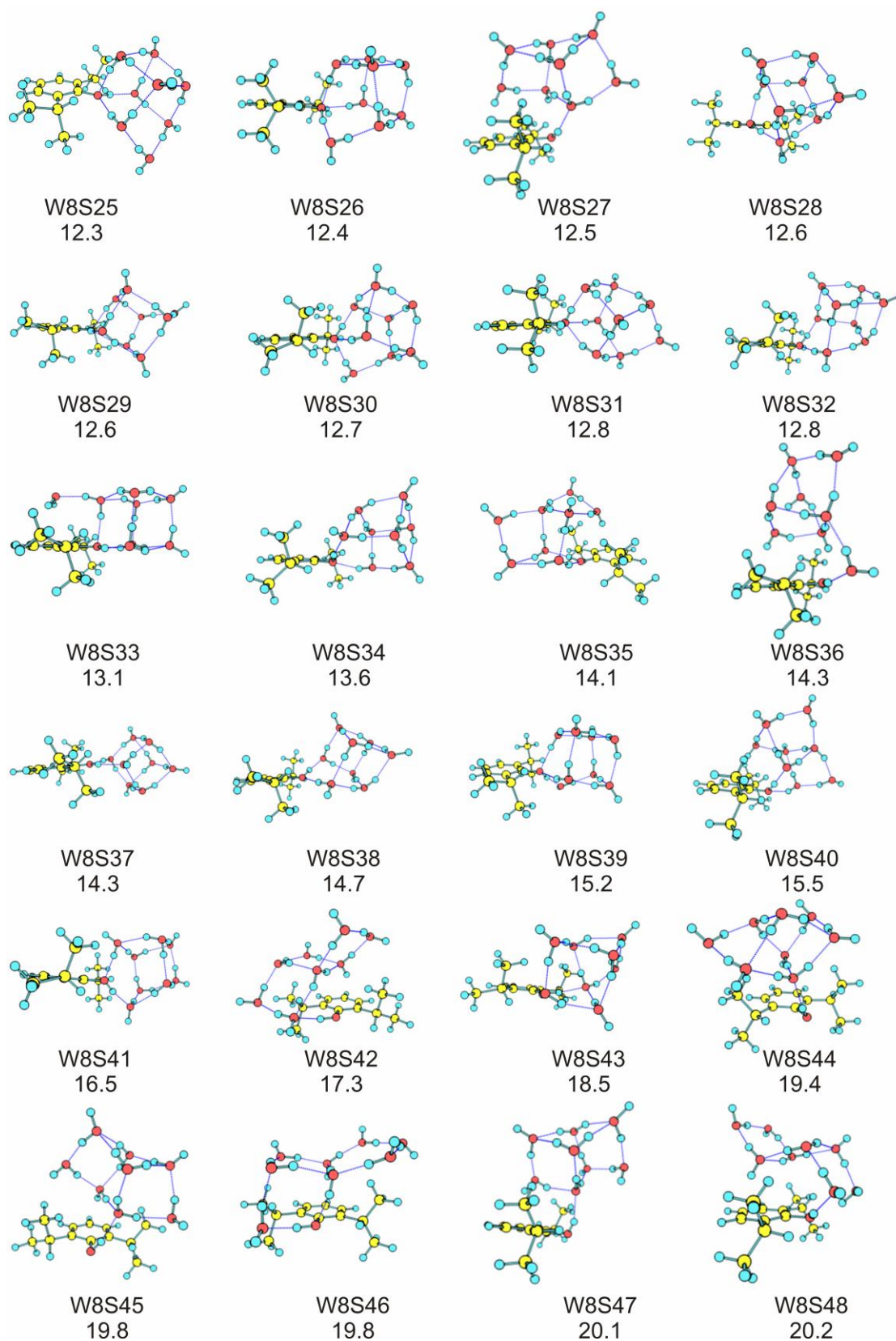


Figure S6. (Cont.)

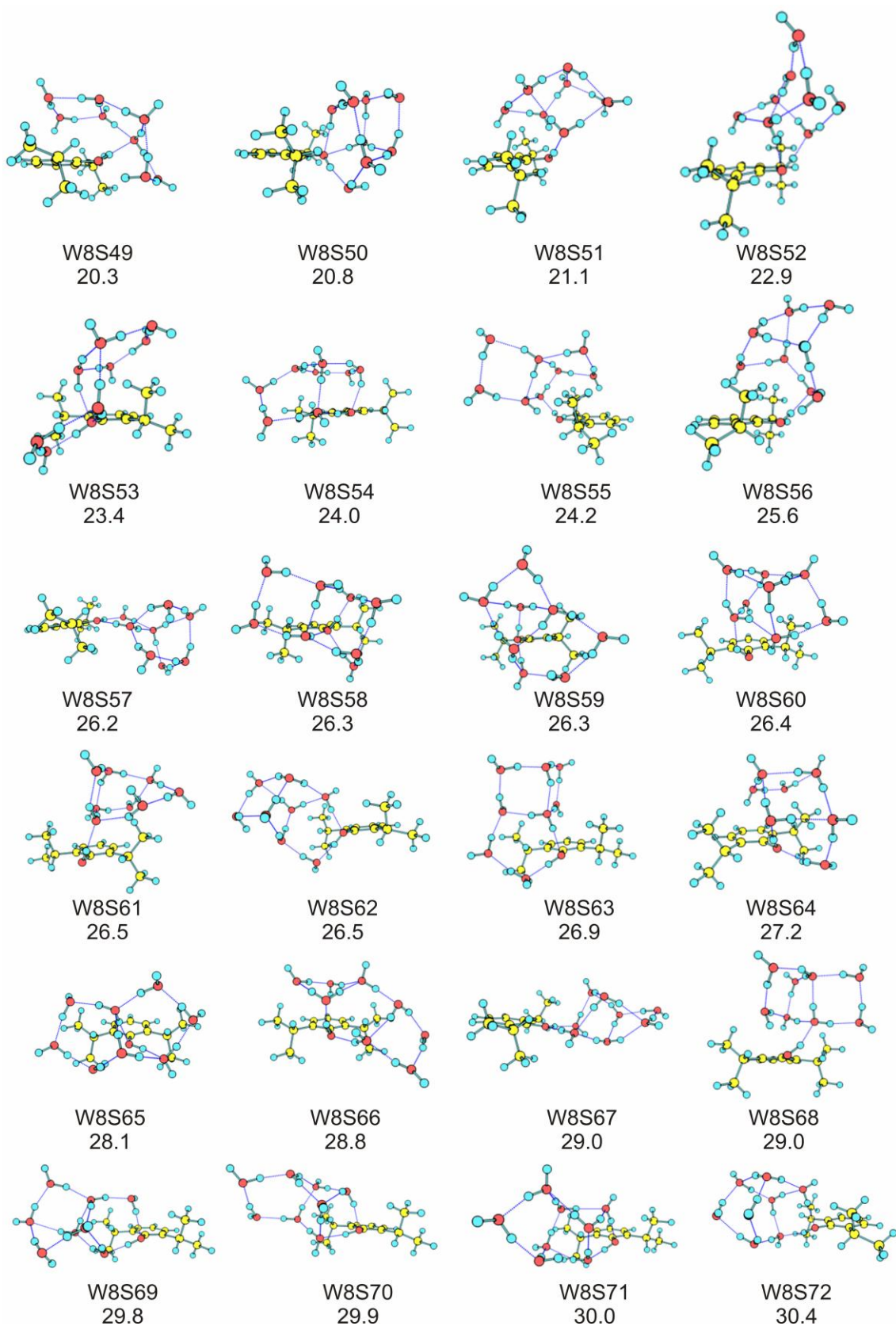


Figure S6. (Cont.)

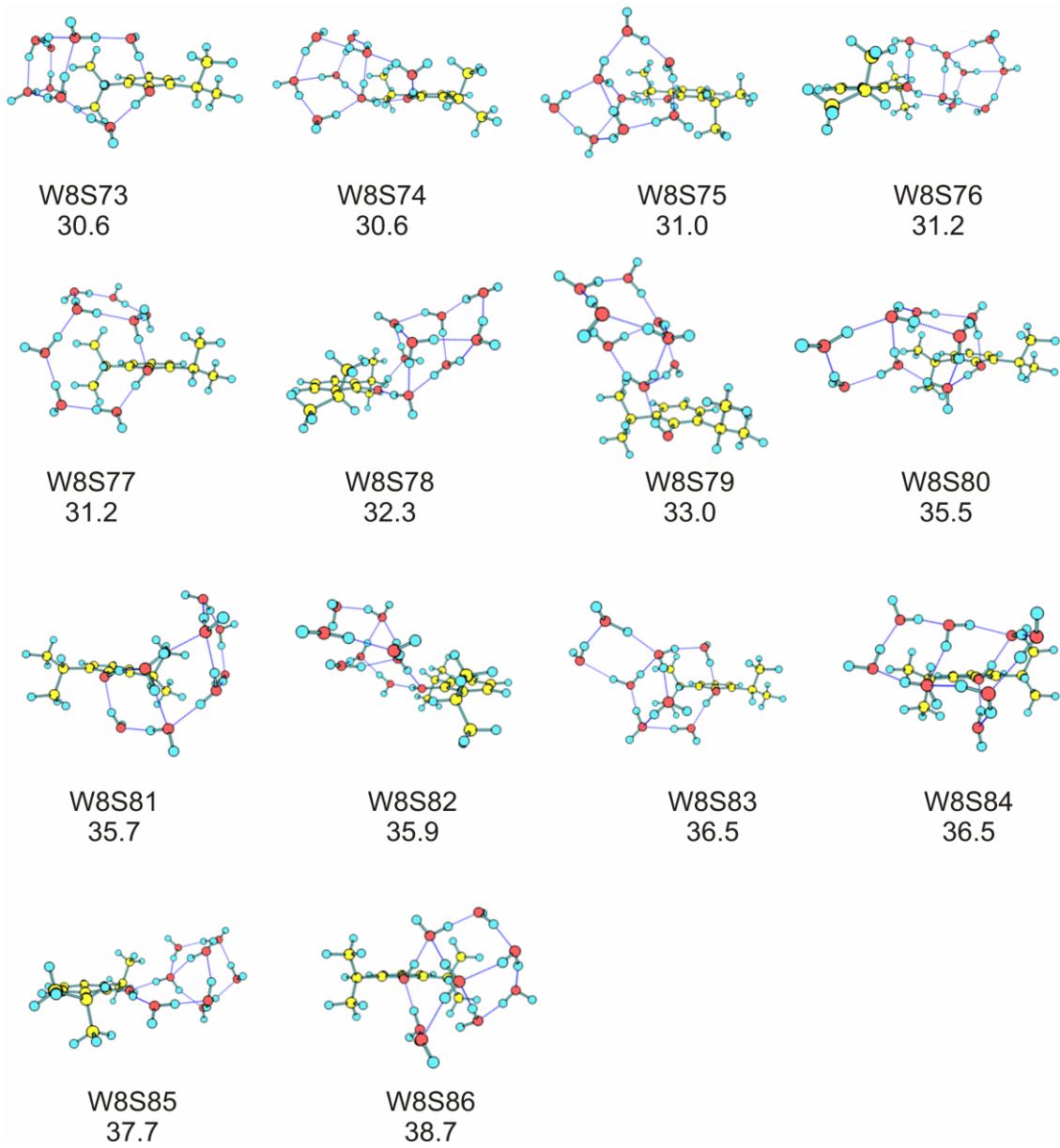


Figure S7. Experimental IDIRS for propofol₁(H₂O)₈ (upper trace) together with the predicted frequencies for each calculated structure. A correction factor of 0.938 was employed.

