Supporting Information

for

Convergent synthetic methodology for the construction of self-adjuvanting lipopeptide vaccines using a novel carbohydrate scaffold

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Experimental part

Table of Contents	Page
Experimental (General)	S 3
Synthesis of Compound 3	S 5
Synthesis of Compound 1	S 5
Characterization of Compounds 4	S 6
Characterization of Compounds 5	S 7
Synthesis of Compound 8	S 7
Synthesis of Peptides and Lipopeptides (Standard Fmoc SPPS conditions)	S 8
N ₃ -J8 epitope	S 9
Glyco-lipopeptide 9	S9
Glyco-lipopeptide 11	S10
Copper Catalyzed Cycloaddition "Click" Reaction	S10
Vaccine candidate 10	S10
Vaccine candidate 12	S 11
NMR Compound 4	S12
NMR Compound 5	S15
NMR Compound 1	S18
Analytical RP-HPLC N ₃ -J8	S21
Mass Spectrum N ₃ -J8	S22
Analytical RP-HPLC Vaccine Candidate 10	S23
Mass Spectrum Vaccine Candidate 10	S24
Analytical RP-HPLC Vaccine Candidate 12	S25
Mass Spectrum Vaccine Candidate 12	S26
References	S27

Experimental

General

Methanol, dimethylformamide (DMF), diethyl ether (Et₂O), acetonitrile, ethyl acetate (EtOAc), hexane, trifluoroacetic acid (TFA), piperidine and O-benzotriazol-1-yl)-N.N.N',N'tetramethyluronium hexafluorophosphate (HBTU) were obtained from Merck KGaA (64271 Darmstadt, Germany). Dichloromethane, dimethyl sulfoxide (DMSO), NaOH and sodium hydrogen carbonate (NaHCO₃) were purchased from Chem-Supply (Adelaide, 5013 Australia). Alfa Aesar (Heysham LA3 2XY, England) supplied NaH 57–63% oil dispersion. O-(7-Azabenzotriazol-1-yl)-N,N,N',N'-tetramethyluronium hexafluorophosphate (HATU) and all N-a-Fmoc-protected amino acids were supplied by Mimotopes (Victoria, 3168 Australia). Rink amide 4-methylbenzhydrylamine (MBHA) resin was purchased from Novabiochem (Läufelfingen, Switzerland). Ultra-pure N₂ gas was supplied by BOC Gases (Brisbane, QLD, Australia). Automated flash chromatography was carried out using a Reveleris[®] X2 flash chromatography system (Grace Materials Technologies, Discovery Sciences) with UV and Evaporative Light Scattering Detection (ELSD). Davisil[®] Chromatographic Silica Media (LC60A 20-45 micron) was purchased from Grace Materials Technologies, Discovery Sciences (Victoria 3178, Australia) and used for dry column vacuum chromatography (DCVC). Deuterated chloroform (CDCl₃) was purchased from Cambridge Isotope Laboratories Inc. (Andover, MA, USA). All other reagents were purchased in analytical grade or higher purity from Sigma-Aldrich (Castle Hill, VIC, Australia). All solvents and reagents were used as purchased unless otherwise stated.

Microwave-assisted peptide couplings were performed using a CEM Discover Bio System manual peptide synthesiser (Matthews, USA). Mass spectra (MS) were obtained from a quadrupole electrospray (Perkin Elmer Sciex API 3000 instrument) in the positive ion mode using Analyst 1.4 (Applied Biosystems/MDS Sciex, Toronto, Canada) software. High resolution mass spectra (HRMS) were run on an ABSCIEX Triple TOF 5600 Mass Spectrometer. Optical rotations were performed on a JASCO P-2000 polarimeter. Nuclear magnetic resonance (NMR) spectra were recorded on a Bruker AM 300 MHz (1D spectra) or Bruker AM 500 MHz (2D spectra) instrument, using CDCl₃ as deuterated solvent with reference to tetramethylsilane as internal standard, at 297 K. Coupling constants are given in Hertz (Hz).

Analytical RP-HPLC was performed on a Shimadzu (Kyoto, Japan) instrument (LC-20AT liquid chromatograph, CMB-20A communication bus module, SIL-20A HT auto sampler, CTO-20 column oven, SPD-M20A diode array detector, ELSD-LT II low temperature evaporative light scattering detector, LabSolutions software), with C18 or C4 Grace Vydac[®] (Columbia, Maryland, USA) columns (10 μ m, 4.6 mm internal diameter × 250 mm,) using a stated gradient of CH₃CN/10% H₂O/0.1% TFA (solvent B) and H₂O/0.1% TFA (solvent A) with a 1 mL/min flow rate and detection at 214 nm.

