

Supporting Information

IR spectrum of the protonated neurotransmitter 2-phenylethylamine: dispersion and anharmonicity of the $\text{NH}_3^+ - \pi$ interaction

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Table T1. Relative energies of stationary points on the potential of H^+PEA calculated at the B3LYP and B3LYP-D3 levels using the aug-cc-pVTZ basis set.

Figure S1. Structures of the most stable isomers of H^+PEA (**G**, **A**) compared to those of the most stable neutral PEA conformers (**B**, **C**) evaluated at the B3LYP-D3/aug-cc-pVTZ level. Bond distances are given in Å.

Figure S2. NBO charge distributions (in me) of the most stable isomers of H^+PEA (**G**, **A**) compared to those of the most stable neutral PEA conformers (**B**, **C**) evaluated at the B3LYP-D3/aug-cc-pVTZ level.

Figure S3. (top) Lengths and scaled harmonic stretch frequencies of the three N-H bonds of H^+PEA (and corresponding average values) as a function of the dihedral angle $\theta = \theta(\text{C1C7C8N})$. The plots are symmetric with respect to $\theta = 0^\circ$. (bottom) Scaled harmonic linear IR spectra of H^+PEA as a function of selected dihedral angles $\theta = \theta(\text{C1C7C8N})$.

Figure S4. Expanded view of the linear harmonic and anharmonic IR absorption stick spectra of the **G** conformer of H^+PEA (unscaled) calculated at the B3LYP-D3/cc-pVTZ level.

Figure S5. IRPD spectrum of $\text{H}^+\text{PEA-Ne}_2$ in the C-H/N-H stretch range compared to linear harmonic IR absorption stick spectrum of the **G** conformer of H^+PEA at the B3LYP-D3/aug-cc-pVTZ level. Isomer-selective IR (M. Schütz et al., unpublished) and Raman spectra (Golan et al., J. Chem. Phys. 131, 024305, 2009) of the **C** conformer of PEA compared to linear harmonic IR absorption stick spectrum at the B3LYP-D3/aug-cc-pVTZ level.

Table T1. Relative energies of stationary points on the potential of H⁺PEA calculated at the B3LYP-D3 and B3LYP (in parentheses) levels using the aug-cc-pVTZ basis set.

	E ₀ in cm ⁻¹	E ₀ in kJ/mol
G	0 (0)	0 (0)
A	1564 (1202)	18.7 (14.4)
TS1	1090 (1169)	13.0 (14.0)
TS2	2786 (2392)	33.3 (28.6)

