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Analysis of hydrogen bond energies and hydrogen bonded networks in water clusters \((\text{H}_2\text{O})_{20}\) and \((\text{H}_2\text{O})_{25}\) using the charge-transfer and dispersion terms

Suehiro Iwata*α

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The hydrogen bonds and their networks in the water clusters \((\text{H}_2\text{O})_{20}\) and \((\text{H}_2\text{O})_{25}\) are characterized using the charge-transfer \((E_{\text{CT}}^{W_a,W_d})\) and dispersion \((E_{\text{Disp}}^{W_a,W_d})\) terms for every pair of the water molecules \((W_a,W_d)\) in the clusters. The terms are evaluated by the perturbation theory based on the ab initio locally projected molecular orbitals (LPMO PT) developed by the present author. The relative binding energies among the isomers evaluated by the LPMO PT agree with those of the high levels of ab initio wave function based theories. A strong correlation between \(E_{\text{CT}}^{W_a,W_d}\) and \(E_{\text{Disp}}^{W_a,W_d}\) for the hydrogen bonded pairs are found. The pair-wise interaction energies are characterized by the types of the hydrogen-donor \((W_d)\) and hydrogen-acceptor \((W_a)\) water molecules. The strongest pair is that of the D2A1 water as a hydrogen-acceptor and the D1A2 water as a hydrogen-donor, where the \(\text{DnAm}\) water implies that the water molecule has \(n\) hydrogen bonding \(\text{O-H}\) and \(m\) accepting \(\text{H} \cdots \text{O}\). The intra-molecular deformation as well as the \(\text{O} \cdots \text{O}\) distance is also dependent on the types of hydrogen bonded pairs. The ring structures in the cluster are classified by the pattern of the alignment of the hydrogen bonds. The lengthening of the hydrogen-bonding \(\text{OH}\) of \(W_d\) is strongly correlated with the charge-transfer \((E_{\text{CT}}^{W_a,W_d})\) energy.

1 Introduction

The properties of hydrogen bonded networks in water clusters have been extensively studied both experimentally and theoretically. Xantheas and his coworkers reported a series of the high levels of computational works on finite sizes of water clusters, and they recently reviewed the works.1 The stable isomers of water clusters \((\text{H}_2\text{O})_n, n \geq 3\), contain the ring (cyclic) structures of the hydrogen bonded networks. The isomers of \((\text{H}_2\text{O})_n, n = 17 \sim 21\) reported by Lagutchenkov, Fanourgakis and Xantheas consist of 4-membered and 5-membered rings,2 but they do not have 6-membered rings. On the other hand, all of the isomers of \((\text{H}_2\text{O})_{25}\), reported by Furtado et al has 6-membered rings, and one of them has a 7-membered ring.3 It is well-known that the 6-membered ring structures are the main unit of various phases of water assemblies, such as liquid water, amorphous waters and crystal ice. In the finite sizes of stable water clusters, the 6-membered ring structures start to appear around \(n \sim 24\). Because every water molecule has two donating OH bonds and two O\(\cdots\)H-accepting cites, even a single 6-membered ring \((\text{H}_2\text{O})_6\) has several distinct isomers of different alignments of the hydrogen bonded molecules. McDonald et al4 counted the 30,026 distinct isomers of the dodecahedron \((\text{H}_2\text{O})_{20}\), which differ by the direction of the hydrogen bonds. Kirov reported that the the number of symmetry-distinct configurations emanating from the different hydrogen positions is 3,043,836 for 5\(^{12}\)6\(^2\) T-cage \((\text{H}_2\text{O})_{24}\) and 61,753,344 for 5\(^{12}\)6\(^4\) H-cage \((\text{H}_2\text{O})_{28}\).5,6

Last decade the author has been developing the ab initio molecular orbital theory suitable for the study on the weak molecular interaction.7–10 The most important target systems of these works are the hydrogen bonded clusters, in particular, of water clusters. In most of the ab initio computation of the non-covalent weak molecular clusters, by both wave function based and density functional theories, the supermolecule approach is commonly used. The computations can be carried out straightforwardly without the intuitive division of the target systems. The interaction energy is evaluated as the difference of large numbers. Because the interaction energy is much smaller than the total electron energy of the system, the well balanced approximation for the separated systems and for the assembled system becomes important. Because of the inconsistent approximation for the one-electron functions (molecular orbitals) and for the many-electron functions (electron configuration), the basis set superposition error (BSSE) sneaks in the binding energy. To avoid the BSSE, there are several attempts. Our approach is to use the local basis sets for the molecular orbitals of each component molecule. The method is called the Locally Projected Molecular Orbital

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(LPMO) perturbation theory (PT), and several applications are published.\textsuperscript{7–10} One of the advantages of the method is that the charge-transfer and dispersion terms between each pair of molecules can be directly evaluated from the first order perturbation wave function.

