

'Click peptide': a novel 'O-acyl isopeptide method' for peptide synthesis and chemical biology-oriented synthesis of amyloid β peptide analogues

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Abstract: After over a decade of studies on aspartic protease inhibitors and water-soluble prodrugs, we have been developing a novel method, since 2003, called 'O-acyl isopeptide method', for the synthesis of peptides containing difficult sequences. With our recent discoveries of 'O-acyl isodipeptide unit' and the 'racemization-free segment condensation method', this method has further evolved as a general synthetic method for peptides. Moreover, 'Click Peptide', which could be a powerful tool for identifying the pathological functions of amyloid β peptides in Alzheimer's disease, represents a valuable use of the isopeptide method in Chemical Biology-oriented research. Copyright © 2006 European Peptide Society and John Wiley & Sons, Ltd.

Keywords: O-acyl isopeptide method; Alzheimer's disease; amyloid β peptide; click peptide; difficult sequence

INTRODUCTION

Since 2003, we have been developing a novel method, called 'O-acyl isopeptide method', for the synthesis of peptides containing difficult sequences in which a native amide bond at a hydroxyamino acid residue, such as Ser being isomerized to an ester bond, is followed by an O–N intramolecular acyl migration reaction [1–10]. Recently, the 'O-acyl isopeptide method' began to be widely utilized by several other groups [11–15]. In chemical biology-oriented research, we developed a novel 'Click Peptide' based on the O-acyl isopeptide method to study the inherent biological functions of native peptides or proteins (Figure 1) [3–7,9].

O-ACYL ISOPEPTIDE METHOD

Several years ago, when we tried to synthesize some peptide derivatives, including phenylnorstatine, for the study of aspartic protease inhibitors [16–18], some of the synthesized compounds could not be purified in preparative scale HPLC owing to their extremely low solubility in various solvents (Figure 2(A)). Thus, these peptide derivatives were considered to be the so-called 'difficult sequence'-containing peptides [19,20].

On the other hand, for over a decade, we studied the pH-dependent 'O–N intramolecular acyl migration [21,22]-type water-soluble prodrugs of the peptide mimetic HIV-1 protease inhibitors [23,24]. These prodrugs, which are O-acyl isoforms of parent drugs possessing α -hydroxy- β -amino acids, had higher water solubility because of a newly formed and ionized amino group. Moreover, migration to the N-acyl parent drugs could be transacted with no side reaction under physiological conditions.

As a consequence, in 2003, we considered that the hydrophilic 'O-acyl isopeptides' derived from the phenylnorstatine-containing peptide derivatives would overcome the solubility problem in HPLC purification (Figure 2(B)). However, we had a surprising discovery in this research, which showed that not only did the 'O-acyl isopeptide' possess a higher solubility in various media, but also that the coupling and deprotection efficacy during solid-phase peptide synthesis (SPPS) was improved by modifying the nature of the difficult sequence [1,2]; namely, the isomerization of the peptide backbone from the N-acyl to O-acyl isopeptide structure, i.e. formation of one single ester bond, significantly changed the unfavorable secondary structure of the native peptides. Thus, this finding led to the development of the 'O-acyl isopeptide method' as a novel synthetic method in the field of Peptide Chemistry.

We also designed an 'O-acyl isodipeptide unit', e.g. Boc-Ser/Thr(Fmoc-Xaa)-OH. The use of O-acyl isodipeptide units, in which the racemization-inducing

