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Syntheses of α -dystroglycan derived sialylated glycosyl amino acids carrying a novel mannosyl serine/threonine linkage

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Abstract

An efficient access to the Fmoc protected glycosyl amino acids **20a**-d which correspond to the unique sequence NeuNAc- α -(2,3)-Gal- β -(1,4)-GlcNAc- β -(1,2)-Man- α -Ser/Thr recently found in laminin-binding protein α -dystroglycan, is described. Tetrasaccharide donor **18** was constructed from monosaccharide blocks **2**, **3**, **8** and **9** and coupled with Ser/Thr derivatives **19a**-d. Resultant glycosyl amino acid **20d** was completely deprotected into **1a**. © 1999 Elsevier Science Ltd. All rights reserved.

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 α -Dystroglycan (α -DG) is a membrane-associated, extracellular glycoprotein. It is anchored to the cell-membrane via binding to the transmembrane glycoprotein β -dystroglycan (β -DG) to form an α/β -DG-complex which is widely expressed in various types of tissues including skeletal muscle^{1a} and schwannoma cells.^{1b} The α/β -DG-complex serves as a transmembrane linker between the extracellular matrix and the intracellular skeleton as α -DG binds with a high affinity to extracellular matrix components such as laminin in the striated muscle whereas the intracellular domain of β -DG binds to cytoskeletal proteins such as dystrophin.^{1c} Disruption of this transmembrane linkage leads to skeletal muscular dystrophies.^{1d}



At present little is known about the biological roles of the carbohydrate moieties of α -DG. Investigations on bovine peripheral nerve α -DG have revealed, that the tetrasaccharide **1a**,**b**, containing a novel *O*-mannosyl serine/threonine moiety,^{1e} is the major constituent of the O-linked carbohydrates and

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contributes to laminin binding.^{1a} To provide molecular probes for elucidating the binding process of bovine peripheral nerve α -DG to laminin, we started the program to synthesize the tetrasaccharidic serine/threonine building blocks **20a**–d. These glycosyl amino acids are designed to be applied to the construction of glycopeptide sequences of α -DG,^{1c} possibly to correlate the laminin binding properties with ligand density or sequence specifity in comparison to the native α -DG.

For the synthesis of the Fmoc-protected glycosyl amino acids 20a-d (Scheme 1) we decided to introduce the amino acid moiety at the end of the synthesis. This option would offer maximum flexibility in choosing the amino acid protection and facilitates the construction process of the tetrasaccharidic moiety by eliminating any technical problem that may arise in the presence of acid or base sensitive amino acid component. The trichloroacetimidate technology² was adopted for the final coupling with serine/threonine derivatives 19a-d,³ which calls for 18 as a suitable glycosyl donor. Compound 18 was constructed via the two disaccharides 7 and 12 (Scheme 2) which were prepared in a multigram scale from easily available building blocks $2,^{4d}$ $3,^5$ 8^6 and $9.^7$



Scheme 1. Retrosynthetic disconnection of the glycosyl amino acids **20a-d** into the trichloroacetimidate **18** and the Fmoc-protected amino acids **19a-d**. The tetrasaccharide **18** was formed by the readily available building blocks **2**, **3**, **8** and **9** (All=allyl, Pfp=pentafluorophenyl, TDMS=thexyldimethylsilyl).



Scheme 2. Formation of the disaccharides 7 and 12; (i) NIS, TfOH, CH₃CN, MS 3 Å, -40° C; (ii) pyridine, Ac₂O (57%, two steps); (iii) TBAF, HOAc, THF (92%); (iv) Cl₃CCN, DBU, CH₂Cl₂, 0°C (93%); (v) NIS, TfOH, CH₃CN, MS 3 Å, -40° C (95%); (vi) ethylenediamine, *n*-BuOH, 100°C; (vii) TfN₃, DMAP, CH₂Cl₂ (82%, two steps); (TCAI=-(C=NH)-CCl₃)

Synthesis of the α -(2,3)-sialylated disaccharide 7 was accomplished by reaction of thiomethylsialoside 2 with the galactose derived triol 3 (Scheme 2). The reaction was performed under standard conditions⁴ (NIS, TfOH, CH₃CN^{4d}) to afford, after acetylation of 4, the α -(2,3)-linked product 5 in 57% yield (Scheme 2, i, ii). Subsequent treatment with tetrabutylammoniumfluoride (TBAF) in the presence of excess acetic acid (Scheme 2, iii) cleaved the thexyldimethylsilyl group. The reaction of the resulting hemiacetal **6** with trichloroacetonitrile and DBU² afforded the trichloroacetimidate **7** (Scheme 2, iv) in 93% yield.

The disaccharide 12, designed to be coupled with trichloroacetimidate 7, was prepared from the monosaccharides 8 and 9. Glycosylation of the mannose derivative 9 with the thioglycoside 8, promoted by NIS/TfOH,^{4a-c} afforded disaccharide 10 in high yield (Scheme 2, v). Compound 10 was subsequently reacted with ethylenediamine in *n*-butanol at 100°C⁸ furnishing the 2-amino disaccharide 11 (Scheme 2, vi) which was subjected to the azidotransfer reaction with trifluoromethanesulfonyl azide⁹ (Tf-N₃) to give the azido compound 12 (Scheme 2, vii) in 82% yield from 10.