In the previous papers,\textsuperscript{11,12} using the charge-transfer terms between the hydrogen bonded pairs, we classified the water molecules by DnAm, where \( n \) is the number of hydrogen-donor OHs and \( m \) is the number of H atoms accepted by the molecule. The hydrogen bonds were characterized by DnAm→Dn′Am′, DnAm (Dn′Am′) being for the hydrogen acceptor (donor) water molecule. The similar characterizations are proposed by various papers. Ohno et al\textsuperscript{13} also classified the pair of the hydrogen bonded waters using the calculated harmonic frequency shifts and compared the observed vibrational spectra of water clusters. Xanthias\textsuperscript{14} defined (da), (aa), (dd), (da) and (aadd) for each water in the local hydrogen bonded network of \( n \leq 6 \). Singer and his coworkers introduced the graph invariants for the water clusters,\textsuperscript{4,15} and the second order graph invariants were used to fit the empirical binding energy. They noted that the energies of these isomers dependent on the number of the nearest-neighbor pairs of double acceptor waters in the clathrate structure.\textsuperscript{15}

The present work is the extension of the previous works\textsuperscript{11,12} to larger water clusters which have more ring structures in the clusters.

\section{Theoretical and Computational Procedure}

\subsection{The binding energy and its analysis}

A series of our papers documented the theoretical basis and the procedure to evaluate the binding energy and its terms in the perturbation theory based on the locally projected molecular orbitals (LPMO PT).\textsuperscript{9–11,16} Here, we describe the basic equations that pertain to the following discussion.\textsuperscript{12} In LPMO, the zero order wave function \( \Psi_{\text{LPMO}} \) for a molecular cluster is a single Slater determinant constructed from sets of local MOs. These MOs are canonical for each molecule but not for the whole cluster, and therefore, the first order wave function starts with the single excitations as

\[
\Psi_{\text{1ST}} = \sum_{\text{Mol}=X}^{X} |\text{LE}_{X}\rangle + \sum_{\text{Mol}=X,Y}^{X \neq Y} |\text{CT}_{X \rightarrow Y}\rangle + \sum_{\text{Mol}=X,Y}^{X \leq Y} |\text{Disp}_{X \rightarrow Y}\rangle
\]

where \( |\text{LE}_{X}\rangle \) stands for the single excitations within molecule \( X \), and \( |\text{CT}_{X \rightarrow Y}\rangle \) stands for the single excitations from molecule \( X \) to \( Y \), whereas \( |\text{Disp}_{X \rightarrow Y}\rangle \) stands for the double excitations of the dispersion type. This expansion is made possible by defining the locally projected excited MOs, most of which are expanded in terms of the basis sets on each molecule \( X \).\textsuperscript{7} Because of the orthogonality condition to the occupied MOs of the other molecules \( \{Y\} \), the coefficient vectors of some of excited MOs of molecule \( X \) have to be partially delocalized over the basis sets on the other molecules \( \{Y\} \). This restricted expansion can avoid the basis set superposition error (BSSE) caused both by the orbital basis inconsistency (OBI) and by the configuration basis inconsistency (CBI).\textsuperscript{9,16}

The calculated binding energy in this approximation can be written as

\[
\begin{align*}
E_{\text{BindE}}^{\text{2&3SPT+Disp}} &= \left( E_{\text{HF}}(\Psi_{\text{LPMO}}) - \sum_{X} E_{\text{HF}}^{X} \right) + \left( E_{\text{2&3SPT}} + E_{\text{3SPT}} \right) \\
&+ E_{\text{2DPT+Disp}} \equiv E_{\text{LPMO}} + E_{\text{2&3SPT}} + E_{\text{Disp}} \\
&= E_{\text{LPMO}} - E_{\text{BindE}}^{\text{2&3SPT}} + E_{\text{Disp}} \tag{2}
\end{align*}
\]

The first parenthesis in equation (2) represents the binding energy \( E_{\text{BindE}}^{\text{LPMO}} \) evaluated by LPMO, which contains the electrostatic, exchange-repulsion and induction (polarization) terms as well as the destabilization energy caused by the geometric deformation. The second parenthesis of equation (2) is the second and third order corrections of the single excitations \( E_{\text{CT+LE}}^{\text{2&3SPT}} \), and is evaluated by the sums

\[
E_{\text{CT+LE}}^{\text{2&3SPT}} = \sum_{\text{Mol}=X,Y}^{X \leq Y} \left( E_{\text{CT}}^{X \rightarrow Y} + E_{\text{CT}}^{Y \rightarrow X} \right) + \sum_{\text{Mol}=X}^{X \leq Y} E_{\text{Disp}}^{X} \tag{3}
\]

in which the contribution from the local excitations \( E_{\text{CT}}^{X} \) is non-zero only at the third order and is always much smaller than the charge-transfer terms. Because the terms in (5) are calculated by the first order wave function (1), each term is directly tabulated for any clusters consisting of any number of molecules when the sum \( E_{\text{CT+LE}}^{\text{2&3SPT}} \) is evaluated. This is contrasted with the ALMO-EDA (absolutely local MO - energy decomposition analysis) of Head-Gordon’s group, in which the CT energy is defined by the difference between the total energies of the HF (or Kohn-Sham DFT) determined with the full basis sets and with the ALMO.\textsuperscript{17,18} The Slater determinant of ALMO is equivalent to the zero order wave function \( \Psi_{\text{LPMO}} \) of our LPMO PT. The ALMO-EDA cannot separate the CT terms to each pair of the molecules when the interacting system consists of more than two molecules. The charge-transfer energy is very much definition-dependent, and therefore, the absolute values have to be used in the analysis with care. The dispersion terms is better defined than the charge-transfer terms because they result from the electron correlation of the inter-molecules. But for the molecular interaction where the orbital overlaps between molecules are significant, the distinction of the intra- and inter-molecule electron correlation becomes not clear.

