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Synthesis of a new conformationally constrained glycoamino acid building block

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Abstract—The synthesis of a suitably protected β -D-glucopyranosyl-(S)- α -methylserine derivative—a new conformationally constrained glycosylated quaternary amino acid analogue of β -D-glucopyranosyl-L-serine—is described. This compound can be used as an attractive building block for the synthesis of new, constrained glycopeptides. © 2003 Elsevier Ltd. All rights reserved.

The synthesis of α,α -disubstituted α -amino acids has received enormous attention because it represents a great advance in the design of modified peptides with altered biological activity.^{1,2} For example, α -methylserine can be regarded as a potential C-terminal α -helix-stabilising building block.³

On the other hand, a vast number of natural proteins contain oligosaccharide side chains—glycoproteins—and the saccharide residue is covalently linked to the peptide backbone. In the majority of cases this linkage is *O*-glycosidic through serine. It is well known that the saccharide moiety plays an important role in biological processes by contributing to the membrane permeability and stabilising certain secondary conformations of the peptide backbone.^{4–7} Nevertheless, while significant research has been undertaken in the design of restricted peptides, the field of conformationally constrained glycopeptides remains relatively unexplored, and most of the restrictions studied are centered in the saccharide moiety.^{8–11}

We have recently combined the aspects mentioned above and focused our attention on the synthesis of conformationally constrained glycosylated amino acids. Such compounds are building blocks for the construction of constrained glycopeptides, which can be Taking into account this observation and the fact that various biologically active O-glycopeptides or O-glycoproteins incorporate the unusual β -D-glucopyranosyl-L-serine (β -D-Glc-L-Ser) substructure, we decided to synthesise a suitably protected β -D-glucopyranosyl- α -methylserine derivative β -D-Glc-(S)- α -MeSer (1) as a conformationally restricted analogue of β -D-Glc-L-Ser (2) (Fig. 1). The novel unnatural glycoamino acid 1 is an attractive building block for incorporation into peptides, since only a limited number of glycosylated amino acid building blocks suitable for glycopeptide synthesis are commercially available.

Figure 1. Protected β-D-glucopyranosyl-(S)-α-methylserine [β-D-Glc-(S)-MeSer] (1) and β-D-glucopyranosyl-L-serine (β-D-Glc-L-Ser) (2).

regarded as potential tools in the elucidation of the conformational requirements for binding to receptors and enzymes. In this sense, a number of glycoamino acids that contain conformational constraints have been synthesised and the majority of unnatural glycosylated amino acids reported are C-glycoside derivatives, $^{12-15}$ including C-glycosylated α -amino acids that possess a quaternary center $(C\alpha)$. $^{16-19}$ Examples of exclusively O-glycosylated quaternary α -amino acids have not been reported to date.

Keywords: amino acids and derivatives; azides; glycosidation; peptide analogues/mimetics.

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Several examples of O-glycopeptides or O-glycoproteins that incorporate the β -D-Glc-L-Ser linkage are present in the first epidermal growth factor-like domains, 20 certain synthetic enkephalin analogues 21 and the blood-clotting factors VII and IX (two plasma glycoproteins that are involved in the blood coagulation cascade). $^{22-24}$ In these proteins the serine residue bears either a disaccharide α -D-Xyl-1,3- β -D-Glc chain in factor VII or a trisaccharide α -D-Xyl-1,3- α -D-Xyl-1,3- β -D-Glc chain in factor IX. More specifically, factor IX consists of a single chain of 416 amino acids and the glycopeptide sequence containing the residues 51–56, including the trisaccharide, has been synthesised. 25

