

THE RELATIVE SUSCEPTIBILITY TO RACEMIZATION OF
L- AND D-RESIDUES IN PEPTIDE SYNTHESIS

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ABSTRACT: An activated D-residue of a protected dipeptide when coupled racemizes twice as much as an activated L-residue, which is contrary to the case for couplings of acylamino acids.

The extent of racemization in peptide synthesis depends on many factors¹ including the coupling method, solvent polarity, and the nature of the residue undergoing activation.^{2,3} Moreover, it depends in particular on the type of N-substituent on the activated residue, namely alkoxycarbonyl [ROCO-] << acyl [RCO-] < acylaminoacyl [R'CONHCHRCO-].^{1,3} Only recently has the influence of the configuration of the activated residue been demonstrated. Arendt *et al.*⁴ showed that Ac-L-Leu racemizes three times as much when coupled with L-Leu-OMe than when coupled with D-Leu-OMe [Table]. The inference is that more racemization will be observed when coupling an L-residue with an L-isomer than when coupling a D-residue with an L-isomer. We have confirmed this hypothesis using benzoylamino acids,⁵ the L-residues racemizing more than the D-residues when coupled with an ester of L-lysine [Table]. However, we now find that this phenomenon holds for acylamino acids only. When protected dipeptides [Z-Gly-Xxx] were coupled, the activated D-residues [Xxx] underwent twice as much stereomutation as did the L-residues [Table]. This was observed for three different residues [Leu, Phe, Val] coupled using DCC alone in dichloromethane or DMF, or in the presence of HOBT⁶ in aqueous DMF. In the latter case, water was added in order to increase the racemization³ so that a clear difference would be observed. The conclusion to be drawn from the data is that unless an adjacent chiral residue instead of glycylic in this case would drastically change the course of events, the danger of racemization is twice as great when a peptide bond is formed between residues of opposite configuration than in normal peptide synthesis.

We have already described³ cases where conclusions based on model tests implicating activated benzoylamino acids did not apply to the couplings of activated dipeptides [Z-Gly-Xxx]. The additional example presented here serves to emphasize the caution which should be exercised in interpreting racemization data obtained using acylamino acids.

Experimental: To a mixture containing Z-Gly-Xxx⁷ [0.50 mmol], L-Lys(Z)-OBzl.HCl⁸ [0.50 mmol] and N-methylmorpholine [0.45 mmol] in 25 ml of solvent was added HOBT⁶ [0.50 mmol] where pertinent followed by DCC [0.50 mmol]. The diastereomeric Gly-D/L-Xxx-Lys peptides were analyzed with an amino acid analyzer⁷ after appropriate work-up and deprotection of the coupling mixtures.³

Abbreviations: Ac, acetyl; Bz, benzoyl; Bzl, benzyl, DCC, N,N'-dicyclohexylcarbodiimide; DMF, N,N-dimethylformamide; HOBT, 1-hydroxybenzotriazole; Z, benzyloxycarbonyl.

Table. Racemization during coupling with an amino acid ester^a

Substrate	Coupling method	Xxx					
		L-Leu	D-Leu	L-Phe	D-Phe	L-Val	D-Val
Z-Gly-Xxx ^b	DCC/CH ₂ Cl ₂	24	56	28	58	16	34
	DCC/DMF	35	70	42	62	64	106
	DCC-HOBT/DMF:H ₂ O (2:1)	12	23	16	36	34	64
Bz-Xxx ^c	DCC/CH ₂ Cl ₂	104	65	115	71	72	31
Ac-Xxx ^d	DCC/dioxane	80	26 ⁺				

a) Data given as % racemization = $2 \times [D-L \text{ or } L-L]/[L-L + D-L]$.

b) Coupling of Z-Gly-Xxx with L-Lys(Z)-OBzl.

c) Coupling of Bz-Xxx with L-Lys(Z)-OMe. Data of Benoiton *et al.*⁵

d) Coupling of Ac-L-Leu with L- and D-Leu-OMe⁺. Data of Arendt *et al.*⁴

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