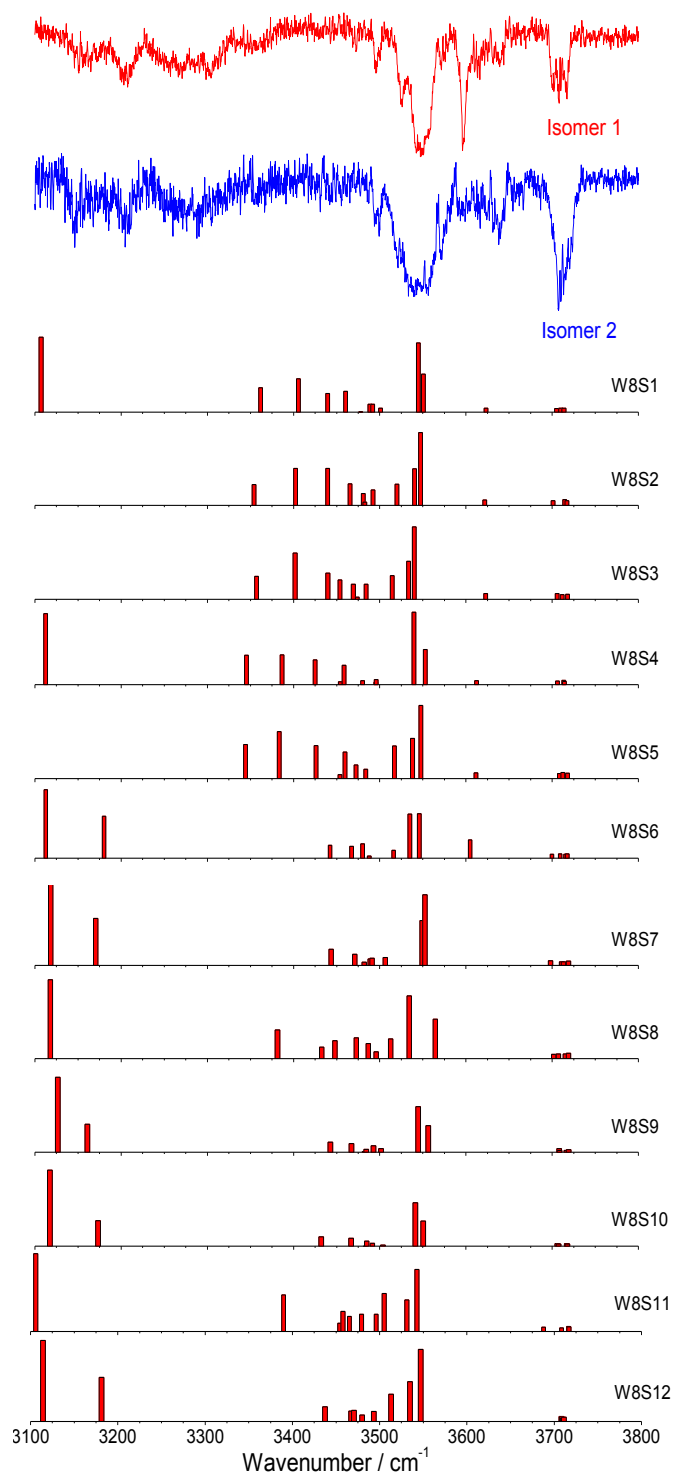


Figure S7.(Cont.)

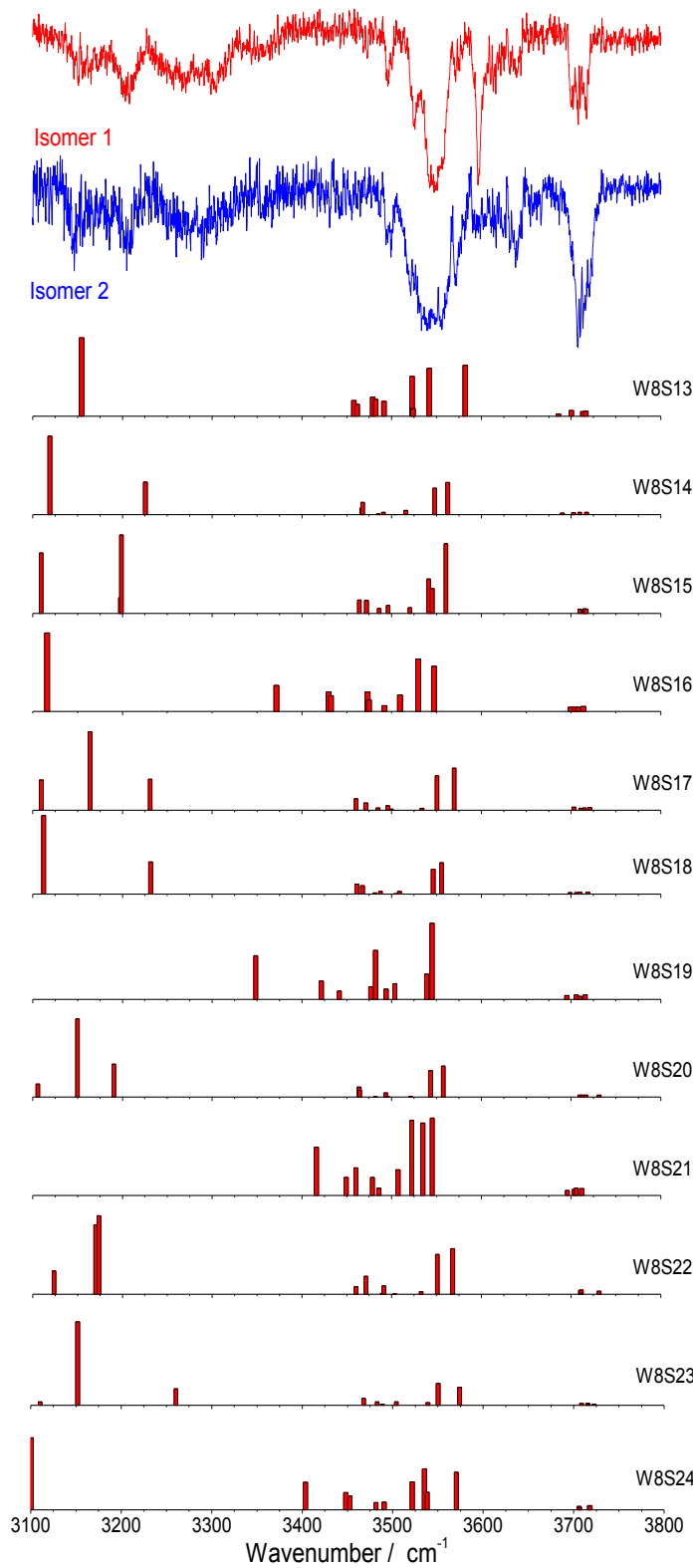


Figure S7.(Cont.)

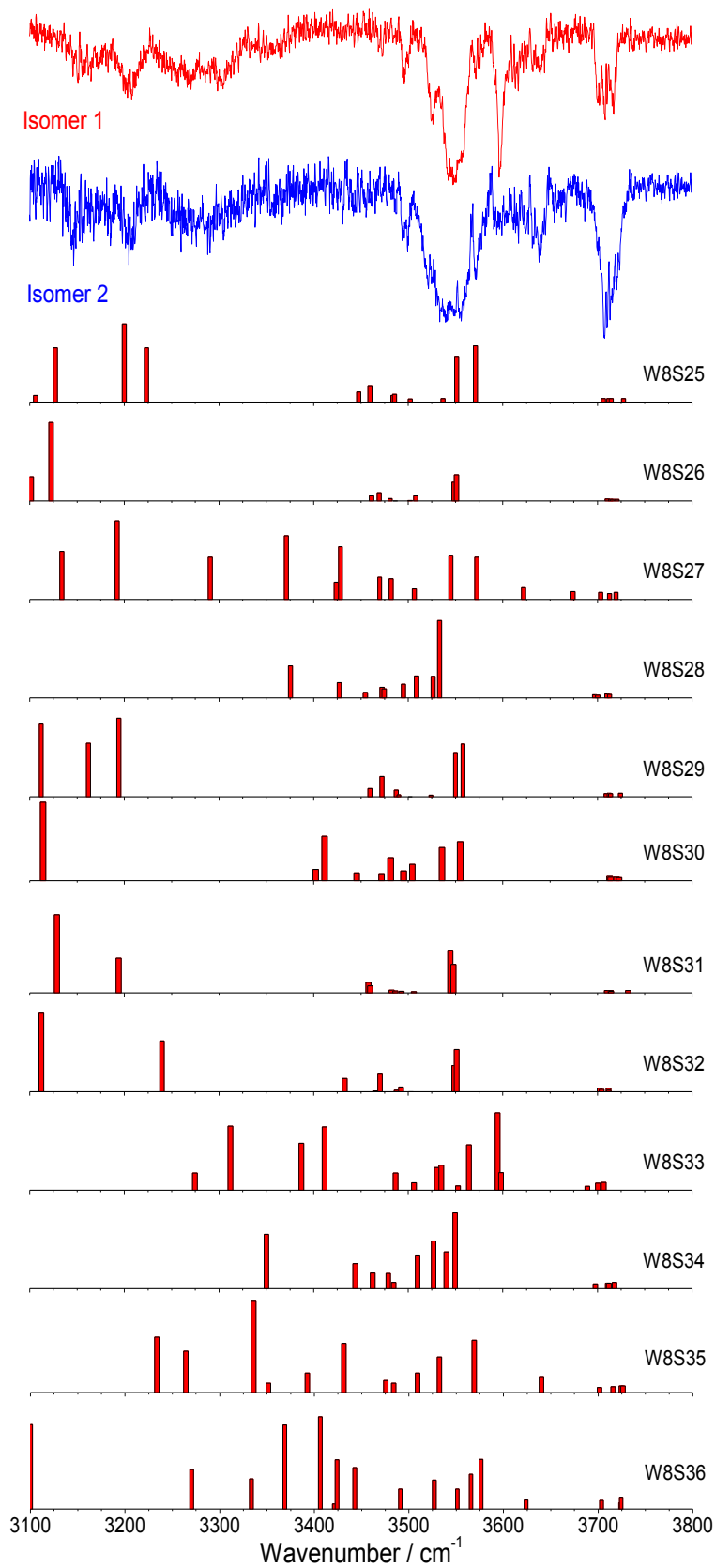


Figure S7.(Cont.)