Semi-preparative RP-HPLC was performed using a Waters Delta 600 system (Milford, Massachusetts, USA; model 600 controller, 490E UV detector). Purifications were achieved using Vydac[®] C4 or C18 preparative columns (10 μ m, 10 mm i.d. × 250 mm) using a stated gradient quoted as % solvent B at 4 mL/min flow rate, and detection at 214 nm. Fractions containing pure compound were pooled and lyophilized overnight.

Preparative RP-HPLC was performed on a Shimadzu (Kyoto, Japan) instrument (LC-20AP liquid chromatograph, CMB-20A communication bus module, SPD-20 UV–vis detector, FRC-10A fraction collector), with C18 or C4 Grace Vydac[®] (Columbia, Maryland, USA) columns (10 μ m, 22 mm internal diameter × 250 mm,) using a stated gradient quoted as % solvent B with a 20 mL/min flow rate and detection at 214 nm.

Methyl 6-hydroxyhexanoate (3):

Concentrated sulfuric acid (1 mL) was added to a solution of ε -caprolactone (10.00 g, 87.61 mmol) in CH₃OH (30 mL) and the mixture was stirred at room temperature for 30 min. The mixture was diluted with Et₂O (100 mL), and the organic phase was washed with sat. NaHCO₃ (2 x 50 mL) and H₂O (50 mL). The combined organic extracts were dried (MgSO₄) and evaporated to give compound **3** as a colorless oil (10.37 g, 81%); ¹H NMR (CDCl₃, 300 MHz) δ 3.63 (s, 3H, OCH₃), 3.60 (t, *J* = 6.5 Hz, 2H, CH₂-6), 2.29 (t, *J* = 7.4 Hz, 2H, CH₂-2), 1.67-1.50 (m, 4H, 2 x CH₂), 1.41-1.30 (m, 2H, CH₂); ¹³C NMR (CDCl₃, 75 MHz) δ 172.0 (C=O), 62.5 (HOCH₂), 51.5 (OCH₃), 34.0, 32.3, 25.3, 24.6 (all CH₂) [1].

2,3,4,6-Tetra-*O*-propargyl-1-*O*-(6-hydroxy-6-oxohexyl)-β-D-glucopyranose (1):

Tetra-*O*-acetyl- α -D-glucopyranosyl bromide **2** [2], methyl 6-hydroxyhexanoate **3** (1.71 g, 11.69 mmol), CH₂Cl₂ (30 mL) and molecular sieves (4 Å, approx. 4 g) were stirred under N₂ at room temperature overnight. Silver(I) oxide (3.61 g, 15.58 mmol) was added and the mixture was stirred at room temperature for 5 h, which resulted in a mixture of orthoester by-product **4** and the desired glycosylation product **5**. TMSOTf (0.14 mL, 0.79 mmol) was added to the reaction mixture and, after 5 min stirring at room temperature, the solids were removed by filtration through a bed of Celite[®]. The filtrate was washed with sat. NaHCO₃ (2 x 50 mL), dried (MgSO₄) and evaporated. The residue was dissolved in CH₃OH (50 mL), Na (0.30 g, 13 mmol) was added and the mixture was stirred under N₂ for 20 min. The mixture was neutralized using Amberlite[®] IR-120H ion-exchange resin, filtered and the solvent evaporated. Water (50 mL) was added to the residue and the aqueous phase was washed with Et₂O (2 x 50 mL). The aqueous solution was lyophilized and the residue was dissolved in dry THF (50 mL). Sodium hydride (57-63% oil dispersion, 3.12 g, 77.90 mmol) was added and the mixture under N₂ for 20 min. Following addition of propargyl

