

Structure of Sch 218157, a Cyclodepsipeptide with Neurokinin Activity

M. CHU, T. M. CHAN, P. DAS, R. MIERZWA,
M. PATEL and M. S. PUAR*

Schering-Plough Research Institute,
2015 Galloping Hill Road, Kenilworth,
New Jersey 07033-0539, USA

(Received for publication March 10, 2000)

Neurokinins (NKs), also known as tachykinins (TKs), are members of a family of 9~11 amino acid peptides which are involved in various pathological conditions including inflammation^{1,2}, pain transmission, cancer, anxiety, asthma, and vasodilation³. The peptides act mainly through specific 7-transmembrane G-protein-coupled receptor domains named NK₁, NK₂, and NK₃. In the course of our screening program for novel neurokinin receptor inhibitors, we have isolated a cyclodepsipeptide (**1**) from an unidentified fungal fermentation culture broth (MYCO-2838)⁴ with selective NK₂ antagonist activity.

The fermentation broth was extracted with EtOAc at harvest (pH ~6.5). The crude extract was purified by high speed centrifugal partition chromatography (CPC) with solvent system of hexane:EtOAc:MeOH:H₂O (3:5:3:5, v/v/v/v). The combined bioactive fractions were further purified by normal phase HPLC (semi-preparative YMC PVA-Sil column, 20×250 mm, S-5, 5~25% MeOH in CH₂Cl₂ with a linear gradient in 30 minutes, UV=218 nm,

12 ml/minute). Pure compound **1** was obtained as a white/pale yellow solid.

The molecular formula was established as C₅₇H₈₉N₁₁O₁₃ by HRFABMS [(M+H)⁺ at *m/z* 1136.6720, calcd.; 1136.6740, measured] and ¹H and ¹³C NMR data (Table 1) were indicative of a peptide with a blocked *N*-terminus (ninhydrin-negative).

Amino acid analysis indicated the presence of one mole each of glycine, valine, D-phenylalanine, proline, and D,L-isoleucine. The CIMS of the acid hydrolysis products supported the presence of the above amino acids and also indicated the presence of pipercolic acid (*m/z* 130) and *N*-methylglutamine (a strong *m/z* 144 (M+1)⁺; *N*-methylglutamic acid anhydride).

Extensive analysis of the COSY, HMBC, HMQC, HMQC-TOCSY NMR data revealed the spin systems of the amino acids; Phe, Pro, Gly, MeVal, two MeGln, Ile, Pip (pipercolic acid), and Val. In addition an acid containing component was assigned as 2-hydroxy-3-methylpentanoic acid (HMP).

The proton NMR data in CDCl₃ and DMSO-*d*₆ (25° and 50°C) indicated the presence of an extra pair of amide protons at δ 7.30 and 6.80 when compared with the data of Sch 217048⁵. Further analysis indicated an upfield shift of ~0.2 ppm for the γCH₂ of MeGlu. Similarly minor changes were observed for the carbon-13 chemical shifts around the MeGlu moiety. The NMR data suggested that the MeGlu moiety in Sch 217048 is transformed to MeGln (MeGlu→MeGln). Thus compound **1** contains two units of MeGln as shown in Figure 1. Attempts to hydrolyze the ester bond under basic condition failed. The rationale behind this failure is the lack of δCOO⁻ group in MeGln that was

Fig. 1. Sch 218157

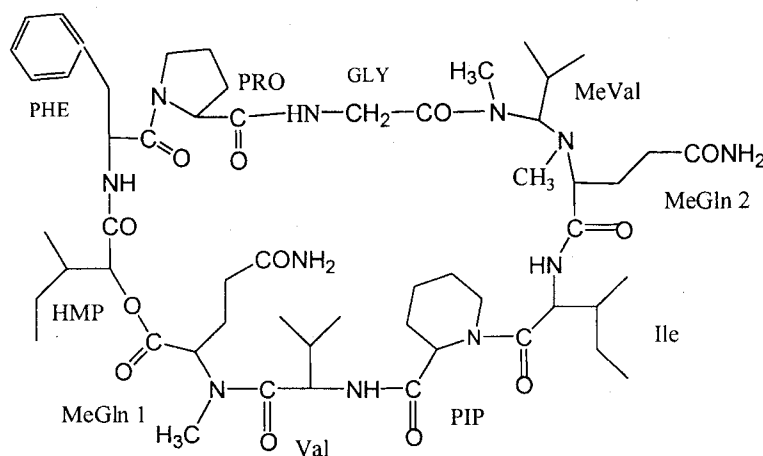


Table 1. ^1H and ^{13}C NMR data for Sch 218157 in DMSO- d_6 ^a.