We demonstrated for several types of the molecular interaction that, if the aug-cc-pVXZ type basis sets\textsuperscript{19} are used, the
binding energy $E_{\text{Bind}}^{\text{LPMO-2SPT}}$ evaluated with the third order single excitation wave function approximates the counterpoise (CP) corrected Hartree-Fock energy $E_{\text{Bind}}^{\text{HF}}$. Thus, $E_{\text{Bind}}^{\text{LPMO-2SPT+Disp}}$ in equation (4) is the “approximate BSSE-free HF + dispersion” energy.

2.2 Computational Details

The in-house ab initio MO program, MOLYX, was partially parallelized for the present study by OpenMP. The new revision enabled us to compute the energy, $E_{\text{Bind}}^{\text{LPMO-2SPT+Disp}}$, of (H$_2$O)$_{25}$ with the aug-cc-pVDZ set on the SGI UV2000 at the RCCS, Okazaki Research facilities of National Institutes of Natural Science (NINS). The load modules, compiled by the intel compiler with MKL, are available both for Mac OS X and for Linux on request.

The geometries of most of water clusters are those reported by Xantheas and his coworkers, and they are reviewed in a recent article, and are cited in the previous paper. They are optimized with MP2/aug-cc-pVDZ. The geometries of the isomers for (H$_2$O)$_{17}$ (H$_2$O)$_{18}$ (H$_2$O)$_{19}$ and (H$_2$O)$_{20}$ are those reported by Xantheas and his coworkers. More recently Xantheas extended the search of the low-lying isomers and the global minima of (H$_2$O)$_{20}$ and compared the energies of many isomers of the pentagonal dodecahedron. In addition, for (H$_2$O)$_{20}$, six isomers determined by Gadre and his coworkers are also studied. To find the isomers, they made combined use of temperature basin paving (TBA) procedure and molecular tailoring approach (MTA). Six isomers of (H$_2$O)$_{25}$ similarly determined by them are also studied.

The charge-transfer term $E_{\text{CT}}^{X,Y}$ for the pair of the hydrogen bonded water molecules (X, Y) is used to identify the hydrogen bond and its direction. Subsequently the hydrogen bonding matrix, defined by Miyake and Aida, is set up. By manipulating the matrix, the water molecules and the hydrogen bonds are classified, and then the $r$-membered ring structures in the clusters are identified.

3 Results and Discussion

Relative binding energy of (H$_2$O)$_{20}$ and (H$_2$O)$_{25}$

Tables 1 and 2 compare the relative binding energy of (H$_2$O)$_{20}$ and (H$_2$O)$_{25}$. Because the optimization procedures of the references are slightly differed from each other, the energies of two groups of isomers of (H$_2$O)$_{20}$ are compared separately. The most stable isomers of two groups, edge-sharing pentagon prism in the former and G20E in the latter, share the “almost” same geometric structure (The difference is described below). As is seen in Tables, the LPMO PT results agree well with the MP2 energies (Note that the energy unit is $kJ mol^{-1}$). The similar agreement was found for (H$_2$O)$_{16}$, (H$_2$O)$_{17}$ and (H$_2$O)$_{18}$ in the previous work. In general, the relative energies of $E_{\text{Bind}}^{\text{LPMO-2SPT+Disp}}$ are close to those of MP2/aug-cc-pVTZ. It is because HF/aug-cc-pVTZ is also nearly BSSE-free as $E_{\text{Bind}}^{\text{LPMO-2SPT+Disp}}$. The total binding energies (E_{\text{Bind}}^{\text{LPMO-2SPT+Disp}}) are $-853.3 kJ mol^{-1}$ for the prism edge-sharing isomer of (H$_2$O)$_{20}$, and $-1071.0 kJ mol^{-1}$ for the G25D isomer of (H$_2$O)$_{25}$. The average hydrogen bond energy is $-25.1 kJ mol^{-1}$ for the former, and $-25.5 kJ mol^{-1}$ for the latter. The largest discrepancy of $E_{\text{Bind}}^{\text{LPMO-2SPT+Disp}}$ from the MP2 energies is found for isomer G25D, which is a unique isomer having a 7-membered ring and no 6-membered rings. As shown in Figure 1, the other isomers of (H$_2$O)$_{25}$ studied in the present paper have 6-membered rings.

Here we have to recall the fact that there are numerous isomers which have the exactly same topological configuration of the oxygen atoms in the water clusters. Using the empirical potential functions, Kuo et al evaluated the binding energies of these 30,026 isomers of pentagonal dodecahedron (H$_2$O)$_{20}$, and the energy difference of the maximum and minimum isomers is more than 130 $kJ mol^{-1}$ in their empirical potential. Tokmachev et al also reported the binding energy of all of 30,026 isomers using the other empirical potential, and their maximum and minimum difference is 148 $kJ mol^{-1}$. Chibaia et al studied also the dodecahedron (H$_2$O)$_{20}$ isomers with the ab initio HF and DFT. By selecting six isomers of the dodecahedron (H$_2$O)$_{20}$ using the empirical total dipole moment as a measure, they optimized the isomers by HF/6-311G*, and evaluated the binding energy by B3LYP/6-311+G** without the BSSE correction. Their calculated energies range from -1149 to -880 $kJ mol^{-1}$ with B3LYP/6-311+G** and from

![Fig. 1] The schematic structures and hydrogen-bonded networks of (H$_2$O)$_{25}$. The arrowed chains show the 6- and 7-membered rings.

