This article was downloaded by: On: 23 January 2011 Access details: Access Details: Free Access Publisher Taylor & Francis Informa Ltd Registered in England and Wales Registered Number: 1072954 Registered office: Mortimer House, 37-41 Mortimer Street, London W1T 3JH, UK



Journal of Carbohydrate Chemistry

Publication details, including instructions for authors and subscription information: http://www.informaworld.com/smpp/title~content=t713617200

An Aminated GDP-Fucose Analog Useful in the Fucosyltransferase Catalyzed Addition of Biological Probes onto Oligosaccharide Chains¹ Christer Hällgren^a; Ole Hindsgaul^a

^a Department of Chemistry, University of Alberta, Alberta, CANADA

To cite this Article Hällgren, Christer and Hindsgaul, Ole(1995) 'An Aminated GDP-Fucose Analog Useful in the Fucosyltransferase Catalyzed Addition of Biological Probes onto Oligosaccharide Chains', Journal of Carbohydrate Chemistry, 14: 4, 453 - 464

To link to this Article: DOI: 10.1080/07328309508005350 URL: http://dx.doi.org/10.1080/07328309508005350

PLEASE SCROLL DOWN FOR ARTICLE

Full terms and conditions of use: http://www.informaworld.com/terms-and-conditions-of-access.pdf

This article may be used for research, teaching and private study purposes. Any substantial or systematic reproduction, re-distribution, re-selling, loan or sub-licensing, systematic supply or distribution in any form to anyone is expressly forbidden.

The publisher does not give any warranty express or implied or make any representation that the contents will be complete or accurate or up to date. The accuracy of any instructions, formulae and drug doses should be independently verified with primary sources. The publisher shall not be liable for any loss, actions, claims, proceedings, demand or costs or damages whatsoever or howsoever caused arising directly or indirectly in connection with or arising out of the use of this material.

AN AMINATED GDP-FUCOSE ANALOG USEFUL IN THE FUCOSYLTRANSFERASE CATALYZED ADDITION OF BIOLOGICAL PROBES ONTO OLIGOSACCHARIDE CHAINS¹

Christer Hällgren and Ole Hindsgaul*

Department of Chemistry, University of Alberta, Edmonton, Alberta, CANADA T6G 2G2

Received September 13, 1994 - Final Form February 13, 1995

ABSTRACT

An analog (1) of GDP-fucose, where C-6 is derivatized with an eight-atom spacer terminating in a primary amino group, was chemically synthesized. This amino group in sugar nucleotide 1 can be acylated using an N-hydroxysuccinimide ester of biotin and it can be coupled to another molecule that also contains an amino group using squaric acid diethyl ester as the coupling reagent. In this way, biotin and a blood group A-active trisaccharide were linked to C-6 of fucose in GDP-fucose. Both complex sugar nucleotides thus prepared were active as donors for a human milk fucosyltransferase, which transferred the derivatized α -linked fucose residue to a glycoside of N-acetyllactosamine, thus labeling this sequence with either biotin or the blood-group A trisaccharide. Compound 1 is proposed as a general and versatile reagent which should permit the addition of biological probes to the sugar chains of cell surface glycoproteins or glycolipids.

INTRODUCTION

A tool allowing the specific addition of biological probes to the oligosaccharide chains of glycolipids and glycoproteins, both isolated and when present at the cell surface, would greatly assist research on the function of glycosylation.^{2,3} For example, if addition of specific oligosaccharide sequences to proteins were possible, this could affect not only their circulation lifetimes, but also result in targeting of specific cells via carbohydrate-recognizing protein-receptors.² Also, if designed biological probes could be added in a mild manner to the cell membrane, the behavior of the resulting tagged cells could be similarly affected.³



Figure 1. Linking of a biological probe of interest to C-6 of the fucose residue in GDPfucose followed by transfer using a fucosyltransferase is proposed as a general method for covalently labeling the sugar chains of glycoproteins and glycolipids.

