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A straightforward method for automated Fmoc-based synthesis of bio-inspired peptide crypto-thioesters†

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Despite recent advances, the direct Fmoc-based solid phase synthesis of peptide α -thioesters for the convergent synthesis of proteins *via* native chemical ligation (NCL) remains a challenge in the field. We herein report a simple and general methodology, enabling access to peptide thioester surrogates. A novel C-terminal *N*-(2-hydroxybenzyl)cysteine thioesterification device based on an amide-to-thioester rearrangement was developed, and the resulting peptide crypto-thioesters can be directly used in NCL reactions with fast *N* \rightarrow *S* shift kinetics at neutral pH. These fast kinetics arise from our bio-inspired design, *via* intein-like intramolecular catalysis. Due to a well-positioned phenol moiety, an impressive >50 fold increase in the kinetic rate is observed compared to an *O*-methylated derivative. Importantly, the synthesis of this new device can be fully automated using inexpensive commercially available materials and does not require any post-synthetic steps prior to NCL. We successfully applied this new method to the synthesis of two long naturally-occurring cysteine-rich peptide sequences.

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Introduction

Peptide α -thioesters are key intermediates for the convergent synthesis of proteins through native chemical ligation (NCL), a reaction that has revolutionized the field.¹ While peptide thioesters can be directly synthesized through Boc-based solid-phase peptide synthesis (SPPS), their access *via* the more widely used Fmoc-based strategy is not straightforward due to the instability of the thioester moiety to repeated piperidine treatments used for Fmoc deprotection. Most Fmoc-based methodologies developed to date rely on a post-SPPS conversion of a piperidine-stable precursor into a thioester, but despite considerable efforts, no universal strategy has emerged.² Thanks to their simple implementation, methods such as Dawson's *N*-acylurea (Nbz)³ and Liu's hydrazides⁴ for the generation of thioester precursors have recently become popular, but are still associated with several limitations.⁵ Thus, a reliable and straightforward route to peptide α -thioesters *via* Fmoc-SPPS is highly desirable.

A number of strategies based on β - or γ -mercapto amide thioesterification devices have recently appeared that exploit an amide-to-thioester rearrangement^{6,7} (Fig. 1A), which is reminiscent of the first step of intein-promoted *in vivo* protein splicing.⁸ In most cases, the *N* \rightarrow *S* acyl shift is reversible and

requires acidic conditions to proceed efficiently, meaning that the peptides have to be first converted into thioesters prior to NCL, which is optimal at pH \sim 7. A few systems, referred to as crypto-thioesters, are able to operate under NCL conditions, thus enabling one-pot reactions (thioester formation and NCL). Cysteine-prolyl esters (CPEs) developed by Aimoto^{7d} (Fig. 1B-b) rely on an elegant intramolecular *O* \rightarrow *N* shift to displace the amide-thioester equilibrium under NCL conditions, but with slow ligation kinetics. Although simple to implement, the α -methyl cysteine crypto-thioesters recently introduced by Offer^{7j} (Fig. 1B-e) are impaired by similarly slow kinetics. In both cases, the limiting factor is likely the disfavored *trans*-to-*cis* isomerization of the amide bond to generate the geometrically feasible intramolecular attack of the thiolate. Use of *N,N*-disubstituted β -mercapto amides can overpass this limitation by disfavoring the unproductive *trans* conformer. The symmetrical bis-sulfanylethylamides (SEAs, Fig. 1B-c) introduced by Melnyk^{7e} and Liu^{7h} can be used in NCL at pH 6–7 with relatively fast kinetics, but work better at lower pH. Otaka's *N*-sulfanylethylamides^{7e} (Sealides, Fig. 1B-d) exploit the electronic effect of the anilide to enhance the electrophilicity of the amide bond. They are efficient NCL substrates at neutral pH only when using a phosphate-based buffer, probably arising from pseudo-intramolecular acid–base catalysis, but the exact mechanism remains elusive.⁹ Interestingly, while we were preparing this paper, a method appeared based on an *N*-ethyl cysteine device featuring a free carboxylic acid; peptides equipped with this device can act as crypto-thioesters. The α -acid group was shown to accelerate the reaction compared to an α -amide-containing