bromide (80% in toluene, 13.02 mL, 116.85 mmol), the reaction mixture was stirred at room temperature overnight. Sodium hydroxide (2 M, 50 mL) was added and the mixture was stirred at room temperature for 1 h. The reaction mixture was transferred to a separating funnel and the basic aqueous phase was washed with Et₂O (3 x 50 mL). The basic phase was acidified with glacial acetic acid and the product was extracted into CH₂Cl₂ (3 x 50 mL). The organic phase was washed with water (50 mL), dried (MgSO₄), filtered and evaporated. The residue was purified by automated flash chromatography (0-10% EtOAc in CH₂Cl₂ over 50 mins) to give compound 1 (1.07 g, 31%) as a colorless oil; $[\alpha]^{24,WI}$ +3 (c = 0.6, CHCl₃); R_f 0.42 (EtOAc/CH₂Cl₂, 1:4); ¹H NMR (CDCl₃, 300 MHz) δ 4.58-4.34 (m, 6H, 3 x OCH₂CCH), 4.29-4.16 (m, 3H, H-1, OCH₂CCH), 3.94-3.83 (m, 2H, OCHH, CHH-6), 3.77 (dd, J = 10.7, 4.7 Hz, 1H, CHH-6), 3.58-3.29 (m, 5H, H-2, H-3, H-4, H-5, OCHH), 2.48-2.44 (m, 4H, 4 x CCH), 2.37 (t, J = 7.4 Hz, 2H, CH₂CO₂CH₃), 1.72-1.59 (m, 4H, 2 x CH₂), 1.48-1.38 (m, 2H, CH₂); ¹³C NMR (CDCl₃, 75 MHz) δ 179.1 (C=O), 103.0 (C-1), 83.2 (C-3), 81.3 (C-2), 80.0, 79.9, 79.8 (all CCH), 79.5 (CCH), 77.3 (CCH), 76.1 (C-4), 74.8, 74.5, 74.4 (all CCH), 74.0 (C-5), 69.7 (OCH₂), 68.4 (C-6), 60.3, 60.1, 59.4, 58.7 (all CH₂CCH), 33.8 (CH₂CO₂CH₃), 29.2, 25.5, 24.4 (all CH₂); HRMS-ESI (m/z): [M + NH₄]⁺ calcd for C₂₄H₃₄NO₈, 464.2284; found 464.229.

Although orthoester byproduct **4** and glycosylation product **5** were not purified, analytical samples were obtained for characterization;

3,4,6-Tri-O-acetyl-1,2-O-[1-(6-methoxy-6-oxohexyloxy)ethylidene]-a-D-glucopyranose

(4); colorless oil; $[\alpha]^{24,WI} + 26$ (c = 0.6, CHCl₃); $R_f 0.42$ (EtOAc/hexane, 1:1); ¹H NMR (CDCl₃, 300 MHz) δ 5.71 (d, J = 5.4 Hz, 1H, H-1), 5.19 (t, J = 2.9 Hz, 1H, H-3), 4.91 (ddd, J = 9.6, 2.7, 0.9 Hz, 1H, H-4), 4.30 (ddd, J = 5.4, 3.0, 0.9 Hz, 1H, H-2), 4.21-4.19 (m, 2H, CH₂-6), 3.98-3.91 (m, 1H, H-5), 3.67 (s, 3H, OCH₃), 3.47 (t, J = 6.6 Hz, 2H, OCH₂), 2.31 (t,

J = 7.5 Hz, 2H, CH₂CO₂CH₃), 2.12, 2.10, 2.09 (all s, 3H, COCH₃), 3.05 (s, 3H, CH₃), 1.69-1.54 (m, 4H, 2 x CH₂), 1.43-1.33 (m, 2H, CH₂); ¹³C NMR (CDCl₃, 75 MHz) δ 174.0, 170.7, 169.7, 169.2 (all C=O), 121.29 (CCH₃), 96.9 (C-1), 73.1 (C-2), 70.14 (C-3), 68.2 (C-4), 67.0 (C-5), 66.3 (OCH₂), 63.1 (C-6), 51.5 (OCH₃), 33.9 (CH₂CO₂CH₃), 29.3, 25.6, 24.6 (all CH₂), 20.8 (4 x CH₃); HRMS-ESI (*m*/*z*): [M + Na]⁺ calcd for C₂₁H₃₂NaO₁₂, 499.1791; found 499.1780.

