



of racemization relative to ring size are being determined.

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## MASS SPECTRA OF ORGANIC MOLECULES. II. AMINO ACIDS<sup>2</sup>

Sir:

In our efforts to extend the applicability of mass spectrometry to organic molecules of extremely low volatility, we have been able to determine the mass spectra of amino acids without prior conversion to more volatile derivatives required for our earlier work.<sup>1,3</sup> Using the same technique which had made it possible to obtain mass spectra of nucleosides,<sup>4</sup> volatilization of the sample directly into the ion source and close to the ionizing electron beam gave excellent spectra of free amino acids and even their hydrochlorides.<sup>5</sup>

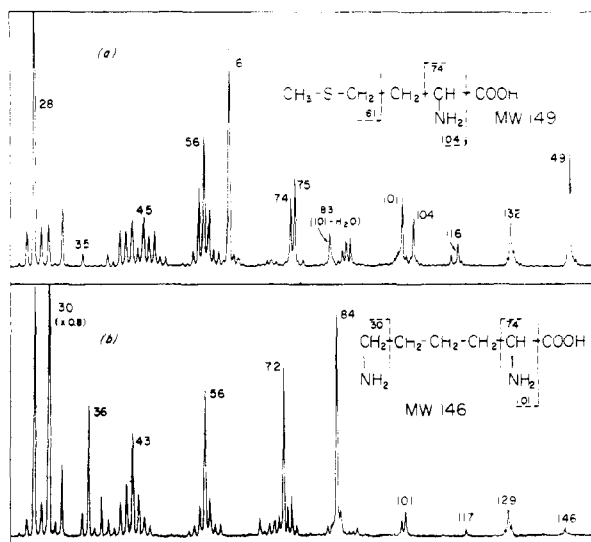


Fig. 1.—Reproductions of mass spectra of amino acids: (a) methionine; (b) lysine monohydrochloride (mass 28 and 32 due to air; mass 30 in (b)  $1.25 \times$  as abundant as shown).

These spectra<sup>6</sup> (determined with samples ranging from 0.25–10  $\mu\text{g}$ .) were quite similar to those of the corresponding ethyl esters,<sup>1</sup> indicating that free amino acids exist in the gas phase as the un-

(1) Paper I: K. Biemann, J. Seibl and F. Gapp, *J. Am. Chem. Soc.*, **83**, 3795 (1961).

(2) This investigation was supported by a grant from the National Aeronautics and Space Administration (NSG 211-62). We wish to thank Mr. M. Munroe for invaluable help with the instrumentation.

(3) K. Biemann, J. Seibl and F. Gapp, *Biochem. Biophys. Res. Commun.*, **1**, 307 (1959).

(4) K. Biemann and J. A. McCloskey, *J. Am. Chem. Soc.*, **84**, 2005 (1962).

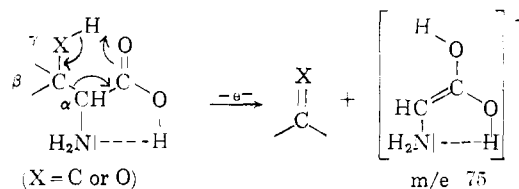
(5) W. L. Baun and D. W. Fischer [*Anal. Chem.*, **34**, 294 (1962)] reported the introduction of free amino acids into a spark source mass spectrometer. Our spectrum of lysine hydrochloride (Fig. 1b) indicates that the much more gentle conditions of sublimation of the sample at relatively low temperatures into an electron beam of 70 ev. as contrasted to sparking at high frequency with 100 Kv. lead to spectra which are much more characteristic of the original molecule.

(6) For experimental conditions see footnote 6 in reference 4. Samples were vaporized at temperatures between 80° and 200°.

dissociated amino carboxylic acids or possibly hydrogen-bonded forms thereof. It will, therefore, suffice to discuss only those peaks not present as such in the mass spectra of the amino acid ethyl esters.

The lack of the ester group leads to a shift of 28 mass units in peaks due to the elimination of a group other than the acid moiety (e.g.,  $m/e$  102 in ethyl esters vs.  $m/e$  74 in the acids). There seem to be only two new modes of fragmentation: First, there is always found a peak of significant intensity at  $m/e$  45 corresponding to the carboxyl group while the corresponding one at  $m/e$  73 in ethyl esters is absent. This we attribute to the stability of a positively charged carboxyl fragment ( $\text{O}=\text{C}=\text{O}^+-\text{H}$ ) which is equivalent to a protonated carbon dioxide molecule while the corresponding carboxy ion of mass 73 ( $\text{O}=\text{C}=\text{O}^+-\text{C}_2\text{H}_5$ ) is energetically less favored.

The second difference is found in the presence of a peak at  $m/e$  75 in the spectra of a number of amino acids, namely, all those containing a hydrogen atom in a  $\gamma$ -position. This rearrangement is well known<sup>7</sup> for fatty acids and their esters, but is not observed in  $\alpha$ -amino esters, because of the availability of the free electron pair on nitrogen.<sup>1</sup> In the free acids hydrogen bonding seems to decrease this effect, thus favoring this rearrangement which is very sensitive to the electron density at the atom attached to  $\text{C}_\alpha$ .<sup>8</sup>



The mass spectrum of methionine (Fig. 1a) illustrates the similarity to that of the ester.<sup>1</sup> The peaks at mass 74, 101, 132, and 149 (mol. wt.) are those which occur 28 mass units higher in the ethyl ester. Fragments of mass 75, 57, and 45 correspond to the additional fragmentation modes discussed above; ( $m/e$  75 is only partly due to  $\text{CH}_2-\text{S}-\text{CH}_2-\text{CH}_2^+$ , as evidenced by deuteration experiments).

It is thus possible to interpret the mass spectra of free amino acids based on the behavior of amino esters under electron impact as discussed previously.<sup>1</sup> Glutamic acid appears to dehydrate to pyroglutamic acid prior to sublimation, but hydroxyamino acids vaporize without decomposition.

The fragmentation processes discussed here and earlier<sup>1</sup> are corroborated by the mass spectra of  $\text{N}^{15}$ -labeled amino acids and of  $\text{N},\text{O}$ -perdeuterio derivatives which can easily be obtained.<sup>9</sup>

Even salts are amenable to this technique if both the corresponding base and acid are sufficiently volatile and thermally stable at the

(7) F. W. McLafferty, *Anal. Chem.*, **31**, 82 (1959).

(8) For a detailed discussion of the interpretation of mass spectra see K. Biemann, "Mass Spectrometry," McGraw-Hill Book Co., Inc., New York, N. Y., 1962, Chap. 3.

(9) For experimental details see footnote 9 in reference 4.