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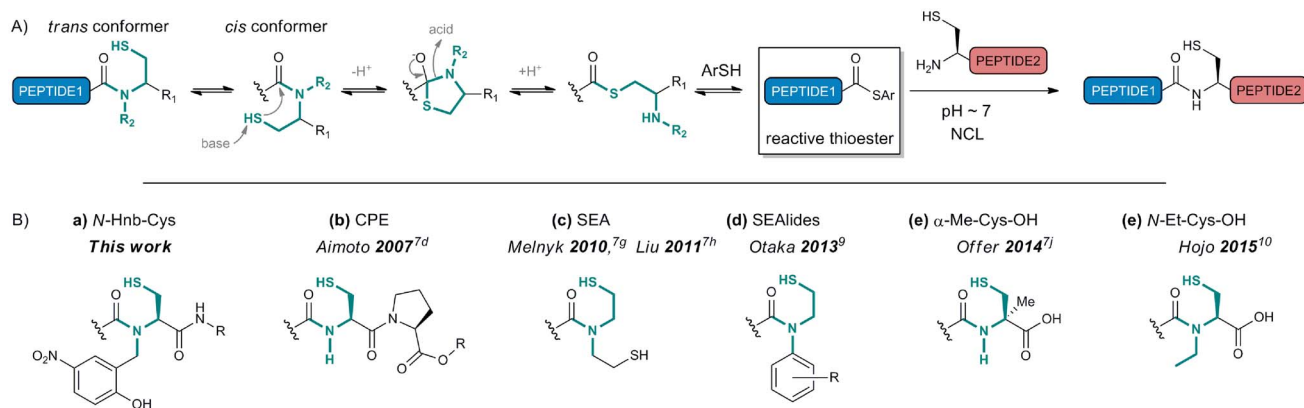


Fig. 1 $N \rightarrow S$ shift-based *in situ* synthesis of peptide thioesters for NCL. (A) Mechanism. (B) Crypto-thioesters bearing thioesterification devices operating under NCL conditions.

analogue, with optimum ligation kinetics at pH 5.5, but with very slow kinetics at pH 7.¹⁰

We herein report the design and optimization of a simple and general methodology based on an N -(2-hydroxybenzyl) cysteine device, which can be automatically assembled in the solid-phase using inexpensive commercially available materials. Peptides bearing this device show fast $N \rightarrow S$ shift kinetics under NCL conditions around neutral pH, likely arising from a biomimetic intein-like intramolecular catalysis mechanism. The synthesis of these novel crypto-thioesters is straightforward and does not require an additional step prior to NCL. We believe that this accessible and robust Fmoc-based thioesterification device provides a significant advance to chemical protein synthesis.

Results and discussion

We based our design on N -alkyl cysteine^{7c} as a scaffold for the design of our thioesterification device, as it can be readily assembled in the solid-phase from inexpensive and commercially available materials.¹¹ However, its $N \rightarrow S$ shift is usually very slow at neutral pH,¹² preventing its direct use in NCL. We focused our efforts on addressing this challenge by taking inspiration from the mechanism of *in vivo* protein splicing promoted by class 1 inteins. The first step of this process consists of a self-catalyzed $N \rightarrow S$ acyl shift of an amine into a thioester,⁸ where a highly conserved histidine residue plays a pivotal role. This N -protonated His is believed to catalyze C–N bond scission through polarization and then protonation of the nitrogen leaving group (Fig. 2A).¹³ We reasoned that the N -alkyl group of cysteine could be functionalized by an appropriate intramolecular proton donor able to mimic the catalytic histidine side chain, resulting in a much faster $N \rightarrow S$ shift at neutral pH. Accordingly, we designed an N -(2-hydroxybenzyl) cysteine device (Fig. 2B) leading to a hypothetical thermodynamically favored six-membered transition state for intramolecular N -protonation. The key to this design is a phenol group with a pK_a close to neutral, as this would result in a low energy cost for the intramolecular proton transfer at pH 7.