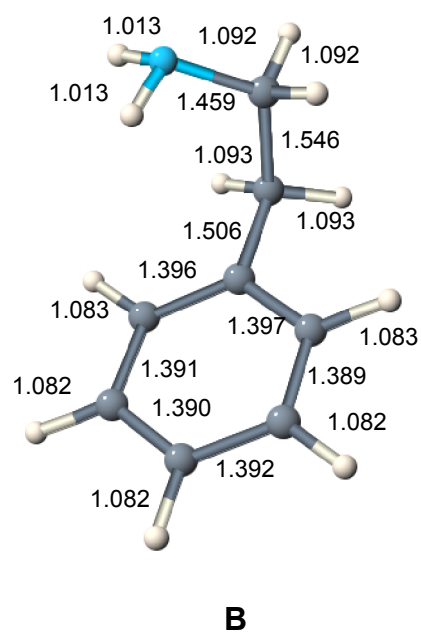
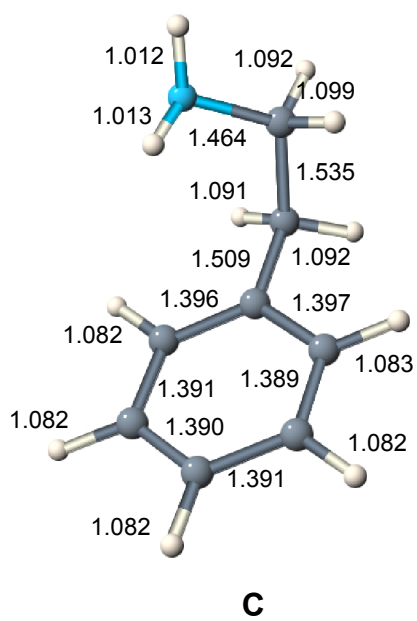
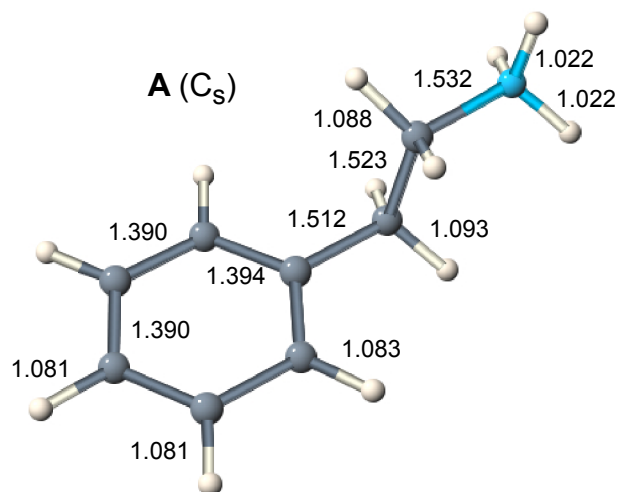
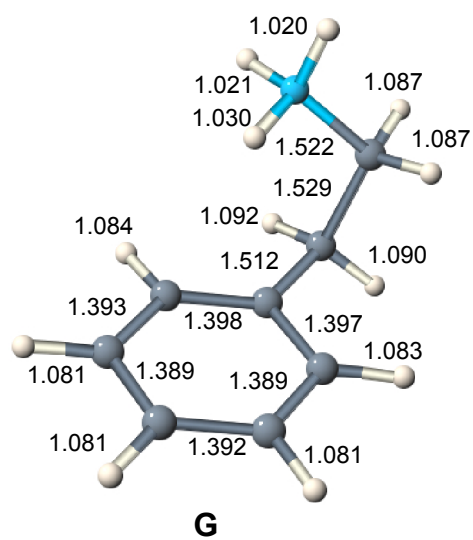