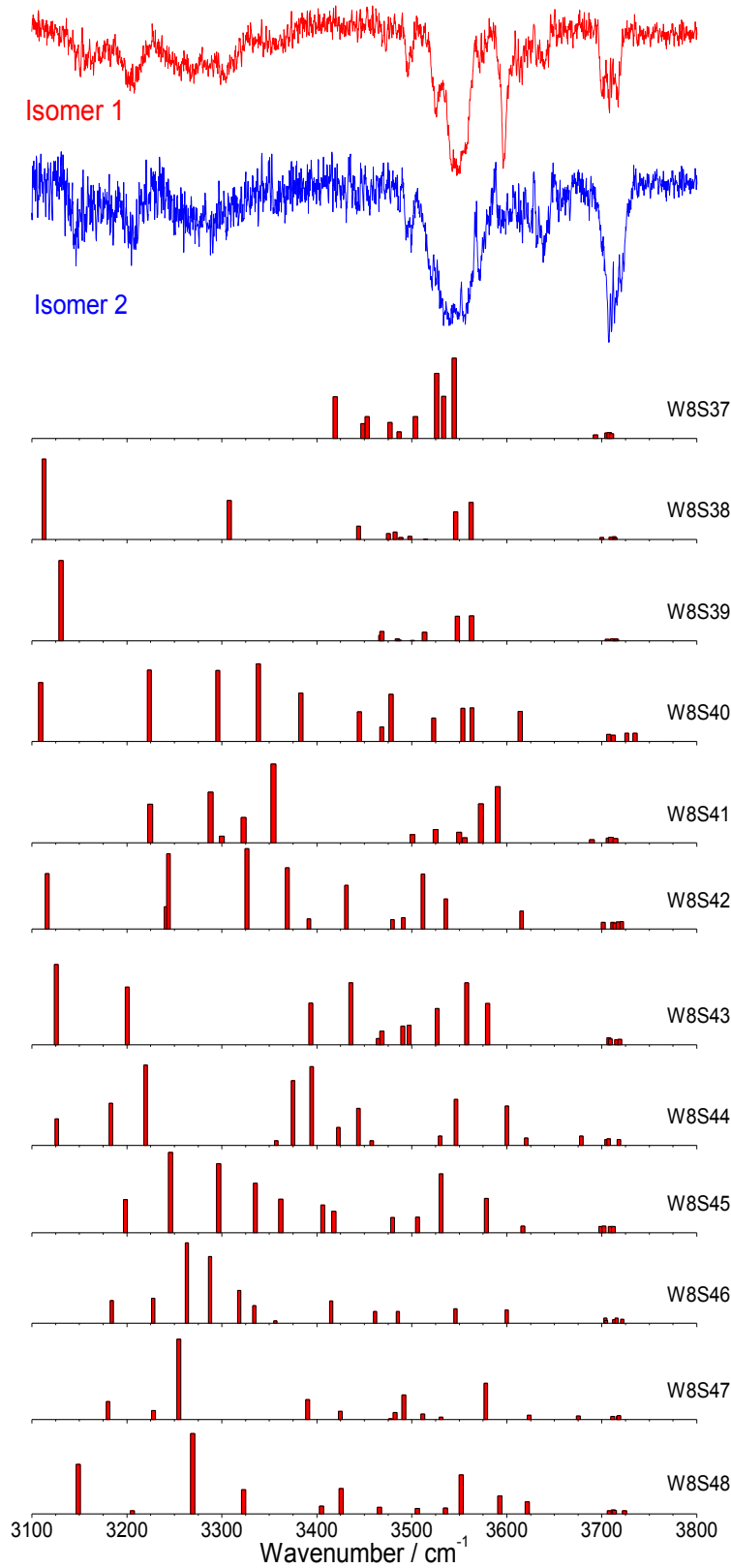


Figure S7.(Cont.)

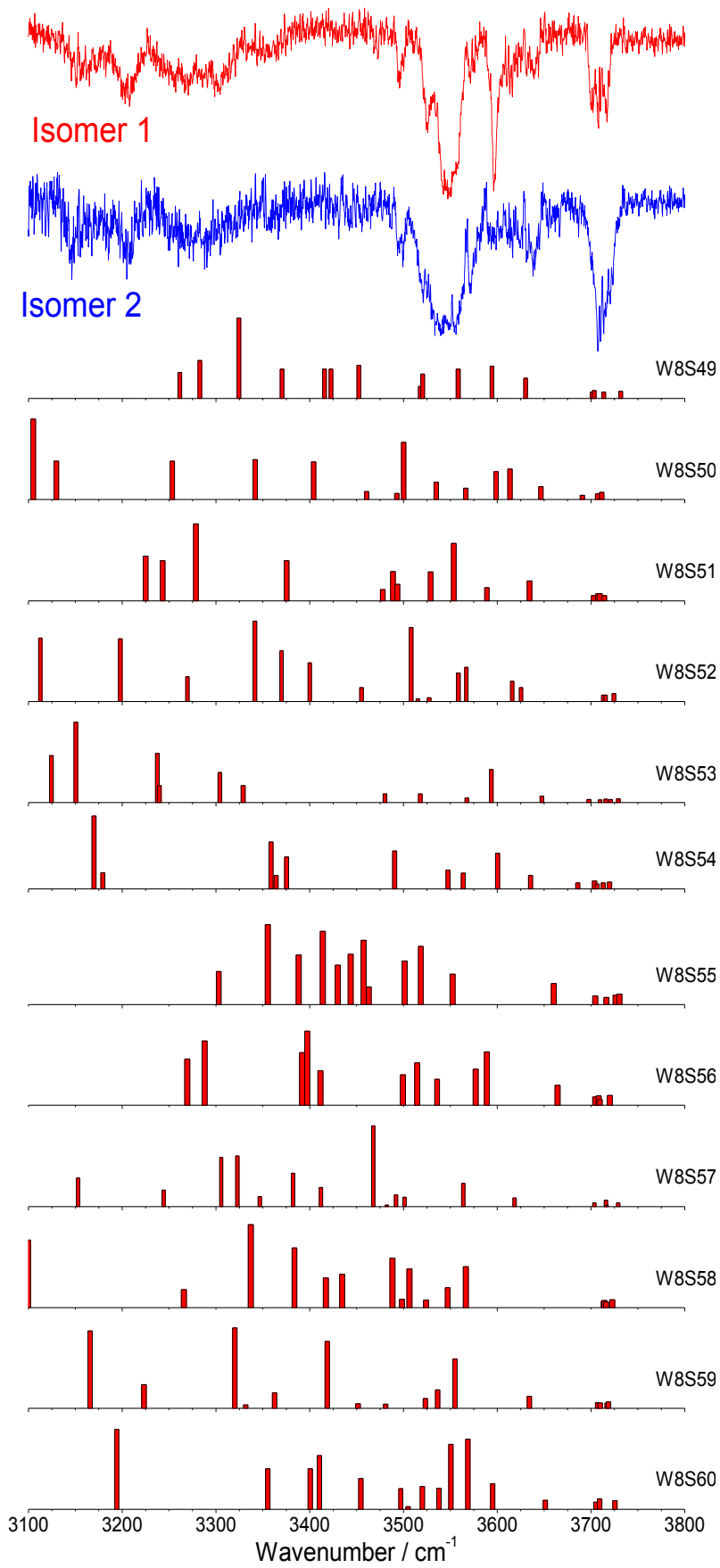


Figure S7.(Cont.)

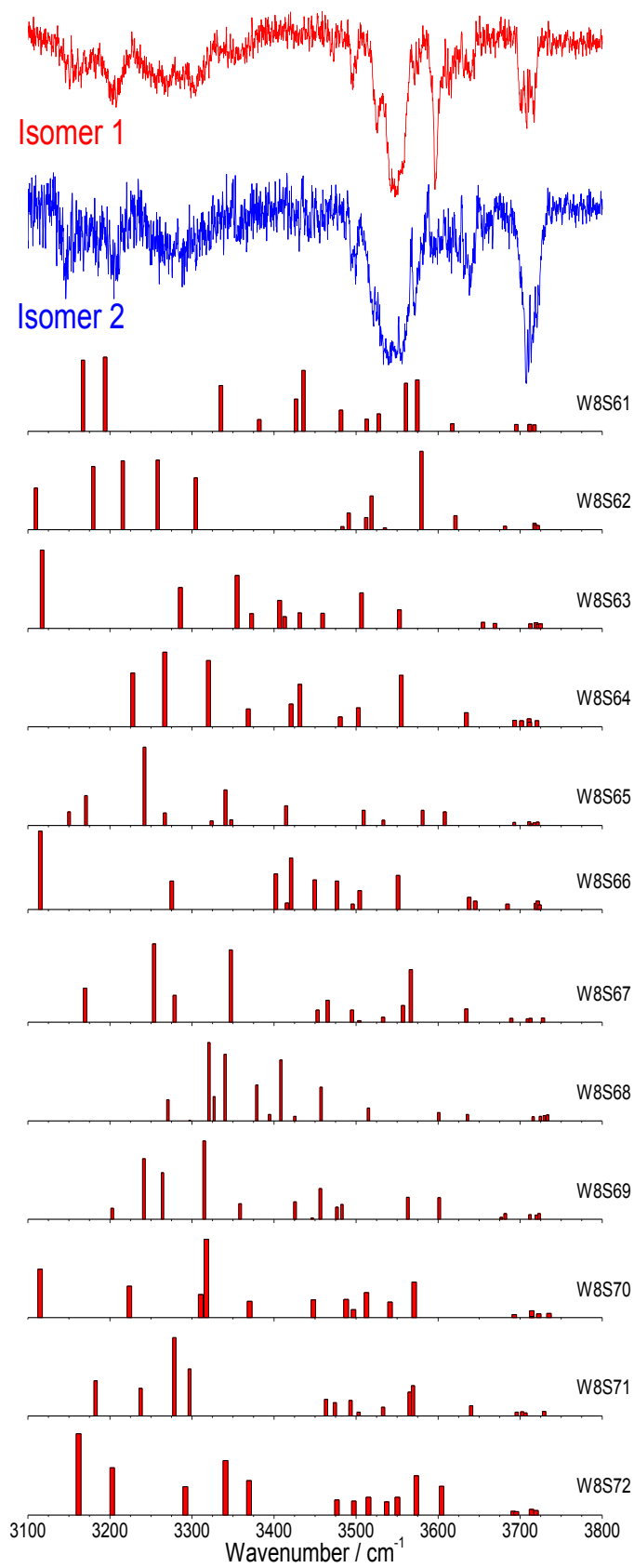


Figure S7.(Cont.)

