

Factors influencing on-resin depsipeptide bond formation: Case studies on daptomycin- and brevicidine-derived sequences

Dennise Palpal-latoc, Margaret A. Brimble, Paul W. R. Harris,* Aimee J. Horsfall*

Table of Contents

Abbreviations	3
Experimental	4
General Information	4
Chemicals	4
Analytical equipment	4
Amino-acid synthesis	5
<i>Scheme S1: Alloc-L-Ser(tBu)-OH synthetic scheme</i>	5
Methyl-(2S)-2-((allyloxycarbonyl)amino)-3-(tert-butyloxy)propanoate (Alloc-Ser(tBu)-OMe)	5
(2S)-2-((allyloxycarbonyl)amino)-3-(tert-butyloxy)propanoic acid (Alloc-Ser(tBu)-OH)	5
<i>Figure S1. ¹H NMR spectrum of methyl (2S)-2-((allyloxycarbonyl)amino)-3-(tert-butyloxy)propanoate (Alloc-Ser(tBu)-OMe)</i>	7
<i>Figure S2. ¹³C NMR spectrum of methyl (2S)-2-((allyloxycarbonyl)amino)-3-(tert-butyloxy)propanoate (Alloc-Ser(tBu)-OMe)</i>	8
<i>Figure S3. ¹H NMR spectrum of (2S)-2-((allyloxycarbonyl)amino)-3-(tert-butyloxy)propanoic acid (Alloc-Ser(tBu)-OH)</i>	9
<i>Figure S4. ¹³C NMR spectrum of (2S)-2-((allyloxycarbonyl)amino)-3-(tert-butyloxy)propanoic acid (Alloc-Ser(tBu)-OH)</i>	10
Peptide synthesis	11
General procedure for loading of HMP linker and subsequent Fmoc-amino-acid attachment	11
General procedure for loading first amino-acid onto 2-chlorotrityl chloride functionalised resin	11
General procedure for peptide synthesis	11
Kaiser test	12
Fmoc liberation test	12
General procedure for peptide cleavage from resin and global deprotection	12
General procedure for esterification of resin bound peptides	13
Depsipeptide conversion calculation	13
Characterisation of peptides	14
<i>Table S1: Characterisation of starting material peptides</i>	14
Supplementary Information	18
Peptidyl resin 1	18
<i>Scheme S2: General synthetic scheme for reaction of peptidyl resin 1</i>	18
<i>Table S2: Additional peak characterisation following reaction of peptidyl resin 1.</i>	18
<i>Table S3: RP-HPLC</i>	20
Peptidyl resin 2	22
<i>Scheme S4: General synthetic scheme for reaction of peptidyl resin 2</i>	22
<i>Table S4: Additional peak characterisation.</i>	22
<i>Table S5: HPLC data following reaction of peptidyl resin 2</i>	24
Peptidyl resin 3	26
<i>Scheme S3: General synthetic scheme for reaction of peptidyl resin 3</i>	26
<i>Table S6: Additional peak characterisation.</i>	26
<i>Table S7: HPLC data following reaction of peptidyl resin 3</i>	27

Peptidyl resin 4	28
<i>Scheme S5: General synthetic scheme for reaction of peptidyl resin 4</i>	28
<i>Table S8: Additional peak characterisation.</i>	28
<i>Table S9: HPLC data following reaction of peptidyl resin 4</i>	29
Peptidyl resin 5	30
<i>Scheme S6: General synthetic scheme for reaction of peptidyl resin 5</i>	30
<i>Table S10: Additional peak characterisation.</i>	30
<i>Table S11: HPLC data following reaction of peptidyl resin 5</i>	31
Peptidyl resin 6	32
<i>Scheme S7: General synthetic scheme for reaction of peptidyl resin 6</i>	32
<i>Table S12: Additional peak characterisation.</i>	32
<i>Table S13: HPLC data following reaction of peptidyl resin 6</i>	33
Peptidyl resins 7 & 12	34
<i>Scheme S8: General synthetic scheme for reaction of peptidyl resins 7</i>	34
<i>Table S14: Additional peak characterisation.</i>	34
Peptidyl resin 8	37
<i>Scheme S16: General synthetic scheme for reaction of peptidyl resin 8</i>	37
<i>Table S16: Additional peak characterisation.</i>	37
<i>Table S17: HPLC data following reaction of peptidyl resin 8</i>	38
Peptidyl resin 9	39
<i>Scheme S10: General synthetic scheme for reaction of peptidyl resin 9</i>	39
<i>Table S18: Additional peak characterisation.</i>	39
<i>Table S19: HPLC data following reaction of peptidyl resin 9</i>	40
Peptidyl resin 10	41
<i>Scheme S11: General synthetic scheme for reaction of peptidyl resin 10</i>	41
<i>Table S20: Additional peak characterisation.</i>	41
<i>Table S21: HPLC data following reaction of peptidyl resin 10</i>	42
Peptidyl resin 11	43
<i>Scheme S14: General synthetic scheme for reaction of peptidyl resin 11</i>	43
<i>Table S22: Additional peak characterisation.</i>	43
<i>Table S23: HPLC data following reaction of peptidyl resin 11</i>	45
Peptidyl resin 13	46
<i>Scheme S12: General synthetic scheme for reaction of peptidyl resin 13</i>	46
<i>Table S24: HPLC data following reaction of peptidyl resin 13</i>	46
Peptidyl resin 14	47
<i>Scheme S13: General synthetic scheme for reaction of peptidyl resin 14</i>	47
<i>Table S25: Additional peak characterisation.</i>	47
<i>Table S26: HPLC data following reaction of peptidyl resin 14</i>	48
References	48

Abbreviations

6-ClHOBt, 1-hydroxy-6-chlorobenzotriazole; **Ac**, acetyl; **Ala (A)**, alanine; **All**, allyl; **Arg (R)**, arginine; **Asn (N)**, asparagine; **Asp (D)**, aspartic acid; **Boc**, *tert*-butoxycarbonyl; **Cys (C)**, cysteine; **DCM**, dichloromethane; **DIC**, *N,N'*-diisopropylcarbodiimide; **DIPEA**, *N,N'*-diisopropylethylamine; **DMAP**, 4-dimethylaminopyridine; **DMF**, *N,N'*-dimethylformamide; **Fmoc**, 9-fluorenylmethoxycarbonyl; **Gln (Q)**, glutamine; **Glu (E)**, glutamic acid; **Gly (G)**, glycine; **h**, hour/s; **HCTU**, O-(1H-6-chlorobenzotriazole-1-yl)-1,1,3,3-tetramethyluronium hexafluorophosphate; **His (H)**, histidine; **HMP**, 4-(hydroxymethyl)phenoxyacetic acid; **HRMS**, high resolution mass spectrometry; **Leu (L)**, leucine; **Lys (K)**, lysine; **MeCN**, acetonitrile; **Met (M)**, methionine; **min**, minute/s; **MS**, mass spectrometry; **NMP**, *N*-methyl-2-pyrrolidone; **Pbf**, 2,2,4,6,7-pentamethyldihydrobenzofuran-5-sulfonyl; **PG**, protecting group; **Phe (F)**, phenylalanine; **ppm**, parts per million; **Pro (P)**, proline; **PS**, polystyrene; **RP-HPLC**, reverse phase high performance liquid chromatography; **rt**, room temperature; **sec**, second/s; **Ser (S)**, serine; ***t*Bu**, *tert*-butyl; **TFA**, trifluoroacetic acid; **Thr (T)**, threonine; **TIPS**, triisopropylsilane; **TMS**, tetramethylsilane; **TritonX100™**, *tert*-octylphenoxyethoxyethanol; **Trp (W)**, tryptophan; **Trt**, trityl; **Tyr (Y)**, tyrosine; **UV**, ultra-violet; **Val (V)**, valine.

Experimental

General Information

Chemicals

All reagents were purchased from commercial sources and used without further purification unless otherwise stated from the following sources:

Sigma Aldrich: *N,N'*-diisopropylethylamine (DIPEA), *N,N'*-diisopropylcarbodiimide (DIC), formic acid,
AK Scientific: 4-(hydroxymethyl)phenoxyacetic acid (HMP) linker, triisopropylsilane (TIPS), Fmoc-L-Asp(OH)-OAll, Fmoc-D-Ser(*t*Bu)-OH

Aapptec: 1-hydroxy-6-chlorobenzotriazole (6-ClHOBt), Fmoc-L-Trp(Boc)-OH, 4-dimethylaminopyridine (DMAP), Boc-L-Trp(H)-OH, O-(1H-6-Chlorobenzotriazole-1-yl)-1,1,3,3-tetramethyluronium hexafluorophosphate (HCTU), Fmoc-D-Asn(Trt)-OH, Fmoc-L-Asp(*t*Bu)-OH, Fmoc-D-Thr(OH)-OH, Fmoc-L-Ser(*t*Bu)-OH, Fmoc-D-Trp(Boc)-OH

Rapp Polymere: Tentagel S resin

Avantor Performance Materials: Acetic anhydride (Ac₂O), diethyl ether (Et₂O)

Oakwood Chemicals: Trifluoroacetic acid (TFA)

Scharlau: *N,N'*-Dimethylformamide (DMF, AR grade), acetonitrile (MeCN, HPLC grade), acetonitrile (MeCN, MS grade)

ECP Limited: Dichloromethane (DCM), pyridine

AusPep: Fmoc-L-Trp(H)-OH

Polypeptide laboratories: Fmoc-L-Thr(OH)-OH

Chem Supply: piperidine,

ChemPep®: 2-chlorotrityl chloride polystyrene (2-ClTrt PS)resin

Chem Impex Int'l Ltd.: Fmoc-D-Ala-OH

CS Bio (Shanghai) Ltd.: Fmoc-Gly-OH, Fmoc-L-Ile-OH, Fmoc-L-Lys(Boc)-OH

GL Biochem: Ser(*t*Bu)-OMe•HCl

Alloc-L-Ser(*t*Bu)-OH was prepared as described below, according to literature procedures.^{1,2}

Analytical equipment

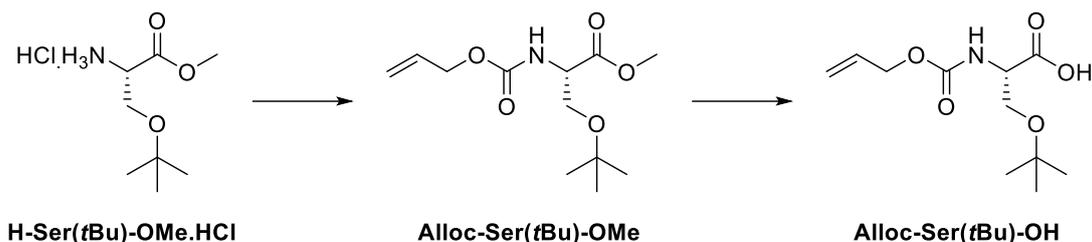
Analytical HPLC was collected on an Agilent 1100 Compact, equipped with an Agilent Zorbax SB300-C8 column (150 x 4.6 mm, 5 μm) using a linear gradient of MeCN with 0.1% (v/v) TFA (Buffer B), in H₂O with 0.1% (v/v) TFA (Buffer A) at 1 mL/min and the spectra visualised at 214 nm.

LCMS and MS (ESI+) data was collected using an Agilent 1260 Infinity liquid chromatography system equipped with an Agilent Zorbax SB300-C3 column (150 x 3 mm, 3.5 μm), over a linear gradient of MeCN with 0.1% (v/v) formic acid, in H₂O with 0.1% (v/v) formic acid at 0.3 mL/min and visualised at 214 nm; and attached to an Agilent 6120 Quadrupole electrospray mass spectrometer.

UV readings for the '*Fmoc liberation test*' were collected on a Shimadzu (Kyoto, Japan) UV-1280 system, in quartz cuvettes with a 10 mm pathlength.

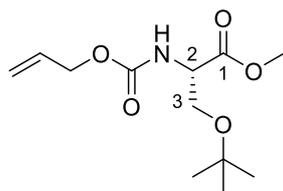
Nuclear magnetic resonance (NMR) spectra were recorded at 298 K on a Bruker AVANCE 400 spectrometer. All chemical shifts are reported in parts per million (ppm) from referenced to tetramethylsilane (TMS) at 0 ppm.

Amino-acid synthesis



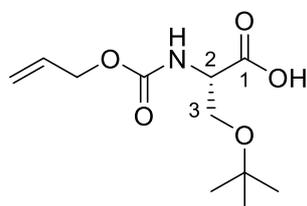
Scheme S1: Alloc-L-Ser(tBu)-OH synthetic scheme

Methyl-(2S)-2-((allyloxycarbonyl)amino)-3-(tert-butyloxy)propanoate (**Alloc-Ser(tBu)-OMe**)



A solution of Na_2CO_3 (2.5 g, 23.6 mmol, 5 eq.) in H_2O (7 mL) was added slowly to an ice-water cooled solution of H-Ser(tBu)-OMe.HCl (1.0 g, 4.72 mmol, 1 eq.) in THF (7 mL). The mixture was stirred for 10 min at 0 °C, then allyl chloroformate (1 mL, 9.44 mmol, 2 eq.) was slowly added. The resulting mixture was stirred for 12 h at rt then concentrated *in vacuo*. The reaction was extracted with EtOAc (3 × 20 mL) and the combined organic extracts were washed with brine (30 mL), dried over anhydrous Na_2SO_4 , filtered and concentrated *in vacuo*. The crude residue was purified by flash chromatography on silica gel (EtOAc-PET ether 15:85) to afford **Alloc-Ser(tBu)-OMe** as a colourless oil (1.1 g, 95%). R_f 0.35 (EtOAc-PET ether 15:85); $^1\text{H NMR}$ (400 MHz, CDCl_3): δ 5.97-5.88 (ddt, $J = 17.1, 10.9, 5.6$ Hz, 1H, Alloc), 5.58 (d, $J = 8.4$ Hz, 1H, NH), 5.32 (dd, $J = 17.2, 1.3$ Hz, 1H, Alloc), 5.22 (dd, $J = 10.4, 1.1$ Hz, 1H, Alloc), 4.59 (ddd, $J = 5.6, 1.3, 1.3$ Hz, 2H, Alloc), 4.44 (ddd, $J = 8.9, 3.0, 2.9$ Hz, 1H, H-2), 3.80 (dd, $J = 9.0, 2.8$ Hz, 1H, H-3), 3.74 (s, 3H, OCH_3), 3.57 (dd, $J = 9.0, 3.3$ Hz, 1H, H-3), 1.12 (s, 9H, tBu) ppm; $^{13}\text{C NMR}$ (100 MHz, CDCl_3): δ 171.1 (CO_2CH_3 , C-1), 156.0 (NHCO_2 , Alloc), 132.7 ($\text{CH}=\text{CH}_2$, Alloc), 117.8 ($\text{CH}=\text{CH}_2$, Alloc), 73.4 ($\text{C}(\text{CH}_3)_3$, tBu), 65.8 (CO_2CH_2 , Alloc), 62.0 (CH_2 , C-3), 54.6 (CH , C-2), 52.3 (CH_3 , OMe), 27.2 ($\text{C}(\text{CH}_3)_3$, tBu) ppm; **HRMS** (ESI+) m/z : 282.1312 (calcd for $[\text{C}_{12}\text{H}_{21}\text{NO}_5+\text{Na}]^+$, 282.1312). The spectroscopic data were in agreement with those reported in the literature.²

(2S)-2-((allyloxycarbonyl)amino)-3-(tert-butyloxy)propanoic acid (**Alloc-Ser(tBu)-OH**)



LiOH (0.23 g, 9.6 mmol, 2 eq.) was dissolved in H_2O (7 mL) and added to a solution of **Alloc-Ser(tBu)-OMe** (1.1 g, 4.8 mmol, 1 eq.) in THF (7 mL). The reaction mixture was allowed to stir at rt for 1 h and concentrated *in vacuo*. The solution was adjusted to pH 2 with 1 M HCl and extracted with EtOAc (3 × 20 mL). The combined organic extracts were washed with brine (30 mL), dried over anhydrous Na_2SO_4 , filtered and concentrated *in vacuo*. The crude residue was purified by flash chromatography on silica gel (EtOAc-PET ether 3:7 and 0.1% AcOH) to afford **Alloc-Ser(tBu)-OH** as a colourless oil (0.36 g, 50%). R_f 0.57 (EtOAc-PET ether 3:7 and 0.1% AcOH); $^1\text{H NMR}$ (400 MHz, CDCl_3): δ 5.97-5.88 (ddt, $J = 22.8, 10.9, 5.6$ Hz, 1H, Alloc), 5.58 (d, $J = 8.1$ Hz, 1H,

NH), 5.32 (dd, $J = 17.2, 1.3$, 1H, Alloc), 5.22 (dd, $J = 10.4, 1.3$, 1H, Alloc), 4.59 (d, $J = 5.6$, 2H, Alloc), 4.46 (ddd, $J = 11.6, 7.5, 3.3$ Hz, 1H, H-2), 3.90 (dd, $J = 8.8, 3.0$ Hz, 1H, H-3), 3.57 (dd, $J = 8.9, 4.9$ Hz, 1H, H-3), 1.19 (s, 9H, *t*Bu) ppm; $^{13}\text{C NMR}$ (100 MHz, CDCl_3): δ 175.1 (CO_2H , C-1), 156.4 (NHCO_2 , Alloc), 132.7 ($\text{CH}=\text{CH}_2$, Alloc), 118.2 ($\text{CH}=\text{CH}_2$, C-8), 74.3 ($\text{C}(\text{CH}_3)_3$, *t*Bu), 66.2 (CO_2CH_2 , Alloc), 61.9 (CH_2 , C-3), 54.4 (CH , C-2), 27.4 ($\text{C}(\text{CH}_3)_3$, *t*Bu) ppm; **HRMS** (ESI+) m/z : 268.1155 (calcd for $[\text{C}_{11}\text{H}_{19}\text{NO}_5+\text{Na}]^+$, 268.1155). The spectroscopic data were in agreement with those reported in the literature.^{1,2}

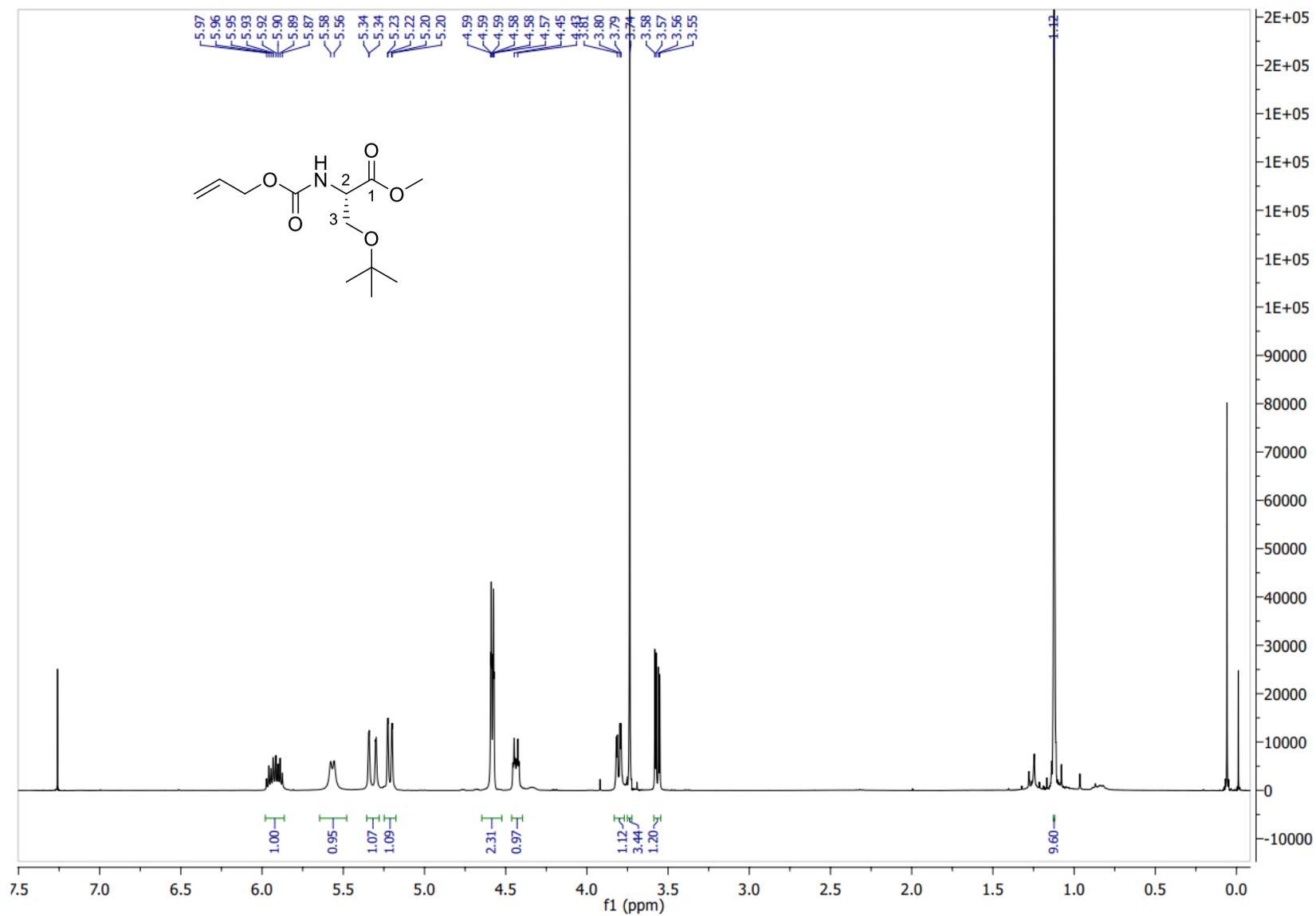


Figure S1. ¹H NMR spectrum of methyl (2S)-2-((allyloxycarbonyl)amino)-3-(tert-butyloxy)propanoate (Alloc-Ser(tBu)-OMe) (CDCl₃, 400 MHz, 298 K).

