

Segue between Favorable and Unfavorable Solvation

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Solvation of small and large clusters are studied by simulation, considering a range of solvent–solute attractive energy strengths. Over a wide range of conditions, both for solvation in the Lennard-Jones liquid and in the SPC model of water, it is shown that the mean solvent density varies linearly with changes in solvent–solute adhesion or attractive energy strength. This behavior is understood from the perspective of Weeks' theory of solvation (*Annu. Rev. Phys. Chem.* **2002**, 53, 533) and supports theories based upon that perspective.

1. Introduction

Some solute species dissolve in a liquid solvent over a large range of conditions. Others demix or phase separate from the liquid solvent even at relatively low solute concentrations. This paper focuses on mesoscopic and microscopic manifestations of these macroscopic phenomena. Our motivation is to understand the range of solvation behaviors that are possible in aqueous solutions when hydrophobic species are altered to hydrophilic and vice versa. We demonstrate that various physical quantities vary smoothly under such alterations, metaphorically akin to a segue that smoothly bridges two distinct musical movements. While our motivation concerns aqueous solutions, the physical principles involved are relatively general.

The basic theory has been formulated by Weeks.¹ One first considers the effect of the solute in excluding a region of space from the solvent. The average effect is described by the response of the solvent to a so-called unbalancing potential. Unbalancing arises because the excluded volume presents a region where solvent molecules lose adhesive forces. In a homogeneous fluid, these adhesive forces tend to be in balance, often allowing their effects to be described as a uniform field or chemical potential shift.^{2,3} Introducing a large enough excluded volume, however, destroys this homogeneity, and the response can be very large, particularly so if the liquid exists near coexistence with its vapor. Indeed, in the macroscopic limit, the average liquid density adjacent to excluded regions bounded by a plane is βp , where p is the pressure, and $\beta^{-1} = k_B T$ is temperature times Boltzmann's constant.⁴ In this limit, therefore, the mean liquid density at the surface is a bulk property. It exhibits a phase transition reflecting the bulk liquid–vapor phase transition. Weeks' unbalancing potential captures the effects of this transition in a mean molecular field approximation.

A typical solute does not exclude liquid from a macroscopic volume. Nevertheless, the unbalancing potential can be substantial and the response to it significant, though not singular, provided the solvent-exposed surface area of the solute has low enough curvature and large enough size. Weeks' expression for the unbalancing potential shows that its strength is roughly proportional to the surface area, A . Thus, to linear order in the response,

$$\langle \rho(\mathbf{r}) \rangle_0 \approx a(\mathbf{r}) - A b(\mathbf{r}) \quad (1)$$

Here, $\rho(\mathbf{r})$ denotes the solvent density field at position \mathbf{r} , which we take to be outside the excluded volume, and the pointed

brackets indicate equilibrium ensemble average, with the subscript 0 implying that only excluded volume forces act between solute and solvent. The function $a(\mathbf{r})$ and $b(\mathbf{r})$ are properties of the solvent and the solute position and shape: $a(\mathbf{r})$ gives the response in the absence of unbalancing forces; $A b(\mathbf{r})$ is significant in size only when the liquid solvent is close to phase coexistence with its vapor and A is comparable to or larger than the surface area of the critical nucleus for the phase transition; for positions \mathbf{r} that are beyond a bulk correlation length of the solvent, $a(\mathbf{r})$ approaches the bulk liquid density and $b(\mathbf{r})$ approaches zero.

Fluctuations around the average and the effects of forces beyond those of excluded volume are captured in the second step of Weeks' treatment. In this step, one notes that solvent density fluctuations in a homogeneous liquid obey Gaussian statistics^{5–7} and assumes this statistic is more generally valid for all small length scale fluctuations in a liquid, even in the presence of inhomogeneities. Therefore, if the solute attracts the solvent molecules with a potential strength proportional to λ , the effect of the attraction on the mean solvent density is linear in λ , i.e., $\langle \rho(\mathbf{r}) \rangle_\lambda \approx \langle \rho(\mathbf{r}) \rangle_0 + \lambda S(\mathbf{r})$, where $S(\mathbf{r})$ is the net or integrated response of the solvent density field at \mathbf{r} to the attractions from the solute. If these attractions emanate from interaction sites in the solute and are finite ranged, then $S(\mathbf{r})$ will be approximately linear in A , leading to