The color figures, which can be seen in the online version, show the 6- and 7-membered rings connected with the arrows.
Table 1 Comparison of the relative binding energy (/kJ mol⁻¹) of the isomers of (H₂O)₂₀

<table>
<thead>
<tr>
<th>Isomers</th>
<th>E₃SP1+Disp BindE ref(1) eq.(4)</th>
<th>MP2</th>
<th>MP2</th>
<th>MTA-MP2 a</th>
<th>MTA-MP2 a</th>
<th>MTA-MP2 a</th>
<th>R₄: R₅ c</th>
</tr>
</thead>
<tbody>
<tr>
<td>ref [1]</td>
<td>apVDZ  b</td>
<td>apVDZ  b</td>
<td>apVTZ  b</td>
<td>CBS d</td>
<td>apVTZ  b</td>
<td>apVDZ  b</td>
<td></td>
</tr>
<tr>
<td>Edge-sharing</td>
<td>0.0</td>
<td>0.0</td>
<td>0.0</td>
<td>0.0</td>
<td>0.0</td>
<td>0.0</td>
<td>(4)²(31)³(1111)²(5)²(41)(32)²</td>
</tr>
<tr>
<td>fused-cube</td>
<td>6.2</td>
<td>7.4(9.0)c</td>
<td>(10.5)c</td>
<td>6.5</td>
<td>4.9(5.8)</td>
<td>(7.9)</td>
<td></td>
</tr>
<tr>
<td>face-sharing</td>
<td>47.6</td>
<td>52.3(53.5)</td>
<td>(46.7)</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Dodecahedron</td>
<td>0.0</td>
<td>0.0</td>
<td>0.0</td>
<td>0.0</td>
<td>0.0</td>
<td>0.0</td>
<td>(4)²(31)³(1111)²(5)²(41)(32)²</td>
</tr>
<tr>
<td>ref [1]</td>
<td>apVDX  b</td>
<td>apVDX  b</td>
<td>apVTZ  b</td>
<td>CBS d</td>
<td>apVTZ  b</td>
<td>apVDX  b</td>
<td></td>
</tr>
<tr>
<td>G20A</td>
<td>10.5</td>
<td>7.1</td>
<td>2.1</td>
<td>6.7</td>
<td>6.3</td>
<td>(4)²(31)³(1111)²(5)²(41)(32)²</td>
<td></td>
</tr>
<tr>
<td>G20B</td>
<td>6.4</td>
<td>6.7</td>
<td>10.9</td>
<td>5.9</td>
<td>9.2</td>
<td>(4)²(31)³(1111)²(5)²(41)(32)²</td>
<td></td>
</tr>
<tr>
<td>G20C</td>
<td>8.0</td>
<td>7.2</td>
<td>7.5</td>
<td>7.5</td>
<td>15.5</td>
<td>(4)²(31)³(22)²(5)(41)(32)²</td>
<td></td>
</tr>
<tr>
<td>G20D</td>
<td>14.2</td>
<td>14.8</td>
<td>15.5</td>
<td>15.5</td>
<td>15.5</td>
<td>10.9</td>
<td></td>
</tr>
<tr>
<td>G20F</td>
<td>10.5</td>
<td>10.0</td>
<td>10.5</td>
<td>11.3</td>
<td>10.9</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>


Table 2 Comparison of the relative binding energy (/kJ mol⁻¹) of the isomers of (H₂O)₂₅

<table>
<thead>
<tr>
<th>Isomers</th>
<th>E₃SP1+Disp BindE ref(1) eq.(4)</th>
<th>MP2</th>
<th>MTA-MP2 a</th>
<th>MTA-MP2 a</th>
<th>MTA-MP2 a</th>
<th>R₄: R₅ : R₆ : R₇ d</th>
</tr>
</thead>
<tbody>
<tr>
<td>ref [1]</td>
<td>apVDZ  b</td>
<td>apVDZ  b</td>
<td>CBS c</td>
<td>apVTZ  b</td>
<td>apVDZ  b</td>
<td></td>
</tr>
<tr>
<td>G25A</td>
<td>0.0</td>
<td>0.0</td>
<td>0.0</td>
<td>0.0</td>
<td>0.0</td>
<td>(4)²(31)³(1111)²(5)²(41)(32)²(2111)²(33)²</td>
</tr>
<tr>
<td>G25B</td>
<td>-3.1</td>
<td>3.0</td>
<td>0.8</td>
<td>2.1</td>
<td>4.2</td>
<td>(4)²(31)³(1111)²(5)²(41)(32)²(2111)²(6)(51)²(42)²</td>
</tr>
<tr>
<td>G25C</td>
<td>1.9</td>
<td>4.1</td>
<td>1.3</td>
<td>2.5</td>
<td>5.4</td>
<td>(4)²(31)³(1111)²(5)²(41)(32)²(2111)²(6)(51)²(3111)²</td>
</tr>
<tr>
<td>G25D</td>
<td>-8.7</td>
<td>4.5</td>
<td>2.1</td>
<td>2.9</td>
<td>5.0</td>
<td>(4)²(31)³(1111)²(5)(41)(32)²(2111)² ;(61)²</td>
</tr>
<tr>
<td>G25E</td>
<td>-2.2</td>
<td>3.7</td>
<td>2.9</td>
<td>3.4</td>
<td>4.2</td>
<td>(4)²(31)³(1111)²(5)(41)(32)²(2111)²(6)(51)²(42)²</td>
</tr>
<tr>
<td>G25F</td>
<td>-0.5</td>
<td>9.4</td>
<td>6.3</td>
<td>7.5</td>
<td>10.0</td>
<td>(4)²(31)³(1111)²(5)(41)(32)²(2111)²(5)(31)²(33)²</td>
</tr>
</tbody>
</table>

a) reference [1] b) apVXZ = aug-cc-pVXZ c) The two-point extrapolation to "Complete Basis Set". d) See text for the definition.