2,3,4,6-Tetra-*O*-acetyl-1-*O*-(6-methoxy-6-oxohexyl)-β-D-glucopyranose (5);

Colorless oil; $[\alpha]^{24,W1}$ -16 (*c* = 0.3, CHCl₃); *R*_f 0.34 (EtOAc/hexane, 1:1); ¹H NMR (CDCl₃, 300 MHz) δ 5.20 (t, *J* = 9.5 Hz, 1H, H-3), 5.08 (t, *J* = 9.6 Hz, 1H, H-4), 4.98 (dd, *J* = 9.5, 8.0 Hz, 1H, H-2), 4.48 (d, *J* = 7.8 Hz, 1H, H-1), 4.26 (dd, *J* = 12.3, 4.8 Hz, 1H, CHH-6), 4.13 (dd, *J* = 12.3, 2.4 Hz, 1H, CHH-6), 3.87 (dt, *J* = 9.6, 6.2 Hz, 1H, OCHH), 3.71-3.66 (m, 4H, H-5, OCH₃), 3.47 (dt, *J* = 9.6, 6.8 Hz, 1H, OCH*H*), 2.30 (t, *J* = 7.4 Hz, 2H, C*H*₂CO₂CH₃), 2.08, 2.04, 2.02, 2.00 (all s, 3H, COCH₃), 1.68-1.55 (m, 4H, 2 x CH₂), 1.40-1.32 (m, 2H, CH₂); ¹³C NMR (CDCl₃, 75 MHz) δ 174.0, 170.7, 170.3, 169.4, 169.3 (all C=O), 100.8 (C-1), 72.9 (C-3), 71.8 (C-5), 71.3 (C-2), 69.8 (OCH₂), 68.5 (C-4), 62.0 (C-6), 51.5 (OCH₃), 33.9 (CH₂CO₂CH₃), 29.1, 25.4, 24.5 (all CH₂), 20.8, 20.64, 20.63, 20.61 (all COCH₃) [3]; HRMS-ESI (*m*/*z*): [M + Na]⁺ calcd for C₂₁H₃₂NaO₁₂, 499.1791; found 499.1802.

Fmoc-Lysine(lauroyl)-OH (K(C₁₂)) (8):

Molecular sieves (4 Å, approx. 4 g) were added to a solution of *para*-toluenesulfonic acid monohydrate (1.36 g, 7.15 mmol) and DIPEA (3.10 mL, 17.80 mmol) in CH₂Cl₂ (30 mL), and the mixture was stirred under N₂ at room temperature for 5 h. Lauroyl chloride (1.40 mL, 6.05 mmol) was added and stirring was continued for 15 min, followed by addition of Fmoc-Lys-OH (2.00 g, 5.43 mmol) and stirring overnight. The solids were removed by filtration through a bed of Celite[®] and the filtrate was washed with acetic acid (5%, 3 x 50 mL), H₂O (50 mL x 2), dried (MgSO₄) and evaporated. Recrystallization from acetonitrile gave compound **8** (1.96 g, 66%) as a white solid; $[\alpha]^{24,W1}$ +22 (c = 1.0, CHCl₃); ¹H NMR (CDCl₃, 300 MHz) δ 7.75 (d, J = 7.5 Hz, 2H, 2 x Ar-H, 7.62-7.58 (m, 2H, 2 x Ar-H), 7.34 (t, J = 7.4, 2H, 2 x Ar-H), 7.31 (d, J = 7.2, 2H, 2 x Ar-H), 5.80-5.75 (m, 2H), 4.40-4.32 (m, 2H), 4.21 (t, J = 7.2 Hz, 1H), 3.31-3.22 (m, 2H), 2.17 (t, J = 7.5, 2H), 1.98-1.71 (m, 2H), 1.64-1.50 (m, 4H), 1.46-1.40 (m, 2H), 1.29-1.18 (m, 16H), 0.88 (t, J = 6.6 Hz, 3H); ¹³C NMR (CDCl₃, 75 MHz) δ 174.8, 174.4, 156.3 (all C=O), 143.7, 141.3 (2 x Ar-C), 127.7, 127.1, 125.1, 119.9 (all Ar-CH), 67.1 (CH₂), 53.6 (CH), 47.1 (CH), 39.1, 36.8, 31.9, 29.7, 29.6, 29.5, 29.4, 29.33, 29.29, 29.0, 25.8, 22.7, 22.2 (all CH₂), 14.1 (CH₃) [4]; HRMS-ESI (m/z): [M + H]⁺ calcd for C₃₃H₄₇N₂O₅, 551.3485; found 551.3475.