$$\langle \rho(\mathbf{r}) \rangle_\lambda \approx \langle \rho(\mathbf{r}) \rangle_0 + \lambda A s(\mathbf{r}) \quad (2)$$

where $s(\mathbf{r})$ is a property of the solvent and the functional form of the attractive interaction. Finally, as both $b(\mathbf{r})$ and $s(\mathbf{r})$ describe responses of the solvent near the solute and per unit exposed area of the solute, one may expect that integrated or coarse-grained effects of the two are approximately proportional. That is to say, $\bar{b}(\mathbf{r}) \approx \lambda^* \bar{s}(\mathbf{r})$, where the over-bar indicates an integral or coarse-graining over some specified microscopic length scale. The value of the proportionality constant, λ^* , depends upon the specific solvent, the specific solute–solvent interactions, and the specific coarse-graining prescription. Adopting this proportionality gives

$$\langle \bar{\rho}(\mathbf{r}) \rangle_\lambda \approx \bar{a}(\mathbf{r}) + (\lambda - \lambda^*) A \bar{s}(\mathbf{r}) \quad (3)$$

Equations 1, 2, and 3 are not generally true, as they do not properly account for saturation in either the limit of very large A or the limit of very large λ . The point of this paper, however, is that these equations are good approximations in a variety of physically pertinent circumstances. We demonstrate their ac-

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curacy in the next section by presenting results of computer simulation calculations of solvent density fields and solvation free energies for solute clusters of various sizes and solvent–solute attractive energy strengths, considering both the Lennard-Jones liquid solvent and water solvent. The results indicate that variations from favorable to unfavorable solvation, i.e., the changes in solvation in passing from large λ to small λ , occur smoothly and in ways that are easy to anticipate. Further, the results provide confidence in theories built from Weeks' perspective. These implications are discussed in Section 3.

2. Simulation Results

To elucidate the response of a liquid to the presence of a solute of general shape and interaction strength, we consider a series of models for both solutes and solvent of increasing complexity.

A. Spherical Solute in Lennard-Jones Solvent. First, we consider the case of a spherical solute immersed in the Lennard-Jones solvent, i.e., a fluid of particles that interact through the pair potential

$$w_{\text{LJ}}(r) = 4\epsilon \left[\left(\frac{\sigma}{r} \right)^{12} - \left(\frac{\sigma}{r} \right)^6 \right] \quad (4)$$

In this equation r denotes the distance between two solvent particles, and ϵ and σ define the energy and length scales of the simulation. The solvent–solvent potentials are shifted so as to be continuous and zero beyond the cutoff distance $r_c = 2.5\sigma$. The reduced thermodynamic state conditions for the solvent are $k_B T/\epsilon = 0.85$, $\rho\sigma^3/\epsilon = 0.022$, and $\rho\sigma^3 = 0.7$, where ρ is the bulk density. These conditions place this Lennard-Jones liquid with truncated and shifted interaction potentials at or very close to liquid–vapor coexistence.^{8,9}

The solute is a sphere of radius R , and its interaction with the solvent is that of a sphere consisting of a Lennard-Jones material with homogeneous density equal to that of the solvent. A solvent particle at distance r from the solute center has the potential energy¹⁰

$$\Phi(r) = \pi\epsilon\rho\sigma^3 \left[\frac{4}{5}\sigma^9 \left(\frac{1}{8rr_+^8} - \frac{1}{9r_+^9} - \frac{1}{8rr_-^8} + \frac{1}{9r_-^9} \right) - 2\sigma^3 \left(\frac{1}{2rr_+^2} - \frac{1}{3r_+^3} - \frac{1}{2rr_-^2} + \frac{1}{3r_-^3} \right) \right] \quad (5)$$

where $r_{\pm} = r \pm R$, if $r \geq R$, and $\phi(r) = \infty$ otherwise. This potential is minimal at a distance r_{\min} , and its minimum value $\Phi_{\min} = \Phi(r_{\min})$ depends on the solute size R .