We expected an additional advantage from the 2-hydroxybenzyl group: acceleration of N -acylation *via* O -acylation followed by the intramolecular $O \rightarrow N$ shift (Fig. 2C), as in the N -acylation of 2-hydroxy-5-methoxybenzyl (Hmb) protected amines.¹⁴ This could overcome a common drawback encountered with most $N \rightarrow S$ acyl shift-based thioesterification devices developed so far, the difficult N -acylation of

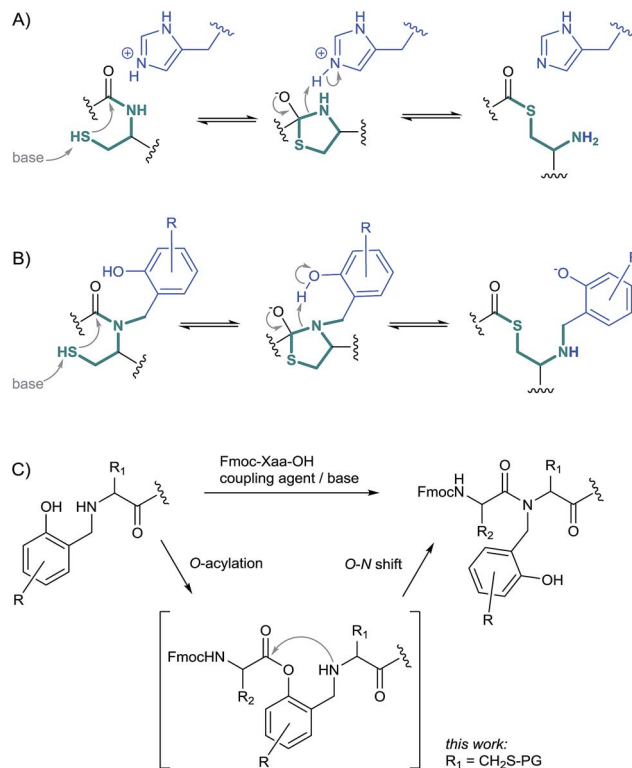


Fig. 2 Rational design of N -(2-hydroxybenzyl)cysteine thioesterification devices. (A) Mechanism of the intein self-catalyzed $N \rightarrow S$ shift in water. (B) Bio-inspired putative mechanism for the self-catalysis of the $N \rightarrow S$ shift in N -acyl- N -(2-hydroxybenzyl)cysteine peptides in water. (C) Mechanism of the intramolecular $O \rightarrow N$ shift in organic solvents shared by N -Hmb peptides and the new N -(2-hydroxybenzyl) cysteine thioesterification devices. PG: protecting group.

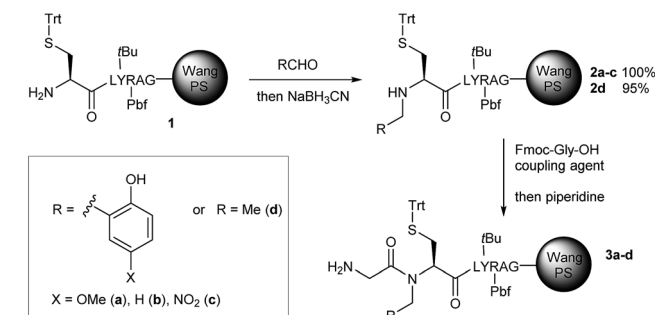


a secondary amine,^{7g,h,15} to introduce the first residue of the peptide sequence.

