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Separation of ϵ -carboxymethyl-lysine from methionine on 60-cm columns of ion-exchange resins

Some peaks of carboxymethyl-(CM-)amino acids overlap the free amino acid peaks in the amino acid analysis performed according to or derived from the method of SPACKMAN *et al.*¹⁻⁴. GOREN *et al.*⁴ proposed an alternative procedure to the ninhydrin detection of CM-amino acids of alkylated proteins based on the use of radioactive alkylating agents and the detection of radioactive products with a flow cell for scintillation counting. In this case, the overlap of CM-amino acid peaks with free amino acid peaks is no longer a problem. When, however, this method cannot be used, it is necessary to have procedures for the separation of CM-amino acids from the free amino acids.

CM-histidine can be separated from the other ninhydrin-positive peaks if the pH of the first buffer of amino acid analysis is lowered from 3.25 to 3.19 (ref. 2). ϵ -CM-lysine is resolved from methionine on a 150-cm column system; however, they overlap on a 60-cm column system³⁻⁵. It must be pointed out that many amino acid analysers are equipped only with 60-cm columns.

In this note, the conditions for the resolution of ϵ -CM-lysine from methionine in a 60-cm column system are described. ϵ -CM-lysine and ϵ,ϵ -di-CM-lysine were prepared from poly-L-lysine (mol. wt. 50-100000; Cyclo Chem. Co.) as reported by GUNDLACH *et al.*⁵ S-CM-cysteine was purchased from Mann. All the other reagents were of analytical grade. A Beckman Unichrom amino acid analyser, equipped with 60-cm columns filled with Beckman custom-research resin type AA-15, was used. The standard working conditions of this apparatus are those described by BENSON AND PATTERSON⁶. The following operation conditions were used: flow rate of the buffer, 67.5 ml/h; flow rate of the buffer plus ninhydrin, 101.5 ml/h; first elution buffer, 0.2 N Na citrate (pH 3.25); second elution buffer, 0.2 N Na citrate (pH 3.8 instead of 4.25); buffer change after 120 min. The analysis was started at 42°, and immediately

TABLE I

ELUTION VOLUME OF SOME FREE AND CARBOXYMETHYLATED AMINO ACIDS

These elution volumes apply exactly only to our conditions (see the text). Each elution volume is the average of that in a number of similar chromatograms. The elution volume, calculated from the chromatograms, includes the volumes of the buffer and of the ninhydrin reagent.

| Amino acid | Elution volume (ml) |
|-----------------------------------|---------------------|
| ϵ,ϵ -Di-CM-lysine | 61.5 |
| S-CM-cysteine | 72.5 |
| Aspartic acid | 80.5 |
| Alanine | 166.5 |
| Half-cystine | 184.0 |
| Valine | 212.0 |
| Methionine | 256.0 |
| ϵ -CM-lysine | 273.5 |
| Isoleucine | 288.0 |
| Phenylalanine | 373.0 |

after starting the temperature was raised to 60°. This temperature was reached with our apparatus and with a room temperature of 20–22° in about 1 h.

The total analysis time for acidic and neutral amino acids under these operation conditions was 3 h 50 min instead of 3 h 10 min under standard conditions.

In Table I, the elution volume of some peaks of free amino acids and CM-amino acids is reported. Under the working conditions reported above, the ϵ -CM-lysine peak appears between methionine and isoleucine and is well separated from both these amino acids. ϵ,ϵ -Di-CM-lysine, S-CM-cysteine and aspartic acid peaks are also well separated from each other, and half-cystine peak is found between alanine and valine peaks.

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*Istituto di Chimica Biologica,
University of Pisa,
Pisa (Italy)*

G. RONCA
R. CHITI
A. LUCACCHINI

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