# Supporting Information for:

## Electronic Structures of Intermolecular Hydrogen Bond Contacts on Solute in Aqueous Solution: Glycine as a Working Prototype

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#### I. Computational details

The geometrical and vibrational features of all modeling clusters were investigated by the density functional theory (DFT) calculations. The hybrid functional of BHandHLY<sup>1</sup> is used since the functional has been shown to be accurately describing the amino acid systems that are rich in H-bonds.<sup>2</sup> The 6-31++G(d,p) basis set is used in most cases, but the 6-31++G(d) basis set may also be selected for better comparability with the literature studies.<sup>3–5</sup> The BHandHLYP vibrational frequencies are scaled by a uniform factor of 0.92 in order to account for the anharmonicity effect and be comparable with the experimental data, as suggested in the literatures.<sup>6–8</sup>

To test the basis set dependence of the computational results, two additional basis sets,  $6-311++G^*$  and  $6-311++G^{**}$  have been used. The differences in the structural parameters and relative energies obtained by different basis sets are small. For example, corresponding to Figure 4 in the main text, Figure S1 illustrates the uncertainty for the computed effective energies and potentials. The bars for the numerical uncertainty refer to the difference between the  $6-31++G^*$  results and the averages of the three basis sets of  $6-31++G^{**}$ ,  $6-311++G^*$  and  $6-311++G^{**}$ . It is clear from Figure S1 that the numerical uncertainty is negligible for the energy scale of interest. The numerical uncertainties for the interested IR modes are also small, but not negligible. They are indicated in Figure 3 of the main text.

All calculations reported here were performed using the Gaussian 03/09 suite of programs.<sup>9,10</sup>



**FIG. S1**. The basis set dependence of the relative effective energies and potentials of modeling clusters. Fig. S1 corresponds to Figure 4 of the main text. More description may be found in the main manuscript.

### II. Tables : Structures and IR frequencies of relevant modeling clusters

### **Table Overview:**

Table S1 is used to support the H-bond models of  $\Pi_1(H)$ ,  $\Pi_3(N)$ , and  $\Pi_2(O)$  discussed in the main text.

Table S2 supports the detailed discussion on  $\Pi_2(O)$  of Figure 3 in the main text.

Table S3 and S4 are the basis for the discussion on zGly clusters in the main text.

Table S5 provides the testing results for phenylalanine and alanine in aqueous solution.

**Table S1:** Structures and characteristic IR frequencies of representative modeling clusters, dGlyW<sub>n</sub>, pGlyW<sub>n</sub>. Frequencies (cm<sup>-1</sup>) are calculated at the BHandHLYP/6-31++G(d,p) level.  $v_{ad}$ (NH<sub>3</sub><sup>+</sup>) and  $v_{sd}$ (NH<sub>3</sub><sup>+</sup>) denote the NH<sub>3</sub><sup>+</sup> asymmetric and symmetric deformations,  $v_{sc}$ (NH<sub>2</sub>) denotes the NH<sub>2</sub> scissor, and v(C=O) and v(C-O(H)) denote the C=O and C-O(H) stretching vibrations, respectively. The experiment data are from Refs [11,12]. Note that due to the sensitivity of the N-H vibrations to the H-bond arrangement, the most satisfying agreement between the theory and experiment is found in clusters with a second hydration shell.

Geometries	Parameters (Degree)	Frequencies
	pGlyW <sub>2</sub> → Π <sub>1</sub> (H): ∠(O-H···O <sub>W</sub> )=179 ∠(N-H···O <sub>W</sub> )=173	
	$dGlyW_1 \rightarrow \Pi_3(N):$ $\angle (C-N\cdots H_W)=116$ $\angle (H-N\cdots H_W)=109,114$	
	pGlyW <sub>9</sub> → Π <sub>2</sub> (O): ∠(C-O···H <sub>W</sub> )=124 ∠(H-O···H <sub>W</sub> )=123 Φ[H-O-C···H <sub>W</sub> ]=165	v(C-O(H)) <sub>calc</sub> =1257, <sub>exp.</sub> ~1258

$pGlyW_{10} \rightarrow \Pi_{1}(H)\Pi_{2}(O)$ $\angle (N-H_{3}\cdots O_{W})=167,163,151$ $\angle (O-H\cdots O_{W})=162$ $\angle (C=O\cdots H_{W})=120,154$ $\Phi[H_{W}\cdots O=C\cdots H_{W}]=175$ $\angle (C-(H)O\cdots H_{W})=114$ $\Phi[H-O-C\cdots H_{W}]=145$	$v_{ad}(NH_3^+)_{calc} = 1617/1631, exp. \sim 1615$ $v_{sd}(NH_3^+)_{calc} = 1511, exp. \sim 1512/1521$ $v(C=O)_{calc} = 1745, exp. \sim 1740$ $v(C=O(H))_{calc} = 1245, exp. \sim 1258$
$dGlyW_{11} \rightarrow \Pi_{1}(H)\Pi_{3}(N)\Pi_{2}(O)$ $\angle (N-H_{2}\cdots O_{W})=149,146$ $\angle (C-N\cdots H_{W})=121$ $\angle (C=O\cdots H_{W})=119,124,132,144$ $\Phi[H_{W}\cdots O=C\cdots H_{W}]=172,158$	$v_{sc}(NH_2)_{calc} = ~1588, exp. ~1590-1600$ $v_{asym}(CO_2^{-})_{calc} = 1567, exp. ~1561$ $v_{sym}(CO_2^{-})_{calc} = 1409, exp. ~1404$

