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New isocysteine building blocks and chemoselective peptide ligation

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Boc-, Fmoc- and Cbz-protected isocysteine building blocks were prepared by a concise three-step procedure starting from thiomalic acid. The use of Boc/Trt-protected isocysteine provided convenient access to isocysteinyl peptides that allow the chemoselective ligation of unprotected peptide fragments in water. The pH-dependency of the isocysteinemediated ligation was compared with that of cysteine-mediated native chemical ligation.

Introduction

The replacement of naturally occurring amino acids by isosteric or artificial amino acids is commonly employed for probing the interaction of a protein with its ligand. Thiolcontaining building blocks are of interest due to their particular reactivity which mediates inhibitory interactions with protein metallo centers, conformational restriction through disulfide bond formation and site-selective tagging reactions among other effects. The β-mercapto amino acid cysteine is by far the most frequently used thiol-containing building block. Its 1,2 aminothiol structure also makes it an invaluable tool for the efficient and convenient joining of two unprotected peptide fragments.**1–5**

The introduction of α -mercapto acids provides the opportunity to alter thiol acidity and/or thiolate basicity which can allow the fine-tuning of metal–ligand interactions. For example, some inhibitors of metalloproteases such as angiotensin-converting enzymes contain α-mercapto acyl peptides.**6,7** The β-amino-αmercapto amino acid isocysteine is a useful cysteine analogue and has been shown to increase the inhibitory activity of peptidic stromelysin binders when incorporated as a replacement of cysteine.**⁸** Isocysteine also might serve as a replacement of β**3** -H cysteine in β-peptides which opens the possibility of altering thiol reactivity without changing the backbone length.**⁹** There is, however, only one report that describes the preparation of an isocysteine derivative that suits the needs of solidphase peptide synthesis.**⁸** We here present convenient syntheses of Boc-, Fmoc- and Cbz-protected isocysteine building blocks. Furthermore we show the utility of isocysteine to support the ligation of unprotected peptide segments by a native chemical ligation-like reaction. To illustrate the effects of alterations of the thiol acidity the pH-dependency of the isocysteine ligation was compared with that of cysteine-mediated native chemical ligation.

Results and discussion

Hanglows' synthesis of isocysteine **7** and also our route to N,Sprotected building blocks **8**–**11** was based on commercially available DL-thiomalic acid 1 as described by Gustavson.^{8,10,11} According to this, simultaneous protection of both the α-mercapto and the α-carboxylic group was achieved upon reaction with cyclopentanone in the presence of *p*-toluenesulfonic acid (Scheme 1). The β-carboxyl group in the formed oxathiolanone **2** remained unaffected and was subsequently converted to the acyl azide. In contrast to Hanglows' Fmoc/ Trt-Isocys synthesis, in which the isocyanate **3** formed after the Curtius rearrangement was hydrolysed to obtain isocysteine in

Scheme 1 a) cyclopentanone, *p*-TsOH, PhCH**3**, reflux, 68%; b) 1. diphenylphoshoryl azide, TEA, PhCH₃; 2. 85 °C; c) PhCH₂OH, 85 °C, 86% (3→4); 9-fluorenylmethanol, 85 °C, 87% (3→5); *t*BuOH, 75 °C, 77% (**3→6**); d) 1. 0.2 M LiOH, THF, 0 °C; 2. Trt-Br, 50% (**5→7**); 1. 0.2 M LiOH, THF, 0 °C; 2. PMB-Cl, 80% (**5→8**); e) 1. 1 M LiOH, THF, RT; 2. PMB-Cl, 96%; f) 1. 1 M LiOH, THF; 2. Trt-Br, 78% (**6 10**); 1. 1 M LiOH, THF, RT. 2. PMB-Cl, 77% (**6 11**).

unprotected form, it was preferred to fashion a more direct establishing of Fmoc-, Cbz- and Boc-protecting groups. The isocyanate **3** was allowed to react with 9-fluorenylmethanol, benzyl alcohol or *tert*-butanol, which completed a three-step sequence and furnished the fully protected isocysteine derivatives **4**, **5** and **6** in 77–87% yield. We next sought a method that would liberate both the carboxyl group and the α-mercapto group without detriment to the acid- and base-sensitive protecting groups in **4**–**6**. The opening of the cyclic ketal was initiated by ester hydrolysis with aqueous LiOH in THF, which proved straightforward with Cbz- and Boc-protected isocysteines **4** and

6. The presence of the base-sensitive Fmoc-group in **5** rendered this step challenging. It was found that careful addition of exactly one equivalent of lithium hydroxide in THF, however, enabled a clean and quantitative removal of the ketal group. The liberated α -mercapto groups were subsequently masked by omitting any attempts to isolate the *N*-protected isocysteine derivatives. Instead, it was preferred to directly add *p*-methoxybenzyl chloride or trityl bromide to the basic THF solution obtained after ester hydrolysis. This convenient one-pot procedure afforded the known Fmoc/Trt-protected isocysteine **7** and the novel Fmoc/PMB-, Cbz/PMB-, Boc/Trt- and Boc/ PMB-protected building blocks **8**–**11** in 50–96% yield. It is noted that the same strategy would allow for the synthesis of enantiomerically pure isocysteine if D- or L-thiomalic acid were used.**¹²**