To systematically investigate the role of attractive interactions, we employ the scheme introduced by Weeks, Chandler, and Andersen (WCA) to partition the energy into the short-ranged repulsive and the long-ranged attractive part,³

$$\Phi_{\lambda}(r) = \Phi_0(r) + \lambda\Phi_1(r) \quad (6)$$

where

$$\Phi_0(r) = \begin{cases} \Phi(r) - \Phi(r_{\min}) & \text{if } r \leq r_{\min} \\ 0 & \text{if } r > r_{\min} \end{cases} \quad (7)$$

and

$$\Phi_1(r) = \begin{cases} \Phi(r_{\min}) & \text{if } r \leq r_{\min} \\ \Phi(r) & \text{if } r > r_{\min} \end{cases} \quad (8)$$

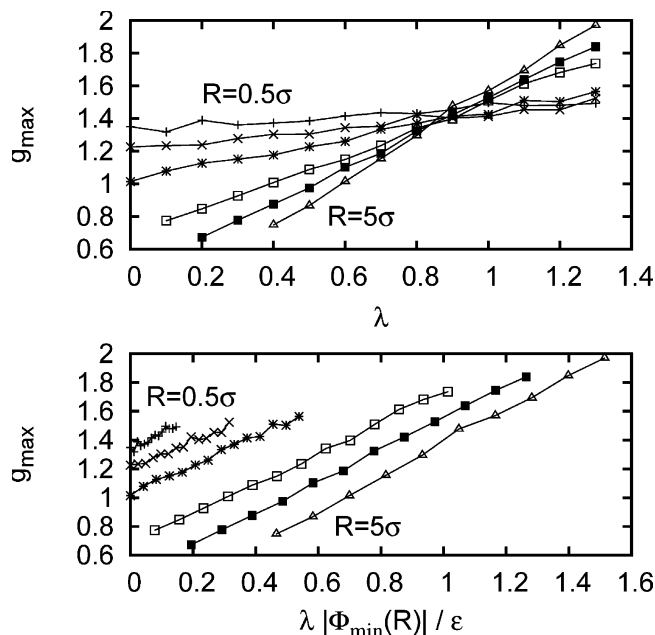


Figure 1. Magnitude of the first peak of the solvent density distribution function around homogeneous Lennard-Jones solutes of radii $R/\sigma = 0.5, 0.7, 1, 2, 3, 5$, plotted as a function of the bare attraction strength λ (top) and the minimum of the solvent–solute potential (bottom). For large solute sizes and small values of λ , the reduced density distribution function did not have a well-defined first peak, and no data is shown for those systems.

The parameter λ controls the strength of the attractive solute–solvent interactions.

We performed computer simulations using the Monte Carlo method at fixed pressure and temperature for different values of the solute size R and interaction strength λ . The number N of solvent particles ranged from 864 to 8192 depending the size of the solute. The solute–solvent interaction was truncated and shifted at a distance $r = R + 2.5\sigma$. We calculate the average reduced solvent density at position r ,

$$g(r) = \langle \rho(r) \rangle / \rho \quad (9)$$

in an ensemble where the solute position is fixed at $r = 0$. Because of the isotropic nature of the solute, this density depends only on the distance $r = |r|$ from the solute center.

We find that the reduced density $g(r)$ has generally a well-defined first peak corresponding to the first solvation shell of the solute. The height of this peak, denoted g_{\max} , is shown in the top panel of Figure 1 for different values of R and λ . This graph encompasses many of the important effects of our understanding of solvophobic and solvophilic solvation.

First, for a fixed value of λ less than 0.9, the solvent density near the solute decreases with increasing solute radius. It has been observed previously that the contact density near a hard sphere solute decreases from values above the bulk liquid density at small solute sizes to the equilibrium vapor density at large solute sizes, both for the Lennard-Jones solvent¹¹ and water.¹⁰ The creation of a liquid–vapor interface near large enough solutes is the basis of our understanding of the hydrophobic effect.¹² Our simulations show that this behavior persists for solutes with weak but finite attraction strengths λ .