Figure S1

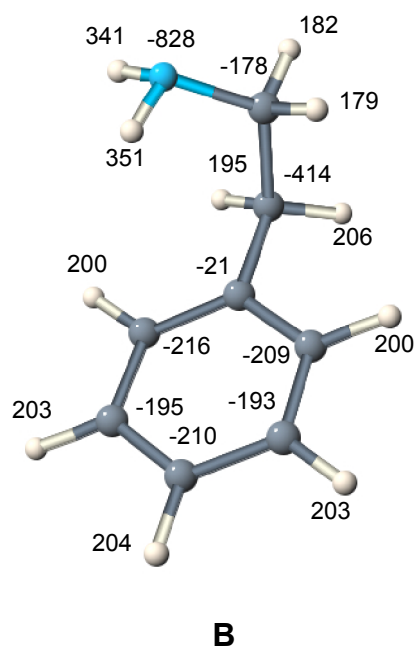
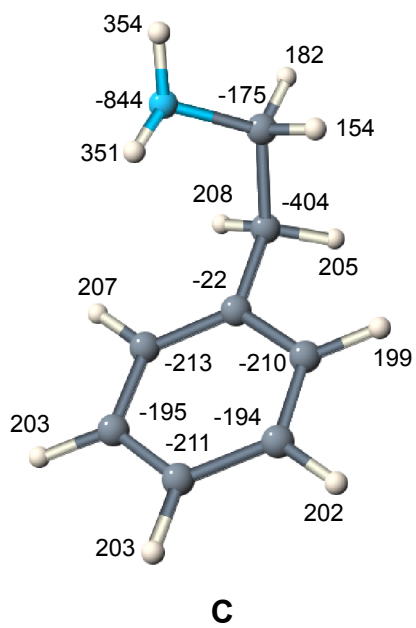
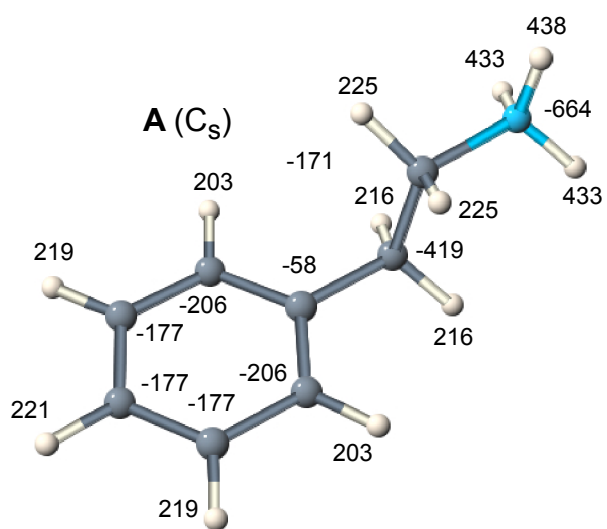
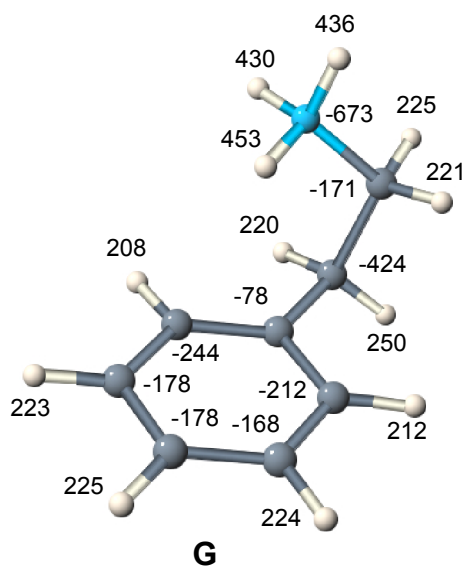