Having established expedient access to various protected isocysteine derivatives we next set out to explore their utility in the solid-phase synthesis of isocysteine containing peptides. Previous work had demonstrated the usefulness of the Fmoc/ Trt-protected building block **7**. **8** We envisaged investigating the use of isocysteine as a ligation handle (*vide infra*). Our particular interest, therefore, concerned the synthesis of peptides such as **16** bearing isocysteine as N-terminal amino acid. We reckoned that Cbz/PMB- and Boc/PMB-protected isocysteine building blocks **9** and **11** would enable the isocysteinyl-peptide **16** to be assembled by applying the Boc-strategy. For the final cleavage trifluoromethanesulfonic acid was employed as a nontoxic alternative to HF. The crude peptide obtained after cleavage, however, failed to precipitate upon addition of ether which hampered its purification. After repeated unsuccessful attempts with various scavengers, we chose to couple Boc/Trt-protected isocysteine **10** to resin-bound peptide **14** which was assembled by using Fmoc/*t*Bu-protected amino acids (Scheme 2). It was expected that the application of the Fmoc-strategy and TFA as cleaving agent would be beneficial. TFA in contrast to TFMSA (trifluoromethanesulfonic acid) is volatile and allows vacuum concentration thereby facilitating ether precipitation. Indeed, in this case purification was straightforward providing peptide **16** after HPLC purification in 69% overall yield.

Isocys-Arg-Ala-Glu-Tyr-Ser

16

Scheme 2 a) 10 eq. Fmoc-Ser(*t*Bu)-OH, 5 eq. DIC, 0.1 eq. DMAP, DMF; b) 1. DMF/piperidine (4 : 1), 2. 4 eq. Fmoc-Aa-OH, 4 eq. HOBt, 3.6 eq. HBTU, 8 eq. DIPEA; 3. pyridine/Ac**2**O (5 : 1); c) 1. DMF/ piperidine (4 : 1), 2. 4 eq. Boc-Isocys(Trt)-OH, 4 eq. HOBt, 3.6 eq. HBTU, 8 eq. DIPEA; 3. pyridine/Ac**2**O (5 : 1); d) CF**3**COOH/PhSMe/ Et**3**SiH/H**2**O (92 : 3 : 3 : 2), 69%.

Terminally modified peptides such as bromoacetamido-,**¹³** maleinimido-,**¹⁴** hydroxylamino- **¹⁵** and azidopeptides **16,17** among others¹⁸ are invaluable tools that can allow chemoselective ligation of two unprotected peptide fragments to be performed in aqueous solution. The 1,2-aminothiol structure of N-terminal cysteine is one of the most suitable entities for achieving such segment couplings demonstrated best by the numerous protein syntheses that have been enabled by the native chemical ligation technology.**2,19** This reaction (Scheme 3) is initiated by a reversible thiol-exchange reaction of cysteine conjugate **17** with thioester **18**. **20** The formed thioester intermediate **19** is subject to a spontaneous S–N-acyl shift, which establishes the α-peptide bond in product **20**. In isocysteine, the 1,2-aminothiol structure is retained. Accordingly, isocysteine is expected to support fragment ligations by a native chemical ligation-like reaction mechanism.**²¹** The S–N-acyl shift in thioester-intermediate **22** would lead to the formation of a β-peptide bond in **23**. The isocysteine-mediated native chemical ligation would hence offer prospects for a convergent assembly of peptides with β-amino acid architecture.**9,22,23**

> Native-Chemical-Ligation Isocysteine-Chemical-Ligation

To assess the feasibility of isocysteine-mediated native chemical ligation we explored model peptides and ligation conditions as described by Kent and co-workers.**3,24** First, peptide mercaptopropionamide thioester **24**, which was obtained through solid-phase synthesis on a mercaptopropionic acid loaded Rink-resin, was converted to the more reactive mercaptobenzylester **25**. Peptide thioester **25** was allowed to react with the isocysteine-peptide **16**. Reactions were carried out in argon saturated 0.1 M phosphate buffer and in the presence of 4% benzylmercaptan which helped to prevent thioester hydrolysis and to maintain a reducing environment. At pH 7 and 1 mM concentration of reactants the isocysteine-mediated native chemical ligation proceeded smoothly. HPLC/ESI-MS analysis showed a new peak that appeared at 16 min with an *m*/*z* that corresponded to ligation product **26** (Fig. 1). The ligation product formed lacked susceptibility to attempted hydrolysis by NaOH-treatment which provided support for the formation of a non-hydrolysable peptide bond. It can be concluded that isocysteine supports a native chemical ligation-like reaction. It is interesting to note that the $S \rightarrow N$ acyl transfer likely to be involved in the isocysteine ligation proceeds *via* main chain expansion as opposed to $S \rightarrow N$ acyl transfer involved in cysteine and auxiliary mediated ligations **25–29** that are characterised by main chain contraction. This feature might be of less importance in convergent peptide assembly, but we presume that it could affect ligation reactions that proceed under geometrical constraints such as peptide- **30,31** and DNAtemplate controlled reactions.**32,33**

Fig. 1 Analytical HPLC trace of the ligation reaction at $\lambda = 280$ nm $(16 + 25)$ and pH 7.