Second, for any given solute size R , the density near the solute increases approximately linearly with the attraction strength λ . This smooth response of the density to an attractive field over a wide range of interaction strengths shows that the solvent

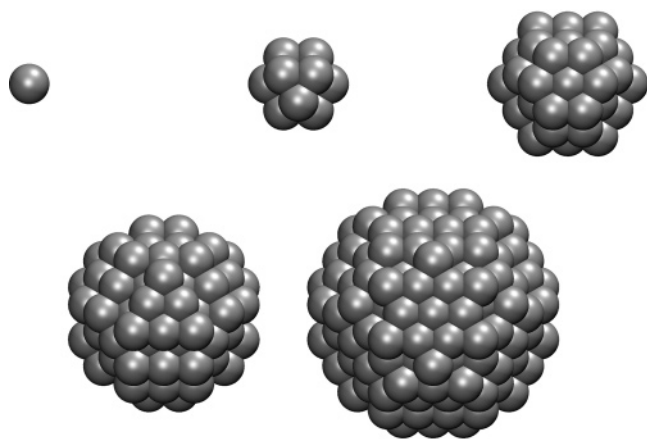


Figure 2. Space-filling view of the solute clusters with particle numbers $n = 1, 13, 57, 135, 305$. The clusters are spherical cuts of a hexagonally close-packed crystal.

properties do not change significantly as λ increases. In particular it is consistent with Gaussian density fluctuations for all attraction strengths.

Third, it is apparent that the slope $\partial g_{\max}/\partial \lambda$, while independent of λ , depends on the radius R of the solute. In particular, there is a value $\lambda^* \approx 0.9$ of the interaction strength where the mean density itself seems to be independent of the solute size. A similar result is suggested by Ashbaugh and Paulaitis for the density near methane clusters in water.¹³ This behavior can be easily understood within the framework introduced in Section 1. The numerical value of λ by itself is not an accurate parameter to describe the strength of the solute–solvent interactions. For the special case of the homogeneous Lennard-Jones solute, a much more reliable indicator is the magnitude of the minimum in the potential energy, $\lambda|\Phi_{\min}(R)|$. The bottom panel of Figure 1 shows that the density response is the same for all solute sizes if the control parameter is chosen to be the actual magnitude of the solute–solvent attractions.

Our results for this simple model solute serve as a useful illustration for the theory outlined in Section 1. We now turn to more complicated solutes, where the lack of isotropy demands the analysis of more complicated density distribution functions. While these complications can obscure the underlying physics, we will see that the basic picture of solvation presented here remains correct.

B. Cluster Solutes in Lennard-Jones Solvent. We consider clusters of n particles solvated in the same Lennard-Jones fluid as before. The clusters are the model solutes of Ashbaugh and Paulaitis.¹³ Space-filling views are shown in Figure 2. The clusters are composed of $n = 1, 13, 57, 135$, and 305 hexagonally close-packed particles with nearest-neighbor distance $d = 2^{1/6}\sigma$. The solute interacts with the solvent molecules with the potential energy

$$W_\lambda = \sum_{i=1}^N \sum_{\alpha=1}^n w_\lambda(|\mathbf{r}_\alpha - \mathbf{R}_i|) \quad (10)$$

where \mathbf{r}_α is the position of the α th cluster particle, \mathbf{R}_i is the position of the i th solvent particle, and

$$w_\lambda(r) = w_0(r) + \lambda w_1(r) \quad (11)$$

Here, $w_0(r)$ and $w_1(r)$ are, respectively, the repulsive branch and the attractive branch of the Lennard-Jones potential,³

$$w_0(r) = \begin{cases} 4\epsilon_s \left[(\sigma_s/r)^{12} - (\sigma_s/r)^6 + \frac{1}{4} \right] & \text{if } r \leq 2^{1/6}\sigma_s \\ 0 & \text{if } r > 2^{1/6}\sigma_s \end{cases} \quad (12)$$

and

$$w_1(r) = \begin{cases} -\epsilon_s & \text{if } r \leq 2^{1/6}\sigma_s \\ 4\epsilon_s [(\sigma_s/r)^{12} - (\sigma_s/r)^6] & \text{if } r > 2^{1/6}\sigma_s \end{cases} \quad (13)$$

We choose $\sigma_s = \sigma$ and $\epsilon_s = \epsilon$ so that the interaction between a cluster particle and a solvent particle is the same as that between two solvent particles at $\lambda = 1$.

