

## IRMPD Spectroscopy Shows That AGG Forms an Oxazolone $b_2^+$ Ion

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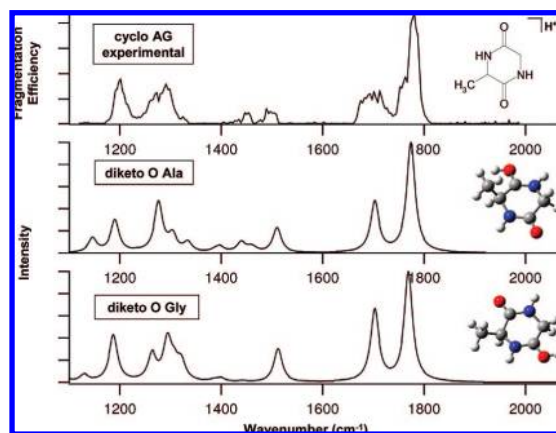
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The determination of structural features at the molecular level can provide fundamental insights into the activity, behavior, and function of biological systems. While techniques such as NMR and X-ray crystallography have stood as benchmarks for determining biological structure, mass spectrometry is playing an increasing role in investigating structural features.<sup>1,2</sup> With the advent of soft ionization techniques such as MALDI and ESI, biological systems from small molecules to peptides to macromolecular protein complexes can be examined using a range of mass spectrometric techniques.<sup>3</sup> Because analysis occurs in a solvent-free environment, mass spectrometry also introduces the opportunity to examine intrinsic features and interactions that may not be present in an aqueous solvent system.<sup>4</sup>

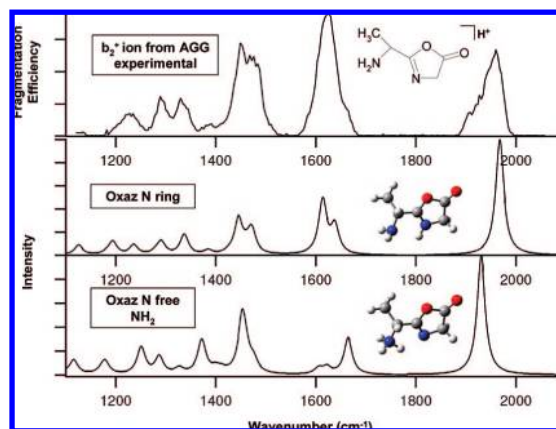
Structures of amino acids, peptides, and peptide fragments have also been a focus of mass spectrometry research.<sup>5</sup> In particular, the formation of  $b_2^+$  ions and their structure has prompted numerous studies. Two main structures have been proposed. If cyclization occurs by attack of the N-terminal amino group on the carbonyl carbon of the second amino acid residue, then a diketopiperazine structure is formed.<sup>6</sup> If the carbonyl oxygen of the first amino acid of a peptide attacks the carbonyl carbon of the second amino acid, an oxazolone structure is formed.<sup>5</sup> Peptide or fragment ion structures can be probed using MS/MS, isotope labeling, gas-phase hydrogen/deuterium exchange, ion mobility, and computational modeling. While many recent studies have shown that the  $b_2^+$  ion of simple peptides is an oxazolone, the studies were performed using indirect methods.<sup>7,8</sup> Studies have also shown that some  $b_2^+$  ions do not form oxazolone or diketopiperazine structures; for example, the involvement of side chains may lead to other structures.<sup>7</sup>

Infrared multiple photon dissociation (IRMPD) combined with theoretical vibrational spectra can provide spectroscopic evidence for the structure of ions. Recently, IRMPD spectroscopy has been used to investigate the structure and site of protonation in amino acids and peptides,<sup>9</sup> to probe alkali-metal complexes of amino acids and peptides,<sup>10</sup> and to investigate the structure of the  $b_4^+$  ion from leu-enkephalin<sup>9c</sup> and c-type ions formed by electron capture dissociation.<sup>11</sup> In the present work, we compare spectroscopy of the  $b_2^+$  ion from protonated AGG by CID in the hexapole collision cell with protonated cyclic AG (cAG), which has a known diketopiperazine structure as a neutral.

IRMPD spectroscopy experiments were performed using the CLIO Free Electron Laser (FEL) in Orsay, France, coupled to an electrospray ionization-Fourier transform ion cyclotron resonance mass spectrometer (7.0 T Bruker Apex Qe).<sup>12</sup> Figure 1 shows the IRMPD spectrum for protonated cAG. The major fragment produced by IRMPD appears at  $m/z$  101 and corresponds to the



**Figure 1.** IRMPD spectrum of protonated cAG plotted as fragmentation efficiency for formation of the most abundant fragment ion vs wavenumber and calculated spectra of two stable diketopiperazine isomers.



**Figure 2.** IRMPD spectrum of the  $b_2^+$  ion produced by fragmentation of AGG plotted as fragmentation efficiency for formation of the most abundant fragment ion vs wavenumber and calculated spectra of two oxazolone isomers.

loss of CO. Fragments also appear at  $m/z$  84 (101-NH<sub>3</sub>) and  $m/z$  73 (101-CO). The spectra obtained for each fragment were similar, but due to the lower signal-to-noise ratio for low abundance fragment ions, only the intensity of the most abundant fragment ion was plotted.