It was planned to use the suitably protected αmethylserine 5 (N-Fmoc-MeSer-OMe) as the starting material for the glycosidation reaction with the donor 2,3,4,6-tetra-O-acetyl- α -D-glucopyranosyl bromide²⁶ **6** using the Koenigs-Knorr methodology with silver triflate (AgOTf). Compound 5 was obtained from methyl α -methylserinate 4, which was synthesised using our methodology involving the Sharpless asymmetric dihydroxylation of the Weinreb amide of 2-methyl-2propenoic acid 3, followed by several transformations (Scheme 1).²⁷ However, this approach proved unsuccessful and further attempts were made using more conveniently protected α -methylserine 9 (N-Fmoc-MeSer-OPfp). Compound 9 was synthesised from α methylserine hydrochloride 7 by protection of the amino group with FmocCl, followed by formation of the Pfp-ester. However, glycosidation of N-Fmoc-Ser-

Me Me NH2
N OMe

3 ii

Me NHFmoc
HO
$$CO_2Me$$

5 $NH_2 \cdot HCI$
HO CO_2He

7 $NH_2 \cdot HCI$
HO CO_2He

7 $NH_2 \cdot HCI$
HO CO_2He

1 iv

NHFmoc
HO CO_2Re

1 iv

NHFmoc
HO CO_2Re

1 iv

 RO_RO
 RO_RO

Scheme 1. Reagents and conditions: (i) Ref. 27: (a) AD-mix α, MeSO₂NH₂, 'BuOH/H₂O (1:1), 0°C, 12 h, 81%; (b) LiOH·H₂O, H₂O/MeOH (1:3), rt, 2 h; (c) AcCl, MeOH, reflux, 12 h, 85%; (d) SOCl₂, CCl₄, reflux, 4 h, 85%; (e) NaN₃, DMF, 50°C, 2 days, 75%; (f) H₂/Pd–C, MeOH, rt, 24 h, 83%; (ii) Fmoc-OSu, TEA, MeCN, rt, 12 h, 55%; (iii) 6N HCl, reflux, 12 h, 100%; (iv) FmocCl, K₂CO₃, H₂O/THF (1:1), 0°C to rt, 12 h, 70%; (v) PfpOH, DCC, AcOEt, 0°C, 2 h, 70%.

OPfp **9** with the donors 2,3,4,6-tetra-O-benzoyl- α -D-glucopyranosyl bromide²⁸ **10** with AgOTf or 2,3,4,6-tetra-O-acetyl- α -D-glucopyranosyl trichloroacetimidate²⁹ **11** with BF₃·Et₂O failed (Scheme 1).

glycosidation reaction with these classical approaches has given good results in the case of serine.²⁵ However, given the disappointing results obtained here, we decided to change the methodology and start from compound 15—a precursor of the α-methylserine 7. Compound 15 was obtained by a modified version of the procedure cited above²⁷ and required formation of the benzyl ester instead of the methyl ester. Diol 12 was prepared by Sharpless asymmetric dihydroxylation of 3 with AD-mix α and the product was hydrolysed with HCl under reflux, treated with Cs₂CO₃ in MeOH and esterified with BnBr to give compound 13. Diol 13 was transformed into its 2,3-cyclic sulfite 14 with thionyl chloride and the ring-opening reaction of this sulfite with azide ions in dimethylformamide gave a mixture of the azido esters 15 and 16. A regioselectivity of 4.5:1 in favour of compound 15 was obtained and this corresponds to attack at the quaternary centre with clean inversion of configuration³⁰ (Scheme 2).

The glycosidation of 15, once separated from 16 by column chromatography, with donor 6 using the Koenigs-Knorr methodology with a mixture of silver carbonate (Ag₂CO₃) and silver perchlorate (AgClO₄) gave 70% of the undesired orthoformiate derivative 17 with 25% of unreacted 6 recovered. Fortunately, the glycosidation of 15 using trichloloracetimidate 11 gave 50% yield of the desired derivative 18. This yield was later improved by using the donor 2,3,4,6-tetra-O-acetyl-α-D-glucopyranosyl fluoride^{31,32} **19** in the presence of BF₃·Et₂O and triethylamine (TEA),³³ a process that gave 18 in 60% yield. The azido group of this compound was readily transformed into the amino group by the action of Ph₃P and water in tetrahydrofuran. This reaction gave β -D-Ac₄Glc-(S)-MeSer-OBn (20), which was easily converted into the required building block 1 by protection of the amino group as the Fmoc carbamate (Scheme 2).34