-950 to -712 kJ mol⁻¹ with HF/6-311G*.

In the present study only one dodecahedron isomer was calculated, and as is shown in Table 1, this dodecahedron isomer is the least stable among ten isomers studied. This isomer is not necessarily the most stable among the 30,026 distinct isomers of dodecahedron. Because Xantheas and his coworkers performed the extensive sampling of the isomers using TIP4P potential, the dodecahedron isomer they obtained is expected to be one of the most stable isomers. Recently Xantheas extended the computational studies of the pentagonal dodecahedron (H₂O)₂₀ isomers, and compared the binding energies of the lowest 20 isomers obtained with various empirical potentials and with the DFT and MP2 calculations.

As was reported by Xantheas and his coworkers, and as is also seen in Table 1, the other isomers of (H₂O)₂₀ are more stable than the dodecahedron. The number of hydrogen bonds in the cluster, which is given at the second column of Table 3, is the smallest for dodecahedron isomers (30) among the isomers studied. But it is not the number of hydrogen bonds that determines the order of the stability. Figure 2 shows the terms in equations (3) and (4). The dispersion term (the rightest column for each isomer) of the dodecahedron isomer is distinctly smaller than that of the other isomers. Similarly in (H₂O)₆, the dispersion term for the cyclic-chair isomer is much smaller than that for the other isomers such as cage and prism forms of the isomers. Without the dispersion term, the cyclic isomer is the most stable. Because of the long-range nature of the dispersion interaction, the substantial dispersion terms are found between the non-neighboring water molecules in the cage and prism isomers of (H₂O)₆. The dispersion interaction between non-neighboring molecules in the more compact forms of the isomers of (H₂O)₂₀ contributes to the total binding energy. The dodecahedron isomer studied in this work has another characteristic: the largest E_CT and smallest E₃SP1+Disp BindE; the latter contains the deformation energy caused by forming the strict regular pentagons. The two terms are inter-related to each other; the water molecules are deformed to favor the CT interaction in twelve 5-membered rings of the dodecahedron.
Table 3 Some characteristic indexes of the isomers of (H₂O)₂₀ and (H₂O)₂₅

<table>
<thead>
<tr>
<th>isomers</th>
<th>HB</th>
<th>N&lt;sub&gt;DIA2&lt;/sub&gt;,N&lt;sub&gt;D2A1&lt;/sub&gt;,N&lt;sub&gt;D2A2&lt;/sub&gt;</th>
<th>D2A1←D1A2&lt;sup&gt;c&lt;/sup&gt;</th>
<th>D2A2←D2A2&lt;sup&gt;c&lt;/sup&gt;</th>
<th>D1A2←D2A1&lt;sup&gt;c&lt;/sup&gt;</th>
<th>r&lt;sup&gt;v&lt;/sup&gt;, r = 4 ~ 7&lt;sup&gt;d&lt;/sup&gt;</th>
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<tbody>
<tr>
<td>edge-sharing</td>
<td>34</td>
<td>[6, 6, 8]</td>
<td>6</td>
<td>2</td>
<td>6</td>
<td>4&lt;sup&gt;12&lt;/sup&gt;, 5&lt;sup&gt;6&lt;/sup&gt;</td>
</tr>
<tr>
<td>fused-cube</td>
<td>36</td>
<td>[4, 4, 12]</td>
<td>4</td>
<td>20</td>
<td>4</td>
<td>4&lt;sup&gt;21&lt;/sup&gt;</td>
</tr>
<tr>
<td>face-sharing</td>
<td>35</td>
<td>[5, 5, 10]</td>
<td>6</td>
<td>2</td>
<td>6</td>
<td>4&lt;sup&gt;13&lt;/sup&gt;, 5&lt;sup&gt;4&lt;/sup&gt;</td>
</tr>
<tr>
<td>dodecahedron</td>
<td>30</td>
<td>[10, 10, 0]</td>
<td>17</td>
<td>0</td>
<td>7</td>
<td>5&lt;sup&gt;12&lt;/sup&gt;</td>
</tr>
<tr>
<td>G20E: edge-sharing</td>
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a) The number of hydrogen bonds in the cluster. b) N_DnAm: the number of water molecules of DnAm type. c) The number of hydrogen bonds between the hydrogen acceptor DnAm and hydrogen donor Dn’Am’. d) r<sup>v</sup>: the number of r-membered rings in the cluster.

configuration.

