

An SFG Study of Interfacial Amino Acids at the Hydrophilic SiO₂ and Hydrophobic Deuterated Polystyrene Surfaces

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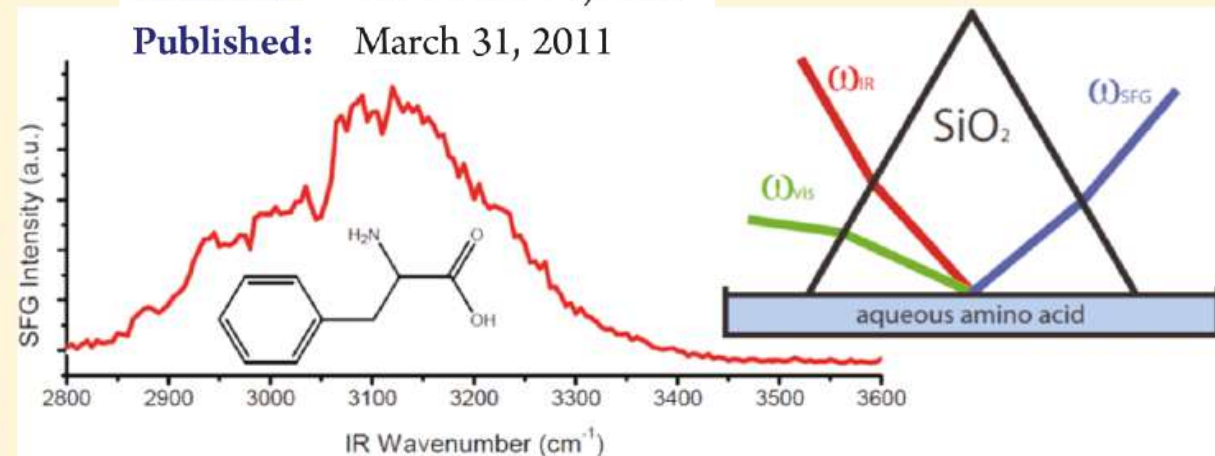
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ABSTRACT: Sum frequency generation (SFG) vibrational spectroscopy was employed to characterize the interfacial structure of **eight individual amino acids**—L-phenylalanine, L-leucine, glycine, L-lysine, L-arginine, L-cysteine, L-alanine, and L-proline—in aqueous solution **adsorbed at model hydrophilic and hydrophobic surfaces.** Specifically, SFG vibrational spectra were obtained for the amino acids at the solid–liquid interface between both **hydrophobic d₈-polystyrene (d₈-PS)** and **SiO₂ model surfaces** and phosphate buffered saline (PBS) **at pH 7.4.**

At the hydrophobic d₈-PS surface, seven of the amino acids solutions investigated showed clear and identifiable C–H vibrational modes, with the exception being L-alanine. In the SFG spectra obtained at the hydrophilic SiO₂ surface, no C–H vibrational modes were observed from any of the amino acids studied. However, it was confirmed by quartz crystal microbalance that amino acids do adsorb to the SiO₂ interface, and the amino acid solutions were found to have a detectable and widely varying influence on the magnitude of SFG signal from water at the SiO₂/PBS interface. This study provides the first known SFG spectra of several individual amino acids in aqueous solution at the solid–liquid interface and under physiological conditions.

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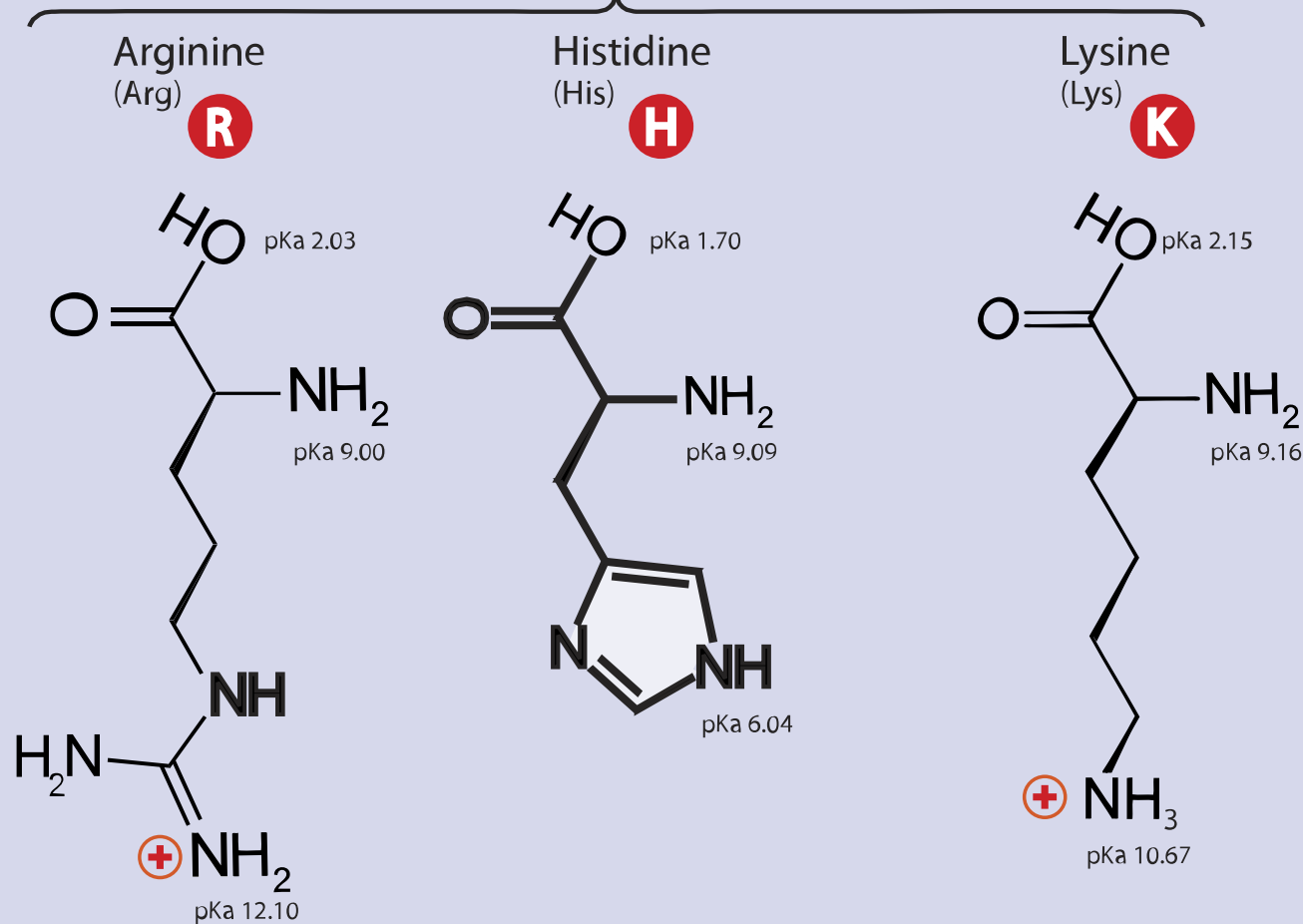


Twenty-One Amino Acids

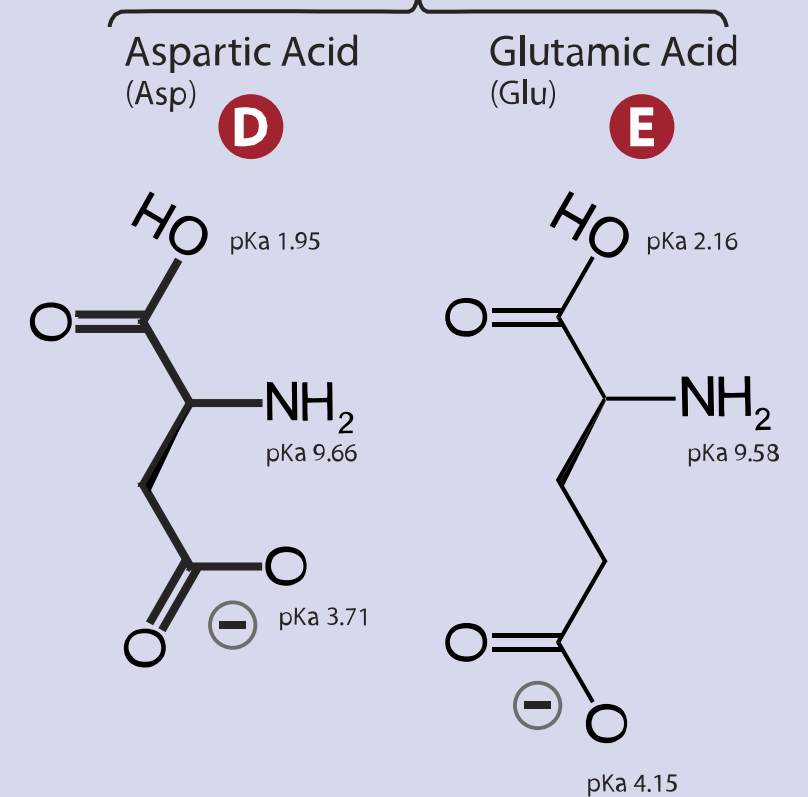
⊕ Positive ⊖ Negative
• Side chain charge at physiological pH 7.4

A. Amino Acids with Electrically Charged Side Chains

Positive

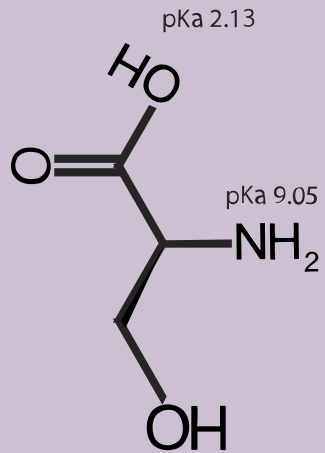


Negative

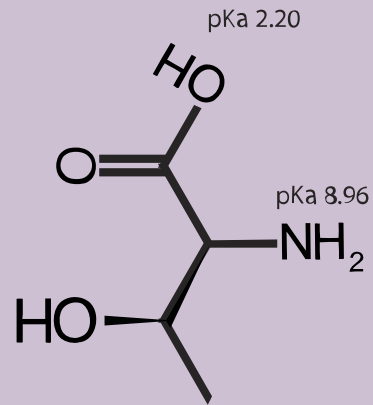


B. Amino Acids with Polar Uncharged Side Chains

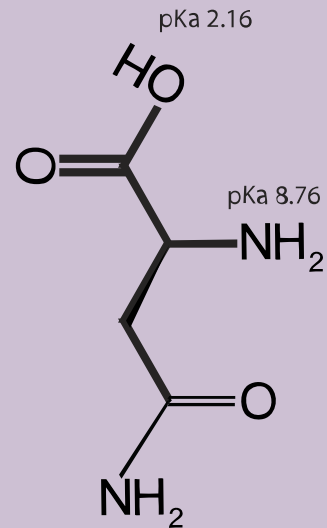
Serine
(Ser) **S**



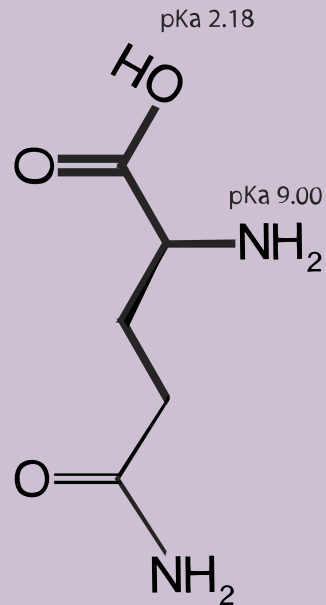
Threonine
(Thr) **T**



Asparagine
(Asn) **N**

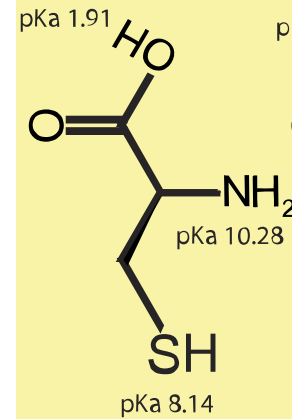


Glutamine
(Gln) **Q**

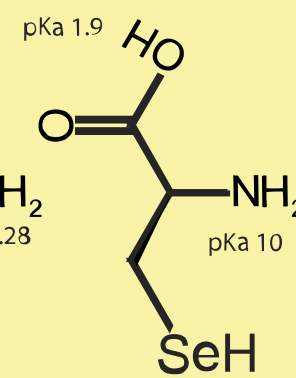


C. Special Cases

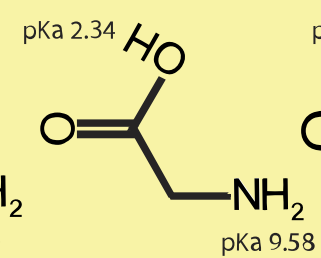
Cysteine
(Cys) **C**



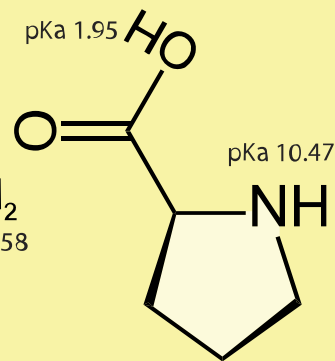
Selenocysteine
(Sec) **U**



Glycine
(Gly) **G**



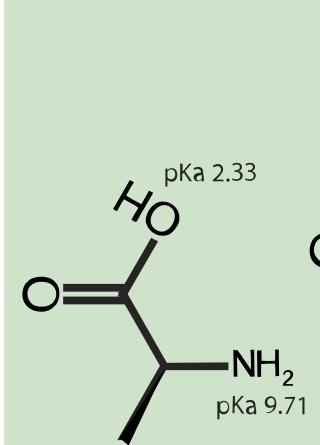
Proline
(Pro) **P**



D. Amino Acids with Hydrophobic Side Chain

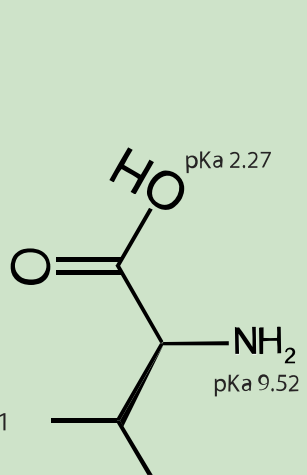
Alanine
(Ala)

A



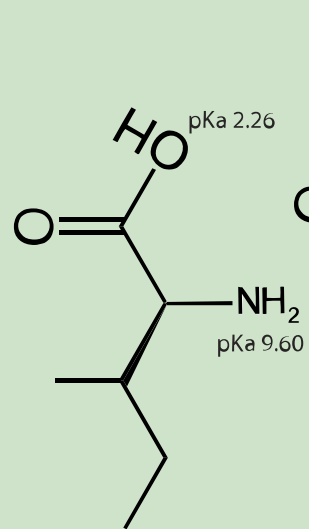
Valine
(Val)

V



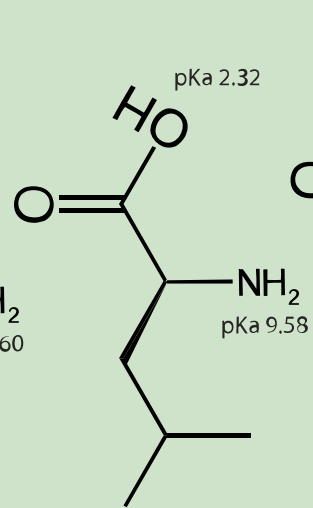
Isoleucine
(Ile)

I



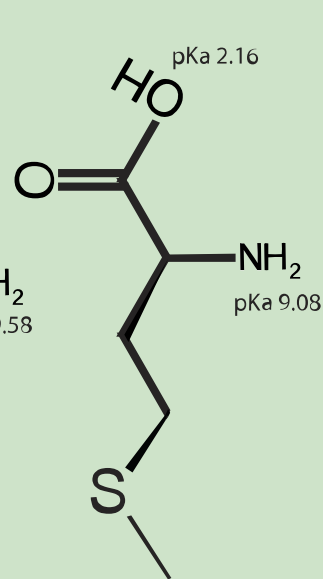
Leucine
(Leu)

L



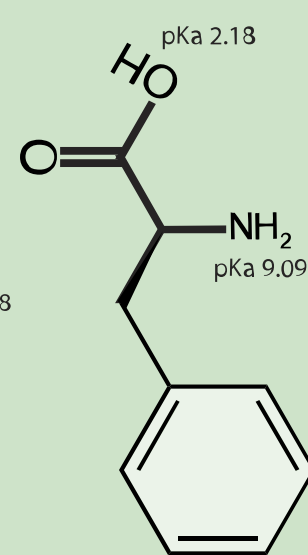
Methionine
(Met)

M



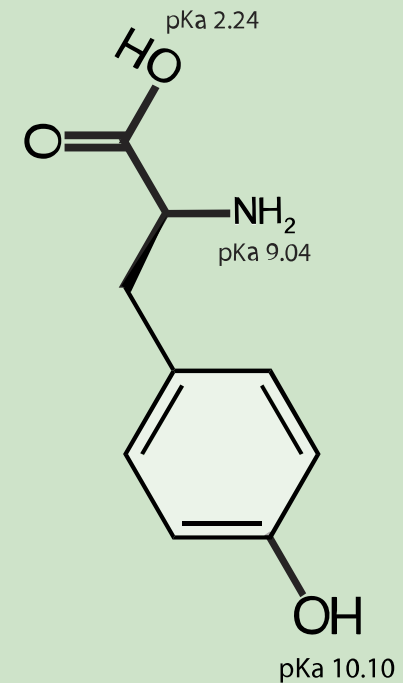
Phenylalanine
(Phe)

F



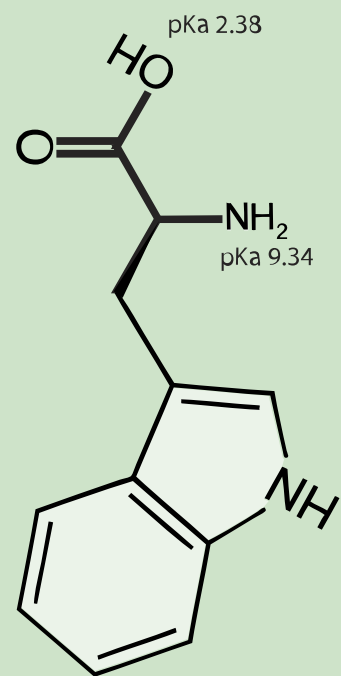
Tyrosine
(Tyr)

Y

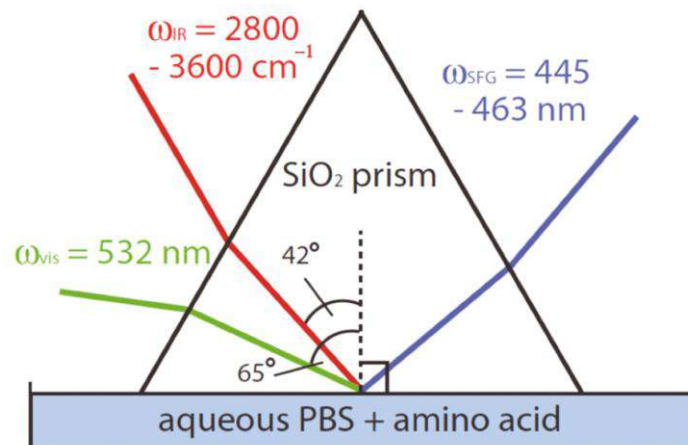


Tryptophan
(Trp)

