Investigating the Elusive Mechanism of Glycosaminoglycan Biosynthesis

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Supplemental Section:

Structural data:

For xylosides 1 through 25, the structural data can be found in the following reference B.

Kuberan et. al., (2008) *Chembiochem*, **9**, 198-200. The data for remaining xylosides are provided below.

Xyloside 26: 1H NMR (CD₃OD): δ 8.436 (1H, s, triazolyl H), 8.111 (1H, d, J = 7.815, Ar-H),

7.346 (1H, dd, J = 7.03, 8.55 Hz, Ar-H), 7.109 (1H, d, J = 8.2 Hz, Ar-H), 7.055 (1H, dd, J = 7.42,

8.59 Hz, Ar-H), 5.556 (1H, d, J = 9.38 Hz, H-1), 4.057-4.004 (2H, m, H-2, H-5a), 3.962 (3H, s),

3.745-3.682 (1H, m, H-4), 3.540-3.468 (2H, m, H-3, H-5b); Mass (EI): calculated for

C₁₄H₁₇N₃O₅+H 308.12465, found 307.9333

Xyloside 27: 1H NMR (CD₃OD): δ 8.408 (1H, s, triazolyl H), 7.754 (2H, d, J = 8.99 Hz, Ar-H), 6.993 (2H, d, J = 8.98 Hz, Ar-H), 5.533 (1H, d, J = 9.37 Hz, H-1), 4.033 (1H, dd, J = 5.47, 11.33 Hz, H-5a), 3.937 (1H, d, J = 9.37 Hz, H-2), 3.824 (3H, s), 3.735-3.673 (1H, m, H-4), 3.574-3.464 (2H, m, H-3, H-5b); Mass (EI): calculated for C₁₄H₁₇N₃O₅ +H 308.12465, found 308.0667 **Xyloside 28**: 1H NMR (CD₃OD): δ 8.544 (1H, s, triazolyl H), 7.022 (2H, d, J = 2.34 Hz, Ar-H), 6.477 (1H, t, J = 2.34 Hz, Ar-H), 5.540 (1H, d, J = 9.37 Hz, H-1), 4.034 (1H, dd, J = 5.47, 11.33 Hz, H-5a), 3.937 (1H, d, J = 9.37 Hz, H-2), 3.824 (6H, s), 3.709-3.669 (1H, m, H-4), 3.541-3.465 (2H, m, H-3, H-5b); Mass (EI): calculated for C₁₅H₁₉N₃O₆ +H 338.13521, found 338.0000 **Xyloside 29**: 1H NMR (CD₃OD): δ 8.871 (1H, s, triazolyl H), 8.470 (2H, s, Ar-H), 7.949 (1H, s, Ar-H), 5.591 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, d, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, dd, J = 9.38 Hz, H-1), 4.051 (1H, dd, J = 5.47, 11.32 Hz, H-5a), 3.937 (1H, dd, J = 9.38 Hz, H-1), 4.

= 9.38 Hz, H-2), 3.740-3.677 (1H, m, H-4), 3.554-3.482 (2H, m, H-3, H-5b); Mass (EI): calculated for $C_{15}H_{13}F_6N_3O_4$ +H 414.08885, found 413.9333

Xyloside 30: 1H NMR (CD₃OD): δ 8.50 (1H, s, triazolyl H), 8.25 (1H, dd, J = 3.51, 6.25 Hz, Ar-

H), 7.93-7.95 (2H, m, Ar-H), 7.70 (1H, dd, *J* = 1.1, 7.0 Hz, Ar-H), 7.52-7.57 (3H, m, Ar-H),

5.64 (1H, d, J = 9.3 Hz, H-1), 4.07 (1H, dd, J = 5.47, 11.13 Hz, H-5a), 4.03 (1H, t, J

= 9.37, H-2), 3.71-3.77 (1H, m, H-4), 3.51-3.58 (2H, m, H-3, H-5b); Mass (EI): calculated

for C₁₇H₁₇N₃O₄ +H 328.12, found 327.93

Figure Legends

Figure S1. Calibration of size exclusion column with polystyrene sulfonate standards. Polystyrene sulfonate standards of various molecular weights, 65000 Da, were analyzed by size exclusion chromatography as described in the "Material and Methods" section. The migration times of the various polystyrene sulfonate species were plotted against the molecular weight to obtain a calibration curve. The migration time of GAG chains primed by various xylosides were compared to the calibration curve to determine the molecular weight.

Figure S2. Chain length analysis of xyloside-primed GAG chains. The molecular weight of the GAG chains synthesized on various primers was determined by measuring their migration time on size exclusion column as described in the "Material and Methods" section. V_0 and V_t represent the void volume and total volume, respectively. The average migration time was determined by using peak width at half maximum. The average molecular weight was determined using the migration time in comparison to the calibration curve obtained for polystyrene sulfonate standards performed under similar conditions.

Figure S3. Disaccharide profiles of xyloside-primed HS chains. GAG chains (~500,000 cpm) were digested with heparitinases and resulting disaccharides were analyzed by SAX-HPLC with inline flow scintillation analyzer as described in the Experimental section. The SAX elution chromatograms of representative of two independent experiments. I: Δ UA-GlcNAc; II: Δ UA-GlcNS6S; IV: Δ UA2S-GlcNS; V: Δ UA2S-GlcNAc6S and IV: Δ UA2S-GlcNS6S.

Figure S4. Disaccharide profiles of xyloside-primed DS chains. GAG chains (~500,000 cpm) were digested with chondroitinase ABC enzyme and resulting disaccharides were analyzed by SAX-HPLC with inline flow scintillation analyzer as described in the Experimental section. The SAX elution chromatograms of representative of two independent experiments. I: Δ UA-GalNAc; and II: Δ UA-GalNAc6S.

Figure S5. Disaccharide profiles of xyloside-primed CS chains. GAG chains (~500,000 cpm) were digested with chondroitinase ABC enzyme and resulting disaccharides were analyzed by SAX-HPLC with inline flow scintillation analyzer as described in the Experimental section. The SAX elution chromatograms of representative of two independent experiments. I: Δ UA-GalNAc; and II: Δ UA-GalNAc6S.

Figure S6. Long term priming ability of GAG chains by xyloside. The long term priming ability of click-xyloside was examined using xylosyl transferase deficient CHO cells (pgsA-745). 100,000 cells were seeded per well of 24-well plates and treated with xyloside **5** at 100 μ M concentration in the presence of 50 μ Ci ³⁵S-SO₄²⁻ or D-[6-³H]-glucosamine. The medium was removed from the well at 24, 48, 96 and 120 h, GAG chains were purified and quantified as described under "Experimental Methods".



FIGURE S1



FIGURE S2A



FIGURE S2B



FIGURE S2C



FIGURE S2D



FIGURE S2E



FIGURE S2F



FIGURE S3A



FIGURE S3B



FIGURE S4A



FIGURE S4B



FIGURE S5A



FIGURE S5B



FIGURE S6