Since all animal cells contain cell surface glycoproteins and glycolipids with their sugar chains exposed to interact with their external environment, these sugar chains are ideal candidates to which biological probes could be attached. Such probes have been previously chemically attached to cell-surface glycoconjugates (by periodate oxidation followed by reductive amination) in order to add the molecule of interest covalently.⁴ The feasibility of much milder and specific derivatizations has been demonstrated and involves the enzymatic addition of monosaccharide residues tagged with the biological probe of interest. The pioneering work in this area involved the use of sialyltransferases.⁵ Later, fucosyltransferases were used for the same purpose, the strategy being summarized in Figure 1.³ The advantage of using fucosyltransferases is that they can transfer to several oligosaccharide sequences that are already sialylated.

We report here an important simplification of our earlier strategy for labeling cellsurface sugar chains using fucosyltransferases.³ A GDP-fucose derivative labeled at C-6 with the biological probe of interest is essential in this approach (Figure 1). However, the synthesis of such probes was very cumbersome and many chemical steps would need to be repeated for each new probe. In the present work, we prepare a single GDP-fucose analog, the aminated derivative (1), and we demonstrate that complex molecules possessing either an activated carboxyl-group or another amine can be added to 1 in simple high-yielding steps. Furthermore, the GDP-fucose derivatives prepared in this manner are shown to be enzymatically active.

RESULTS AND DISCUSSION

The synthesis of the GDP-fucose analog 1 was straightforward using chemistry well established in the D-galactose series. Briefly, acetonation of L-galactose gave the 1,2:3,4-diacetonide⁶ which was then 6-O-allylated to give 2. Hydrolysis of the isopropylidene group in 2, followed by acetylation gave 3. Attempted conversion of 3 to the glycosyl bromide with HBr failed, but was successful using TiBr₄. Reaction of the resulting glycosyl bromide (not isolated) with tetrabutylammonium phosphate, followed by de-O-acetylation, gave the required 6-O-allyl- β -L-galactopyranosyl phosphate 4. Conventional coupling of 4 with GMP-morpholidate⁷ yielded the sugar nucleotide 5. Reaction of 5 with cysteamine (2-aminothioethanol) under UV-irradiation gave the target 1.

Compound 1 contained the required 6'-amino group which allowed further very simple derivatization. The simplest derivatization methods are reaction with an activated ester and coupling with another amine using a heterobifunctional reagent. To demonstrate that 1 could be derivatized as expected, it was treated with a water soluble *N*-hydroxysuccinimide activated ester of biotin (6) which resulted in the production of 7. To show that 1 could be coupled to other amines, we selected the method of Tietze and Arlt,⁷ using commercial diethyl squarate, which shows extraordinary promise as a reagent for the very simple coupling of two amines. The blood-group-A active trisaccharide, as its synthetic 8-methoxycarbonyloctyl glycoside 8,⁹ was first treated with ethylenediamine to produce the required amine derivative 9. Reaction of 9 with diethyl squarate gave the monoamide 10 which was further reacted with 1 to give the desired A-blood-group-tagged-GDP-Fuc derivative 11 (Scheme 2).

Finally, both of the GDP-Fuc derivatives 7 and 11 proved to be active as donors for a partially purified preparation of fucosyltransferase from human milk.¹⁰ For demonstration purposes, incubations were performed using the *N*-acetyllactosamine derivative 12 which we use as a model for the corresponding terminal oligosaccharide of cell-surface glycoconjugates. Enzymatic glycosylation of 12 with either 7 or 11 as the donor produced the expected tagged oligosaccharides 13 and 14 which were characterized



Scheme 1

by ¹H NMR spectroscopy. The molecular weights of **13** and **14** were confirmed using MALDI-TOF mass spectrometry. The enzyme specificity requires the addition to occur at OH-3 of the GlcNAc residue as shown.

CONCLUSION

In summary, the GDP-fucose derivative 1, reported in this work, is shown to react in a predictable way with active esters and amino-containing molecules to produce







AMINATED GDP-FUCOSE ANALOG

derivatives that can be transferred enzymatically to oligosaccharide chains. This means that, in principle, any molecule containing an activatable carboxyl group, or a free amino group, can be covalently added to the carbohydrate chains of cell-surface glycoconjugates in a very simple and mild manner. Several such experiments are in progress.