Figure S2

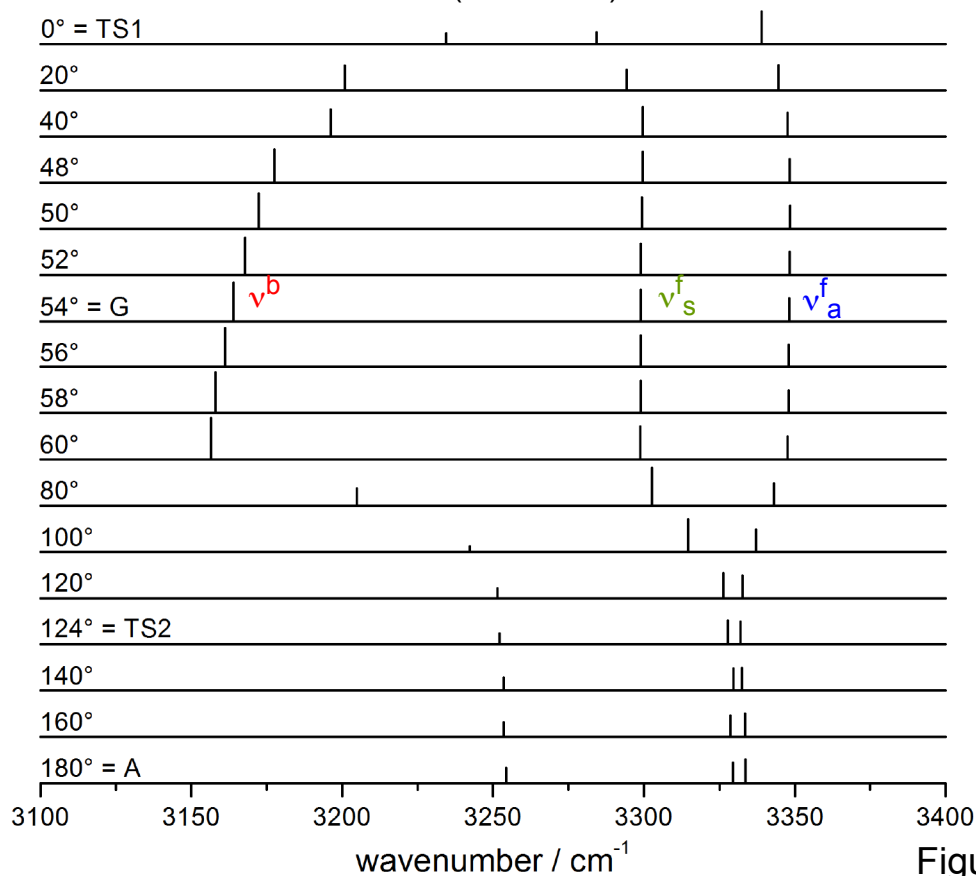
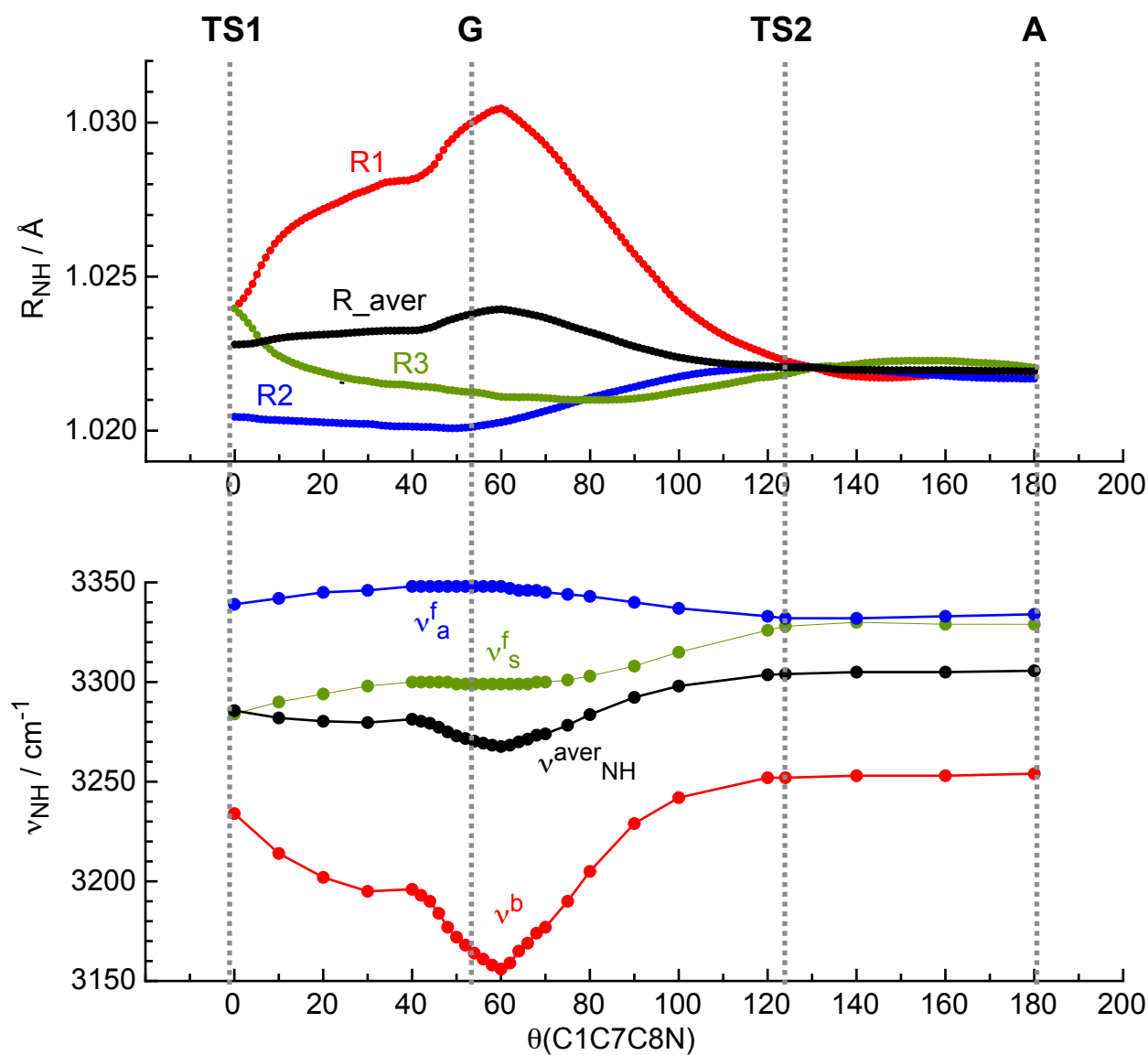


