## RESOLUTION OF a-METHYL AMINO ESTERS BY CHYMOTRYPSIN

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Abstract: The resolution of DL- $\alpha$ -MeTrp-OMe HCl, DL- $\alpha$ -MePhe-OMe HCl and DL- $\alpha$ -Me-pF-Phe-OMe HCl have been achieved using  $\alpha$ -chymotrypsin at pH 5.0.

 $\alpha$ -Alkyl amino acids are gaining importance in structure-activity relationship studies of peptide hormones<sup>1</sup>. Introduction of  $\alpha$ -methyl residues into the naturally occurring sequence gives rise to peptide analogs having constrained conformation. In addition, analogs with  $\alpha$ -methyl residues are often resistant to hydrolysis by proteolytic enzymes and thus are active for a longer time.

Resolution of several  $\alpha$ -methyl-Di-amino acids has been carried out by fractional crystal-lization of diastereoisomeric salts<sup>2,3</sup> and by the action of carboxypeptidase A on the N-tri-fluoroacetyl derivatives<sup>1</sup>. The latter method works well but N-TFA derivatives are not always easy to prepare in good yield. Acylase, often used to resolve acyl amino acids, acts very slowly on most  $\alpha$ -methyl substrates<sup>2</sup>.

We have found that chymotrypsin hydrolyzes the L-isomers of some  $\alpha$ -methyl amino acid esters having an unacylated amino group. The rate is slow compared to the corresponding simple  $\alpha$ -amino acid esters, but sufficiently rapid to be a practical method for resolving a DL mixture. The hydrolysis of unacylated amino acid esters by chymotrypsin has been studied  $^{4-5}$ , but its action on unacylated  $\alpha$ -methyl amino acid esters apparently has not been noted. Introduction of an  $\alpha$ -methyl group into a good substrate for chymotrypsin, e.g. acetyl-L-tyrosine amide lowers the rate of enzymatic hydrolysis by a factor of  $10^5$ , which is attributed not so much to decreased binding affinity but to an incorrect orientation of the bond to be cleaved  $^2$ . We find that the introduction of an  $\alpha$ -methyl group into an unacylated amino acid ester lowers the rate of hydrolysis also, but much less dramatically.

The stereospecificity of enzymatic action on  $\alpha$ -alkyl amino acid derivatives cannot be assumed to be the same as for the natural amino acids<sup>6</sup>. In the present case, we show that chymotrypsin hydrolyzes only the L (S) isomer of  $\alpha$ -methylphenylalanine methyl ester; the absolute configuration of L- $\alpha$ -methylphenylalanine has been determined<sup>3,7</sup>, and we assume that this specificity is extended to  $\alpha$ -methyltryptophan methyl ester.

DL- $\alpha$ -methyltryptophan methyl ester was prepared by the method of Brana et al. <sup>8</sup>. Although they claim a stereoselective synthesis, their product has a very low rotation and in our hands their procedure gave only the DL-mixture. DL- $\alpha$ -methylphenylalanine and the p-fluoro derivative were prepared by following the procedure of Stein et al. <sup>9</sup>.

The resolution is described in detail for DL- $\alpha$ -methyltryptophan methyl ester. A solution of DL- $\alpha$ -methyltryptophan methyl ester (10.2 g, 40 mmoles) in 500 ml water at 37°C was adjusted to pH 5.0 by the addition of 0.5 N LiOH.  $\alpha$ -Chymotrypsin<sup>10</sup> (500 mg) was added and the solution was stirred gently while the pH was maintained at 5.0 by the addition of 0.5 N LiOH with pH-stat. The uptake of base stopped after 40 hours; the solution was stirred with 1 g of charcoal and filtered to remove the enzyme, then adjusted to pH 10.0 with LiOH. Extraction with ethyl acetate yielded D- $\alpha$ -methyltryptophan methyl ester, 88% yield, crystallized as its hydrochloride. The aqueous solution was lyophilized and the solid residue, containing L- $\alpha$ -methyltryptophan together with about 4% of D-isomer from non-enzymatic hydrolysis, was reconverted to the methyl ester with thionyl chloride and methanol. Treatment with  $\alpha$ -chymotrypsin as above for 30 hours gave L- $\alpha$ -methyltryptophan essentially free of the D-isomer. The results are summarized in the Table.

Table	e
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Product	Reaction Time (h)	Yield, %	m.p.	$[\alpha]_{\mathrm{D}}^{25}$
D-o-MeTrp-OMe HCl	40	88	160° dec.	-26.3° (C9.9, MeOH)
L-G-MeTrp-OMe HCl	30	72	159° dec.	+27.8° (C9.9, MeOH)
D-o-MePhe-OMe HCl	35	79	112-115°	-23.6° (C.8, NHCl)
L-o-MePhe-OMe HCl		66		
D-a-Me-p-F-Phe-Me HCl	35	78	142-144°	-11.2° (C1.0, MeOH)
D-0-Me-Phe				+19.3° (C1.0, H <sub>2</sub> 0) <sup>11</sup>

## References and Notes

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- 11. D- $\alpha$ -MePhe was obtained by refluxing the methyl ester with 6 N HCl and neutralizing. ACKNOWLEDGMENT: This work was supported by a grant from the National Institutes of Health (R01 HD12776) and by the Grace M. Showalter Residuary Trust. We wish to thank Mr. Patrick Edwards for technical assistance.

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