Electronic Supplementary Information for

# Evidence for Cooperative Na<sup>+</sup> and Cl<sup>-</sup> Binding by Strongly Hydrated L-Proline

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### 1 Methods — Statistical Mechanics

For 1D-RISM site-specific coordination numbers,  $n_{\alpha\beta}$ , were obtained from radial pair distribution functions (PDFs),  $g_{\alpha\beta}(r)$ , as

$$n_{\alpha\beta} = 4\pi\rho_{\beta} \int_{0}^{r_{\min}} g_{\alpha\beta}(r)r^2 \,\mathrm{d}r \tag{S1}$$

where  $\rho_{\beta}$  is the average number density of solvent sites/atoms of kind  $\beta$ . Such partial coordination numbers give the average number of sites/particles of type  $\beta$  in a coordination sphere of radius  $r_{\min}$  around an  $\alpha$  site. The integration limit,  $r_{\min}$ , is defined by the position of the first minimum of the PDF and corresponds to the radius of the first coordination shell around reference site  $\alpha$ .

In the case of 3D-RISM the spatial distribution functions (SDFs),  $g_{\beta}(\mathbf{r})$ , give the spatial distribution of solvent molecules around a solute molecule, which is commonly represented by isodensity surfaces at a selected probability level (see Fig. 6 of the main manuscript). From such SDFs the associated total coordination number of sites  $\beta$  within a shell of volume  $V_{\rm s}$  in contact with the reference molecule can be calculated as

$$n_{\rm t} = \rho_\beta \int_0^{V_{\rm s}} g_\beta(\boldsymbol{r}) \,\mathrm{V} \tag{S2}$$

Since Pro is not a spherical molecule, its hydration shell is also not spherical. Thus, for the calculation of the total water coordination number the previously suggested approximation for the first hydration shell as a closed surface with arbitrary shape was used.<sup>1</sup>

The SDFs can also be used to calculate cylindrical distribution functions (CDFs),  $g_{pyr-\alpha}(z)_R$ , originally derived for describing the local atom density adjacent to planar surfaces of molecules.<sup>2</sup> These CDFs give the probability of finding a specific site at a distance, z, orthogonal to the reference plane xy defined by the central molecule, within a cylinder of radius R.<sup>1,3</sup> For the present investigation such CDFs are a useful tool to characterize the distribution of water molecules above and below the plane of the pyrrolidine ring and to calculate corresponding coordination numbers, see Fig. S1.

#### 2 Methods — Correction for kinetic depolarization

Due to their electric field ions orient surrounding solvent dipoles to some extent. When moving in an external electric field,  $\vec{E}$ , ions therefore exert a torque on surrounding solvent dipoles opposing the tendency of the latter to align with  $\vec{E}$ . This purely dynamical effect, with a finite magnitude at zero frequency, leads to a depolarization of the bulk solvent in addition to solvation effects and accordingly, the experimentally detected bulk-water amplitude,  $S_{\rm b}$ , is reduced by the kinetic dielectric decrement,  $\Delta \varepsilon_{\rm kd}$ , compared to the equilibrium amplitude,  $S_{\rm b}^{\rm eq}$ , reached in the absence of ionic motion.<sup>4</sup> The latter is also reduced compared to the pure solvent because of solvent dilution and, mainly, ion solvation.<sup>5</sup> In electrolytes of salt concentration c the amplitude relevant for calculating the DRS-detected concentration of bulk-like solvent,  $c_{\rm b}$ , and thus of the total effective solvation number,  $Z_{\rm t}$ , is therefore given by

$$S_{\rm b}^{\rm eq}(c) = S_{\rm b}(c) + \Delta \varepsilon_{\rm kd} \tag{S3}$$

The original continuum theory of Hubbard and Onsager<sup>4</sup> (HO) was derived for vanishing salt concentration and thus is problematic when correcting experimental  $S_{\rm b}$  values but recently Sega *et al.*<sup>6</sup> published a phenomenological modification valid also at finite salt concentrations, yielding

$$\Delta \varepsilon_{\rm DD} = p \times \frac{\varepsilon_{\rm w}(0) - \varepsilon_{\infty}(0)}{\varepsilon_{\rm w}(0)} \times \tau_{\rm w}(0) \times \frac{\kappa}{\varepsilon_0} \times \exp[-\kappa_{\rm D}R] \times (\kappa R + 2)/2 \tag{S4}$$

where  $\varepsilon(0) [= 78.368]$  is the static permittivity of the pure solvent,  $\varepsilon_{\infty}(0) [= 3.52]$  is the infinite-frequency permittivity,  $\tau_{w}(0) [= 8.35 \text{ ps}]$  its relaxation time,  $\kappa$  is the solution conductivity,  $\varepsilon_{0}$  the electric field constant and  $\kappa_{D}$  the reciprocal Debye length.<sup>7</sup> For the effective ion size  $R = (r_{+} + d_{w} + r_{-})/2 = 0.284 \text{ nm}$  was chosen, where  $r_{+}$  and  $r_{-}$  are the radii of Na<sup>+</sup> and Cl<sup>-</sup>, and  $d_{w}$  is the diameter of a water molecule.<sup>8</sup> For the hydrodynamic parameter the value for slip boundary conditions, p = 2/3, was chosen as this yielded consistent limiting ionic hydration numbers when used with HO theory for vanishing electrolyte concentrations.<sup>5</sup>