AA		^{13}C ^b	^1H (mult, J in Hz) ^b	AA		^{13}C ^b	^1H (mult, J in Hz) ^b
MeGln 1	CO	169.5		MeVal	CO	169.7	
	α	62.0	4.13 (dd, 9.0, 4.0)		α	57.0	5.14 (d, 10.0)
	β	23.9	2.25 m		β	27.3	2.29 m
	γ	30.7	2.15 m		γ -Me	17.9	0.75 (d, 6.5)
	δ -CONH ₂	173.4			γ -Me	19.1	0.83 (d, 6.5)
	NMe	38.5	3.22 s		NMe	28.1	2.90 s
Val	CO	172.2		Gly	CO	170.4	
	α	54.2	4.57 (t, 8.0, 9.0)		α	41.1	4.40 (dd, 17.0, 8.0)
	β	31.0	1.95 m				4.24 (d, 17.0)
	γ -Me	17.7	0.80 (d, 6.5)		Pro	CO	170.6
	γ -Me	19.2	0.82 (d, 6.5)	α		59.4	4.55 (dd, 8.0, 5.0)
		NH		8.65 (d, 8.0)	β	29.1	2.12 m, 1.74 m
Pip	CO	170.2		γ	24.7	2.05 m, 1.92 m	
	α	52.4	5.12 (dd, 4.0, 2.5)	δ	47.0	3.70 m	
	β	27.0	1.75 m	Phe	CO	169.7	
	γ	19.4	1.41 m, 1.15 m		α	52.2	4.69 (dt, 8.0, 8.0, 4.0)
	δ	24.4	1.72 m		β	36.3	2.93 m
	ϵ	43.1	3.81 m, 3.58 m		γ -C ₁	137.4	
Ile	CO	170.2		-C ₂ , C ₆	129.2	7.32 m	
	α	52.5	4.84 (dd, 8.0, 3.0)	-C ₃ , C ₅	128.4	7.22 m	
	β	36.1	1.76 m	-C ₄	126.4	7.22 m	
	γ -Me	16.0	0.86 (d, 6.5)	NH		7.60 (d, 8.0)	
	γ -CH ₂	22.3	1.26 m	HMP	CO	168.0	
	δ -Me	11.3	0.77 (d, 6.5)		α	74.4	5.00 (d, 1.0)
	NH		6.22 (d, 8.0)		β	35.6	1.95 m
MeGln 2	CO	168.4		γ -Me	13.8	0.65 (d, 6.5)	
	α	58.9	4.88 (dd, 8.0, 3.0)	γ -CH ₂	25.3	1.20 m	
	β	23.8	2.05 m, 1.75 m	δ -Me	11.4	0.76 (d, 6.5)	
	γ	30.8	2.08 m, 1.95 m				
	δ -CONH ₂	173.1					
	δ -CONH ₂		7.29 bs, 6.80 bs				
	NMe	29.1	2.65 s				

a Instruments: Varian XL-400, GE-400.

b The chemical shifts are in ppm with reference to internal TMS and coupling constants, J, are in Hertz.

envisioned to participate intramolecularly in the hydrolysis of the ester bond.

Compound **1** showed selective binding to the NK₂ receptor with an IC₅₀ value of 68 nM and an IC₅₀ value was >1000 nM for the NK₁ receptor. These values are comparable to those reported for Sch 217048.

References

- 1) LOW, J. A., III. & R.M. SNIDER: The role of tachykinins in pulmonary diseases. *Ann. Rep. Med. Chem.* 28: 199~207, 1993
- 2) LONGMORE, J.; C. J. SWAIN & R. G. HILL: Neurokinin receptors. *Drug News and Perspect.* 8: 5~23, 1995

- 3) GERSPACHER, M. & A. von. SPRECHER: Dual neurokinin NK₁/NK₂ receptor antagonists. *Drugs of the Future* 24: 883~892, 1999
- 4) The microorganism was supplied by Dr. B. KATZ from MYCOsearch Lab. Sch 218157 is not a co-product of Sch 217084 which was produced by a different microorganism, namely MYCO-2475
- 5) HEGDE, V. R.; M. S. PUAR, T. M. CHAN, P. DAI, P. R. DAS & M. PATEL: Sch 217048: A novel cyclodepsipeptide with neurokinin antagonist activity. *J. Org. Chem.*, 63: 9584~9586, 1998.