SUPPORTING INFORMATION

Synthesis and Assessment of Glycosaminoglycan Priming Activity of Cluster-xylosides for Potential Use as Proteoglycan Mimetics

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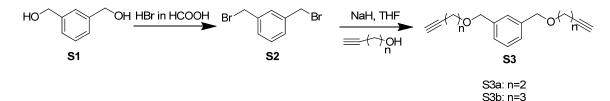
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Experimental Sections:

Anhydrous solvents were purchased and used directly or dried over standard drying agents and freshly distilled prior to use. Reactions were monitored by TLC on silica gel 60 F-254 with detection by Von's reagent. Intermediate compounds were purified by flash chromatography columns using silica gel 60 (230–400 mesh) and were run under medium pressure at 5–7 psi. Final products were purified by high performance liquid chromatography (HPLC) on reverse phase C18 column (VYDAC 2.2 cm×25 cm) with solvent A (25 mM formic acid in water) and solvent B (95% Acetonitrile) at a flow rate of 5 ml/min in a linear gradient over 120 min starting with 0% B. All synthetic compounds were characterized by NMR using a Varian Mercury 400 MHz spectrometer. The compounds were also confirmed for their final structures by electrospray ionization mass spectrometry (ESI-MS) using a Finnigan LCQ mass spectrometer or a Bruker Q-Tof mass spectrometer in either positive or negative ion mode .



Scheme 1: Preparation of precursors carrying alkyne group for click-chemistry

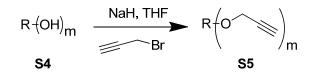
Compound **S2**: 1, 3-Benzenedimethanol (3.618 mmol, 500 mg) was treated with 31 % HBr in acetic acid (5 ml). The reaction was stirred at room temperature for 2 hours. After addition of dichloromethane (10 ml), the organic solvent was washed with water, saturated bicarbonate solution, dried over Na₂SO₄ and evaporated. The crude material was purified by flash chromatography on silica gel to give the compound (**S2**) (900mg, 90% yield). ¹H NMR (CDCl₃): δ 7.42 (1H, s), 7.33 (3H, d, *J* = 1.2 Hz), 4.48 (4H, s) Compound **S3**: To an anhydrous THF solution (10 ml) of alcohol carrying an alkyne group (2 mmol) was added sodium hydride (3 mmol). The reaction mixture was stirred for 30 minutes at room temperature under argon atmosphere followed by the addition of (**S2**) (1 mmol). The mixture was stirred overnight at room temperature. THF solvent was evaporated. The crude material was dissolved in ethyl acetate, washed with water and then saturated sodium chloride solution, dried over Na₂SO₄ and evaporated. The residue was purified by column chromatography to give the compound (**S3**)

n	S3	Yield (%)	Reference
2	S3a	84%	N/A
3	S3b	97%	N/A

Compound **S3a**: ¹H NMR (CDCl₃) δ 7.34-7.25 (4H, m), 4.51 (4H, s), 3.58 (4H, t, *J* = 6.6 Hz), 2.51 (4H,dt, *J* = 6.8, 2.7 Hz), 2.00 (2H, t, *J* = 2.7 Hz)

Compound **S3b**: ¹H NMR (CDCl₃) δ 7.34-7.25 (4H, m), 4.51 (4H, s), 3.58 (4H, t, *J* = 6.3 Hz), 2.32 (4H,dt, *J* = 7.0, 2.7 Hz), 1.94 (2H, t, *J* = 2.7 Hz), 1.84 (4H, p, *J* = 6.3 Hz)

Scheme 2: Preparation of propargylated precursors for the synthesis of bis- and tris- xylosides



Compound **S5**: To an anhydrous THF solution (10 ml) of compound (**S4**) (1 mmol) was added sodium hydride ([m+1] mmol). The reaction mixture was stirred for 30 minutes at room temperature under argon atmosphere followed by addition of propargyl bromide ([m+1] mmol). The mixture was stirred overnight at room temperature. Solvent was evaporated. The crude material was dissolved in ethyl acetate and washed with water and saturated sodium chloride solution, dried over Na_2SO_4 and evaporated. The residue was purified by column chromatography to give the compound (**S5**)