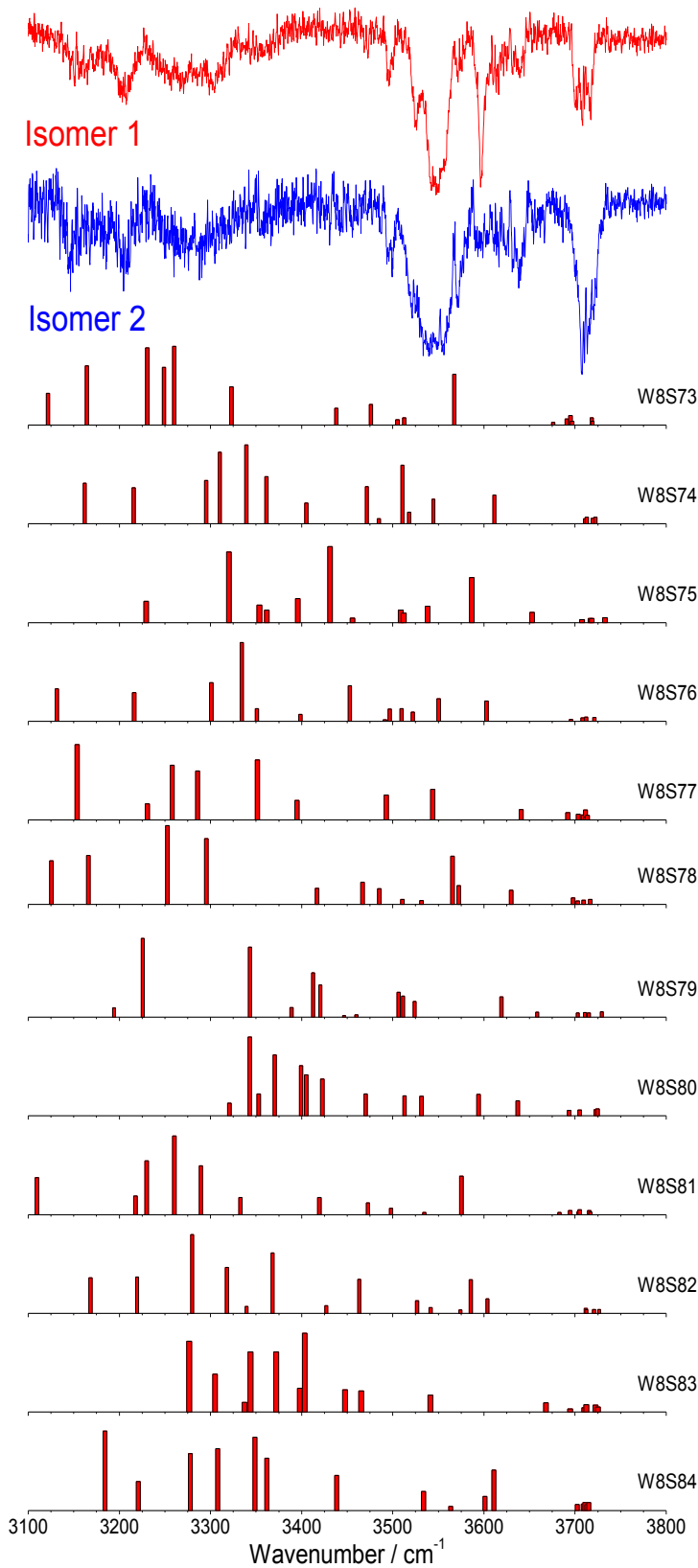


Figure S7.(Cont.)

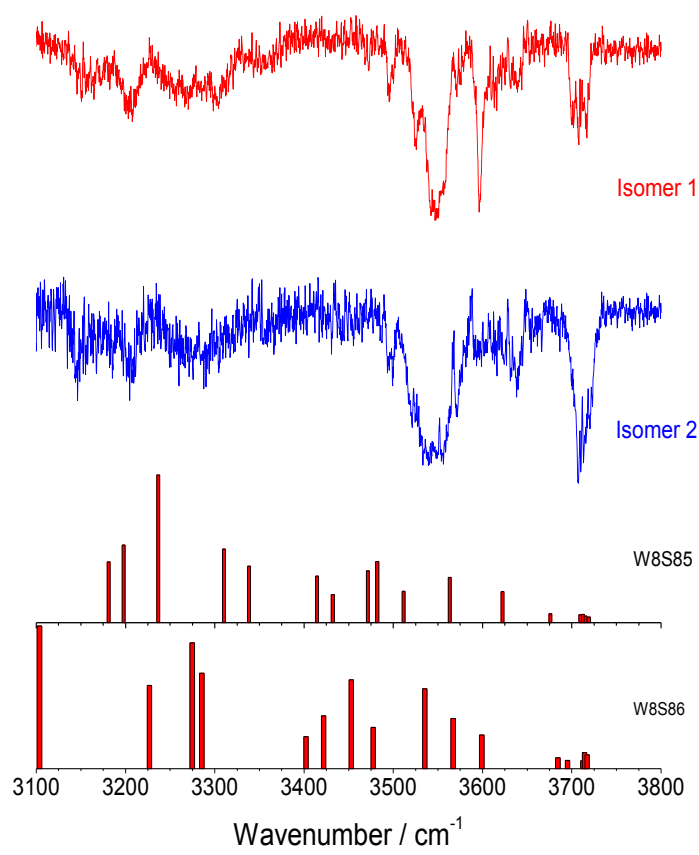


Figure S8. Experimental IDIRS for propofol₁(H₂O)₈ isomer 1 (upper trace) obtained in the fingerprint region, together with the predicted frequencies for the 11 lowest energy calculated structure. A correction factor of 0.938 was employed

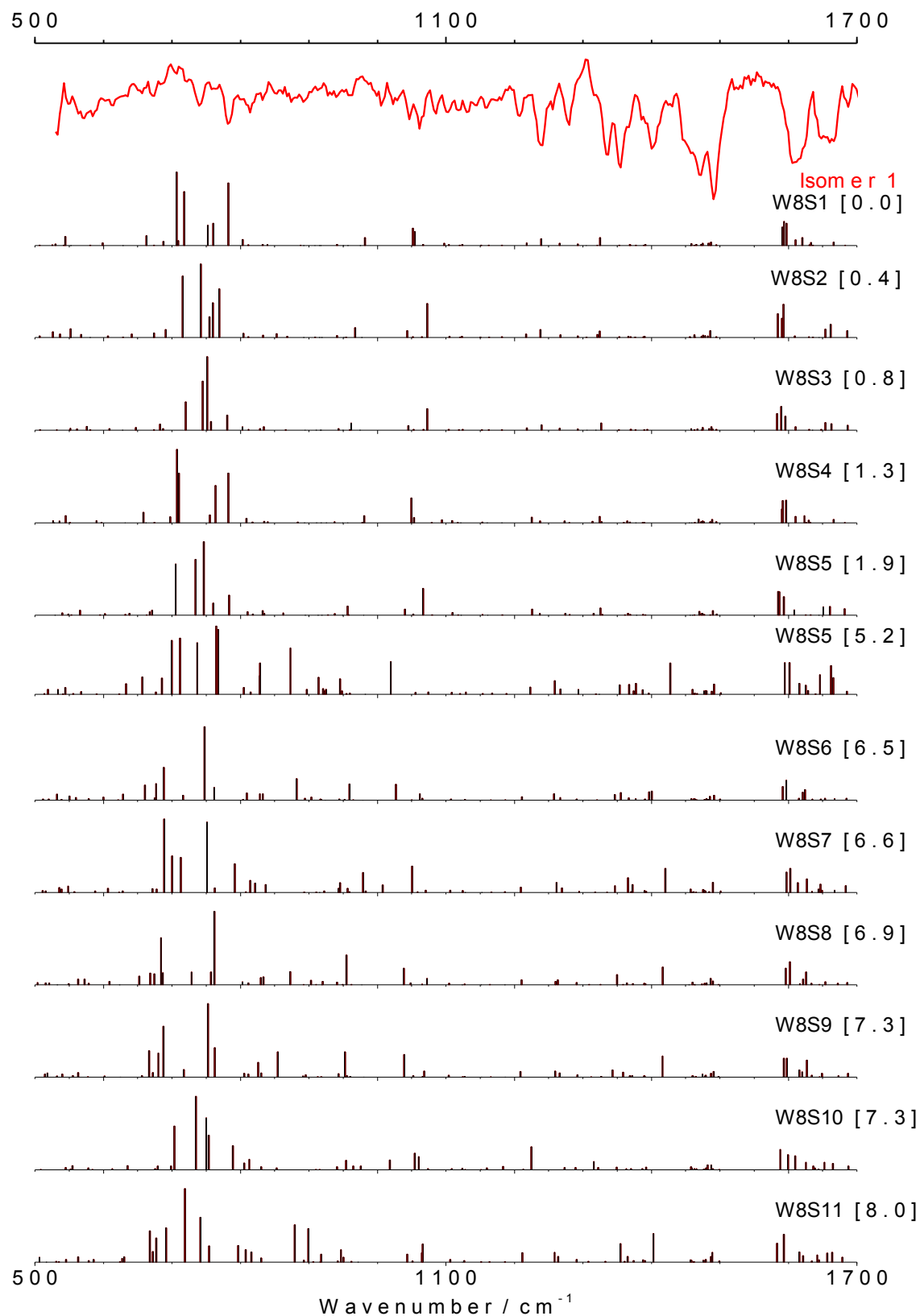


Figure S8. Calculated structures for propofol₁(H₂O)₉ at M06-2X/6-311++G(d,p) level, together with their relative stability in kJ/mol. ZPE correction was applied to all the energy values