Optimization of the hydroxybenzyl group

Three different *N*-2-hydroxybenzyl groups were introduced on a solid supported cysteine derivative through an automated reductive amination protocol (compounds **2a–c**) and were compared with a reference *N*-ethyl cysteine device (**2d**). For this preliminary study, a reporter peptide (LYRAG-NH₂) was introduced C-terminal to cysteine to facilitate the analysis using LC-MS after TFA-mediated deprotection and cleavage from the resin. Excellent results were obtained with the 2-hydroxy-5-nitrobenzyl (Hnb) group,¹⁶ giving quantitative *N*-acylation yields for a Gly residue under mild amide coupling conditions (HBTU/HOBt) (Table 1, entry 5), while the ethyl-substituted compound (Table 1, entry 7) was unreactive. Hmb or unsubstituted 2-hydroxybenzyl groups (Table 1, entries 1 and 3) showed only moderate assistance. These results are in line with the literature:¹⁶ Alewood and collaborators already observed the much superior acyl transfer efficiencies of 2,6- and 2,5-Hnb compared to Hmb. Moreover, the p*K*_a of the hydroxyl group of Hnb is close to 7 (~6.5, see ESI, Fig. S11†), in line with our requirements discussed above for efficient biomimetic catalysis. We accordingly selected *N*-Hnb-Cys as our preferred scaffold.

Table 1 Screening of different *N*-(2-hydroxybenzyl) groups for their ability to enhance the *N*-acylation yield of a solid-supported *S*-trityl-protected *N*-substituted cysteine. PS: polystyrene



Entry	Resin	Coupling agent ^a	Product	Yield ^b [%]
1	2a	HBTU/HOBt	3a	<5 ^c
2	2a	HATU	3a	45 ^c
3	2b	HBTU/HOBt	3b	11
4	2b	HATU	3b	12
5	2c	HBTU/HOBt	3c	>99
6	2c	HATU	3c	>99
7	2d	HBTU/HOBt	3d	<1
8	2d	HATU	3d	12
9	2d	PyBrop	3d	1
10	2d	HATU/DMAP ^d	3d	13

^a All reactions were conducted for 2 h, in DMF, using 10 equiv. Fmoc-Xaa-OH, 9.5 equiv. coupling agent and 20 equiv. iPr₂NEt. ^b Yields determined after TFA-mediated cleavage of the peptide resin using HPLC analysis of the amine-to-amide conversion. ^c Estimated yields due to closely eluting HPLC peaks. ^d 0.1 equiv. DMAP.

Optimization of the thiol protecting group of cysteine and linkage to the resin

When exploring the scope of peptides bearing our *N*-Hnb-Cys thioesterification device, we observed an instability upon purification and storage due to a premature spontaneous *N* → *S* shift. Fortunately, the protection of the cysteine thiol as an *S*-*S*tBu disulfide, stable to Fmoc-SPPS and readily removed under NCL conditions,¹⁷ resolved this problem (see ESI, pS24–S29†).¹⁸ As the use of a reporter peptide was no longer needed at this point, we looked at introducing Cys(*S*tBu) as either the α -acid or the amide, using a Wang-type or a Rink's amide linker, respectively. Disappointingly, in the former case we were confronted with Cys epimerization¹⁹ and β -elimination, followed by the addition of piperidine.²⁰ In the latter case, we observed a TFA-catalyzed hydrolysis²¹ reaction, leading to a mixture of amide-(7) and acid-terminated (8) compounds (see ESI pS27–S29† for details). Simple incorporation of a Gly spacer between the Rink's linker and the device solved these problems. Preliminary evaluation of NCL reactions with the model cysteinyl peptide **6** showed that this additional residue (compound **9**) did not affect the NCL kinetics compared to cysteine α -amide **7**, the reaction being even slightly faster than for the α -acid **8** (ESI, Fig. S57†). As expected, cysteine deprotection was very fast, showing the complete removal of the *S*tBu group within minutes.