It is worthwhile noting that both the isocysteine containing starting material **16** and the ligation product **26** exist as a mixture of diastereomers due to the use of racemic isocysteine. Nevertheless, under the selected conditions the HPLC-traces showed single albeit broadened peaks and it was therefore not possible to explore whether one of the diastereomers conferred higher ligation rates than the other.

Fig. 2 shows the time-dependent formation of ligation product **26** at pH 7 and pH 6. It became evident that the ligation at pH 6 is slower than the ligation at pH 7. For example, product **26** was formed in 63% yield after 240 min as opposed to 44% yield obtained at pH 6.

Fig. 2 Isocysteine-mediated chemical ligation at pH 7 and pH 6.

It has been shown that the cysteine-mediated native chemical ligation proceeds rapidly around pH 7 but is rendered less efficient at pH < 5.5.**³⁴** To explore whether the isocysteine-mediated native chemical ligation would exhibit a similar pH-dependence we compared the initial rate of formation of isocysteinepeptide **26** and cysteine-containing peptide **28** as a function of pH at 1 mM concentrations of reactants and at identical reaction conditions (Fig. 3). RP-HPLC-analysis revealed that the initial rate of cysteine native chemical ligation was reduced by a factor of 0.25 when lowering the pH from pH 7 to pH 6. This pH decrease reduced the initial rate of isocysteine native chemical ligation only by a factor of 0.56. Further pH decreases to pH 5 led to a diminution of initial rates such that the formation of ligation products almost ceased. At pH 7, the ligation of cysteine-peptide **28** was 3-times faster than ligation of the isocysteine-peptide **26** while at pH < 6, the isocysteine native chemical ligation was as efficient as the cysteine native chemical ligation.

Fig. 3 pH-Profile of initial ligation rates of the cysteine ligation (**25** Cys-Arg-Ala-Glu-Tyr-Ser, 27 → Leu-Tyr-Lys-Ala-Gly-Cys-Arg-Ala-Glu-Tyr-Ser, 28) and the isocysteine ligation $(16 + 25 \rightarrow 26)$. Ligation conditions as specified in Fig. 1.

It became apparent that the isocysteine-mediated ligation is less pH-dependent than the cysteine-mediated native chemical ligation. We assume that the increased acidity of the thiol group is the decisive factor that leads to this attenuation of the pH-dependence. This assumption is based on the notion that native chemical ligation proceeds through the thiolate form of the cysteine-peptide and that the thiol-exchange reaction is rate-limiting.³ In isocysteine, the α-thiol group should be more acidic than the β-thiol group of cysteine due to the inductive effect of the carboxyl group. This influence becomes manifest in the pK_a -value of thioglycolic amide ($pK_a = 8.2$) which is considerably lower than that of β-mercaptopropionic amide $(pK_a = 9.4)$.³⁵ The calculated pK_a -values of isocysteine amide $(pK_a = 8.2)$ and cysteine amide $(pK_a = 9.1)$ suggest a similar difference of thiol acidity.**³⁶** The more acidic the thiol group, the more thiolate is provided even under acidic conditions and hence the faster the ligation reaction. On the other hand, inductive and steric effects are expected to decrease the nucleophilicity of the α-thiolate in isocysteine when compared with the β-thiolate of cysteine which might explain why at $pH > 6$ the cysteine native chemical ligation proved faster than the isocysteine native chemical ligation.

Conclusion

In conclusion, we have provided convenient synthetic access to various isocysteine building blocks for solution and solid-phase peptide synthesis. The replacement of cysteine by isocysteine allows for an alteration of thiol reactivity as evidenced by the attenuated pH-dependence of isocysteine-mediated native

chemical ligation of unprotected peptide fragments. Isocysteine being a β-amino acid, also might be incorporated in β-peptides, structurally well-defined peptide analogues that show interesting biological activities. It is, for example, conceivable that the replacement of $β³$ -H-cysteine by isocysteine allows the thiol reactivity to be altered while maintaining the backbone length and the ability to confer chemoselective fragment condensations. In this regard, the ability of isocysteine to support native chemical ligation reactions might be of advantage for the convergent assembly of modified β-peptides.

Experimental

General materials and methods

Reagents and chemicals were obtained from Acros, Aldrich, Biosolve, Fluka, Novabiochem or Senn Chemicals and were used without further purification. All solvents were freshly distilled. Proton and carbon NMR spectra were recorded on a Varian Mercury 400 or a Bruker DRX-500 spectrometer at room temperature. The signals of the residual protonated solvent (CDCl₃ or [D₆]DMSO) were used as reference signals. Coupling constants *J* are reported in Hz. Flash chromatography was performed using Merck silica gel 60. TLC was performed with aluminium-backed silica gel 60 F**254** plates (Merck). Melting points were measured using a Büchi B-540 apparatus in open capillary tubes and are not corrected.