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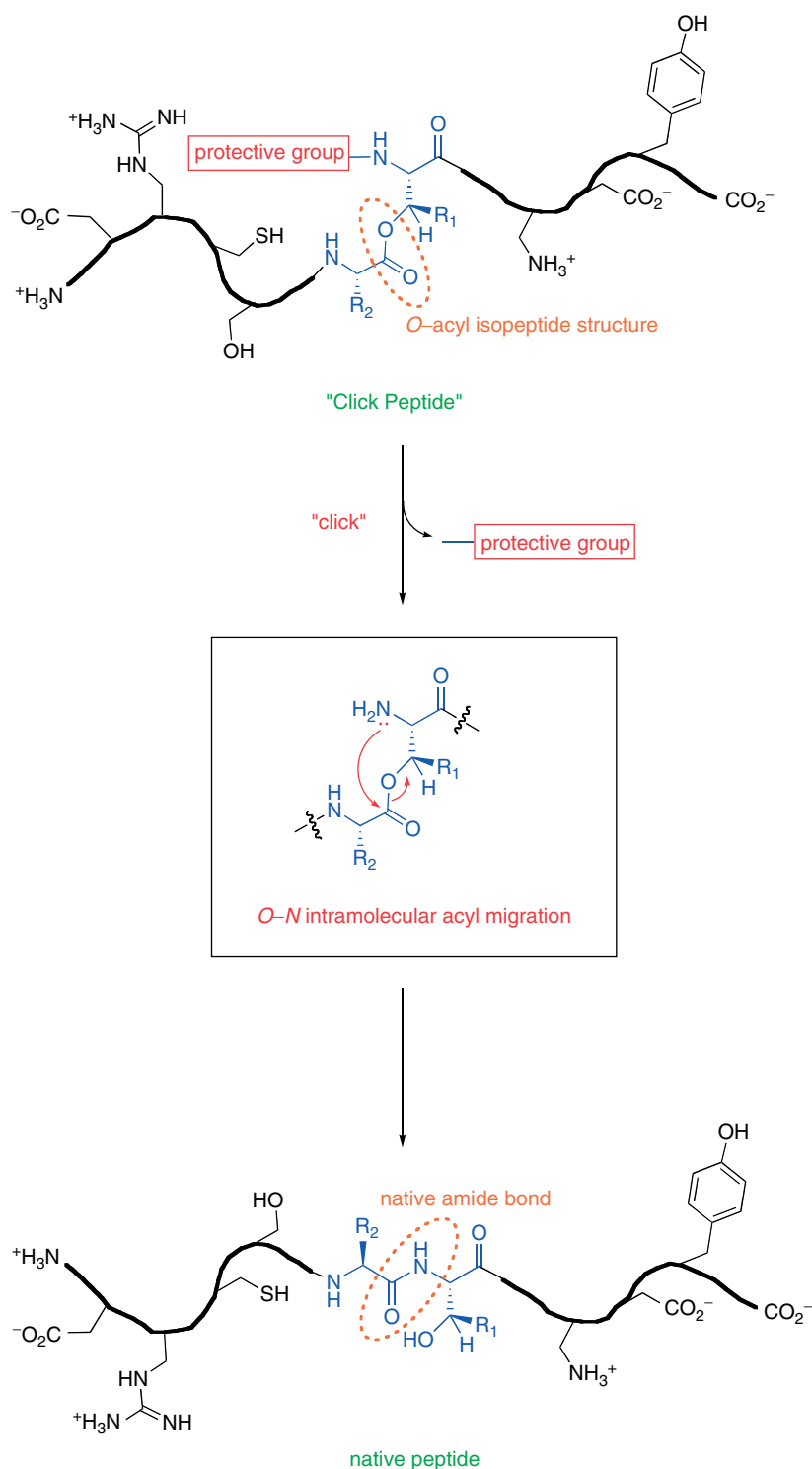


Figure 1 'Click Peptide' based on the 'O-acyl isopeptide method'.

esterification reaction on the resin could be omitted, allows the application of the 'O-acyl isopeptide method' to fully automated protocols for the synthesis of long peptides or proteins (Figure 3) [8]. Additionally, very recently, we developed a novel 'racemization-free segment condensation' based on the 'O-acyl isopeptide method' [10].