Study of the *N*-acylation of the device

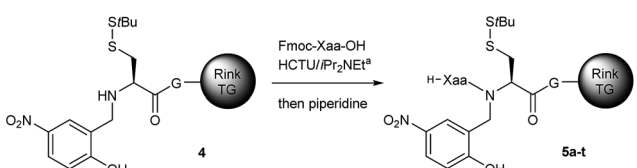
N-Acylation of the optimized device was then examined for the twenty proteogenic amino acids, using a standard SPPS protocol (Table 2). In most cases the coupling yields were good to excellent. No significant epimerization was observed, except for Fmoc-Cys(Trt) (7%), which is well known for its propensity for racemization.²² Six residues showed modest *N*-acylation yields (<50%). In these difficult cases, we looked at further optimizing the coupling, avoiding the use of sensitive or costly reagents, and keeping in mind an easy-to-automate procedure. Three successive couplings of Fmoc-Ser(*t*Bu) resulted in an excellent yield (85%, entry 16). The five-fold coupling of the more demanding Fmoc-Val led to a good 65% yield (entry 21). Alternatively, single coupling under microwave heating at 70 °C led to an excellent 92% yield for Fmoc-Val (entry 22) and 78% for Fmoc-Ile (entry 24). Importantly, the unreacted secondary amine could be quantitatively capped by acetylation, giving a byproduct that can be readily removed during the peptide precipitation step following TFA-based cleavage from the resin.²³

Mechanistic insights

Next, we wanted to assess the validity of our intein-inspired design hypothesis that should result in the self-catalysis of the *N* → *S* shift at neutral pH through intramolecular protonation. Here, we synthesized a model peptide equipped with a 2-methoxy-5-nitrobenzylcysteine device (**12**), designed to inhibit the acid–base properties of the phenol group while not being expected to lead to major changes in terms of steric hindrance



Table 2 Acylation of the resin **4** with the 20 different protected pro-teogenic amino acids under automated coupling conditions. TG: Tentagel



Entry	Xaa	Product	Nb of couplings ^b	Yield ^c [%]
1	Gly	5a	1	97
2	Asp(OtBu)	5b	1	94
3	Ala	5c	1	90
4	Glu(OtBu)	5d	1	89 ^d
5	Cys(Trt)	5e	1	86 ^e
6	Met	5f	1	85
7	Asn(Trt)	5g	1	80 ^f
8	His(Trt)	5h	1	77
9	Phe	5i	1	77
10	Gln(Trt)	5j	1	72
11	Leu	5k	1	71
12	Trp(Boc)	5l	1	69
13	Arg(Pbf)	5m	1	67
14	Tyr(tBu)	5n	1	58
15	Ser(tBu)	5o	1	49
16	Ser(tBu)	5o	3	85
17	Lys(Boc)	5p	1	45
18	Thr(tBu)	5q	1	19
19	Pro	5r	1	19
20	Val	5s	1	16
21	Val	5s	5	65
22	Val	5s	1	92 ^g
23	Ile	5t	1	12
24	Ile	5t	1	78 ^g

^a Reactions were conducted at a 0.025 mmol scale using a Prelude synthesizer at RT for 30 min in NMP/DMF 5 : 1, using 10 equiv. Fmoc-Xaa-OH, 9.5 equiv. HCTU and 20 equiv. *i*Pr₂NEt. ^b When indicated, the coupling step was repeated under the same conditions. ^c Yields determined after TFA-mediated cleavage of the peptidyl resin and subsequent HPLC analysis of the amine-to-amide conversion. ^d Concomitant formation of pyroglutamate during TFA cleavage. ^e 93 : 7 mixture of L-Cys/D-Cys. ^f Incomplete deprotection of the Trt group. ^g Coupling at 70 °C.

