ing radiations on the constitutive amino acids of DNase I with that on amino acid solutions seems unwarranted since the experimental conditions are not comparable. The ionic yield in terms of ammonia formation is a function of amino acid concentration, pH and radiation intensity and, in this respect, there seems no real basis for comparison. Furthermore, the respective state in which the amino acids are irradiated are obviously quite different.

#### SUMMARY

1. Significant changes in the amino acid composition of DNase I were observed after exposure of the enzyme to irradiation. Radiation-induced deamination appears to be the most important factor in the production of this change.

2. Exposure to large doses of ionizing radiation resulted in a significant change in the ultraviolet spectrum of DNase I. Although the absorption at 250 m $\mu$  increased linearly with radiation dose, the enzyme activity decreased exponentially.

3. These studies indicate that large doses of ionizing radiations sufficient to destroy the enzymic properties of this protein did not bring about an equally extensive change in the amino acid composition.

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# THE ACTION OF CHYMOTRYPSIN ON N-ALKYL DERIVATIVES OF PHENYLALANINE ETHYL ESTER

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Studies by NEURATH et al.<sup>1</sup> have shown that chymotrypsin can act on ester linkages as well as on amide bonds, provided that the other specificity requirements are met. Thus, chymotrypsin readily hydrolyzes the ester group of such compounds as benzoyl L-tyrosine ethyl ester, benzoyl L-phenylalanine ethyl ester, acetyl L-tyrosine ethyl ester, etc. It was first found that the replacement of the "secondary peptide bond", such as the benzoyl amino or the acetyl amino group of the above compounds, by a References p. 185.

free amino group greatly decreases the susceptibility of the resulting compound to chymotryptic action. This and other observations were explained by the assumption that the higher susceptibility of compounds bearing a "secondary peptide bond" is the result of the formation of two hydrogen bridges between this peptide bond and a complementary peptide bond on the enzyme surface. However, it was found by GOLDENBERG AND GOLDENBERG<sup>2</sup> that L-phenylalanine ethyl ester was readily hydrolyzed by chymotrypsin if the cleavage was carried out at pH 6.4 instead of the slightly alkaline medium (pH 7.8) usually employed in experiments with chymotrypsin. In view of these findings, it appeared of interest to study the action of chymotrypsin on those derivatives of phenylalanine ethyl ester in which one hydrogen of the amino group was replaced by an alkyl group.

#### EXPERIMENTAL

We prepared the following compounds: DL-N-ethylphenylalanine ethyl ester, DL-N-methylphenylalanine ethyl ester, and L- and D-N-methylphenylalanine ethyl ester.

D-N-Methylphenylalanine ethyl ester hydrochloride was prepared from L-phenylalanine in the same manner as the L-isomer. M.p. 133-134°C. Found: N, 5.8; neutral equivalent, 239 (Sørensen formol titration). Calculated: N, 5.7; neutral equivalent, 244.  $[\alpha]_{D}^{\infty} \rightarrow 9.6^{\circ}$  (5% in water).

DL-N-Methylphenylalanine ethyl ester hydrochloride was prepared from DL-phenylalanine in a way similar to the L- and D-compounds. Found: N, 5.8; neutral equivalent, 240 (Sørensen formol titration). Calculated: N, 5.7; neutral equivalent, 244.

DL-N-Ethylphenylalanine ethyl ester hydrochloride was prepared from DL-a-bromo-β-phenylpropionic acid<sup>10</sup> by treating it with ethylamine and by esterification of the obtained product by FISCHER'S method. The substance was recrystallized from ethanol. Found: N, 5.3; neutral equivalent, 257 (Willstätter-Waldschmidt-Leitz titration). Calculated: N, 5.4; neutral equivalent, 258. DL-Phenylalanine ethyl ester hydrochloride was prepared from DL-phenylalanine by

DL-Phenylalanine ethyl ester hydrochloride was prepared from DL-phenylalanine by FISCHER's method. The substance was recrystallized from ethyl acetate. Found: neutral equivalent, 228 (Willstätter-Waldschmidt-Leitz titration). Calculated: 230.

L-Phenylalanine ethyl ester hydrochloride was prepared from L-phenylalanine in a similar way to the DL-compound. It was recrystallized from alcohol-ether. Found: neutral equivalent, 228 (Willstätter-Waldschmidt-Leitz titration). Calculated: 230.

Worthington crystalline, salt-free chymotrypsin was used. As can be seen from the results summarized in Table I, at a pH range varying between 5.7 and 7.7 none of the N-alkyl compounds mentioned was susceptible to the action of chymotrypsin.

#### DISCUSSION

Considering the fact that N-benzoyl and N-acetyl derivatives of phenylalanine ethyl ester, as well as phenylalanine ethyl ester itself, are readily attacked by chymotrypsin, the resistance of N-methyl and N-ethyl derivatives of phenylalanine ethyl ester to chymotryptic activity is somewhat surprising. This resistance could be explained by a diminished tendency of the alkyl-substituted amino group to form a hydrogen bond, by proton donation, with an appropriate group on the enzyme surface. The following *References p. 185*.