The Application of O-Acyl Isopeptide Method

Moreover, we have successfully applied the 'O-acyl isopeptide method' to the chemical biology-oriented synthesis of the Alzheimer's disease (AD)-related amyloid β peptide ($A\beta$) 1–42 analogues, leading to the development of pH- or photo-triggered 'Click Peptide'

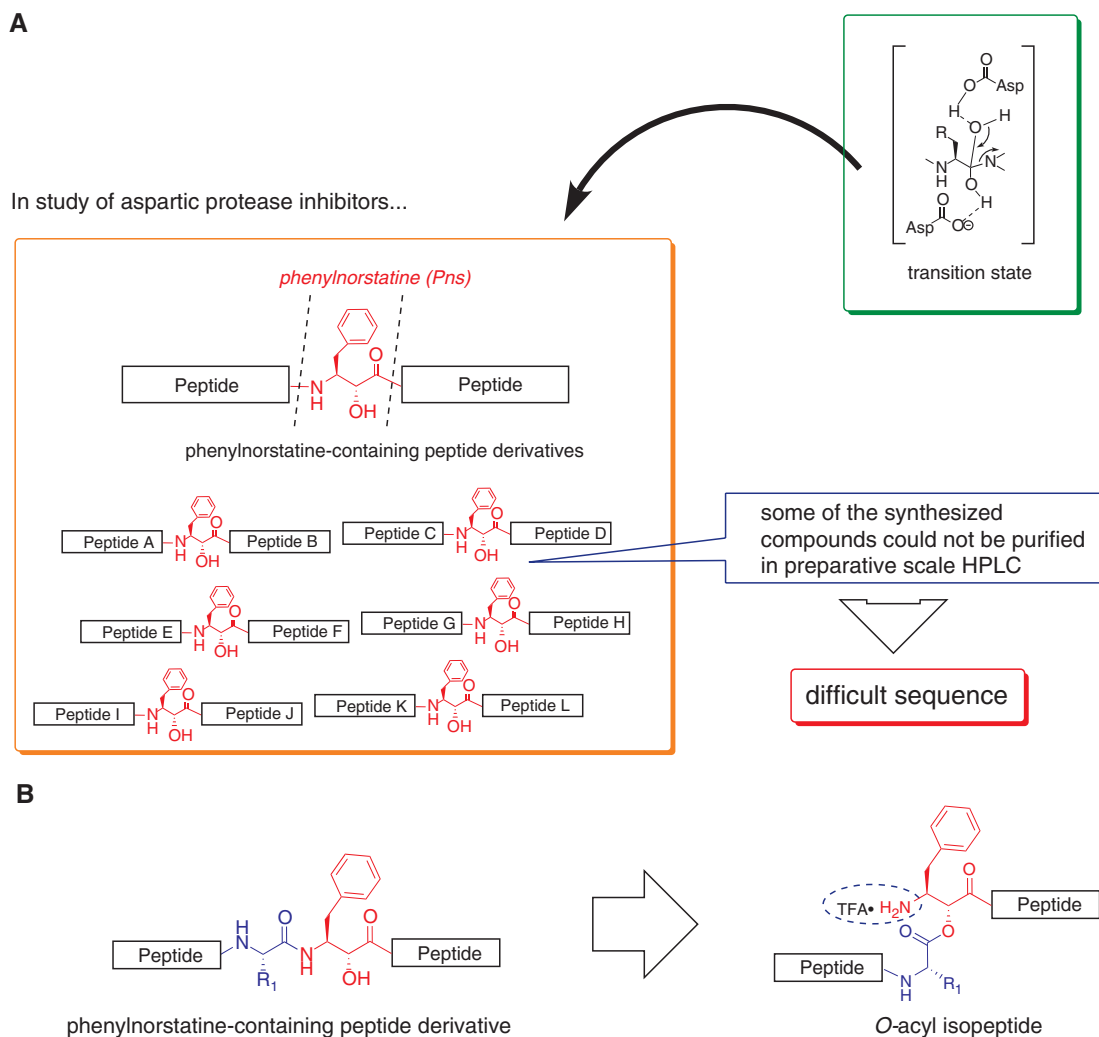


Figure 2 (A) 'Difficult sequence' in our studies of aspartic protease inhibitors, (B) design of *O*-acyl isopeptide derived from phenylnorstatine-containing peptide derivative.

