

possibly because of steric hindrance. HUGHES *et al.*¹ found that exhaustive reaction with O-methylisourea led to the reaction of 54–57 out of 64–68 amino groups, determined by Van Slyke amino nitrogen analysis. Partial reaction with GDMP occurs with the N-terminal amino groups of these proteins. About 60 % of the N-terminal aspartic acid of serum albumin reacted, while about 30 % of N-terminal leucine of lactoglobulin appeared to react.

More complete details and further studies on this reagent will be reported later.

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The C-terminal amino acid of carboxypeptidase-A

Although it has been shown that CPase-A consists of a single polypeptide chain with N-terminal asparagine² and a molecular weight of 34,000^{3,4}, no information has yet been made concerning its C-terminal residue. The C-terminal amino acid of CPase-A has now been found to be asparagine both by the enzymic method, in which the native crystalline CPase-A was allowed to act upon the denatured enzyme, and by the catalytic-hydrazinolysis method.

CPase-A used in this experiment was a specimen crystallized from frozen bovine pancreas and recrystallized 6–10 times according to NEURATH's method⁵, but including one DFP treatment. It was shown to be homogeneous ultracentrifugally and to yield aspartic acid (0.3–0.9 mole/mole) and serine (0.05–0.1 mole/mole) in the N-terminal determination by the DNP method, supporting the result obtained by THOMPSON². The amino acid composition determined by the DNP procedure⁶ was almost in agreement with that obtained by SMITH *et al.*⁴.

The CPase-A was denatured by incubating at room temperature (23–25°) for 2–3 h with 0.2 % aq. sodium dodecyl sulfate (pH about 7) in the presence of 0.04 *M* β -phenyl propionate, a strong competitive inhibitor of the enzyme⁷. After dialysis

Abbreviations: CPase-A, carboxypeptidase-A (Anson's¹); CPase-B, basic carboxypeptidase; DFP, diisopropyl fluorophosphate; DNP, dinitrophenyl.

overnight in a refrigerator against 0.05 *M* Veronal buffer, pH 7.5, the solution of the denatured CPase-A (0.4 %) was incubated at 25° with crystalline CPase-A in the presence of 0.1 *M* NaCl and 10⁻³ *M* DFP (substrate: enzyme = 40:1, molar ratio). Aliquots removed from the digestion mixture at various time intervals were dinitrophenylated in aq. NaHCO₃ at 40° and analyzed for amino acids. The release of asparagine reached 1 mole/mole CPase-A after about 2-h incubation, followed by the release of valine (0.85 mole/mole), threonine (0.7), leucine (0.7), glutamic acid (0.5) and others. No change was found in the N-terminal residue of the protein before and after the digestion. When performic acid-oxidized CPase-A⁸, hardly soluble in the buffer solution, was incubated at 25° for 20 h with CPase-A (substrate: enzyme = 50:1), an incomplete release of asparagine (about 0.4 mole/mole), threonine, valine, leucine and others could be observed. In contrast, no amino acid was liberated by the action of DFP-treated CPase-B (a fraction containing a strong CPase-B activity⁹) upon the oxidized CPase-A.

Catalytic hydrazinolysis at 60°¹⁰ of crystalline CPase-A also revealed the C-terminal asparagine (about 1 mole/mole as its di-DNP- β -hydrazide after 24 h), but gave no appreciable amount of other amino acids. The C-terminal asparagine was successfully determined for the first time from the hydrazinolysate of protein¹¹, by modifying the extraction procedure¹² followed by identification on LEVY's two-dimensional paper chromatogram⁶.

All these results show that CPase-A has asparagine in its C-terminal. Combining the result of THOMPSON with ours, it is concluded that the enzyme is composed of a single polypeptide chain, Asp(NH₂). ---- (Glu, Leu, Thr, Val)·Asp(NH₂). It may further be inferred that trypsin is not responsible for the presence of the C-terminal residue after the activation of procarboxypeptidase-A, because no basic amino acid but asparagine was found in the C-terminus. This finding must be kept in mind in considering the course of the reaction process.

After completion of this manuscript, a paper by GRASSMANN AND RIEDEL¹³ appeared stating that no C-terminus was found in CPase-A by the hydrazinolysis method. It is possible that these authors might have missed asparagine β -hydrazide from the C-terminus due to the aldehyde treatment.

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