LOCALIZATION OF THE ASPARTIC ACID RESIDUE PRESENT IN THE ACTIVE CENTER OF PORCINE PEPSIN

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It is known that the catalytic center of porcine pepsin includes the β -carboxy group of aspartic acid. The use of diazoacetyl inhibitors has enabled the amino-acid sequence containing the "active" aspartic acid to be shown [1, 2]: Ile-Val-Asp-Thr-Gly-Thr-Ser. However, the part of the polypeptide chain of pepsin in which this sequence is present has not hitherto been known.

In an investigation of a thermolysin hydrolyzate of the B-3 fragment [3, 4] formed by the cleavage of carboxymethylated pepsin with cyanogen bromide, two peptides were isolated with the following structures:

The sequence 1-7 of peptide B3Th624 coincides with the sequence given above and, therefore, the aspartic acid residue participating in the formation of the active center of pepsin is present in the B-3 fragment. The peptide B3Th611 contains homoserine and is C-terminal in the B-3 fragment. Tang and Hartley [5] have previously reported that pepsin contains a sequence including a methionine residue: Tyr-Gly-Thr-Gly-Ser-Met-Asp-Val-Pro-Thr-Ser. However, Chen and Tang [6] have recently found a different and, in their opinion, more reliable methionine-containing sequence in pepsin:

 Phe-Gly-Gly-Met-Asp-Val-Pro-Thr-Ser-Ser-Gly-Gly-Leu.

 1
 2
 3
 4
 5
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 13

The sequence of residues 5-13 coincides with the N-terminal sequence of fragment B-1, which occupies the C-terminal position in the pepsin molecule [3]; the sequence 2-4 coincides with the C-terminal sequence of the fragment B-4 [7]. Apparently, in the molecule of porcine pepsin the B-4 fragment precedes the B-1 fragment. Tang and Hartley showed that pepsin contains the following methionine-including sequence:

> Asp-Ser-Ile-Tre-Met (Asp, Gly, Gly) (Ala, Tyr) [5]. 1 2 3 4 5 6 7 8 9 11

Sequence 1-5 coincides with the C-terminal sequence of the N-terminal fragment of pepsin B-2 [3]. The sequence 6-10 partially agrees with the N-terminal sequence of the fragment B-5; NH_2 -Asp-Gly-Glu-Thr-Ile [3]. In our laboratory, L. P. Revina, I. B. Pugacheva, and É. A. Vakhitova have isolated the following methionine-containing peptide:

Sequence 1-3 in it is the C-terminal sequence of fragment B-2, and sequence 4-7 coincides with the N-terminal sequence of fragment B-5. On the basis of these facts, it may be assumed that in the pepsin molecule fragment B-2 is followed by fragment B-5, so that the peptides formed in the cleavage of porcine pepsin by cyanogen bromide are arranged in the polypeptide chain of the enzyme in the following order:

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$$\frac{\text{NH}_2}{\text{B}=2} \xrightarrow{\text{Met}} \frac{\text{Met}}{\text{B}=3} \xrightarrow{\text{Met}} \frac{\text{Met}}{\text{B}=4} \xrightarrow{\text{Met}} \frac{\text{Met}}{\text{B}=1} \xrightarrow{\text{Ala-COOH}}$$

Thus, it can be seen that the aspartic acid residue in the active center of pepsin and reacting selectively with diazoacetyl inhibitors is located close to the C end of the molecule.

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