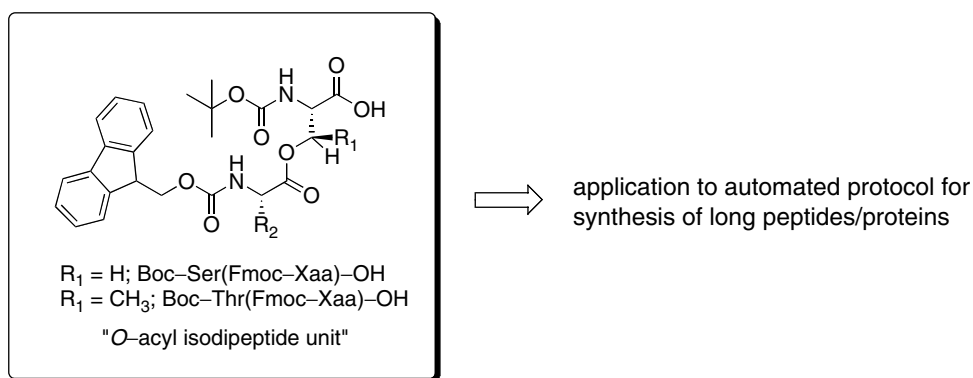


Figure 3 General structure of *O*-acyl isodipeptide unit: application of the '*O*-acyl isopeptide method' to fully automated protocol.

(Figure 1) [3–7,9]. The 'Click Peptide' did not exhibit the self-assembling nature under physiological conditions because of one single ester, and could migrate to the original $A\beta_{1-42}$ with a quick and easy one-way conversion reaction (so-called 'click') via the *O*-*N*

intramolecular acyl migration. A clear understanding of the pathological mechanism of $A\beta_{1-42}$, a currently unexplained process, would be of great significance in the discovery of novel drug targets against AD [25–28]. Currently, the difficulties in handling $A\beta_{1-42}$, because

