# Supporting Information for "Many-Body Contributions in Water Nano-Clusters"

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### Results for the DC model

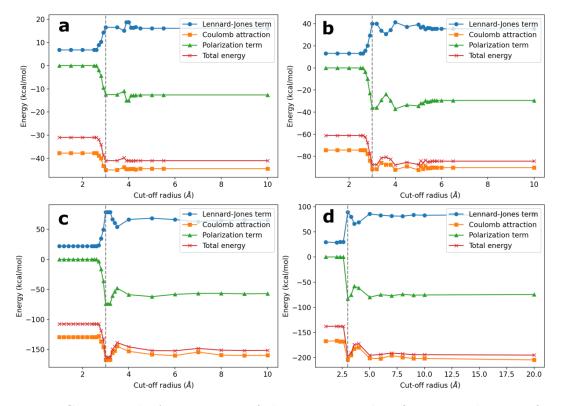


Figure 1: DC potential: As in Fig. 2 of the main text, but for water clusters of 6 (a), 10 (b), 16 (c), and 20 (d) molecules.

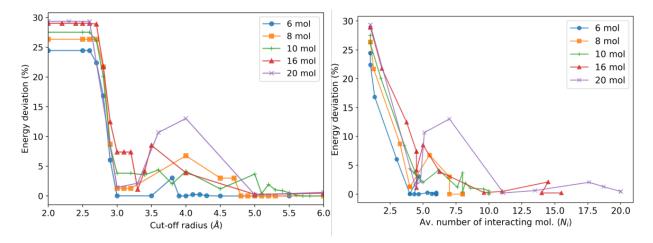


Figure 2: DC potential: Energy deviation from the reference DC value as a function of the cut-off radius r (left) and as a function of the average number of interacting molecules  $\langle N_i \rangle$  (right) for clusters of 6 (blue circles), 8 (orange squares), 10 (green pluses), 16 (red triangles), and 20 (purple crosses) DC water molecules. The deviations are within 5% when the cut-off radius coincides with the first coordination shell ( $r \sim 3$  Å). We take the absolute value of the energy deviations for the artificial minimum-energy configurations–at r between two consecutive coordination shells.

## Minimum-energy configurations with the cut-off at the first coordination shell for the DC model

In Fig.3, we show that, when we include many-body effects until the first coordination shell, the minimum energy configurations are more similar to the results in the DC limit rather than the TIP4P-like limit.

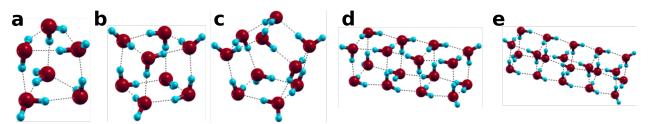


Figure 3: DC potential: The lowest-energy configuration for a cluster with 6 (a), 8 (b), 10 (c), 16 (d), and 20 (e) water molecules calculated for our model with the cut-off corresponding to the 1rst coordination shell ( $r \sim 3$  Å).

#### Results for the MB-pol potential

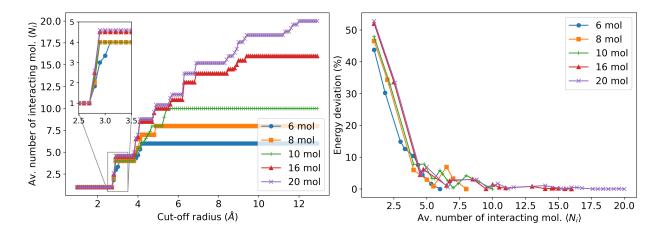


Figure 4: MB-pol potential: Average number of interacting molecules  $\langle N_i \rangle$  as a function of the cut-off radius r (left) and energy deviation from the reference DC value as a function of  $\langle N_i \rangle$  (right) for clusters of 6 (blue circles), 8 (orange squares), 10 (green pluses), 16 (red triangles), and 20 (purple crosses) MB-pol water molecules. The deviations are within 5% when the cut-off radius coincides with the first coordination shell ( $r \sim 3$  Å). We take the absolute value of the energy deviations for the artificial minimum-energy configurations-at r between two consecutive coordination shells.

Among the available many-body models, MB-pol<sup>1-3</sup> has been shown to correctly perform the many-body analyses of small clusters.<sup>4,5</sup> Here we use the MBX calculator<sup>6</sup> to produce the results in Fig. 4. Due to the expensive computational cost, we adopt the minimum-energy configurations for the DC potential as a starting point for the MB-pol analysis.

#### Results for the KJ potential

The many-body KJ potential<sup>7</sup> is a rigid and four-site model similar to the DC potential. In both models, the polarizable site M is located on the bisector of the H-O-H bond angle. The most crucial difference between the two models is the sites associated with the dispersion force: the oxygen atom for the DC model and the M site for the KJ potential. Due to the expensive computational cost, we computed only two cluster sizes for the KJ potential: 8 and 20 molecules (Fig. 5).

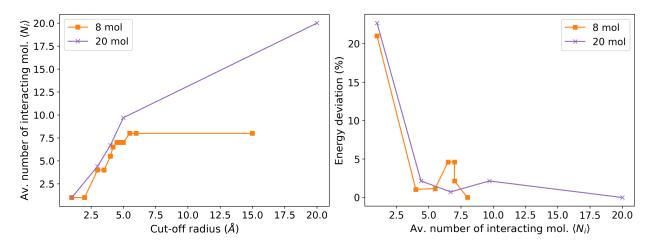


Figure 5: KJ potential: As in Fig. 4 but for the KJ water model for clusters of 8 (orange squares) and 20 (purple crosses).

#### References

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