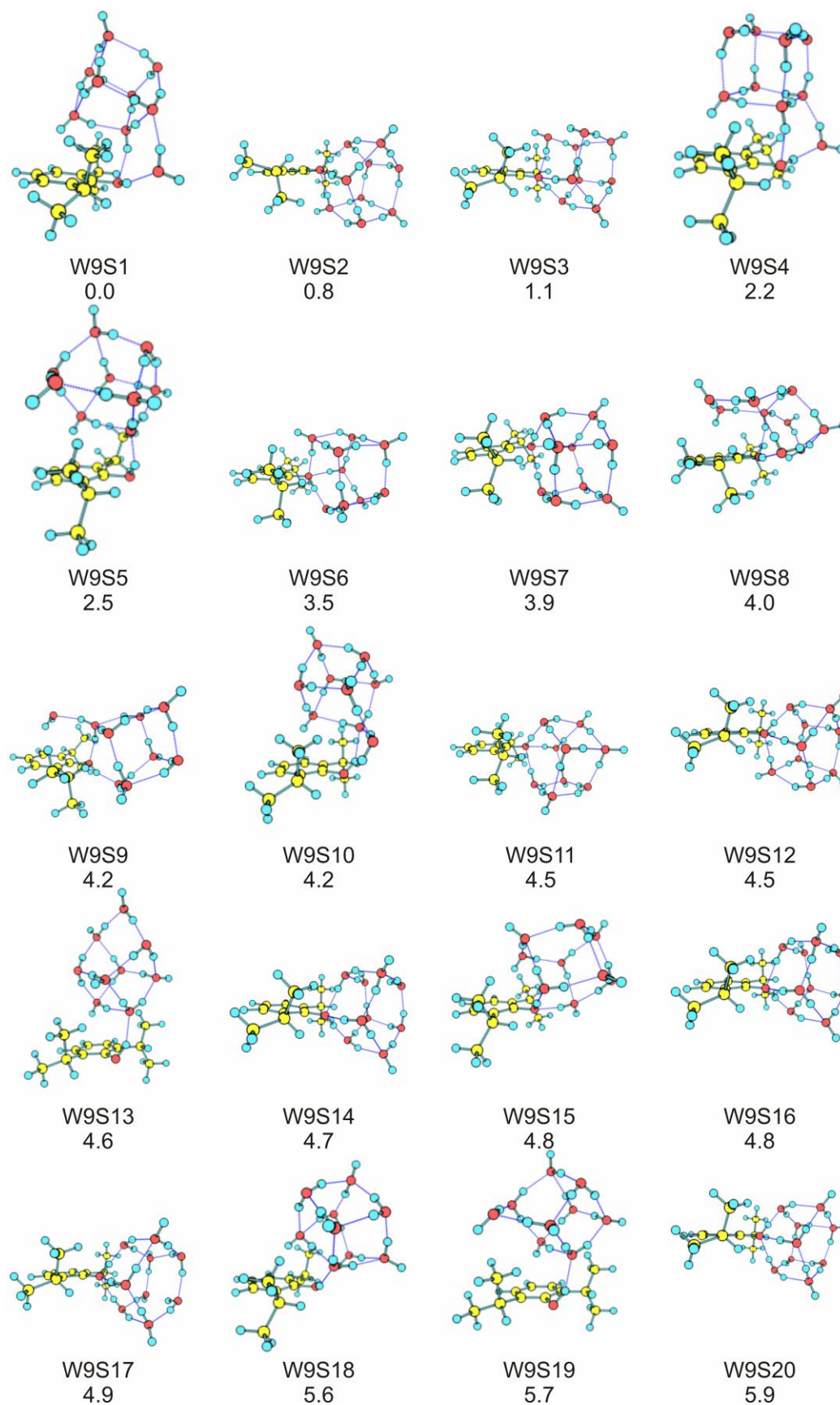


Figure S9. (Cont.)

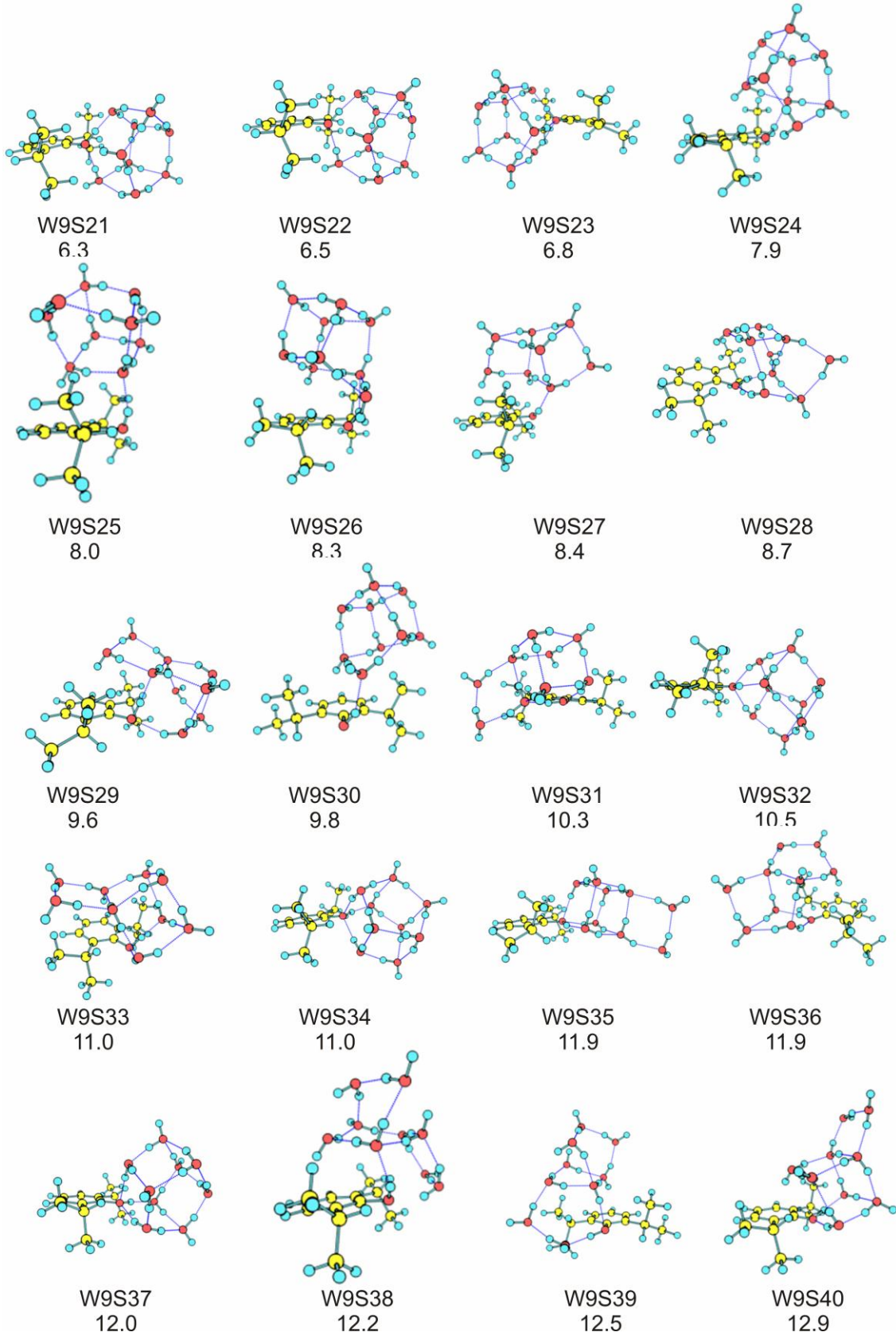


Figure S9. (Cont.)

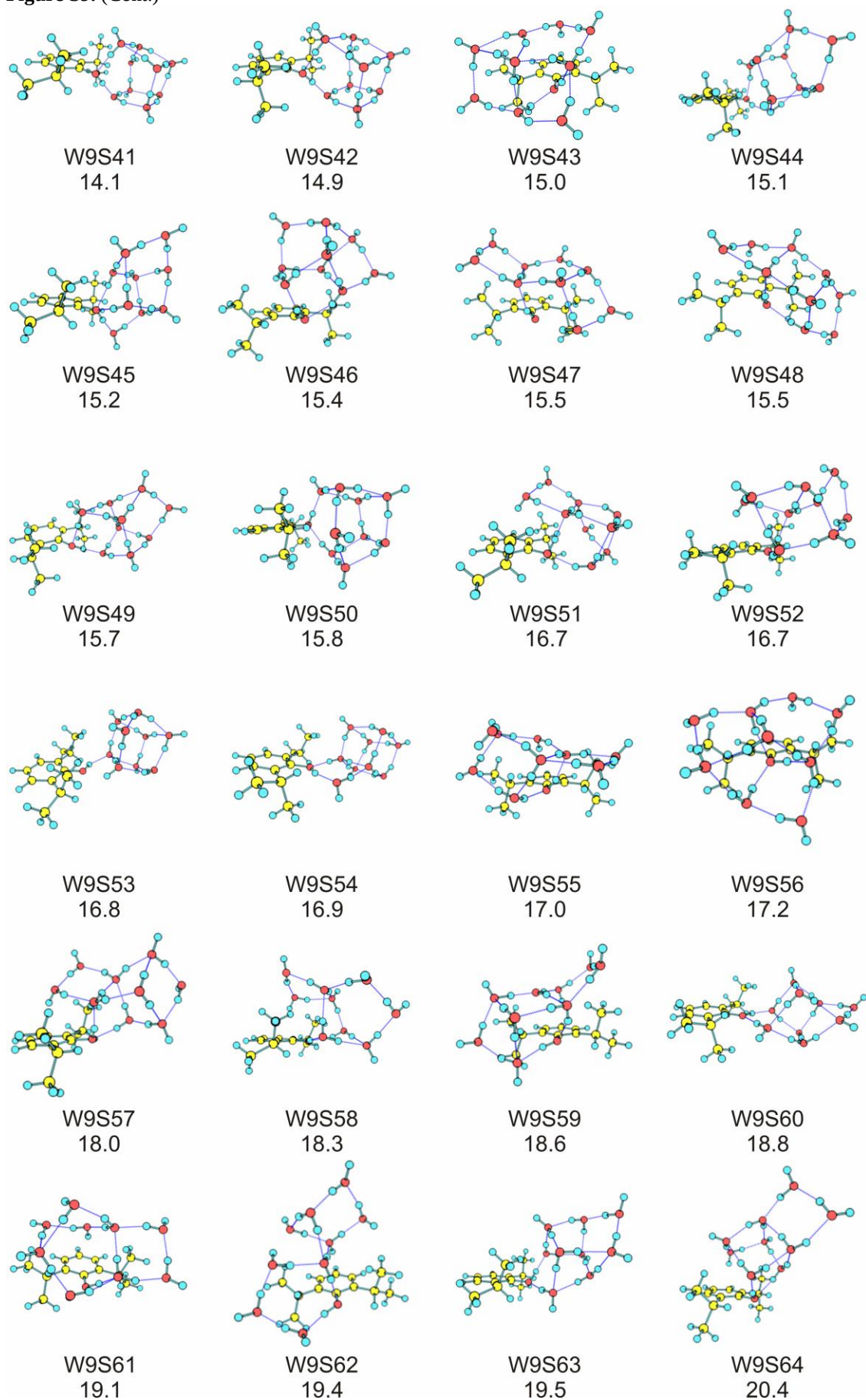


Figure S9. (Cont.)

