Supporting information

Expanding the scope of $N \rightarrow S$ acyl transfer in native peptide sequences

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¹³C analysis data for thioester formation (Gly-Cys-OH)



Alpha-C-OH to MESNa Thioester

¹³C analysis data for thioester formation (Xaa-**D-Cys**-OH)



¹³C analysis data for thioester formation (Xaa-Cys-NH₂)



$^{\rm 13}{\rm C}$ analysis data for thioester formation (Xaa-D-Cys-NH_2)



¹³C analysis data for thioester formation at pH 1.8 (Xaa-Cys-OH)



¹³C analysis data for thioester formation at pH 3.3 (Xaa-Cys-OH)



¹³C analysis data for thioester formation at pH 3.8 (Xaa-Cys-OH)



Alpha-C-OH to MESNa Thioester at pH 3.8

¹³C analysis data for thioester formation at pH 4.0 (Xaa-Cys-OH)



 $^{\rm 13}{\rm C}$ analysis data for thioester formation at pH 4.3 (Xaa-Cys-OH)



LSC-206(0) 60 1 C:\Bruker\TOPSPIN 600 Alpha-C-OH to MESNa Thioester at pH 4.3

¹³C analysis data for thioester formation at pH 4.5 (Xaa-Cys-OH)



 $^{\rm 13}{\rm C}$ analysis data for thioester formation at pH 1.3 (Xaa-Cys-NH_2)



Alpha-C-NH2 to MESNa Thioester at pH 1.3

 $^{\rm 13}{\rm C}$ analysis data for thioester formation at pH 1.8 (Xaa-Cys-NH_2)



Alpha-C-NH2 to MESNa Thioester at pH 1.8

 $^{\rm 13}{\rm C}$ analysis data for thioester formation at pH 3.3 (Xaa-Cys-NH_2)



 $^{\rm 13}{\rm C}$ analysis data for thioester formation at pH 3.5 (Xaa-Cys-NH_2)



Alpha-C-NH2 to MESNa Thioester at pH 3.5 LSC-218(0) 60 1 C:\Bruker\TOPSPIN 600

 $^{\rm 13}{\rm C}$ analysis data for thioester formation at pH 3.8 (Xaa-Cys-NH_2)



Alpha-C-NH2 to MESNa Thioester at pH 3.8

 $^{\rm 13}{\rm C}$ analysis data for thioester formation at pH 4.5 (Xaa-Cys-NH_2)



Alpha-C-NH2 to MESNa Thioester at pH 4.5 LSC-222(0) 60 1 C:\Bruker\TOPSPIN 600

 $^{\rm 13}{\rm C}$ analysis data for thioester formation at pH 4.8 (Xaa-Cys-NH_2)



Summary of pH data:



HPLC Analysis of Thioester formation and cyclisation of Agardhipeptin analogues:



<u>GlyCys</u>









PheCys







Selected LC-MS data for Agardhipeptin thioester formation/Cyclisation studies

LeuCys: THIOESTER FORMATION









whole TIC @ t48



Leu-Cys: CYCLISATION

t0



starting material 9b



Product 12 b

CAhpLC t24 2014-09-30 194 (3.582) Sm (SG, 50x1.00); Cm (190:197) 1: Scan ES+ 100



PheCys

THIOESTER FORMATION









AhpFC t24 2014-10-01 171 (3.156) Sm (SG, 50x1.00); Cm (171:178) 1: Scan ES+ 100 → 2.97e7



combined TIC @ t48

AhpFC t48 2014-10-02 167 (3.082) Sm (SG, 50x1.00); Cm (157:190) 1: Scan ES+ 1007 7.27e6



CYCLISATION





Product 12e

CAhpFC t24 2014-10-01 199 (3.673) Sm (SG, 50x1.00); Cm (197:205) 1: Scan ES+ 100 → 914.82 3.77e7



GlyCys: THIOESTER FORMATION









CYCLISATION



starting material 9a



Product 12a

CAhpGC t24 2014-09-30 162 (2.990) Sm (SG, 50x1.00); Cm (157:166) 1: Scan ES+ 100_□ 5.40e7



SFTI-1 synthesis



mass spectra of linear SFTI-1 precursor (A), cyclised (B) and cyclised/ oxidised (C) SFTI-1.



Analytical HPLC of cyclisation reactions performed at 60 $^{\circ}$ C (**A**) and 50 $^{\circ}$ C (**B**).



Separation of unmodified and aspartimide-modified cyclic SFTI-1 using preparative RP-HPLC.



A) overlaid ${}^{1}H^{-}H$ NOESY and ${}^{1}H^{-}H$ TOCSY spectra of wild-type SFTI-1 showing sequential assignment of backbone amide chemical shifts. **B)** Annotated ${}^{1}H^{-}H$ TOCSY spectrum of wild-type SFTI-1 showing complete side-chain resonances for each amino acid.