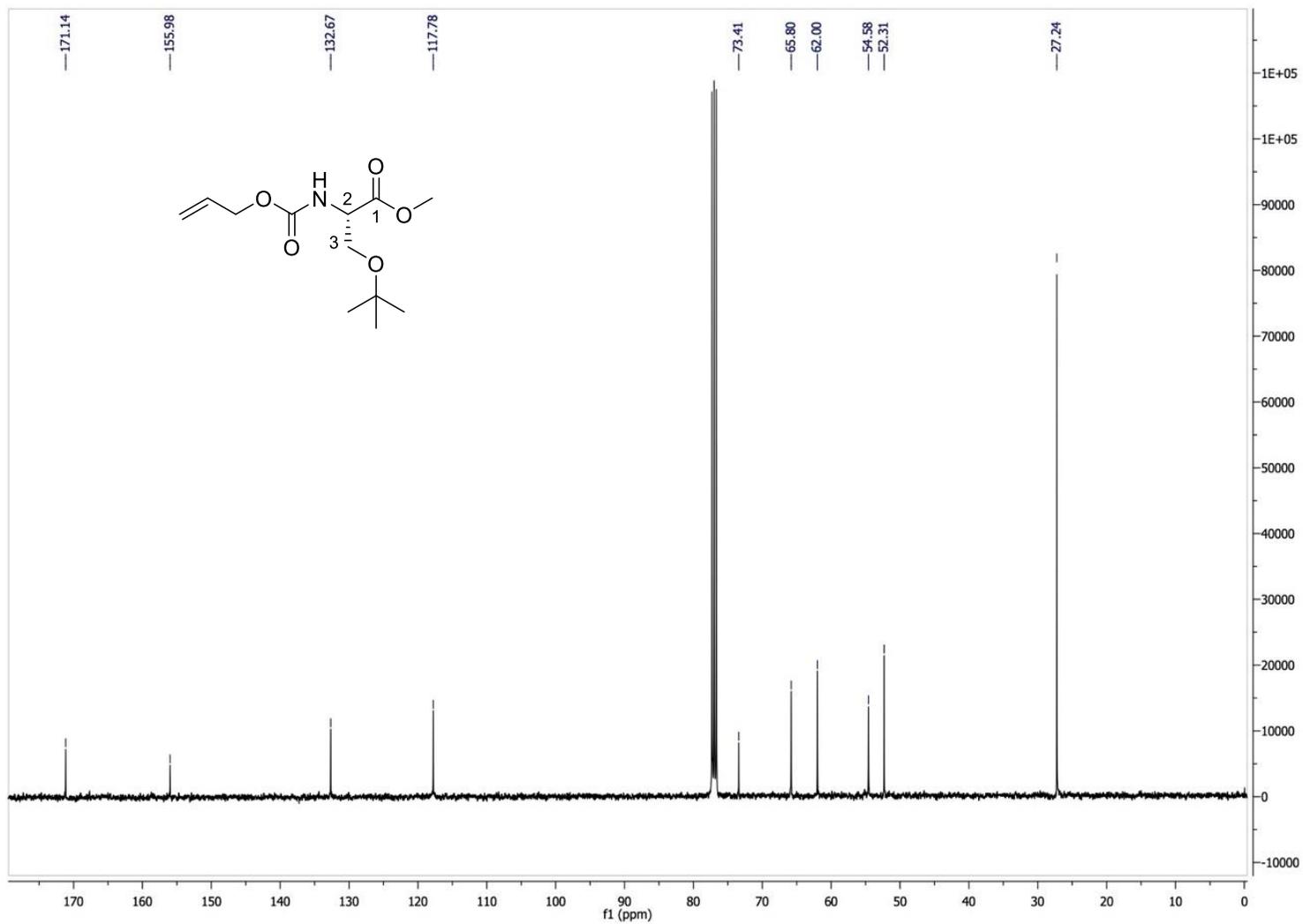


Figure S2. ¹³C NMR spectrum of methyl (2S)-2-((allyloxy carbonyl)amino)-3-(tert-butoxy)propanoate (**Alloc-Ser(tBu)-OMe**) (CDCl₃, 100 MHz, 298 K)

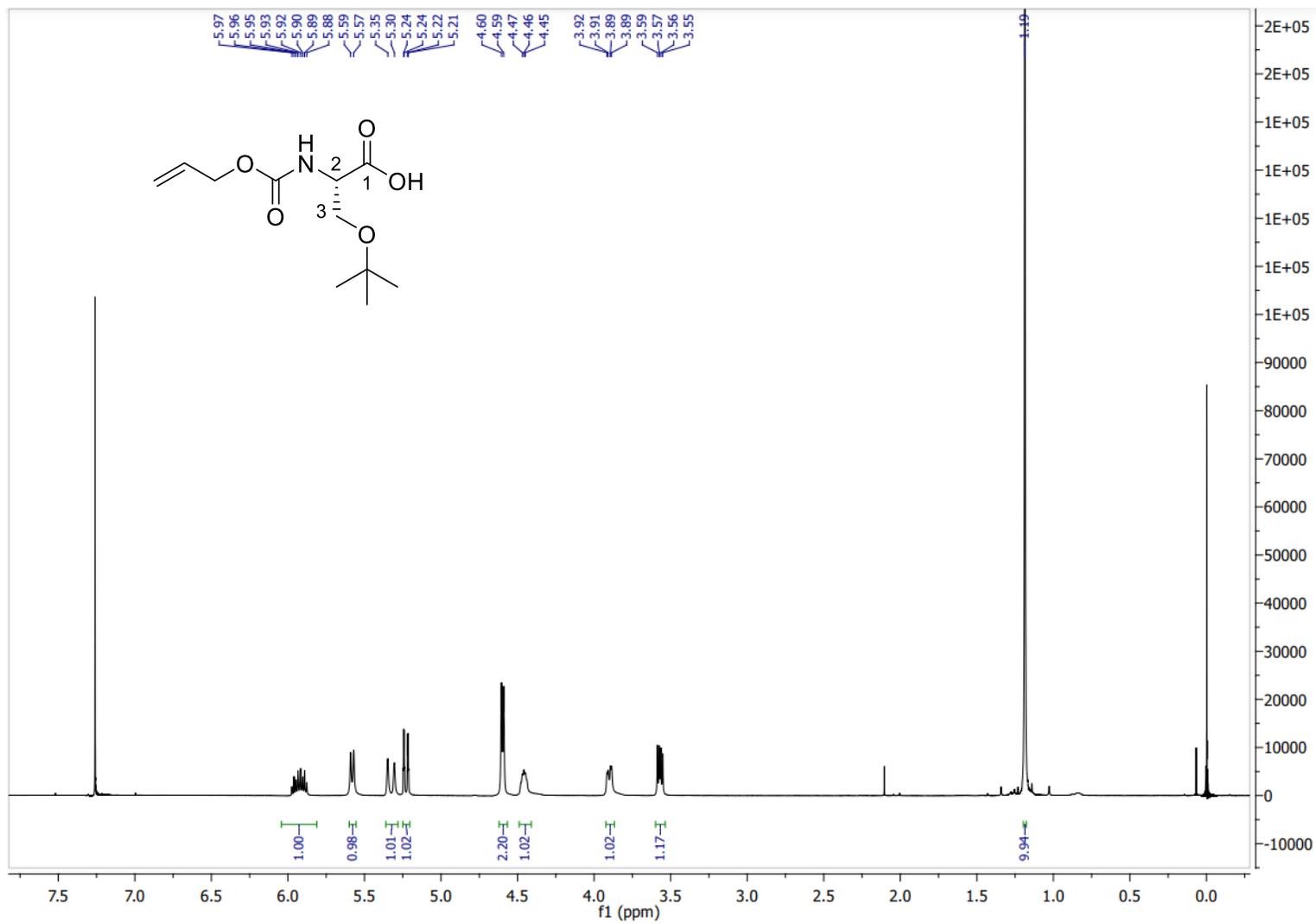


Figure S3. ¹H NMR spectrum of (2S)-2-((allyloxycarbonyl)amino)-3-(tert-butyloxy)propanoic acid (**Alloc-Ser(tBu)-OH**) (CDCl₃, 400 MHz, 298 K).

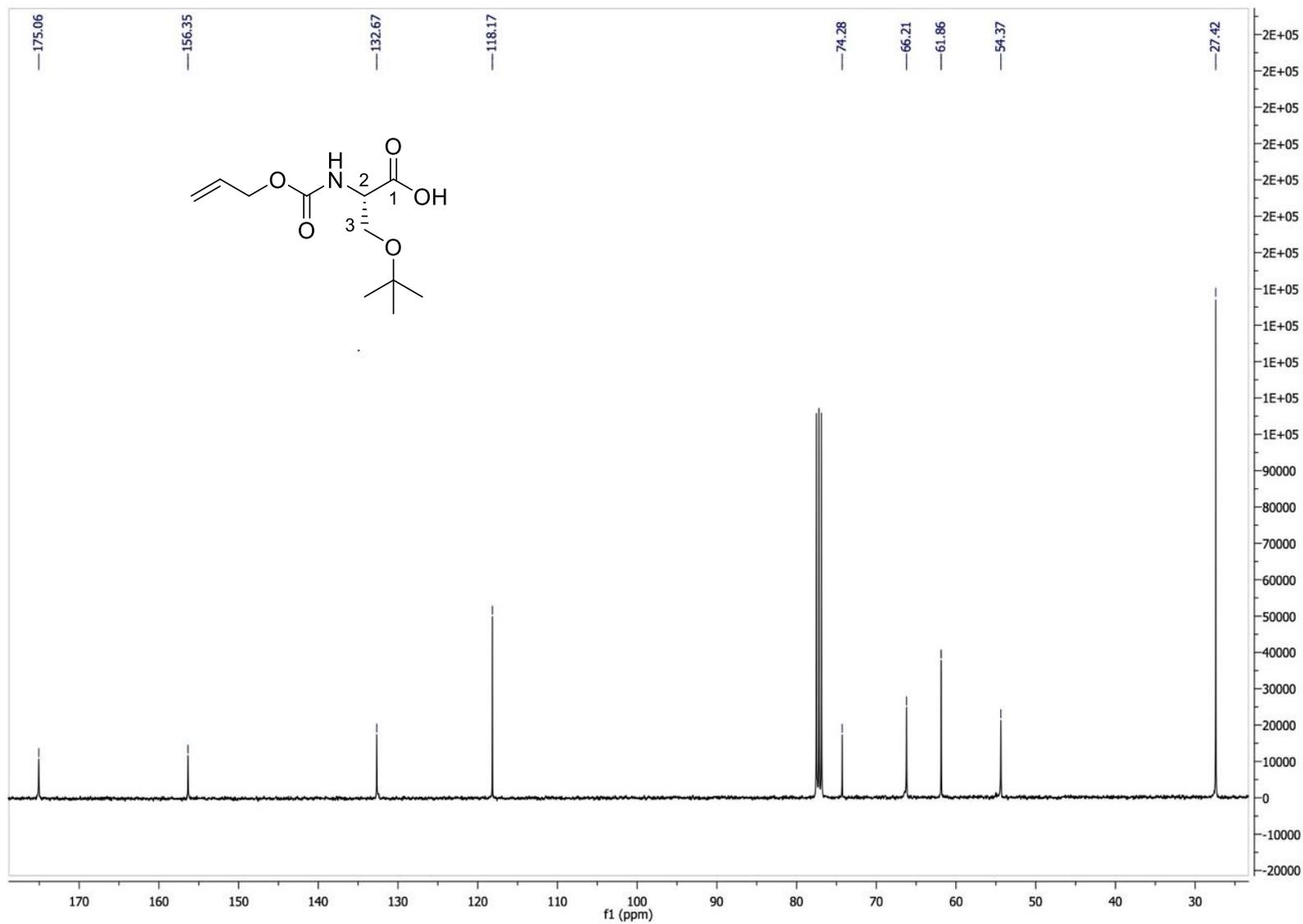


Figure S4. ^{13}C NMR spectrum of (2S)-2-((allyloxycarbonyl)amino)-3-(tert-butyloxy)propanoic acid (**Alloc-Ser(tBu)-OH**) (CDCl_3 , 100 MHz, 298 K).

Peptide synthesis

All peptides were synthesised, per the general procedures listed below, using the following protected amino-acids, unless otherwise specified: Fmoc-D-Ala-OH, Fmoc-D-Asn(Trt)-OH, Fmoc-L-Asp(OH)-OAll, Fmoc-L-Asp(tBu)-OH, Fmoc-Gly-OH, Fmoc-L-Ile-OH, Fmoc-L-Lys(Boc)-OH, Fmoc-L-Thr(OH)-OH or Fmoc-D-Thr(OH)-OH, Fmoc-L-Trp(Boc)-OH or Fmoc-L-Trp(H)-OH, and Fmoc-L-Ser(tBu)-OH. Identity of the linear peptide sequence, prior to esterification, was confirmed by RP-HPLC and MS (ESI+) by a 'minicleave' of a small portion of resin. Peptidyl resin was then split, and subjected to esterification reaction conditions. Amino-acids building blocks used in esterification reactions included Fmoc-L-Trp(Boc)-OH, Fmoc-L-Trp(H)-OH, Boc-L-Trp(H)-OH, Fmoc-D-Trp(Boc)-OH, Fmoc-L-Ser(tBu)-OH, Fmoc-D-Ser(tBu)-OH and Alloc-L-Ser(tBu)-OH.

General procedure for loading of HMP linker and subsequent Fmoc-amino-acid attachment

Tentagel S resin (0.26 mmol/g) was swelled in DMF (7 mL/g resin) for 15 min and then the solution drained from the resin. A solution of 4-(hydroxymethyl)phenoxyacetic acid (HMP linker, 4 equiv), 1-hydroxy-6-chloro-benzotriazole (6-ClHOBt, 4 equiv) and *N,N'*-diisopropylcarbodiimide (DIC, 4 equiv) in DMF (4 mL/g resin) was added to the preswelled resin and agitated for 2 h. The solution was drained from the resin, and the resin was washed with DMF (3 × 8 mL/g resin) and DCM (2 × 8 mL/g resin). A 'Kaiser test' was performed to determine if all amine groups were acylated. The resin was immediately reswelled in DCM, and the solution drained from the resin. A solution of Fmoc-amino-acid (Fmoc-L-Asp(OH)-OAll, Fmoc-Gly-OH or Fmoc-L-Ile-OH) (5 equiv) and DIC (5 equiv) in DCM (4 mL/g resin) was then added to the resin, followed by *N,N*-dimethylaminopyridine (DMAP, 0.1 equiv) and the mixture agitated for 4 h, or overnight. Acetic anhydride (250 µL/g resin) was then added to the mixture and agitated for a further 1 h, to ensure any remaining hydroxyl groups were acylated. The solution was then drained from the resin, and the resin washed with DMF (3 × 8 mL/g resin) and DCM (3 × 8 mL/g resin). An 'Fmoc liberation test' was performed to determine the loading of the Fmoc-amino-acid to the resin. Final resin loadings ranged between 0.10-0.19 mmol/g.

General procedure for loading first amino-acid onto 2-chlorotrityl chloride functionalised resin

2-chlorotrityl chloride functionalised polystyrene resin (0.5 g, 0.89 mmol/g) was swelled in DCM (10 mL) for 15 min. Fmoc-protected amino-acid (0.45 mmol, 1 equiv) was dissolved in DCM (5 mL) and DIPEA (75.6 µL, 2.25 mmol, 5 equiv) added. The amino-acid solution was added to the resin, and the solution shaken (HCl gas evolved) and vented. The mixture was placed on rocker for 2 h and then MeOH (5 mL) added and rocked for a further 30 min to cap any free reactive sites. The solvent was removed and the resin washed with DMF (5 × 8 mL), DCM (2 × 8 mL), DMF (2 × 8 mL) and DCM (5 × 8 mL), then the resin dried by suction. An 'Fmoc liberation test' was then performed to determine the loading of the Fmoc-amino-acid to the resin. Final resin loadings ranged between 0.077-0.095 mmol/g.

General procedure for peptide synthesis

Resin preloaded with the first Fmoc-amino-acid (0.1 mmol) was preswelled in DMF (5 mL) for 15 min, and the solvent then drained from the resin. Solid-phase peptide chain elongation was carried out on an Activotec Activo-P14 peptide synthesiser. A solution of 20% (v/v) piperidine in DMF (3.8 mL) was added to the resin, and the mixture agitated at rt for 3 min. The solution was then drained from the resin, and a fresh portion of 20% (v/v) piperidine in DMF (3.8 mL) was added to the resin, and the mixture agitated at rt for a further 9 min. The

solution was then drained from the resin and the resin washed with DMF (5 × 1 min, 4 mL). A solution of Fmoc-amino-acid (5 equiv), HCTU (4.9 equiv, 980 µL, 0.5 M in DMF) and DIPEA (10 equiv, 174 µL) in DMF (made up to 4 mL total) was added to the resin and the mixture agitated at rt for 20 min. The solution was then drained from the resin and the resin washed with DMF (4 × 1 min, 4 mL). The deprotection and coupling cycles were repeated until the desired sequence was achieved. Following the final *N*-terminal Fmoc deprotection, the resin was removed from the synthesiser, and a solution of acetic anhydride (470 µL, 50 equiv) and DIPEA (870 µL, 50 equiv) in DMF (2.5 mL) was added to the resin and agitated for 30 s. The solution was then drained from the resin, and the resin washed with DMF (3 × 5 mL), then DCM (3 × 5 mL) and then airdried.