## 3 Supplementary Tables

Table S1: Site-specific coordination numbers,  $n_{\alpha\beta}$ , and corresponding distances,  $r_{\alpha\beta}$  (in brackets), of L-proline in aqueous solutions as a function of Pro concentrations, c(Pro), from 1D-RISM calculations

$\begin{array}{c ccccccccccccccccccccccccccccccccccc$									
$\begin{array}{c cccccccccccc} Carboxylate group & & & & & & & & & & & & & & & & & & &$	6.0								
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	Carboxylate group								
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	4.43								
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	(0.310)								
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	3.94								
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	(0.310)								
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	1.66								
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	(0.175)								
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	1.41								
$\begin{array}{ccccccc} \mathrm{NH}_2^+ \ \mathrm{group} \\ n_{\mathrm{N1Ow}} & 4.48 & 4.23 & 3.94 & 3.69 & 3.40 & 3.14 \\ (r_{\mathrm{N1Ow}} /\mathrm{nm}) & (0.300) & (0.298) & (0.298) & (0.298) & (0.298) & (0.295) \\ n_{\mathrm{H8Ow}} & 0.90 & 0.87 & 0.82 & 0.78 & 0.75 & 0.70 \\ (r_{\mathrm{H8Ow}} /\mathrm{nm}) & (0.175) & (0.175) & (0.175) & (0.175) & (0.175) \\ n_{\mathrm{H9Ow}} & 0.83 & 0.79 & 0.75 & 0.70 & 0.66 & 0.62 \\ (r_{\mathrm{H9Ow}} /\mathrm{nm}) & (0.173) & (0.170) & (0.170) & (0.170) & (0.170) \\ \end{array}$	(0.178)								
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$\rm NH_2^+$ group								
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	2.84								
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	(0.295)								
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	0.65								
$egin{array}{ccccccc} n_{ m H9Ow} & 0.83 & 0.79 & 0.75 & 0.70 & 0.66 & 0.62 \ (r_{ m H9Ow}/ m nm) & (0.173) & (0.170) & (0.170) & (0.170) & (0.170) & (0.170) \end{array}$	(0.175)								
$(r_{\rm H9Ow}/{\rm nm})$ (0.173) (0.170) (0.170) (0.170) (0.170) (0.170)	0.57								
	(0.170)								
Ring carbon atoms									
$n_{\rm C2Ow}$ 3.88 3.77 3.49 3.13 2.87 2.64	2.37								
$(r_{ m C2Ow}/ m nm)$ (0.340) (0.338) (0.338) (0.338) (0.338) (0.338)	(0.335)								
$n_{\rm C3Ow}$ 5.44 5.31 4.94 4.50 4.14 3.84	3.49								
$(r_{ m C3Ow}/ m nm)$ (0.355) (0.355) (0.355) (0.353) (0.350) (0.350)	(0.350)								
$n_{\rm C4Ow}$ 6.68 6.37 5.97 5.64 5.38 5.01	4.59								
$(r_{ m C4Ow}/ m nm)$ (0.350) (0.350) (0.348) (0.348) (0.348) (0.348) (0.345)	(0.345)								
$n_{\rm C50w}$ 6.96 6.60 6.15 5.80 5.36 4.97	4.51								
$(r_{ m C5Ow}/ m nm)$ (0.323) (0.323) (0.323) (0.323) (0.320) (0.320)	(0.320)								
$H_2O$ sandwiching pyrrolidine ring <sup>b</sup>									
$n_{\rm pyr-Ow}$ 1.04 0.98 0.93	0.56								
Hydrophilic sites: $n_{\rm h} = n_{\rm O10w} + n_{\rm O20w} + n_{\rm N10w}$									
$n_{\rm h}$ 19.53 17.95 16.25 15.39 13.88 12.77	11.21								
Total number of first-shell $H_2O$ molecules, $n_t$ (3D-RISM)									
$n_{\rm t}$ 25.4 24.0 22.4	13.2								

<sup>*a*</sup> Data for hydrophilic sites taken from Fedotova and Dmitrieva<sup>9</sup>; <sup>*b*</sup> from CDF peaks at z = 0.360 nm (6M: 0.340 nm) and -0.340 nm, see Fig. S1.