W



Tool: Sum frequency
(Solid/Liquid Interface)



$$I_{\text{ssp}}^{\text{SFG}} = \frac{32\pi^3 \omega_s^2}{c^3} \frac{\text{Re}[\epsilon_1^{1/2}(\omega_s)]}{|\epsilon_1(\omega_s)| \text{Re}[\epsilon_1^{1/2}(\omega_{\text{Vis}})\epsilon_1^{1/2}(\omega_{\text{IR}})]} \times \left| \chi_{\text{yyz}}^{(2)}(\omega_s) \sec \beta \sin \beta_{\text{IR}} L^S L_T^{\text{Vis}} L_T^{\text{IR}} \frac{\epsilon_1^{1/2}(\omega_{\text{IR}})\epsilon_2^{1/2}(\omega_{\text{IR}})}{\epsilon_M(\omega_{\text{IR}})} \right|^2 I^{\text{Vis}} I^{\text{IR}} \quad (1)$$

where

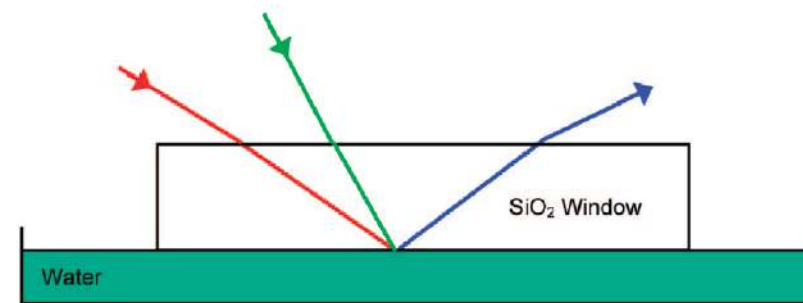
$$L^S = \frac{2\epsilon_1^{1/2}(\omega_s)\cos \beta}{\epsilon_2^{1/2}(\omega_s)\cos \gamma + \epsilon_2^{1/2}(\omega_s)\cos \beta} \quad (2)$$

$$L_T^{\text{Vis}} = \frac{2\epsilon_2^{1/2}(\omega_{\text{Vis}})\cos \beta_{\text{Vis}}}{\epsilon_2^{1/2}(\omega_{\text{Vis}})\cos \gamma_{\text{Vis}} + \epsilon_1^{1/2}(\omega_{\text{Vis}})\cos \beta_{\text{Vis}}} \quad (3)$$

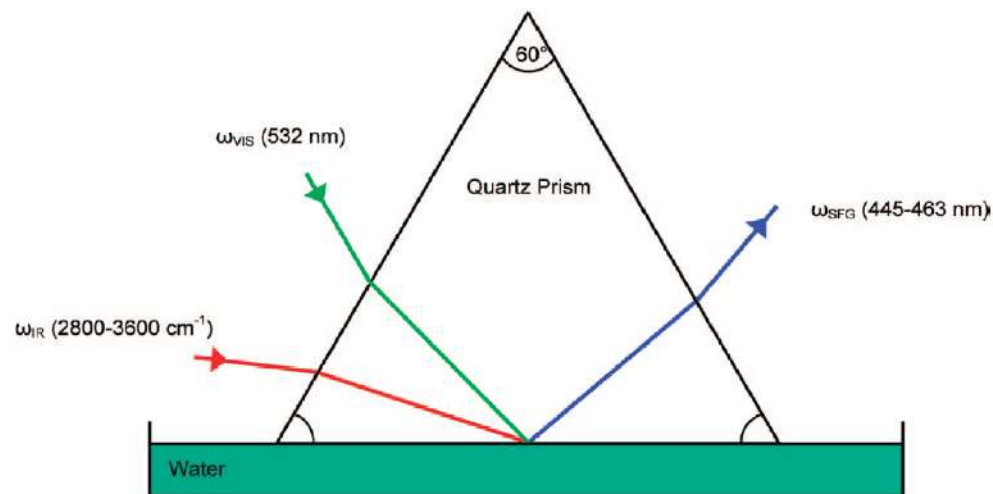
$$L_T^{\text{IR}} = \frac{2\epsilon_1^{1/2}(\omega_{\text{IR}})\cos \beta_{\text{IR}}}{\epsilon_2^{1/2}(\omega_{\text{IR}})\cos \beta_{\text{IR}} + \epsilon_1^{1/2}(\omega_{\text{IR}})\cos \gamma_{\text{IR}}} \quad (4)$$

Chosen angles:

- maximize the collection of SFG signal
- While minimizing wavelength dependent



(a)



(b)

Figure 2. Scheme of the (a) “slab” geometry and (b) “prism” geometry used in this work. See Table 1 for the respective angles of incidence.

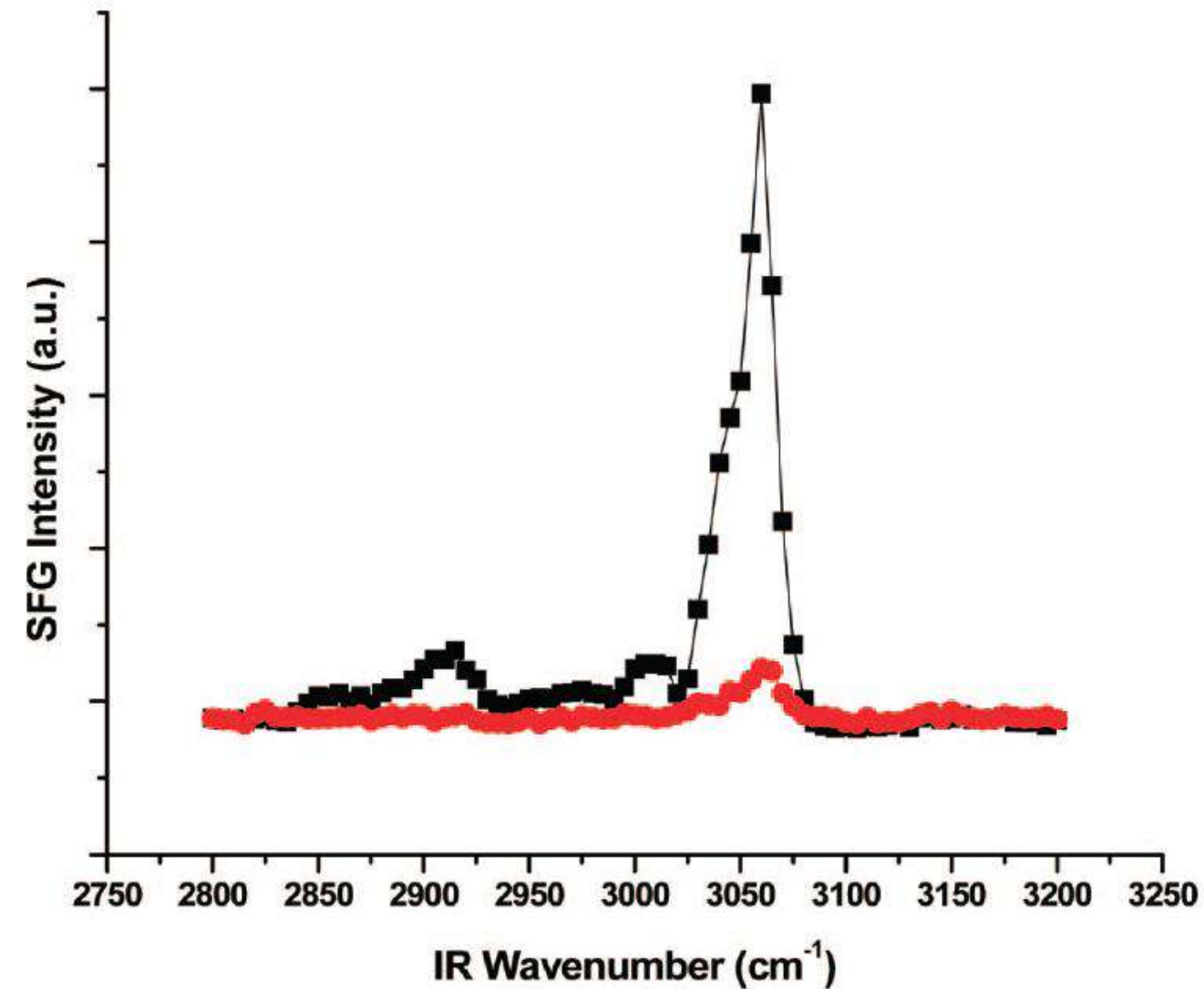


Figure 5. SFG spectra of the polystyrene/air interface in the slab (red circle) and prism (black squares) geometries. Note the increase in the SFG signal in the prism geometry. Additionally, vibrational modes not clearly seen in the slab geometry (due to the low signal-to-noise ratio) are clearly present in the prism geometry (e.g., the mode around 2915 cm^{-1}).

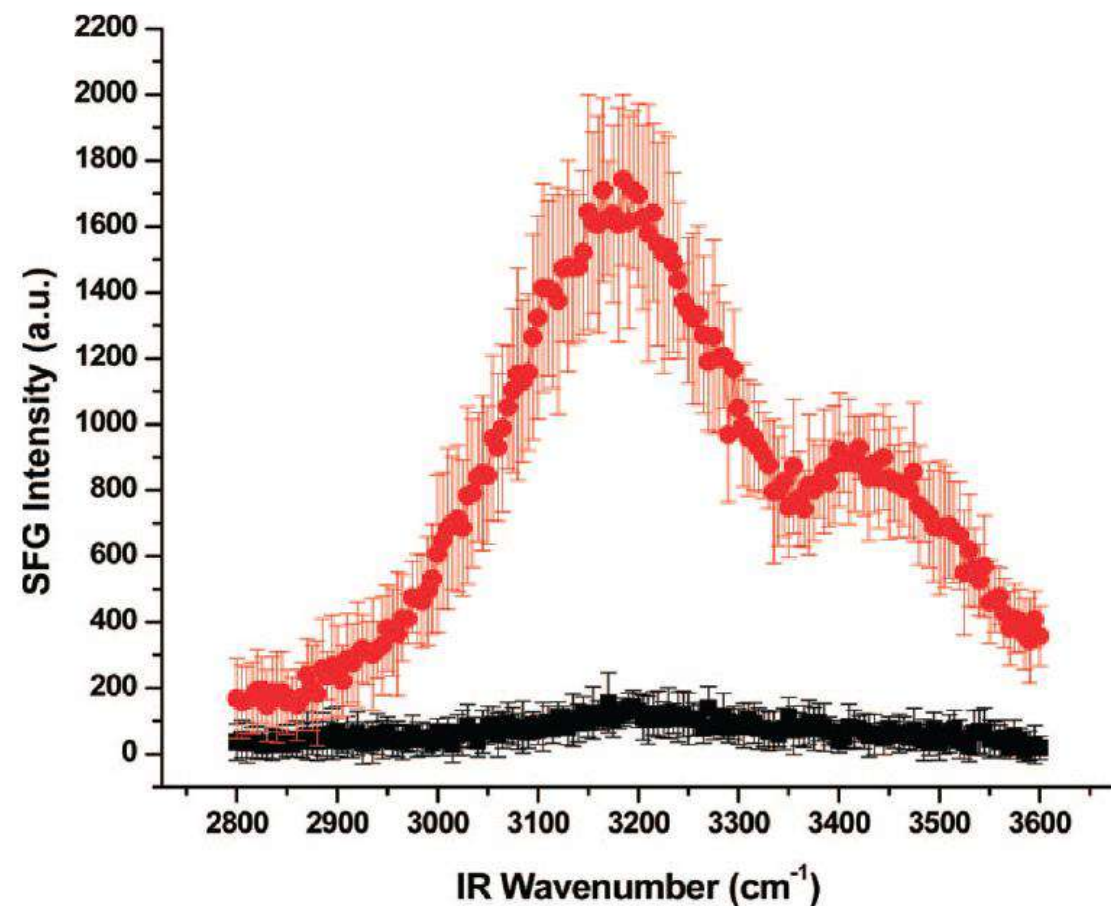


Figure 6. SFG spectra of the pure water/silica interface for both the prism (red circles) and the slab (black squares). Error bars come from the average of 5 scans of 200 shots per data point (for the prism) and 8 scans of 200 shots per data point (for the slab).

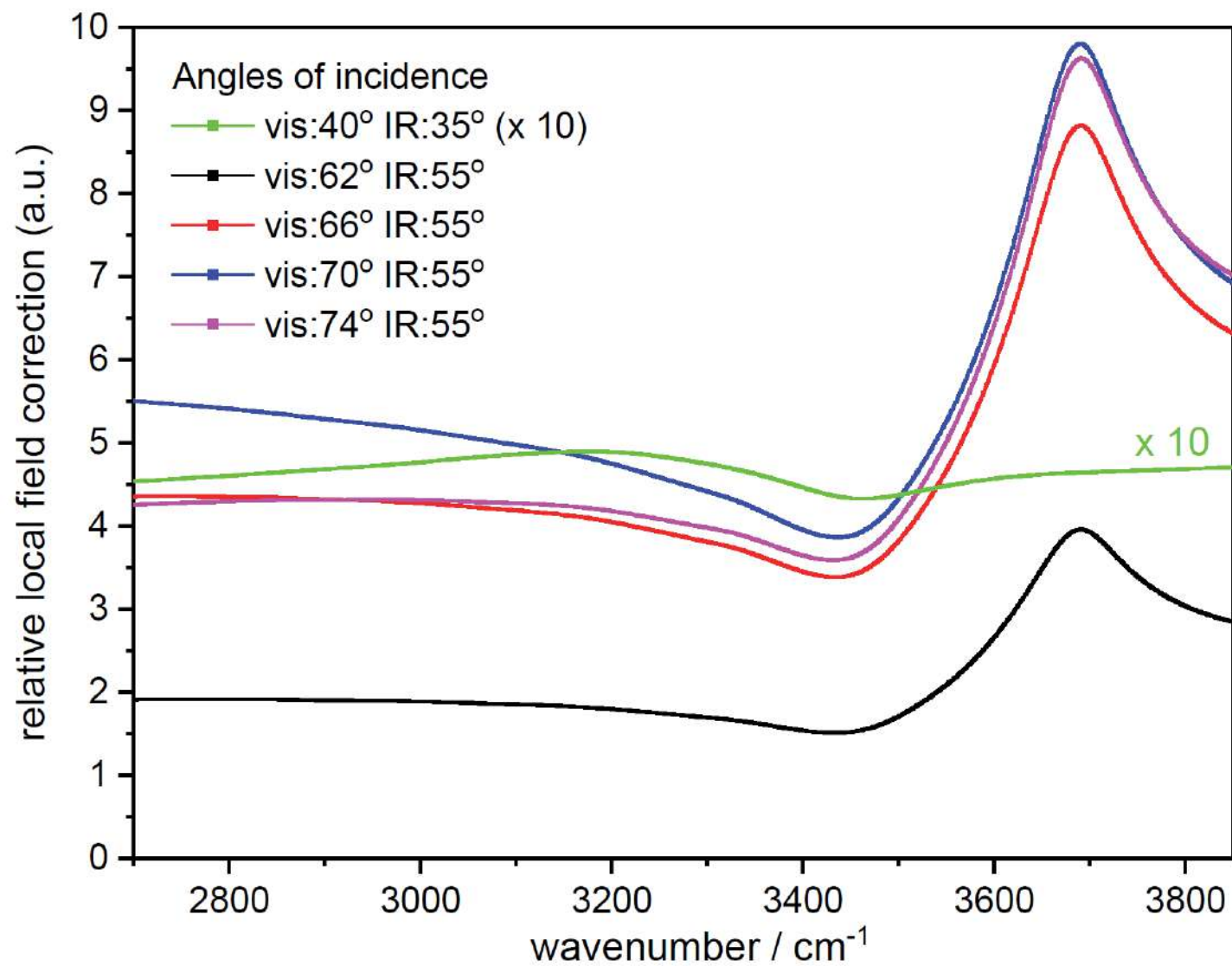


Fig. S5. IR dependence of the local field corrections for each of the experimental geometries studied. Note that the values for 40° visible AOI and 35° IR AOI experimental geometry, were multiplied by 10 for ease of comparison. The maximum enhancement in the overall spectra intensity is attained close to the critical angles.

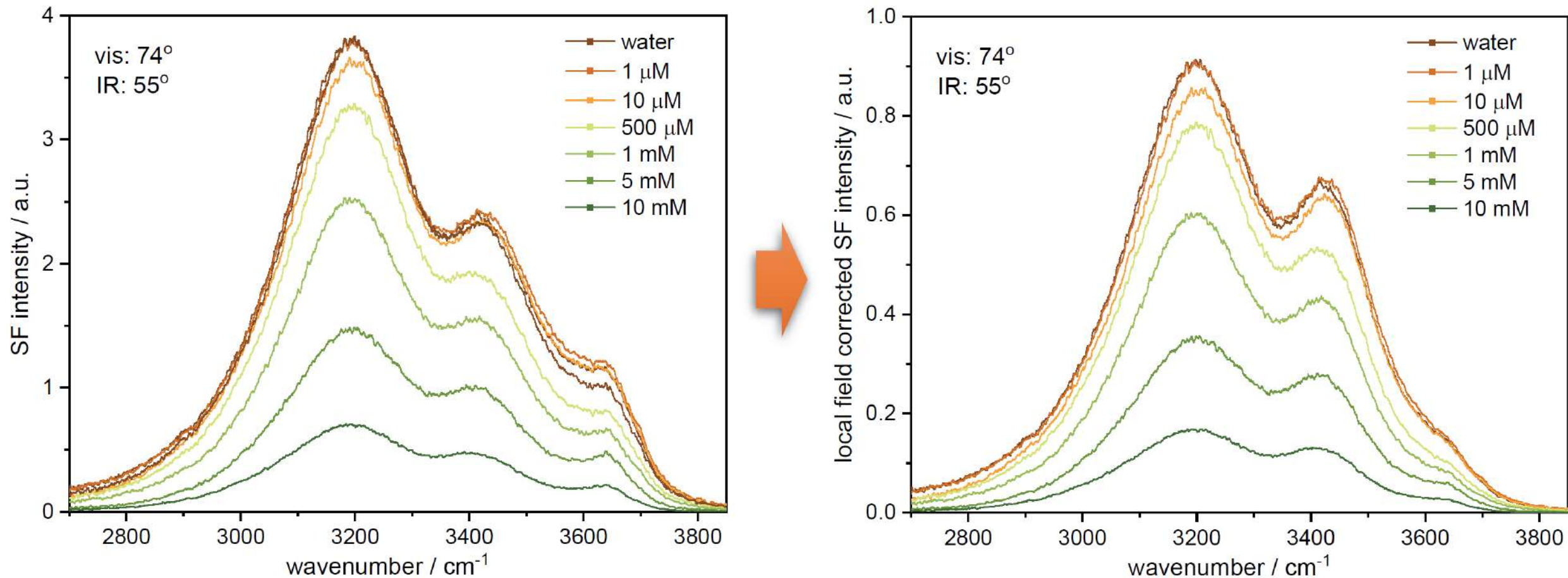
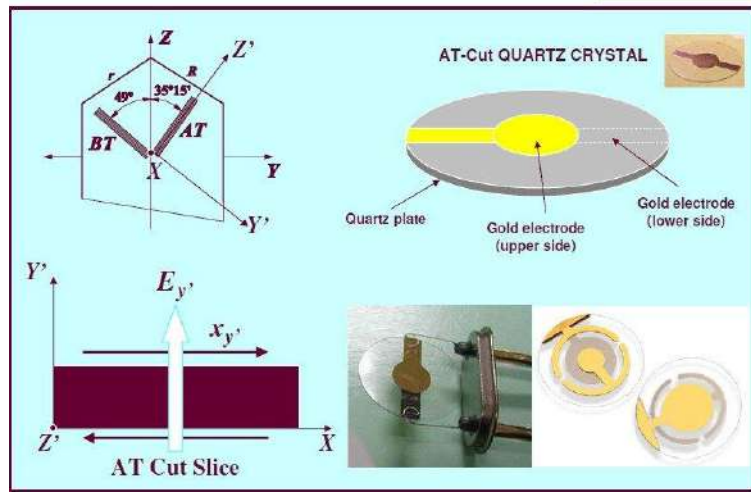


Fig. S6. (left) SF spectra collected in the ssp polarization combination as a function of NaCl for an IR angle of incidence of 55° and visible angle of 74°. (right) Local field corrected spectra at the same experimental geometry.