Kaiser test

A small sample of resin (~2-5 mg) was washed thoroughly with DCM, and then 2 drops of each 80% (w/v) phenol in *n*-butanol (40 g in 20 mL), 5% (w/v) ninhydrin in *n*-butanol (1.0 g in 20 mL) and 200 µM KCN in pyridine (16.5 mg KCN in 25 mL H₂O, 1 mL KCN aq. solution in 49 mL pyridine) added to the resin. The mixture was then heated at 100°C for 5 min and the colour assessed. Blue beads (positive result) indicate the presence of free amines, and colourless/yellow beads (negative result) indicate the absence of free amines.

Fmoc liberation test

Immediately following coupling of the Fmoc-protected building block to the resin, two samples of resin (~1-5 mg) were thoroughly washed with DCM (6 × 300 µL) and airdried with suction. The resin samples were transferred to eppendorfs and the accurate mass of the resin recorded, then freshly prepared 20% (v/v) piperidine in DMF (1.0 mL) added to the resin. The mixture was agitated for 20 min at rt. A sample of the supernatant (200 µL) was then taken and diluted 10-fold in 20% (v/v) piperidine in DMF (1.80 mL), and the absorbance at 290 nm recorded. Absorbance at 290 nm was first zeroed against a sample of 20% (v/v) piperidine in DMF. The loading of the resin was then calculated for each sample, using the following equation:

$$\text{Resin loading} \left(\frac{\text{mmol}}{\text{g}} \right) = \frac{(A_{290} \times DF \times V_{rxn})}{(\epsilon_{290} \times m_{resin})}$$

Where, A_{290} is the absorbance at 290 nm as recorded by a UV-Vis spectrophotometer

DF is the dilution factor (here, 10)

V_{rxn} is the volume (in mL) of the deprotection reaction (here, 1 mL)

ϵ_{290} is the extinction coefficient of the fulvene-piperidine adduct at 290 nm ($\epsilon_{290} = 6089$)³

m_{resin} is the mass of resin in reaction sample (in g)

The two values were averaged (mean) to give the average loading of the resin.

General procedure for peptide cleavage from resin and global deprotection

A solution of 95:2.5:2.5 TFA:TIPS:H₂O (~300 µL) was added to a small portion of the peptidyl resin (~10 mg) and agitated for 1.5 h to cleave the peptides from the resin. The TFA solution was then concentrated under a stream of nitrogen to ~100-200 µL, and cold diethyl ether (1.0 mL) added to precipitate the peptides. The mixture was centrifuged to pellet the precipitate, the solvent was decanted and the pellet dried under a stream of nitrogen. The pellet was resuspended in 50% aq. MeCN with 0.1% TFA (~200 µL), syringe filtered (0.45 µm) analysed by RP-HPLC over a linear gradient at 214 nm, and peptide identity confirmed by MS analysis.

General procedure for esterification of resin bound peptides

Reagents, tryptophan/serine, DIC and DMAP, were freshly made up as stock solutions in the reaction solvent specified (DMF, DCM, NMP, or 1:1:1 DMF:DCM:NMP). A 6-ClHOBt stock, where applicable, was made up in DMF regardless of reaction solvent due to limited solubility in DCM. An *example* of the typical stock solutions prepared is provided,

DIC stock in reaction solvent (5 equiv/reaction, 0.05 mmol/20 μ L): 156.3 μ L DIC in 243.7 μ L solvent

DMAP stock in reaction solvent (0.1 equiv/reaction, 0.001 mmol/20 μ L): 36.6 mg DMAP in 6.0 mL solvent

Example of amino-acid stock, Fmoc-L-Trp(Boc)-OH in reaction solvent (5 equiv/reaction, 0.05 mmol/50 μ L): 263 mg Fmoc-L-Trp(Boc)-OH in 0.5 mL solvent

Peptidyl resin (0.01 mmol) was preswelled in the reaction solvent (0.5 mL) for 15 min then the solvent pipetted from the resin. Reaction solvent (final reaction volume 0.5 mL) was added to the resin, followed by amino-acid stock (5 equiv, 50 μ L), then optional additives as specified (e.g. TritonX100, or 6-ClHOBt as a stock solution in DMF). DIC (5 equiv, 20 μ L), and finally, DMAP (0.1 equiv, 20 μ L) were added to the mixture, which were then agitated for 1 h, and the solvent mixture then pipetted from the resin and fresh reagents added to the resin. The mixtures were agitated for a further 1 h, the solvent mixture then pipetted from the resin, and the resin was washed with DCM (2 \times 0.5 mL). A portion of the resin was taken, and the peptide cleaved from the resin as detailed in '*General procedure for peptide cleavage from resin and global deprotection*'.

Depsipeptide conversion calculation

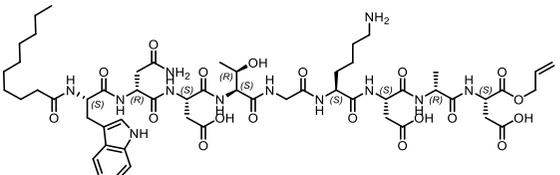
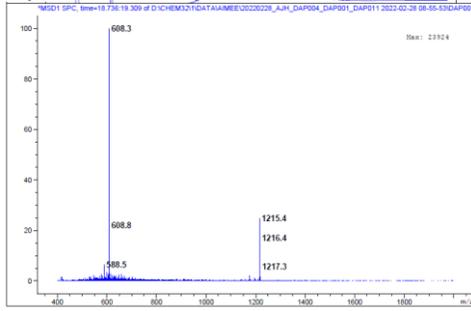
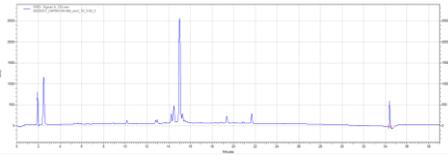
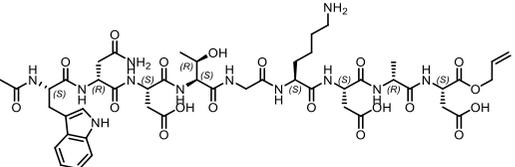
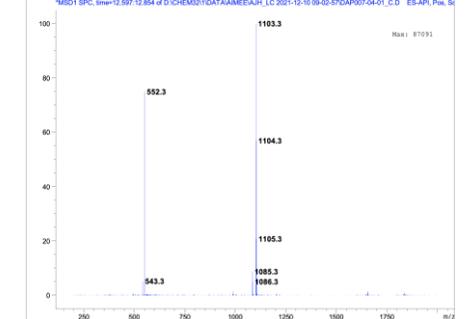
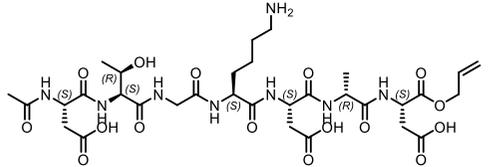
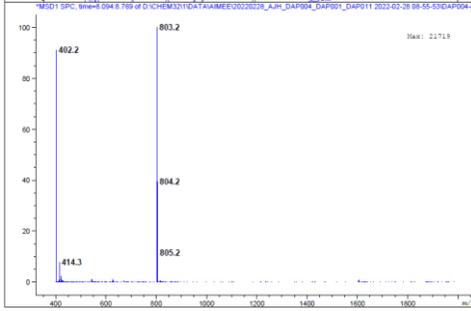
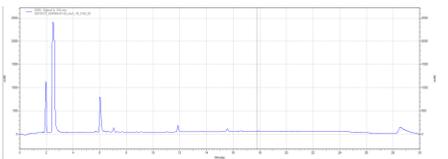
A HPLC spectrum of the reaction mixture was collected and visualised at 214 nm. The resulting spectrum was analysed in the Agilent EZChrom software, and integrated over the relevant gradient time. The percentage (%) area under the curve of the product peak, and any relevant adducts, (e.g. tryptophan CO₂ adducts, as specified in Tables S3, S5, S7, S9, S11, S13, S15, S17, S19, S21, S23, S26), was summed and divided by the sum of the % area of the starting material and product peaks (see generalised equation below). The mean average conversion of at least two experiments is reported.

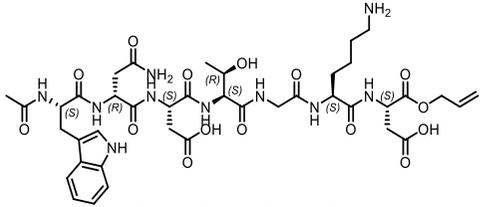
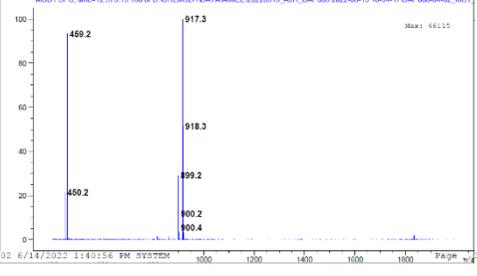
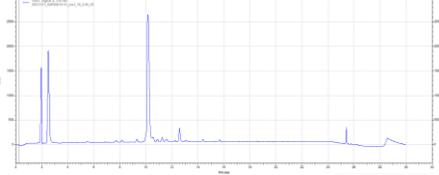
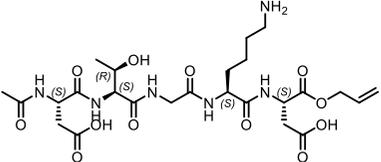
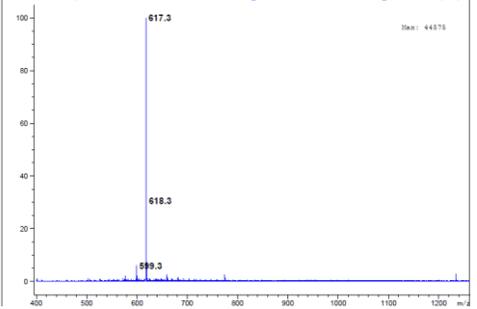
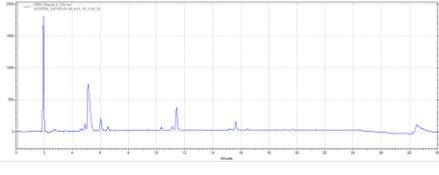
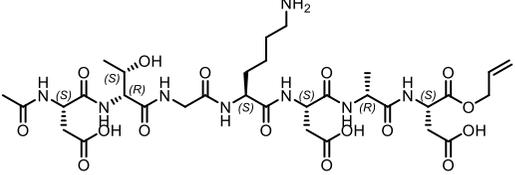
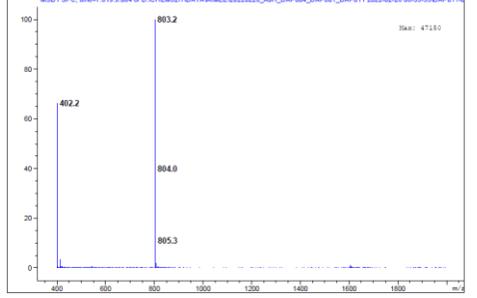
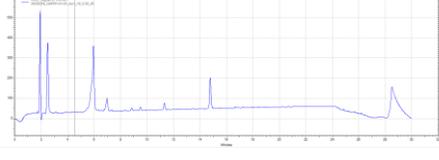
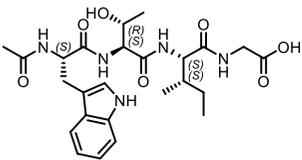
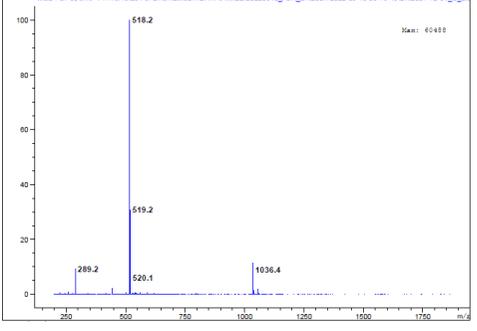
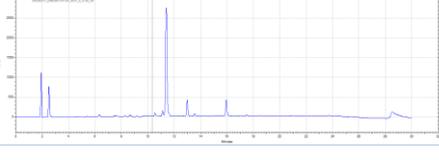
$$\%conversion\ to\ depsipeptide = \frac{\sum Area\ under\ product\ peaks}{\sum area\ under\ product\ and\ starting\ material\ peaks} \times 100$$

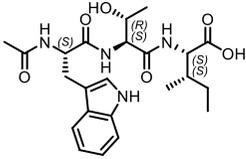
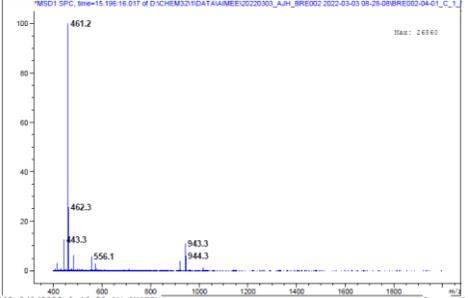
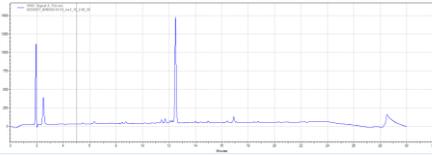
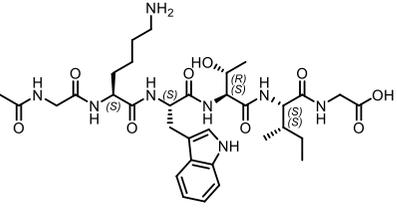
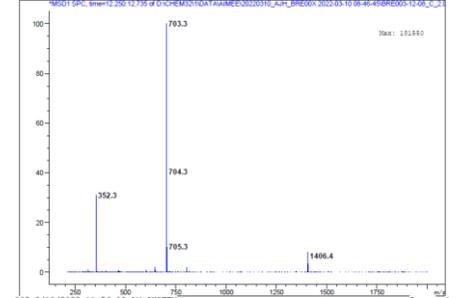
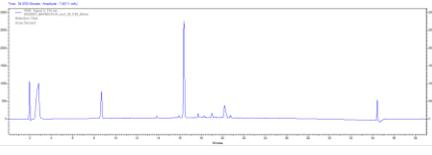
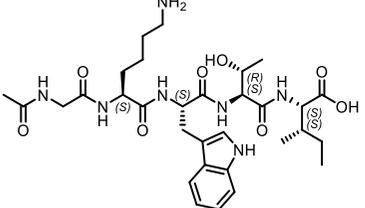
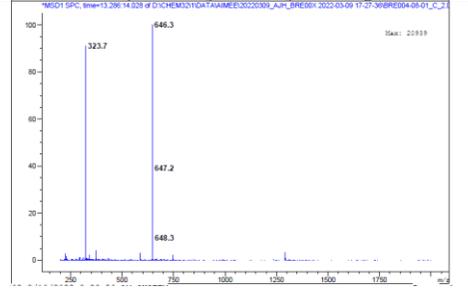
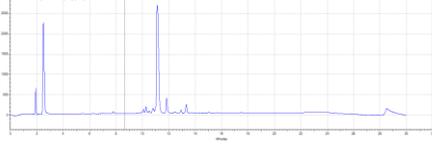
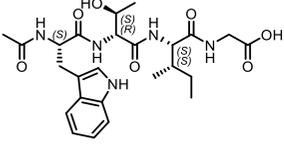
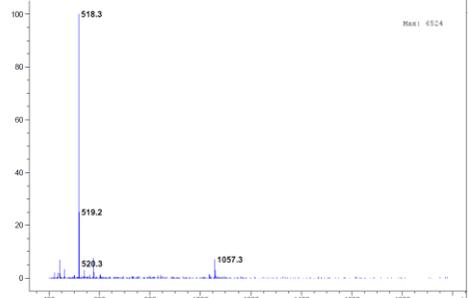
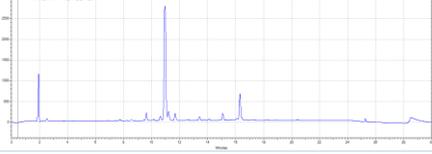
Note: The dried precipitate from the cleavage, was resuspended in 50% aq. MeCN with 0.1% TFA, filtered, and analysed by RP-HPLC within 2 h of redissolution. Extended time in solution, resulted in hydrolysis of the depsipeptide bond for some sequences. Omission of the acid or MeCN resulted in solubility problems for some sequences.

