A Simple & Convenient Solid Phase Synthesis of Bacterial Origin Octapeptide Sequence, Glu-Asp-Gly-Asn-Lys-Pro-Gly-Lys-OH

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Accepted: 28 January 2010/Published online: 3 March 2010 © The Author(s) 2010. This article is published with open access at Springerlink.com

Abstract The repeating octapeptide sequence, Glu-Asp-Gly-Asn-Lys-Pro-Gly-Lys-OH derived from the glycoprotein found in *Staphylococcus aureus* cell wall is assembled by simple solid phase peptide synthesis methodology using a base labile linker.

Keywords Phencyl bromide · Protein-A fragment · *Staphylococcus aureus* · Bacterial cell wall · Octapeptide · Antibacterial · Base labile linker · Manual SPPS

Protein-A, a glycoprotein obtained from Staphylococcus aureus Cowen 1. bacterial cell wall (Kronvall and Gewurz 1970) appears to be working against toxicity of certain chemical exposure (Dohadwala and Ray 1985; Singh et al. 1990) and is a potent immunomodulator (Catalona et al. 1981). It has been shown to ameliorate toxicity of cyclophosphamide, carbon tetrachloride, bacterial endotoxins (Ray et al. 1984; Kerbel and Davies 1974) and have been demonstrated to regress primary tumors in rodent and mammalian models through modulating the reactivity of cellular and humoral factors (Atherton and Sheppard 1985). The octapeptide Glu-Asp-Gly-Asn-Lys-Pro-Gly-Lys-OH was found to be the repeating sequence in protein-A structure. Toxicology and other biological studies prompted to synthesize the sequence *albeit* using a very simple and convenient method inspired from Merrifield's' original synthesis.

The octapeptide sequence was assembled (Scheme 1) using a base labile linker (Katti et al. 1992). The first amino

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acid lysine was anchored from C-terminal to amino methyl resin as its t-Boc derivative. The linker was synthesized starting from phencyl bromide, which, in turn, was prepared from acetophenone and bromine using aluminium chloride under extra cooled conditions $(-78^{\circ}C)$. The phencyl bromide (2) was coupled with chloroacetic acid and it was further extended by reacting with 2-mercaptoethanol to give the full-length liker (6). The linker (6) was coupled with t-Boc-Lys^Z using dicyclohexylcarbodiimide (DCC) and 4-dimethyl aminopyridine (DMAP) to give *t*-Boc-Lys^Z-Linker (7) and it was oxidized to sulphone (9) via reduction and subsequent oxidation of the product (8) with potassium per oxomonosulphate (oxone). The oxidized t-Boc-Lys^Z-Linker (9) was coupled to aminomethyl resin using DCC and 1-hydroxybenzotriazole (HOBt) in dichloromethane-dimethyl formamide (CH₂Cl₂-DMF), quantified by picric acid method (Rich and Sing 1979; Stewart and Young 1984). The first loading of t-Boc-Lys^Z-Linker (97.12 micromoles per gram) on solid support was sequentially extended towards n-terminus by single addition of 3-molar equivalents of suitably protected amino acids activated by DCC-HOBt. The protected amino acids were prepared during the study, purified by recrystallization and well characterised by thorough comparison with authentic reference samples. The *t*-Boc, Z, benzyl, methyl ester groups were employed in protecting the amino and carboxyl groups in amino acids i.e., Lys (Klieger and Gibian 1962; Schwyzer and Rittel 2004; Bezas and Zervas 1961), Glu (Bodanszky and Bodanszky 1982; Hirs and Timasheff 1979), Asp (Hirs and Timasheff 1979; Fujino and Hatanaka 1967, Asn (Broadbent et al. 1967), Gly (Moroder et al. 1976) and Pro (Beyerman et al. 1968). The requisite time for each coupling was about 1 h. The assembled peptide sequence was cleaved from the solid support using 5% aq. sodium hydroxide in dioxane-

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methanol in 68% yield based on first lysine loading. The protected peptide obtained as yellowish syrup was thoroughly washed with organic solvents CH_2Cl_2 , $CHCl_3$, $CHCl_3$ –MeOH (5%), brine and dichloromethane and was deprotected in 5N HCl-dioxane with thioanisole and ethanedithiol (EDT) as scavengers and crude yield of the peptide was 87%. The octapeptide showed remarkable order of antifungal activity at micromolar dose levels in initial testing against *Candida albicans*.