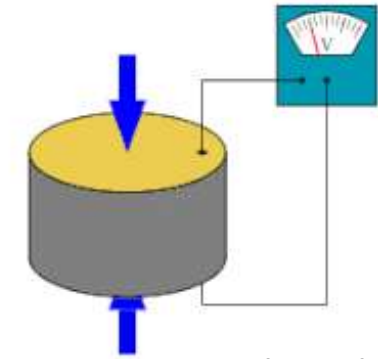
Complementary Tool:
Quartz Crystal Microbalance (QCM)

AT cut Quartz

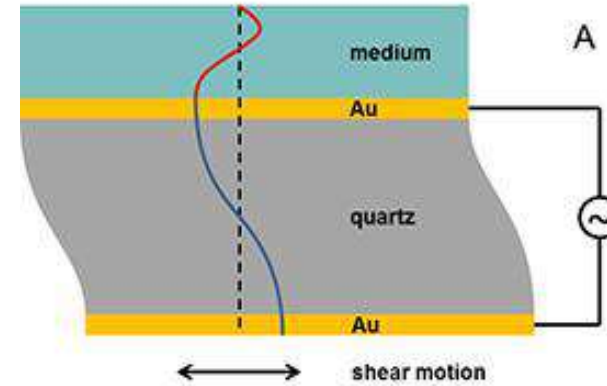


<https://www.tradekorea.com/product/detail/P416979/Electrochemical-research-Biosensor-crystal-.html>

Piezoelectric



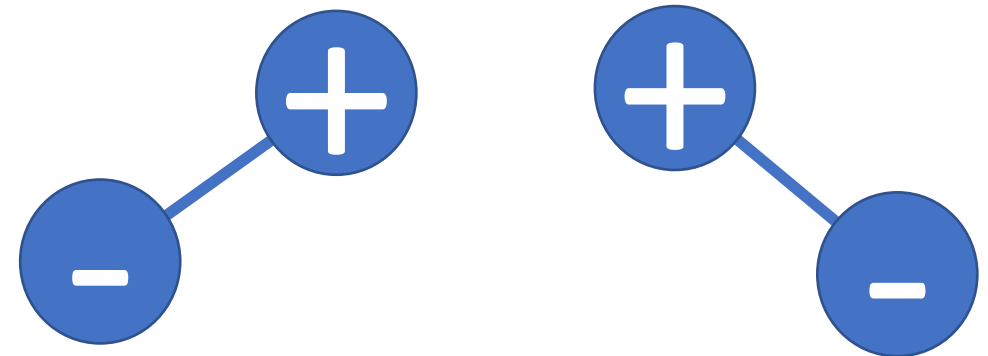
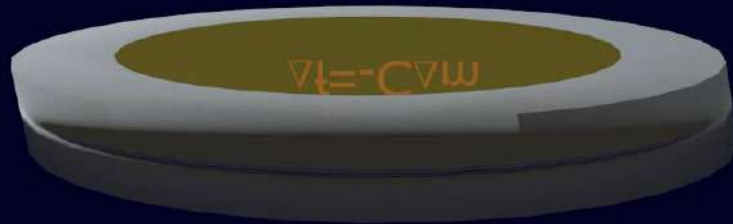
Wikipedia



<https://www.3t-analytik.de/technologies/qcm-d/what-qcm-d>

crystal oscillation frequency is proportional to mass

$$\Delta f = -C \Delta m$$

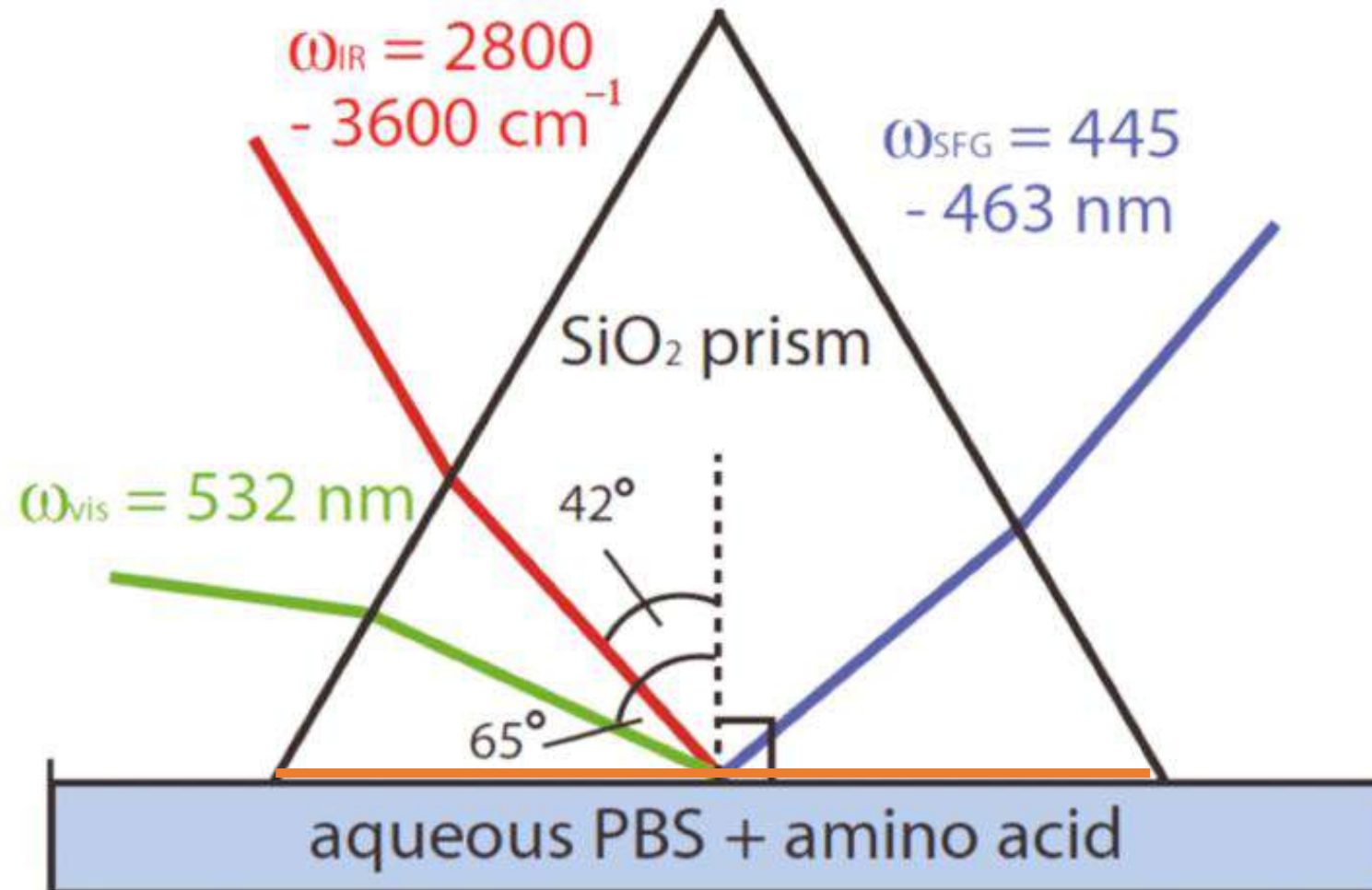


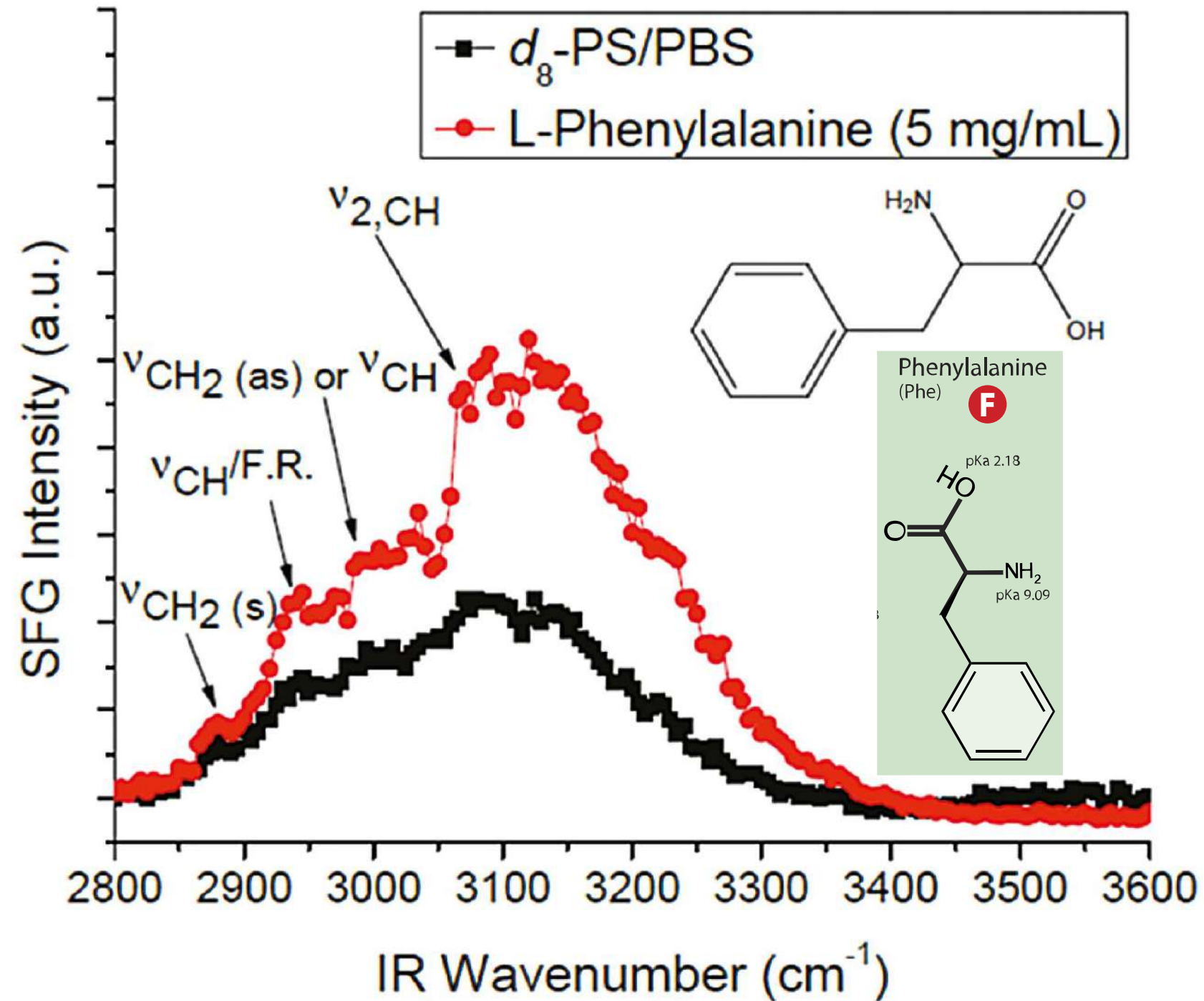
QCM results

Table 1. Adsorbed Mass at the Aqueous Liquid–Solid Interface Determined by QCM

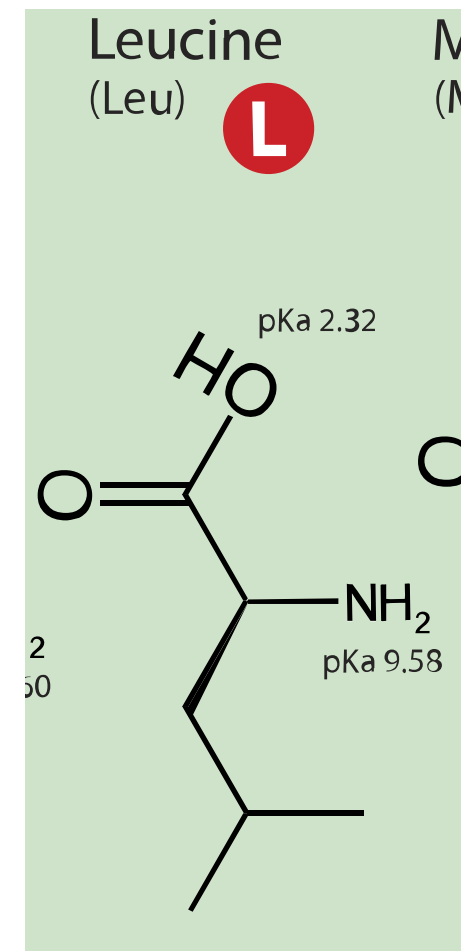
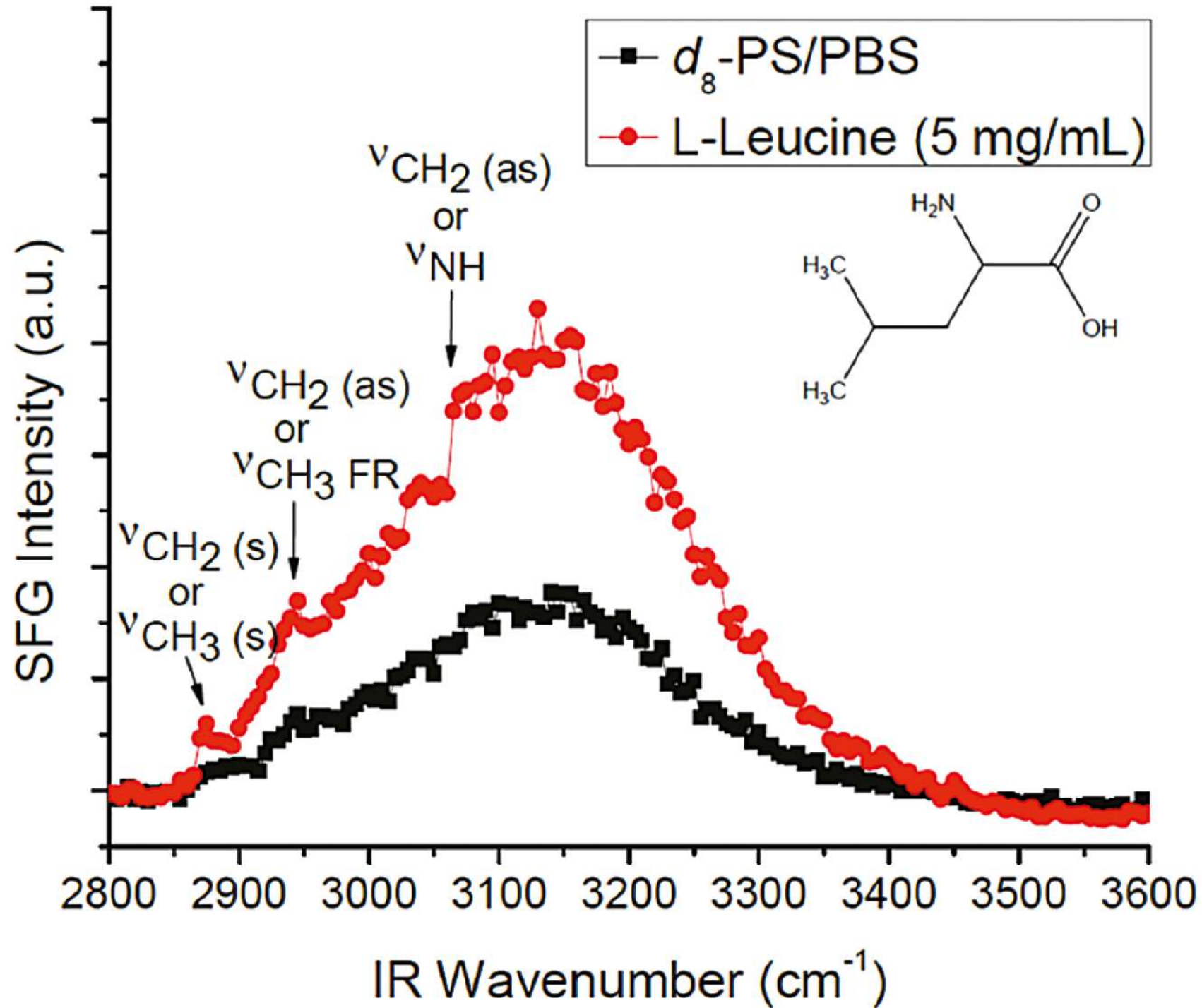
amino acid	concentration (mg/mL)	interface	adsorbed mass (ng/cm ²)	95% confidence interval (ng/cm ²)
L-phenylalanine	5	PS	57	4
L-phenylalanine	5	SiO ₂	63	4
L-arginine	15	SiO ₂	196	20
L-alanine	40	PS	469	33
L-alanine	40	SiO ₂	467	20
L-cysteine	25	SiO ₂	245	22

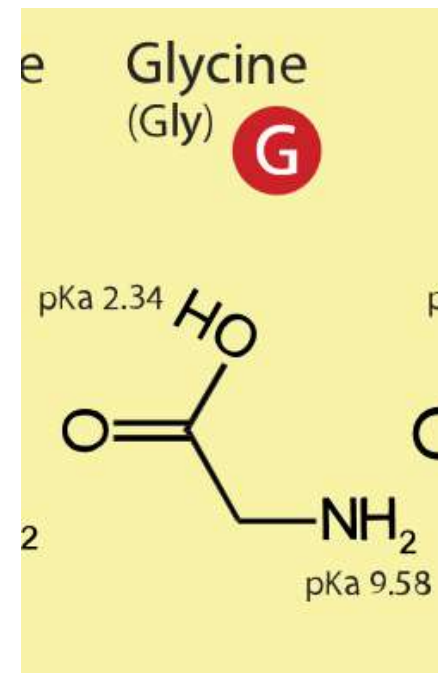
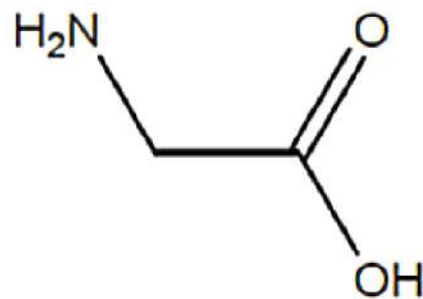
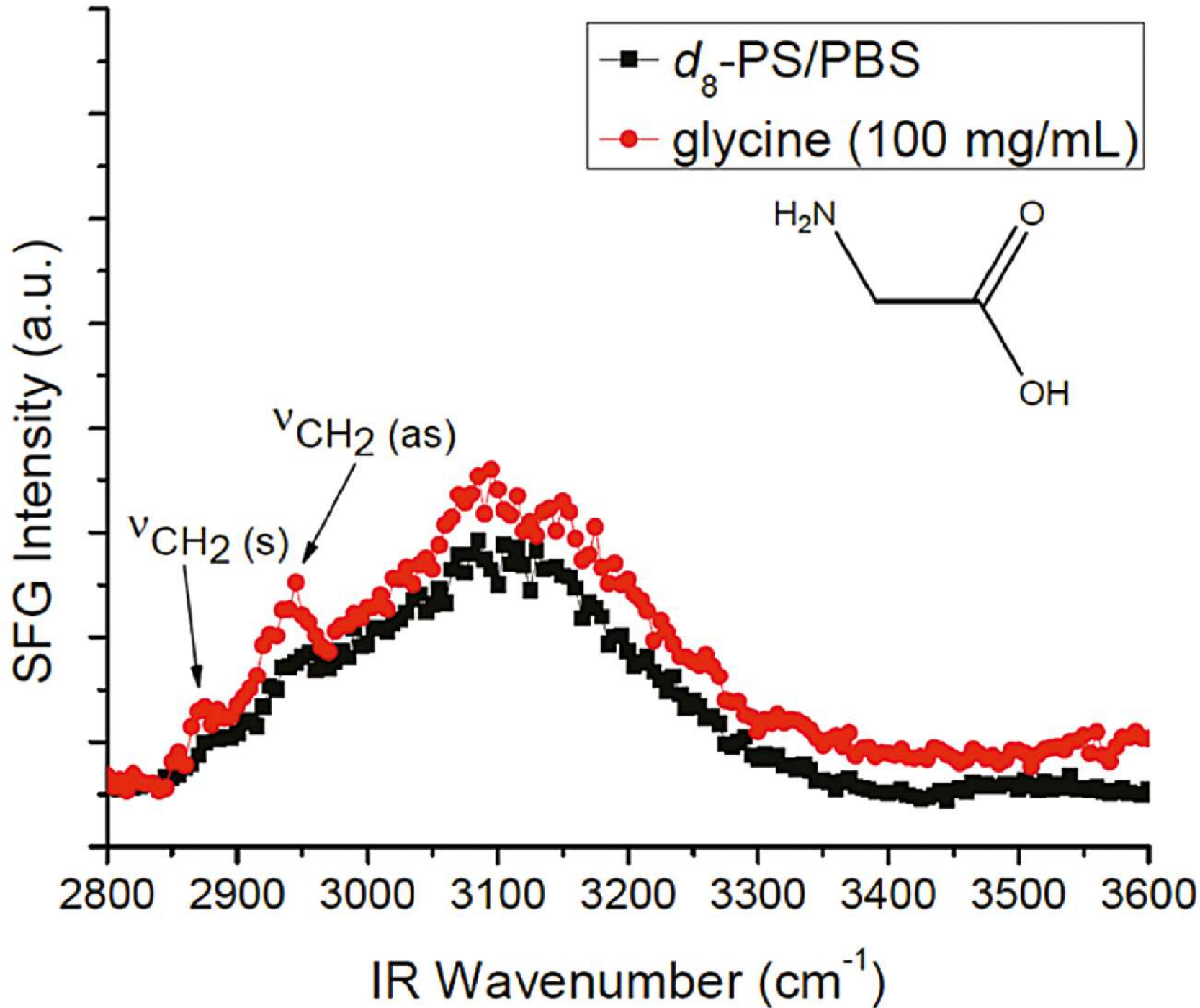
SFG results d-PS coated



