

## ***O*-(*N*-Succinimidyl)-1,1,3,3-tetramethyluronium Tetrafluoroborate–*N*-Hydroxysuccinimide–CuCl<sub>2</sub>: A Facile and Reliable System for Racemization-Free Coupling of Peptides Having a Carboxy-terminal *N*-Methylamino Acid<sup>1)</sup>**

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**Simultaneous use of *N*-hydroxysuccinimide (HOSu) and CuCl<sub>2</sub> with a HOSu-based uronium coupling reagent, *O*-(*N*-succinimidyl)-1,1,3,3-tetramethyluronium tetrafluoroborate, has been found to eliminate the racemization of the carboxy-terminal *N*-methylamino acid residue during the segment condensation.**

**Key words** *N*-methylamino acid; racemization-free coupling; copper(II) chloride; racemization; segment condensation; peptide synthesis

In peptide synthesis by the convergent strategy the carboxy-terminal amino acid residue of the peptide segment is exposed to the risk of racemization.<sup>3)</sup> Extensive studies on the prevention of racemization have thus been continued, and some compounds, *e.g.*, *N*-hydroxysuccinimide (HOSu),<sup>4)</sup> 1-hydroxybenzotriazole (HOBt),<sup>5)</sup> *N*-hydroxy-5-norbornene-2,3-dicarboximide,<sup>6)</sup> 1-hydroxy-7-azabenzotriazole (HOAt),<sup>7)</sup> and CuCl<sub>2</sub>,<sup>8,9)</sup> were found to reduce the racemization in the carbodiimide-mediated coupling effectively. However, the situation is quite different when the carboxy-terminal residue is an *N*-methylamino acid, which is considerably more susceptible to racemization than a common amino acid. For instance, McDermott and Benoiton reported that the dicyclohexylcarbodiimide (DCC)–HOSu coupling of (*N*-benzyloxycarbonyl-alanyl)-*N*-methylleucine and H–Gly–OBzl (Bzl, benzyl) in tetrahydrofuran at room temperature was accompanied by 11% racemization of the *N*-methylleucine residue.<sup>10)</sup> Though the racemization decreased to an undetectable level on cooling to –5 °C,<sup>11)</sup> more convenient and reliable methods are desirable.

Recently, we have reported that CuCl<sub>2</sub> could eliminate the racemization of the carboxy-terminal *N*-methylamino acid residue in the 1-ethyl-3-(3-dimethylaminopropyl)carbodiimide hydrochloride (WSCl)-mediated segment coupling at room temperature,<sup>12)</sup> but could not sufficiently suppress the racemization in the coupling with onium-based reagents including benzotriazolyl-*N*-oxytris(dimethylamino)phosphonium hexafluorophosphate (BOP)<sup>13)</sup> and *O*-(7-azabenzotriazol-1-yl)-1,1,3,3-tetramethyluronium hexafluorophosphate (HATU).<sup>7)</sup> This class of coupling reagents is popularly used in peptide synthesis because of its advantages over carbodiimides; compared to carbodiimides, onium reagents usually complete couplings in a short time without any serious side reactions, give no troublesome by-products such as urea derivatives, and are safe to use from the standpoint of allergenicity. If racemization can be suppressed sufficiently, therefore, onium-based reagents would be convenient for use in the segment coupling of peptides having the carboxy-terminal *N*-methylamino acid and could facilitate efficient synthesis of *N*-methylamino acid-containing peptides by the convergent strategy. We have thus directed our study to the development of a racemization-free coupling system using an

onium-based coupling reagent.

Leading studies by McDermott and Benoiton<sup>10)</sup> and Davies and Mohammed<sup>11)</sup> suggested that racemization of the carboxy-terminal *N*-methylamino acid in the DCC–HOBt coupling occurs by two major routes as outlined in Fig. 1. Elimination of the racemization by CuCl<sub>2</sub> in the carbodiimide-mediated coupling<sup>12)</sup> reveals that CuCl<sub>2</sub> inhibits almost completely the oxazolonium pathway (Fig. 1a). On the contrary, insufficient suppression of the racemization by CuCl<sub>2</sub> in the BOP-mediated coupling<sup>12)</sup> implies that CuCl<sub>2</sub> cannot inhibit the active ester pathway (Fig. 1b). Racemization by pathway b, however, is known not to appear in the HOSu ester coupling.<sup>11)</sup> This suggests that pathway b can possibly be omitted when a HOSu-based onium salt is used as a coupling reagent. Taking this knowledge of racemization-mechanisms into consideration, we have designed a novel coupling system consisting of *O*-(*N*-succinimidyl)-1,1,3,3-tetramethyluronium tetrafluoroborate (TSTU)<sup>14)</sup> and CuCl<sub>2</sub> to achieve racemization-free segment coupling at the *N*-methylamino-acyl bond, and evaluated it in a simple model reaction between Boc–Phe–MeAla–OH (Boc, *tert*-butoxycarbonyl; MeAla, *N*-methylalanine) and H–Phe–OBzl.