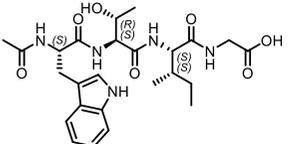
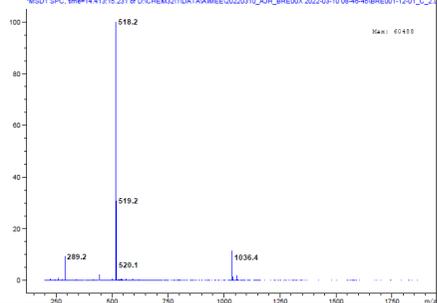
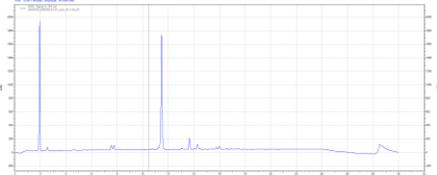
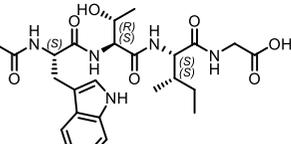
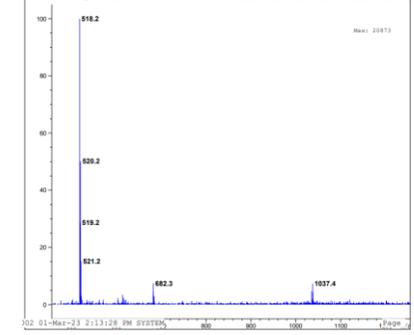
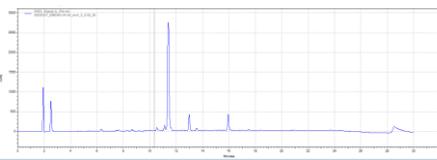
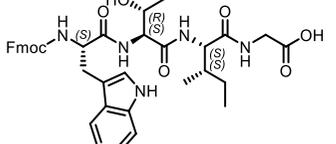
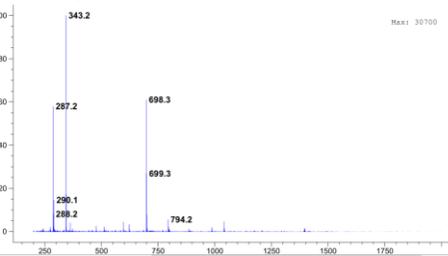
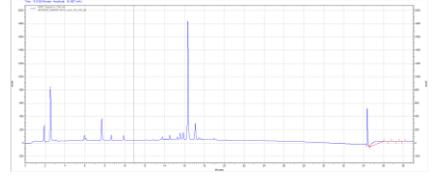
Characterisation of peptides

Table S1: Characterisation of starting material peptides

Peptidyl resin code	Peptide code	Structure	Mass found [M+H] ⁺	MS	R _t	HPLC Representative starting material (crude)
1 (TG)	S1	 <p>Chemical Formula: C₅₅H₈₂N₁₂O₁₉ Exact Mass: 1214.5819</p>	1215.7		15.1 min	<p>5-95% B over 30min</p> 
3 (TG)	S3	 <p>Chemical Formula: C₄₇H₆₆N₁₂O₁₉ Exact Mass: 1102.4567</p>	1103.6		10.2 min	<p>5-50% B over 20min</p> 
2 (TG)	S2	 <p>Chemical Formula: C₃₂H₅₀N₈O₁₆ Exact Mass: 802.3345</p>	803.3		6.0 min	<p>5-50% B over 20min</p> 

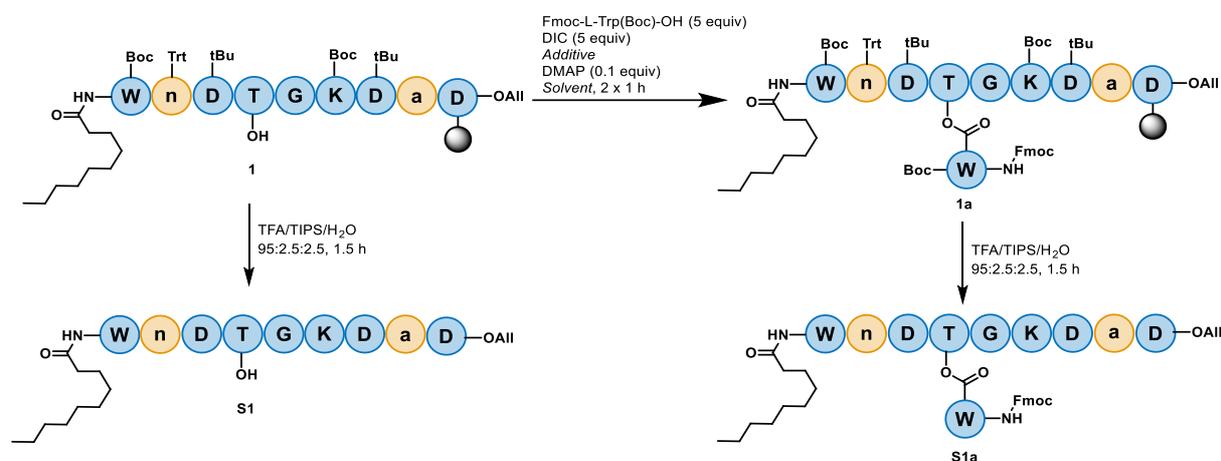
4 (TG)	S4	 <p>Chemical Formula: C₄₀H₅₆N₁₀O₁₅ Exact Mass: 916.3927</p>	917.5	 <p>MS/1 SPC: Inset 12.573.13.180 of D:\CHEM3\1\DATA\AMEE\20220513_A.H_DAP09_2022-06-13 10:56:17\DAP09-04-02_MCI</p>	10.1 min	<p>5-50% B over 20min</p> 
5 (TG)	S5	 <p>Chemical Formula: C₂₅H₄₀N₆O₁₂ Exact Mass: 616.2704</p>	617.3	 <p>MS/1 SPC: Inset 0.652715 of D:\CHEM3\1\DATA\AMEE\JULY_2021\08-03 16:03-ORNDP05-05-01_MCI.D_ES-APR_Pos.02</p>	5.2 min	<p>5-50% B over 20min</p> 
6 (TG)	S6	 <p>Chemical Formula: C₃₂H₅₀N₈O₁₆ Exact Mass: 802.3345</p>	803.2	 <p>MS/1 SPC: Inset 7.918.984 of D:\CHEM3\1\DATA\AMEE\OCTOBER_A.H_DAP08_DAP07_DAP011_2022-02-28 08:59-53\OAP011</p>	6.0 min	<p>5-50% B over 20min</p> 
7 (TG)	S7	 <p>Chemical Formula: C₂₅H₃₅N₅O₇ Exact Mass: 517.2536</p>	518.2	 <p>MS/1 SPC: Inset 14.15.16.251 of D:\CHEM3\1\DATA\AMEE\20220510_A.H_BR000_2022-05-10 06:48:49\BR001-13-01_C_2</p>	11.4 min	<p>5-50% B over 20min</p> 

<p>8 (TG)</p>	<p>S8</p>	 <p>Chemical Formula: C₂₃H₃₂N₄O₆ Exact Mass: 460.2322</p>	<p>461.2</p>	 <p>MS1 SPC: time=15.181616 of D:\CHEM2\1\DATA\AMEE\20220303_AH_BRE02\2022-03-03-08-38-09\BRE02-04-01_C_1</p> <p>Peak: 24940</p> <p>461.2, 462.3, 443.3, 556.1, 943.3, 944.3</p>	<p>12.5 min</p>	<p>5-50% B over 20min</p> 
<p>9 (TG)</p>	<p>S9</p>	 <p>Chemical Formula: C₃₃H₅₀N₈O₉ Exact Mass: 702.3701</p>	<p>703.3</p>	 <p>MS1 SPC: time=12.261215 of D:\CHEM2\1\DATA\AMEE\20220310_AH_BRE01\2022-03-10-08-44-09\BRE01-12-04_C_1</p> <p>Peak: 351940</p> <p>352.3, 703.3, 784.3, 785.3, 1406.4</p>	<p>10.1 min</p>	<p>5-50% B over 20min</p> 
<p>10 (TG)</p>	<p>S10</p>	 <p>Chemical Formula: C₃₁H₄₇N₇O₈ Exact Mass: 645.3486</p>	<p>646.3</p>	 <p>MS1 SPC: time=13.261403 of D:\CHEM2\1\DATA\AMEE\20220309_AH_BRE01\2022-03-09-11-21-39\BRE01-09-01_C_1</p> <p>Peak: 20939</p> <p>323.7, 646.3, 647.2, 648.3</p>	<p>11.2 min</p>	<p>5-50% B over 20min</p> 
<p>11 (TG)</p>	<p>S11</p>	 <p>Chemical Formula: C₂₅H₃₅N₅O₇ Exact Mass: 517.2536</p>	<p>518.3</p>	 <p>MS1 SPC: time=0.7031468 of D:\CHEM2\1\DATA\AMEE\20220705_AH_TAR01\BRE01\2022-07-05-14-46-32\BRE01-01-02</p> <p>Peak: 6524</p> <p>518.3, 519.2, 520.3, 1057.3</p>	<p>11.0 min</p>	<p>5-50% B over 20min</p> 

12 (PS)	S7	 <p>Chemical Formula: C₂₅H₃₅N₅O₇ Exact Mass: 517.2536</p>	518.2		11.4 min	<p>5-50% B over 20min</p> 
13 (PS)	S7	 <p>Chemical Formula: C₂₅H₃₅N₅O₇ Exact Mass: 517.2536</p>	518.2		11.5 min	<p>5-50% B over 20min</p> 
14 (PS)	S12	 <p>Chemical Formula: C₃₈H₄₃N₅O₈ Exact Mass: 697.3112</p>	698.3		16.3 min	<p>5-95% B over 30min</p> 

Supplementary Information

Peptidyl resin 1



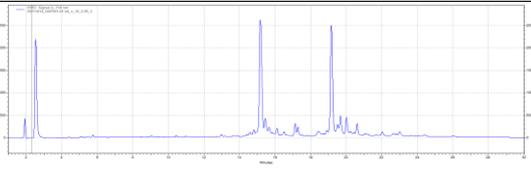
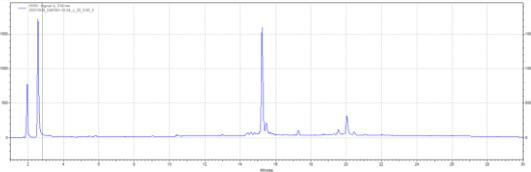
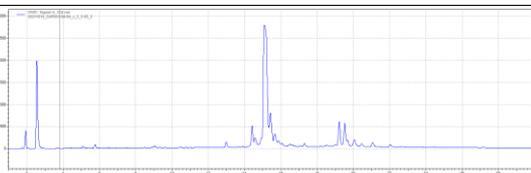
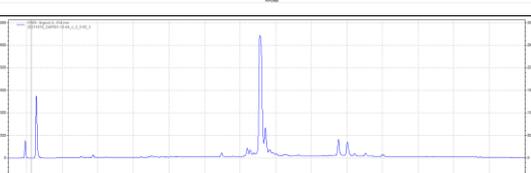
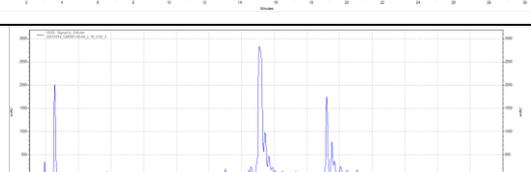
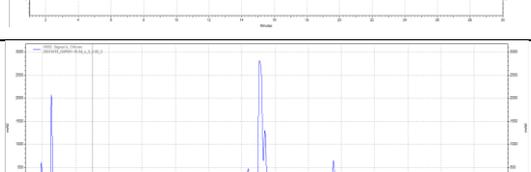
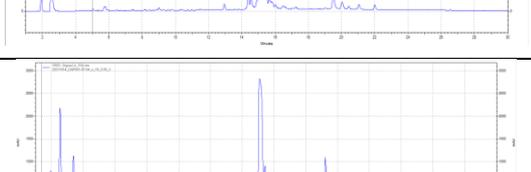
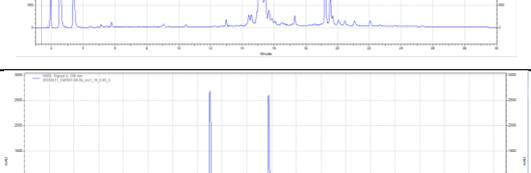
Scheme S2: General synthetic scheme for reaction of peptidyl resin 1 (HMP-functionalised Tentagel®) with Fmoc-L-Trp(Boc)-OH for 2 x 1 h, or 20 h (2 x 1 h, + 18 h) as described in the Experimental Section. The resulting reaction mixtures were characterised by RP-HPLC and MS and products identified in Table S2. The experimental conditions were varied and the resulting RP-HPLC traces are summarised in Table S3.

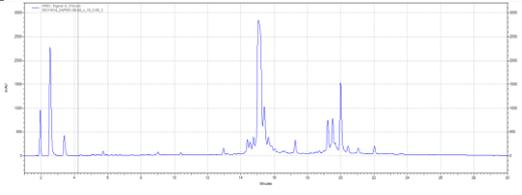
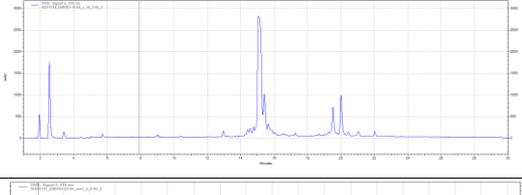
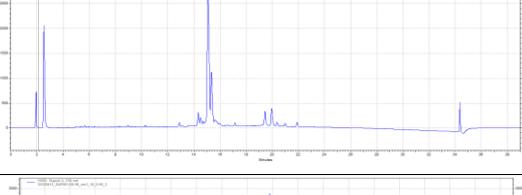
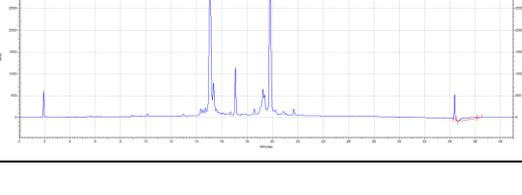
Table S2: Additional peak characterisation following reaction of peptidyl resin 1. Representative trace following reaction of 1 with Fmoc-L-Trp(Boc)-OH, as shown in Scheme S2, illustrating key products and potential by-products as identified by ESI+ MS.

Peak	Extracted LCMS ESI+	R _t (5-95% B)	Observed m/z	Identification (Mw)
A	<p>MS/MS SPC, time=18.385 of D:\CHEM32\TIDA\AGAMEE\20220803_AJH_DAP001_DAP007_2022-08-03 11-34-49\DAP001-08-06_ML</p> <p>Max: 17176</p> <p>302 8/4/2022 10:49:42 AM SYSTEM Page 2/5</p>	14.4 min	[M+H] ⁺ 1260.4 [M+2H] ²⁺ 630.4	S1+CO ₂ (1259.3)

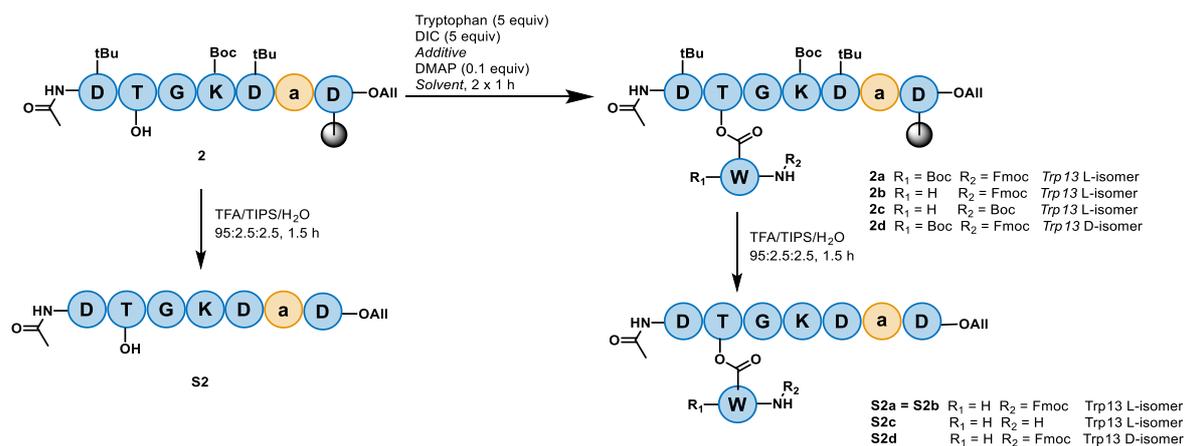
B		15.0 min	$[M+H]^+$ 1215.4 $[M+2H]^{2+}$ 608.2	S1 (1214.6)
C / D		16.1 min 17.0 min	$[M+H]^+$ 471.0 $[M+H]^+$ 427.2	C) Fmoc-Trp-OH + CO ₂ (470.2) D) Fmoc-Trp-OH (426.2)
E		18.2 min	$[M+2H]^{2+}$ 856.8 $[M+H]^+$ 1712.4	S1a+2CO ₂ (1711.8)
F		18.9 min	$[M+H]^+$ 834.4 $[M+2H]^{2+}$ 1667.4	S1a+CO ₂ (1666.7)
G		19.8 min	$[M+H]^+$ 1624.3 $[M+2H]^{2+}$ 812.3	S1a (1622.7)

Table S3: RP-HPLC data following reaction of peptidyl resin 1 with Fmoc-L-Trp(Boc)-OH to give depsipeptidyl resin 1a, as described in the Experimental section and depicted in Scheme S2. Entries 1-7 show data for Table 2 (entries 1-7), and entries 8-9 show data for Table 3 (entry 1). The conversion of peptidyl resin 1 to 1a was determined as described in the Experimental section using the peaks of the linear (S1) and branched depsipeptide (S1a), and any corresponding adducts, specified.

Entry	Reaction time	Additive	Solvent	Conversion of 1 to 1a (%; 214 nm)	HPLC (5-95% B, 3%B/min, 214 nm)	R _t of peaks used for conversion calculation
1	2 x 1 h	-	DMF	10.6%		S1 15.2 min S1a 20.0 min
	2 x 1 h + 18 h	-	DMF	19.0%		S1 15.2 min S1a 20.0 min
2	2 x 1 h	1% TritonX100	DMF	4.8%		S1 15.1 min S1a 20.0 min
	2 x 1 h + 18 h	1% TritonX100	DMF	8.5%		S1 15.1 min S1a 20.0 min
3	2 x 1 h	10% TritonX100	DMF	4.9%		S1 15.1 min S1a 20.0 min
	2 x 1 h + 18 h	10% TritonX100	DMF	4.6%		S1 15.1 min S1a 20.0 min
4	2 x 1 h	-	NMP	3.5%		S1 15.1 min S1a 20.1 min
5	2 x 1 h	-	DCM	47.1%		S1+CO ₂ 14.4 min S1 14.9 min S1a+CO ₂ 19.0 min S1a 19.8 min

6	2 x 1 h	-	1:1:1 DMF:DCM: NMP	19.5%		S1 15.1 min S1a 20.0 min
7	2 x 1 h	1% TritonX100	1:1:1 DMF:DCM: NMP	14.3%		S1 15.2 min S1a 20.0 min
8	2 x 1 h	-	DMF	8.2%		S1 15.1 min S1a 20.0 min
9	2 x 1 h	-	DCM	48.2%		S1 15.1 min S1a 19.8 min

Peptidyl resin 2



Scheme S4: General synthetic scheme for reaction of peptidyl resin 2 (HMP-functionalised Tentagel®) with appropriately protected tryptophan (specified in Table S5) for 2 x 1 h as described in the Experimental Section. The resulting reaction mixtures were characterised by RP-HPLC and MS and the products identified in Table S4. The experimental conditions were varied and the resulting RP-HPLC traces are summarised in Table S5.

Table S4: Additional peak characterisation. Representative trace following reaction of **2** with Fmoc L-Trp(Boc)-OH, as shown in Scheme S4, illustrating key products and potential by-products as identified by ESI+ MS.