compared to Hnb (peptide **9**). We were delighted to see that the hydroxyl group very efficiently catalyzes the *N* → *S* acyl shift: >50-fold rate enhancement of NCL was measured for **9** compared to **12** under the NCL conditions (Fig. 3), supporting our hypothesis.²⁴

Moreover, the ligation of **9** at different pH values showed the *N* → *S* acyl shift/NCL process to be fastest at pH ~ 6.5, and slightly slower at lower pH (ESI, Fig. S67†). This fits with the p*K*_a of the Hnb group (6.5), suggesting the need for a subtle balance between the phenol group protonation and the thiol group deprotonation (cysteine p*K*_a ~ 8.5) in influencing the *N* → *S* shift. To eliminate a possible *N* → *O* shift mechanism, the thiol was protected with a stable acetamidomethyl (Acm) group (**13**). This NCL-unreactive compound confirms the crucial role of the SH group and provides further data to suggest an intramolecular *N* → *S* shift mechanism. Critically, this latter finding

also highlights the potential of the *N*-Hnb-Cys thioester precursors for successive ligations in the *N*-terminus-to-*C*-terminus (*N*-to-*C*) direction. We are currently evaluating latent crypto-thioesters that could be activated by deprotection of a NCL-stable cysteine protecting group, as has been exploited for related *N*-methyl Cys,^{11,25} CPE,²⁶ SEA¹⁷ and Sealide²⁷ thioester precursors.

Study of the NCL with model peptides

Next, we turned our attention to the effectiveness of our new thioesterification device in NCL reactions for a range of thioesters bearing different amino acids C-terminal to the *N*-Hnb-Cys device (Fig. 3). Note that we used relatively dilute conditions for the peptide reactants (~1 mM) and low concentrations of the 4-mercaptophenylacetic acid catalyst (MPAA, 25 mM).²⁸ Relatively fast NCL kinetics were observed using Ala (peptide **9**) and Ser (**10**) crypto-thioesters, as the reactions were complete after 24 h at 37 °C. For comparison with the NCL kinetics data available in the literature, we determined the apparent second order kinetic constants, with results in the 0.03–0.06 M⁻¹ s⁻¹ range, in line with the reported values for standard NCL with preformed thioesters.²⁹ These data validate the rapid *N* → *S* shift promoted by our device. Ligation at valine (crypto-thioester **11**) was 10 times slower, as we had anticipated from the steric hindrance of its side chain.³⁰ However, increasing the MPAA concentration to 300 mM, lowering the pH close to the optimum (6.6) and heating to 50 °C led to kinetics comparable to Ala and Ser crypto-thioesters. The latter result suggests that the method could be applied in routine NCL even for peptide sequences known to be kinetically demanding.

Potential epimerization during the *N* → *S*-shift/NCL process was rigorously examined by synthesizing a D-Ala-containing product (**15**) as an HPLC standard. Only trace epimerization (<0.4%) was detected in the crude NCL mixture starting from the L-Ala crypto-thioester **9**. We also examined the hydrolysis of the crypto-thioesters into the corresponding acids. All reactions were clean and did not show significant amounts of hydrolysis (<4% for **9** after 24 h and using excess crypto-thioester).

Application of the methodology to two cysteine-rich peptides

Importantly, our new method was applied to the synthesis of two long naturally-occurring cysteine-rich peptide sequences, MT7 and Cg-BigDef1. Muscarinic toxin 7 (MT7), 65 residues, was isolated from green mamba venom.³¹ MT7 naturally forms a three-finger-fold stabilized by four disulfide bonds. Big defensin 1 (Cg-BigDef1) is a 93-residue host-defense peptide isolated from Japanese oysters and is composed of a hydrophobic N-terminal domain and a β-defensin-like cationic C-terminal domain containing six cysteine residues involved in three disulfide bridges.³²