Experimental

Amino acids and aminomethyl resin were from purchased from Sigma Chemical Co. USA and Merck Darmstadt, Germany, respectively. Amino acids except Gly were of L configuration. Homogeneity of amino acids and peptides was established on TLC silica gel G plates (precoated from Merck) in BuOH:AcOH:H₂O (25:5:10), BuOH:Pyridine:AcOH:H₂O (9:8:6:7), CHCl₃:MeOH:H₂O (25:3:2). Synthesis was carried out in 3-necked sintered and suction outlets fitted 500 ml RB flask with manual solid phase synthesis (Merrifield 1963).

Preparation of Phencyl Bromide (2)

To a solution of acetophenone (1, 49 g), cooled in acetone– liquid nitrogen at -78° C was added bromine (20 ml) drop wise with vigorous stirring over 30 min. When color of bromine disappeared and a pinkish reaction mixture with crystals appeared, it was filtered, water and ether washed quickly, dried under vacuum to give the phencyl bromide (2, 46 g), m. p. 48°C.

Preparation of C₆H₅C(O)CH₂C-O-CH₂-S-(CH₂)₂-OH (6)

Prepared according to Katti et al. (1992).

Preparation of Boc-Lys^Z-O-(CH₂)₂-S-CH₂-(O)-C–O-CH₂-C(O)-C₆H₅ (7)

Boc-Lys^Z (5.8 g) in dry CH_2Cl_2 was mixed with Linker (**6**, 3.8 g) and DCC (3.1 g), DMAP (250 mg) under stirring at 0°C and stirring continued for 2 h at 0°C. A syrupy material was obtained and was devoid of CH_2Cl_2 under vacuum and the solid residue was taken in EtOAc, water washed, dried over Na₂SO₄, EtOAc evaporated to give Boc-Lys^Z-Linker (**7**, 7.1 g).

Scheme 1 Schematic	$C_6H_5-C(0)-CH_3$ (1)
representation of synthesis	AICI D- 78.0
	(ACI, 3DI, -700C)
	$C_6 n_5 - C_1 (0) - C n_2 D I_1 (2)$
	$\bigvee_{2} \text{Cl-CH}_{2} - \text{C} \text{ (O)-OH} (3)$
	$C_6H_5-C(O)-CH_2-O-C(O)-CH_2Cl$ (4)
	\checkmark SH-(CH ₂) ₂ -OH (5), DIEA in EtOAC
	$C_6H_5-C(O)-CH_2-O-C(O)-CH_2-S-(CH_2)_2-OH$ (6)
	Boc-LysZ, DCC-DMAP in CH CL -DMF
	Boc-Lys ^Z -O-(CH ₂) ₂ -S-CH ₂ -(O)-C-O-CH ₂ -C(O)-C ₆ H ₅ (7)
	Zn-AcOH, MeOH
	Boc-Lys ^Z -O-(CH ₂) ₂ -S-CH ₂ -CO ₂ H (8)
	Voxone, MeOH-H ₂ O
	Boc-Lys ^Z -O-(CH ₂) ₂ - $\overset{2}{SO_2}$ -CH ₂ -CO ₂ H (9)
	$\mathbf{H}_{\mathbf{N}}$ -CHC_H $\mathbf{e}_{\mathbf{e}_{\mathbf{e}_{\mathbf{e}_{\mathbf{e}}}}}$, DCC-HOBt, 0
	Boc-Lys ^Z -O-(CH ₂) ₂ -SO ₂ -CH ₂ -CONH-CH ₂ -C ₆ H4-® (10)
	↓ 1N-HCI-Dioxane,0C,20min.,DIEA-DMF,
	Protected Amino Acid (3 equiv.), HOBt (1 equiv.) in DMF, 0
	Boc-Glu ^{gOBzl} -Asp ^{gOBzl} -Gly-Asn-Lys ^Z -Pro-Gly-Lys ^Z -O-(CH ₂) ₂ -SO ₂ -CH ₂ -CONH-CH ₂ -C ₆ H ₄ -®
	$\bigvee_{5\% \text{ aq. NaOH: Dioxane: MeOH (1:2:1), 0}} $ (11)
	0C, 30 min
	Boc-Glu ^{gOBzl} -Asp ^{gOBzl} -Gly-Asn-Lys ^Z -Pro-Gly-Lys ^Z -OH (12)
	5N HCI-Dioxane, EDT, Thioanisole
	Glu-Asp-Gly-Asn-Lys-Po-Gly-Lys-OH (13)