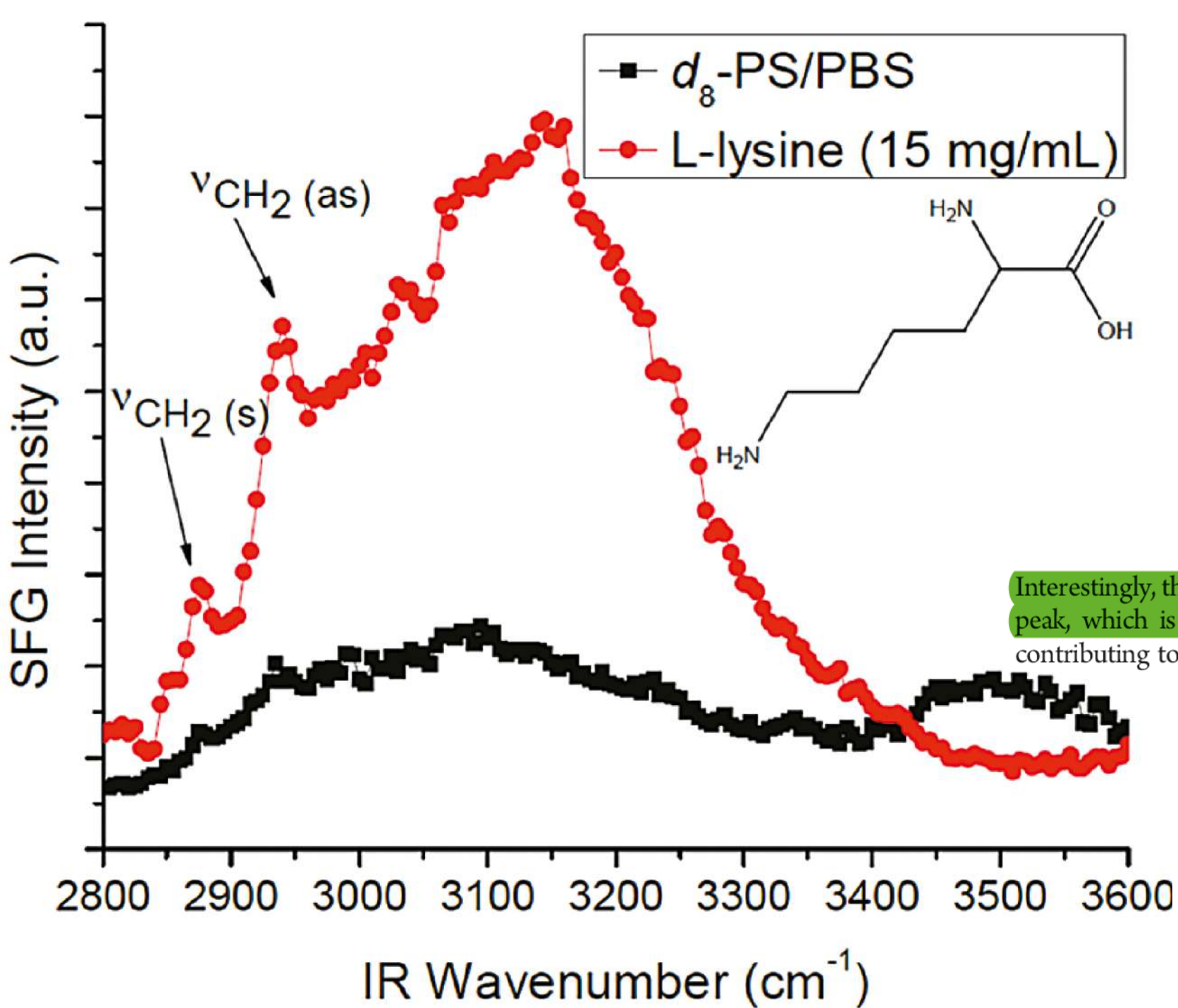


- Fig2: L-Phenylalanine
- 1points=500 laser shots.
- Larger OH due to adsorption of L-PA
- 2875/ cm^{-1} $\text{CH}_{2(s)}$
- 2939/ cm^{-1} CH and FR, CH connecting phenyl and amino acid backbone
- 2991/ cm^{-1} C-H combination band or $\text{CH}_{2(as)}$
- 3049/ cm^{-1} Aromatic C-H ν_2 stretch from phenyl
- QCM: adsorbed mass 57 $\text{ng}/\text{cm}^2 = 48\text{\AA}^2$ per molecule

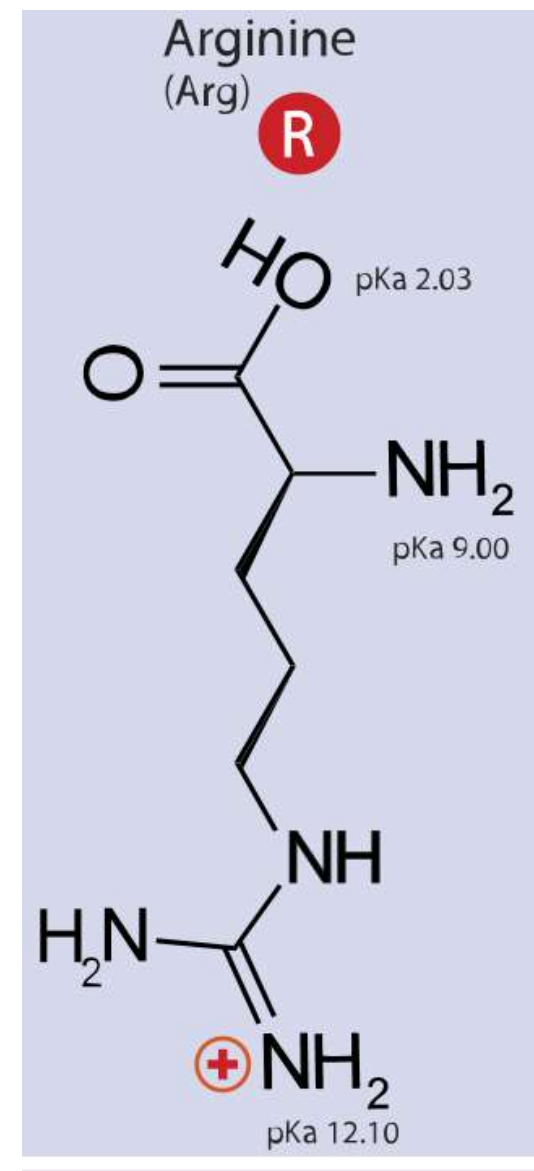
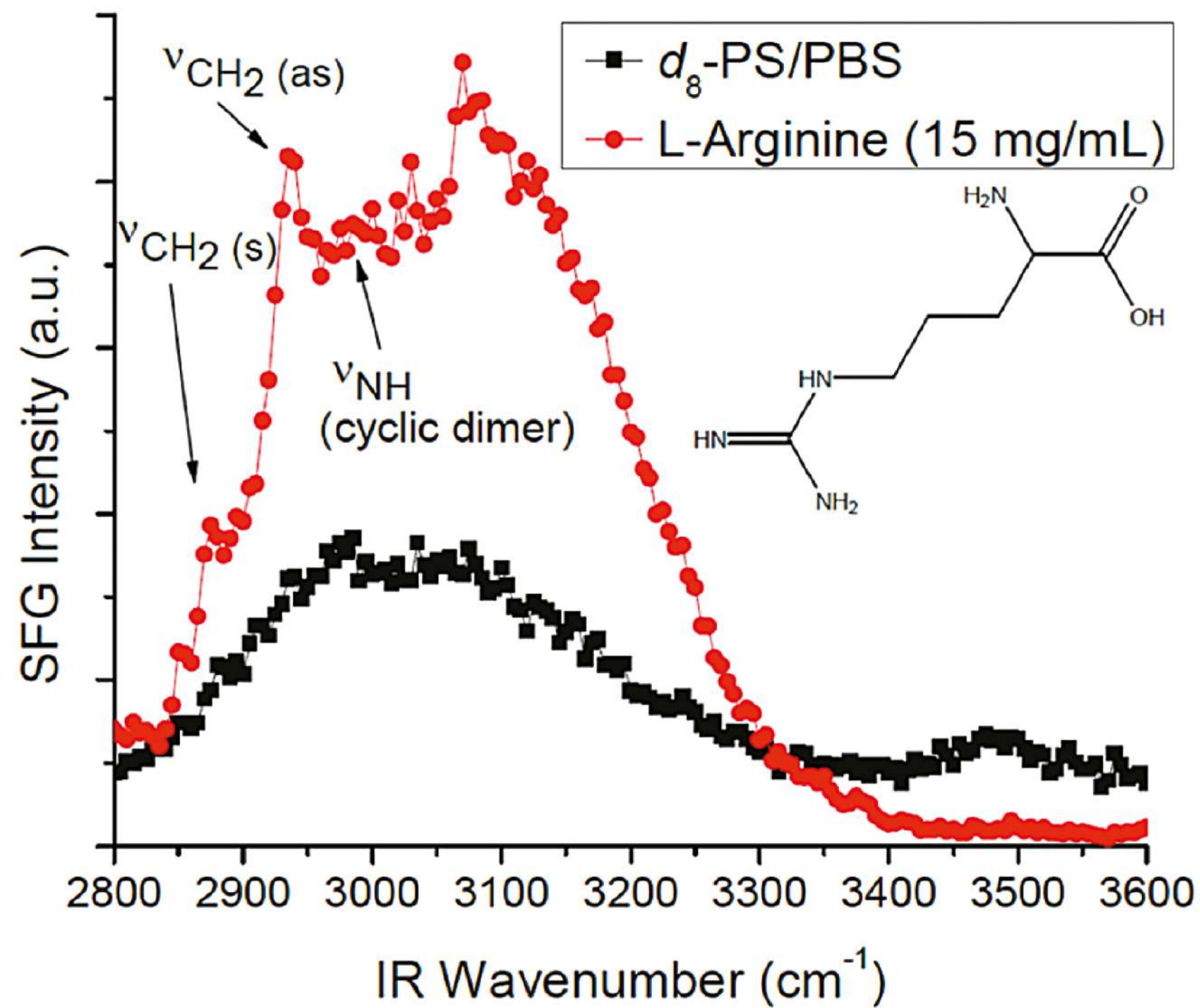


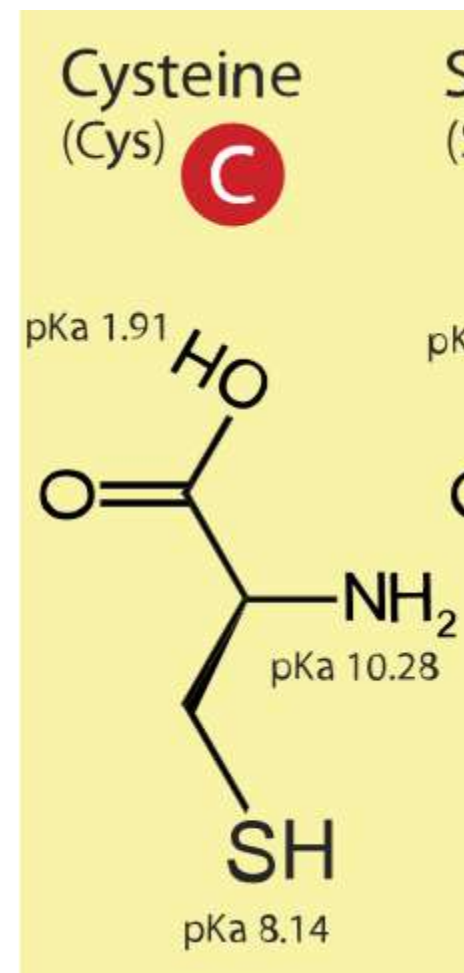
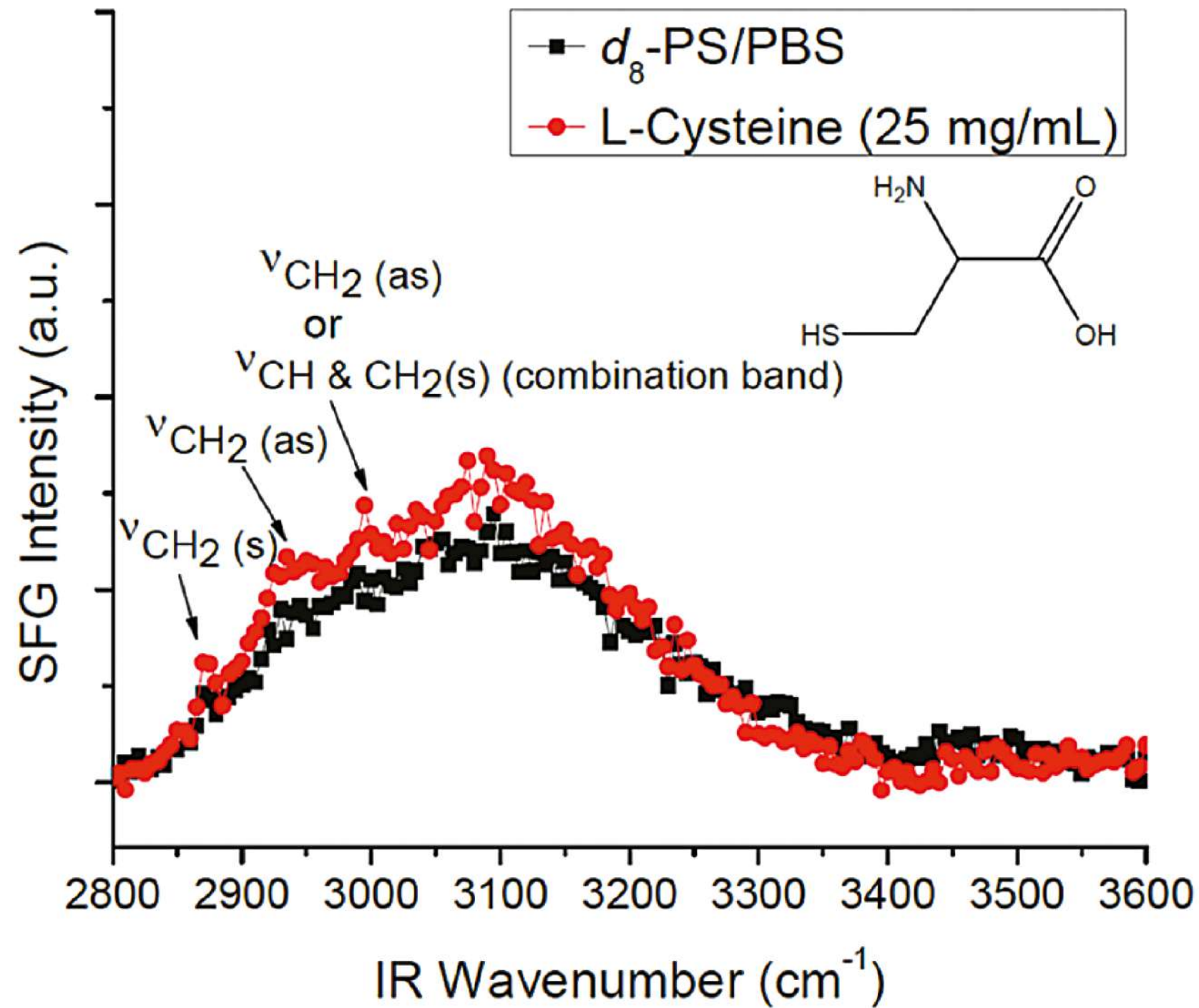


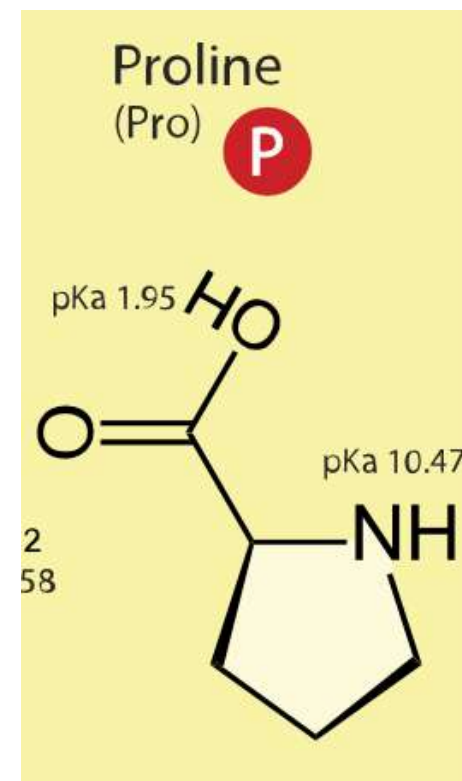
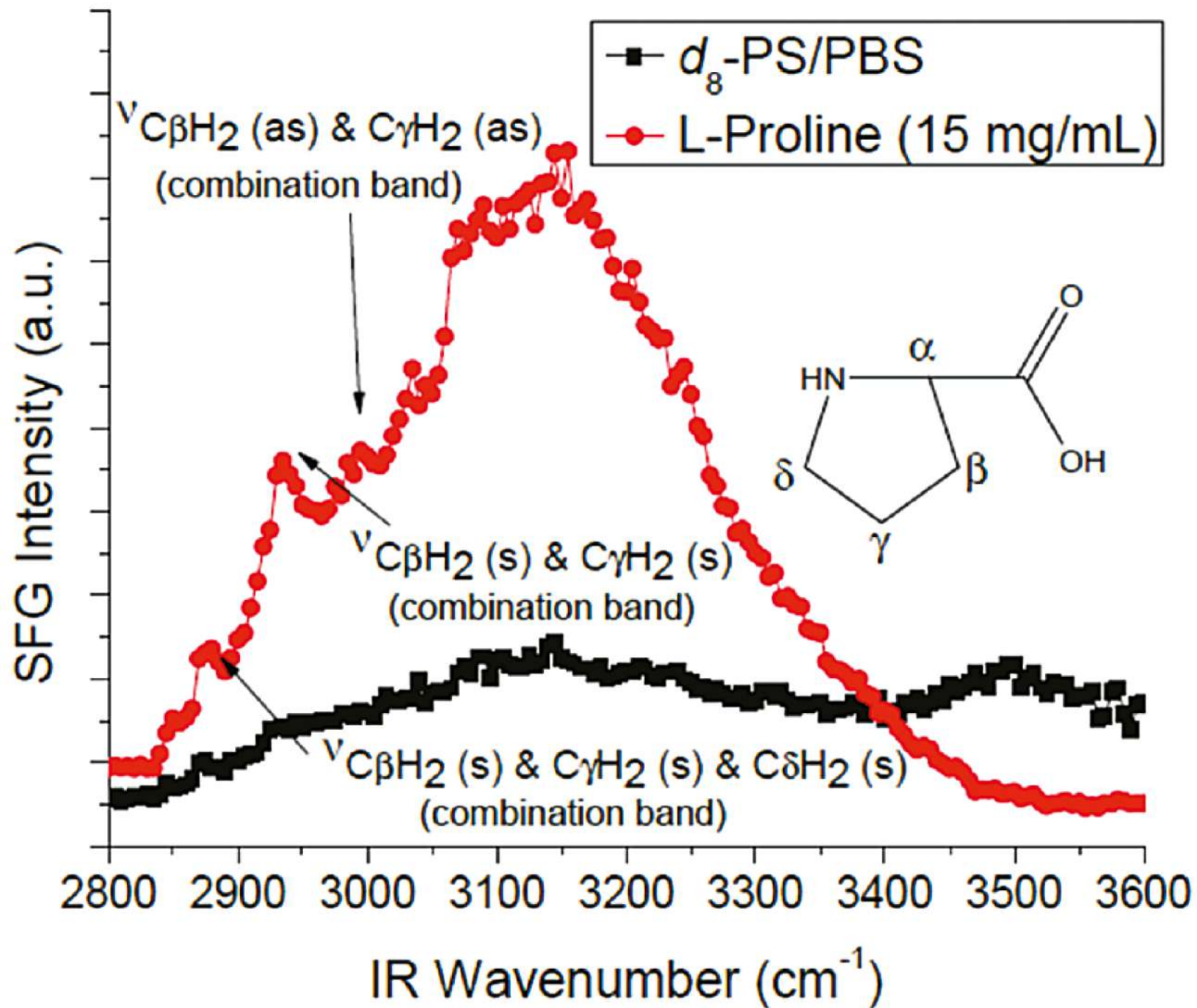
Similar OH due to higher concentration of amino acid.