Synthesis of Peptides and Lipopeptides

Stepwise SPPS (manual and microwave-assisted) was performed using rink amide MBHA resin (substitution ratio: 0.60 mmol/g) with standard Fmoc SPPS chemistry:

Fmoc-amino acids (4.2 equiv.) were pre-activated with HATU (0.4 M in DMF, 4 equiv.), and DIPEA (6.2 equiv.). Coupling cycles consisted of Fmoc deprotection with piperidine (20% in DMF, 2 x 15 min (manual) or 2 x 2 min, 70 °C (microwave)), DMF washing, followed by coupling with pre-activated Fmoc-amino acids (2 x 30 min (manual) or 2 x 5 min, 70 °C (microwave)). After assembly of the required sequence, the peptidyl-resin was washed with DMF, CH_2Cl_2 and dried under vacuum. Peptides were cleaved from the resin by stirring in a solution of TFA, triisopropylsilane (TIPS) and H_2O (95:2.5:2.5) for 3 h. The resin was removed by filtration and the excess TFA was evaporated by a stream of N_2 . The peptides were precipitated from ice-cold Et_2O , centrifuged and the Et_2O decanted.

N₃-J8 epitope:

The J8 B cell epitope (QAEDKVKQSREAKKQVEKALKQLEDKVQ) was synthesized using the above manual standard Fmoc SPPS conditions. Azidoacetic acid [5] was coupled in triplicate to the N-terminus of the J8 B cell epitope using the same conditions as those used for Fmoc-amino acid couplings. Following cleavage from the resin and precipitation, the crude N_3 -J8 epitope was dissolved in distilled H₂O, lyophilized and purified by preparative RP-HPLC on a C18 column with a gradient of 0-25 % over 10 min, 25-30% solvent B over 30 min (0.633 g, 47% purified yield).

HPLC analysis (C18 column): $t_R = 28.39 \text{ min} (0\% \text{ for 5 min}, 0-20\% \text{ over 5 min}, 20-50\%$ solvent B over 30 min); MS-ESI (*m*/*z*): $[M + 2H]^{2+}$ calcd 1683.4, found 1683.9; $[M + 3H]^{3+}$ calcd 1122.6, found 1122.8; $[M + 4H]^{4+}$ calcd 842.2, found 842.6; $[M + 5H]^{5+}$ calcd 674.0, found 674.2; MW 3364.8 g/mol.

Glyco-lipopeptide (9):

The LLCP lipopeptide (K(C12)GK(C12)K(C12)G) was synthesized using microwaveassisted standard Fmoc SPPS conditions. Carbohydrate carrier **1** (2.5 equiv.) was preactivated with HBTU (2.4 equiv.) and DIPEA (4.0 equiv.), and coupled to the LLCP lipopeptide with microwave assistance (2 x 10 min at 70 °C), to give glyco-lipopeptide **9**. Following cleavage from the resin and precipitation, the crude glyco-lipopeptide **9** was dissolved in a minimum volume of DMF, diluted with solvent A and B (1:1, 100 mL), and lyophilized. Glyco-lipopeptide **9** was used without purification in the proceeding step. HPLC analysis (C4 column): $t_R = 25.31$ (0-50% over 5 min, 50-100% solvent B over 30 min); MS-ESI (*m/z*): [M + H]⁺ calcd 1491.0, found 1491.6; [M + 2H]²⁺calcd 746.1, found 746.5; MW 1490.0 g/mol.

Glyco-lipopeptide (11):

The T-helper-LLCP lipopeptide (KFVAAWTLKAA-K(C12)GK(C12)K(C12)G) was synthesized using microwave-assisted standard Fmoc SPPS conditions. Carbohydrate carrier **1** was coupled as described above, to give glyco-lipopeptide **11**. Following cleavage from the resin and precipitation, the crude glyco-lipopeptide **11** was dissolved in a minimum volume of DMF, diluted with solvent A and B (1:1, 100 mL), and lyophilized. Glyco-lipopeptide **11** was used without purification in the proceeding step. HPLC analysis (C4 column): $t_R = 33.50$ (0-50% over 10 min, 50-80% solvent B over 30 min); MS-ESI (*m/z*), [M + H]⁺ *mz* calcd 2679.5, found 2679.9; [M + 2H]²⁺ calcd 1340.3, found 1340.6), [M + 3H]³⁺ calcd 893.8, found 894.3; MW 2678.5 g/mol.

