Supporting Information

Kastelic et al. 10.1073/pnas.1507303112

Calculation of Measurable Properties

The free energy of solution A is in our approach given by the sum of three contributions:

$$A = A^{\mathrm{id}} + A^{\mathrm{hs}} + A^{\mathrm{ass}},$$
 [S1]

where A^{id} is the ideal part, A^{hs} the hard-sphere contribution (1), and A^{ass} the protein–protein associative part (2–4). The ideal and the hard-sphere contributions are given by the next two expressions:

$$\frac{\beta A^{\rm id}}{N} = \ln(\Lambda^3 \rho) - 1, \qquad [S2]$$

$$\frac{\beta A^{\rm hs}}{N} = \frac{\eta (4 - 3\eta)}{(1 - \eta)^2},$$
 [S3]

where Λ is the de Broglie thermal wavelength (5) and η is the packing fraction, related to the density, $\eta = \pi \rho \sigma^3/6$. Note that A^{ass} is defined by Eq. **6** of the main text.

Once the free energy is known other thermodynamic quantities can easily be calculated (5). Chemical potential μ of the protein molecule and pressure *P* needed in calculations of the coexistence lines are obtained via these equations:

$$\mu = \left[\frac{\partial(A/V)}{\partial\rho}\right]_{T,V},$$
[S4]

$$P = \rho \mu - \frac{A}{V}.$$
 [85]

Note that the ideal and the hard-sphere free-energy terms were differentiated analytically, whereas the associating term was evaluated numerically.

In the one-component system of two coexisting phases with equilibrium densities ρ_a and ρ_b , additional restrictions must be fulfilled. For every temperature *T* the equilibrium conditions read

$$\mu(T, \rho_{\rm a}) = \mu(T, \rho_{\rm b}),$$
 [S6]

$$P(T, \rho_{\rm a}) = P(T, \rho_{\rm b}).$$
 [S7]

To compute the number concentrations of the coexisting phases ρ_a and ρ_b efficiently, the iterative Newton-Ralphson method was used. The convergence criterium, requesting for the

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sum of relative errors in ρ_a and ρ_b to be less than 10^{-4} , was typically reached in fewer than five iterations.

The results for the reduced chemical potential $\beta\mu$ and dimensionless pressure $\beta P \sigma^3$ showing the coexisting points for lysozyme at T = 270 K and γ IIIa-crystallin at T = 300 K are shown in Fig. S1. Dashed horizontal lines connect the number densities ρ_a and ρ_b of the two coexisting phases. The differences between densities of the coexisting phases, $\rho_b - \rho_a$, observed in Fig. S1 increase with the number of binding sites M. It is clear from our computation that larger number of sites M (solid line, γ IIIacrystallin, M = 14) yields a broader liquid–liquid phase diagram. For a lower number of sites (dotted and dashed line, lysozyme, M = 10) a somewhat deeper potential well would be needed to obtain the same critical temperature, determined by the $\beta\mu$ and $\beta P \sigma^3$ curves.

Among thermodynamic properties, the second virial coefficient B_2 is very important. In the framework of Wertheim's theory it is given by the next expression (6):

$$B_2 = B_2^{(\text{hs})} - 2\pi M^2 \int_{\sigma}^{2d+a_W} e_{\text{R}}(r)\overline{f}_{\text{ass}}(r)r^2 dr, \qquad [\textbf{S8}]$$

where $e_{\rm R}(r) = \exp[-\beta u_{\rm R}(r)]$. $\bar{f}_{\rm ass}(r)$ is the solid-angle average of the Mayer function defined as

$$\bar{f}_{\rm ass}(r) = \int f_{\rm AB}(\mathbf{x}_{\rm AB}(\mathbf{r})) d\Omega_{\rm A} d\Omega_{\rm B},$$
[S9]

 $f_{AB}(\mathbf{x}_{AB}(\mathbf{r})) = \exp[-\beta u_{AB}(\mathbf{x}_{AB}(\mathbf{r}))] - 1$ is the site-site Mayer function, and Ω_A and Ω_B are the solid angles. Wertheim (7) obtained the result in analytical form

$$\overline{f}_{ass}(r) = \frac{\exp(\beta e_{\rm W}) - 1}{24d^2r} (a_{\rm W} + 2d - r)^2.$$

$$\times (2a_{\rm W} - 2d + r)$$
[S10]

Another measurable quantity, closely related to B_2 , is the osmotic compressibility χ_{osm} (8):

$$\chi_{\rm osm} = \left(\frac{\partial\beta P}{\partial\rho}\right)_{N,T} = 1 + 2B_2\rho + \dots$$
 [S11]

 $\chi_{\rm osm}$ was obtained by numerical differentiation of the pressure as suggested in the first part of Eq. **S11**.

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Fig. S1. Calculated reduced chemical potential $\beta\mu$ and dimensionless pressure $\beta P \sigma^3$ for lysozyme (dotted and dashed line, T = 270 K) and γ llla-crystallin (solid line, T = 300 K), based on parameters from Table 1 of the main text.

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S A