Peak	Extracted LCMS ESI+	R _t (5-50% B)	Observed m/z	Identification (Mw)
A		5.9 min	[M+H] ⁺ 803.2 [M+2H] ²⁺ 462.2	S2 (802.8)

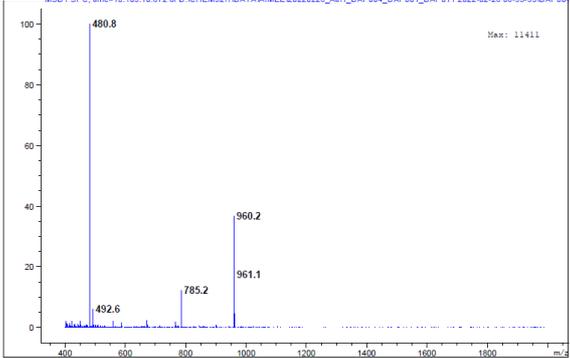
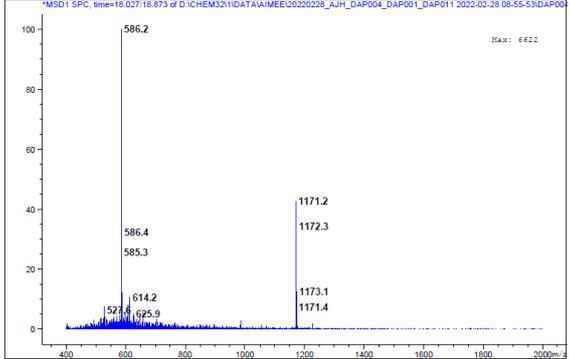
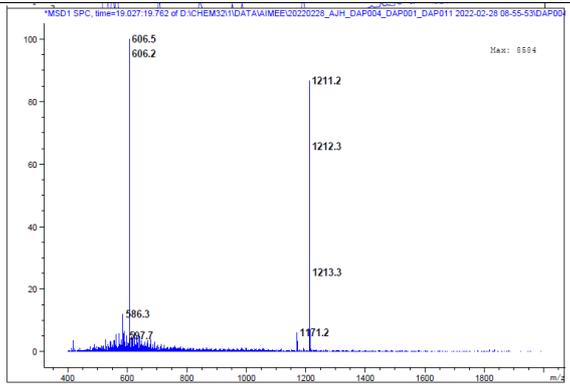
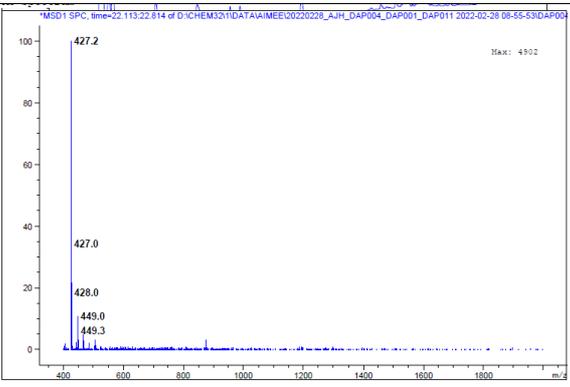
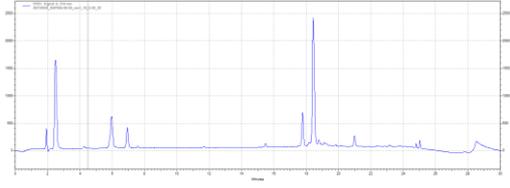
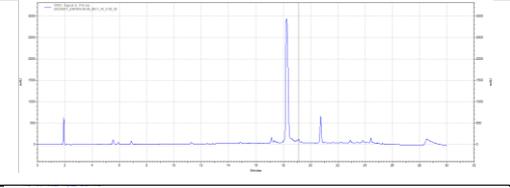
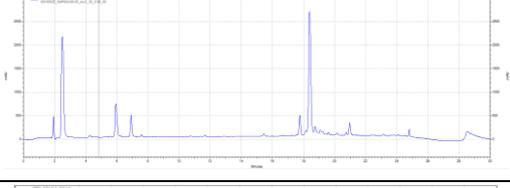
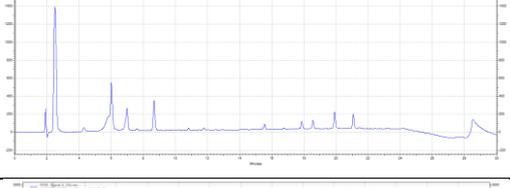
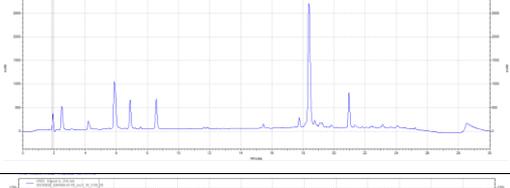
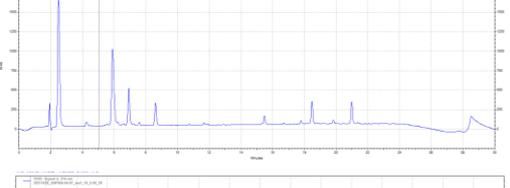
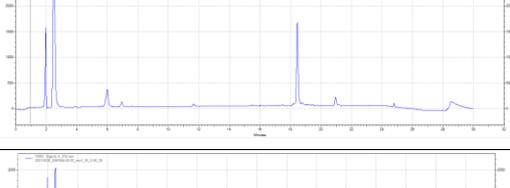
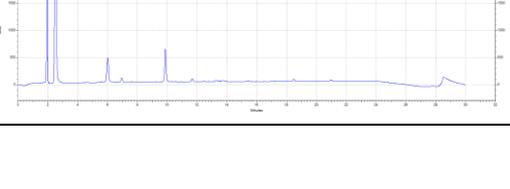
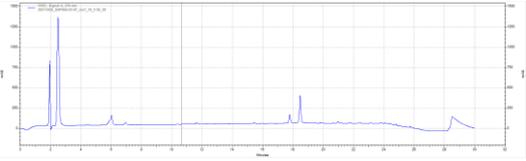
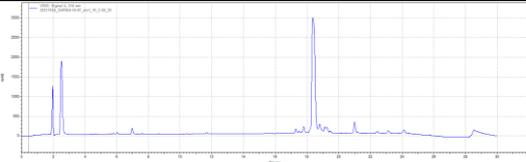
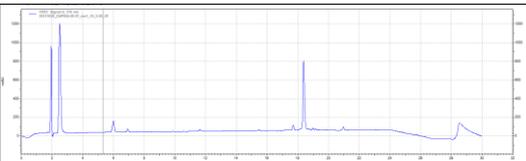
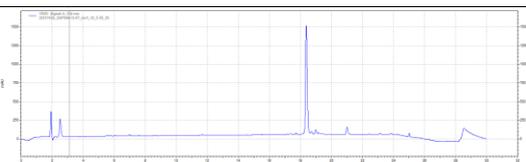
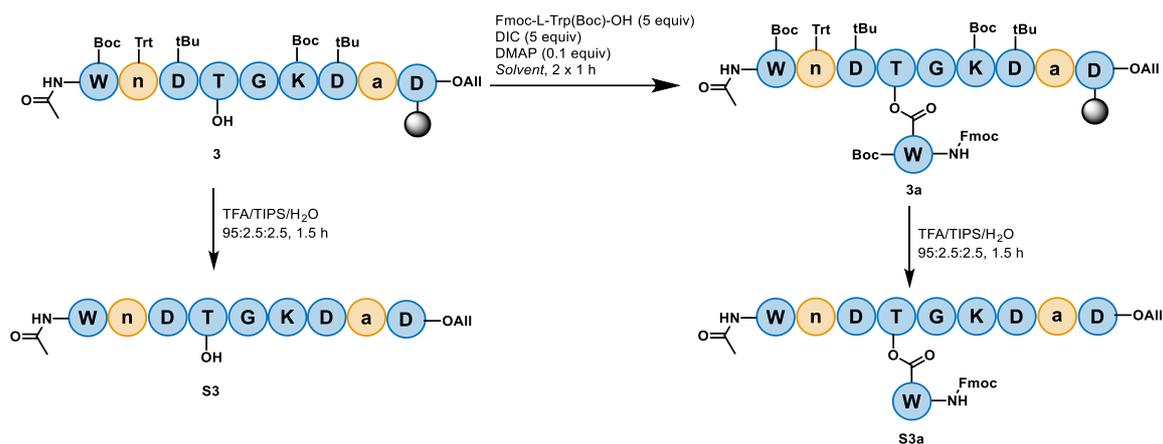
B		6.9 min	<p>[M+H]⁺ 960.2 [M+2H]²⁺ 480.8</p>	S2 +156
C	-	8.5 min	Identified by retention time against standard	6-ClHOBt (169)
D	Decays readily in aqueous media. Identified in crude MS and by disappearance of HPLC peak over time.	17.7 min		S2a+CO ₂ (1254.2)
E		17.1 min	<p>[M+H]⁺ 1171.2 [M+2H]²⁺ 586.2</p>	S2a-OAll (1171.2)
F		18.4 min	<p>[M+H]⁺ 1211.2 [M+2H]²⁺ 606.5</p>	S2a (1211.2)
G/H		20.8 min	[M+H] ⁺ 427.2	Fmoc-Trp-OH (426.5)

Table S5: HPLC data following reaction of peptidyl resin 2 with an appropriately protected tryptophan to give the corresponding depsipeptidyl resin as described in the Experimental section and depicted in Scheme S4. Entries 1-7 show data for Table 2 (entries 8-15), and entries 8-11 show data for Table 3 (entries 2 & 6). The conversion was determined as described in the Experimental section using the peaks of the linear (**S2**) and branched depsipeptide, and any corresponding adducts, specified below.

Entry	Tryptophan	Additive	Solvent	Depsipeptidyl resin	Conversion of 2 (% _{214 nm})	HPLC (5-50% B over 20 min, 214 nm)	R _t of peaks used for conversion calculation
1	Fmoc-L-Trp(Boc)-OH	-	DMF	2a	79.8%		S2 5.9 min S2a+CO₂ 17.8 min S2a 18.4 min
2	Fmoc-L-Trp(Boc)-OH	-	DCM	2a	98.2%		S2 6.0 min S2a 18.3 min
3	Fmoc-L-Trp(Boc)-OH	-	Dry DMF	2a	82.5%		S2 5.9 min S2a+CO₂ 17.8 min S2a 18.4 min
3	Fmoc-L-Trp(Boc)-OH	6ClHOBT	DMF	2a	10.1%		S2 6.0 min S2a+CO₂ 17.9 min S2a 18.6 min
4	Fmoc-L-Trp(Boc)-OH	6ClHOBT	DCM*	2a	57.9%		S2 5.9 min S2a+CO₂ 17.8 min S2a 18.4 min
5	Fmoc-L-Trp(Boc)-OH	6ClHOBT	Dry DMF	2a	17.1%		S2 5.9 min S2a 18.5 min
6	Fmoc-L-Trp(H)-OH	-	DMF	2b	67.6%		S2 6.0 min S2a 18.5 min
7	Boc-L-Trp(H)-OH	-	DMF	2c	51.2%		S2 6.0 min S2c 9.9 min

8	Fmoc-L-Trp(Boc)-OH	-	DMF	2a	62.5%		S2 6.0 min S2a+CO₂ 17.8 min S2a 18.5 min
9	Fmoc-L-Trp(Boc)-OH	-	DCM	2a	98.8%		S2 6.0 min S2a 18.4 min
10	Fmoc-D-Trp(Boc)-OH	-	DMF	2d	73.9%		S2 6.0 min S2a+CO₂ 17.7 min S2d 18.4 min
11	Fmoc-D-Trp(Boc)-OH	-	DCM	2d	99.0%		S2 6.0 min S2d 18.4 min

Peptidyl resin 3



Scheme S3: General synthetic scheme for reaction of peptidyl resin 3 (HMP-functionalised Tentagel®) with Fmoc-L-Trp(Boc)-OH for 2 x 1 h as described in the Experimental Section. The resulting reaction mixtures were characterised by RP-HPLC and MS and the products identified in Table S6. The experimental conditions were varied and the resulting RP-HPLC traces are summarised in Table S7.

Table S6: Additional peak characterisation. Representative trace following reaction of **3** with Fmoc-L-Trp(Boc)-OH, as shown in Scheme S3, illustrating key products and potential by-products as identified by ESI+ MS.

Peak	Extracted LCMS ESI+	R _t (5-50% B)	Observed m/z	Identification (Mw)
A		10.2 min	[M+H] ⁺ 1103.3 [M+2H] ²⁺ 552.3	S3 (1103.1)
B		10.5 min	[M+H] ⁺ 1146.3 [M+2H] ²⁺ 573.3	S3+CO₂ (1146.4)

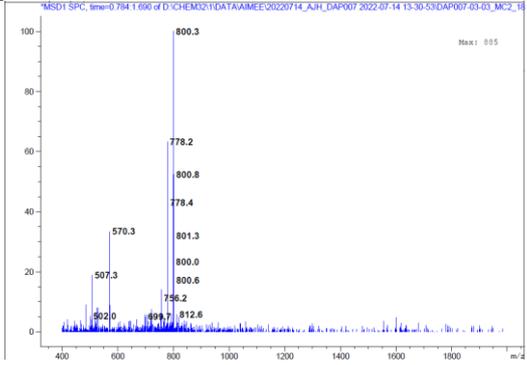
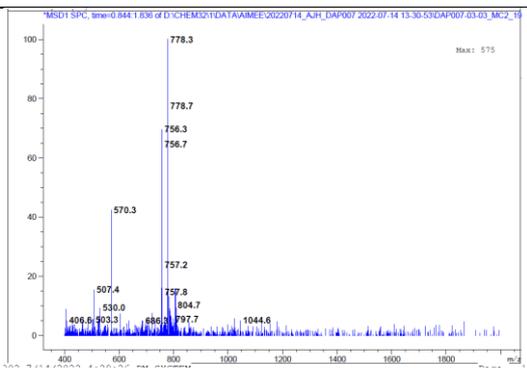
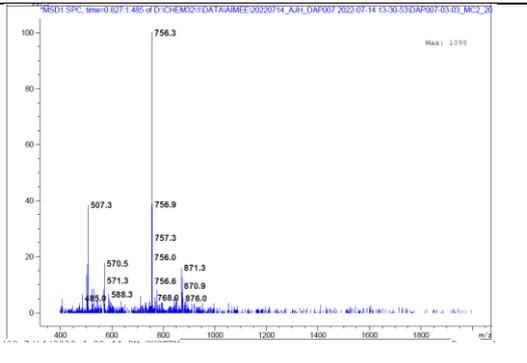
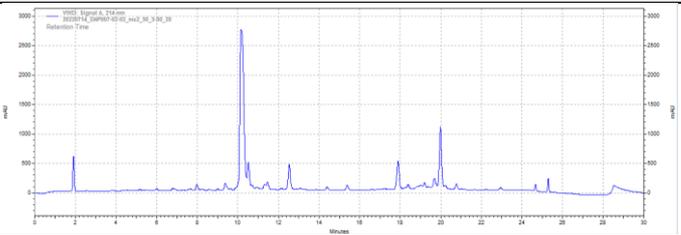
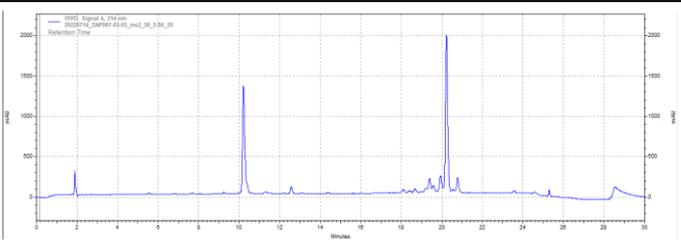
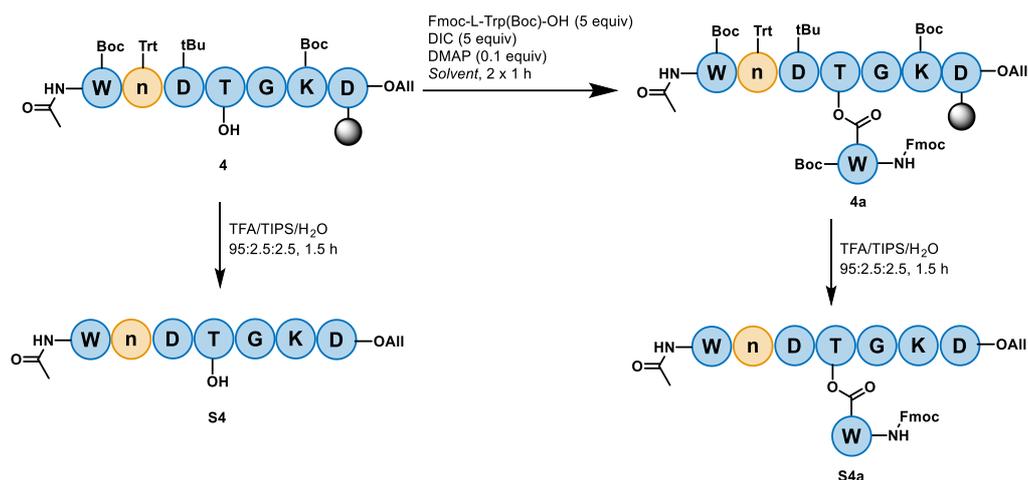
C		18.4 min	<p>$[M+2H]^{2+}$ 800.3</p> <p><i>LCMS overlap/in situ degradation of CO₂ adduct</i></p> <p>$[M+2H]^{2+}$ 778.3</p>	<p>S3a+2CO₂ (1597.6)</p> <p>S3a+CO₂ (1554.6)</p>
D / E		19.1 min / 19.3 min	<p>$[M+2H]^{2+}$ 778.3</p> <p><i>LCMS overlap/in situ degradation of CO₂ adduct</i></p> <p>$[M+2H]^{2+}$ 756.3</p> <p>$[M+3H]^{3+}$ 507.3</p>	<p>D/E) S3a+CO₂ (1554.6)</p> <p>Two stereoisomers of CO₂ adduct possible</p> <p>S3a</p>
F		19.9 min	<p>$[M+2H]^{2+}$ 756.3</p> <p>$[M+3H]^{3+}$ 507.3</p>	S3a (1511.6)

Table S7: HPLC data following reaction of peptidyl resin **3 with Fmoc-L-Trp(Boc)-OH to give depsipeptidyl resin **3a**, as described in the Experimental section and depicted in Scheme S3. The conversion of peptidyl resin **3** to **3a** was determined as described in the Experimental section using the peaks of the linear (**S3**) and branched depsipeptide (**S3a**), and any corresponding adducts, specified.**

Entry	Solvent	Conversion of 3 to 3a (% 214 nm)	HPLC (5-50%B over 20 min, 214 nm)	R_t of peaks used for conversion calculation
1	DMF	17.1%		<p>S3 10.2 min</p> <p>S3a 20.0 min</p>
2	DCM	50.6%		<p>S3 10.2 min</p> <p>S3a 20.2 min</p>

Peptidyl resin 4



Scheme S5: General synthetic scheme for reaction of peptidyl resin **4** (HMP-functionalised Tentagel®) with Fmoc-L-Trp(Boc)-OH for 2 x 1 h as described in the Experimental Section. The resulting reaction mixtures were characterised by RP-HPLC and MS and the products identified in Table S8. The experimental conditions were varied and the resulting RP-HPLC traces are summarised in Table S9.

Table S8: Additional peak characterisation. Representative trace following reaction of **2** with Fmoc-L-Trp(Boc)-OH, as shown in Scheme S5, illustrating key products and potential by-products as identified by ESI+ MS.