EXPERIMENTAL

General methods. Optical rotations were measured with a Perkin-Elmer 241 polarimeter at 22 ± 2 °C. Analytical TLC was performed on Silica Gel 60-F254 (E. Merck, Darmstadt) with detection by quenching of fluorescence and/or by charring with 8% sulfuric acid. The solvent systems used have been designated by letters: A, 3:1 toluene-ethyl acetate; B, 7:3:1 2-propanol-water-concd ammonia; C, 60:35:6 chloroformmethanol-water. All commercial reagents were used as supplied and chromatography solvents were distilled prior to use. Column chromatography was performed on Silica Gel 60 (E. Merck 40-60 µM, Darmstadt). C-18 Silica Gel was from Toronto Research Chemicals Inc. (35-70 µM), Millex-GV (0.22 µM) filter units were from Millipore (Missisuaga, ON), C-18 and QMA Sep-Pak sample preparation cartridges were from Waters Associates (Missisuaga, ON). The biotin active ester 6 was from Pierce (Rockford, IL), ¹H NMR spectra were recorded at 360 MHz (Bruker WM-360), 300 MHz (Bruker AM-300) or 500 MHz (Varian Unity 500) with either internal (CH3)4Si (δ 0, CDCl₃, CD₃OD) or DOH (δ 4.80, D₂O). ¹³C NMR spectra were recorded at 75.5 MHz (Bruker AM-300) with internal (CH3)4Si (δ 0, CDCl3, CD3OD) or external acetone (δ 31.0, D2O). ³¹P NMR spectra were recorded at 81.0 MHz (Bruker WH-200) or 162.0 MHz (Bruker WH-400) with external H₃PO₄ (δ 0, CDCl₃, CD₃OD, D₂O). ¹H NMR data are reported as though they were first order. Protons in the 8-methoxycarbonyloctyl (MCO) spacer are designated H-1 to H-8 with increasing distance from the anomeric center and protons of the allyl group are designated as defined below.



Unless otherwise stated, all reactions were carried out at room temperature. In the processing of reaction mixtures, solutions of organic solvents were washed with half their volumes of aqueous solutions. Organic solutions were dried (sodium sulfate) prior to concentration under vacuum at < 40 °C (bath). Microanalyses were carried out by the analytical services at this department and all samples submitted for elemental analyses were dried overnight under vacuum with phosphorous pentoxide at 56 °C (refluxing acetone).

Matrix assisted laser desorption ionization/time-of-flight (MALDI-TOF) mass spectra were recorded on a home-built instrument using a nitrogen laser and 2,5-dihydroxybenzoic acid as the matrix. The masses recorded have an accuracy of better than ± 1 amu.

6-O-Allyl-1,2:3,4-di-O-isopropylidene- α -L-galactopyranose (2). A solution of 1,2:3,4-di-O-isopropylidene- α -L-galactopyranose (4.25 g, 16.3 mmol) and allyl bromide (1.7 mL, 19.6 mmol) in N,N-dimethylformamide (10 mL) was added dropwise to a vigorously stirred mixture of sodium hydride (735 mg, 24.5 mmol) in $N_{,N}$ dimethylformamide (5 mL). After 30 min, methanol (5 mL) was added and the reaction mixture was stirred for an additional 5 min. The reaction mixture was partitioned between toluene and brine. The organic layer was dried (MgSO4), filtered, and concentrated. Column chromatography (8:1 hexane-ethyl acetate) gave 2 (4.51 g, 92 %): $[\alpha]_{578}$ +72° (c 1.0, chloroform); $R_f 0.61$ (system A); ¹H NMR (CDCl₃) δ 1.33, 1.35, 1.45, 1.55 (s, 3) H each, 4 x CH₃), 3.59 (dd, 1 H, $J_{5,6a} = 6.8$ Hz, $J_{6a,6b} = 10.1$ Hz, H-6a), 3.66 (dd, 1 H, $J_{5,6b} = 5.8$ Hz, H-6b), 3.98 (ddd, 1 H, $J_{4,5} = 1.9$ Hz, H-5) 4.05 (dddd, 2H, $J_{Hd,He}$ 13.5 Hz, $J_{Hc,Hd} = J_{Hc,He} = 6.0$ Hz, $J_{Ha,Hd} = J_{Hb,Hd} = J_{Ha,He} = J_{Hb,He} = 1.5$, H_d , H_e), 4.27 (dd, 1 H, $J_{3,4}$ = 7.9 Hz, H-4), 4.31 (dd, 1 H, $J_{1,2}$ = 5.2 Hz, $J_{2,3}$ = 2.5 Hz, H-2), 4.61 (dd, 1 H, H-3), 5.18 (dddd, 1 H, $J_{Ha,Hc}$ = 10.5 Hz, $J_{Ha,Hb}$ = 1.4 Hz, H_a), 5.28 (dddd, 1 H, $J_{Hb,Hc} = 17.3$ Hz, H_b), 5.55 (dd, 1 H, H-1), 5.93 (dddd, 1H, H_c). Anal. Calcd for C15H24O6: C, 60.0; H, 8.0. Found: C, 60.1; H, 8.1.