The $n = 1$ solute is spherically symmetric, and the mean solvent density near it is described by the isotropic radial distribution function $g(r)$ as before. For the clusters with $n > 1$, however, specification of the mean solvent density requires many more variables than r , and it is natural to consider projections of the mean density. One such projection is the so-called proximal distribution $g_{\text{prox}}(r)$, defined as the reduced density averaged over the surface^{13,14}

$$\{\mathbf{r}' \in \mathbb{R}^3: r_{\text{prox}}(\mathbf{r}') = r\} \quad (14)$$

where

$$r_{\text{prox}}(\mathbf{r}) = \min\{|\mathbf{r} - \mathbf{r}_\alpha|: \alpha = 1, \dots, n\} \quad (15)$$

is the distance from a point \mathbf{r} to the nearest cluster particle. While explicitly a function of only one length, r , it is implicitly a multipoint distribution function¹⁵ except in the case $n = 1$. For that special case, $g_{\text{prox}}(r)$ is the standard radial distribution function $g(r)$.

Figure 3 illustrates the behavior of the proximal distribution function for the model solutes in the Lennard-Jones solvent. Ashbaugh and Paulaitis proposed the height g_{\max} of the first peak in $g_{\text{prox}}(r)$ as a meaningful measure of the solvent density near arbitrary solutes. In the top panel, we show that the behavior of this observable is qualitatively the same as that shown in Figure 1 for isotropic solutes. The response of the mean solvent density to changes in solute–solvent attraction is proportional to that change. This result is as expected according to eq 2. Another important point is that this proportionality leads to a coincidence of curves at $\lambda = \lambda^*$, where in this case $\lambda^* \approx 0.5$. The existence of a co-incident value of λ is anticipated in eq 3.

Ashbaugh and Paulaitis found in their study of methane clusters solvated in water that at λ^* not only the heights of the first peak in $g_{\text{prox}}(r)$ coincide for different cluster sizes, but that the proximal distribution functions themselves are very similar. The bottom panel of Figure 3 establishes a similar result for our model system. From this behavior, Ashbaugh and Paulaitis conclude that the solvation environment of a monomeric methane particle is essentially the same as that of methane confined to a cluster of arbitrary size. Figure 4 shows that this conclusion is not justified. The contour lines of constant density do not coincide with the surfaces of constant proximal distance for the $n = 135$ cluster. In particular, the crevices of the solute, where the attraction strength is large, contain a much higher solvent density than the proximal distribution function suggests. No such behavior exists for the monomeric solute, where the density profile is isotropic.

As the strength of the attractive solute–solvent interaction increases, the clusters lose their hydrophobic character until eventually they become hydrophilic. To quantify this process,

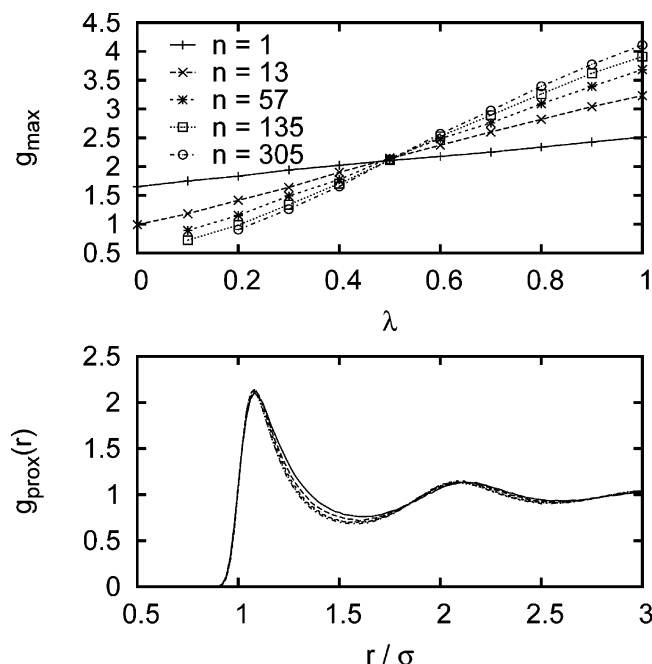


Figure 3. Proximal solvent density distribution function around clusters of Lennard-Jones particles immersed in the Lennard-Jones fluid. (top) Height of the first peak in $g_{\text{prox}}(r)$ for different cluster sizes n and attraction strengths λ . (bottom) Proximal distribution functions for $\lambda = \lambda^* = 0.5$.