In conclusion, we have synthesised the suitably protected glycosylated amino acid β-D-Glc-(S)-α-MeSer (1). This compound is an important building block in the synthesis of constrained glycopeptides in which the conformational restriction is only present in the amino acid moiety (quaternary α -amino acid). To the best of our knowledge this is the first time that a glycosylated amino acid with these features has been synthesised. The nearest previous example is the glycosylated amino acid recently obtained by Lane and Halcomb, 19 although in this case the restriction affects not only the amino acid moiety but also the carbohydrate. We intend to explore the incorporation of other carbohydrates into different hydroxylated quaternary α-amino acids, using the methodology described here, in order to assess the applicability of this approach in the synthesis of conformationally restricted glycopeptides.

Scheme 2. Reagents and conditions: (i) Ref. 27: AD-mix α, MeSO₂NH₂, 'BuOH/H₂O (1:1), 0°C, 12 h, 81%; (ii) (a) 6N HCl, reflux, 1 h, (b) Cs₂CO₃, MeOH, rt, 20 min. (c) BnBr, DMF, rt, 48 h, 84%; (iii) SOCl₂, CCl₄, reflux, 4 h, 81%; (iv) NaN₃, DMF, 50°C, 12 h, column chromatography (hexane/AcOEt = 4/1), 77% of **15** and 17% of **16**; (v) **6**, Ag₂CO₃, AgClO₄, CH₂Cl₂, MS 4 Å, -10°C to rt, 12 h; 70% (vi) **11**, BF₃·EtO₂, CH₂Cl₂, rt, 12 h; 50% (vii) **19**, BF₃·Et₂O, TEA, CH₂Cl₂, rt, 12 h; 60%; (viii) Ph₃P (2 equiv.), H₂O (2 equiv.), THF, reflux, 4 h, 71%; (ix) FmocCl, DIEA, CH₂Cl₂, rt, 12 h, 58%.

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- 34. Compound **20**: Mp: 65–67°C. $[\alpha]_D^{25} = -14.3$ (*c* 1.01, MeOH); ¹H NMR (400 MHz, CDCl₃) δ (ppm): 1.28 (s,

3H), 2.00 (s, 3H), 2.02 (s, 3H), 2.04 (s, 3H), 2.06 (s, 3H), 3.50 (d, 1H, J=9.6 Hz), 3.64 (ddd, 1H, J=10.0 Hz, J=4.6 Hz, J=2.5 Hz), 4.11 (dd, 1H, J=12.3 Hz, J=2.5 Hz), 4.17 (d, 1H, J=9.6 Hz), 4.25 (dd, 1H, J=12.3 Hz, J=4.6 Hz), 4.57 (d, 1H, J=8.0 Hz), 4.96 (dd, 1H, J=9.6 Hz, J=8.0 Hz), 5.07 ('t', 1H, J=9.6 Hz), 5.15 (s, 2H), 5.18 ('t', 1H, J=9.6 Hz), 7.33–7.37 (m, 5H); ¹³C NMR (400 MHz, CDCl₃) δ (ppm): 20.5, 20.6, 22.7, 58.5, 61.9, 67.0, 68.4, 71.3, 71.8, 72.7, 76.1, 101.1, 128.0, 128.3, 128.5, 132.0, 132.1, 135.7, 167.1, 169.3, 170.1, 170.5, 175.3. Anal. calcd for $C_{25}H_{33}NO_{12}$: C, 55.65; H, 6.17; N, 2.60. Found; C, 55.71; H, 6.22; N, 2.84.