Peptide Ligation

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Traceless Ligation of Cysteine Peptides Using Selective Deselenization**

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The synthesis of proteins with a fully native sequence is an ongoing challenge in protein chemistry. Native chemical ligation (NCL) approaches have proven to be generally applicable where cysteine (Cys) residues are appropriately positioned, [1,2] however, the synthesis of many proteins often require ligation at non-Cys sites in the polypeptide sequence. [3-7] Previously, we introduced a reductive strategy for ligation at Ala sites^[7] based on global desulfurization of Cys^[8] that has found widespread utility for the synthesis of complex proteins by NCL. [9,10] Selective desulfurization can be affected by both Rainey Ni and Pd/C/H₂^[7] and, more recently, by the radical initiator VA-044 in combination with the water soluble phosphine TCEP (tris(2-carboxyethyl)phosphine).[11] However, since these conditions result in global desulfurization of all thiols in the protein, the method requires protection and deprotection of all other Cys residues in the native sequence. [6a,11,12] These additional steps complicate the synthesis of larger polypeptides^[13] and limit the use of natural Cys residues for ligation.

Selenocysteine (Sec, U) has been shown to expand the NCL method to Xaa-Sec site, allowing the synthesis of selenoproteins.^[14-16] Additionally, the resulting selenopeptides can be deselenized under similar conditions to that used for Cys containing peptides to yield the corresponding Ala peptide sequences.^[11,17] We reasoned that the high propensity of selenols to form radicals^[18] could be harnessed for selective

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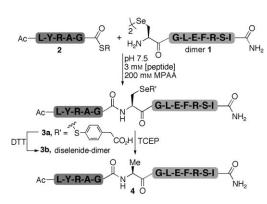
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Scheme 1. Native chemical ligation of N-terminal Sec-peptide **1** with C-terminal thioester-peptide **2** gives the ligated product, **3 a.** Following purification, **3 a** was deselenized to the alanyl peptide **4**, using excess TCEP. R = 3-mercaptopropionyl-Leu. MPAA = 4-mercaptophenylacetic acid; DTT = dithiothreitol; TCEP = tris(2-carboxyethyl) phosphine.

reduction of selenols in the presence of thiols (Scheme 1), thus avoiding the need for protection/deprotection steps. This approach was inspired by our observation that synthetic analogs of glutaredoxin 3 (Grx3) containing Sec were incompatible with reduction by the water soluble reducing agent TCEP,^[19] leading to the generation of significant levels of a deselenized side products. By contrast, the wt-Grx3 was found to be stable to TCEP. Indeed, the sensitivity of Sec in peptides and proteins to reduction by TCEP has been previously noted in the development of selenocysteine ligation methods and in the context of selenoproteins.^[15,16a,20] Importantly, TCEP^[21] is widely used to reduce disulfides to thiols in peptides and proteins without reduction of the C–S bond.^[22-24]

Accordingly, we ligated the N-terminal Sec-peptide 1 (UGLEFRSI-amide, isolated in the form of a diselenide dimer) to the thioester peptide 2 (Ac-LYRAG-SR) (Scheme 1 and Figure S1 in the Supporting Information), to produce 3a. [14-16] The ligation conditions of 6M GdmCl, 200 mm Na₂HPO₄, pH 7.5, saturated with 200 mm 4-mercaptophenylacetic acid (MPAA)^[25] gave the best results. The aromatic thiol acted as both a catalyst to activate the alkyl thioester and as a mild reducing agent to generate a small pool of free selenol to facilitate the ligation reaction. In addition, under these conditions the product was spontaneously converted to selenylsulfide 3a, with MPAA, which was beneficial for the HPLC purification. [26]

The purified selenylsulfide peptide **3a** was treated with 50-fold excess TCEP at pH 5.5 to produce the deselenized alanyl-peptide **4** (Scheme 1, Figure S2 and S3). Under these conditions, the reaction proceeded to completion yet was

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slower than anticipated (17 h). We reasoned that the presence of the aromatic thiol, MPAA (1 equiv), could interfere with the deselenization reaction^[27,28] by acting as a radical scavenger, thus competing with the deselenization of the Sec residue.^[27-29] To remove the aromatic thiol, compound **3a** was reduced with 50-fold excess DTT (dithiothreitol) to produce the free selenol, which subsequently reoxidized to give the dimeric diselenide peptide, **3b**. This reaction sequence was accomplished in one pot due to the high reactivity of the selenol and low redox potential of diselenides.^[19] Deselenization of the diselenide dimer **3b** with excess TCEP proceeded rapidly, generating the Ala peptide **4** in less than 4 h, a time comparable to previously reported deselenization reactions.