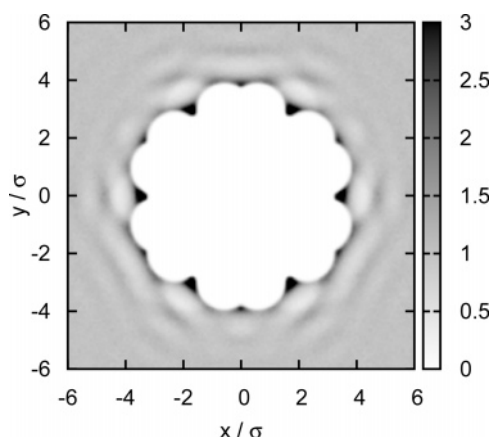


Figure 4. Reduced solvent density around the $n = 135$ solute cluster immersed in the Lennard-Jones fluid at $\lambda = \lambda^* = 0.5$. Shown is a slice through the solute center of width 0.1σ with resolution 0.05σ , demonstrating that the solvent density does not follow the shape of the solute surface.

we calculate the solvation free energy, or excess chemical potential, as a function of λ , which is given by

$$\Delta\mu_\lambda = \Delta\mu_0 + \int_0^\lambda d\lambda' \langle W_1 - W_0 \rangle_{\lambda'} \quad (16)$$

where $\langle \dots \rangle_\lambda$ indicates ensemble average with solute–solvent attractive force strength λ .

Solvation free energies due to excluded volume are approximately proportional to the excluded volume for small solutes. Further, at large excluded volumes with small surface curvature and no attractive solvent–solute forces, solvent density is depleted near the solute. Therefore, for the solutes we have considered, $\Delta\mu_0$ can be well approximated by the solvation free energy for a hard sphere with the same excluded volume as the solute clusters.¹¹ Here we define the excluded volume as those spatial regions where the solute–induced field is greater than the thermal energy $k_B T$.

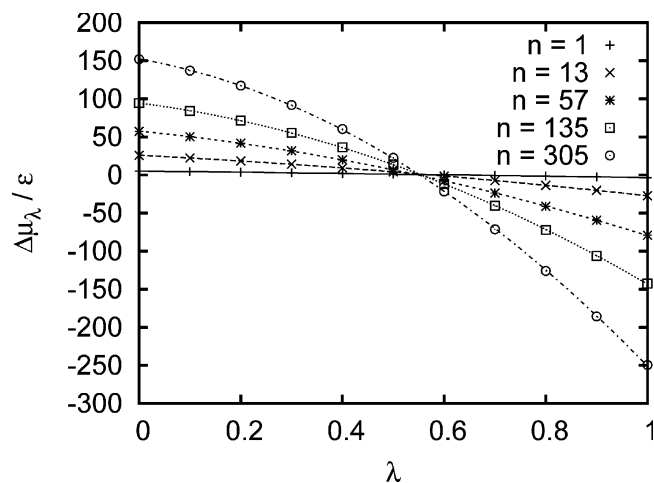


Figure 5. Excess solvation energy of solute clusters in the Lennard-Jones fluid. The symbols are calculated from simulation using eq 16, and the lines are quadratic fits through the data.

The result of this calculation is shown in Figure 5. Over the range of interaction strengths considered, the solvation free energies change from strongly positive to negative. In other words, at large values of λ , the clusters are not solvophobic. This transition occurs smoothly and regularly. The shape of these curves is understood because the attractive energy is approximately proportional to λ times an integral over $\langle \rho(r) \rangle_\lambda$, and the latter is a linear function of λ , as noted in eq 2.

C. Cluster Solutes in SPC Water. In another series of calculations, the solvent is the SPC model for liquid water¹⁶ and the solutes are clusters of methane molecules in the united atom approximation of ref 17. As in ref 13, these clusters are hexagonally close-packed with a particle spacing of $d = 4.19$ Å. The methane molecules interact with the water oxygens through a Lennard-Jones potential with $\sigma = 3.448$ Å and $\epsilon = 0.895$ kJ/mol, which was truncated and shifted at a distance of 9 Å. As before, we use the WCA partition of the interaction potential to obtain control over the strength of the attractions, given by the parameter λ . Electrostatic interactions were calculated using Ewald summation. Molecular dynamics simulations were performed using the LAMMPS program^{18,19} at temperature $T = 298$ K and a box size chosen to give the correct bulk density.¹³

