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Total Synthesis of B-Chain of Human Q2HS Glycoprotein

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Abstract: B-chain of human α 2HS glycoprotein 1, a cystein-containing heptacosapeptide carrying a trisaccharide (sialyl T) side chain, was synthesized for the first time by the solid-phase method utilizing a glycosyl serine building block 11 with benzyl protecting groups. © 1997 Elsevier Science Ltd.

Present-day synthesis of glycopeptides relies heavily on solid-phase methods, which provide not only rapid assembly of peptide backbones but also high coupling efficiency with sophisticated and fully automated instruments.¹ However, synthesis of a longer oligopeptide (>25 amino acid-peptide) possessing both multiple functional side chains and oligosaccharide attachment has rarely been achieved.² Recently we have disclosed a solid-phase synthesis of heptacosapeptide 2^3 carrying an O-linked disaccharide $(Gal\beta 1 \rightarrow 3GalNAc\alpha 1 \rightarrow, Thomsen-Friedenreich antigen)$, an asialo demercapto $[Ala^{18}]$ analogue of B-chain⁴ (1) of human α 2HS glycoprotein, which is a human plasma globulin behaving as one of the negative acutephase reactants.5



The synthesis was performed on the basis of Fmoc method using an automated peptide synthesizer. Only at the coupling step with the benzylated disaccharide-linked serine unit, the reaction was carried out utilizing a mechanical shaker in order to make easy the recovery of the unreacted glycoserine, though efficiency

of the condensation was moderate (55%). The ultimate deprotection was accomplished by hydrogenolysis to give the target compound 2. The fact that a glycoserine-deleted peptide was formed in 45% and the unreacted glycoserine building block was recovered in a reasonable amount even after running the reaction for 64 h led us to conclude that more effective mixing would be necessary to complete the solid-phase reaction.

In this communication, we report a first total synthesis of the native B-chain of α 2HS glycoprotein involving an improved procedure by the use of a vortexing mixer with a suitably protected glycoserine building block 11.

The trisaccharide-linked serine 11 was synthesized as follows. The disaccharide 3^6 was treated with ceric ammonium nitrate in a mixture of toluene:CH₃CN:H₂O (3 : 4 : 3) to afford hemiacetal 4 (77%), which was converted into trichloroacetimidate 5 (CCl₃CN, DBU, α/β 3/1, 97%). The glycosyl acceptor 7 was prepared in 2 steps from the known glycosylated serine derivative 6^7 (1. 80% CF₃CO₂H-CH₂Cl₂, 97%, 2. TBDMSCl, imidazole, DMF, 91%). Glycosylation of 7 with the α -trichloroacetimidate 5 was promoted by BF₃•OEt₂ (0.3 equiv.) in toluene-CH₂Cl₂ at -15 - -5° C to give a β 1 \rightarrow 3 linked product 8⁸ (54%) and its α -isomer (7%). Desilylation of 8 (aq. CF₃CO₂H-CH₂Cl₂, 83%) followed by benzylidenation [PhCH(OMe)₂, p-TsOH, CH₃CN, 95%] afforded 9, which upon treatment with AcSH in pyridine⁹ (\rightarrow 10) and then with Pd(PPh₃)₄-MeNHPh¹⁰ in THF was converted into 11⁸ (70%).



The henicosapeptide (7 - 27)-linked HMP resin (4-hydroxymethylphenoxymethyl-copolystylene-1% divinylbenzene) was synthesized in a 0.25 mmol scale on an automated peptide synthesizer according to the Fmoc protocol, the side-chain functional groups being blocked with Trt groups for cysteine, glutamine, and histidine, Boc group for lysine, and Pmc (2,2,5,7,8-pentamethylchroman-6-sulfonyl) group for arginine. The coupling reactions were performed with DCC-HOBt in NMP (N-methylpyrrolidone), while the N-terminal Fmoc group was removed with piperidine in NMP. Coupling yield in each step was determined by ninhydrin method and the estimated overall yield after 20 couplings was 88%. The glycoserine building block 11 (2.6 equiv) was thus activated with DCC-HOBt and condensed to the deFmoc henicosapeptide-resin (14 µmol). The mixture was shaken on a vortex test tube-mixer for 24 h at room temperature.