Reversed phase HPLC

Analytical reversed phase HPLC was performed with an Agilent 1100 series system and a DAD-detector recording at 280 nm. A Macherey & Nagel Nucleosil (100-5) type CC 250/4 Nautilus RP-18 column was used at room temperature. Semipreparative reversed phase HPLC was performed with a Gilson Nebula Series with automated fraction collection and Gilson 156 UV-detector. A Macherey & Nagel SP 125/10 Nucleodur Gravity RP-18 column was used at 55 \degree C. Gradients of solvent A $(98.9\% \text{ water}/1\% \text{ acetonitrile}/0.1\% \text{ TFA})$ and B $(98.9\% \text{ m} \text{C})$ acetonitrile/1% water/0.1% TFA) were adjusted according to the separation needs.

Mass spectrometry

Matrix assisted laser desorption ionisation analysis was carried out with a Voyager-DE Pro BioSpectrometer™ from PerSeptive Biosystems using a 2,5-dihydroxybenzoic acid (DHB) matrix. ESI-MS was carried out using a Agilent 1100 series system with a Macherey & Nagel Nucleosil (100-5) type CC 250/4 Nautilus RP-18 column and a Finnigan Thermoquest LCQ. FAB-MS was recorded on a JEOL JMS-SX102A machine, applied matrices were given for each compound.

Solid-phase peptide synthesis

Reactions were carried out at room temperature and performed manually using 5 ml polyethylene syringe reactors which were equipped with a fritted disc. All peptides were synthesised using HBTU/HOBt chemistry. Typically 4 eq. of the corresponding amino acid were allowed to react for 2–3 min with 3.6 eq. of HBTU, 4 eq. of HOBt and 8 eq. of DIPEA in DMF/CH₂Cl₂ (1 : 1). The solution was added to the resin and agitated for 2 h at room temperature. Cysteine was coupled using DIC/HOBt chemistry in order to avoid extensive racemisation. Typically 4 eq. of the amino acid were reacted with 4 eq. of HOBt and 4 eq. of DIC in DMF for 20 min and then added to the resin. Unreacted amino groups were capped by a 15 min treatment of pyridine/ $Ac₂O$ (5 : 1) after every coupling step. Fmoc cleavage was achieved using two times a solution of 20% piperidine in DMF for 10 min. The Boc group was cleaved using two times pure TFA for 5 min. The resins were washed after every step

with CH₂Cl₂ (5×), DMF (5×) and CH₂Cl₂ (5×) to avoid any side reactions.

(2-Oxo-1-oxa-4-thia-spiro[4.4]non-3-yl)-acetic acid (2)

To a solution of 2-mercaptosuccinic acid **1** (5.0 g, 33.3 mmol) in toluene (70 ml) were added cyclopentanone (3.54 ml, 40.2 mmol) and *p*-toluenesulfonic acid (76 mg, 0.40 mmol). The reaction vessel was equipped with a Dean–Stark trap and was refluxed for 8 h. The solution was concentrated under reduced pressure, and the residue was dissolved in sat. Na $HCO₃(30 \text{ ml})$. The aqueous solution was washed with CH₂Cl₂ (2 \times 20 ml), adjusted to pH 2 with concentrated HCl and extracted with $CH₂Cl₂$ (3 × 30 ml). The organic phase was dried (MgSO₄) and evaporated under reduced pressure. The residue was purified by silica gel flash chromatography to furnish 4.90 g (68%) of **2** as a white solid. $Mp = 93$ °C. $R_f = 0.21$ (EtOAc : cyclohexane) = 2 : 5 1% AcOH). **¹** H-NMR (CDCl**3**): δ 1.75 (m, 4H); 1.94 (m, 2H); 2.24 (m, 2H); 2.84 (dd, 1H, *J* = 9.16, *J* = 17.69); 3.21 (dd, 1H, *J* = 4.02, *J* = 17.57); 4.38 (dd, 1H, *J* = 4.14, *J* = 9.16); 11.43 (s, 1H). **¹³**C-NMR (CDCl**3**): 24.89 (CH**2**); 25.17 (CH**2**); 39.15 (CH**2**); 42.63 (CH**2**); 43.94 (CH**2**); 44.93 (CH); 97.20 (Cq); 174.83 (Cq); 177.59 (Cq). HRMS [FAB, *m*-NBA, pos.]: calcd. 216.0456; found 216.0468.

(2-Oxo-1-oxa-4-thia-spiro[4.4]non-3-ylmethyl)-carbamic acid benzyl ester (4)

A solution of **2** (1.02 g, 4.73 mmol) and triethylamine (728 µl, 5.20 mmol) in toluene (20 ml) was stirred 30 min at room temperature. The reaction mixture was cooled to 0° C and diphenylphosphoryl azide (1.12 ml, 5.10 mmol) was added dropwise. After stirring for 2 h at room temperature, the solution was heated to 85 °C until the evolution of N_2 ceased. The reaction mixture was cooled to room temperature and EtOAc (30 ml) was added. The organic phase was washed with sat. NaHCO₃ (20 ml), dist. H₂O (20 ml), dried (MgSO₄) and concentrated under reduced pressure. The residue was dissolved in toluene (20 ml) and benzyl alcohol (460 μ l, 4.45 mmol) was added. The solution was kept at 80° C for 14 h and then allowed to cool to room temperature. The solvent was evaporated under reduced pressure and the residue was purified by silica gel chromatography to afford 1.32 g (87%) of **4** as an yellow oil. R_f $= 0.24$ (EtOAc : cyclohexane $= 4 : 1$). ¹H-NMR (CDCl₃): δ 1.75 (m, 4H); 1.94 (m, 2H); 2.24 (m, 2H); 3.65 (m, 2H); 4.19 (dd, 1H, *J* = 5.65); 5.07 (s, 2H); 7.31 (m, 5H). **¹³**C-NMR (CDCl**3**): 23.58 (CH**2**); 23.84 (CH**2**); 41.34 (CH**2**); 42.53 (2 × CH**2**); 48.29 (CH**2**); 67.12 (CH**2**); 95.85 (Cq); 128.24–128.73 (4 × CH); 136.33 (Cq); 156.42 (Cq); 173.43 (Cq). HRMS [FAB, *m*-NBA, pos.]: calcd. 321.1035; found 321.1020.