Boc–Phe–MeAla–OH<sup>12,15)</sup> was coupled with H–Phe–OBzl in *N,N*-dimethylformamide (DMF) by TSTU in the presence or absence of additives, HOSu, HOBt, and CuCl<sub>2</sub>. After storage of each reaction mixture at room temperature for >24 h, a portion was subjected to HPLC analysis, and % L–D–L isomer was determined as (peak area of the L–D–L isomer) × 100 / [(peak area of the L–L–L isomer) + (peak area of the L–D–L isomer)]. Coupling yields were determined from the sum of peak areas of diastereomers in comparison with a standard solution of the L–L–L authentic sample. Partial HPLC profiles of samples from coupling with TSTU, TSTU–CuCl<sub>2</sub>, and TSTU–HOSu–CuCl<sub>2</sub> are shown in Fig. 2.

The results are listed in Table 1 with some data selected from our previous report<sup>12)</sup> for comparison. A significant amount of the L–D–L product was detected in the coupling with TSTU alone, and also in the presence of HOSu and HOBt. However, CuCl<sub>2</sub> could reduce the amount of the L–D–L product in the TSTU coupling, whereas it was not sufficiently effective in the BOP- and HATU-mediated couplings. These results indicate that the racemization by both pathways

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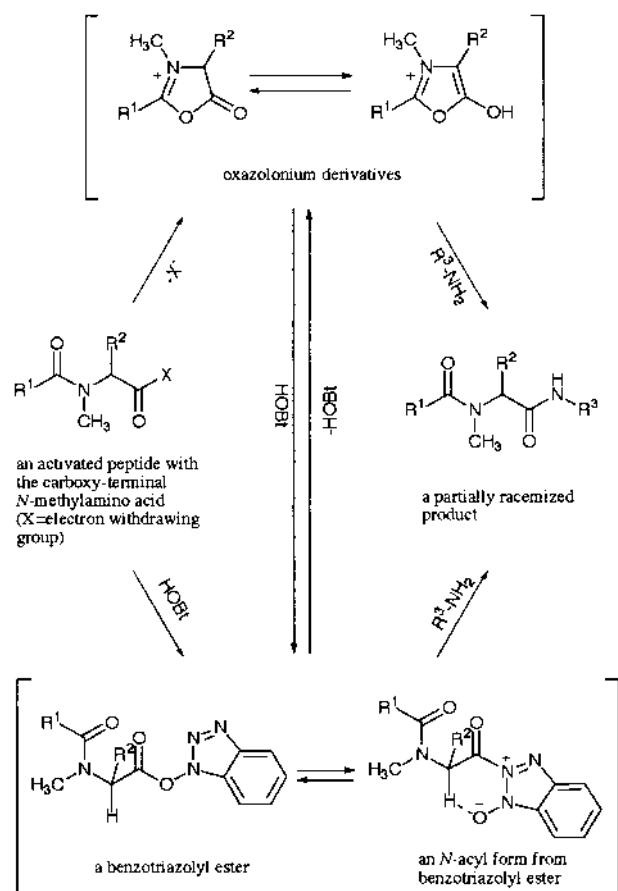
**(a) oxazolonium pathway****(b) active ester pathway**

Fig. 1. Two Major Pathways for Racemization of the Carboxy-Terminal *N*-Methylamino Acids in Segment Condensation

Table 1. Extents of Racemization during the Coupling of Boc-Phe-MeAla-OH with H-Phe-OBzl