Peak	Extracted LCMS ESI+	R _t (5-50% B)	Observed m/z	Identification (Mw)
A		10.2 min	[M+H] ⁺ 917.3 [M+2H] ²⁺ 459.2 [M+H] ⁺ 899.2 [M+2H] ²⁺ 450.2	S4 (916.4) S4-H₂O (898.4) Aspartimide
B		10.4 min	[M+H] ⁺ 961.3 [M+2H] ²⁺ 481.3 [M+H] ⁺ 917.3 [M+2H] ²⁺ 459.2 [M+H] ⁺ 899.2	S4+CO₂ (960.0) CO ₂ adduct on tryptophan <i>Overlapping signal gives mixed MS spectrum</i> S4 (916.4) S4-H₂O (898.4)

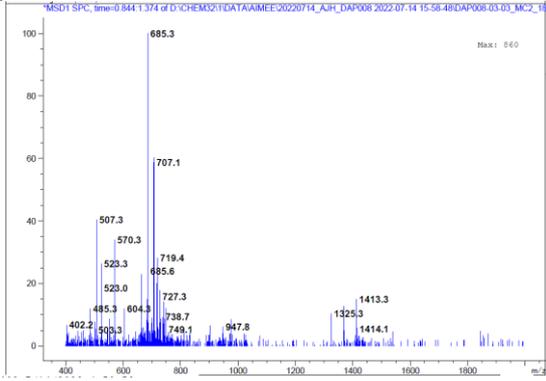
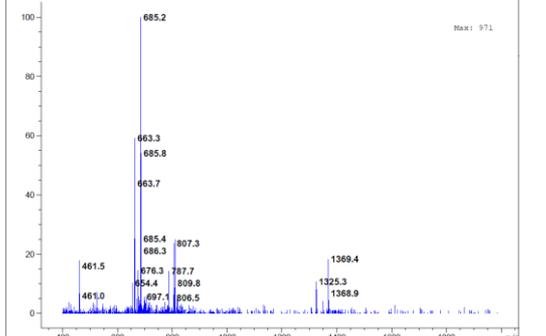
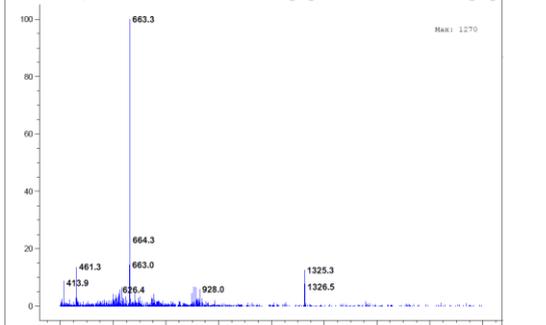
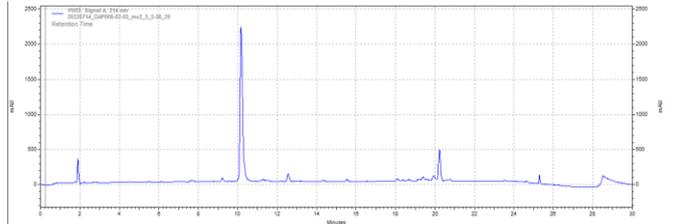
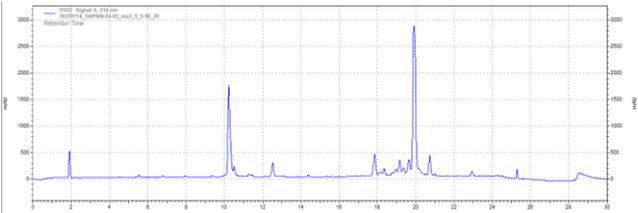
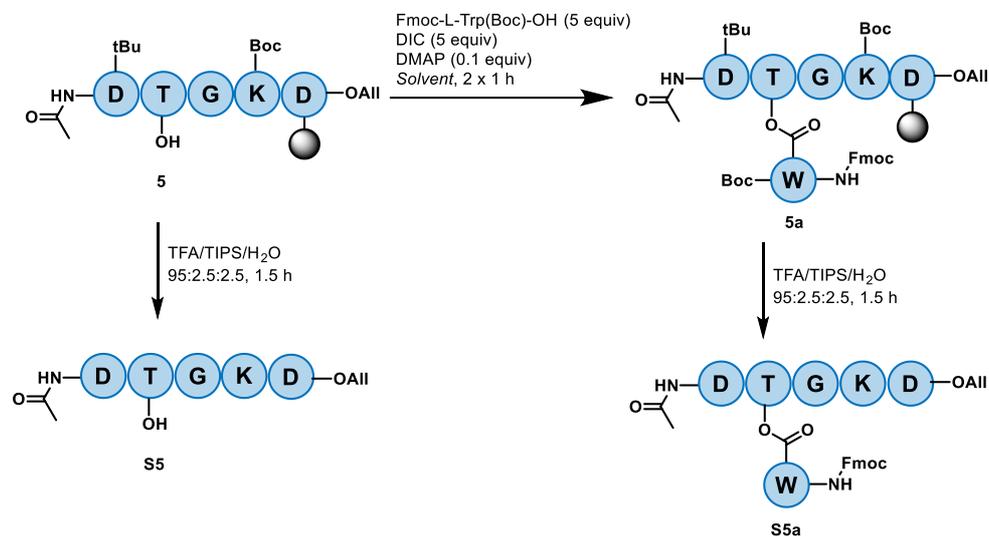
C		18.7 min	<p>[M+2H]²⁺ 450.2</p> <p>[M+H]⁺ 1413.3</p> <p>[M+2H]²⁺ 707.1</p> <p>[M+H]⁺ 1369.4</p> <p>[M+2H]²⁺ 685.3</p> <p>[M+H]⁺ 1325.3</p>	<p>Aspartimide</p> <p>S4a+2CO₂ (1411.4)</p> <p>CO₂ adduct on both tryptophans</p> <p><i>Overlapping MS signals & continual CO₂ adduct decay results in mixed MS spectrum</i></p> <p>S4a+CO₂ (1368.4)</p> <p>S4a (1325.4)</p>
D/E		19.3 / 19.5 min	<p>[M+H]⁺ 1369.4</p> <p>[M+2H]²⁺ 685.3</p> <p>[M+H]⁺ 1325.3</p> <p>[M+2H]²⁺ 663.3</p>	<p>D/E S4a+CO₂ (1368.4)</p> <p><i>Two stereoisomers of CO₂ adduct possible</i></p> <p><i>Overlapping MS signals</i></p> <p>S4a (1325.4)</p>
F		20.1 min	<p>[M+H]⁺ 1325.1</p> <p>[M+2H]²⁺ 663.3</p>	<p>S4a (1325.4)</p>

Table S9: HPLC data following reaction of peptidyl resin 4 with Fmoc-L-Trp(Boc)-OH to give depsipeptidyl resin 4a, as described in the Experimental section and depicted in Scheme S5. The conversion of peptidyl resin 4 to 4a was determined as described in the Experimental section using the peaks of the linear (S4) and branched depsipeptide (S4a), and any corresponding adducts, specified.

Entry	Solvent	Conversion of 4 to 4a (% _{214 nm})	HPLC (5-50% B over 20 min, 214 nm)	R _t of peaks used for conversion calculation
1	DMF	12.2%		<p>S4 10.2 min</p> <p>S4a 20.2 min</p>

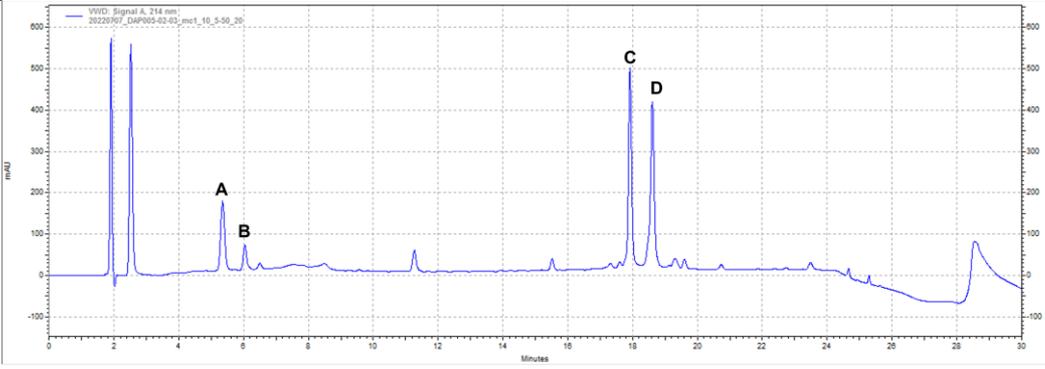
2	DCM	65.9%		S4 10.2 min S4a 19.9 min S4a+2CO₂ 19.4 min S4a+CO₂ 19.6 min
---	-----	-------	--	--

Peptidyl resin 5



Scheme S6: General synthetic scheme for reaction of peptidyl resin 5 (HMP-functionalised Tentagel®) with Fmoc-L-Trp(Boc)-OH for 2 x 1 h as described in the Experimental Section. The resulting reaction mixtures were characterised by RP-HPLC and MS and the products identified in Table S10. The experimental conditions were varied and the resulting RP-HPLC traces are summarised in Table S11.

Table S10: Additional peak characterisation. Representative trace following reaction of **5** with Fmoc-L-Trp(Boc)-OH, as shown in Scheme S6, illustrating key products and potential by-products as identified by ESI+ MS.

				
Peak	Extracted LCMS ESI+	R _t (5-50% B)	Observed m/z	Identification (Mw)
A		5.3 min	[M+H] ⁺ 617.3	S5 (616.6)

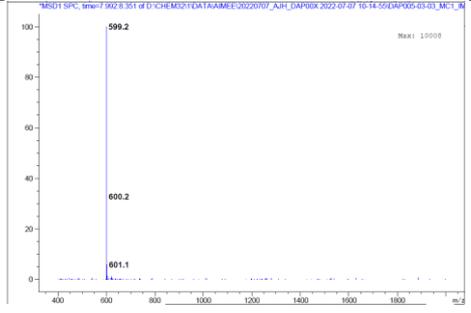
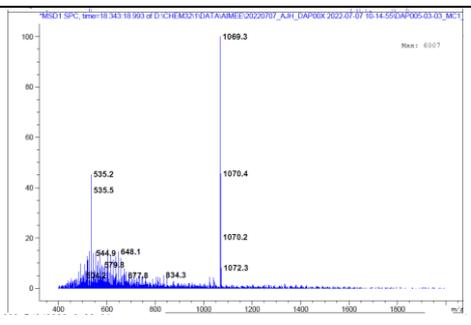
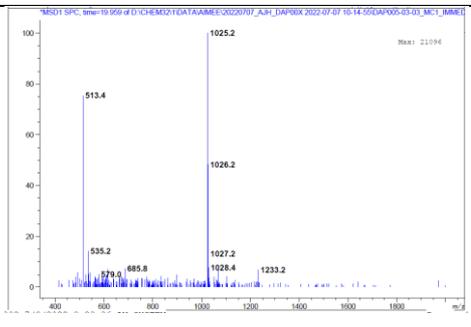
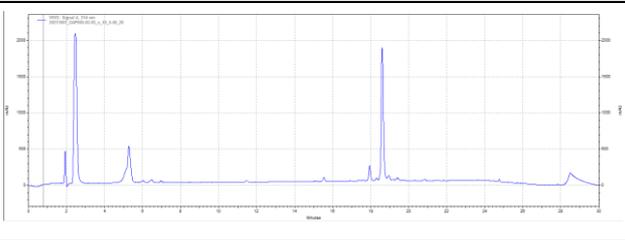
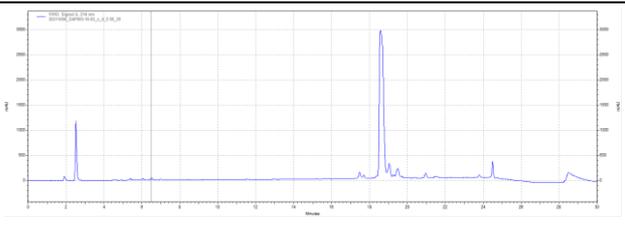
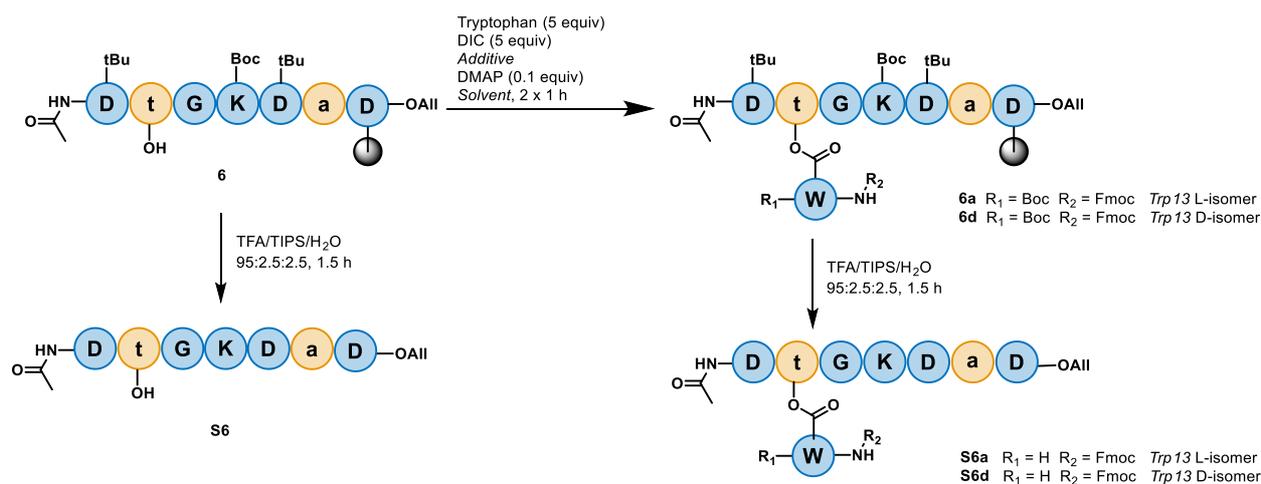
B		6.0 min	[M+H] ⁺ 599.2	S5-H₂O (598.6) Aspartimide
C		17.9 min	[M+H] ⁺ 1069.3 [M+2H] ²⁺ 535.2	S5a+CO₂ (1068.1) CO ₂ adduct on tryptophan
D		18.5 min	[M+H] ⁺ 1026.2 [M+2H] ²⁺ 513.4	S5a (1025.1)

Table S11: HPLC data following reaction of peptidyl resin 5 with Fmoc-L-Trp(Boc)-OH to give depsipeptidyl resin 5a, as described in the Experimental section and depicted in Scheme S6. The conversion of peptidyl resin 5 to 5a was determined as described in the Experimental section using the peaks of the linear (S5) and branched depsipeptide (S5a), and any corresponding adducts, specified..

Entry	Solvent	Conversion of 5 to 5a (% 214 nm)	HPLC (5-50% B, 214 nm)	R _t of peaks used for conversion calculation
1	DMF	74.2%		S5 5.3min S5a+CO₂ 17.8 min S5a 18.6 min
2	DCM	99.3%		S5 5.1 min S5a 18.6 min

Peptidyl resin 6



Scheme S7: General synthetic scheme for reaction of peptidyl resin 6 (HMP-functionalised Tentagel®) with Fmoc-Trp(Boc)-OH for 2 x 1 h as described in the Experimental Section. The resulting reaction mixtures were characterised by RP-HPLC and MS and the products identified in Table S12. The experimental conditions were varied and the resulting RP-HPLC traces are summarised in Table S13.

Table S12: Additional peak characterisation. Representative trace following reaction of **6** with Fmoc-L-Trp(Boc)-OH, as shown in Scheme S7, illustrating key products and potential by-products as identified by ESI+ MS.

Peak	Extracted LCMS ESI+	R _t (5-50% B)	Observed m/z	Identification (Mw)
A		5.9 min	[M+H] ⁺ 402.2 [M+2H] ²⁺ 803.2	S6 (802.8)
B		7.0 min	[M+H] ⁺ 960.3 [M+2H] ²⁺ 480.8 [M+H] ⁺ 785.2	S6+157 S6-H₂O (784.8) <i>Aspartimide</i>

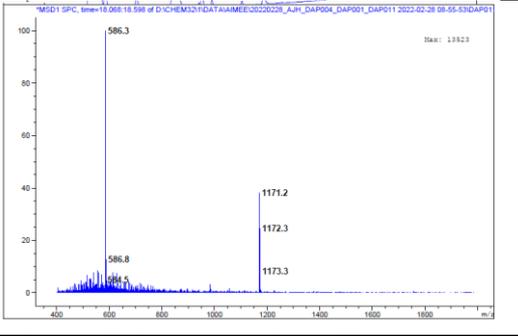
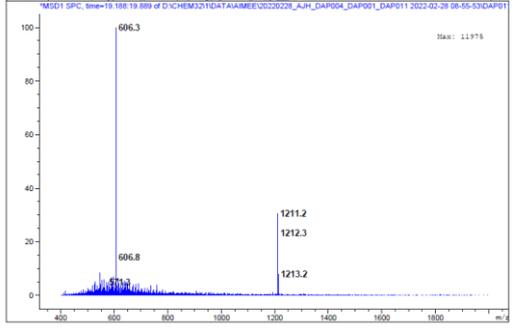
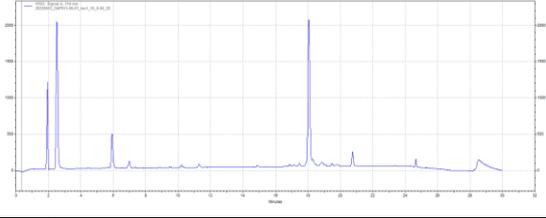
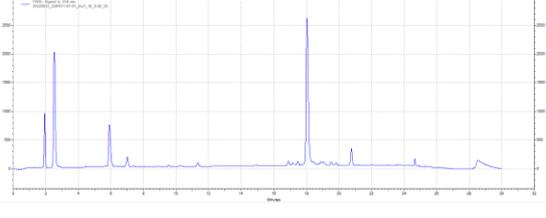
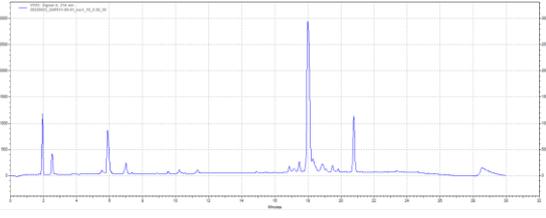
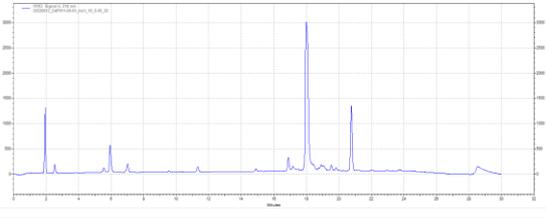
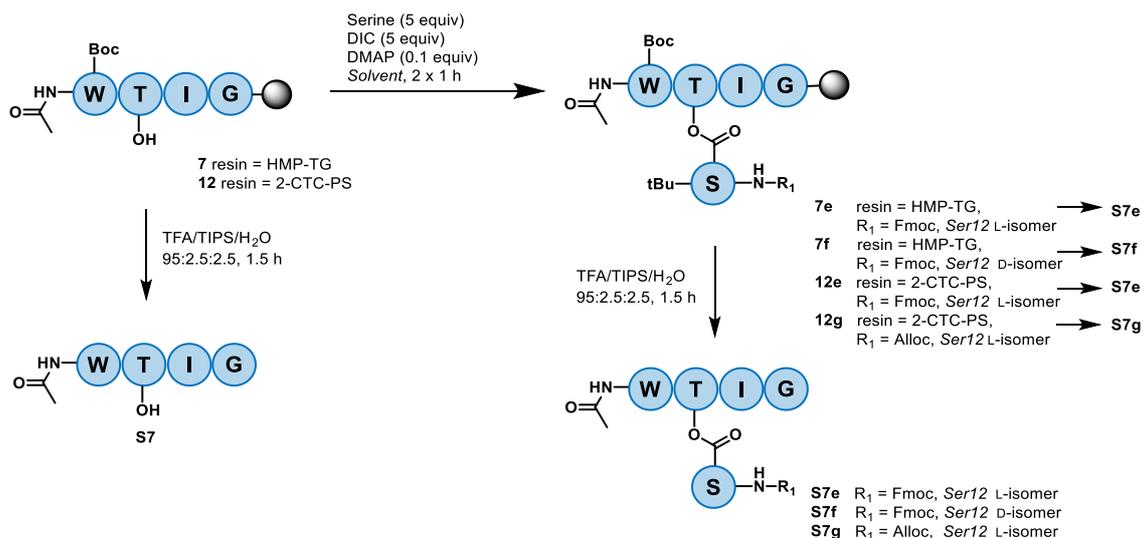
C		16.8 min	<p>[M+H]⁺ 1171.2 [M+2H]²⁺ 586.3</p>	<p>S6a-OAlI (1170.5) <i>Hydrolysis of allyl ester in solution</i></p>
D		18.0 min	<p>[M+H]⁺ 1211.1 [M+2H]²⁺ 606.3</p>	<p>S6a (1211.2)</p>

Table S13: HPLC data following reaction of peptidyl resin 6 with Fmoc-L-Trp(Boc)-OH to give depsipeptidyl resin 6a, or with Fmoc-D-Trp(Boc)-OH to give depsipeptidyl resin 6d, as described in the Experimental section and depicted in Scheme S7. The conversion of peptidyl resin 6 to 6a or 6d was determined as described in the Experimental section using the peaks of the linear (S6) and branched depsipeptide (S6a or S6d, respectively), and any corresponding adducts, specified.