Figure S3

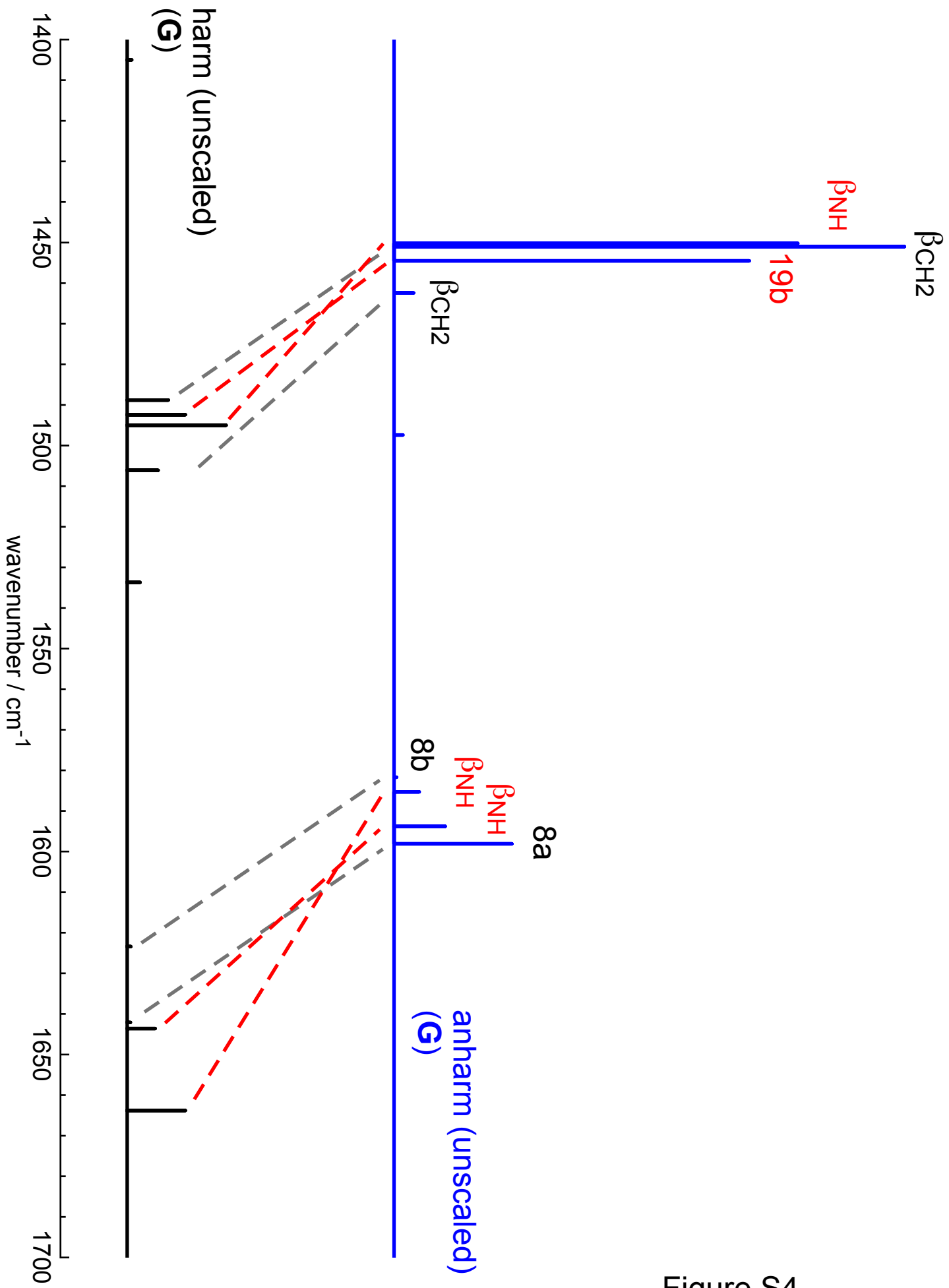