Interestingly, there is an apparent blue-shift in the broad O-H bond peak, which is consistent with the presence of an N-H stretch contributing to the shoulder of the O-H peak.^{88,89}





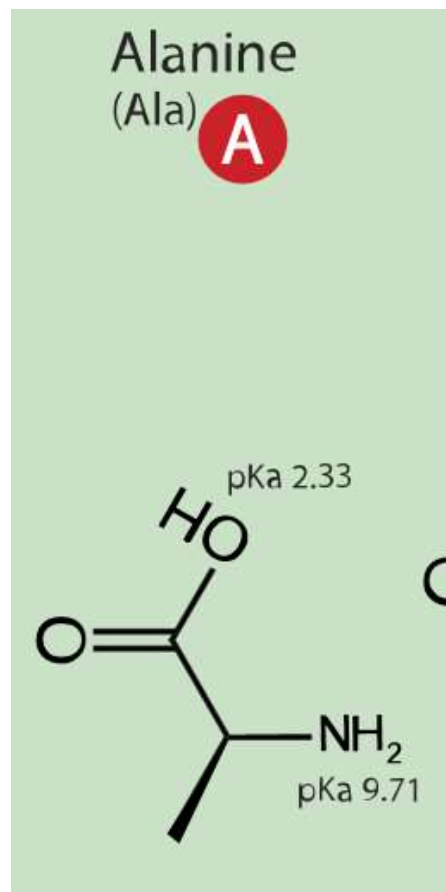
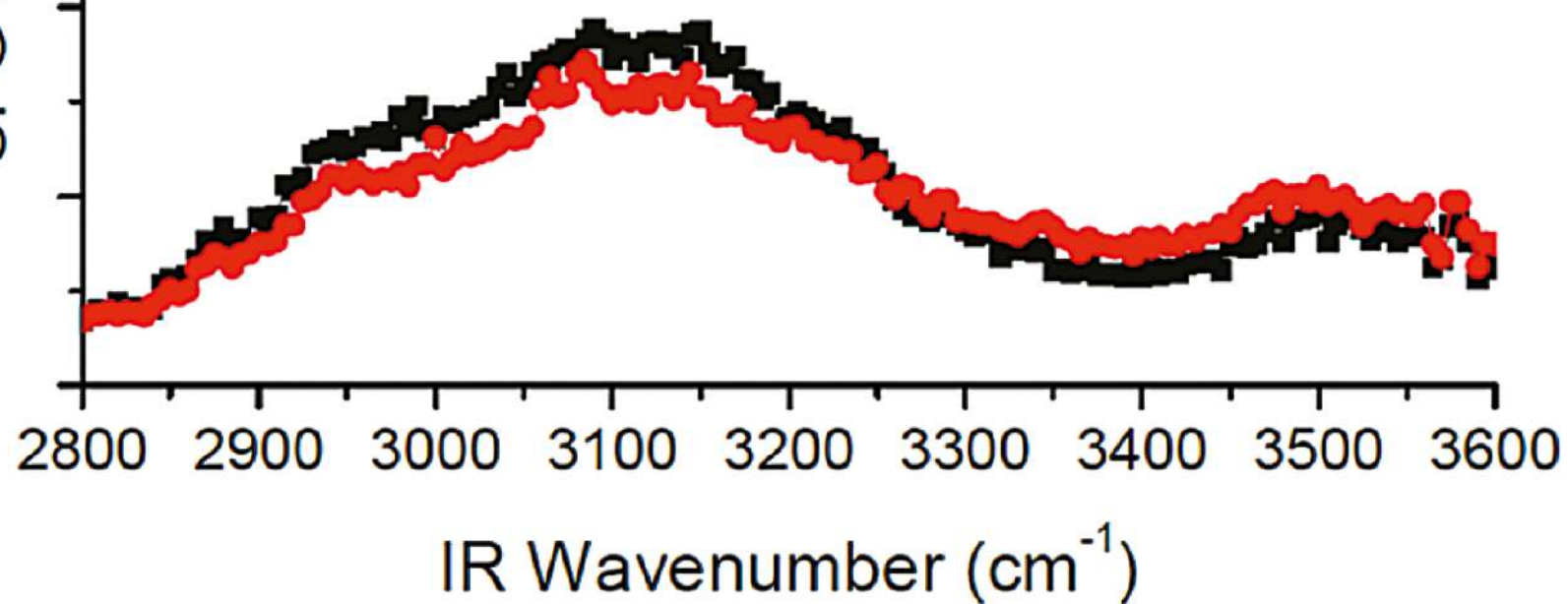
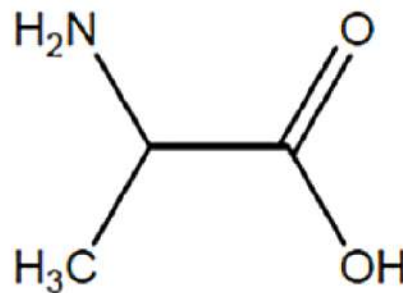


3000-3400/ cm region

OH and NH contribution

SFG Intensity (a.u.)

■ d_8 -PS/PBS
● L-Alanine (40 mg/mL)



- No CH band due to molecular arrangement, not to absence.
- QCM: 469 ng/cm²

SiO₂/PBS: Black spectrum

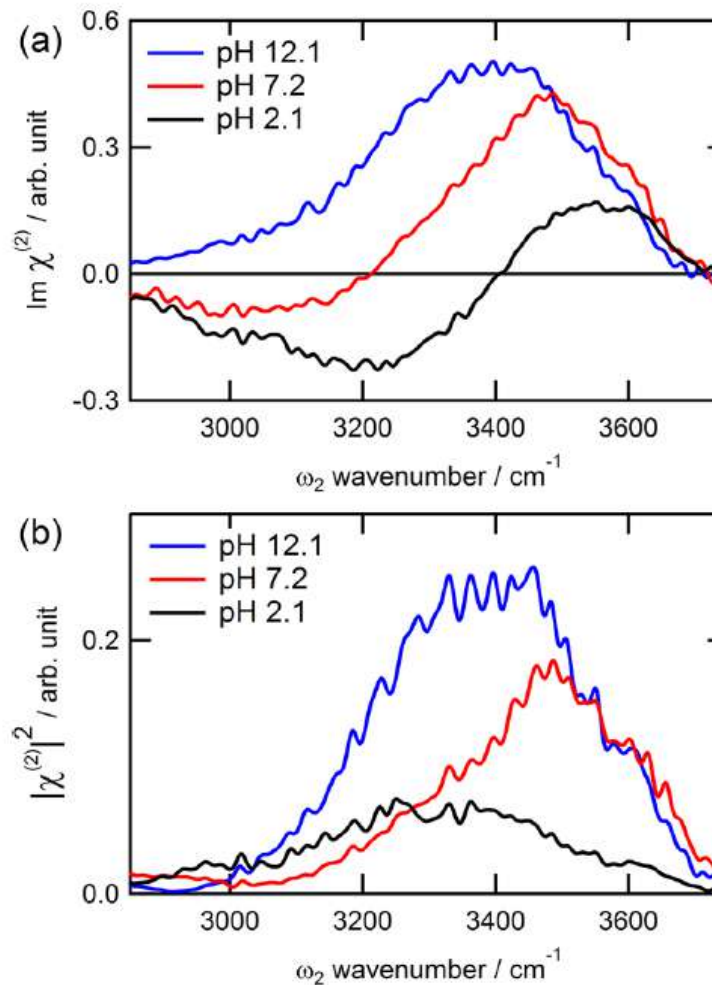
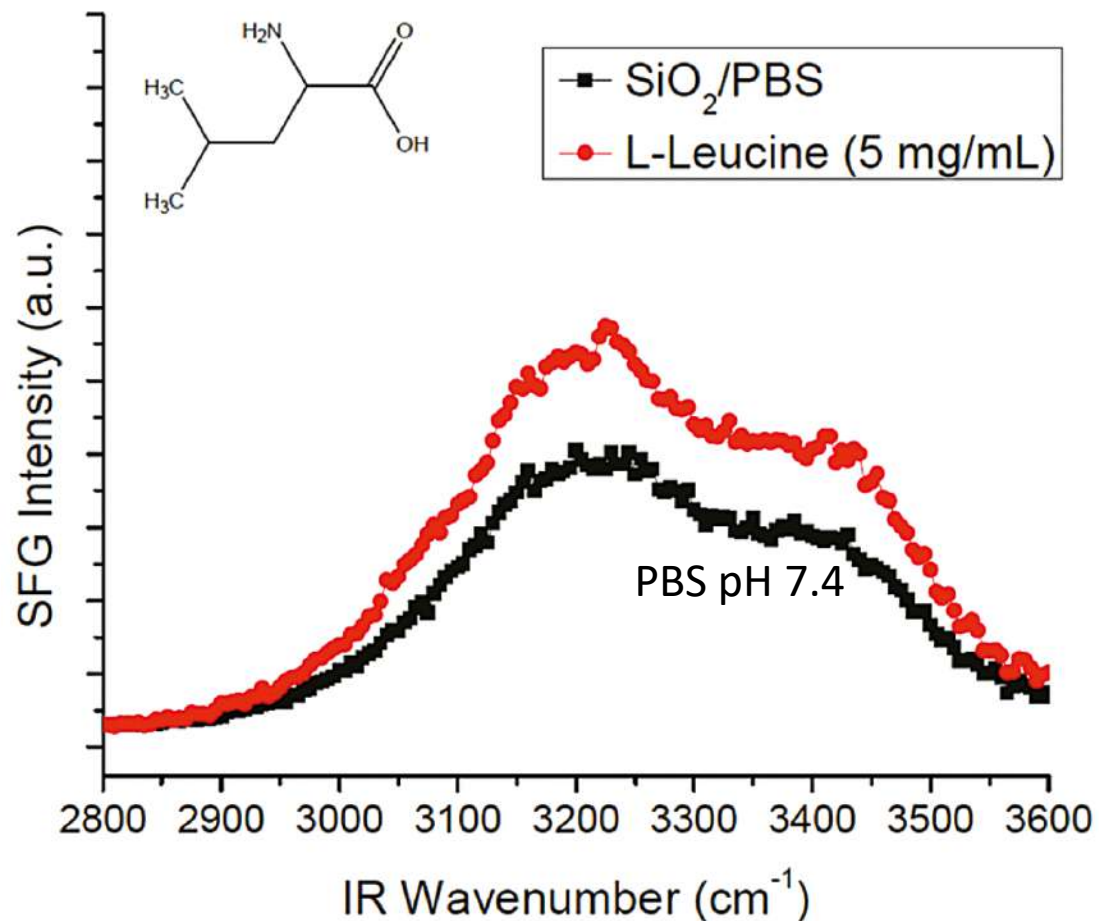
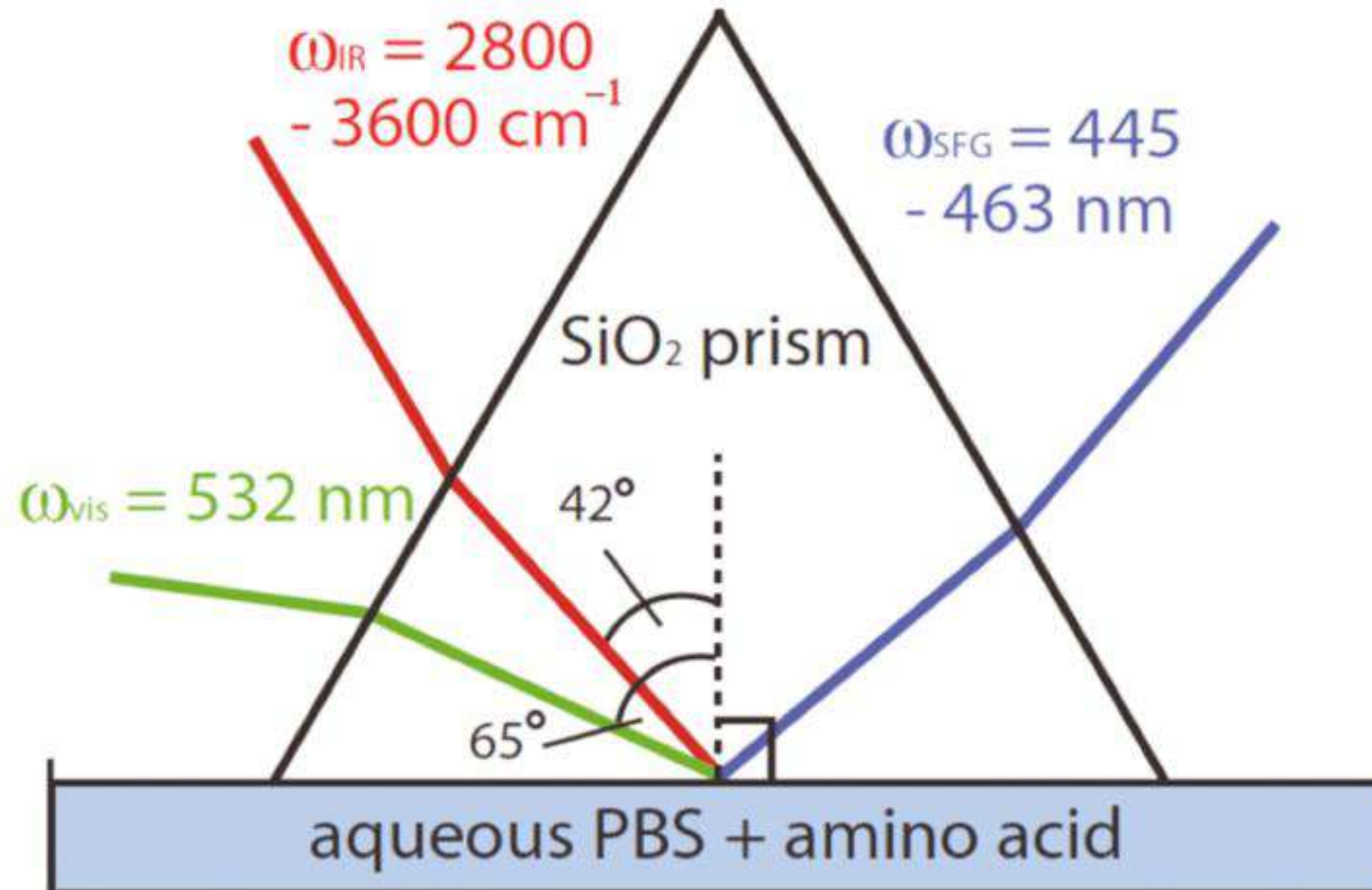