As noted above, Ashbaugh and Paulaitis¹³ found that the proximal solvent density distribution functions are very similar for different cluster sizes at the specific attraction strength $\lambda^* \approx 1$, from which they concluded that the solvent density near a cluster of solute particles is basically the same as that near a single solute particle. In Figure 6, we show the reduced water density near the $n = 135$ solute with an interaction strength $\lambda = \lambda^*$. For this specific system, the solvent density is indeed approximately constant along contours of constant proximal distance, so that

$$\langle \rho(r) \rangle \approx \rho g_{\text{prox}}(r_{\text{prox}}(r)) \quad (17)$$

is a rather good approximation for the range of parameters considered. However, this agreement is not exact, as deviations are visible in the crevices. In these regions adjacent to the cluster, the solvent interacts with many solute particles, and the attractive potential well reaches values as deep as -5 kJ/mol. This value is to be compared with the average energy of a water molecule in the bulk fluid, which is approximately 40 kJ/mol.²⁰ Thus, the cluster crevices are approaching hydrophilic character

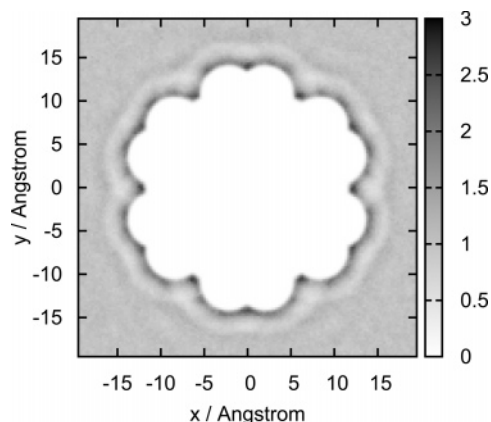


Figure 6. Reduced water density around the $n = 135$ solute cluster at $\lambda = \lambda^* = 1$. Shown is a slice through the solute center of width 0.5 \AA with resolution 0.25 \AA . To a good approximation, the solvent density follows the shape of the solute cluster.

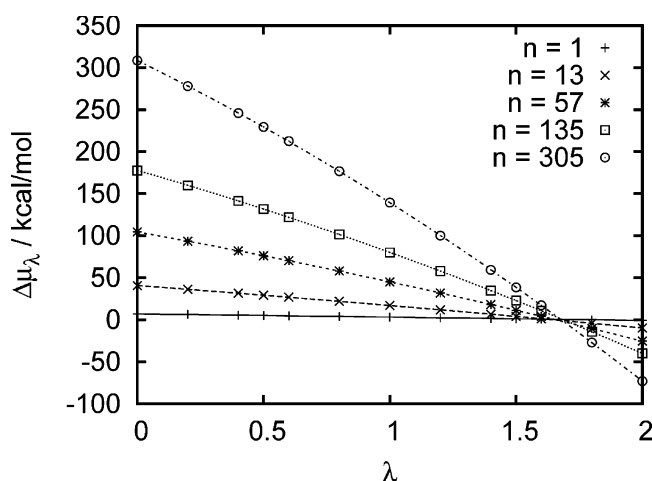


Figure 7. Excess solvation energy of methane clusters in water. The symbols are computed from simulation with the aid of eq 16 and known solvation free energies of hard sphere solutes in water.²⁸ The lines are quadratic fits to the data.

and can compensate for a fraction of the average water–water interaction energy.

Figure 7 shows the solvation free energy of the methane clusters in water, calculated as before. Again the clusters show a smooth transition from hydrophobic solvation at small values of λ to hydrophilic solvation at large attraction strengths. For this particular system, the crossover between these two regimes occurs at values of λ greater than λ^* .