Additive <sup>a)</sup>	% L-D-L <sup>b)</sup>			% yield <sup>b)</sup>		
	TSTU <sup>c)</sup>	Bop <sup>c,d)</sup>	HATU <sup>c,d)</sup>	TSTU <sup>c)</sup>	Bop <sup>c,d)</sup>	HATU <sup>c,d)</sup>
—	18.1	13.6	9.5	80	95	103
HOSu	14.0	14.5	11.1	73	95	97
HOBt	18.9	15.9	20.9	92	97	102
CuCl <sub>2</sub>	0.4	2.7	4.7	47	35	52
HOSu-CuCl <sub>2</sub>	<0.2	3.5	8.1	45 <sup>e)</sup>	27	46

a) One eq. b) Average of two independent experiments. c) Coupling reagents, 1.25 eq.; diisopropylethylamine, 2 eq. d) Ref. 12. e) 60% with 2.5 eq. of TSTU.

a and b can be suppressed effectively by a combination of TSTU and CuCl<sub>2</sub>. A small amount of the L-D-L product was, however, detected even in the presence of CuCl<sub>2</sub>. In the segment condensation of common peptides, BOP-HOBt was reported to result in a lower racemization level than BOP alone.<sup>16)</sup> We thus examined the simultaneous use of HOSu and CuCl<sub>2</sub> in the TSTU coupling. No appreciable racemization was detected by HPLC at an identification limit of 0.2%, and TSTU-HOSu-CuCl<sub>2</sub> proved to be an appropriate combination to suppress the racemization in the coupling of peptide segments having a carboxy-terminal *N*-methylamino

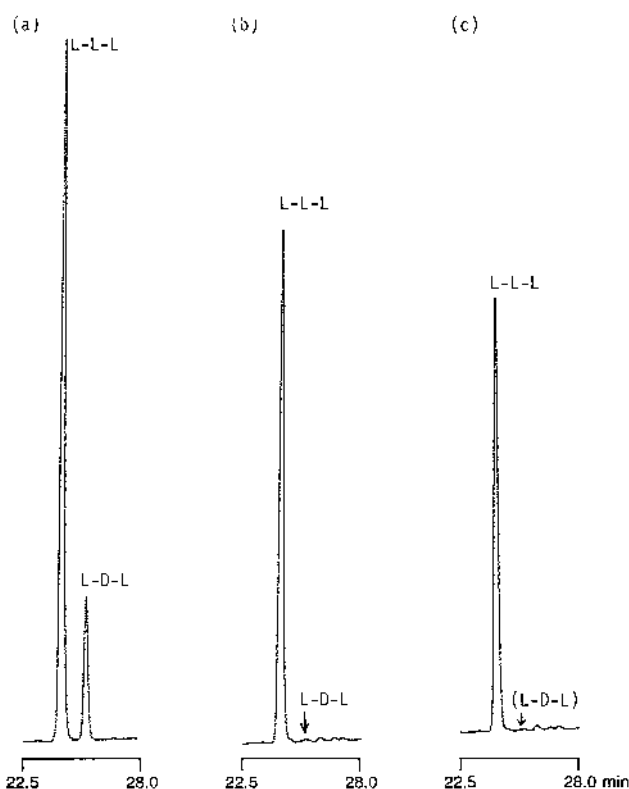


Fig. 2. Partial HPLC Profiles of Samples from Coupling with (a) TSTU, (b) TSTU-CuCl<sub>2</sub>, and (c) TSTU-HOSu-CuCl<sub>2</sub>

acid. In the WSCI-CuCl<sub>2</sub> coupling of Boc-Phe-MeAla-OH and H-Phe-OBzl, a small amount of the L-D-L product (0.2%) was detected, and the yield was similar (46%) to that of TSTU-HOSu-CuCl<sub>2</sub>.<sup>12)</sup> When HOSu and CuCl<sub>2</sub> were simultaneously used in the WSCI-coupling, racemization was not detected, but the yield decreased to 33%.<sup>12)</sup> Thus the TSTU-HOSu-CuCl<sub>2</sub> system would be more favorable than WSCI-CuCl<sub>2</sub> or WSCI-HOSu-CuCl<sub>2</sub> systems.

In conclusion, a novel coupling system consisting of TSTU, HOSu, and CuCl<sub>2</sub> has proved to completely eliminate the racemization of the carboxy-terminal *N*-methylamino acid residue during segment condensation. This is the first example of an onium-based coupling system which enables the racemization-free segment condensation at the *N*-methylaminoacyl bond. Optimization of the reaction conditions is now under way,<sup>17)</sup> facilitating the efficient convergent synthesis or head-to-tail cyclization of *N*-methylamino acid-containing peptides.