Copper Catalyzed Cycloaddition "Click" Reaction

Vaccine candidate (10):

Copper wire (0.100 g) was added to a solution of N₃-J8 (44.0 mg, 13.0 μ mol) epitope and crude glyco-lipopeptide **9** (2.0 mg, 1.3 μ mol) in DMF (3 mL), and the mixture was stirred at 45 °C for 1 h. The copper wire was removed by filtration and the DMF filtrate was loaded directly onto a preparative C4 column (RP-HPLC). The product **10** was eluted with a gradient of 0% for 25 min, 0-20% over 5 min, 20-60% solvent B over 60 min (4.1 mg, 32% purified yield).

HPLC analysis (C4 column): $t_R = 28.59 (0\%$ for 5 min, 0-20% over 5 min, 20-60% solvent B over 30 min); MS-ESI (*m/z*): $[M + 5H]^{5+}$ calcd 2991.0, found 2990.2; $[M + 6H]^{6+}$ calcd 2492.7, found 2494.2; $[M + 7H]^{7+}$ calcd 2136.7, found 2137.9; $[M + 8H]^{8+}$ calcd 1869.8, 1870.5; $[M + 9H]^{9+}$ calcd 1662.1, found 1663.2; $[M + 10H]^{10+}$ calcd 1496.0, found 1497.3; $[M + 11H]^{11+}$ calcd 1360.1, found 1360.6; $[M + 12H]^{12+}$ calcd 1246.9, found 1247.7; $[M + 13H]^{13+}$ calcd 1151.0, found 1151.5; MW 14950.2 g/mol.

Vaccine candidate (12):

Copper wire (0.100 g) was added to a solution of N₃-J8 (30.3 mg, 9.0 μ mol) epitope and crude glyco-lipopeptide **11** (2.4 mg, 0.9 μ mol) in DMF (3 mL), and the mixture was stirred at 45 °C for 1 h. The copper wire was removed by filtration and the DMF filtrate was loaded directly onto a preparative C4 column (RP-HPLC). The product **12** was eluted with a gradient of 0% for 25 min, 0-20% over 5 min, 20-60% solvent B over 60 min (3.5 mg, 24% purified yield).

HPLC analysis (C4 column): $t_R = 31.85$ (0% for 5 min, 0-20% over 5 min, 20-60% solvent B over 30 min); MS-ESI (*m*/*z*): $[M + 9H]^{9+}$ calcd 1794.1, found 1794.9; $[M + 10H]^{10+}$ calcd 1614.8, found 1614.9; $[M + 11H]^{11+}$ calcd 1468.1, found 1468.5; $[M + 12H]^{12+}$ calcd 1345.8, found 1346.3; $[M + 13H]^{13+}$ calcd 1242.4, found 1243.1; $[M + 14H]^{14+}$ calcd 1153.7, found 1154.2; $[M + 15H]^{15+}$ calcd 1076.8, found 1077.3; $[M + 16H]^{16+}$ calcd 1009.6, found 1010.2; $[M + 17H]^{17+}$ calcd 950.3, found 950.5; MW 16137.7 g/mol.

3,4,6-Tri-*O*-acetyl-1,2-*O*-[1-(6-methoxy-6-oxohexyloxy)ethylidene]-α-D-glucopyranose (4)



¹H NMR (CDCl₃, 300 MHz)



 $\textbf{3,4,6-Tri-}\textit{O}-acetyl-\textbf{1,2-}\textit{O}-[\textbf{1-(6-methoxy-6-oxohexyloxy)ethylidene}]-\alpha-D-glucopyranose~(4)$



¹³C and DEPT NMR (CDCl₃, 300 MHz)



 $\textbf{3,4,6-Tri-}\textit{O}-acetyl-\textbf{1,2-}\textit{O}-[\textbf{1-(6-methoxy-6-oxohexyloxy)ethylidene}]-\alpha-D-glucopyranose~(4)$



COSY and HSQC 2D NMR (CDCl₃, 500 MHz)



2,3,4,6-Tetra-*O*-acetyl-1-*O*-(6-methoxy-6-oxohexyl)-β-D-glucopyranose (5)



¹H NMR (CDCl₃, 300 MHz)



2,3,4,6-Tetra-*O*-acetyl-1-*O*-(6-methoxy-6-oxohexyl)-β-D-glucopyranose (5)



¹³C and DEPT NMR (CDCl₃, 300 MHz)