6-*O*-Allyl-1,2,3,4-tetra-*O*-acetyl-α,β-L-galactopyranose (3). Compound 2 (3.15 g, 10.5 mmol) was treated with 90% aqueous trifluoroacetic acid (100 mL) for 15 min and then concentrated. The aliquot was coevaporated three times with toluene (100 mL), dissolved in pyridine (75 mL) and treated with acetic anhydride (25 mL) at 60 °C for one h and then concentrated. Column chromatography (6:1 toluene-ethyl acetate) gave **3** (1.82 g, 45 %) as a mixture of anomers: R_f 0.38 (system A); ¹H NMR (CDCl₃) δ 2.00, 2.01, 2.02, 2.04, 2.11, 2.15, 2.16 (s, 3 H each, COC<u>H₃</u>), 5.71 (d, 1 H, J_{1,2} = 8.2 Hz, H-1β), 5.82 (dddd, 1 H, Hc), 6.39 (d, 1 H, J_{1,2} = 3.2 Hz, H-1α).

Anal. Calcd for C17H24O10: C, 52.6; H, 6.2. Found: C, 52.8; H, 6.2.

Bis(triethylammonium) 6-O-Allyl- β -L-galactopyranos-1-yl Phosphate (4). Titanium tetrabromide (3.2 g, 8.8 mmol) was added to a solution of 3 (1.14 g, 2.93 mmol) in dichloromethane-ethyl acetate (10:1, 50 mL) and the reaction mixture was stirred at room temperature for 48 h. Sodium acetate (1 g) was then added, and stirring was continued for 15 min. The mixture was filtered through Celite, diluted with dichloromethane and washed with ice cold water, dried and concentrated. The residue was dissolved in toluene (30 mL) and added to a stirred mixture of tetrabutylammonium phosphate (2.4 g, 6 mmol) in acetonitrile (60 mL) containing ground molecular sieves (4Å). The reaction was stirred an additional 4 h and then concentrated. The residue was

dissolved in water, washed with dichloromethane, concentrated, redissolved in water and passed through Dowex 50W-X8 (NH4⁺ form) resin. The carbohydrate containing fractions were pooled, concentrated, redissolved in a minimum amount of water, loaded on a C-18 Silica-gel column (1.5x36 cm). The column was eluted with water with a flow of 1 mL/min, 7 mL fractions were collected and those fractions containing carbohydrate but not inorganic phosphate (determined by a molybdate-color assay)¹¹ were pooled and lyophilized. The residue was dissolved in methanol-cyclohexylamine (1:1, 60 mL) and refluxed until TLC (system B) indicated complete conversion to a slower moving product. (2 h, $R_f 0.75 \rightarrow 0.68$, system B). The solution was concentrated and partitioned between water and dichloromethane. The water phase was concentrated, redissolved in water and passed through Dowex 50W-X8 (Et₃NH⁺ form) resin. The carbohydrate containing fractions were pooled and lyophilized to give 4 (418 mg, 28%) as a colorless powder: R_f 0.68 (system B); ¹H NMR (D₂O, 400 MHz) δ 1. 28 (t, 18 H, 7.3 Hz, Et₃N), 3.19 (q, 12 H, Et₃N), 3.55 (dd, 1 H, $J_{1,2}$ = 7.7 Hz, $J_{2,3}$ = 10.0 Hz, H-2), 3.65-3.77 (m, 3 H, H-3, H-6a, H-6b), 3.86 (ddd, 1 H, H-5), 3.89 (dd, 1 H, $J_{3,4} = 3.2$ Hz, $J_{4,5} < 1$ Hz, H-4), 4.09 (dddd, 2H, $J_{Hd,He}$ 13.5 Hz, $J_{Hc,Hd} = J_{Hc,He} = 6.0$ Hz, $J_{Ha,Hd} = J_{Hb,Hd} = J_{Ha,He} = J_{Hb,He}$ = 1.5 Hz, H_d, H_e), 4.85 (dd, 1 H, $J_{1,P}$ = 7.8 Hz, H-1), 5.28 (dddd, 1 H, $J_{Ha,Hc}$ = 10.5 Hz, $J_{Ha,Hb} = 1.4$ Hz, H_a), 5.35 (dddd, 1 H, $J_{Hb,Hc} = 17.2$ Hz, H_b), 5.95 (dddd, 1H, H_c); ³¹P NMR (D₂O; 162.0 MHz) δ 1.5