TCEP is a common reagent in protein chemistry that is selective for disulfide reduction. [21-24] The chemoselectivity of the deselenization conditions was further evaluated with peptide **5a**, H-Ala-**Sec**-Gly-Ser-**Cys**-Lys-Trp-Thr-Met-Ala-NH₂ which contains the potentially sensitive amino acids Cys, Met and Trp. Peptide **5a** oxidized under ambient conditions to generate the cyclic selenylsulfide peptide **5b** (Scheme 2). Treatment of **5b** with only TCEP (40-fold excess)

Scheme 2. Selective deselenization of peptide **5 b**. The deselenized peptide **6** was the major product with only traces of side product, **7**. $PB = NaH_2PO_4$.

at pH 5.5 resulted in rapid deselenization of the Sec to yield peptide **6**. Surprisingly, analysis of the minor side product identified peptide **7**, H-Ala-Ala-Gly-Ser-Ala-Lys-Trp-Thr-Met-Ala-NH₂, consistent with a slow additional desulfurization of the Cys residue. [30,31] However, when the reaction was run (200 mm NaH₂PO₄, pH 5.1) with 9.4 equivalents DTT to reduce the selenylsulfide, addition of just 2.2 equivalents TCEP led exclusively to deselenization. Monoreduced deselenized product **6**, H-Ala-Ala-Gly-Ser-Cys-Lys-Trp-Thr-Met-Ala-NH₂, was obtained cleanly in 2 h with only traces of the deselenized–desulfurized peptide **7** detected (Scheme 2, Figure 1 and Figure S4).

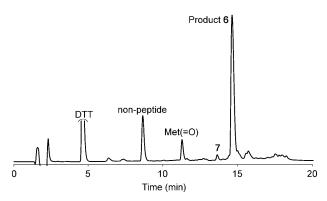


Figure 1. HPLC chromatogram for the deselenization reaction mixture of peptide 5 b, which was carried out with 9.4 equivalents DTT for 1 h, and 2.2 equivalents TCEP for 2 h to give major product, 6, with traces of the deselenized—desulfurized product, 7. Met(=0) corresponds to deselenized product with oxidized Met present in the starting material.

We propose that this deselenization of Sec to Ala occurs through a radical-mediated mechanism (Scheme 3), similar to the desulfurization of thiols by trialkyl phosphines (and phosphites) under elevated temperatures or UV light proposed first by Walling et al^[27] and recently by Wan and Danishefsky (in the presence of a radical initiator such as VA-044).^[11,32]

Scheme 3. Proposed deselenization mechanism of Sec-containing peptides with TCEP in aqueous solution.

In principle, loss of selenium could occur through an elimination mechanism to generate dehydroalanine.[33] To rule out this possibility, we carried out two experiments. First, the diselenide 1 was treated with either hydrogen peroxide^[34] or TCEP (Scheme 4). Under hydrogen peroxide the selenium atom was converted to the corresponding selenoxide with subsequent β -syn elimination to produce the modified peptide 8 where the Sec has been converted to dehydroalanine (Scheme 4, bottom).^[34] Peptide 8 was rapidly hydrolyzed under the reaction conditions to the corresponding pyruvoyl peptide 9a, which was observed together with its hydrate, 9b (Figure S6B). In contrast, when diselenide dimer 1 was reacted with TCEP under the deselenization conditions, the corresponding Ala peptide, 10, was obtained without any traces of either 9a or 9b (Scheme 4 top, Figure S6C).[33] Second, we carried out the deselenization reaction of peptide

Scheme 4. Reaction of dimer peptide 1 with H_2O_2 produced the dehydroalanine analog, 8, which was rapidly hydrolyzed under the reaction conditions to the corresponding pyruvoyl peptide 9a (in equilibrium with its hydrate 9b). In contrast, reaction of peptide 1 with TCEP gave only the deselenized product, 10, with an N-terminal Ala.