These two peptide sequences were chosen as challenging examples to demonstrate the applicability of our methodology, as their syntheses *via* the NCL of two segments require long thioesters, 41 and 56 amino acids, respectively. In addition, the N-terminal Cg-BigDef1 domain is a “difficult sequence”,³³ *i.e.* hard to synthesize using Fmoc-SPPS.³⁴ Both of the crypto-



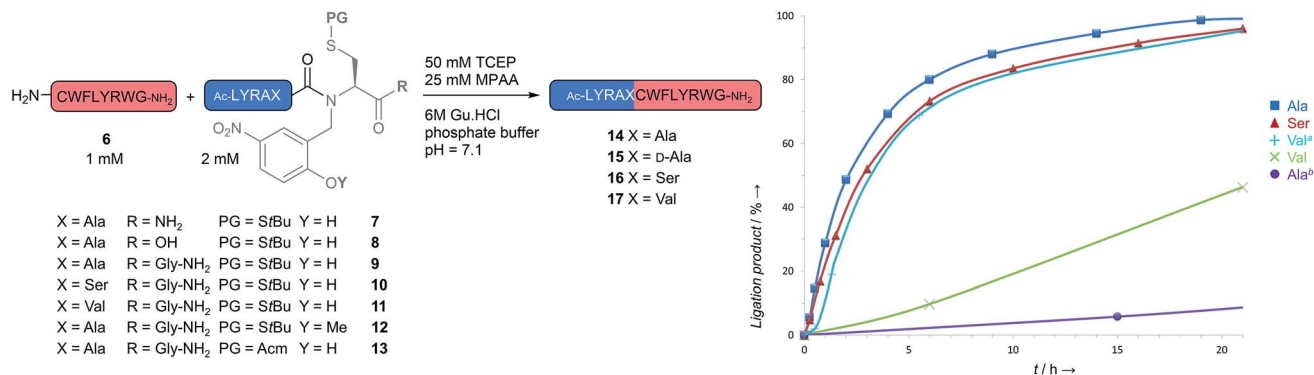


Fig. 3 NCL of the model crypto-thioester peptides 7–11 bearing an *N*-Hnb-cysteine device, and an analog (12) with the hydroxyl group masked as a methyl ether. ^a 300 mM MPAA, pH 6.6, 50 °C. ^b Hnb methyl ether (12).