The IRMPD experimental results were compared with quantum chemical calculations to determine the structures that produce the best IR spectral match with the experimental spectrum. Both geometry optimizations and frequency calculations were performed at the B3LYP/6-311++G(d,p) level for different  $b_2^+$  oxazolone and diketopiperazine structures with protonation at oxygen or nitrogen atoms.<sup>13</sup> The acylium structure was excluded due to its higher energy compared to the two cyclic structures. Theoretical

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**Table 1.** Relative Calculated Enthalpy and Free Energy Changes for the Different Structures of the AG  $b_2^+$  Ion

	B3LYP/6-311++G(d,p)	
	$\Delta\Delta H_{298.15}$ (kJ/mol)	$\Delta\Delta G_{298.15}$ (kJ/mol)
diketo O Gly	0.0	0.0
diketo O Ala	2.2	3.1
oxaz N ring (Gly)	16.8	16.7
oxaz N free NH <sub>2</sub> (Ala)	31.4	31.6
diketo N Ala	56.1	56.9
diketo N Gly	63.1	63.8

IR frequencies were scaled with a factor of 0.978 and were convoluted with a Lorentzian profile with a 20  $\text{cm}^{-1}$  full width at half-maximum. The most pertinent spectra and geometries are compared with the experimental spectra in Figures 1 and 2, and the energies are summarized in Table 1 (see Supporting Information for other spectra and geometries).

The  $b_2^+$  diketopiperazine structure protonated on the glycine amide oxygen (diketo O Gly) has the lowest energy among all the diketopiperazine and oxazolone structures. The amide oxygen protonated diketopiperazines (diketo O Gly and diketo O Ala) have lower free energies than the amide nitrogen protonated forms (diketo N Ala and diketo N Gly) by 56.9 and 63.8 kJ/mol, respectively, which suggests that the amide oxygen protonated diketopiperazines are the major isomers rather than the nitrogen protonated ones. Among the oxazolones, the oxazolone ring nitrogen protonated isomer (oxaz N ring) is 14.9 kJ/mol lower in free energy than the free amino protonated form (oxaz N free NH<sub>2</sub>). Oxygen protonated oxazolone isomers were unstable according to the computational results; therefore only nitrogen protonated oxazolone isomers will be discussed further.

Cyclic AG neutral has a diketopiperazine structure, and IRMPD experiments and quantum chemical computations clearly demonstrate a diketopiperazine structure for protonated cAG. Among all the structures, the two diketopiperazine isomers protonated at amide oxygens (diketo O Ala and diketo O Gly) show the best agreement to the experimental spectrum of protonated cAG in Figure 1. The experimental band at 1775  $\text{cm}^{-1}$  corresponds to unprotonated carbonyl amide stretching (amide I mode). Protonation at a carbonyl group causes the C=O to stretch simultaneously with the C–N bond, which causes a shift in the carbonyl stretching band to lower wavenumbers (1705  $\text{cm}^{-1}$ ). The band at 1512  $\text{cm}^{-1}$  is an amide II mode. The band at 1202  $\text{cm}^{-1}$  is from the OH bending of the protonated carbonyl oxygen. The diketopiperazine isomer protonated on the Gly amide oxygen shows similar band patterns to that protonated on the Ala amide oxygen, with a slight shift in the amide I mode, which contributes to a broad band. The 3.1 kJ/mol free energy difference, similar band patterns, and broadened experimental bands suggest that protonated cAG is a combination of the alanine and glycine amide oxygen protonated isomers.

The  $b_2^+$  ion IR action spectrum of AGG shown in Figure 2 has a very different band pattern compared with the cAG spectrum. The carbonyl stretching bands are blue-shifted by  $\sim 200$  to 1970  $\text{cm}^{-1}$  and exist as a main band with a shoulder rather than the distinct doublet of protonated cAG. The  $b_2^+$  ion spectrum is in excellent agreement with quantum chemical calculations for the oxazolone protonated on the ring nitrogen (oxaz N ring). The long shoulder on the C=O stretching band indicates a minor contribution from an oxazolone protonated on the free NH<sub>2</sub> nitrogen (oxaz N free NH<sub>2</sub>). In addition to the carbonyl stretching band, there is a strong band at the 1630  $\text{cm}^{-1}$  region for the  $b_2^+$  ion which is absent for cAG. The band at 1630  $\text{cm}^{-1}$  and its shoulder are characteristic of the scissoring mode of free NH<sub>2</sub>. This band with a long shoulder

to the blue side in the experimental spectrum indicates that the feature is most likely a combination of two adjacent computed bands. The band at 1339  $\text{cm}^{-1}$  is from the twisting mode of free NH<sub>2</sub>. Because the diketopiperazine structure does not have a free NH<sub>2</sub>, the 1630 and 1339  $\text{cm}^{-1}$  bands are characteristic markers for an oxazolone structure. Although the protonated diketopiperazine exhibits a strong OH bending feature near 1202  $\text{cm}^{-1}$ , there is no OH functional group in the two nitrogen protonated oxazolones and no band at 1202  $\text{cm}^{-1}$  is observed in Figures 2 (top).

In conclusion, the experimental IRMPD data and quantum chemical calculations show that protonated cAG is a diketopiperazine and the  $b_2^+$  ion from CID of protonated AGG is an oxazolone. cAG retains its cyclic form upon protonation and is most likely a mixture of isomers protonated at alanine and glycine residues. In contrast, the  $b_2^+$  ion from AGG is an oxazolone with protonation at the ring nitrogen as the major isomer. This is the first time that an IR spectrum of a  $b_2^+$  ion has been obtained, and for this AG  $b_2^+$  ion, the oxazolone structure previously proposed based on calculations and experiments was confirmed.

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**Supporting Information Available:** Figures S1, S2, S3, additional band assignments, and complete ref 13. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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