**Table S2:** Structures and characteristic IR frequencies of representative modeling clusters,  $dGlyW_n$ , for deprotonated (anionic) Gly. Frequencies (cm<sup>-1</sup>) are calculated at the BHandHLYP/6-31++G(d,p) level.  $v_{as}(CO_2^{-})$  and  $v_{ss}(CO_2^{-})$  denote the CO<sub>2</sub><sup>-</sup> asymmetric and symmetric stretching vibrations, respectively. Band gaps in the Table refers the differences in  $v_{as}(CO_2^{-})$  and  $v_{ss}(CO_2^{-})$ ; The structure of  $dGlyW_{4d}$  is from Ref [13]; The experiment data are taken from Ref [11].

Chustans		Frequencies		
Clusters	Geometries	$v_{\rm as}(\rm CO_2^-)$	$v_{\rm ss}(\rm CO_2^{-})$	Band gap
dGlyW <sub>0</sub>		1619	1351	268
<i>d</i> GlyW <sub>1a</sub>		1600	1369	231
dGlyW <sub>1b</sub>		1616	1355	260
dGlyW <sub>1c</sub>		1605	1349	257
dGlyW <sub>2a</sub>		1599	1383	216
dGlyW <sub>2b</sub>	8.000	1583	1370	213

dGlyW <sub>3a</sub>	1385	1386	200
dGlyW <sub>3b</sub>	1576	1386	190
dGlyW <sub>4a</sub>	1578	1392	186
dGlyW <sub>4b</sub>	1574	1393	181
dGlyW <sub>4c</sub>	1574	1387	187
dGlyW <sub>4d</sub>	1575	1381	194
dGly₩₅	1579	1409	170

dGlyW <sub>6</sub>	1573	1410	163
Exp.	1561	1404	157

**Table S3:** IR frequencies (freq) and intensities (inten) for zGlyW<sub>8</sub> of Fig. 2 of the main text. Frequencies (cm<sup>-1</sup>) are calculated at the BHandHLYP/6-31++G(d) level. Only frequencies in the range of 1300–1700 cm<sup>-1</sup> that are relevant to the CO<sub>2</sub><sup>-</sup> vibrations are shown here. The experiment data are from refs [3,11].

	Exp.		
freq	inten	Assignment	
1667	56	$NH_3$ ad. + $H_2O$ bend + $CO_2$ as.	
1660	43	$NH_3$ ad. + $H_2O$ bend + $CO_2$ as.	
1639	258	$NH_3$ ad. + $CO_2$ as. + $H_2O$ bend	1637
1623	61	$CO_2$ as. + $NH_3$ ad. + $H_2O$ bend	
1583	528	$CO_2$ as. + $H_2O$ bend	1598/1599
1545 1433	135 13	$NH_3$ sd + $H_2O$ bend $CO_2$ ss. + $CH_2$ sc.	1510
1412	118	$CO_2$ ss. + $CH_2$ sc.	1412/1413
1329	108	$CH_2 r. + CO_2 ss.$	1331

**Table S4:** Relative cluster total energies ( $E_{tot}$ ), solute energies ( $E_{eff}$ ) and solute-solvent interaction energies ( $V_{GW}$ ) of model clusters of zGly and nGly. All energies are in kcal/mol and calculated at the BHandHLYP/6-31++G(d,p) level with the BSSE corrections. The structure of zGlyW8 (z1) is obtained through our proposed H-bond model. The structure of reference modeling clusters, 8*Z*-*c*, 8*Z*-*a*, 8*Z*-*e*, 8*N*8-*a*, 8*N*1-*a*, 8*N*6-*a*, and 8*N*1-*d*, are from Ref [14].

No.	Clusters	E <sub>tot</sub>	$E_{\rm eff}$	V <sub>GW</sub>
<i>z</i> 1	zGlyW <sub>8</sub> (z1)	0.0	0.0	-97.7
<i>z</i> 2	8Z-c	-4.7	8.9	-88.6
<i>z</i> 3	8Z-a	-6.0	13.7	-86.2
<i>z</i> 4	8Z-e	-5.3	15.3	-80.1
<i>c</i> 1	8N8-a	-7.4	34.3	-36.5
<i>c</i> 2	8N1-a	-6.0	39.6	-26.9
<i>c</i> 3	8N6-a	-6.5	40.3	-27.2
<i>c</i> 4	8N1-d	-5.5	39.7	-26.7

<i>P</i> henylalanine			Alanine		
			e e e		
∠(C=C	)…H <sub>W</sub> )=123,1	26,127,130	∠(C=0	····H <sub>w</sub> )=124,1	25,131,132
$\Phi[H_W \cdots O = C \cdots H_W] = 161,157$		$\Phi[H_W$	···O=C···H <sub>w</sub> ]	=171,168	
	$v_{as}(C-O)$	$v_{\rm ss}(C-O)$		$v_{as}(C-O)$	$v_{\rm ss}({\rm C-O})$
Theo.	1563	1398	Theo.	1571	1402
Exp.	1560	1410	Exp.	~1560	~1410

**Table S5:** Testing results for the proposed  $\Pi_2(O)$  state of solvated phenylalanine and alanine. The experiment data are from Refs [15,16].

**Table S6:** Testing results for formamide in aqueous solution and in solid/liquid phase. The experiment data are from Refs [17-19].



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