(2-Oxo-1-oxa-4-thia-spiro[4.4]non-3-ylmethyl)-carbamic acid 9*H***-fluoren-9-ylmethyl ester (5)**

A solution of $2(200 \text{ mg}, 0.92 \text{ mmol})$ and triethylamine $(142 \mu l,$ 1.01 mmol) in toluene (10 ml) was stirred 30 min at room temperature. The reaction was cooled to 0° C and diphenylphosphoryl azide (198 µl, 0.92 mmol) was added dropwise. After stirring over 2 h at room temperature, the solution was heated up to 85 \degree C until the evolution of N₂ ceased. The reaction was cooled to room temperature and EtOAc (20 ml) was added. The organic phase was washed with sat. NaHCO₃ (15 ml), dist. H**2**O (15 ml), dried (MgSO**4**) and concentrated under reduced pressure. The dried residue was dissolved in toluene (10 ml) and 9-fluorenylmethanol (157 mg, 0.8 mmol) was added. The solution was kept at 80 $^{\circ}$ C for 24 h and then allowed to cool to room temperature. The solvent was evaporated under reduced pressure and the residue was purified by silica gel chromatography to afford 320 mg (85%) of **5** as an yellow oil. $R_f = 0.22$ (EtOAc : cyclohexane = 4 : 1). ¹H-NMR (CDCl**3**): δ 1.75 (m, 4H); 1.94 (m, 2H); 2.24 (m, 2H); 3.66 (m,

2H); 4.20 (m, 1H); 4.37 (m, 2H); 7.28 (t, 2H, *J* = 7.40); 7.37 (t, 2H, *J* = 7.53); 7.56 (d, 2H, *J* = 7.53); 7.73 (d, 2H, *J* = 7.53). **¹³**C-NMR (CDCl**3**): 23.67 (CH**2**); 23.95 (CH**2**); 41.42 (CH**2**); 42.53 (CH**2**); 42.66 (CH); 47.32 (CH); 48.36 (CH**2**); 67.28 (CH**2**); 95.70 (Cq); 120.22–127.93 (8 \times CH); 130.01 (2 \times Cq); 141.50 (Cq); 143.95 (Cq); 156.51 (Cq); 173.62 (Cq). HRMS [FAB, *m*-NBA, pos.]: calcd. 409.1348; found 409.1335.

(2-Oxo-1-oxa-4-thia-spiro[4.4]non-3-ylmethyl)-carbamic acid *tert***-butyl ester (6)**

A solution of $2(200 \text{ mg}, 0.70 \text{ mmol})$ and triethylamine $(108 \mu\text{J},$ 0.77 mmol) in toluene (10 ml) was stirred 30 min at room temperature. The reaction was cooled to 0° C and diphenylphosphoryl azide (160 µl, 0.77 mmol) was added dropwise. After stirring over 2 h at room temperature, the solution was heated to 85 °C until the evolution of N_2 ceased. The reaction was cooled to room temperature and EtOAc (20 ml) was added. The organic phase was washed with sat. Na $HCO₃$ (15 ml), dist. H**2**O (15 ml), dried (MgSO**4**) and concentrated under reduced pressure. The residue was dissolved in toluene (10 ml) and *tert*-butanol (881 µl, 7.7 mmol) was added. The solution was kept at 75 °C for 40 h and then allowed to cool to room temperature. The solvent was evaporated under reduced pressure and the residue was purified by silica gel chromatography to afford 183 mg (73%) of 6 as an yellow oil. $R_f = 0.26$ (EtOAc : cyclohexane = 4 : 1). ¹H-NMR (CDCl₃): δ 1.42 (s 9H); 1.75 (m 2H); 1.94 (m, 2H); 2.24 (m, 2H); 3.59 (m, 2H); 4.18 (m, 1H). **¹³**C-NMR (CDCl**3**): 23.57 (CH**2**); 23.83 (CH**2**); 28.42 (CH**3**); 41.44 (CH**2**); 42.19 (CH**2**); 42.53 (2 × CH**2**); 48.35 (CH); 80.02 (Cq); 95.79 (Cq); 173.54 (Cq). HRMS [FAB, *m*-NBA, pos.]: calcd. 287.1191; found 287.1186.