Scheme 2

Then the resin was put back into a small scale reaction vessel for the synthesizer and the N-terminal amino acid residues (1 - 5) were coupled under the automatic program. After completing the peptide chain elongation, treatment of the deFmoc glycopeptide-resin with a TFA-based cleavage reagent (95% aq. TFAphenol-thioanisole-1,2-ethanedithiol, 42:3:2:1) followed by gel-permeation chromatography afforded the In contrast to the prior experiment³, HPLC analysis on a reversed-phase column resin-free glycopeptide. demonstrated that no detectable amount of glycoserine-deleted byproduct was formed in the reaction and that the product consisted mainly of hexa- (52%, MALDI-TOF+MS : 3920 [M+1]*), penta- (31%, 3829 [M+1]*), and tetrabenzylated trisaccharide-bound peptides (4%, 3740 [M+1]*). The latter two were formed by partial debenzylation during the TFA-cleavage procedure. The glycopeptides were collected by preparative HPLC and treated with TMSOTf/TFA¹¹ in the presence of thioanisole to give the debenzylated product, which upon gel-permeation chromatography afforded two fractions corresponding to the dimeric (58%, 6760 [M+1]*) and the monomeric (42%, 3363 [M+1]⁺) glycopeptides, respectively. Both fractions were further purified by reversed-phase HPLC. The dimeric compound retained about half of the lactone linkage which was evidenced by the characteristic proton signal at δ 5.30 ppm for Gal H-4, while the monomer had no lactone and The dimer was treated with 0.2M NaHCO₃/D₂O (pH 7.5) for 4 days to cleave was the target compound 1. the lactone⁶ and then with 1,4-dithiothreitol overnight to give 1 exclusively (85% total yield for deprotection). The 1 H-NMR signals for the carbohydrate part of synthesized 1⁸ were in good agreement with those reported for the trisaccharide derived from natural 1.6 ESI•MS showed m/z 1133.1 as $[(M+3)/3]^{3+}$. Finally, it was observed that the isolated 1 was prone to dimerize in the absence of antioxidant (e.g. dithiothreitol).

In conclusion, the B-chain of α 2HS glycoprotein, a heptacosapeptide carrying an O-linked sialotrisaccharide, cystein, and several basic amino acid residues, was synthesized for the first time, in good

yield by solid-phase method. The benzyl groups on the carbohydrate residues were efficiently removed under acidic condition without any significant side-reaction on the carbohydrate linkages.

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- 8. Selected ¹H-NMR data are given below. Compound 8 (270 MHz, CDCl₃): δ 7.75 (d, 2 H, *J* = 7.3 Hz, Ar), 7.61 (d, 2 H, *J* = 7.3 Hz, Ar), 7.4-7.1 (m, 34 H, Ar), 5.93 (m, 1 H, -CH=CH₂), 5.83 (d, 1 H, *J* = 8.6 Hz, NH), 5.34 (brd, 1 H, *J* = 17.2 Hz, =CH₂), 5.26 (brd, 1 H, *J* = 10.2 Hz, =CH₂), 5.21 (d, 1 H, *J* = 4.0 Hz, H-4b), 4.95 (d, 1 H, *J* = 3.3 Hz, H-1a), 2.18 (dd, 1 H, *J* = 4.6, 11.9 Hz, H-3c eq), 1.71 (s, 3 H, Ac), 0.85 (s, 9 H, t-Bu), 0.03 and 0.02 (2 s, 6 H, 2 Me); ¹³C NMR (68 MHz): δ 103.0 (C-1b), 99.4 (C-1a), 95.3 (C-2c). Compound 11 (270 MHz, CDCl₃): δ 7.72 (d, 2 H, *J* = 7.6 Hz, Ar), 7.54 (m, 4 H, Ar), 7.4-7.1 (m, 37 H, Ar), 6.12 (brd, 1 H, *J* = 6.9 Hz, NH), 5.93 (d, 1 H, *J* = 7.3 Hz, NH), 5.39 [brs, 1 H, PhCH(O)₂], 5.12 (d, 1 H, J = 3.6 Hz, H-4b), 5.04 (brs, 1 H, H-1a), 2.13 (brd, 1 H, *J* = 13.0 Hz, H-3c eq), 1.82 and 1.72 (2 s, 6 H, 2 Ac). Compound 1 [600 MHz; D₂O, 25 °C (or 60 °C), *t*-BuOH (δ 1.23)]: δ 7.62 (brs, 1 H, His), 7.31 (brt, 2 H, Phe), 7.27 (brt, 1 H, Phe), 7.19 (d, 2 H, *J* 7.3 Hz, Phe), 6.85 (brs, 1 H, His), 4.89 (d, 1 H, *J* 2.4 Hz, H-1:GalNAc, 60 °C), 4.45 (d, 1 H, *J* 4.3, 12.2 Hz, H-3eq;NeuAc), 2.02 and 1.99 (2s, 6 H, 2Ac;GalNAc and NeuAc), 1.77 (t, 1 H, *J* 12.2 Hz, H-3ax; NeuAc).
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