Dilithium 6-0-Allyl-β-L-galactopyranos-1-yl Guanosine 5'-Diphosphate (5). Compound 4 (418 mg, 832 µmol) was dissolved in dry pyridine (10 mL) and added to a suspension of guanosine 5'-monophosphomorpholidate 4-morpholine-N,N'-dicyclohexylcarboxamidine salt (604 mg, 832 µmol, Sigma). The activated nucleoside monophosphate and 4 were previously dried by coevaporating with dry pyridine three times. The mixture was coevaporated with pyridine one more time, then dissolved (suspended) in dry pyridine and vigorously stirred for 3.5 d. The mixture was concentrated, dissolved in a minimum amount of water and loaded onto a Dowex 1X2-200 (Cl⁻ form, 2.5x18 cm) column. The column was eluted with a linear LiCl gradient ($0 \rightarrow$ 0.5 M) made of water (800 mL) and of 0.5 M LiCl (800 mL) at 2 mL/min and 20 mL fractions were collected. The appropriate fractions (eluted at 0.4-0.5 M) were pooled, concentrated and desalted on a Biogel P-2 column (3x90 cm) to give 5 (226.4 mg, 41.4%) as a white powder: $R_f 0.60$ (system B); ¹H NMR (D₂O, 400 MHz) δ 3.62 (dd, 1 H, $J_{1",2"} = 7.5 \text{ Hz}, J_{2",3"} = 10.0 \text{ Hz}, \text{ H-2"}, 3.67-3.75 \text{ (m, 3 H, H-2", H-6")}, 3.86 \text{ (ddd, 1)}$ H, H-5"), 3.91 (dd, 1 H, $J_{3",4"} = 3.3$ Hz, $J_{4",5"} = 0.8$ Hz, H-4"), 4.06 (dddd, 2H, $J_{Hd,He}$ 13.5 Hz, $J_{Hc,Hd} = J_{Hc,He} = 5.9$ Hz, $J_{Ha,Hd} = J_{Hb,Hd} = J_{Ha,He} = J_{Hb,He} = 1.5$ Hz, H_d , H_e), 4.23 (dd, 2H, $J_{4',5a'}$ = 3.6 Hz, $J_{4',5b'}$ = 5.4 Hz, H-5'), 4.37 (ddd, 1 H, $J_{3',4'}$ = 3.1 Hz, H-4'), 4.55 (dd, 1H, $J_{2',3'}$ = 5.2 Hz, H-3'), 4.91 (dd, 1 H, $J_{1'',P}$ = 8.4 Hz, H-

1"), 5.26 (ddd, 1 H, $J_{Ha,Hc} = 10.5$ Hz, $J_{Ha,Hb} = 1.6$ Hz, H_a), 5.33 (ddd, 1 H, $J_{Hb,Hc} = 17.3$ Hz, H_b), 5.93 (ddd, 1H, H_c), 5.94 (d, 1 H, $J_{1',2'} = 7.2$ Hz, H-1'), 8.12 (s, 1H, H-8 guanosine); ³¹P NMR (D₂O; 81.015 MHz) δ -12.6 (d, ³ $J_{P,H}$ 20.6 Hz, Fuc-P), -10.7 (d, Rib-P).