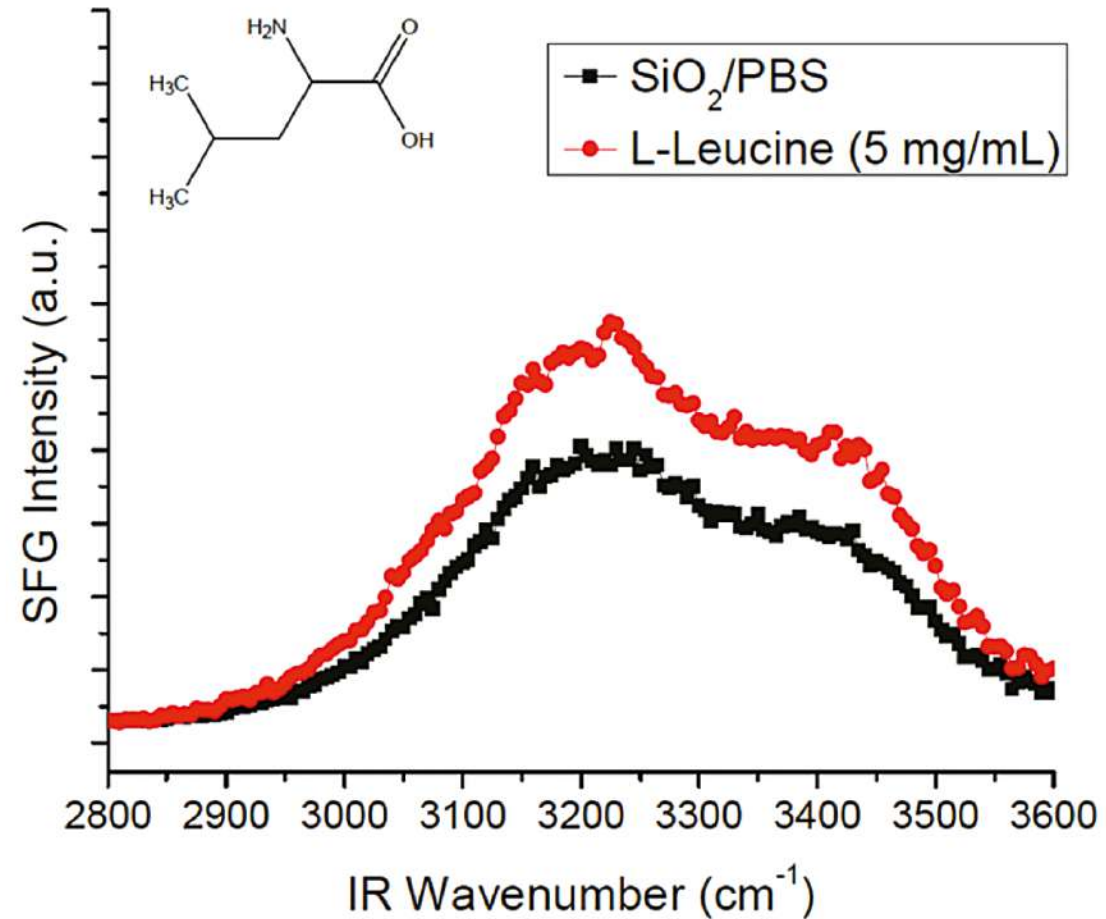
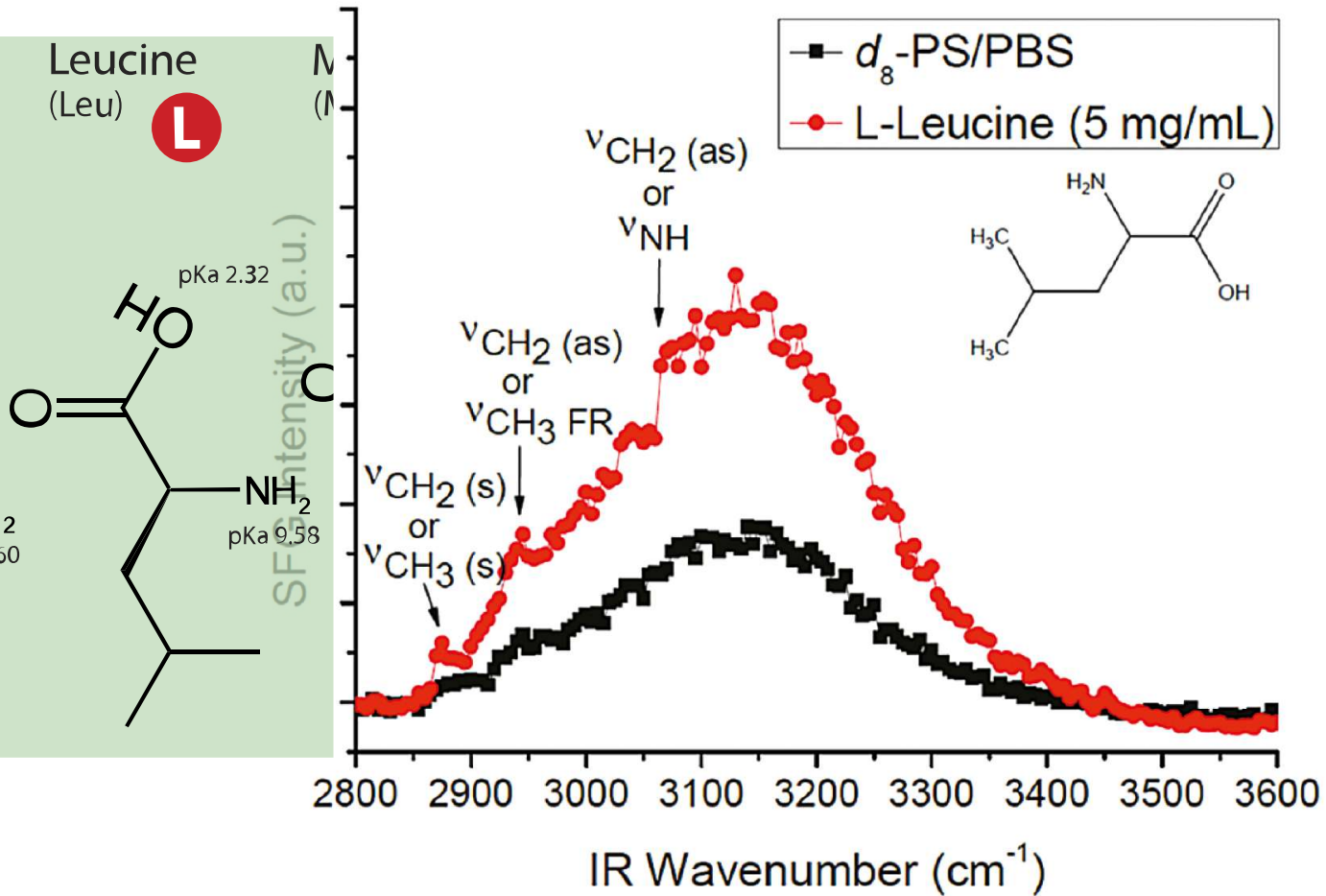
Figure 3. (a) Imaginary part of the $\chi^{(2)}$ spectra of the buried silica/HOD-D₂O interface for different pH in phosphate buffer ($I = 10$ mM, ratio of H₂O:HOD:D₂O = 1:8:16). (b) Corresponding magnitude square of the $\chi^{(2)}$ spectra ($|\chi^{(2)}|^2$ spectra) of the buried silica/HOD-D₂O interface for different pH in phosphate buffer. The pH of the solution for each spectrum is indicated in the Figure.

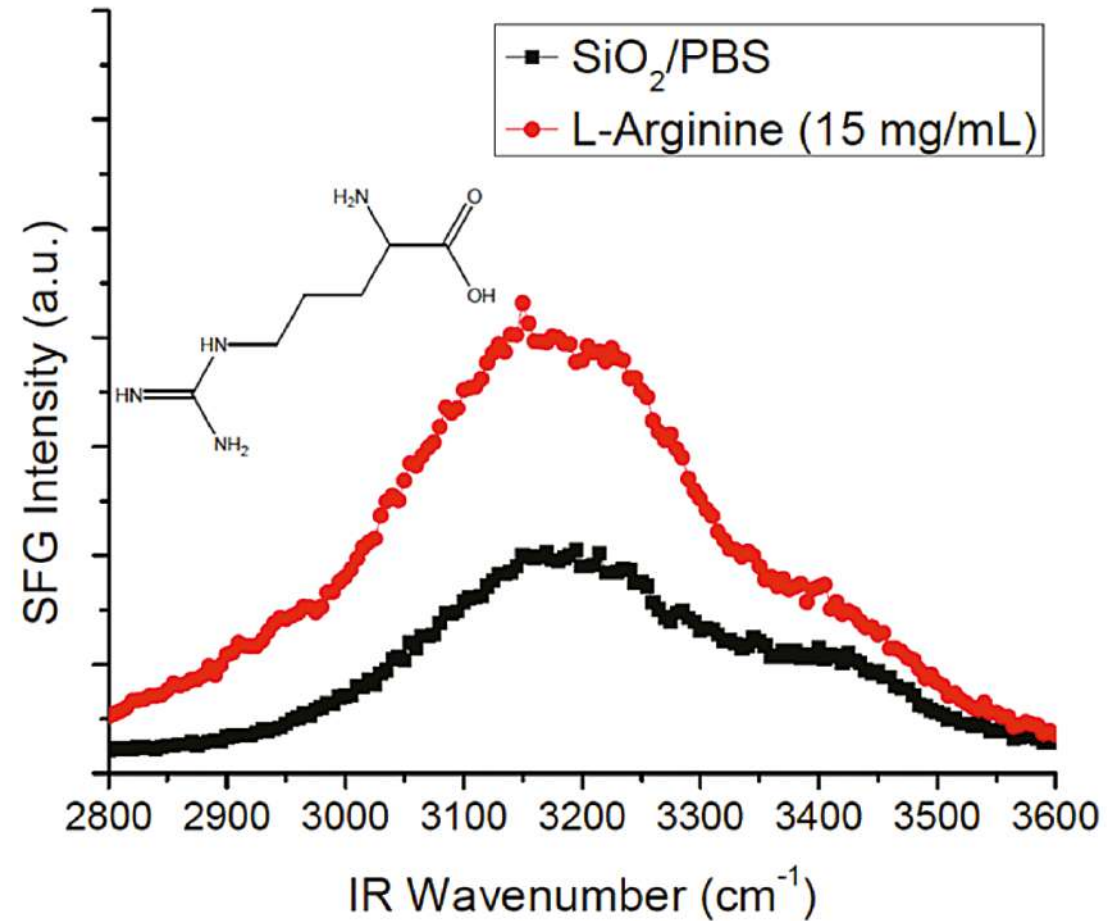
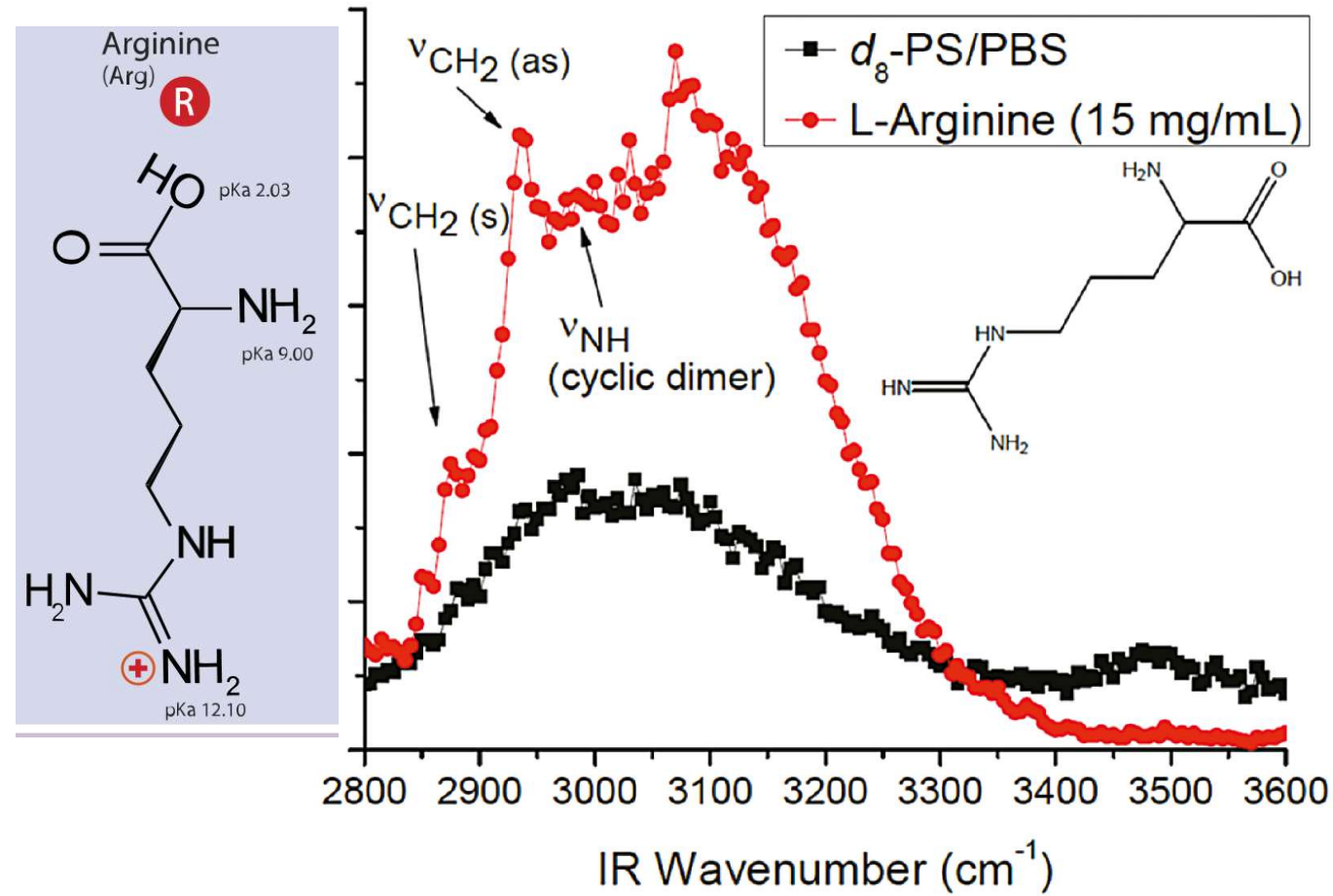
SFG results

No d-PS coated

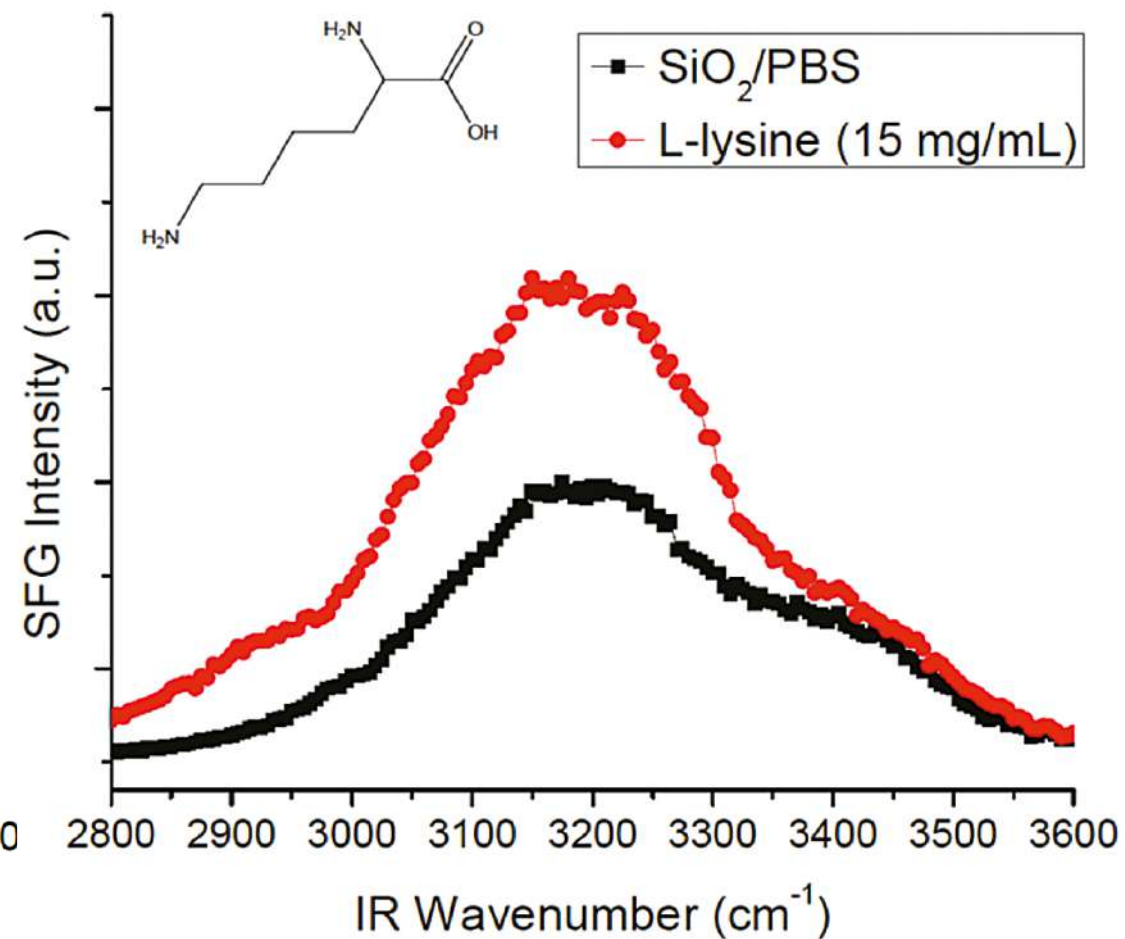
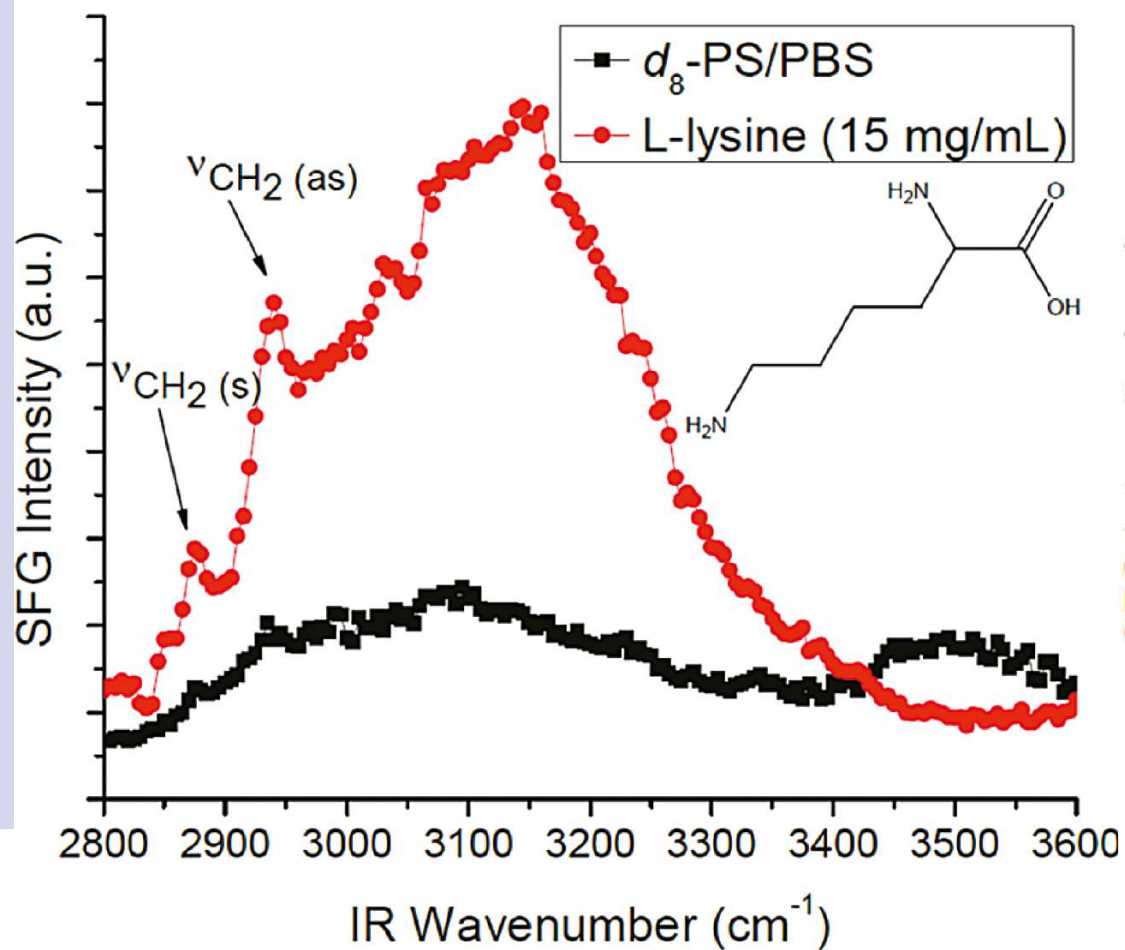
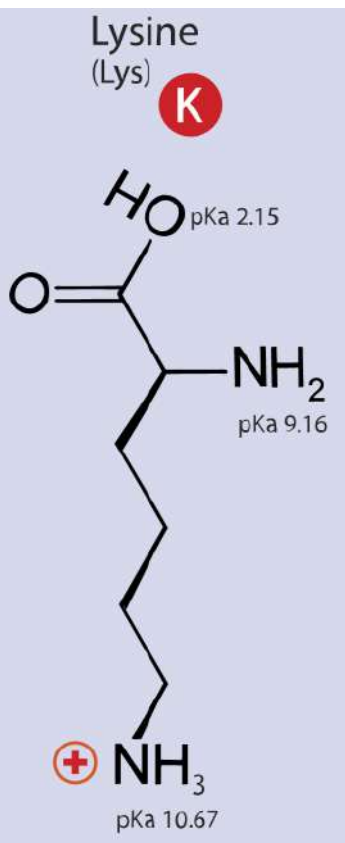