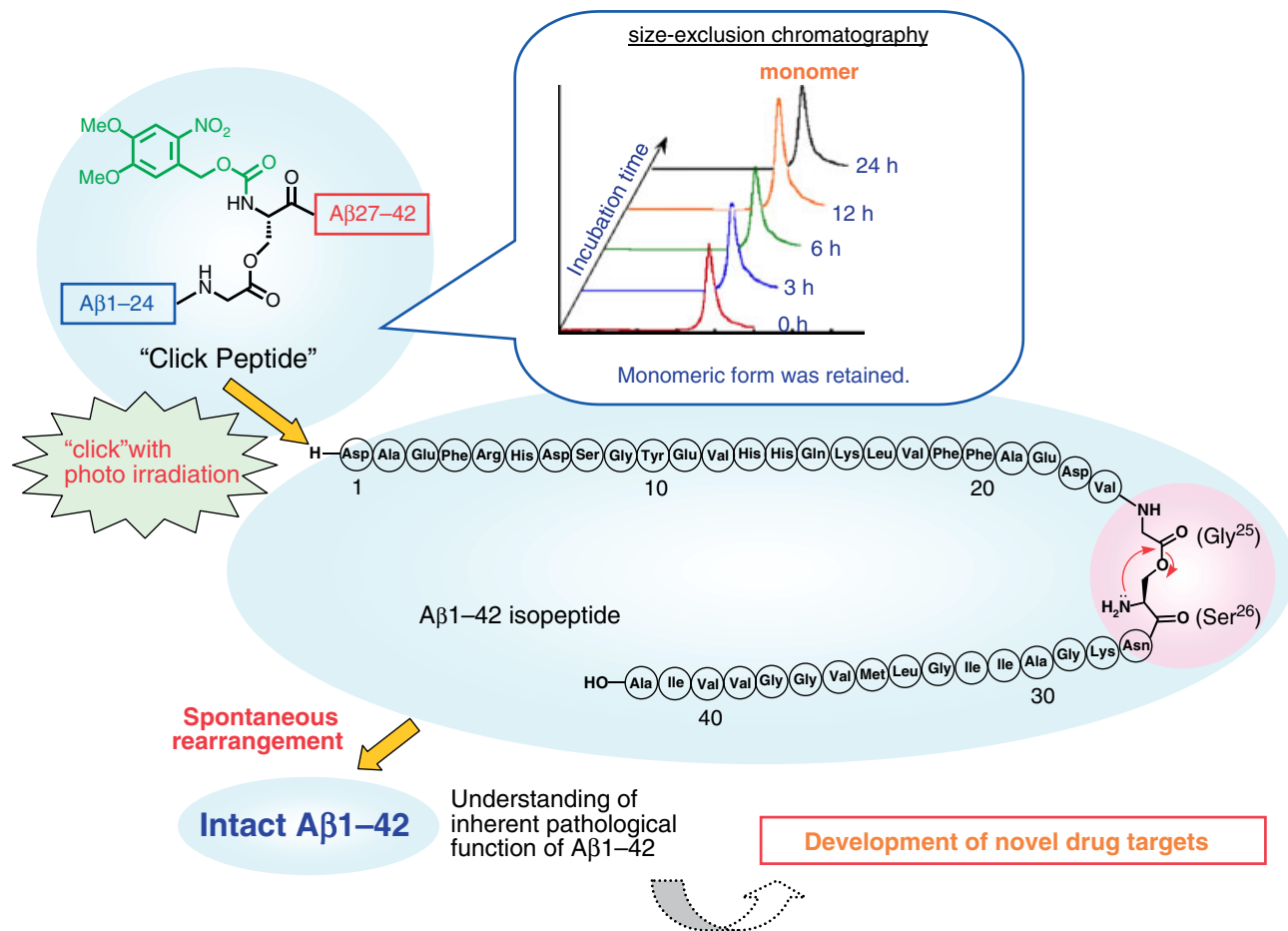


Figure 4 Photo-triggered click peptide. The production of A β 1–42 by photo-triggered click followed by *O*–*N* intramolecular acyl migration reaction of A β 1–42 isopeptide.

of its highly aggregative nature, hamper the progress of A β 1–42-related AD research [29–32]. The ‘Click Peptide’ method would open the doors for investigation of the biological functions of A β 1–42 in AD by inducible activation of A β 1–42 self-assembly (Figure 4).

Interestingly, shortly after we disclosed the ‘*O*-acyl isopeptide method’ [1,2], several other groups also confirmed the efficacy of this method. Carpino *et al.* synthesized the Jung–Redemann 26-residue peptide efficiently by utilizing the ‘*O*-acyl isopeptide method’, whereas this peptide could not be synthesized by standard SPPS [12]. By carefully evaluating the appropriate protecting group, stability of the ester bond during assembly, and occurrence of side reactions, they concluded that the efficacy of the ‘*O*-acyl isopeptide method’ was comparable to that of the pseudoproline method [15]. Börner *et al.* also synthesized the *O*-acyl isopeptide for efficient preparation of poly(ethylene oxide)–peptide conjugates [14]. Moreover, Mutter *et al.* confirmed by circular dichroism (CD)-based analyses that the secondary structure of *O*-acyl isopeptide structure is significantly different from that of the corresponding *N*-acyl native peptides [11,13],

which agrees with our hypothesis. These reports indicate that the ‘*O*-acyl isopeptide method’ is widely advantageous for peptide preparation by disrupting the unfavorable secondary structures of the native peptides.

CONCLUSION

Classical *O*–*N* intramolecular acyl migration was revived by our group as a powerful key reaction in the field of modern medicinal chemistry in the development of water-soluble prodrugs. After more than a decade of prodrug studies, we recently disclosed the ‘*O*-acyl isopeptide method’ as a novel synthetic method in the field of peptide chemistry and its application to chemical biology-oriented synthesis of A β analogues, leading to the development of ‘Click Peptide’ (Figure 5). We hope that the strategy using the ‘*O*-acyl isopeptide method’, in which a simple isomerization to an *O*-acyl isopeptide remarkably and temporarily changes the physicochemical properties of the native peptide and an *O*–*N* intramolecular acyl migration triggers the native amide bond formation under physiological conditions,