Figure S4

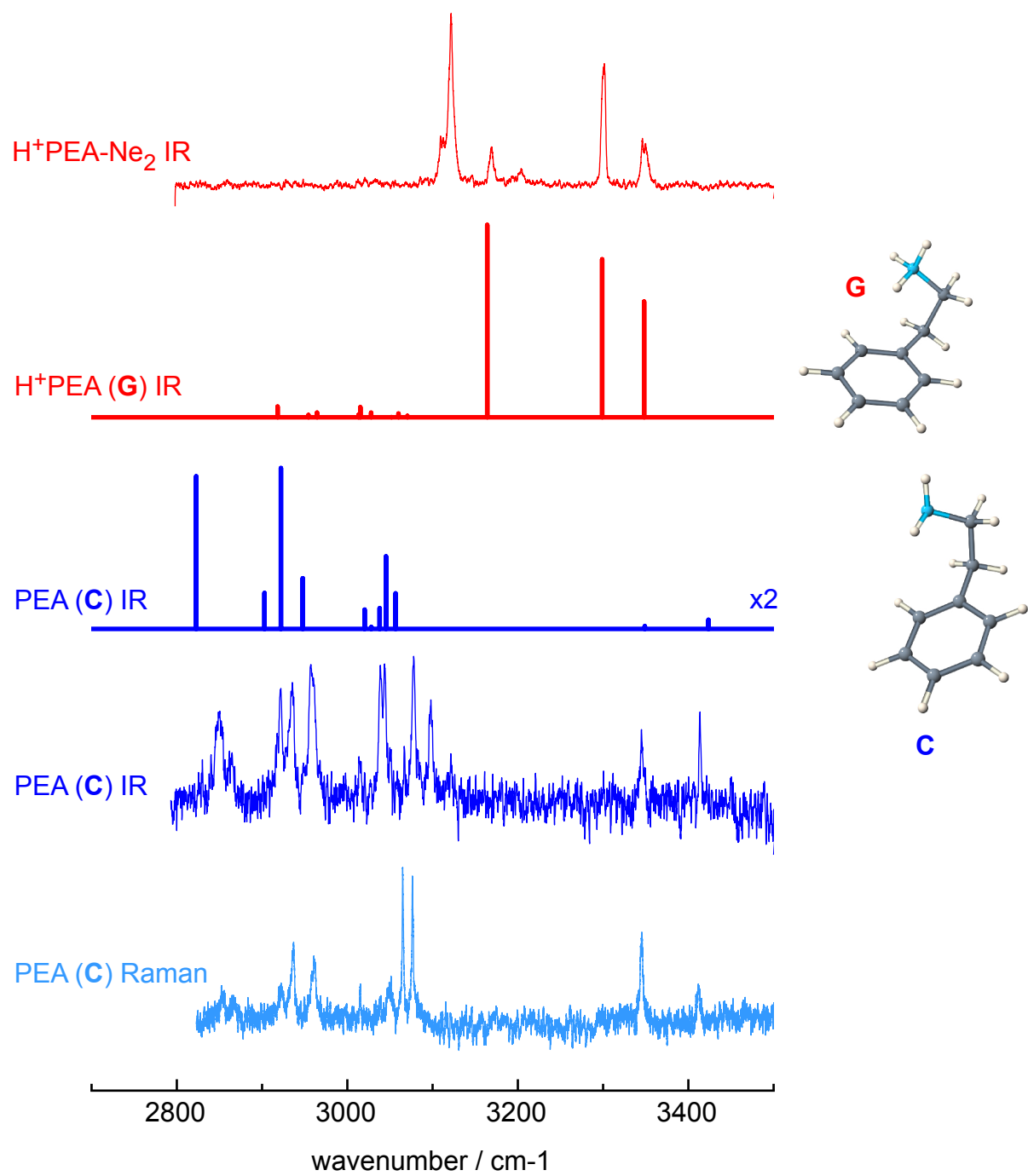


Figure S5