After preparation of the disaccharides 7 and 12, the formation of the tetrasaccharide 13 was next examined. The best results were obtained with equimolar amounts of 7 and 12 using TMSOTf (0.2 equiv.) as promoter to give tetrasaccharide 13 in 83% yield (Scheme 3, i). Conversion of compound 13 into the peracetylated trichloroacetimidate 18 was accomplished by a series of standard functional group manipulations (Scheme 3, ii–vi). Thus, the azido group of 13 was reduced by 1,3-propanedithiol¹⁰ to afford the corresponding amine which was treated with acetic anhydride to furnish the *N*-acetylated compound 14 in 95% yield (Scheme 3, ii). Complete removal of the benzyl groups (Scheme 3, iii) afforded heptaol 15 which was acetylated to 16 (Scheme 3, iv) and subsequently treated with hydrazine acetate in DMF at 0°C (Scheme 3, v) to give the hemiacetal 17. Finally, the reaction of 17 with trichloroacetonitrile/DBU² (Scheme 3, vi) afforded the designed trichloroacetimidate 18. The stereoselective attachment of the Fmoc amino acids 19a–d to 18 was performed at –20°C in CH₂Cl₂:toluene (1:1) by activation with TMSOTf (0.2 equiv.). These conditions (Scheme 3, vii) exclusively provided the α -linked glycosyl amino acids 20a–d¹¹ in good yields, which are designed to be incorporated into Fmoc-based solid phase peptide synthesis.



Scheme 3. Formation of the glycosyl amino acids **20a**–d and deprotection of compound **20d**; (i) TMSOTf, CH_2Cl_2 , MS 4 Å, $-20^{\circ}C$, (83%); (ii) (a) 1,3-propanedithiol, DIPEA, pyridine: H_2O (7:3); (b) Ac_2O (95%, two steps); (iii) H_2 , Pd(OH)₂, MeOH, HOAc; (iv) pyridine, Ac_2O , 0°C (95%, two steps); (v) $H_2NNH_2 \times HOAc$, DMF, 0°C; (vi) Cl_3CCN , DBU, CH_2Cl_2 , 0°C (91%, two steps); (vii) TMSOTf, CH_2Cl_2 :toluene (1:1), MS 4 Å, $-20^{\circ}C$; (viii) Pd(Ph₃P)₄, dimedone, THF (90%); (ix) morpholine, CHCl₃; (x) NaOMe/MeOH, pH≈9.0; (xi) 5 mM NaOH/aq. MeOH (90%, three steps); (DIPEA=diisopropylethylamine, TCAI=–(C=NH)–CCl₃)

Completely deblocked tetraosyl serine 1a was obtained from 20d as follows (Scheme 3, viii–xi). First, the allyl ester was cleaved by $Pd(Ph_3P)_4$ in the presence of dimedone (Scheme 3, viii). Subsequent

removal of the Fmoc group was effected by morpholine (Scheme 3, ix) followed by complete removal of acetyl groups (NaOMe/MeOH, pH \approx 9.0, (Scheme 3, x)) and saponification of methyl ester (5 mM NaOH/aq. MeOH, (Scheme 3, xi)). Purification by size exclusion chromatography (Bio-Gel[®] P-2) furnished the deprotected compound **1a** in 81% overall yield from **20d**. The structure of **1a** was confirmed by NMR and ESI mass spectroscopy.¹²

In summary, we have established a versatile and efficient synthetic route to the α -(2,3)-sialylated tetrasaccharidic glycosyl amino acids **20a-d** containing a novel *O*-mannosyl serine/threonine linkage.

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- 5. (a) Compound 3 was prepared from 2,3,4-Tri-O-acetyl-6-O-benzyl-D-galactopyranosyl^{5b} acetate in three steps: (i) N₂H₄×HOAc, DMF, 0°C (85%); (ii) TDMSCl, imidazole, DMF, 0°C-rt (91%); (iii) NaOMe/MeOH, -10°C (93%); for the use of the TDMS group for anomeric protection, see (c). (b) Dekany, G.; Wright, K.; Toth, I. J. Carbohydr. Chem. 1997, 16, 983-999. (c) Hummel, G.; Schmidt, R. R. Tetrahedron Lett. 1997, 38, 1173-1176.
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- 11. Selected ¹H and ¹³C NMR data (CDCl₃): **20a** ¹H NMR (500 MHz): δ 5.83 (d, 1H, J=8.1 Hz, NH^{Ser}), 4.74 (d, 1H, J<1.0 Hz, H-1 α^{M}), 4.67 (d, 1H, J=8.2 Hz, H-1 β^{Gal}), 4.57 (d, 1H, J=8.1 Hz, H-1 β^{GN}); ¹³C NMR (125 MHz): δ 100.90 (C-1^{Gal}), 99.66 (C-1^{GN}), 98.27 (J_{C,H}=169.9 Hz, C-1^M), 96.74 (C-2^{SA}); **20b** ¹H NMR (500 MHz): δ 6.07 (d, 1H, J=8.8 Hz, NH^{Ser}), 4.83 (d, 1H, J<1.0 Hz, H-1 α^{M}), 4.66 (d, 1H, J=8.3 Hz, H-1 β^{Gal}), 4.62 (d, 1H, J=7.8 Hz, H-1 β^{GN}); ¹³C NMR (125 MHz):