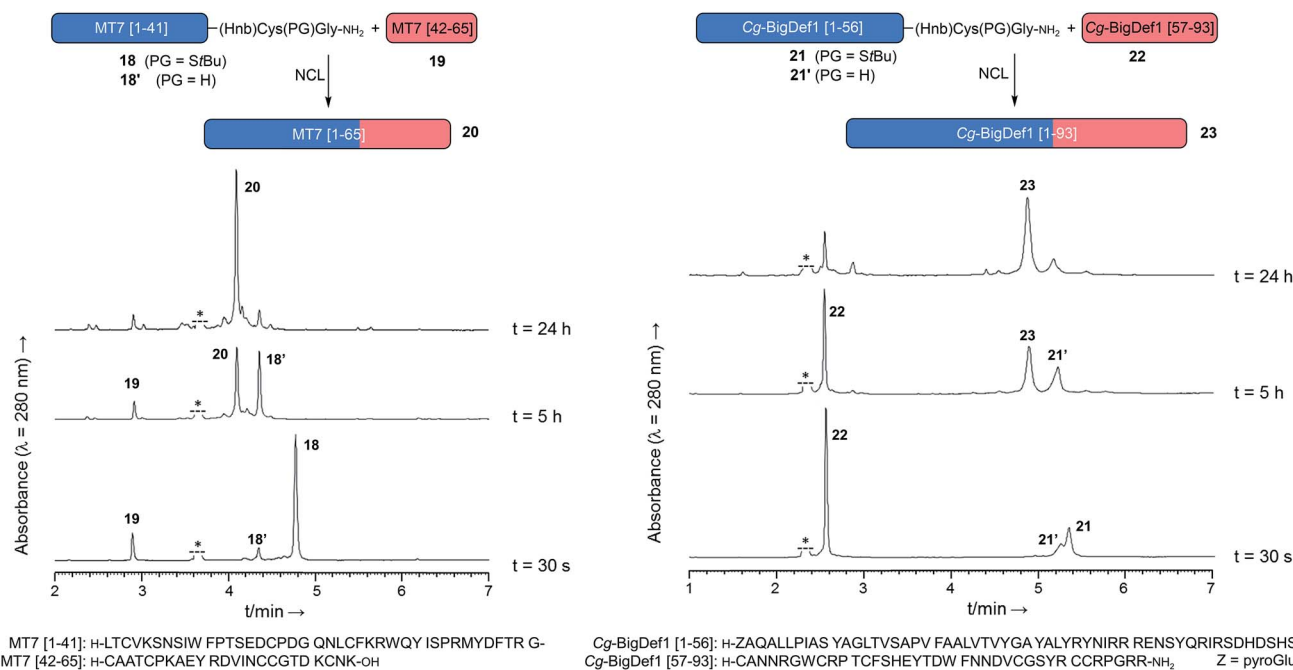


Fig. 4 Application of the *N*-(2-hydroxy-5-nitrobenzyl)cysteine thioesterification device to the NCL-based syntheses of two long and demanding peptide sequences.

thioesters (**18** and **21**) were synthesized and purified in good yields (13% and 7%, see ESI, pS86 and S93[†]). The two ligations were complete overnight at 1 and 2 mM concentrations, respectively, and showed clean HPLC profiles (Fig. 4). Finally, pure reduced forms of MT7 (**20**) and Cg-BigDef1 (**23**) were isolated after HPLC purification in 14% and 18% yields, respectively³⁵ (ESI, Fig. S85 and S93[†]), demonstrating the utility of the *N*-Hnb-Cys method.

Conclusions

In conclusion, we introduced a novel C-terminal thioesterification device, *N*-Hnb-Cys(*S*t*Bu*), that can rapidly rearrange into a thioester at neutral pH, allowing its use in NCL even under demanding conditions. Fast kinetics likely result from an

internal catalysis mechanism provided by a well-positioned phenol moiety arising from a bio-inspired intein-like design. Synthesis from inexpensive commercially available materials and further peptide elongation are straightforward, and can be fully automated on a peptide synthesizer without requiring a post-synthetic step, thus constituting a major advance in the field. Applicability of the strategy was demonstrated by the synthesis of two long naturally occurring cysteine-rich peptide sequences. Further work is ongoing in our laboratory for its application to other synthetically demanding targets.

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- Note that a simple adjustment of the initial amount of resin allows the SPPS of the crypto-thioester sequence to be conducted at the desired scale.
- The hydroxyl group could assist the reaction *via* a different mechanism to the one we postulated, such as stabilization of the oxyanion moiety of the tetrahedral intermediate. Note that the activation of the amide by hydrogen bonding between the phenol group and the carbonyl oxygen seems unlikely: NMR analysis of a small model peptide showed no significant differences in the *cis/trans* ratio between the



- hydroxy and methoxy compounds (see ESI pS14–S23†). A hydrogen bond would be expected to stabilize the *cis* form and thus modify the ratio. DFT studies are planned to help better understand the mechanism.
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- 34 Standard Fmoc-SPPS of Cg-BigDef [1–42] C-terminal α -amide gave extremely poor results and low elongation yields, probably arising from its difficult hydrophobic sequence. Considering these results, we optimized the synthesis of the cryptothioester **21** by incorporating multiple coupling steps.
- 35 These isolated yields can appear rather low considering the excellent conversion yield, as determined from the HPLC chromatograms of the crude NCL mixtures, see Fig. 4. We attribute this loss of material to the non-optimized workup and HPLC purification protocols.