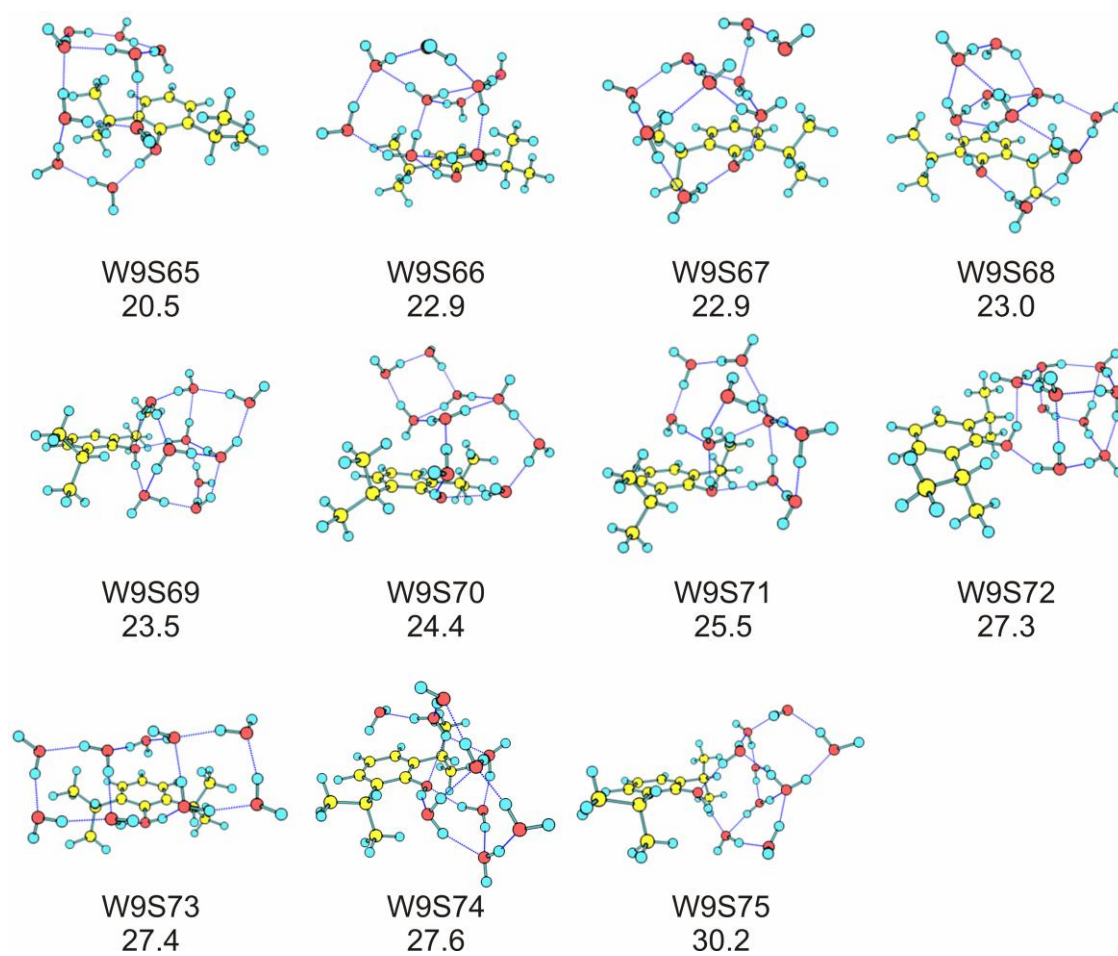


Figure S10. Experimental IDIRS in the fingerprint region for propofol₁(H₂O)₉ (upper trace) together with the predicted frequencies for the eight most stable calculated structures. A correction factor of 0.938 was employed.

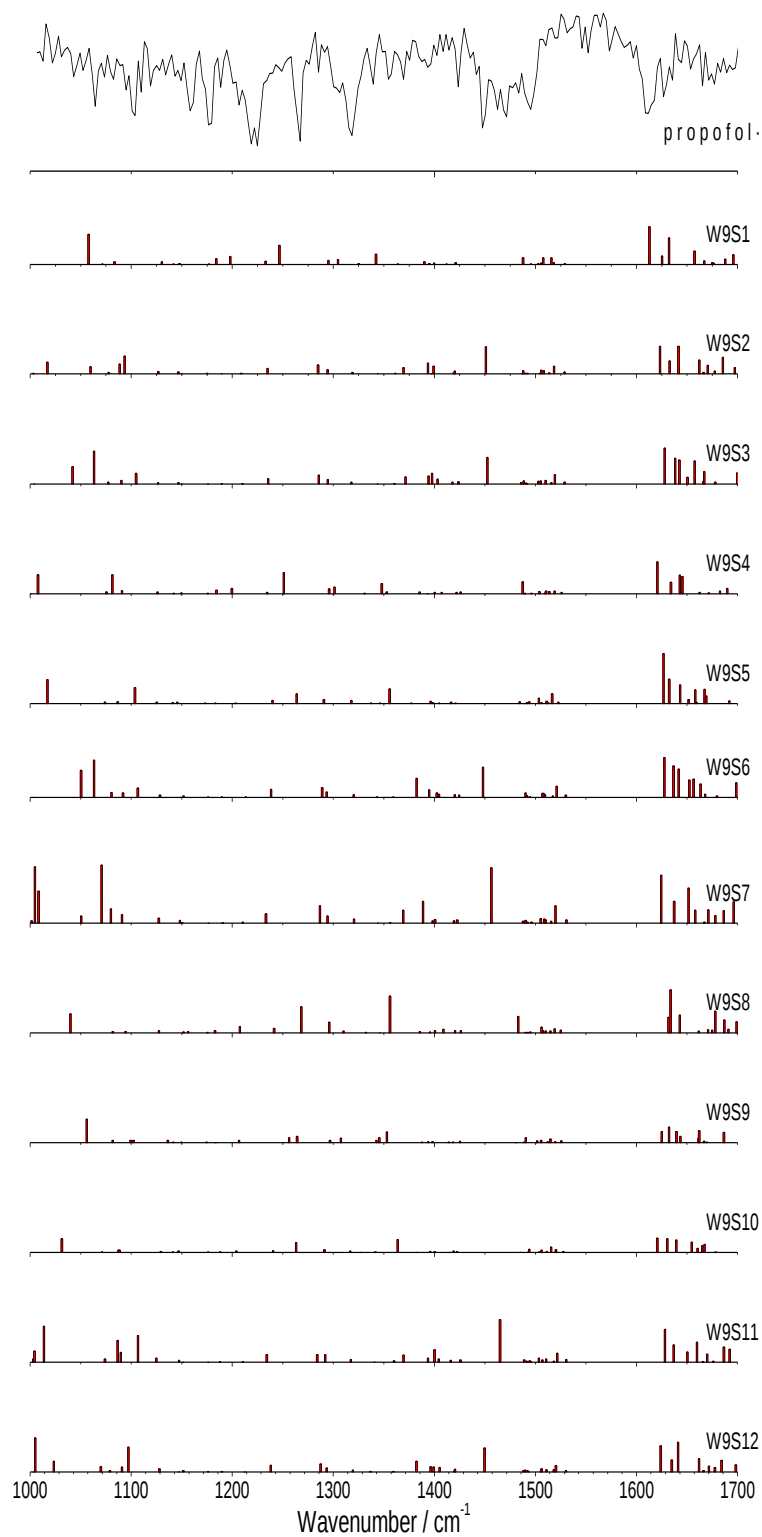


Figure S11. Experimental IDIRS for propofol₁(H₂O)₉ (upper trace) together with the predicted frequencies for each calculated structure. A correction factor of 0.938 was employed.

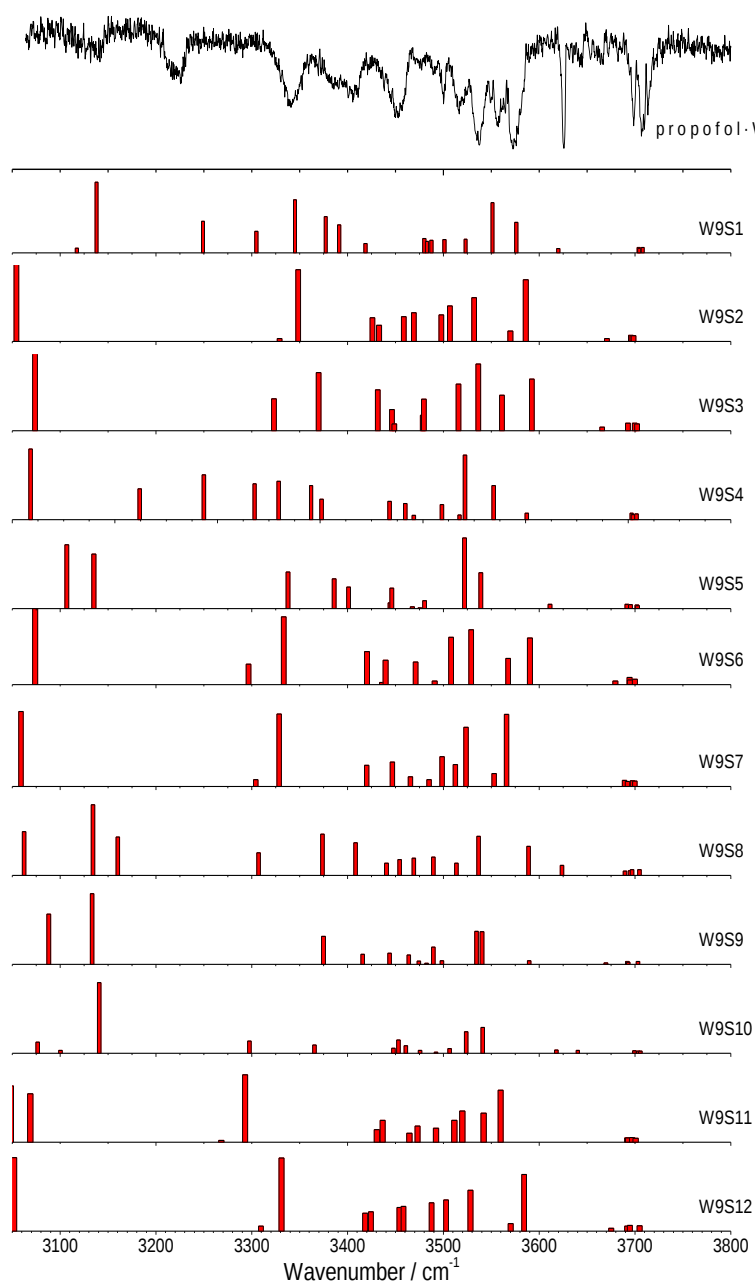


Figure S11. (Cont.)