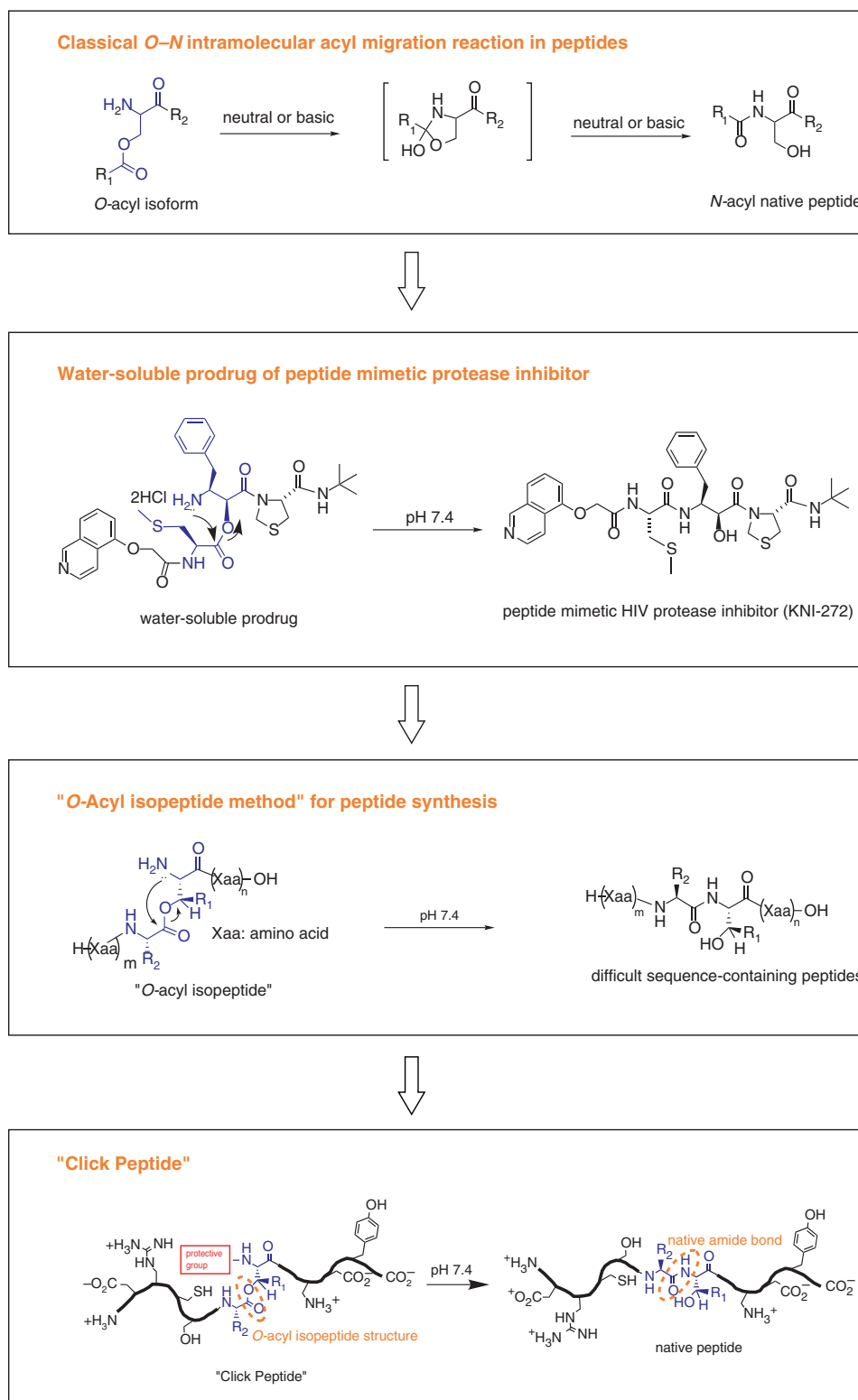


Figure 5 Our workflow on peptide science based on the 'O-N intramolecular acyl migration reaction'.

will further contribute to the study of peptides and proteins. Examples of such studies include the studies of membrane peptides/proteins that are difficult to handle in various conditions because of their high self-assembling characters.

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