Boc-Lys^Z-O-(CH₂)₂-S-CH₂-CO₂H (8)

To a solution of (7, 2.2 g) in MeOH (30 ml) was added acetic acid (2 ml) and a pinch of Zinc dust under stirring at RT After 1 hr. the mixture was filtered and syrup obtained was dissolved in EtOAc, washed with water and organic layers combined, dried over anhydrous Na_2SO_4 and evaporated to give (8) as oil, 1.0 g.

Boc-Lys^Z-O-(CH₂)₂-SO₂-CH₂-CO₂H (9)

To the stirring solution of (8, 1 g) in MeOH was added oxone (0.60 g in 5 ml water) and stirring continued for 1 h at RT. Reaction mixture was extracted with EtOAc, H₂O washed and organic layer concentrated in vacuum to yield (9, 1.2 g).

Boc-Lys²-O-(CH₂)₂-SO₂-CH₂-CONH-CH₂-C₆H₄-
$$\otimes$$
 (10)

Aminomethyl resin (2 g) swelled in CH_2Cl_2 was coupled with Boc-Lys^Z-O-(CH₂)₂-SO₂-CH₂-CO₂H (**9**, 1.0 g) in CH₂Cl₂-DMF (1:1, 50 ml) under vigorous stirring using DCC (500 mg) and HOBt (400 mg) dissolved in CH₂Cl₂ at 0°C for 4 h. The coupling was repeated by adding more DCC (500 mg) and HOBt (400 mg) with stirring for 2 more hours. The unreacted Boc-Lys^Z-O-(CH₂)₂-SO₂-CH₂CO₂H was detected by TLC after final loading and the coupled resin was washed with DMF, CH₂Cl₂, MeOH and DMF (25 ml each).

Sequence Loading and Quantitation

Picric acid method was used to quantify the loading of resin. The unbound amino endings were capped with Ac₂O-DMAP. The resin was washed with CH₂Cl₂ and DMF (20 ml × 2) and quantified to 91% (712 mM/g) ready for coupling. The suitably protected amino acids were coupled to solid support in the following manner, (1) the cleavage of *t*-Boc group was carried out using 1N HCl in dioxane at 0°C, neutralized with DIEA and coupled using DCC-HOBt in DMF:CH₂Cl₂ (1:1) at 0°C, (2) after each coupling and deprotection, the resin was washed with DMF:CH₂Cl₂ (1:1, 15 ml × 2, 5 min each), (3) prior to every deprotection, the coupling was monitored by picric acid method and allowed to proceed to completion generally within 2 h.

Sequence Release and Purification

The peptide was released from solid support with 5% aq. NaOH-dioxane-MeOH (1:2:1, 20 ml) at 0°C for 0.5 h. The

crude peptide was kept in 5N HCl-dioxane with ethanedithiol and thioanisole for 0.5 h to yield the (naked) peptide which was washed with dichloromethane, diethyl ether and purified by column chromatography using Sephadex LH-20 in MeOH and water:MeOH (65:35) with fractions pooled and lyophilized to give a transparent oil, 4.41 g, 73%, TLC:single spot, BuOH:AcOH:H₂O (25:5:10), Rf 5, Ninhydrin. FAB-MS (m/z): 843 (M+) corroborated the structure.

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