S4	m	S5	Yield (%)	Reference*
НООН	2	S5a	48%	N/A
НОСОН	2	S5b	65%	1
HO~O~OH	2	85c	54%	2
HONO ₂ HONO ₂	3	S5d	27%	N/A

* the procedure was slightly modified from the published reports

Compound **S5a**: ¹H NMR (CDCl₃) δ 7.39-7.30 (4H, m), 4.61 (4H, s), 4.18 (4H, d, *J* = 2.3 Hz), 2.47 (2H, t, *J* = 2.3 Hz)

Compound **S5b**: ¹H NMR (CDCl₃) δ 7.35 (4H, s), 4.61 (4H, s), 4.17 (4H, d, *J* = 2.3 Hz),

2.45 (2H, t, *J* = 2.3 Hz)

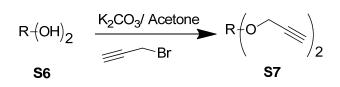
Compound **S5c**: ¹H NMR (CDCl₃) δ 4.12 (4H, d, *J* = 2.4 Hz), 3.61-3.58 (12H, m), 2.38

(2H, s)

Compound **S5d**: ¹H NMR (CDCl₃) δ 4.12 (6H, d, *J* = 2.4 Hz), 3.51 (6H, t, *J* = 5.9 Hz),

2.42 (3H, t, *J* = 2.4 Hz), 2.03-1.99 (6H, m), 1.55-1.48 (6H, m)

Scheme 3: Preparation of additional propargylated precursors



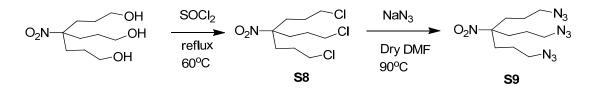
Compound **S7**: To the solution (10 ml) of compound (**S6**) (1 mmol) in acetone, was added potassium carbonate (3 mmol). The reaction mixture was stirred for 30 minutes at room temperature. Propargyl bromide (3 mmol) was then added and the mixture was stirred overnight. The reaction mixture was concentrated. The resulting crude material was dissolved in ethyl acetate, washed with water and saturated sodium chloride solution, dried over Na_2SO_4 and rotary evaporated under reduced pressure. The residue was purified by column chromatography to give the compound (**S7**).

S6	S7	Yield (%)	Reference
HO OH	S7a	20%	3
НОСОН	S7b	56%	4

* The procedures were modified from the published reports

Compound **S7a**: ¹H NMR (CDCl₃) δ 7.89 (2H, d, *J* = 7.8), 7.38 (2H, t, *J* = 7.8), 6.97 (2H, d, *J* = 7.0), 4.88 (4H, s), 2.53 (2H, s)

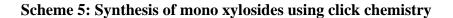
Compound **S7b**: ¹H NMR (CDCl₃) δ 7.68 (2H, d, *J* = 8.6 Hz), 7.19 (4H, d, *J* = 12.1 Hz), 4.78 (4H, s), 2.54 (2H, s)

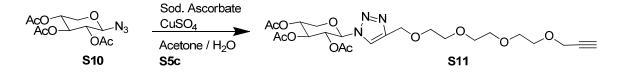


Scheme 4: Preparation of the precursor for the synthesis of tris-xyloside 9

Compound **S8**: Nitromethanetrispropanol (2.127 mmol) was dissolved in SOCl₂ (42.54 mmol). The reaction mixture was refluxed at 60 °C. After the completion of reaction, the reaction mixture was dissolved in ethyl acetate and washed with water, brine and then dried over Na₂SO₄. The final mass is 380 mg (62 %) and the crude product was used in the next step without any further purification. ¹H NMR (CDCl₃): δ 3.54 (6H, t, *J* = 6.4 Hz), 2.10-2.06 (6H, m), 1.74-1.67 (6H, m)