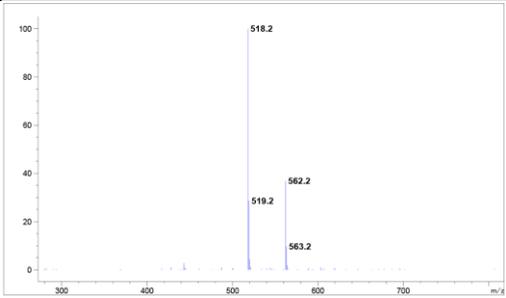
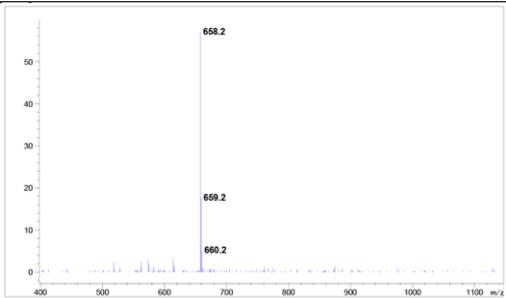
Entry	Fmoc-Trp(Boc)-OH	Solvent	Depsipeptidyl resin	Conversion of 6 (% _{214 nm})	HPLC (5-50% B over 20 min, 214 nm)	R _t of peaks used for conversion calculation
1	Fmoc-L-Trp(Boc)-OH	DMF	6a	81.6%		<p>S6 5.9 min S6a 18.1 min</p>
2	Fmoc-D-Trp(Boc)-OH	DMF	6d	78.2%		<p>S6 5.9 min S6d 18.0 min</p>
3	Fmoc-L-Trp(Boc)-OH	DCM	6a	80.9%		<p>S6 5.9 min S6a 18.0 min</p>
4	Fmoc-D-Trp(Boc)-OH	DCM	6d	87.2%		<p>S6 5.9 min S6d 18.0 min</p>

Peptidyl resins 7 & 12



Scheme S8: General synthetic scheme for reaction of peptidyl resins 7 (HMP-functionalised Tentagel®) & 12 (2-CITrt functionalised polystyrene) with appropriately protected serine (specified in Table S16) for 2 x 1 h as described in the Experimental Section. The resulting reaction mixtures were characterised by RP-HPLC and MS and the products identified in Table S14. The experimental conditions were varied and the resulting RP-HPLC traces are summarised in Table S15.

Table S14: Additional peak characterisation. Representative trace following reaction of 7, 12 or 13 with Fmoc-Ser(*t*Bu)-OH (in blue) and Alloc-L-Ser(*t*Bu)-OH (in green) as shown in Scheme S8, or Scheme 12. Key products **S7e-g** are identified by ESI+ MS.

Peak	Extracted LCMS ESI+	R_t (5-50% B)	Observed m/z	Identification (Mw)
A/B		11.5 min / 11.7 min	[M+H] ⁺ 518.2 [M+H] ⁺ 562.2	Overlapping signals A) S7 (517.6) B) S7+CO₂ (561.6)
C		15.4 min	[2M+H] ⁺ 658.2	Fmoc-Ser-OH (327.1)

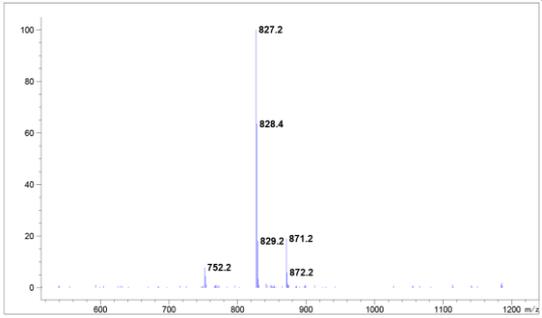
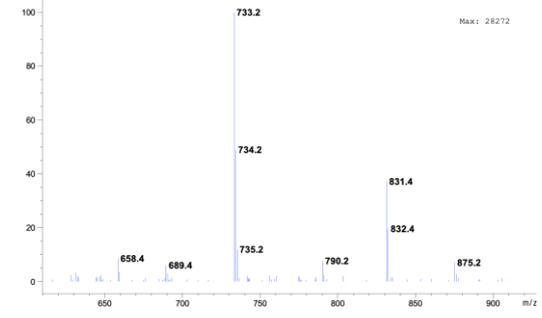
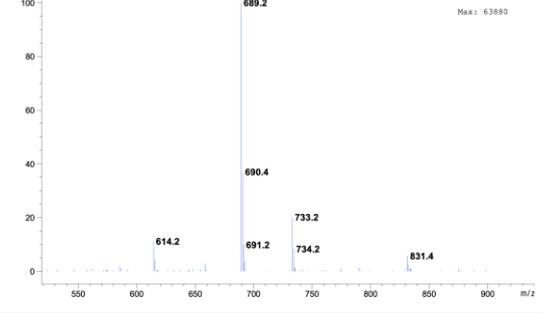
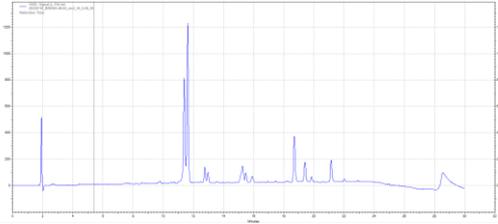
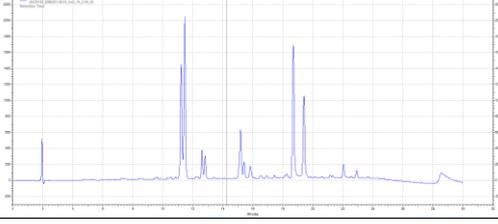
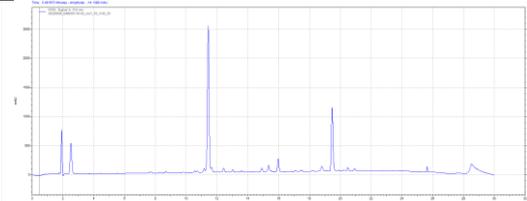
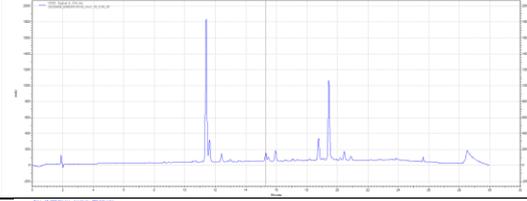
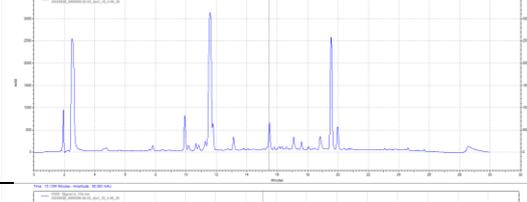
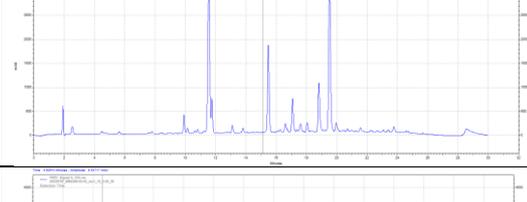
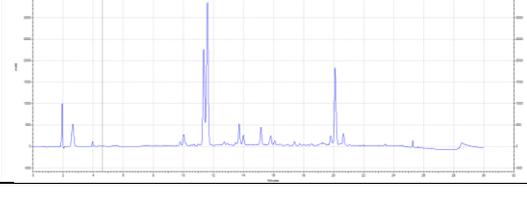
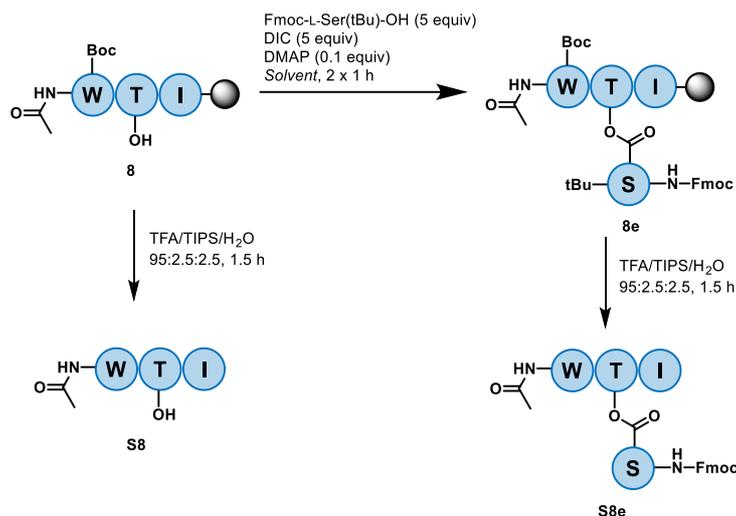
D/E		18.8min/ 19.5 min	[M+H] ⁺ 827.2 [M+H] ⁺ 871.2	Overlapping signals D) S7e (826.9) E) S7e+CO₂ (870.9)
F		19.8 min	[M+H] ⁺ 733.2	S12g+CO₂ (732.7)
G		20.1 min	[M+H] ⁺ 689.2	S12g (688.7)

Table S15: HPLC data following reaction of peptidyl resin 7 or 12 with appropriately protected serine to give the corresponding depsipeptidyl resin as described in the Experimental section and depicted in Scheme S8. The conversion was determined as described in the Experimental section using the peaks of the linear (S7**) and corresponding branched depsipeptide and any corresponding adducts as specified below.**

Entry	Peptidyl resin	Solvent	Serine	Depsipeptidyl resin	Conversion of peptidyl resin (%; 214 nm)	HPLC (5-50% B over 20 min, 214 nm)	R _t of peaks used for conversion calculation
1	7	DMF	Fmoc-L-Ser(<i>t</i> Bu)-OH	7e	20.2%		S7 11.5 min S7+CO₂ 11.7 min S7e+CO₂ 18.8 min S7e 19.5 min
2	7	DCM	Fmoc-L-Ser(<i>t</i> Bu)-OH	7e	47.4%		S7 11.5 min S7+CO₂ 11.7 min S7e+CO₂ 18.8 min S7e 19.5 min

3	7	DMF	Fmoc-D-Ser(tBu)-OH	7f	29.9%		S7 11.4 min S7+CO₂ 11.6 min S7f+CO₂ 18.8 min S7f 19.5 min
4	7	DCM	Fmoc-D-Ser(tBu)-OH	7f	28.1%		S7 11.4 min S7+CO₂ 11.6 min S7f+CO₂ 18.8 min S7f 19.5 min
5	12	DMF	Fmoc-L-Ser(tBu)-OH	12e	26.1%		S7 11.6 min S7+CO₂ 11.8 min S7e+CO₂ 18.8 min S7e 19.5 min
6	12	DCM	Fmoc-L-Ser(tBu)-OH	12e	51.1%		S7 11.6 min S7+CO₂ 11.8 min S7e+CO₂ 18.8 min S7e 19.5 min
7	12	DCM	Alloc-L-Ser(tBu)-OH	12g	17.8%		S7 11.4 min S7+CO₂ 11.6 min S7g+CO₂ 19.8 min S7g 20.1 min

Peptidyl resin 8



Scheme S16: General synthetic scheme for reaction of peptidyl resin 8 with Fmoc-L-Ser(tBu)-OH for 2 x 1 h as described in the Experimental Section. The resulting reaction mixtures were characterised by RP-HPLC and MS and the products identified in Table S16. The experimental conditions were varied and the resulting RP-HPLC traces are summarised in Table S17.

Table S16: Additional peak characterisation. Representative trace following reaction of **8** with Fmoc-L-Ser(tBu)-OH, as shown in Scheme S9, illustrating key products and potential by-products as identified by ESI+ MS.

Peak	Extracted LCMS ESI+	R _t (5-50% B)	Observed m/z	Identification (Mw)
A		12.5 min	[M+H] ⁺ 461.2	S8 (460.5)
B		15.3 min	[M+H] ⁺ 328.0	Fmoc-Ser-OH (327.3)

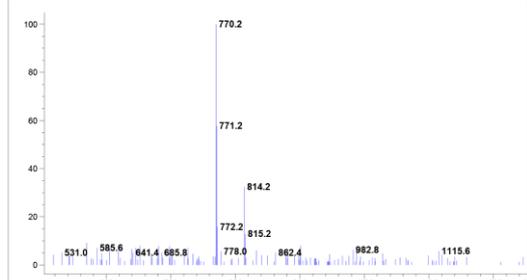
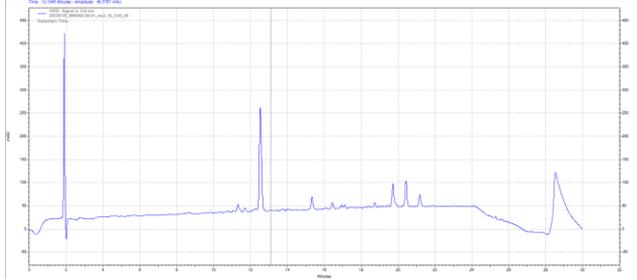
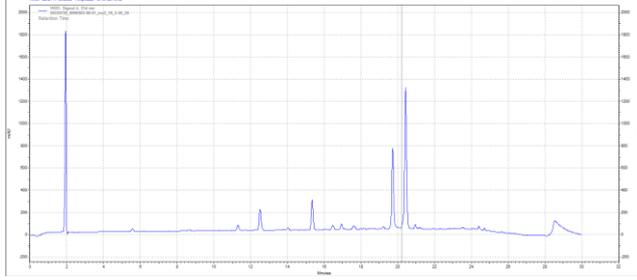
C/D		19.7 min/ 20.4 min	[M+H] ⁺ 814.0 [M+H] ⁺ 770.2	Overlapping signals C) S8e+CO₂ (812.8) D) S8e (769.9)
-----	---	-----------------------	--	--

Table S17: HPLC data following reaction of peptidyl resin **8 with Fmoc-L-Ser(*t*Bu)-OH to give depsipeptidyl resin **8e**, as described in the Experimental Section and depicted in Scheme S9. The conversion of peptidyl resin **8** to depsipeptidyl resin **8e** was determined as described in the Experimental Section using the peaks of the linear (**S8**) and branched depsipeptide (**S8e**), and any corresponding adducts, specified.**

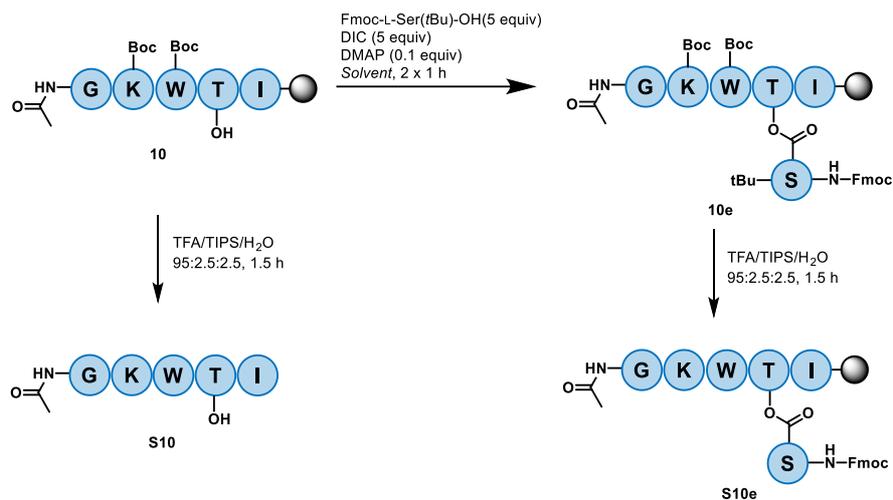
Entry	Solvent	Conversion of 8 to 8e (% _{214 nm})	HPLC (5-50% B over 20 min, 214 nm)	R _t of peaks used for conversion calculation
1	DMF	29.0%		S8 12.5 min S8e+CO₂ 19.7 min S8e 20.4 min
2	DCM	90.8%		S8 12.5 min S8e+CO₂ 19.7 min S8e 20.4 min

D		17.4 min	[M+H] ⁺ 1012.0	S9e (1011.5)
---	--	----------	---------------------------	--------------

Table S19: HPLC data following reaction of peptidyl resin **9 with Fmoc-L-Ser(*t*Bu)-OH to give depsipeptidyl resin **9e**, as described in the Experimental section and depicted in Scheme S10. The conversion of peptidyl resin **9** to depsipeptidyl resin **9e** was determined as described in the Experimental section using the peaks of the linear (**S9**) and branched depsipeptide (**S9e**), and any corresponding adducts, specified.**

Entry	Solvent	Conversion of 9 to 9e (% _{214 nm})	HPLC (5-50% B over 20 min, 214 nm)	R _t of peaks used for conversion calculation
1	DMF	2.7%		S9 10.2 min S9+CO₂ 10.6 min S9e 17.4 min
2	DCM	13.6%		S9 10.2 min S9+CO₂ 10.7 min S9e 17.4 min

Peptidyl resin 10

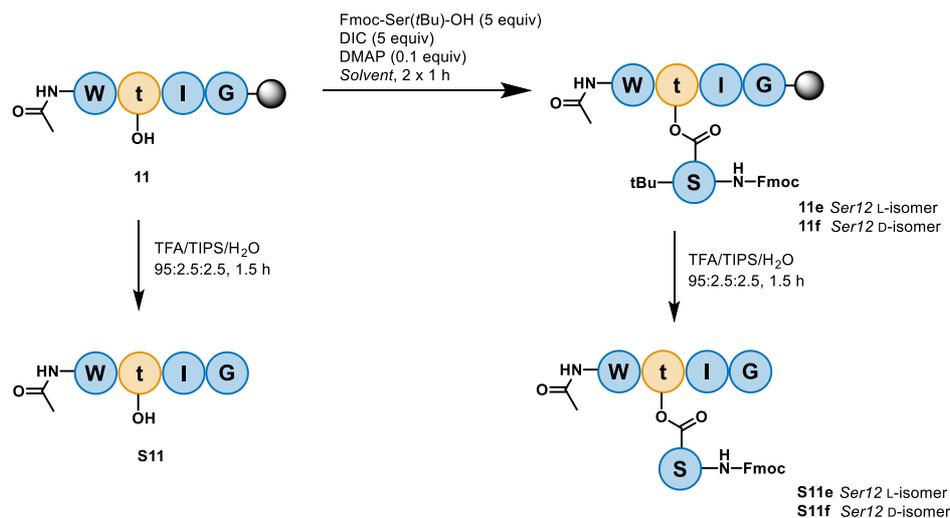


Scheme S11: General synthetic scheme for reaction of peptidyl resin 10 with Fmoc-L-Ser(tBu)-OH for 2 x 1 h as described in the Experimental Section. The resulting reaction mixtures were characterised by RP-HPLC and MS and the products identified in Table S20. The experimental conditions were varied and the resulting RP-HPLC traces are summarised in Table S21.

Table S20: Additional peak characterisation. Representative trace following reaction of **10** with Fmoc-L-Ser(tBu)-OH, as shown in Scheme S11, illustrating key products and potential by-products as identified by ESI+ MS.