Disodium 6-O-3-(S-2-Aminothioethyl)-propyl-β-L-galactopyranos-1-yl Guanosine 5'-Diphosphate (1). Cysteamine (24 mg, 210 μmol) was added to a solution of 5 (14.1 mg, 21.4 μmol) in water (3 mL) and irradiated for 2 h in a quartz vial using a Hanovia UV-lamp ($R_f 0.60 \rightarrow 0.36$ system B). The reaction mixture was passed through a Dowex 50W-X8 (Na⁺ form) resin and the appropriate fractions were collected, concentrated, redissolved in a minimum amount of water and desalted on a Biogel P-2 column (3x90 cm) to give 1 (11.8 mg, 69%): $R_f 0.36$ (system B); ¹H NMR (D₂O, 300 MHz) δ 1.83 (quintet, 2 H, NCH₂CH₂CH₂S), 2.63 (t, 2 H, CH₂), 2.84 (t, 2 H, CH₂), 3.22 (t, 2 H, CH₂), 3.58-3.77 (m, 4 H, H-2", H-3", H-5", H-6"), 3.86 (ddd, 2 H, H-5""), 3.91 (dd, 1 H, $J_{3",4"} = 3.4$ Hz, $J_{4",5"} < 1$ Hz, H-4"), 4.24 (dd, 2 H, $J_{4',5'} = 3.5$ Hz, $J_{4',5'} = 5.2$ Hz, H-5'), 4.37 (ddd, 1 H, H-4'), 4.54 (dd, 1 H, $J_{2',3'} = 5.0$ Hz, $J_{3',4'} = 3.4$ Hz, H-3'), 4.98 (dd, 1 H, $J_{1",2"} = J_{1",P} = 7.5$ Hz, H-1"), 5.95 (d, 1H, $J_{1',2'} = 6.1$ Hz, H-1'), 8.13 (s, 1 H, H-8 guanosine).

Disodium 6-0-3-[*S*-2-{6-(Biotinamido) Hexanoate} Amidothioethyl]propyl- β -L-galactopyranos-1-yl Guanosine 5'-Diphosphate (7). Sulfosuccinimidyl 6-(biotinamido) hexanoate (6, 6.6 mg, 7.3 µmol) was added to a stirred solution of 1 (8.6 mg, 6.6 µmol) in saturated aqueous sodium hydrogencarbonate (0.5 mL, pH 9.45) and stirred overnight (R_f 0.36 \rightarrow 0.31 system B). The reaction mixture was loaded directly on a Biogel P-2 column (3x90 cm) and gave 7 (9.1 mg, 80%): R_f 0.31 (system B); ¹H NMR (D₂O, 360 MHz) δ 1.84 (q, 2H, 7.3 Hz), 2.19 (t, 2 H, 7.5 Hz), 2.26 (t, 2 H, 7.4 Hz), 3.60-3.75 (m, 3 H, H-2", H-3", H-6"a, H-6"b), 3.86 (ddd, 1 H, H-5"), 3.90 (dd, J_{3,4} = 3.3 Hz, J_{4,5} < 1 Hz, H-4"), 4.22 (dd, 2 H, H-5'), 4.36 (ddd, 1 H, H-4'), 4.44 (dd, 1 H, J = 7.9 Hz, J = 4.4 Hz), 4.54 (dd, 1 H, J_{2',3'} = 5.1 Hz, J_{3',4'} = 3.2 Hz, H-3'), 4.62 (dd, 1 H, J = 4.4 Hz, J = 7.8 Hz), 4.96 (dd, 1 H, J_{1",2"} = J_{1",P} = 7.7 Hz, H-1"), 5.94 (d, 1 H, J_{1',2'} = 6.2 Hz, H-1'), 8.12 (s, 1 H, H-8 guanosine).