3-(9*H***-Fluoren-9-ylmethoxycarbonylamino)-2-tritylsulfanylpropionic acid (7)**

To a solution of **5** (80 mg, 0.20 mmol) in THF (5 ml), cooled to 0 °C was added dropwise 0.2 M LiOH (1 ml). After 30 min the starting material was completely consumed and trityl bromide (110 mg, 0.40 mmol) was added. The solution was stirred 3 h at room temperature and adjusted to pH ∼ 3 with AcOH. The aqueous solution was extracted with EtOAc $(3 \times 10 \text{ ml})$. The organic extract was dried (MgSO**4**), filtered and concentrated under reduced pressure. The residue was purified by silica gel flash chromatography to furnish 57 mg (50%) of **7** as a white solid. Mp = 128–130 °C. R_f = 0.36 (DCM : MeOH = 95 : 5 + 1% AcOH). **¹** H-NMR ([D**6**]-DMSO): δ 2.99 (m, 1H); 3.05 (m, 2H); 4.11–4.18 (m, 3H); 7.30–7.37 (m, 19H); 7.64 (t, 2H, *J* = 8.48); 7.87 (d, 2H, *J* = 7.48). **¹³**C-NMR ([D**6**]-DMSO): 36.54 (CH); 46.55 (CH); 67.40 (Cq); 79.80 (CH**2**); 126.89–129.24 (6 × CH); 140.68 (Cq); 143.79 (Cq); 144.17 (Cq); 155.60 (Cq); 176.94 (Cq). HRMS [FAB, *m*-NBA, pos.]: calcd. 585.1974; found 585.1978.

3-(9*H***-Fluoren-9-ylmethoxycarbonylamino)-2-(4-methoxy-benzylsulfanyl)-propionic acid (8)**

To a solution of **5** (80 mg, 0.20 mmol) in THF (5 ml), cooled to 0 °C was added dropwise 0.2 M LiOH (1 ml). After 30 min the starting material was completely consumed and *p*-methoxybenzyl chloride (28 µl, 0.20 mmol) was added. The solution was stirred 2 h at 0 °C and adjusted to pH ~ 3 with AcOH. The aqueous solution was extracted with EtOAc $(3 \times 10 \text{ ml})$. The organic extract was dried (MgSO**4**), filtered and concentrated under reduced pressure. The residue was purified by silica gel flash chromatography to furnish 73 mg (80%) of **8** as a yellow oil. $R_f = 0.35$ (DCM : MeOH = 95 : 5 + 1% AcOH). ¹H-NMR ([D**6**]-DMSO): δ 3.24 (m, 1H); 3.34 (m, 2H); 3.69 (m, 3H); 3.77 (m, 2H); 4.24 (m, 3H); 6.82 (d, 2H, *J* = 8.53); 7.23 (d, 2H, *J* = 8.53); 7.31 (t, 2H, *J* = 7.15); 7.40 (t, 2H, *J* = 7.40); 7.69 (t, 2H, *J* = 6.40); 7.88 (d, 2H, *J* = 7.53). **¹³**C-NMR ([D**6**]-DMSO): 34.31 (CH**2**); 41.98 (CH); 46.25 (CH**2**); 46.98 (CH); 55.30 (CH**3**); 65.92 (CH**2**); 114.12 (2 × CH); 120.46 (2 × CH); 125.58–129.73 $(8 \times CH)$; 130.46 (Cq); 141.05 (2 \times Cq); 144.17 (2 \times Cq); 156.48 (Cq); 158.59 (Cq); 172.73 (Cq). HRMS [FAB, *m*-NBA, pos.]: calcd. 463.1453; found 463.1432.

3-Benzyloxycarbonylamino-2-(4-methoxy-benzylsulfanyl) propionic acid (9)

To a solution of **4** (1.29 g, 4.01 mmol) in 20 ml THF was added at 0 $\rm{°C}$ dropwise 1 M LiOH (15 ml) and the reaction was stirred for 1 h at room temperature. The mixture was treated with *p*-methoxybenzyl chloride (544 µl, 4.01 mmol) and stirred at room temperature for further 3 h. The reaction was adjusted to pH ∼ 3 with AcOH and H**2**O (10 ml) was added. The aqueous solution was extracted with EtOAc $(3 \times 20 \text{ ml})$. The organic phase was dried (MgSO**4**) and evaporated under reduced pressure. The residue was purified by silica gel flash chromatography to obtain 1.45 g (96%) of **9** as a yellow oil. $R_f = 0.34$ (DCM : MeOH = 95 : 5 1% AcOH). **¹** H-NMR ([D**6**]-DMSO): δ 3.38 (m, 2H); 3.47 (m, 2H); 3.75 (s, 3H); 3.83 (m, 2H); 5.05 (s, 2H); 6.81 (d, 2H, *J* = 8.53); 7.23 (d, 2H, *J* = 8.53); 7.32 (m, 5H). **¹³**C-NMR ([D**6**]-DMSO): 35.40 (CH**2**); 41.04 (CH**2**); 44.99 (CH); 55.15 (CH**3**); 66.94 (CH**2**); 113.95 (2 × CH); 128.04– 128.45 (4 \times CH); 130.18 (2 \times CH); 136.07 (Cq); 156.24 (Cq); 158.82 (Cq); 176.28 (Cq). HRMS [FAB, *m*-NBA, pos.]: calcd. 375.1140; found 375.1160.