L-Leucine, L-Arginine, and L-Lysine

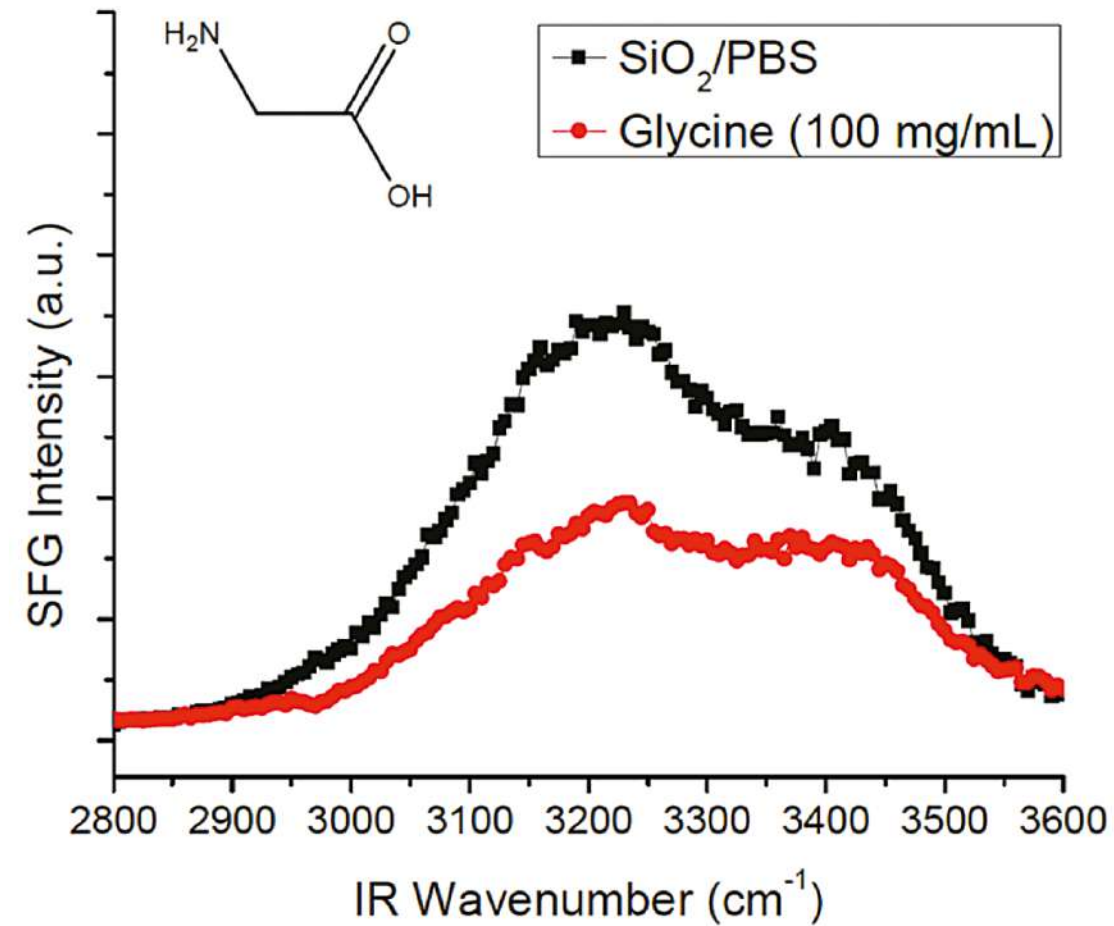
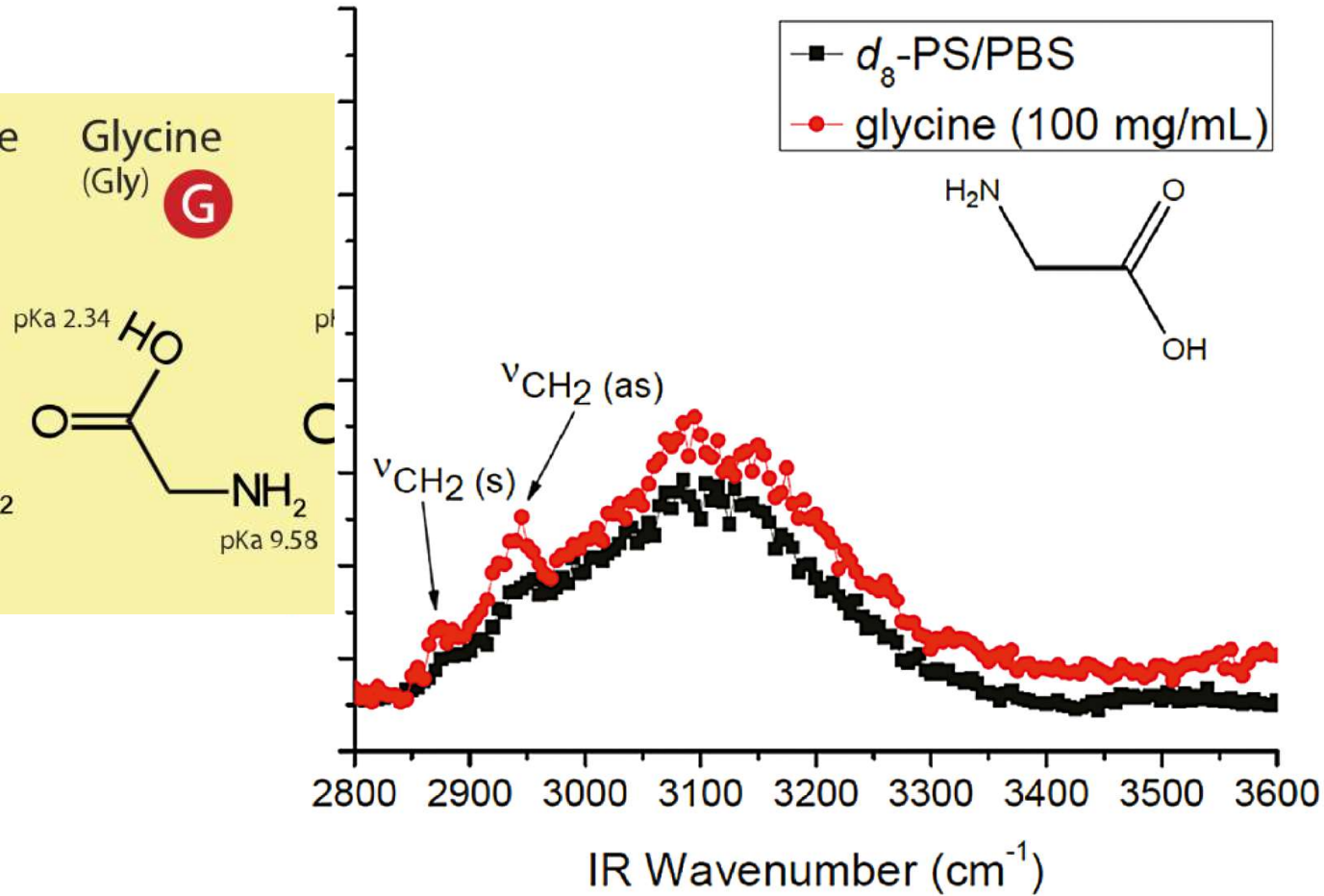




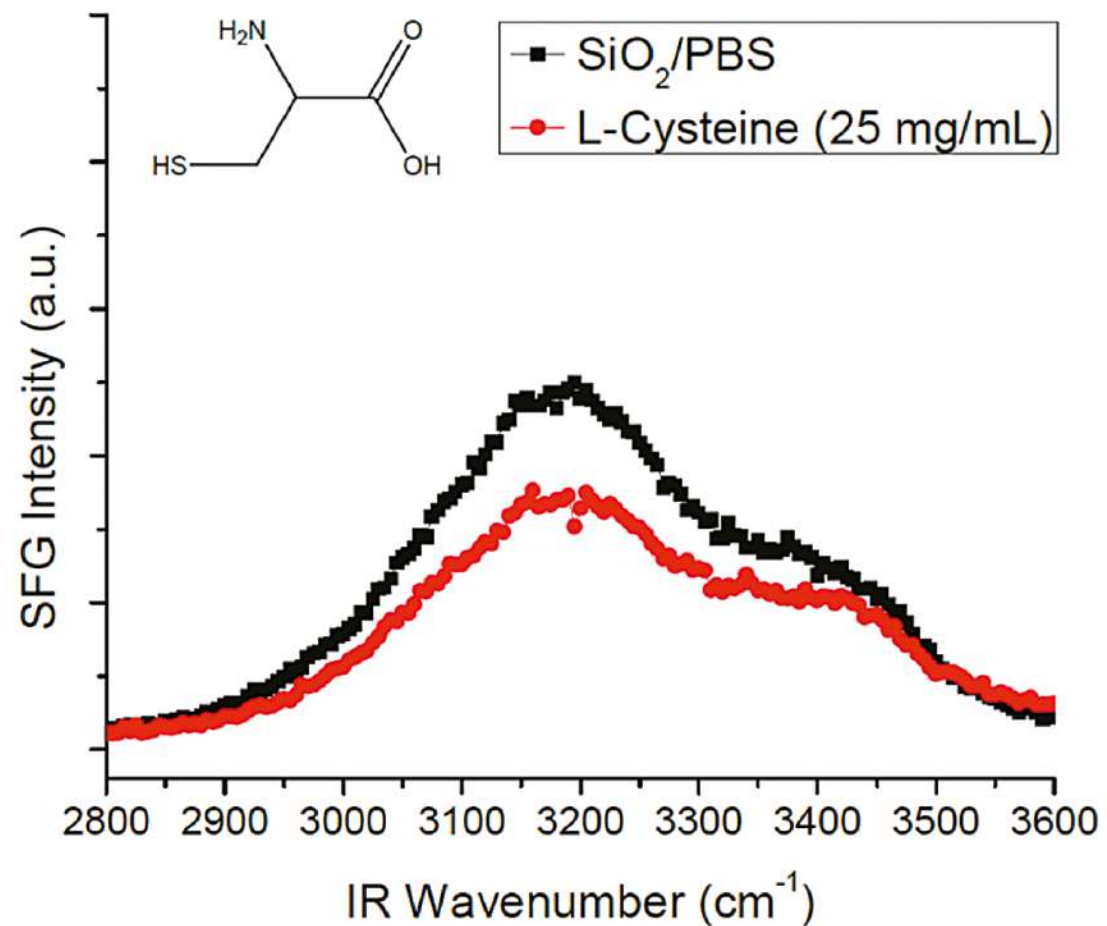
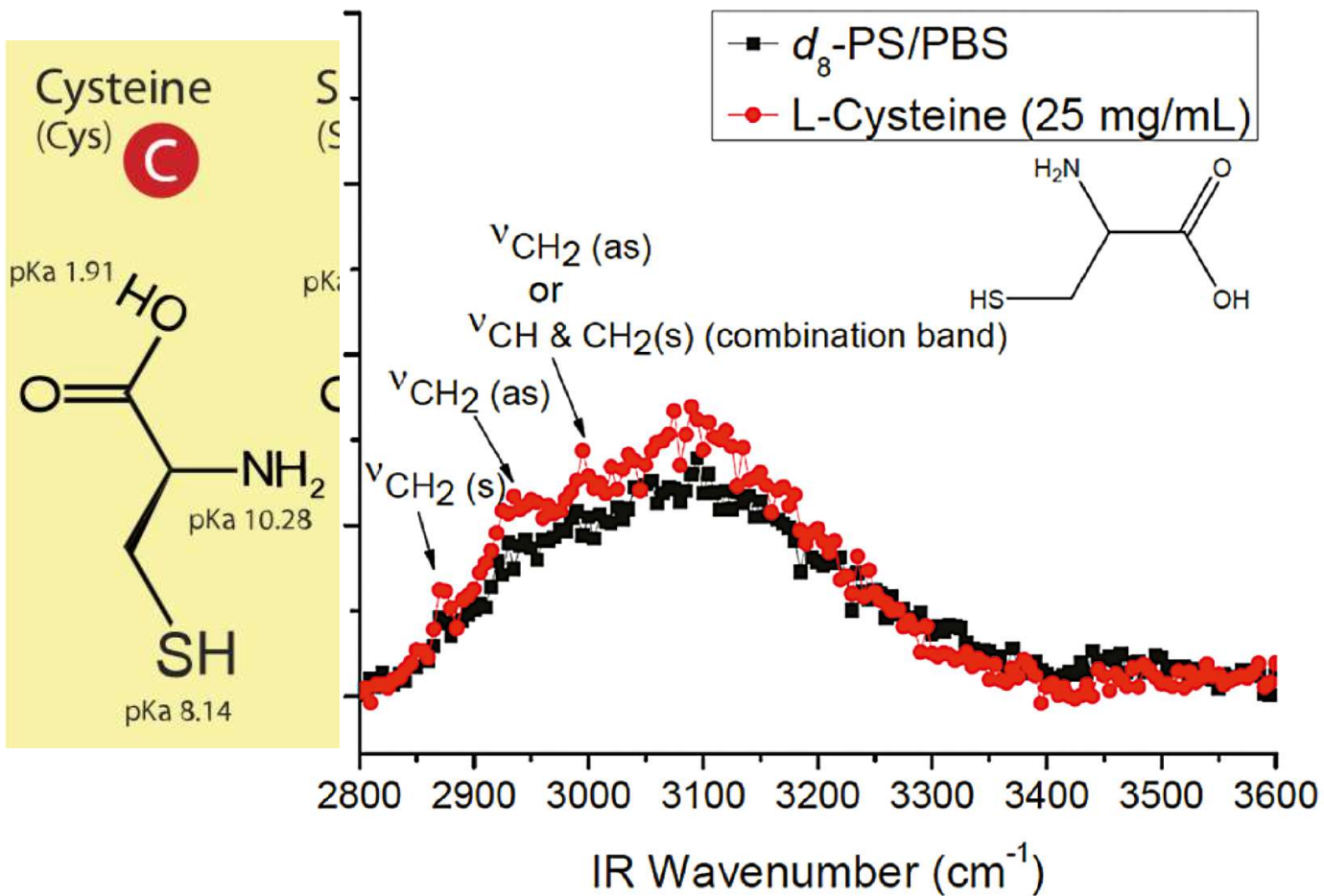
Band Broadening due to NH

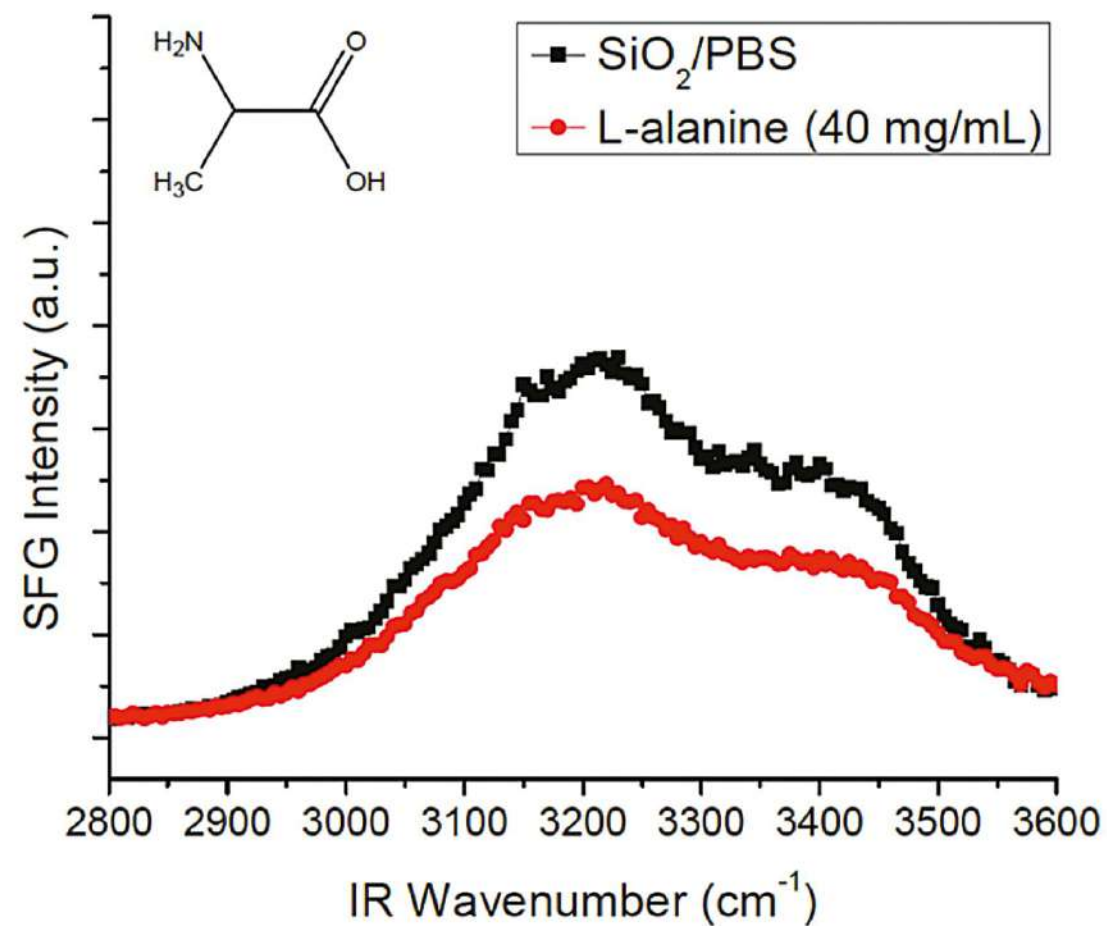
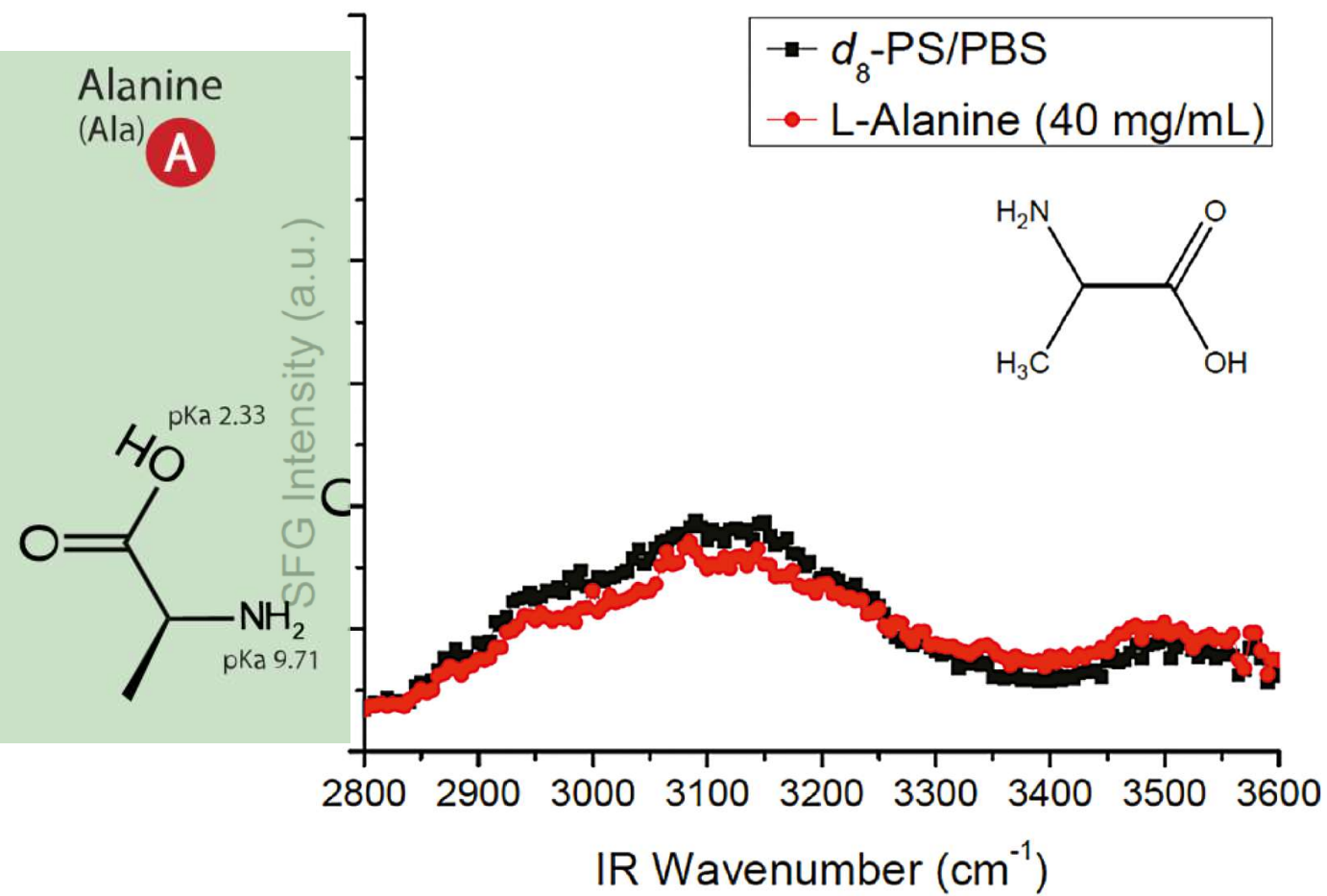