In Table 3, the number of water molecules N_DnAm and the number of the hydrogen bonded pairs DnAm ← Dn’Am’ are given. They take various values both for (H₂O)₂₀ and (H₂O)₂₅. The number of dangling bonds of these clusters is equal to N_DIA2. The last column of Table 3 shows the number of r-membered ring structures {n<sub>r</sub>, r = 4 ~ 7} as r<sup>v</sup>. The dodecahedron (H₂O)₂₀ has twelve 5-membered rings, while the edge-sharing prism isomer (H₂O)₂₀ has twelve 4-membered rings and six 5-membered rings. While counting the number of 6-membered rings, a little care is needed; when two neighboring 6-membered rings share four water molecules, there is another 6-membered ring. The examples are found in isomers G25B and G25E.

The r-membered rings are classified by the directions of the hydrogen bond chains in the ring. For instance, for the 4-membered hydrogen bonded rings, there are three ways forming the rings of waters as:

(4) : W₁ ← W₂ ← W₃ ← W₄ ← W₁
(31) : W₁ ← W₂ ← W₃ → W₄ ← W₁
(1111) : W₁ ← W₂ → W₃ ← W₄ → W₁

The index R<sub>r</sub>, at the last column of Tables 1 and 2 shows the numbers of the ring types in the network. For example, R<sub>4</sub> of the edge-sharing prism isomer is (4) (31) (1111)<sup>3</sup>, which implies that there are five 4-membered rings of type (4), four 4-membered rings of type (31) and three 4-membered rings of type (1111). The geometric structure of isomer G20E of Gadre’s group is almost identical with that of edge-sharing prism except that R<sub>4</sub> of G20E is (4) (31) (1111)<sup>3</sup>. The types of the ring structure in R<sub>r</sub> are closely related to the stability of the clusters. The geometric characteristics of 20 isomers of the pentagonal dodecahedron (H₂O)₂₀ isomers studied by Xantheas are not described in the paper. It should be examined how the relative binding energy of these isomers is related with the index R<sub>r</sub>. More systematic computations for large clusters are required to derive some general rules for the stability of the ring types in the clusters. In smaller cyclic isomers of (H₂O)₂₀, r = 3 ~ 6, the most stable cyclic isomers have the hydrogen bonded chain, aligned to the same direction as type (r) in the above definition. Inside of the large clusters, as seen in Figure 1, the rings are nested to each other. Therefore, not all of the hydrogen bonds are able to align in the same direction for some of the rings.

Interestingly, all of the isomers of (H₂O)₂₅ of Gadre’s<sup>3</sup> have the same numbers of hydrogen bonds, N_DIA2, N_D2A1, and N_D2A2, even though, without any guiding principle for finding the stable configurations, the optimization started from the structures determined by the temperature basin paving (TBA) procedure. As Figure 1 and the last column of Tables 2 and 3 show, however, the hydrogen bonded networks in these isomers are very different from each other. Their total binding energies are close to each other (within 10 kJ mol⁻¹). It seems that TBA of Shanker and Bandyopadhay is an efficient procedure to find many candidates of the stable conformers as they claimed.<sup>21</sup>

Analysis of Hydrogen Bond Energies

Figure 3 is the plots of the charge-transfer E<sub>CT</sub><sup>XY</sup> and dispersion terms E<sub>Disp</sub><sup>XY</sup> as a function of the the O⋯O length for every
pair of the hydrogen bonds in $(\text{H}_2\text{O})_{17} \sim (\text{H}_2\text{O})_{21}$. The different symbols are used for the types of the pair of hydrogen-acceptor and hydrogen-donor waters. The strongest type of the hydrogen bonds is $\text{D}2\text{A}1 \leftrightarrow \text{D}2\text{A}2$ (an upper-ward triangle mark), and almost all of the short $\text{O} \cdots \text{O}$ distance ($\leq 2.7\,\text{Å}$) hydrogen bonds are of this type. The strong correlation between $E_{\text{CT}}^{\text{XY}}$ and $E_{\text{Disp}}^{\text{XY}}$ can be noticed, in particular for this type of the hydrogen bonds. This correlation is more clear in Figure 4, in which the direct correlation between $E_{\text{CT}}^{\text{XY}}$ and $E_{\text{Disp}}^{\text{XY}}$ is shown in Figure 3 Correlation of the $\text{O} \cdots \text{O}$ length with the charge-transfer $E_{\text{CT}}^{\text{XY}}$ and dispersion $E_{\text{Disp}}^{\text{XY}}$ energy for the isomers of $(\text{H}_2\text{O})_{17} \sim (\text{H}_2\text{O})_{21}$ optimized by Xantheas and his coworkers. The different symbols are used, depending on the types of the pair of hydrogen-acceptor and hydrogen-donor waters. The filled points are for $E_{\text{Disp}}^{\text{XY}}$.