Table S2: 1D-RISM results for site-specific coordination numbers,  $n_{\alpha\beta}$ , and corresponding distances,  $r_{\alpha\beta}$  (in brackets), of L-proline in aqueous Pro+NaCl solutions of c(Pro) = 0.6 M, and c(NaCl)

$c({ m NaCl})/{ m M}$	0	0.2	0.5	1.0	1.5	2.0		
Carboxylate group								
$n_{O1Ow}$	7.43	7.44	7.35	7.27	7.20	7.12		
$(r_{ m O1Ow}/ m nm)$	(0.310)	(0.310)	(0.310)	(0.310)	(0.310)	(0.310)		
$n_{\rm O2Ow}$	6.65	6.67	6.60	6.53	6.49	6.42		
$(r_{ m O2Ow}/ m nm)$	(0.313)	(0.313)	(0.313)	(0.313)	(0.313)	(0.313)		
$n_{ m O1Hw}$	2.12	2.11	2.09	2.06	2.04	2.01		
$(r_{ m O1Hw}/ m nm)$	(0.175)	(0.175)	(0.175)	(0.175)	(0.175)	(0.175)		
$n_{ m O2Hw}$	1.83	1.82	1.80	1.77	1.75	1.73		
$(r_{ m O2Hw}/ m nm)$	(0.178)	(0.178)	(0.178)	(0.178)	(0.178)	(0.178)		
$\rm NH_2^+$ group								
$n_{ m N1Ow}$	4.31	4.28	4.23	4.16	4.11	4.04		
$(r_{ m N1Ow}/ m nm)$	(0.298)	(0.298)	(0.298)	(0.298)	(0.298)	(0.300)		
$n_{ m H8Ow}$	0.87	0.86	0.85	0.84	0.82	0.81		
$(r_{ m H8Ow}/ m nm)$	(0.175)	(0.175)	(0.175)	(0.175)	(0.175)	(0.175)		
$n_{ m H9Ow}$	0.79	0.79	0.77	0.76	0.74	0.73		
$(r_{ m H9Ow}/ m nm)$	(0.170)	(0.170)	(0.170)	(0.173)	(0.173)	(0.173)		
		Ring ca	rbon ator	ns				
$n_{ m C2Ow}$	3.91	3.88	3.85	3.80	3.63	3.58		
$(r_{ m C2Ow}/ m nm)$	(0.340)	(0.340)	(0.340)	(0.340)	(0.340)	(0.340)		
$n_{\rm C3Ow}$	5.46	5.27	5.39	5.33	5.29	5.23		
$(r_{ m C3Ow}/ m nm)$	(0.353)	(0.353)	(0.353)	(0.353)	(0.353)	(0.353)		
$n_{\rm C4Ow}$	6.51	6.48	6.42	6.34	6.30	6.23		
$(r_{ m C4Ow}/ m nm)$	(0.350)	(0.350)	(0.350)	(0.350)	(0.350)	(0.350)		
$n_{ m C5Ow}$	6.72	6.68	6.61	6.51	6.45	6.36		
$(r_{ m C5Ow}/ m nm)$	(0.323)	(0.323)	(0.323)	(0.323)	(0.323)	(0.323)		
Hydrophilic sites: $n_{\rm h} = n_{\rm O1Ow} + n_{\rm O2Ow} + n_{\rm N1Ow}$								
$n_{ m h}$	18.39	18.39	18.18	17.96	17.80	17.58		
Na <sup>+</sup> -carboxylate interactions								
$n_{\rm O1Na}$		·	0.04			0.14		
$(r_{ m O1Na}/ m nm)$			(0.258)			(0.258)		
	$\mathrm{Cl}^{2}$	NH <sub>2</sub> <sup>+</sup> gr	oup intera	actions				
$n_{ m H8Cl}$		20	0.02			0.08		
$(r_{ m H8Cl}/~ m nm)$			(0.188)			(0.188)		
$n_{\rm H8Cl}$			0.02			0.0.08		
$(r_{ m H8Cl}/~ m nm)$			(0.183)			(0.183)		

## 4 Supplementary Figures



Figure S1: (a) Cylindrical distribution functions,  $g_{\text{pyr}-\text{Ow}}(z)_R$  and  $g_{\text{pyr}-\text{Hw}}(z)_R$  (R = 0.126 nm), for H<sub>2</sub>O hydrating the pyrrolidine ring of Pro at  $c(\text{Pro}) \rightarrow 0$ ; (b) Concentration dependence of  $g_{\text{pyr}-\text{Ow}}(z)_R$  for aqueous L-proline solutions.



Figure S2: Spectra of (a) relative permittivity,  $\varepsilon'(\nu)$ , and (b) dielectric loss,  $\varepsilon''(\nu)$ , of aqueous L-proline solutions at 25 °C and concentrations c(Pro) / M = 0 (1), 0.395 (2), 0.981 (3), 1.944 (4), 3.805 (5), 5.569 (6). Symbols show experimental data, the lines give fits with the D+D+D model.