L-N-Methylphenylalanine was prepared from D-phenylalanine according to the method of IZUMIVA et al.<sup>6,7</sup>.  $[a]_D^{23} + 49.5^{\circ}$  (in 0.15 N NaOH). FISCHER AND LIPSCHITZ<sup>8</sup> have reported  $[a]_B^{18} + 49.7^{\circ}$ . The substance was converted to the ethyl ester hydrochloride by the method used by FISCHER in the preparation of L-tyrosine ethyl ester<sup>9</sup>. The product obtained was recrystallized from alcoholether. M.p. 133-134°C (Fischer-Johns melting point apparatus). Found: N. 5.7; neutral equivalent, 248 (Willstätter-Waldschmidt-Leitz titration), 247 (Sørensen formol titration). Calculated: N. 5.7; neutral equivalent, 244.  $[a]_D^{20} + 9.4^{\circ}$  (5% in water). IZUMIVA AND FRUTON<sup>7</sup> have reported m.p. 133°C and  $[a]_D^{22} + 7.8^{\circ}$  (2% in water) for this compound.

Time, minutes Colorimeter reading Colorimeter reading Colorimeter reading Colorimeter reading Colorimeter reading Colorimeter reading Colorimeter reading Colorimeter reading Colorimeter reading Colorimeter reading	Cubetrate	Н		5.7			6.6			0.9				
Colorimeter reading**       465       400       355       470       355       296       465       300       244         Colorimeter reading**       470       465       450       455       300       244         Colorimeter reading**       470       460       450       450       455       450       455       455         Colorimeter reading**       435       430       435       420       415       450       50       135       218 <th>C MOST de C</th> <th>Time, minutes</th> <th>0</th> <th>10</th> <th>50</th> <th>0</th> <th>10</th> <th>50</th> <th>0</th> <th>10</th> <th>30</th> <th>0</th> <th>IO</th> <th>20</th>	C MOST de C	Time, minutes	0	10	50	0	10	50	0	10	30	0	IO	20
Colorimeter reading**       470       460       450       450       450       450       455       460       455       465       465       465       465       465       465       465       465       465       465       465       465       465       465       465       465       465       456       450       457       455       450       456       450       45	JL-PEE*	Colorimeter reading**	465	400	365	470	355	296	465	300	244	465	350	294
Colorimeter reading**       440       435       430       435       420       415       450       454       450         Colorimeter reading**       435       430       425       435       420       415       450       50       513       228<	ы-РЕЕ***	Colorimeter reading**	470	460	450	465	460	450	470	465	465	465	460	455
Colorimeter reading**       435       430       425       435       420       445       450       45	DL-N-CH <sub>3</sub> -PEE <sup>*</sup>	Colorimeter reading**	440	435	430	435	420	415	450	454	450	475	480	475
Colorimeter reading**       445       440       460       450       455       460       445       435       435         Colorimeter reading**       435       435       435       435       455       450       455       450       445       435         Colorimeter reading**       236       168       138       220       80       30       228       60       13         Colorimeter reading**       234       232       214       216       214       230       228       228         Colorimeter reading**       218       220       224       216       216       228       228       228         Colorimeter reading**       214       216       216       228       228       228       228         Colorimeter reading**       218       220       224       216       216       228       228       228         Colorimeter reading**       215       218       220       216       214       228       228       228       228         Colorimeter reading**       215       212       218       216       214       226       224       216       214       226       224       216 <td>01-N-CH<sub>3</sub>-PEE<sup>***</sup></td> <td>Colorimeter reading**</td> <td>435</td> <td>430</td> <td>425</td> <td>435</td> <td>420</td> <td>420</td> <td>445</td> <td>450</td> <td>450</td> <td>465</td> <td>475</td> <td>475</td>	01-N-CH <sub>3</sub> -PEE <sup>***</sup>	Colorimeter reading**	435	430	425	435	420	420	445	450	450	465	475	475
Colorimeter reading**       435       435       455       450       455       450       456       450       440         Colorimeter reading**       236       168       138       220       80       30       228       60       13         Colorimeter reading**       234       234       232       214       216       214       230       228       228         Colorimeter reading**       218       220       224       216       216       228       228       228         Colorimeter reading**       214       216       216       228       228       228       228         Colorimeter reading**       214       218       222       218       216       212       228       228       228         Colorimeter reading**       216       218       224       216       212       228       228       228       228       228         Colorimeter reading**       215       212       218       220       216       214       226       224       216	DL-N-C2H5-PEE*	Colorimeter reading**	445	445	440	460	450	455	460	445	435	470	475	475
Colorimeter reading**       236       168       138       220       80       30       228       60       13         Colorimeter reading**       234       232       214       216       214       230       228       228         Colorimeter reading**       234       232       214       216       214       230       228       228         Colorimeter reading**       218       220       224       216       216       228       222         Colorimeter reading**       214       218       222       218       216       230       228       228         Colorimeter reading**       214       216       216       212       218       226       228       228         Colorimeter reading**       215       212       218       220       216       214       226       224       216         Colorimeter reading**       215       212       218       220       216       214       226       224       216	0L-N-C <sub>2</sub> H <sub>5</sub> -PEE <sup>***</sup>	Colorimeter reading**	435	435	435	455	450	455	465	450	440	475	475	470
Colorimeter reading**       234       232       214       216       214       230       228       228         EF*       Colorimeter reading**       218       230       228       228       228       228         EF*       Colorimeter reading**       218       210       216       216       228       222         EE**       Colorimeter reading**       214       218       220       224       216       230       228       223         EE**       Colorimeter reading**       214       218       216       216       228       228       228         EE**       Colorimeter reading**       215       216       214       216       212       228       222         EE**       Colorimeter reading**       215       212       218       220       216       214       226       224       216	-PEE*	Colorimeter reading**	236	168	138	220	80	30	228	60	13	236	120	32
Colorimeter reading**       218       220       224       216       216       228       228       222         Colorimeter reading**       214       218       213       218       212       218       226       224       216       216       214       226       224       216	-PEE***	Colorimeter reading**	234	234	232	214	216	214	230	228	228	222	220	222
Colorimeter reading**       214       218       218       222       218       216       230       228       228       228       228       228       228       228       228       228       228       228       222       200       200       216       214       216       214       216       212       228       222       228       222       222       226       222       222       226       224       216       214       226       224       216       214       226       224       216       214       226       224       216       214       226       224       216       214       226       224       216       214       226       224       216       216       214       226       224       216       216       216       216       224       216       214       226       224       216       216       216       216       216       214       226       224       216       21	-N-CH3-PEF*	Colorimeter reading**	218	218	220	224	216	216	228	228	222	222	220	222
Colorimeter reading** 220 220 216 224 216 212 228 228 222 Colorimeter reading** 215 212 218 220 216 214 226 224 216	-N-CH <sub>3</sub> -PEE***	Colorimeter reading**	214	218	218	222	218	216	230	228	228	222	220	222
Colorimeter reading** 215 212 218 220 216 214 226 224 216	-N-CH3-PEE*	Colorimeter reading**	220	220	216	224	216	212	228	228	222	216	212	216
•	D-N-CH <sub>3</sub> -PEE ***	Colorimeter reading**	215	212	218	220	216	214	226	224	216	218	218	218