3-*tert***-Butoxycarbonylamino-2-tritylsulfanyl-propionic acid (10)**

To a solution of **6** (344 mg, 1.17 mmol) in 10 ml THF was added dropwise 1 M LiOH (5 ml) at 0 °C. The mixture was heated to room temperature, stirred for 1 h before trityl bromide (388 mg, 1.2 mmol) was added. The solution was stirred for further 4 h and adjusted to pH ∼ 3 by AcOH. H**2**O (10 ml) was added and the aqueous solution was extracted with EtOAc $(3 \times 10 \text{ ml})$. The organic phase was dried $(MgSO₄)$, concentrated under reduced pressure and the residue was purified by silica gel flash chromatography to furnish 419 mg (78%) of **10** as a white solid. $Mp = 58-60$ °C. $R_f = 0.32$ (DCM : MeOH $= 95 : 5 + 1\%$ AcOH). ¹H-NMR ([D₆]-DMSO): δ 1.46 (s, 9H); 2.99 (m, 2H); 3.07 (m, 1H); 7.25–7.36 (m, 15H). **¹³**C-NMR ([D**6**]-DMSO): 28.17 (3 × CH**3**); 36.43 (CH**2**); 47.07 (CH); 67.35 (Cq); 77.69 (Cq); 126.93 (3 \times CH); 128.08 (3 \times CH); 129.21 (3 × CH); 144.15 (Cq); 155.01 (Cq); 172.28 (Cq). HRMS [FAB, *m*-NBA, pos.]: calcd. 463.1817; found 463.1835.

3-*tert***-Butoxycarbonylamino-2-(4-methoxy-benzylsulfanyl) propionic acid (11)**

To a solution of **6** (90 mg, 0.31 mmol) in 5 ml THF was added dropwise 1 M LiOH (3 ml) at 0 °C. The mixture was heated to room temperature, stirred for 1 h before *p*-methoxybenzyl chloride (43 µl, 0.31 mmol) was added. The solution was stirred for further 3 h and adjusted to pH ∼ 3 by AcOH. H**2**O (10 ml) was added and the aqueous solution was extracted with EtOAc $(3 \times 10 \text{ ml})$. The organic phase was dried (MgSO₄), concentrated under reduced pressure and the residue was purified by silica gel flash chromatography to furnish 81 mg (77%) of **11** as a yellow oil. $R_f = 0.34$ (DCM : MeOH = 95 : 5 + 1% AcOH). ¹H-NMR ([D₆]-DMSO): δ 1.36 (s, 9H); 3.18 (m, 2H); 3.28 (m, 2H); 3.72 (s, 3H); 3.75 (m, 2H); 6.86 (d, 2H); 7.23 (d, 2H). **¹³**C-NMR ([D**6**]-DMSO): 28.24 (3 × CH**3**); 33.92 (CH**2**); 46.13 (CH**2**); 55.05 (CH**3**); 62.61 (CH); 77.69 (Cq); 77.95 (Cq); 113.46 (CH); 113.83 (CH); 127.97 (Cq); 129.46 (CH); 130.18 (CH); 155.59 (Cq); 158.34 (Cq); 172.44 (Cq). HRMS [FAB, *m*-NBA, pos.]: calcd. 341.1297; found 341.1282.

Isocys-Arg-Ala-Glu-Tyr-Ser (16)

52 mg of Wang resin $(0.98 \text{ mmol g}^{-1}, 0.051 \text{ mmol})$ was used and the peptide assembly proceeded as described above (Fmoc-

Ser(*t*-Bu)-OH, Fmoc-Tyr(*t*-Bu)-OH, Fmoc-Glu(*t*-Bu)-OH, Fmoc-Ala-OH, Fmoc-Arg(Pmc)-OH, Boc-Isocys(Trt)-OH). The peptide was released and deprotected by treating the resin with a solution of CF**3**COOH : PhSMe : Et**3**SiH : H**2**O (92 : 3 : 3 : 2, 2 ml) for 2 h. After filtration the resin was washed two times with pure TFA (1 ml). The combined filtrates were concentrated under reduced pressure and cold ether was added (5 ml). The precipitate was collected by centrifugation and washed two times with ether. Purification by semi-preparative reversed phase HPLC and lyophilization furnished 33 mg (69%) of **16** as a white powder. MS (MALDI-TOF, DHB, pos.): *m*/*z*: calcd. for $C_{29}H_{46}N_9O_{11}S$ [M + H]: 728.8, found: 729.1.