Figure 4b) shows that in smaller clusters the strong $\text{D}2\text{A}1 \leftrightarrow \text{D}2\text{A}2$ (a upper-ward triangle mark) and weak $\text{D}1\text{A}2 \leftrightarrow \text{D}2\text{A}1$ (a rhombus mark) bonds are dominant. They together often make the chains in the cluster. The $\text{D}2\text{A}2 \leftrightarrow \text{D}2\text{A}2$ bonds in this figure are mostly of isomers $(\text{H}_2\text{O})_{16}$, as seen in Table 3 and in Figure 4c), there are many $\text{D}2\text{A}2 \leftrightarrow \text{D}2\text{A}2$ hydrogen bonds in $(\text{H}_2\text{O})_{20}$ and $(\text{H}_2\text{O})_{25}$, and they contribute to the stability of the clusters. The approximate linear relation of $E_{\text{CT}}^{\text{XY}}$ and $E_{\text{Disp}}^{\text{XY}}$ for this type bond can be noticed in Figure 4. Also it should be emphasized that the dispersion term is dominant in the $\text{D}2\text{A}2 \leftrightarrow \text{D}2\text{A}2$ hydrogen bonds, which are the bonds that construct the hydrogen bonded networks in large water clusters as well as in liquid waters and ice. Figures 3 and 4 demonstrate that the interaction energies and the $\text{O} \cdots \text{O}$ lengths depend on the pair of the hydrogen-donor and -acceptor waters, but that their values are distributed in broad ranges; the neighboring hydrogen bonds influence the hydrogen bonds electronically and by the structural constraints. The non-additive and many-body effects, that are not fully accounted for in the classification by the type of the hydrogen bonded pairs $\text{DnAm} \leftrightarrow \text{Dn’Am}$, cause

Fig. 2 The energy analysis of the isomers of $(\text{H}_2\text{O})_{20}$. The total binding energy $E_{\text{BindE}}$ (Note that the right ordinate and its scale differ from those of the left ordinate for the columns). The first of four columns for each isomer is the first term, $E_{\text{LPMO}}^{\text{XY}}$, of equation (3). The second column is the second term, $E_{\text{CT}}^{\text{XY}}+E_{\text{Disp}}^{\text{XY}}$, of equation (3). The third is the sum of these two terms, $E_{\text{LPMO}}^{\text{XY}}-2E_{\text{CT}}^{\text{XY}}+E_{\text{Disp}}^{\text{XY}}$, and is the first term of equation (4). The last column is the dispersion term, $E_{\text{Disp}}^{\text{XY}}$.

Fig. 3 Correlation of the $\text{O} \cdots \text{O}$ length with the charge-transfer $E_{\text{CT}}^{\text{XY}}$ and dispersion $E_{\text{Disp}}^{\text{XY}}$ energy for the isomers of $(\text{H}_2\text{O})_{17} \sim (\text{H}_2\text{O})_{21}$ optimized by Xantheas and his coworkers. The different symbols are used, depending on the types of the pair of hydrogen-acceptor and hydrogen-donor waters. The filled points are for $E_{\text{Disp}}^{\text{XY}}$. 

$E_{\text{Disp}}^{\text{XY}}$ has eight (1111) 4-membered rings. By changing the direction of the hydrogen bonds in these rings, more stable fused-cube isomers than that studied might be found. The dodecahedron isomer studied has only one (2111) 5-membered ring, which suggests that this might be one of the most stable among the 30,026 dodecahedrons.

Figure 4b) shows that in smaller clusters the strong $\text{D}2\text{A}1 \leftrightarrow \text{D}2\text{A}2$ (a upper-ward triangle mark) and weak $\text{D}1\text{A}2 \leftrightarrow \text{D}2\text{A}1$ (a rhombus mark) bonds are dominant. They together often make the chains in the cluster. The $\text{D}2\text{A}2 \leftrightarrow \text{D}2\text{A}2$ bonds in this figure are mostly of isomers $(\text{H}_2\text{O})_{16}$, as seen in Table 3 and in Figure 4c), there are many $\text{D}2\text{A}2 \leftrightarrow \text{D}2\text{A}2$ hydrogen bonds in $(\text{H}_2\text{O})_{20}$ and $(\text{H}_2\text{O})_{25}$, and they contribute to the stability of the clusters. The approximate linear relation of $E_{\text{CT}}^{\text{XY}}$ and $E_{\text{Disp}}^{\text{XY}}$ for this type bond can be noticed in Figure 4. Also it should be emphasized that the dispersion term is dominant in the $\text{D}2\text{A}2 \leftrightarrow \text{D}2\text{A}2$ hydrogen bonds, which are the bonds that construct the hydrogen bonded networks in large water clusters as well as in liquid waters and ice. Figures 3 and 4 demonstrate that the interaction energies and the $\text{O} \cdots \text{O}$ lengths depend on the pair of the hydrogen-donor and -acceptor waters, but that their values are distributed in broad ranges; the neighboring hydrogen bonds influence the hydrogen bonds electronically and by the structural constraints. The non-additive and many-body effects, that are not fully accounted for in the classification by the type of the hydrogen bonded pairs $\text{DnAm} \leftrightarrow \text{Dn’Am}$, cause
a certain distribution of the plots in Figures.

Figure 4c) for (H₂O)₁₀ and (H₂O)₁₅ of Gadre’s group more clearly shows that the relation between $E_{CT}^{X,Y}$ and $E_{Disp}$ in the D2A1$\leftrightarrow$D1A2 bonds is different from that in the other types of hydrogen bonded pairs. The dispersion terms for the D2A1$\leftrightarrow$D1A2 bonds are smaller than those for D2A2$\leftrightarrow$D2A2 bonds. Because the D1A2 water molecule has a dangling bond, the molecule is at the surface of the cluster. Figure 5 shows the dependence of $E_{CT}^{X,Y}$ and the O-H on the O...O length in the isomers of (H₂O)₁₀ and (H₂O)₁₅. Where $H_b$ is the hydrogen bonding hydrogen. The plots clearly show the lengthening of O-H bonds by the hydrogen bond formation. This lengthening is related to the well-known experimental and theoretical observation in the low-frequency shifts of the harmonic frequency of the OH stretching modes. The lengthening of the O-H bond is positively correlated with $E_{CT}^{X,Y}$, and depends on the types of the hydrogen bonds.