8-(N-2-Aminoethyl)-carboxamidooctyl 3-O-(2-Acetamido-2-deoxy- α -D-galactopyranosyl)-[2-O-(α -L-fucopyranosyl)- β -D-galactopyranoside (9). Trisaccharide ethyl ester 8⁹ (57.4 mg, 80.4 mmol) was treated with ethylenediamine (2 mL) at 70 °C for 2 d (Rf 0.64 \rightarrow 0.45 system B). The solution was diluted with water (20 mL), cooled to room temperature and loaded onto two consecutive Sep-Pak C-18 cartridges. The cartridges were washed with water (20 mL) and 9 was eluted with methanol (20 mL), concentrated, redissolved in water (7 mL), and filtered through a Millipore filter. The washings were loaded onto another Sep-Pac cartridge and taken through the same cycle of washing, elution and filtering. Lyophilization of both fractions gave **9** (57.6 mg, 98%) as a white powder: $R_f 0.45$ (system B); ¹H NMR (D₂O, 360 MHz) δ 1.24 (d, 1 H, $J_{5,6}$ = 6.6 Hz, H-6 fucose), 1.33 (m, 8 H, H-3 to H-6, MCO), 1.62 (m, 4 H, H-2, H-7 MCO), 2.06 (s, 3 H, acetamido CH₃), 2.28 (t, 2 H, H-8 MCO), 2.89 (t, 2 H, C<u>H</u>₂NHCO), 3.35 (t, 2 H, C<u>H</u>₂NH₂), 4.49 (q, 1 H, $J_{4,5}$ < 1 Hz, H-5 fucose), 4.55 (d, 1 H, $J_{1,2}$ = 7.8 Hz, H-1 Gal), 5.19 (d, 1 H, $J_{1,2}$ = 3.9 Hz, H-1 GalNAc[†]), 5.34 (d, 1 H, $J_{1,2}$ = 3.6 Hz, H-1 Fuc[†]); [†] may be interchanged.

Squaric acid coupling of 9 and 1 to give 11. 3,4-Diethoxy-3-cyclobutene-1,2-dione (diethyl squarate, Aldrich, 1.88 µL, 13.5 µmol) was added to a stirred mixture of 8 (10.3 mg, 14.1 μ mol) in ethanol (1 mL). After 3 h, TLC showed complete conversion to a faster moving spot (presumably 10, $R_f 0.05 \rightarrow 0.73$ system C). Compound 1 (9.0 mg, 11.2 µmol) dissolved in 1 mL of 0.1 M sodium bicarbonate buffer (pH 9.5) was added and stirring was continued for 16 h. The solution was diluted with water (5 mL) and loaded onto a Sep-Pak QMA cartridge. The cartridge was washed with water (7 mL), then the product was eluted with 3M sodium chloride solution (10 mL). The eluent was loaded onto a Sep-Pak C-18 cartridge and the cartridge was washed with water (10 mL) and 11 was eluted with methanol (10 mL), concentrated, redissolved in water (7 mL), and filtered through a Millipore filter. Lyophilization gave 11 (3.7 mg, 98%) as a white powder: $R_f 0.40$ (system B); ¹H NMR (D₂O, 360 MHz) δ 1.23 (d, 1 H, $J_{5,6} = 6.5$ Hz, H-6 fucose), 1.26 (m, 8 H, H-3 to H-6 MCO), 1.49 (m, 2 H, H-2 MCO), 1.58 (quintet, 2 H, H-7 MCO), 1.82 (quintet, 2 H, NCH₂CH₂CH₂O), 2.05 (s, 3 H, acetamido CH₃), 2.19 (t, 2 H, H-8 MCO), 2.63 (t, 2 H, CH₂), 2.80 (t, 2 H, CH₂), 3.42 (t, 2 H, CH₂), 4.35 (ddd, 1 H, H-4 ribose), 4.47 (q, 1 H, $J_{4,5} < 1$ Hz, H-5 fucose), 4.51-4.54 (m, 2 H, H-1 D-Gal, H-3 Rib), 4.97 (dd, 1 H, $J_{1,2} = J_{1,P} = 7.8$ Hz, H-1 L-Gal), 5.18 (d, 1 H, $J_{1,2}$ = 3.8 Hz, H-1 GalNAc[†]), 5.33 (d, 1 H, $J_{1,2}$ = 3.7 Hz, H-1 Fuc[†]), 5.92 (d, 1 H, $J_{1,2} = 6.2$ Hz, H-1 Rib), 8.11 (s, 1 H, H-8 guanosine); [†] may be interchanged