Leu-Tyr-Lys-Ala-Gly-COS(CH₂), CONH₂ (24)

To 104 mg of MBHA resin $(0.8 \text{ mmol g}^{-1}, 0.083 \text{ mmol})$ was added a solution of 3-tritylsulfanylpropionic acid (1.47 mg, 0.42 mmol), HBTU (159.3 mg, 0.42 mmol) and DIPEA (145 µl, 0.83 mmol) in DMF/DCM $(1:1, 2$ ml). After 14 h the trityl group was removed by means of a one hour treatment with CF**3**COOH : Et**3**SiH (95 : 5, 1 ml). The peptide assembly proceeded as described above (Boc-Gly-OH, Boc-Ala-OH, Boc-Lys(2-Cl-Z)-OH, Boc-Tyr(Bz)-OH, Boc-Leu-OH). The peptide was released and deprotected by treating the resin with a solution of CF**3**COOH : CF**3**SO**3**H : PhSMe : EDT (77 : 10 : 10 : 3, 2 ml) for 2 h. After filtration the resin was washed two times with pure TFA (1 ml). The combined filtrates were concentrated under reduced pressure and cold ether was added (5 ml). The precipitate was collected by centrifugation and washed two times with ether. Purification by semi-preparative reversed phase HPLC and lyophilization furnished 35 mg (49%) of **24** as a white powder. MS (MALDI-TOF, DHB, pos.): calcd. for $C_{29}H_{48}N_7O_7S$ [M + H]: 638.8 found: 639.6.

Cys-Arg-Ala-Glu-Tyr-Ser (27)

52 mg of Wang resin $(0.98 \text{ mmol g}^{-1}, 0.051 \text{ mmol})$ were used and the peptide assembly proceeded as described above (Fmoc-Ser(*t*-Bu)-OH, Fmoc-Tyr(*t*-Bu)-OH, Fmoc-Glu(*t*-Bu)-OH, Fmoc-Ala-OH, Fmoc-Arg(Pmc)-OH, Fmoc-Cys(Trt)-OH). The peptide was released and deprotected by treating the resin with a solution of CF_3COOH : PhSMe: Et_3SH : H_2O (92:3:3) : 2, 2 ml) for 2 h. After filtration the resin was washed two times with pure TFA (1 ml) . The combined filtrates were concentrated under reduced pressure and cold ether was added (5 ml). The precipitate was collected by centrifugation and washed two times with ether. Purification by semi-preparative reversed phase HPLC and lyophilization furnished 38 mg (78%) of **27** as a white powder. MS (MALDI-TOF, DHB, pos.): *m*/*z*: calcd. for C**29**H**46**N**9**O**11**S [M H]: 728.8, found: 729.2.

Chemical ligations

Leu-Tyr-Lys-Ala-Gly-COS(CH**2**)**2**CONH**2** (**24**) (100 µl of 10 mM solution in H**2**O) was added to an argon saturated 0.1 M sodium phosphate buffer (800 μ l) adjusted to the desired pH-value by addition of 2 M NaOH. Benzylmercaptan (40 µl) was added and the mixture was stirred until the starting material was completely consumed. The corresponding C-terminal peptide (100 µl of 10 mM solution) was added and the ligation was monitored by analytical reversed phase HPLC. Gradient: 0–5 min, 5% B; 5–15 min, 5 \rightarrow 20% B; 15–20 min, $20 \rightarrow 50\%$ B; flow 1 ml min⁻¹. The eluted peptides were identified by coupled electrospray mass spectrometry. ESI-MS *m*/*z*: calcd. for Leu-Tyr-Lys-Ala-Gly-COSBn (**25**), C**33**H**48**N**6**O**6**S [M H]: 657.3, found: 657.3; calcd. for Cys-Arg-Ala-Glu-Tyr-Ser (27), $C_{29}H_{46}N_9O_{11}S$ [M + H]: 728.3, found 728.4; calcd. for Isocys-Arg-Ala-Glu-Tyr-Ser (**16**), C**29**H**46**N**9**O**11**S [M H]: 728.3, found: 728.4; calcd. for Leu-Tyr-Lys-Ala-Gly-Cys-Arg-Ala-Glu-Tyr-Ser (**28**), C**55**H**86**N**15**O**17**S [M H]: 1260.6, found: 1260.5; calcd.; for Leu-Tyr-Lys-Ala-Gly-Isocys-ArgAla-Glu-Tyr-Ser (26), $C_{55}H_{86}N_{15}O_{17}S$ [M + H]: 1260.6, found: 1260.4.

Determination of the pH–rate profile

Aliquots (50 μ I) of the ligation mixture were diluted with 20% TFA (30 µl) and analysed by RP-HPLC. The extinction coefficients at 280 nm of **16**, **25** and **27** were equal as judged by the peak areas obtained upon HPLC-analysis of equimolar mixtures. The extinction coefficient of the ligation products **26** and **28** was calculated as the sum of the extinction coefficient of **16** and **25** or **25** and **27**, respectively. These estimations were reasonable since a) extinction coefficients at 280 nm are determined by the Tyr-absorbance with one tyrosine in **16**, **25**, **27** and two tyrosine residues in the ligation products **26** and **28**, b) hydrolysis of the benzylmercaptan thioester **25** furnished a product which showed an unchanged peak area, thereby confirming that the benzylmercaptan is a weak chromophore at 280 nm. The peak areas corresponding to **16**, **25**, **26**, **27**, and **28** were therefore calibrated by weighing factors of 1, 1, 0.5, 1, 0.5, respectively. The relative content of the ligation products was determined and plotted against the reaction time. The initial rates were calculated by determining the slope of the linear range of the resulting curve. A plot of the initial rates of Cysand Isocys-mediated ligations against the pH-value furnished the pH–rate profile depicted in Fig. 3.

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