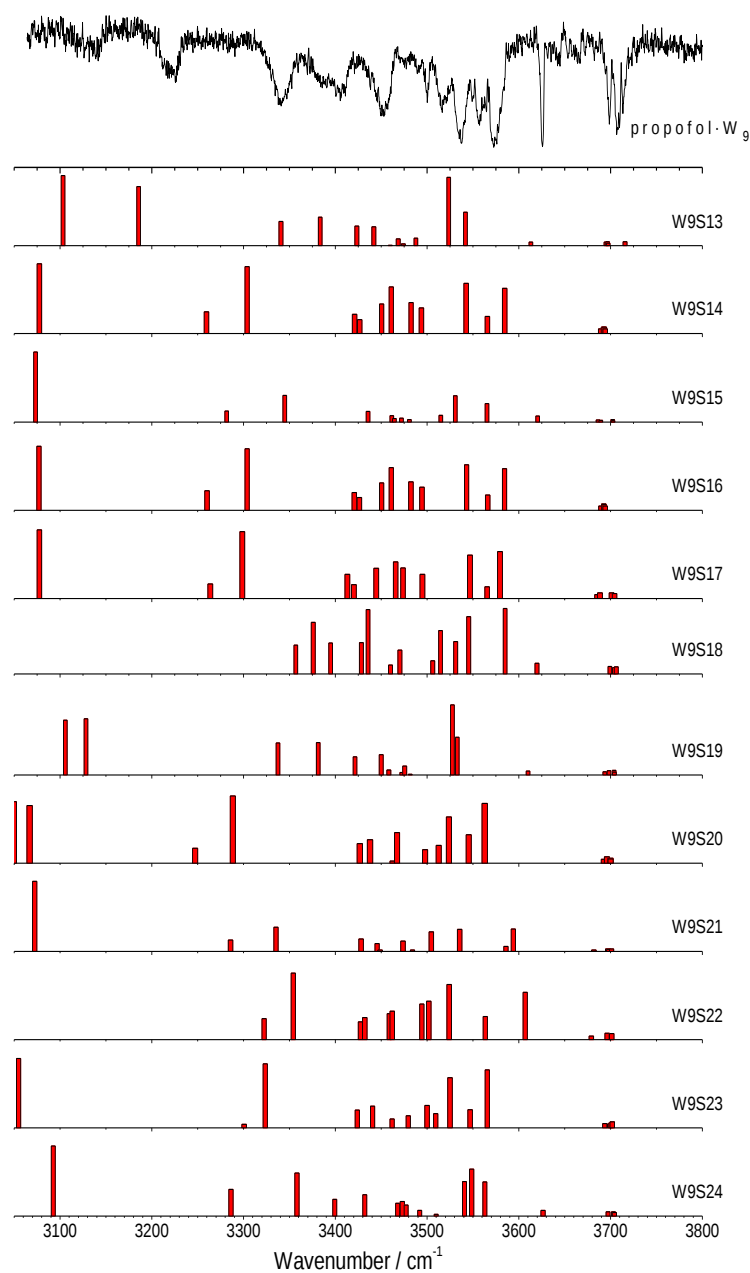


Figure S11. (Cont.)

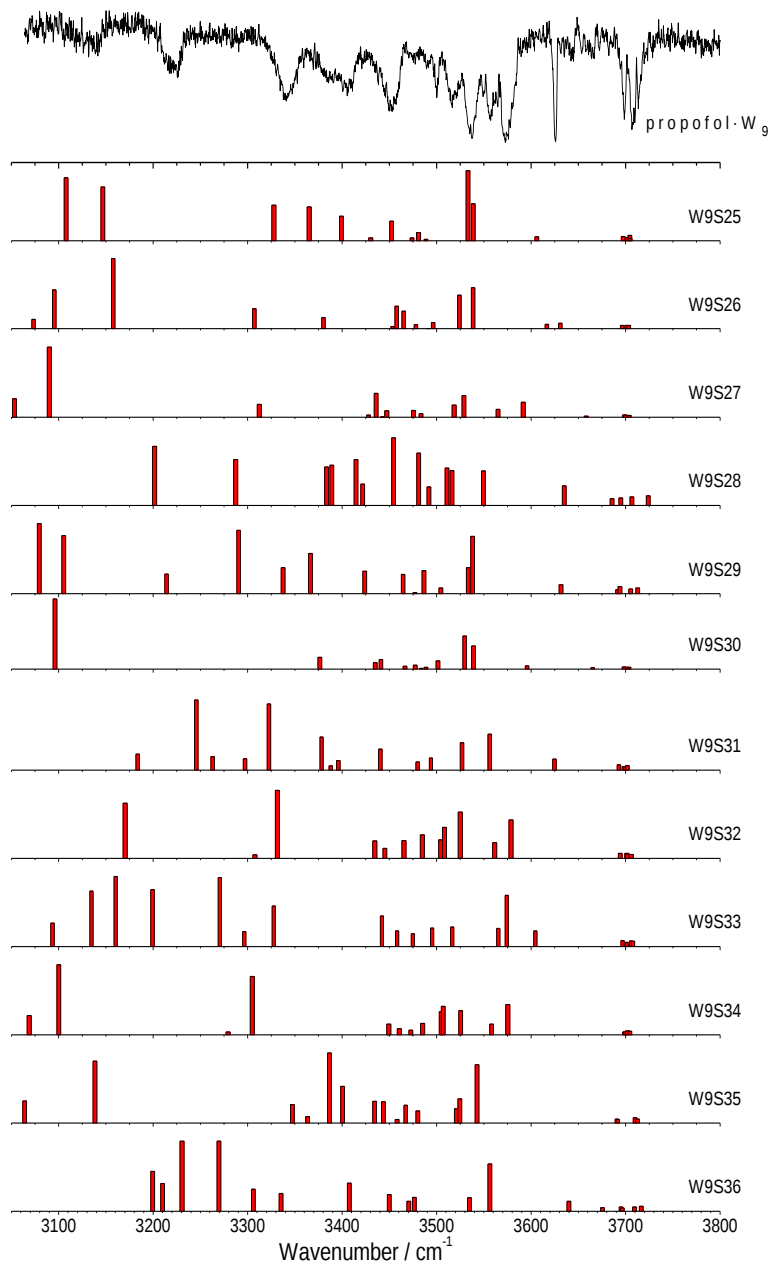


Figure S11. (Cont.)

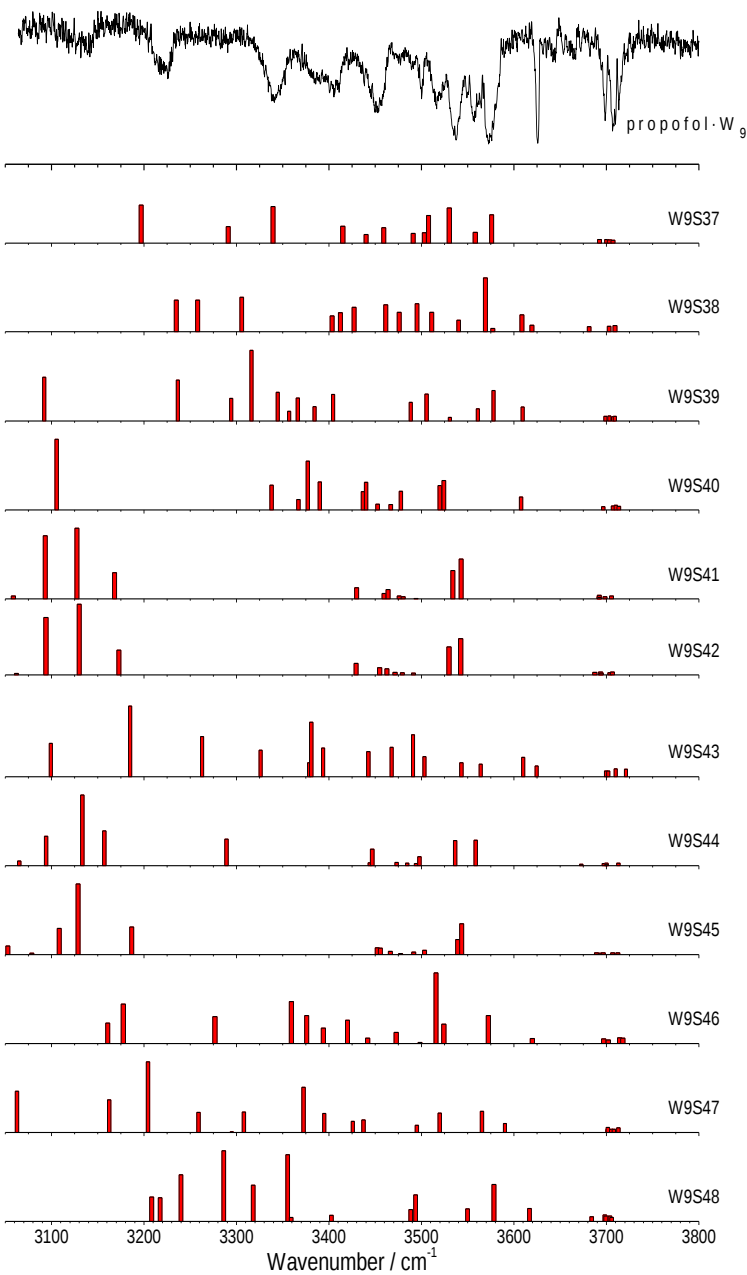


Figure S11. (Cont.)

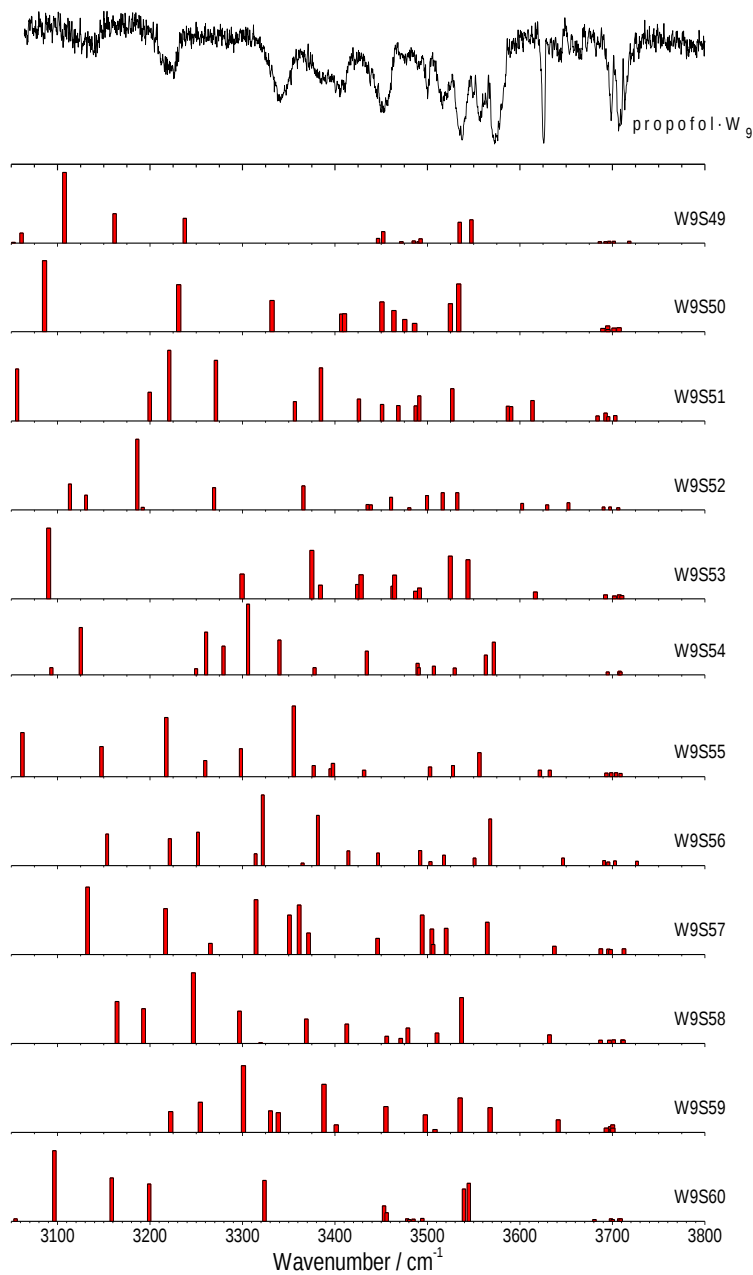


Figure S11. (Cont.)