Enzymatic Synthesis of 13 Using 7 as the Donor. α -3/4-Fucosyltransferase¹⁰ (100 µL, 200 mU/mL) was added to a solution of LacNAc-MCO (12) (0.49 mg, 0.89 µmol) and 7 (0.59 mg, 0.60 µmol) in 440 µL buffer (20 mM HEPES, 20 mM MnCl₂, 0.2 % BSA, pH = 7) and incubated at 37 °C for 10 h (R_f 0.52 \rightarrow 0.45 system C). Extra donor (1.00 mg) and extra enzyme (50 µL) were added and the solution was incubated for an additional 10 h. The solution was diluted with water (3 mL) and loaded onto a Sep-Pak C-18 cartridge, the cartridge was washed with water (30 mL) and 13 was eluted with methanol (10 mL), concentrated, redissolved in water (7 mL), and filtered through a Millipore filter. Lyophilization gave 13 (0.6 mg, 65%) as a white powder: R_f 0.45 (system C); ¹H NMR (D₂O, 360 MHz) δ 5.218 (d, 1H, J_{1,2} = 4.0 Hz, H-1 Fuc), 2.030 (s, 3H, NHAc). MALDI-TOF MS: M + Na⁺ obs. = 1195.4 (calcd C₅₁H₈₉O₂₁N₅S_{2 + Na⁺} = 1194.6). Enzymatic Synthesis of 14 Using 11 as the Donor. α -3/4-Fucosyltransferase¹⁰ (100 µL, 200 mU/mL) was added to a solution of LacNAc-MCO (12) (0.47 mg, 0.85 µmol) and 11 (0.90 mg, 0.60 µmol) in 440 µL buffer (20 mM HEPES, 20 mM MnCl₂, 0.2 % BSA, pH = 7) and incubated at 37 °C for 16 h (R_f 0.52 \rightarrow 0.35 system C). Extra donor (1.00 mg, 0.94 µmol) and extra enzyme (50 µL) were added and the solution was incubated for an additional 16 h. The solution was diluted with water (3 mL) and loaded onto a Sep-Pak C-18 cartridge. The cartridge was washed with water (30 mL) and 14 was eluted with methanol (10 mL), concentrated, redissolved in water (7 mL), and filtered through a Millipore filter. Lyophilization gave 14 (0.9 mg, 65%) as a white powder: R_f 0.35 (system C); ¹H NMR (D₂O, 500 MHz) δ 5.323, 5.225 and 5.175 (each d, 1H, J_{1,2} = 4.0 Hz, H-1 α), 4.530 (J_{1,2} = 7.9 Hz), 4.472 (J_{1,2} = 7.0 Hz) and 4.457 (J_{1,2} = 7.8 Hz) (each H-1 β), 2.044 and 2.031 (each s, 3H, NHAc). MALDI-TOF MS: M + Na⁺ obs. = 1660.3 (calcd C₇₀H₁₁₉O₃₆N₅S + Na⁺ = 1660.7).

ACKNOWLEDGMENTS

This work was supported by the Medical Research Council of Canada. "Stiftelsen Bengt Lundqvists Minne" is gratefully acknowledged for financial support to C.H. We thank Professor M. M. Palcic for a generous gift of the fucosyltransferase and Randy Whittal for recording the MALDI-TOF mass spectra.

REFERENCES AND NOTES

- 1. Presented as Abstract B1.64 at the XVIIth International Carbohydrate Symposium, Ottawa, Canada, July 17-22, 1994.
- 2. A. Varki, Glycobiology, 3, 97 (1993).
- G. Srivastava, K. J. Kaur, O. Hindsgaul and M. M. Palcic, J. Biol. Chem., 267, 22356 (1992).
- 4. C. G. Gamberg and M. Tolvanen, Anal. Biochem., 170, 520 (1988).
- H. J. Gross, J. C. Bunsch, J. C. Paulson and R. Brossmer, Eur. J. Biochem., 168, 595 (1987).
- 6. R. S. Tipson, Methods Carbohydr. Chem., II, 246 (1963).
- 7. S. Roseman, J. J. Distler, J. G. Moffat and H. G. Khorana, J. Am. Chem. Soc., 83, 659 (1961).
- 8. L. Tietze, M. Arlt, M. Beller, K.-H. Glusenkamp, E. Jahde and M. F. Rajewsky, *Chem. Ber.*, **124**,1215 (1991).
- 9. R. U. Lemieux, Chem. Soc. Rev., 7 423 (1978).
- M. M. Palcic, A. P. Venot, R. M. Ratcliffe and O. Hindsgaul, *Carbohydr. Res.*, 190, 1 (1989).
- 11. F. Seela and S. Waldek, Nucleic Acids Res., 2, 2343 (1975).