Figure S3: Relaxation amplitudes of the solute mode,  $S_1$  ( $\blacktriangle$ ), of slow water,  $S_s = S_2$  ( $\blacktriangledown$ ), and of bulk-like water,  $S_b$  ( $\bigcirc$ ) of aqueous L-proline solutions at 25 °C and solute concentrations c(Pro). Lines are guide to the eye; the open symbol is pure water.



Figure S4: Relaxation times of the solute mode,  $\tau_1$  ( $\blacktriangle$ ), of slow water,  $\tau_s = \tau_2$  ( $\triangledown$ ), and of the cooperative relaxation of bulk-like water,  $\tau_3$  ( $\bigcirc$ ) of aqueous L-proline solutions at 25°C and solute concentrations c(Pro). Lines are guide to the eye; the open symbol is pure water.



Figure S5: Minimum-energy structures of  $\text{Pro}\cdot n\text{H}_2\text{O}$  (n = 0...4) complexes and their associated dipole moments,  $\mu$ , obtained with Gaussian (B3LYP/cc-pVDZ level with C-PCM solvation model).<sup>10,11</sup> The arrow indicates the dipole direction.



Figure S6: Pair distribution functions  $g_{\rm N1Ow}(r)$  (solid lines) and  $g_{\rm N1Hw}(r)$  (broken lines) of aqueous L-proline solutions at 25 °C.



Figure S7: (a) Pair distribution functions  $g_{O1Ow}(r)$  (solid lines) and  $g_{O1Hw}(r)$  (broken lines) of aqueous L-proline solutions at 25 °C. (b) Corresponding functions  $g_{O2Ow}(r)$  (solid lines) and  $g_{O2Hw}(r)$  (broken lines).



Figure S8: Pair distribution functions  $g_{\rm H8Ow}(r)$  (a) and  $g_{\rm H9Ow}(r)$  (b) of aqueous L-proline solutions at 25 °C.



Figure S9: Spectra of (a) relative permittivity,  $\varepsilon'(\nu)$ , and (b) dielectric loss,  $\varepsilon''(\nu)$ , of solutions of NaCl in 0.6 M aqueous L-proline at 25 °C and concentrations c(NaCl) / M = 0, 0.205, 0.999, 1.513, 2.023 increasing in arrow direction. Symbols show experimental data (partly omitted for clarity), the lines give fits with the D+D+D+D model.



Figure S10: Experimental bulk-water amplitude,  $S_{\rm b}$  ( $\bullet$ ), and corresponding equilibrium amplitude after correction for kinetic depolarization,  $S_{\rm b}^{\rm eq}$  ( $\blacktriangle$ ), of NaCl solutions of concentration  $c({\rm NaCl})$  in 0.6 M aqueous L-proline at 25 °C. Also included is the amplitude,  $S_{\rm w}$ , expected from the analytical water concentration. The difference  $S_{\rm w} - S_{\rm b}^{\rm eq}$  yields the total concentration of bound water.



Figure S11: Cylindrical distribution functions,  $g_{\text{pyr-Ow}}(z)_R$  (R = 0.126 nm), for H<sub>2</sub>O hydrating the pyrrolidine ring of Pro at c(Pro) = 0.6 M and c(NaCl) = (0, 0.5 and 2.0) M.



Figure S12: (a) Pair distribution functions  $g_{O1Ow}(r)$  (solid lines) and  $g_{O1Hw}(r)$  (broken lines) of NaCl solutions in 0.6 M aqueous L-proline at 25 °C. (b) Corresponding functions  $g_{N1Ow}(r)$  (solid lines) and  $g_{N1Hw}(r)$  (broken lines).



Figure S13: (a) Pair distribution functions  $g_{O1Na}(r)$  and  $g_{O2Na}(r)$  of 0.5 M and 2.0 M NaCl in 0.6 M aqueous L-proline at 25 °C. (b) Corresponding functions  $g_{H8Cl}(r)$  and  $g_{H9Cl}(r)$ .



Figure S14: Effective dipole moment,  $\mu_{\text{eff}}$  ( $\bullet$ ), of L-proline-ion aggregates as a function of NaCl concentration, c(NaCl), in 0.6 M L-proline(aq) at 25 °C obtained from the experimental amplitude  $S_1$  corrected for ion-cloud relaxation. Also shown are the minimum-energy structures of pro·Na<sup>+</sup>, pro·Cl<sup>-</sup> and pro·NaCl aggregates and their dipole moments obtained with Gaussian (B3LYP/cc-pVDZ level with C-PCM solvation model).<sup>10,11</sup> The arrow indicates the dipole direction.

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