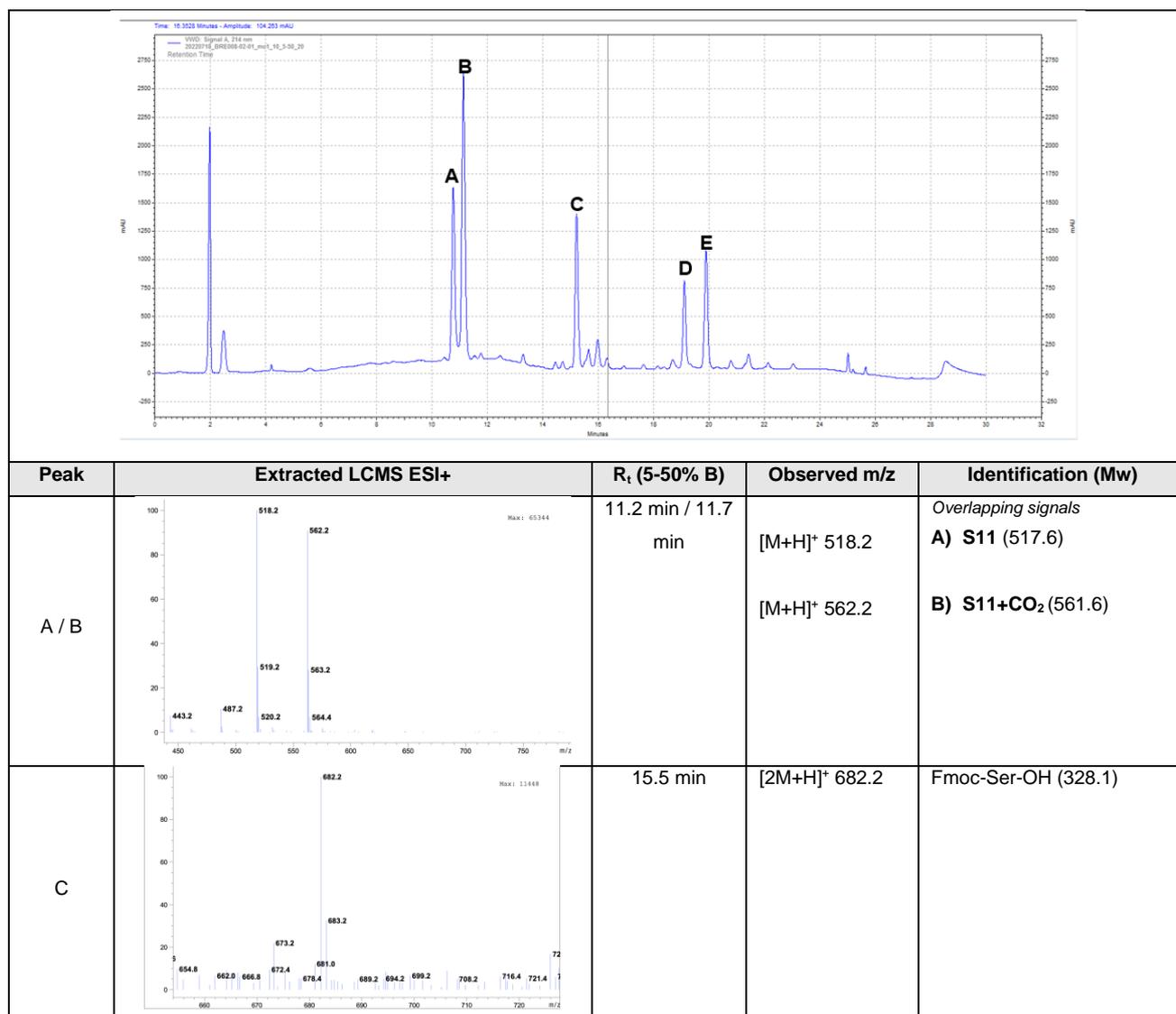
Peak	Extracted LCMS ESI+	R _t (5-50% B)	Observed m/z	Identification
A/B		11.2 min/ 11.6 min	[M+H] ⁺ 646.4 [M+H] ⁺ 690.2	Overlapping signals A) S10 (645.8) B) S10+CO₂ (689.8)
C		15.3 min	[M+H] ⁺ 328.3	Fmoc-Ser-OH (327.1)

Peptidyl resin 11



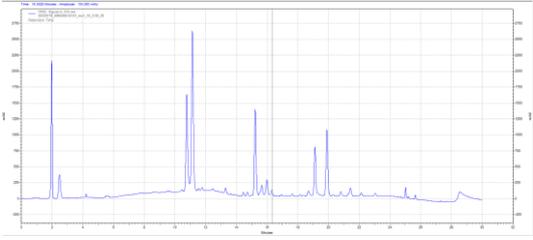
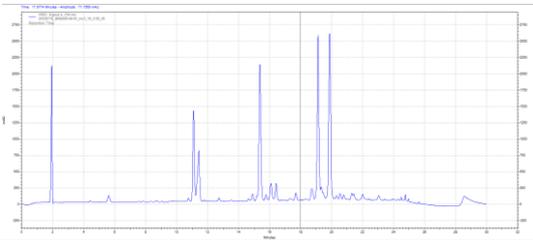
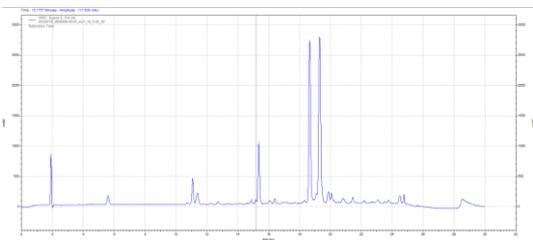
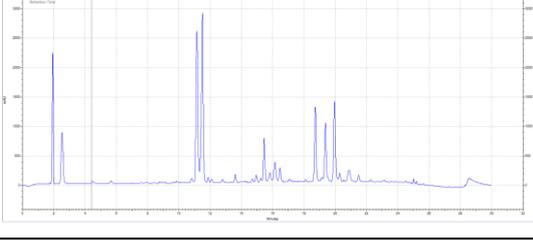
Scheme S14: General synthetic scheme for reaction of peptidyl resin 11 with Fmoc-L-Ser(*t*Bu)-OH or Fmoc-D-Ser(*t*Bu)-OH for 2 x 1 h as described in the Experimental Section. The resulting reaction mixtures were characterised by RP-HPLC and MS and the products identified in Table S22. The experimental conditions were varied and the resulting RP-HPLC traces are summarised in Table S23.

Table S22: Additional peak characterisation. Representative trace following reaction of 11 with Fmoc-Ser(*t*Bu)-OH, as shown in Scheme S14, illustrating key products and potential by-products as identified by ESI+ MS.

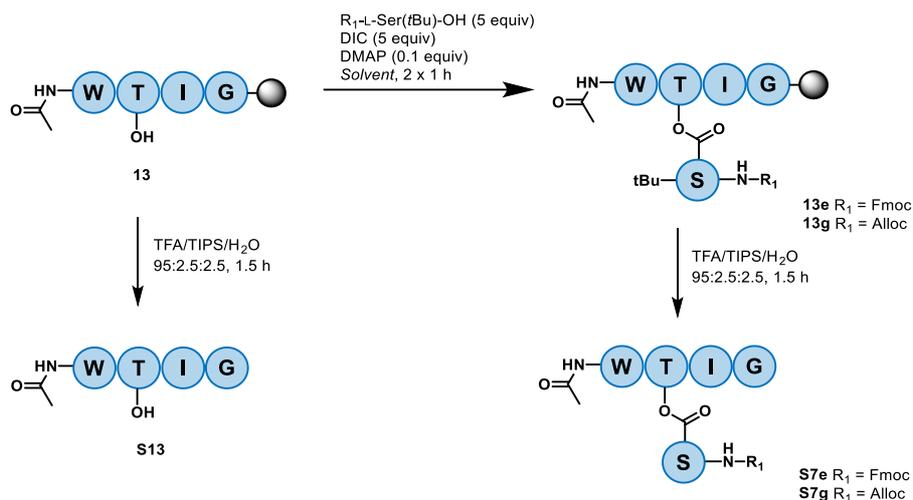


D	<p>Mass spectrum for peak D at 19.1 min. The x-axis is m/z from 750 to 1000, and the y-axis is relative intensity from 0 to 100. The base peak is at m/z 871.2. Other significant peaks are at m/z 872.2, 873.2, 874.2, and 797.0. The maximum intensity is 44640.</p>	19.1 min	[M+H] ⁺ 871.2	S7e+CO₂ (870.9)
E	<p>Mass spectrum for peak E at 19.9 min. The x-axis is m/z from 750 to 1000, and the y-axis is relative intensity from 0 to 100. The base peak is at m/z 827.2. Other significant peaks are at m/z 828.2, 829.4, 871.2, and 872.2. The maximum intensity is 29000.</p>	19.9 min	[M+H] ⁺ 827.2	S7e (826.9)

Table S23: HPLC data following reaction of peptidyl resin 11 with Fmoc-L-Ser(*t*Bu)-OH to give depsipeptidyl resin 11e; or with Fmoc-D-Ser(*t*Bu)-OH to give depsipeptidyl resin 11f, as described in the Experimental section and depicted in Scheme S14. The conversion of peptidyl resin 11 to depsipeptidyl resin 11e or 11f was determined as described in the Experimental section using the peaks of the linear (S11) and branched depsipeptide (S11e or S11f, respectively), and any corresponding adducts, specified.

Entry	Serine	Solvent	Conversion of 11 (% _{214 nm})	Depsipeptidyl resin	HPLC (5-50% B over 20 min, 214 nm)	Peaks used for conversion
1	Fmoc-L-Ser(<i>t</i> Bu)-OH	DMF	31.1%	11e		S11 10.8 min S11+CO ₂ 11.1 min S11e+CO ₂ 19.1 min S11e 19.9 min
2	Fmoc-L-Ser(<i>t</i> Bu)-OH	DCM	62.7%	11e		S11 11.1 min S11+CO ₂ 11.4 min S11e+CO ₂ 19.1 min S11e 19.9 min
3	Fmoc-D-Ser(<i>t</i> Bu)-OH	DMF	31.0%	11f		S11 11.2min S11+CO ₂ 11.5 min S11f+CO ₂ 19.4 min S11f 19.9 min
4	Fmoc-D-Ser(<i>t</i> Bu)-OH	DCM	90.9%	11f		S11 11.1 min S11+CO ₂ 11.4 min S11f+CO ₂ 18.7 min S11f 19.3 min

Peptidyl resin 13

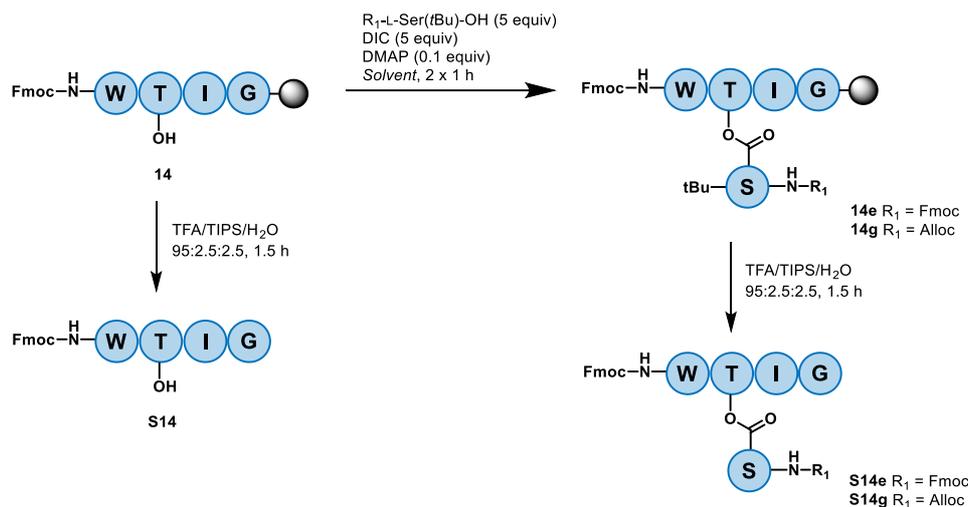


Scheme S12: General synthetic scheme for reaction of peptidyl resin 13 with Fmoc-L-Ser(*t*Bu)-OH or Alloc-L-Ser(*t*Bu)-OH for 2 x 1 h as described in the Experimental Section. The resulting reaction mixtures were characterised by RP-HPLC and MS and the products identified in Table S14. The experimental conditions were varied and the resulting RP-HPLC traces are summarised in Table S24.

Table S24: HPLC data following reaction of peptidyl resin 13 with Fmoc-L-Ser(*t*Bu)-OH to give depsipeptidyl resin **13e**; or with Alloc-L-Ser(*t*Bu)-OH to give depsipeptidyl resin **13g**, as described in the Experimental section and depicted in Scheme S12. The conversion of peptidyl resin **13** to depsipeptidyl resin **13e** or **13g** was determined as described in the Experimental section using the peaks of the linear (**S7**) and branched depsipeptide (**S7e** or **S7g**, respectively), and any corresponding adducts, specified.

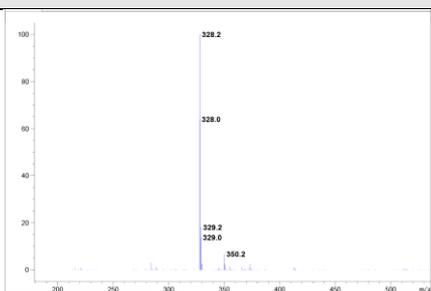
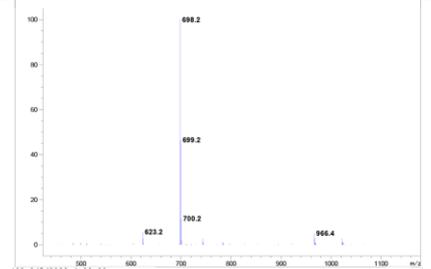
Entry	Solvent	Serine	Deptsipeptidyl resin	Conversion of 13 (% _{214 nm})	HPLC (5-50% B over 20 min, 214 nm)	R_t of peaks used for conversion calculation
1	DMF	Fmoc-L-Ser(<i>t</i> Bu)-OH	13e	30.9%		S7 11.6 min S7e 19.6 min
2	DCM	Fmoc-L-Ser(<i>t</i> Bu)-OH	13e	33.6%		S7 11.6 min S7e 19.6 min
3	DCM	Alloc-L-Ser(<i>t</i> Bu)-OH	13g	11.8%		S7 11.4 min S7g 16.5 min

Peptidyl resin 14



Scheme S13: General synthetic scheme for reaction of peptidyl resin 14 with Fmoc-L-Ser(*t*Bu)-OH or Alloc-L-Ser(*t*Bu)-OH for 2 x 1 h as described in the Experimental Section. The resulting reaction mixtures were characterised by RP-HPLC and MS and the products identified in Table S25. The experimental conditions were varied and the resulting RP-HPLC traces are summarised in Table S26.

Table S25: Additional peak characterisation. Representative trace following reaction of 14 with Fmoc-L-Ser(*t*Bu)-OH, as shown in Scheme S13, illustrating key products and potential by-products as identified by ESI+ MS.

Peak	Extracted LCMS ESI+	R_t (5-95% B)	Observed m/z	Identification
A		13.0 min	$[\text{M}+\text{H}]^+$ 328.2	Fmoc-Ser-OH (327.1)
B		16.4 min	$[\text{M}+\text{H}]^+$ 698.2	S14 (697.8)

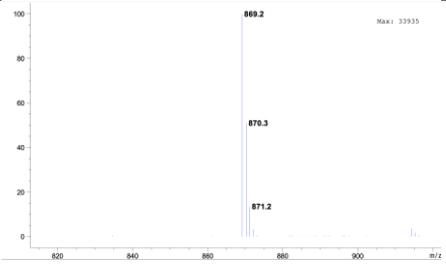
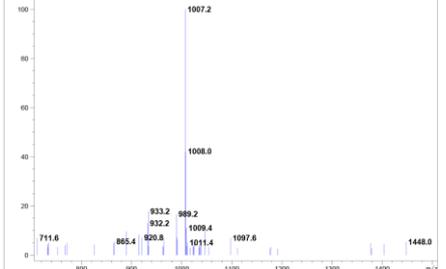
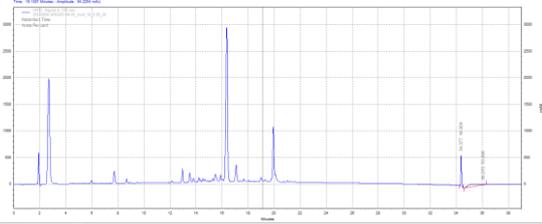
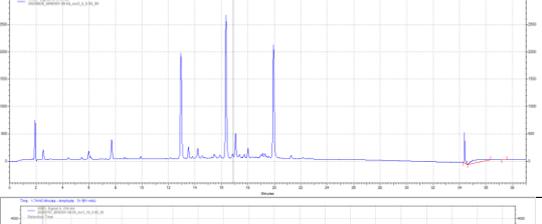
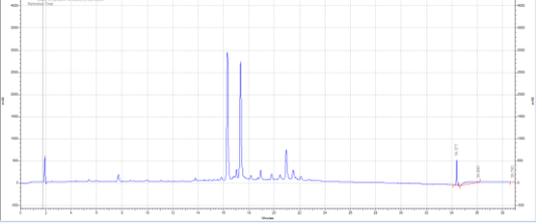
C		17.4 min	[M+H] ⁺ 869.2	S14g (868.9)
D		19.9 min	[M+H] ⁺ 1007.2	S14e (1007.1)

Table S26: HPLC data following reaction of peptidyl resin 14 with Fmoc-L-Ser(*t*Bu)-OH to give depsipeptidyl resin 14e; or with Alloc-L-Ser(*t*Bu)-OH to give depsipeptidyl resin 14g, as described in the Experimental section and depicted in Scheme S13. The conversion of peptidyl resin 14 to depsipeptidyl resin 14e or 14g was determined as described in the Experimental section using the peaks of the linear (S12) and branched depsipeptide (S14e or S14g, respectively), and any corresponding adducts, specified.

Entry	Solvent	Serine	Depsipeptidyl resin	Conversion of 14 (% 214 nm)	HPLC (5-95% B, 3%B/min, 214 nm)	R _t of peaks used for conversion calculation
1	DMF	Fmoc-L-Ser(<i>t</i> Bu)-OH	14e	20.9%		S14 16.4 min S14e 19.9 min
2	DCM	Fmoc-L-Ser(<i>t</i> Bu)-OH	14e	62.3%		S14 16.4 min S14e 19.9 min
3	DCM	Alloc-L-Ser(<i>t</i> Bu)-OH	14g	22.4%		S14 16.3 min S14g 17.4 min

References

- (1) Martin-Gómez, H.; Albericio, F.; Tulla-Puche, J. A Lasso-Inspired Bicyclic Peptide: Synthesis, Structure and Properties. *Chem. – Eur. J.* **2018**, *24* (72), 19250–19257. <https://doi.org/10.1002/chem.201803899>.
- (2) Hermant, Y.; Palpal-Iatoc, D.; Kovalenko, N.; Cameron, A. J.; Brimble, M. A.; Harris, P. W. R. The Total Chemical Synthesis and Biological Evaluation of the Cationic Antimicrobial Peptides, Laterocidine and Brevicidine. *J. Nat. Prod.* **2021**, *84* (8), 2165–2174. <https://doi.org/10.1021/acs.jnatprod.1c00222>.
- (3) Eissler, S.; Kley, M.; Bächle, D.; Loidl, G.; Meier, T.; Samson, D. Substitution Determination of Fmoc-Substituted Resins at Different Wavelengths. *J. Pept. Sci.* **2017**, *23* (10), 757–762. <https://doi.org/10.1002/psc.3021>.