TABLE I

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consideration may serve as an alternative explanation of our findings. GOLDENBERG AND GOLDENBERG<sup>2</sup> found that optimal hydrolysis of phenylalanine ethyl ester occurred at pH 6.4 when the rate of cleavage was measured by titration of the liberated acid. However, when they investigated the action of chymotrypsin on the same compound by measuring the disappearance of the ester, using HESTRIN's hydroxamic acid method<sup>3</sup>, they found an apparent increase in the rate of the reaction and a distinct broadening of the pH range of enzymic activity with a shift towards the alkaline region. This discrepancy can easily be understood if one realizes that HESTRIN'S method measures, in addition to hydrolysis, the disappearance of ester caused by transpeptidation reactions whose pH optimum may differ from that of the hydrolysis<sup>4</sup>. Since we have not found any activity of chymotrypsin with our compounds using HESTRIN'S method, it appears that neither hydrolysis nor replacement reactions occurred. In order to explain the lack of susceptibility of the N-alkyl derivatives to chymotryptic activity we suggest that phenylalanine ethyl ester is not hydrolyzed directly by chymotrypsin but is first converted, by a transpeptidation reaction, to phenylalanylphenylalanine ethyl ester (or to the ester of a higher peptide). This compound bearing a "secondary peptide bond" is then rapidly hydrolyzed by the enzyme. The inability of chymotrypsin to carry out transpeptidation reactions with the Nalkyl-substituted esters would then account for the resistance of such compounds to the action of the enzyme. It may be recalled that TAUBER<sup>5</sup> obtained phenylalanylphenylalanine ethyl ester on treating phenylalanine ethyl ester with chymotrypsin at pH 8.8. He also found that chymotrypsin did not markedly hydrolyze this dipeptide ester at pH 7.7. If our assumption is correct, namely that this ester acts as an intermediary compound in the chymotryptic hydrolysis of phenylalanine ethyl ester, it would seem that the optimal pH for its cleavage must be in the slightly acid region.

Experiments to check the validity of the above explanations are under way.

#### SUMMARY

The action of chymotrypsin on N-methyl and N-ethyl derivatives of phenylalanine ethyl ester at a pH range varying between 5.7 and 7.7 was investigated. Neither hydrolysis nor replacement reactions could be detected.

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