$2,3,4,6\text{-}Tetra\text{-}\textit{O}\text{-}acetyl\text{-}1\text{-}\textit{O}\text{-}(6\text{-}methoxy\text{-}6\text{-}oxohexyl)\text{-}\beta\text{-}D\text{-}glucopyranose}\ (5)$



COSY and HSQC 2D NMR (CDCl₃, 500 MHz)



S17

$2,3,4,6\text{-}Tetra\text{-}\textit{O}\text{-}propargyl\text{-}1\text{-}\textit{O}\text{-}(6\text{-}methoxy\text{-}6\text{-}oxohexyl)\text{-}\beta\text{-}D\text{-}glucopyranose}\ (1)$



¹H NMR (CDCl₃, 300 MHz)



2,3,4,6-Tetra-O-propargyl-1-O-(6-methoxy-6-oxohexyl)-β-D-glucopyranose (1)



¹³C and DEPT NMR (CDCl₃, 300 MHz)



2,3,4,6-Tetra-*O*-propargyl-1-*O*-(6-methoxy-6-oxohexyl)-β-D-glucopyranose (1)



COSY and HSQC 2D NMR (CDCl₃, 500 MHz)



Analytical RP-HPLC chromatograms of N₃-J8 epitope

Blue trace = % solvent B; 5 min at 0%, 0-20% over 5 min, 20-50% over 30 min.

 $t_R = 28.39 \text{ min}$

UV detection ($\lambda = 214$ nm)







Mass Spectrum of N₃-J8 epitope

Molecular Weight: 3364.8

MS-ESI (m/z): $[M + 2H]^{2+}$ calcd 1683.4, found 1683.9; $[M + 3H]^{3+}$ calcd 1122.6, found 1122.8; $[M + 4H]^{4+}$ calcd 842.2, found 842.6; $[M + 5H]^{5+}$ calcd 674.0, found 674.2.



Analytical RP-HPLC chromatograms of Vaccine Candidate 10

Blue trace = % solvent B; 5 min at 0%, 0-20% over 5 min, 20-60% over 30 min.

 $t_R = 28.59 \text{ min}$

UV detection ($\lambda = 214$ nm)







Mass Spectrum of Vaccine Candidate 10

Molecular Weight: 14950.2

MS-ESI (*m/z*): $[M + 5H]^{5+}$ calcd 2991.0, found 2990.2; $[M + 6H]^{6+}$ calcd 2492.7, found 2494.2; $[M + 7H]^{7+}$ calcd 2136.7, found 2137.9; $[M + 8H]^{8+}$ calcd 1869.8, 1870.5; $[M + 9H]^{9+}$ calcd 1662.1, found 1663.2; $[M + 10H]^{10+}$ calcd 1496.0, found 1497.3; $[M + 11H]^{11+}$ calcd 1360.1, found 1360.6; $[M + 12H]^{12+}$ calcd 1246.9, found 1247.7; $[M + 13H]^{13+}$ calcd 1151.0, found 1151.5.



Analytical RP-HPLC chromatograms of Vaccine Candidate 12

Blue trace = % solvent B; 5 min at 0%, 0-20% over 5 min, 20-60% over 30 min.

 $t_R = 31.85 \text{ min}$

UV detection ($\lambda = 214$ nm)







Mass Spectrum of Vaccine Candidate 12

Molecular Weight: 16137.7

$$\begin{split} \text{MS-ESI} \ (m/z): \ [\text{M} + 9\text{H}]^{9+} \text{ calcd } 1794.1, \text{ found } 1794.9; \ [\text{M} + 10\text{H}]^{10+} \text{ calcd } 1614.8, \text{ found } 1614.9; \ [\text{M} + 11\text{H}]^{11+} \text{ calcd } 1468.1, \text{ found } 1468.5; \ [\text{M} + 12\text{H}]^{12+} \text{ calcd } 1345.8, \text{ found } 1346.3; \\ [\text{M} + 13\text{H}]^{13+} \text{ calcd } 1242.4, \text{ found } 1243.1; \ [\text{M} + 14\text{H}]^{14+} \text{ calcd } 1153.7, \text{ found } 1154.2; \ [\text{M} + 15\text{H}]^{15+} \text{ calcd } 1076.8, \text{ found } 1077.3; \ [\text{M} + 16\text{H}]^{16+} \text{ calcd } 1009.6, \text{ found } 1010.2; \ [\text{M} + 17\text{H}]^{17+} \text{ calcd } 950.3, \text{ found } 950.5. \end{split}$$



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