Glycine, L-Cysteine, and L-Alanine

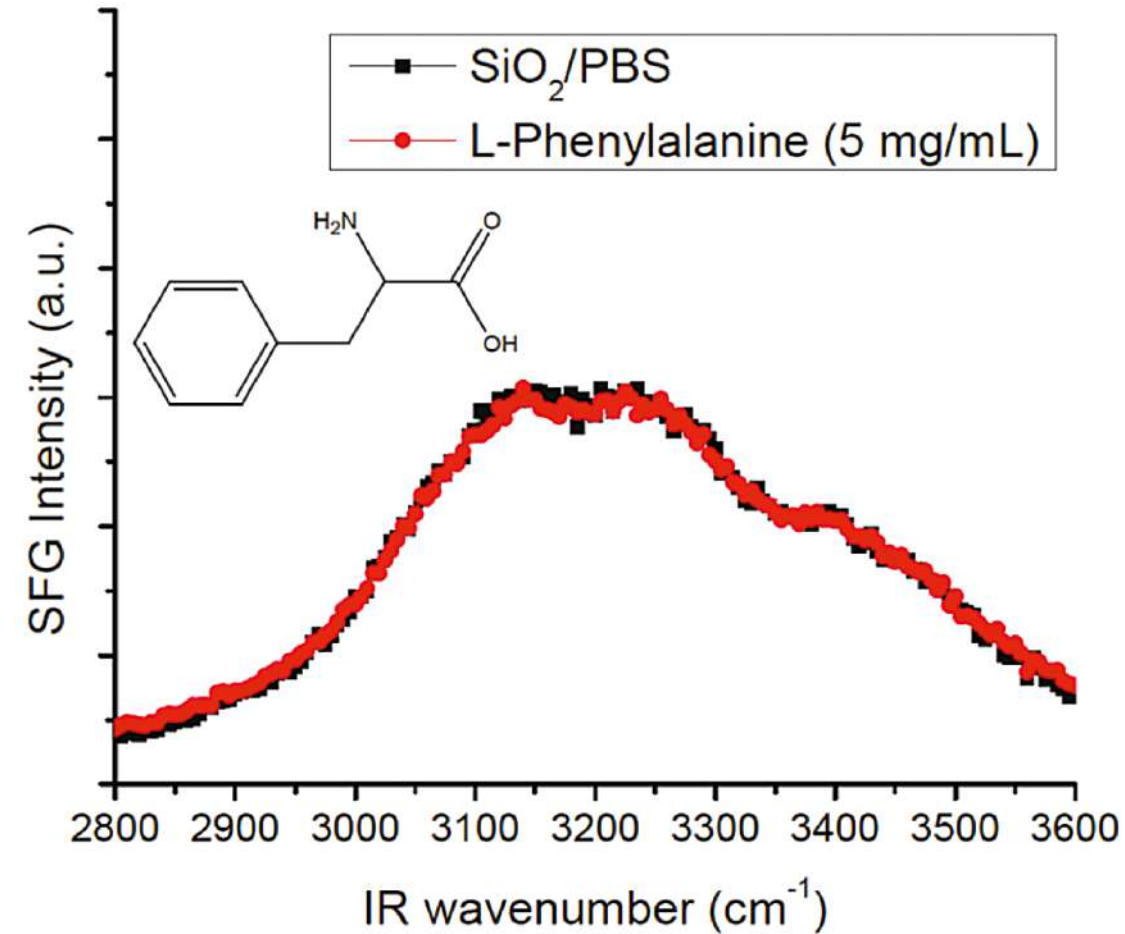
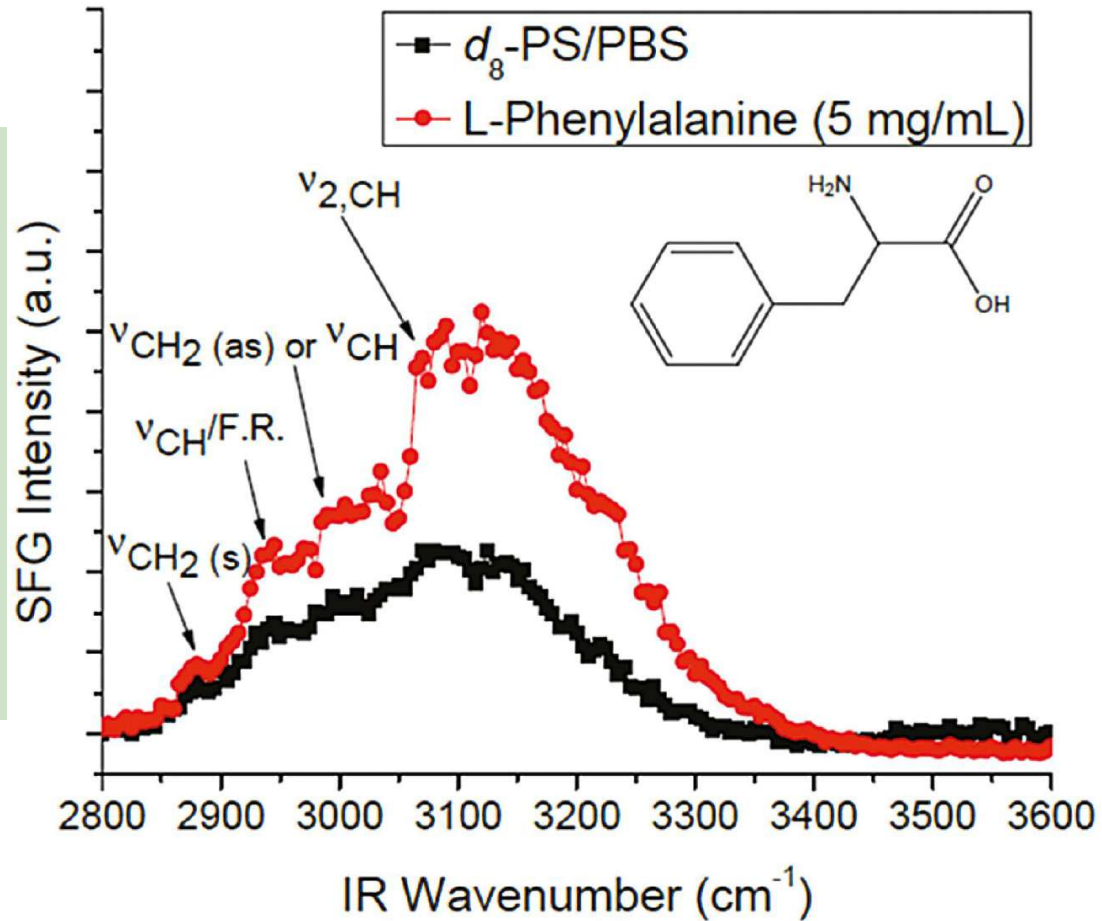
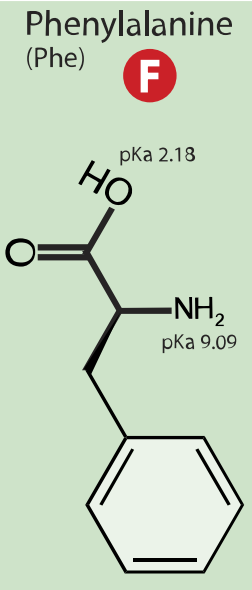


Weaker OH due to
higher solution concentration
displacement of interfacial water molecule.



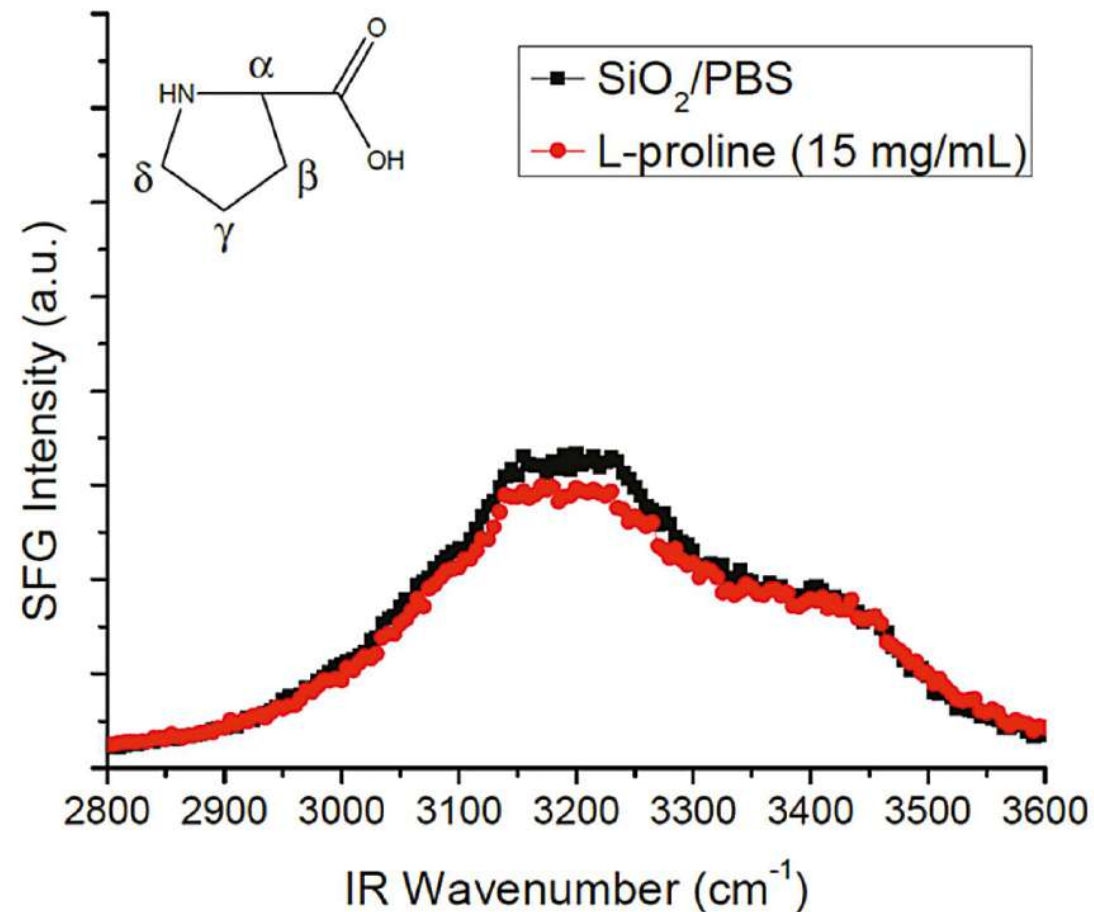
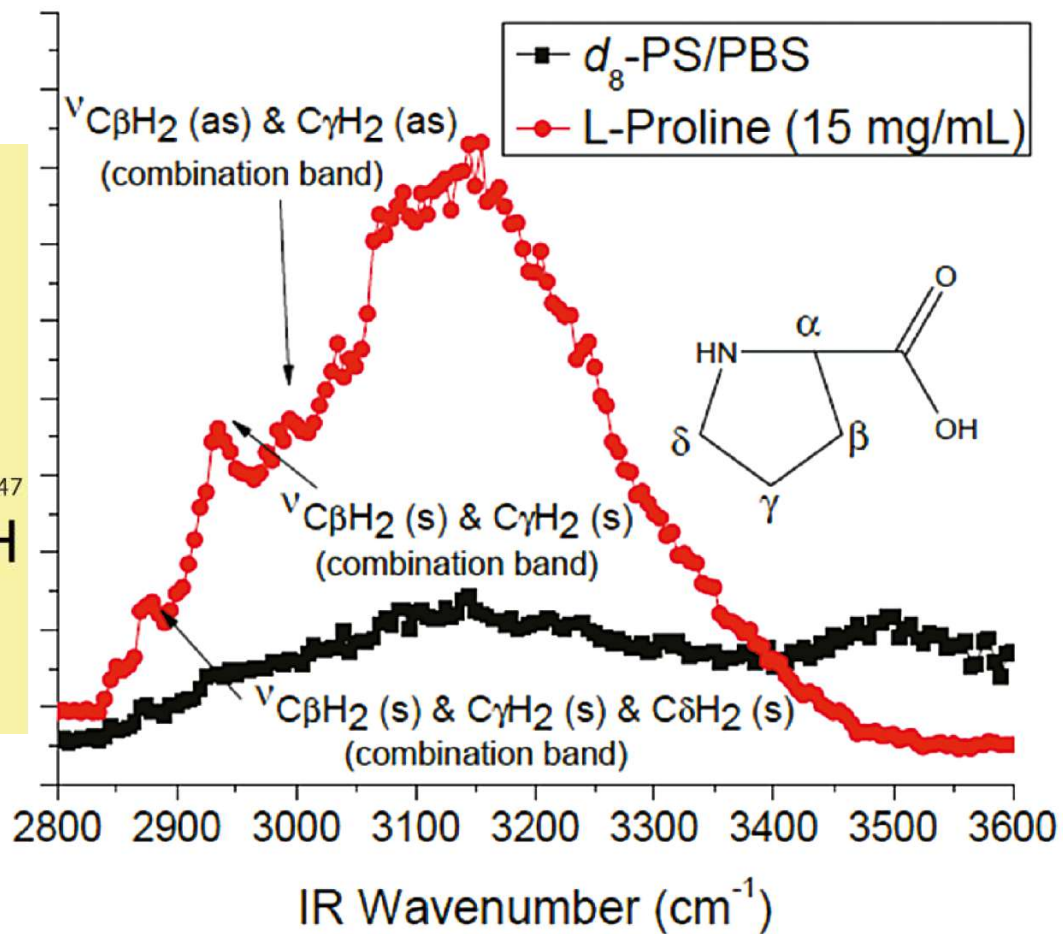
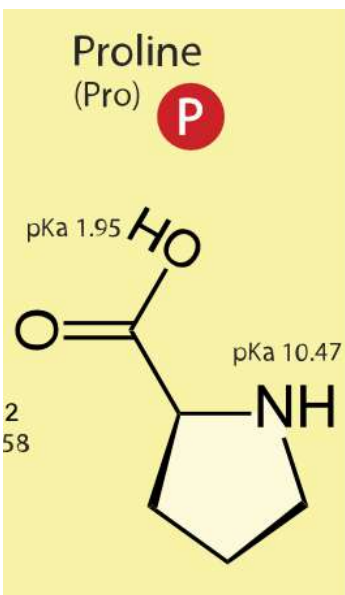


L-Phenylalanine and L-Proline.

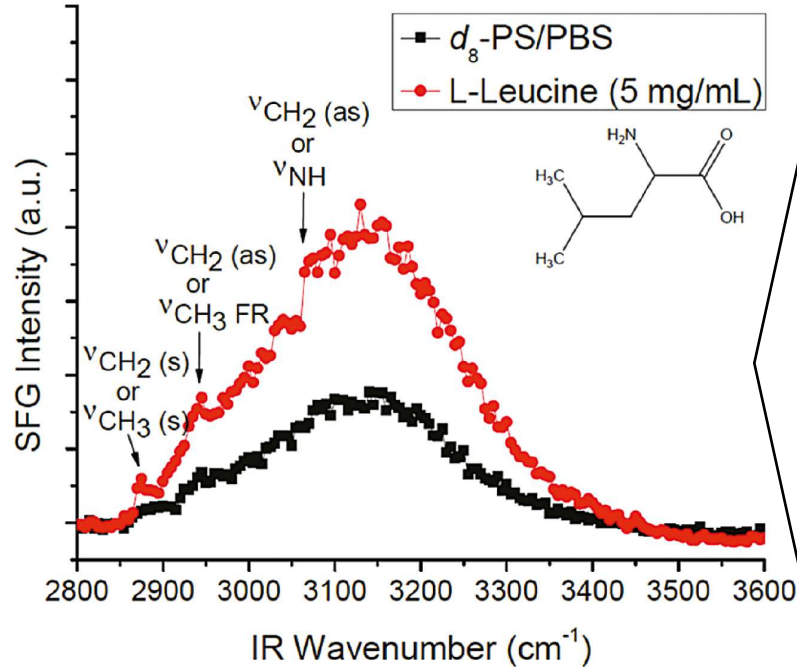


QCM: 63 ng/cm^2 on SiO_2 surface

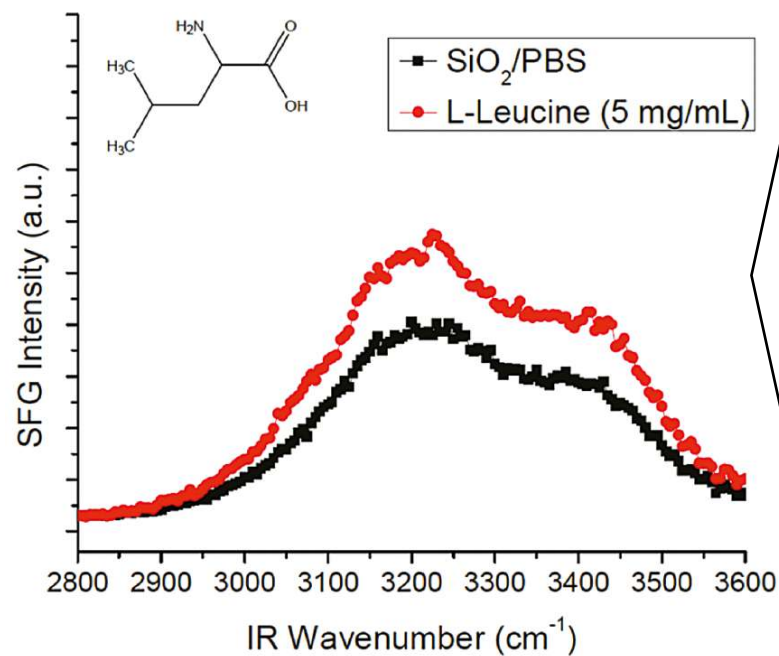
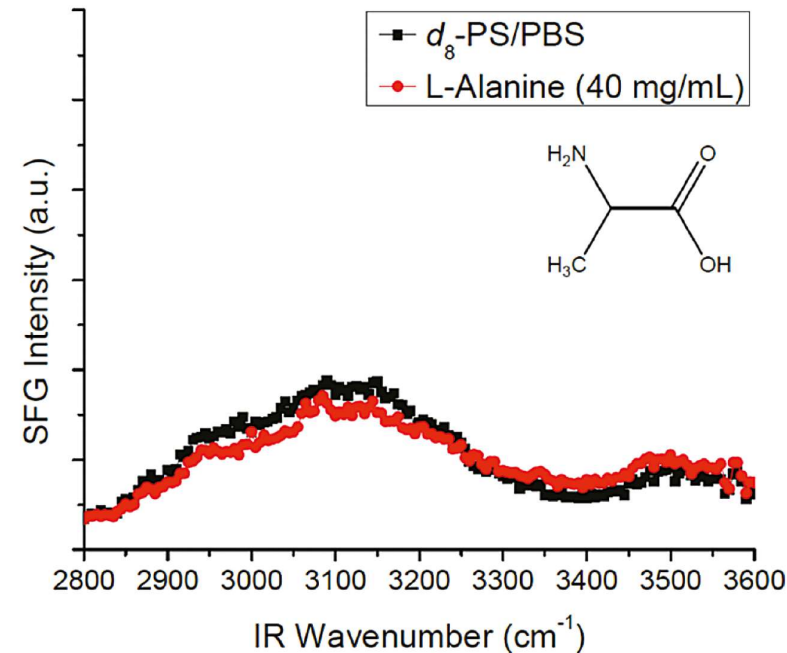
Same spectrum: due to surface organization, not to an absence



Conclusion



7 over 8 show CH except L-Alanine
(due to configuration of molecule at surface)



No CH=not well ordered as in d-PS case
OH: concentration, frequency of interaction