**Experimental**

In the HPLC analysis, elution profiles were monitored at 220 nm by a Waters 480 UV/VIS tunable absorbance detector and recorded with a SIC Chromatocoder-21.

Boc-Phe-MeAla-OH, Boc-Phe-MeAla-Phe-OBzl, and Boc-Phe-D-MeAla-Phe-OBzl were prepared as previously reported.<sup>12)</sup> TSTU was purchased from Novabiochem (U.S.A.). DMF was of peptide synthesis grade, and used without purification. CuCl<sub>2</sub> used in this study was its dihydrate.

**Racemization Test** The coupling reactions were conducted using the following stock solutions: A, 0.04 M Boc-Phe-MeAla-OH in DMF; B, 0.04 M H-Phe-OBzl (prepared from H-Phe-OBzl·HCl in the usual manner) in DMF; C, 0.25 M TSTU in DMF; D, 0.1 M diisopropylethylamine in DMF; E, 0.4 M HOSu in DMF; F, 0.4 M CuCl<sub>2</sub> in DMF.

TSTU Method: To a mixture of A (50 μl, 2 μmol), B (50 μl, 2 μmol), C (10 μl, 2.5 μmol), and DMF (10 μl) was added D (40 μl, 4 μmol).

TSTU-HOSu Method: To a mixture of A (50 μl, 2 μmol), B (50 μl,

2  $\mu\text{mol}$ ), C (10  $\mu\text{l}$ , 2.5  $\mu\text{mol}$ ), E (5  $\mu\text{l}$ , 2  $\mu\text{mol}$ ), and DMF (5  $\mu\text{l}$ ) was added D (40  $\mu\text{l}$ , 4  $\mu\text{mol}$ ).

**TSTU–CuCl<sub>2</sub> Method:** To a mixture of A (50  $\mu\text{l}$ , 2  $\mu\text{mol}$ ), B (50  $\mu\text{l}$ , 2  $\mu\text{mol}$ ), C (10  $\mu\text{l}$ , 2.5  $\mu\text{mol}$ ), F (5  $\mu\text{l}$ , 2  $\mu\text{mol}$ ), and DMF (5  $\mu\text{l}$ ) was added D (40  $\mu\text{l}$ , 4  $\mu\text{mol}$ ).

**TSTU–HOSu–CuCl<sub>2</sub> Method:** To a mixture of A (50  $\mu\text{l}$ , 2  $\mu\text{mol}$ ), B (50  $\mu\text{l}$ , 2  $\mu\text{mol}$ ), C (10  $\mu\text{l}$ , 2.5  $\mu\text{mol}$ ), E (5  $\mu\text{l}$ , 2  $\mu\text{mol}$ ), and F (5  $\mu\text{l}$ , 2  $\mu\text{mol}$ ) was added D (40  $\mu\text{l}$ , 4  $\mu\text{mol}$ ).

**Determination of % L–D–L:** After >24 h reaction at room temperature, each reaction mixture was diluted with DMF (640  $\mu\text{l}$ ), and a portion (10  $\mu\text{l}$ ) of the resultant solution was analyzed by HPLC [column, Waters  $\mu\text{Bondasphere 5C}_{18}$  100 Å (3.9 $\times$ 150 mm); solvent system, 0.05% trifluoroacetic acid (TFA) in water:0.05% TFA in MeCN 60:40 to 20:80 in 40 min at a flow rate of 1.0 ml/min. Boc–Phe–MeAla–Phe–OBzl,  $t_R$  24.26 min; Boc–Phe–D–MeAla–Phe–OBzl,  $t_R$  25.44 min]. % L–D–L and % yield were determined as follows: % L–D–L = (peak area of the Boc–Phe–D–MeAla–Phe–OBzl)  $\times$  100 / [(peak area of the Boc–Phe–MeAla–Phe–OBzl) + (peak area of the Boc–Phe–D–MeAla–Phe–OBzl)]; % yield = [(peak area of the Boc–Phe–MeAla–Phe–OBzl) + (peak area of the Boc–Phe–D–MeAla–Phe–OBzl)]  $\times$  100 / [peak area of standard solution of Boc–Phe–MeAla–Phe–OBzl (2  $\mu\text{mol}$ /800  $\mu\text{l}$  DMF, 10  $\mu\text{l}$  injection)]. The results are summarized in Table 1.

## References and Notes

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