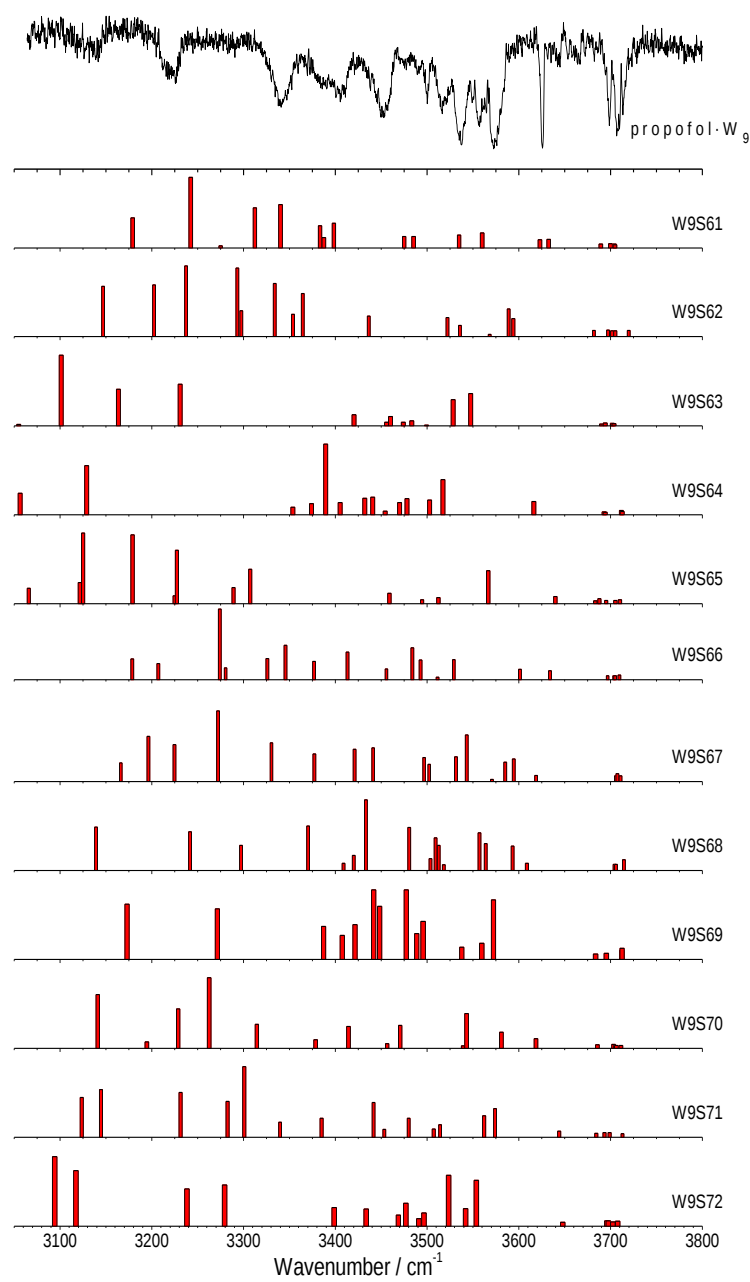
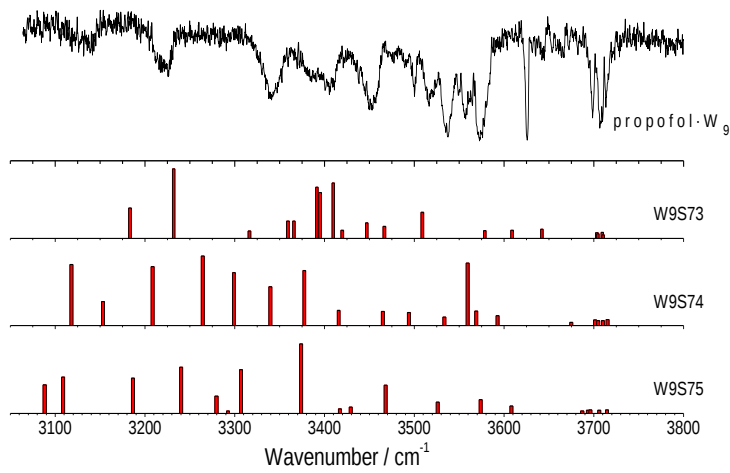
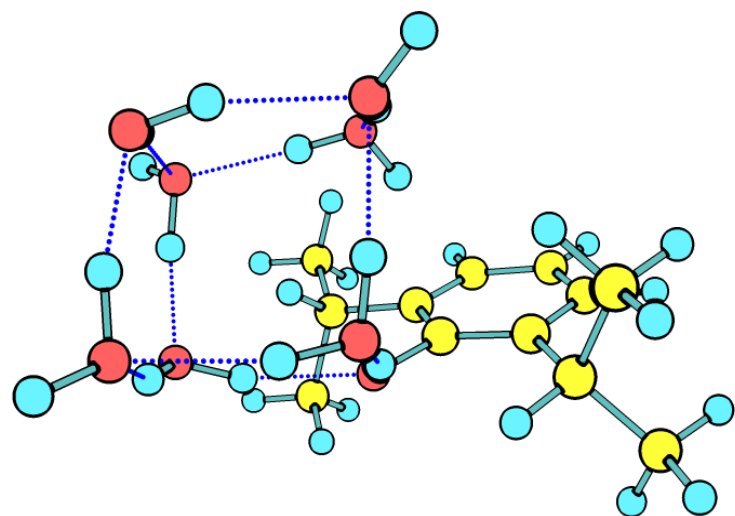


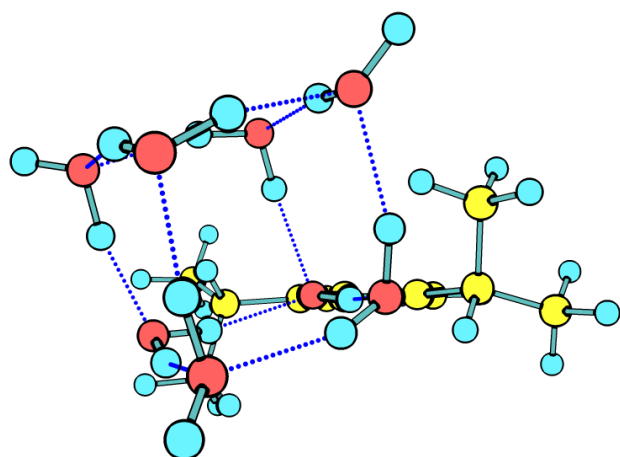
Figure S11. (Cont.)



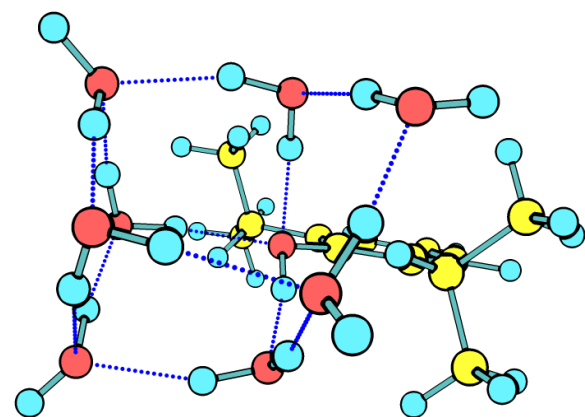
Structures assigned to the experimental conformers in xyz format. An image of the structure is also shown to assist in identification.



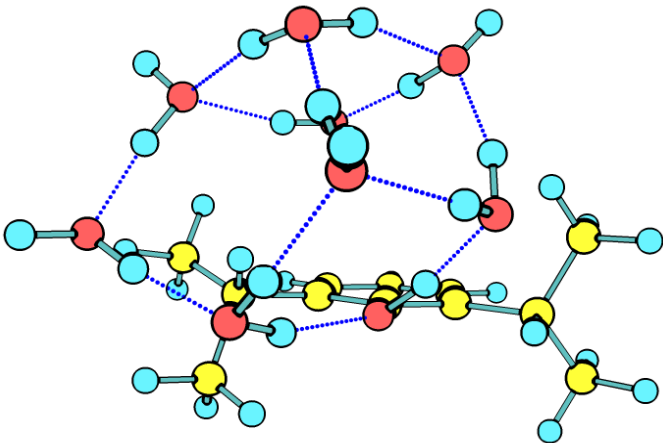
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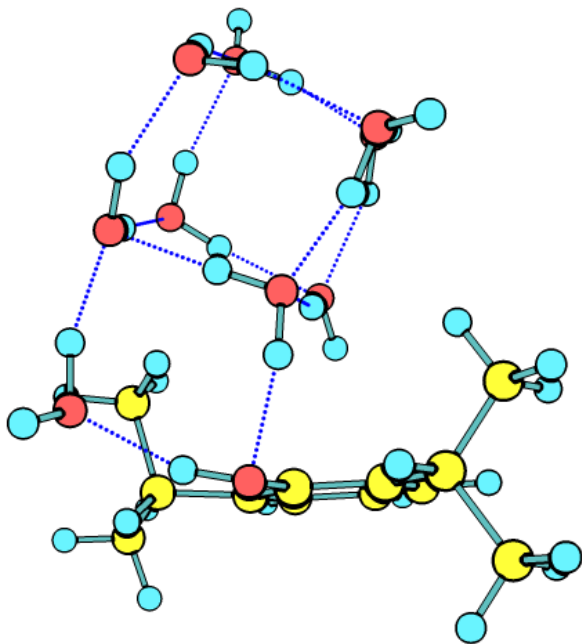
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1	3.994893000	-0.029318000	-0.256750000



6	-1.459006000	-2.659762000	0.797566000
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8	-0.312827000	2.734641000	-0.710184000
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1	0.557446000	2.984664000	-1.061215000
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