5b (Scheme 2) under the regular conditions using D_2O instead of H_2O . The reaction mixture was worked up and analyzed in H_2O to reverse any deuterium incorporation at acidic sites. The resulting deselenization product was found to be consistent with [**D**]6, having a mass of M+1 in comparison to the product from H_2O , 6 (Scheme 5, Figure S5). Furthermore, the chromatographically distinct minor side-product (additional desulfurization) was found to be consistent with [**D**₂]7 having a mass of M+2 in comparison with the above described **7**. Any mechanism involving dehydroalanine formation would lead to an additional deuterium being incorporated at the α-carbon (Scheme 5, Figure S5).^[31]

Epimerization is always a concern in the development of new methods. While the desulfurization of Cys residues by Raney Ni has been experimentally shown to be racemization-free, [7,35] the radical-mediated mechanism could result in epimerization. [11] The proposed deselenization mechanism proceeds through a radical on the β -carbon, which is expected to be the major intermediate. [11,32] However, radical migration from β - to the α -carbon could lead to undesired epimerization at the Sec residue. Furthermore, it has been experimentally confirmed that thiyl radicals can abstract hydrogen from α -carbons in peptides and proteins (as well as acidic protons in lipids and DNA). [36,37] To test for epimerization, the pentapeptide H-Phe-Lys-Sec-Ser-Asp-NH₂ (FKUSD.amide, isolated in the form of a diselenide dimer, **11**) was synthesized

Scheme 5. Deselenization of peptide 5 b in phosphate buffered D₂O by TCEP forms two products; the deselenized monodeuterated product, [D]6 as the major product and the doubly deuterated deselenized—desulfurized minor product, [D₂]7.

together with H-Phe-Lys-(l)Ala-Ser-Asp-NH₂ (FK(l)ASD. amide, **12**) and H-Phe-Lys-(d)Ala-Ser-Asp-NH₂ (FK(d) ASD.amide, **13**, which by HPLC elutes at 8.8 min, 1 min earlier than the l-enantiomer). Deselenization of **11** with TCEP was complete within 2 h, producing only the l-enantiomer product FK(l)ASD, which co-eluted with the synthetic peptide, and has an identical mass. (Scheme 6, Figure S7 and S8).

Scheme 6. The diselenide dimer of FKUSD, **11**, was deselenized with 10 equivalents TCEP in 200 mm phosphate buffer, pH 5.1 to give only FK(I)ASD product, **12**, no traces of the of **13** were observed.

Taken together, these results support the formation of a radical intermediate at the β -carbon of Sec, without migration to the α -carbon, indicating a chemoselective process with complete retention of stereoselectivity.

Finally, to evaluate selective deselenization of a complex polypeptide following ligation, we prepared the 38-mer peptide Grx3(1–38)(Cys11Sec–Cys14Sec–Ala38Cys) containing two Sec and one Cys residues by NCL (Scheme S1), together with the all-Cys control, Grx3(1–38)(Ala38Cys). Following removal of thiophenol by HPLC, deselenization of Grx3(1–38)(Cys11Sec–Cys14Sec–Ala38Cys) by TCEP gave a doubly deselenized product Grx3(1–38)(Cys11Ala–Cys14Ala–Ala38Cys) as the major product with a minor Cys reduction side-product (Figure S9 to S12). [31] In contrast

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the Cys-containing analog Grx3(1–38)(Ala38Cys) was fully stable to the treatment with TCEP.

In conclusion, we have developed a straightforward method for the selective reduction of selenocysteine to alanine in the presence of unprotected cysteine. The deselenization reaction by TCEP is highly chemo- and enantioselective. In effect, this strategy will allow ligation at both Ala and Cys sites in the same protein sequence and maintain native Cys residues without the need for side chain protection. This general approach has the potential to be extended to other amino acids with the appropriate β - or γ - selenol group. $^{[7,9]}$ Furthermore, despite the widespread use of TCEP in protein chemistry, we recommend its use with caution, particularly in the production of selenoproteins. $^{[38]}$

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