 $\delta 100.86 \text{ (C-1}^{Gal}\text{)}, 99.58 \text{ (C-1}^{GN}\text{)}, 98.92 \text{ (}J_{C,H}=168.7 \text{ Hz}, \text{C-1}^{M}\text{)}, 96.75 \text{ (C-2}^{SA}\text{)}; 20c ¹\text{H} NMR (400 \text{ MHz}): \delta 5.92 \text{ (d, 1H, } J=8.3 \text{ Hz}, \text{NH}^{Thr}\text{)}, 4.70 \text{ (d, 1H, } J<1.0 \text{ Hz}, \text{H-1}\alpha^{M}\text{)}, 4.67 \text{ (d, 1H, } J=8.3 \text{ Hz}, \text{H-1}\beta^{Gal}\text{)}, 4.56 \text{ (d, 1H, } J=7.8 \text{ Hz}, \text{H-1}\beta^{GN}\text{)}; \\ ^{13}\text{C} \text{ NMR} (100 \text{ MHz}): \delta 100.85 \text{ (C-1}^{Gal}\text{)}, 99.85 \text{ (C-1}^{GN}\text{)}, 98.79 \text{ (}J_{C,H}=171.7 \text{ Hz}, \text{C-1}^{M}\text{)}, 96.70 \text{ (C-2}^{SA}\text{)}; 20d ¹\text{H} \text{ NMR} (500 \text{ MHz}): \delta 5.92 \text{ (d, 1H, } J=8.3 \text{ Hz}, \text{NH}^{Scr}\text{)}, 4.70 \text{ (d, 1H, } J<1.0 \text{ Hz}, \text{H-1}\alpha^{M}\text{)}, 4.67 \text{ (d, 1H, } J=8.3 \text{ Hz}, \text{H-1}\beta^{Gal}\text{)}, 4.56 \text{ (d, 1H, } J=7.8 \text{ Hz}, \text{H-1}\beta^{Gal}\text{)}, 4.56 \text{ (d, 1H, } J=7.8 \text{ Hz}, \text{H-1}\beta^{Gal}\text{)}, 4.56 \text{ (d, 1H, } J=7.8 \text{ Hz}, \text{H-1}\beta^{Gal}\text{)}, 4.56 \text{ (d, 1H, } J=7.8 \text{ Hz}, \text{H-1}\beta^{Gal}\text{)}, 4.56 \text{ (d, 1H, } J=7.8 \text{ Hz}, \text{H-1}\beta^{Gal}\text{)}, 4.56 \text{ (d, 1H, } J=7.8 \text{ Hz}, \text{H-1}\beta^{GN}\text{)}; \\ ^{13}\text{C} \text{ NMR} (125 \text{ MHz}): \delta 100.86 \text{ (C-1}^{Gal}\text{)}, 99.77 \text{ (C-1}^{GN}\text{)}, 98.39 \text{ (}J_{C,H}=170.4 \text{ Hz}, \text{C-1}^{M}\text{)}, 96.75 \text{ (C-2}^{SA}\text{)}. \\ \end{array}$

Selected data for 1a: ¹H NMR (500 MHz, D₂O, ¹BuOH as internal standard): δ 4.86 (d, 1H, J=1.4 Hz, H-1α^M), 4.56 (d, 1H, J=7.6 Hz, H-1β^{GN}), 4.53 (d, 1H, J=7.9 Hz, H-1β^{Gal}), 2.74 (dd, 1H, J_{vic}=4.6 Hz, J_{gem}=12.4 Hz, H-3_{cq}^{SA}), 2.03, 2.02 (2s, 6H, NAc), 1.79 (dd, 1H, J_{vic}=12.2 Hz, H-3_{ax}^{SA}); ¹³C NMR (125 MHz, D₂O, ¹BuOH as internal standard): δ 175.70, 175.40, 174.55, 172.53 (C=O), 103.30 (C-1^{Gal}), 100.50 (C-2^{SA}), 100.20 (C-1^{GN}), 98.29 (C-1^M), 40.33 (C-3^{SA}), 23.01, 22.73 (CH₃); ESI MS (negative mode): m/z=922.2 (M-H)⁻; calcd for C₃₄H₅₆N₃O₂₆ (M-H): 922.3; [α]_D²²=-3.0 (0.46, H₂O). 1a was recently synthesized chemoenzymatically by Matsuo, I.; Isomura, M.; Ajisaka, K. *Tetrahedron Lett.* 1999, 40, 5047-5050.