Figure 6 are the plots for the OH lengths of the water molecules. The abscissa is the sum $\Sigma_{Y\neq X} E_{CT}^{X,Y}$ for molecule X of the DnAm type. When X is the D1 type water, a single term dominates in the sum $\Sigma_{Y\neq X} E_{CT}^{X,Y}$. The large value of the sum (left-ward at the abscissa) implies that water X is a good hydrogen donor and electron acceptor. Understandably the length of the dangling OH (unfilled down-ward triangles) of the D1A2 water is independent of the sum (the abscissa). It can be noticed that these lengths of the dangling OH in Figure 6a and b are slightly different by about 0.002 Å. It is because the geometries come from the different sources. The hydrox bonding OH (filled down-ward triangles) of the D1A2 water is substantially lengthened, and the weakening of the bond depends on the charge-transfer energy as the hydrogen donor, but the change is not linear. In a simple orbital theory of charge-transfer interaction, the $\sigma^*$ anti-bonding orbital of OHb is an electron acceptor orbital. Both of Figure 6a and b clearly demonstrate the close correlation of the charge-transfer energy and the OHb lengthening. The amount of the charge-transfer, and, in other words, the weakening of the OH bond, is sensitive to the surrounding geometric constraints. The plots in Figure 6 are not able to show the details of the causes. Figures show that both in D2A2 and D2A1 waters, two OHb bonds are not equivalent in most of cases; the longer one of each molecule is shown by the filled mark. In figures the square points for D2A2 and the circle points of D2A1 are located at the abscissa less than twice of those (down-ward triangles) of D1A2, which implies that the lengthening of the OH bonds is not a simple linear function of the charge-transfer interaction.

These plots are another demonstration that $E_{CT}^{X,Y}$ in the present definition is a good measure to characterize the strength of the hydrogen bonds and hydrogen bonded networks.

**Concluding Remark**

The work demonstrates that the LPMO PT is a practical and reasonably accurate ab initio MO method to study the molecular interaction, and can be applied for a large size of water clusters without a big computer resource. Importantly the method provides us not only the binding energy but also the pair-wise interaction terms, which help to characterize the interaction energy pair-wisely. The water molecules in the water clusters are classified by DnAm, and the hydrogen bonds are characterized by the $\text{DnAm}^\text{+}$-$\text{Dn Am}^\text{−}$. The strongest hydrogen bonds are of the type D2A1$\leftrightarrow$D1A2. Another significant finding is the close correlation of the OHb lengthening and the charge-transfer energy. The r-membered rings in the water clusters are also classified by the lengths of the in-phase hydrogen bonds, and the numbers of these rings in the cluster are related to the stability of the cluster. As studied by Singer and his coworkers, there are numerous numbers of isomers of (H₂O)₅ of the same topological configuration of the oxygen atoms, whose hydrogen bonded networks differ only in their directions. The present results provide some guidance for systematic search of the isomers favored both in terms of energy and entropy, and help for further ab initio computations of the large clusters having many 6-membered rings.

**Acknowledgment**

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**References**

Fig. 4 Correlation between charge-transfer $E_{CT}^{X,Y}$ and dispersion $E_{Disp}^{X,Y}$ terms, classified by the types of the hydrogen bonded pairs. a) The isomers of $(\text{H}_2\text{O})_{17}\sim(\text{H}_2\text{O})_{21}$ optimized by Xantheas and his coworkers. b) The isomers of $(\text{H}_2\text{O})_{n}, n = 2 \sim 16$ optimized by Xantheas and his coworkers. c) The isomers of $(\text{H}_2\text{O})_{20}$ and $(\text{H}_2\text{O})_{25}$ optimized by Gadre and his coworkers.
Fig. 5 Correlation of the O···O distance with the charge-transfer energy $E_{CT}^{X,Y}$ and the O-H$_b$ for the isomers of (H$_2$O)$_{20}$ and (H$_2$O)$_{25}$, where H$_b$ is the hydrogen bonding hydrogen. The unfilled points are for $E_{CT}^{X,Y}$.

Fig. 6 Relation of the O-H distance of a hydrogen donor water X with the sum $\sum_{Y \neq X} E_{CT}^{Y \rightarrow X}$. The unfilled marks are for the shorter OH bonds of D2 waters.

a) (H$_2$O)$_{17} \sim$ (H$_2$O)$_{21}$, b) (H$_2$O)$_{20}$ and (H$_2$O)$_{25}$. 

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Pictorial Abstract

Title

Analysis of hydrogen bond energies and hydrogen bonded networks in water clusters \((\text{H}_2\text{O})_{20}\) and \((\text{H}_2\text{O})_{25}\) using the charge-transfer and dispersion terms

Author

Suehiro Iwata

Relation of the charge-transfer and dispersion terms with the \(\text{O—O}\) length for every pair of the hydrogen bonded waters in the isomers of \((\text{H}_2\text{O})_{17}\sim(\text{H}_2\text{O})_{21}\)