3. Discussion

The results of the previous section show that the role of solvent–solute attractions on solvation is significant but simple. In particular, while fluctuations of solvent density are affected by the shape and size of the solute and the thermodynamic state of the solvent, the mean solvent density changes only linearly with changes in the strength of adhesion between solute and solvent. If, for example, a solute is small, the liquid solvent will be constrained to be outside the solute's excluded volume but otherwise behave as if it were bulk. This conception is precisely the physics that underlies the Pratt–Chandler theory of hydrophobicity^{7,21} and its contemporary variant.⁵ On the other hand, if the solute's girth is such to exclude a volume with large low-curvature surface area, fluctuations of the solvent are changed to reflect the formation of an interface.^{1,12,22} To the

extent there is no solvent–solute adhesion, the interface is essentially the same as that between liquid and vapor.²³ The role of attractions between the solute and solvent is not to change the fluctuations, but to affect the average solvent density and the position of this interface. For strong enough attractions, the effect is to pin the interface at specific positions, as seen in Figure 4 and to a lesser extent in Figure 6.

This picture of the role of attractions in the context of hydrophobic solvation has been discussed before.^{10,12} Simulation results of Werder et al. on the hydration of several different models of graphite in water²⁴ can be viewed as illustrations of that role. Namely, Werder et al. establish linear trends for hydration energies as functions of solvent–solute attractive energy strength. The relative range of parameters is smaller in that work than in ours, probably explaining why curvature in the linear trends is not apparent. Along with demonstrating regularity and simplicity, Werder et al. also show that the trends are significant in that modest potential parameter changes result in models of “graphite” that change from hydrophobic to hydrophilic. In one model, Werder et al. consider a Lennard-Jones potential between graphite carbon atoms and water oxygen atoms with energy and length parameters $\epsilon_{\text{CO}} = 0.39 \text{ kJ/mol}$ and $\sigma_{\text{CO}} = 0.32 \text{ nm}$. They establish that, in this model, line tension is positive and the contact angle between water and a planar graphite surface is finite. In contrast, a similar model of a presumed hydrophobic surface but with $\epsilon_{\text{CO}} = 0.48 \text{ kJ/mol}$ and $\sigma_{\text{CO}} = 0.33 \text{ nm}$ yields a negative surface tension between water and the surface.^{25,26} This latter surface is therefore not hydrophobic.

An important consequence of the simplicity we emphasize regards the entropy of solvation. To the extent that density fluctuations obey Gaussian statistics (i.e., the mean density responds linearly to forces of adhesion), the entropy of solvation is independent of adhesive energy strength. This idea is used in ref 27 to replace the hydration entropy of a protein with that of a hard sphere occupying the same volume even though the surface of the folded protein is largely hydrophilic. By demonstrating linear trends for the mean density, our work here provides explicit justification for this replacement.

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- (15) Equivalent to eqs 14 and 15, the proximal distribution function can be written as $\Omega_1 r^2 \rho g_{\text{prox}}(r) = \langle \sum_{i=1}^N \delta(r - |R_i|) \prod_{a=2}^N \theta(|R_i - r_a| - |R_i|) \rangle$. The equilibrium average is performed with solute site 1 at the origin, i.e., $r_1 = 0$, this site being at the surface of the cluster and with subtended

solid angle Ω_1 , $\delta(x)$ is the Dirac δ function, and $\theta(x)$ is the unit Heaviside function. The quantity $\rho_{g_{\text{prox}}}(r)$ is the mean density of solvent that is closest to and within a shell of radius r from cluster particle $\alpha = 1$.

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(26) In ref 24, Werder et al. do not report the oil–water surface tension for these particular parameters, but Choudhury and Pettitt, in effect, do so in ref 25. Specifically, for carbon–carbon potential parameters, the authors of ref 25 cite $\epsilon_{\text{CC}} = 0.36$ kJ/mol and $\sigma_{\text{CC}} = 0.34$ nm and combine these with oxygen–oxygen parameters $\epsilon_{\text{OO}} = 0.65$ kJ/mol and $\sigma_{\text{OO}} = 0.32$ nm, using geometrical and arithmetic means, respectively, to obtain their carbon–oxygen parameters. They compute a potential of mean force by molecular simulation for a pair of plates formed from these “carbon” particles. Figure 1a of their paper shows that the solvent contribution to the free energy for dissociated plates is lower than that of associated plates. The plates are reasonably large, so that the solvation energy is predominantly proportional to the solute surface area exposed to the solvent. Choudhury and Pettitt’s result for the solvent contribution to the potential of mean force therefore shows that the interactions used in their study produce a negative solvation surface free energy for the hydrated plate.

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