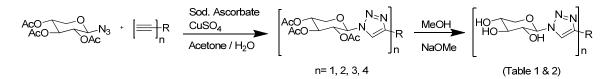
Compound **S9**: The compound (**S8**) (1.275 mmol) was dissolved in dry DMF (10 ml) followed by addition of sodium azide (6.38 mmol). The reaction was heated to 90 °C under stirring. After completion of the reaction as indicated by TLC, the reaction mixture was dissolved in ethyl acetate, washed with water and brine, and then dried over Na₂SO₄. The final mass is (270mg, 66%). ¹H NMR (CDCl₃): δ 3.28 (2H, t, *J* = 6.6 Hz), 1.95-1.91 (2H, m), 1.47-1.39 (2H, m)





Compound **S11**: To a solution of alkyne (**S5c**) (4.98 mmol) and xylosyl azide (**S10**) (1.66 mmol) in 8 ml of acetone and water (1:1) solvent mixture, sodium ascorbate (0.66 mmol) was added, followed by Cu₂SO₄. 5H₂0 (1.33 mmol) at room temperature, and the mixture was stirred until disappearance of compound (**S10**) (as indicated by TLC). At the end of the reaction, the reaction mixture was evaporated using rotary evaporator under reduced pressure to obtain the final product (360 mg, 41%).^{5, 6} ¹H NMR (CD₃OD): δ 8.24 (1H, s), 6.02 (1H, d, *J* = 8.6 Hz), 5.58-5.47 (2H, m), 5.23-5.17 (1H, m), 4.64 (2H, s), 4.25 (1H, dd, *J* = 5.9, 11.3 Hz), 4.18 (2H, d, *J* = 2.3 Hz), 3.77 (1H, t, *J* = 10.9 Hz), 3.66-3.62 (12H, m), 2.83 (1H, t, *J* = 2.3 Hz), 2.04 (3H, s), 2.02 (3H, s), 1.83 (3H, s)

Scheme 6: Synthesis of cluster-xylosides using click chemistry^{5, 6}



Various cluster-xylosides were synthesized using click chemistry and purified as described in the earlier section.

Xyloside **12**: ¹H NMR (CD₃OD): δ 8.60 (2H, s, triazolyl H), 8.34 (1H, s, Ar-H), 7.86 (2H, d, *J* = 7.8 Hz, Ar-H), 7.54 (1H, t, *J* = 7.8 Hz, Ar-H), 5.57 (2H, d, *J* = 9.0 Hz, H-1), 4.05 (2H, dd, *J* = 5.5, 11.1 Hz, H-5a), 3.96 (2H, t, *J* = 9.0 Hz, H-2), 3.68-3.75 (2H, m, H-4), 3.53 (2H, t, *J* = 9.0 Hz, H-3), 3.51 (2H, t, *J* = 10.9 Hz, H-5b); ¹³C NMR (CD₃OD): δ 148.5, 132.4, 130.8, 126.6, 121.6, 90.5, 78.6, 74.0, 70.7, 69.9; ESI-MS: Calcd for C₂₀H₂₄N₆O₈Na: 499.1553 [M+Na]⁺, found 498.9333.

Xyloside **13**: ¹H NMR (CD₃OD): δ 8.58 (1H, s, triazolyl H), 7.92 (1H, s, Ar-H), 5.57 (1H, d, *J* = 9.0 Hz, H-1), 4.05 (1H, dd, *J* = 5.5, 11.3 Hz, H-5a), 3.97(1H, t, *J* = 9.0 Hz, H-2), 3.69-3.75 (1H, m, H-4), 3.54 (1H, t, *J* = 9.0 Hz, H-3), 3.51 (1H, t, *J* = 10.9 Hz, H-5b); ¹³C NMR (CD₃OD): δ 148.4, 131.9, 127.3, 121.5, 90.4, 78.6, 74.0, 70.7, 69.9; ESI-MS: Calcd for C₂₀H₂₅N₆O₈: 477.1734 [M+H]⁺, found 476.8667.

Xyloside **14**: ¹H NMR (CD₃OD): δ 8.16 (2H, s, triazolyl H), 7.37 (1H, s, Ar-H), 7.33-7.28 (3H, m, Ar-H), 5.52 (2H, d, *J* = 9.4 Hz, H-1), 4.65 (4H, s), 4.58 (4H, s), 4.01(2H, dd, *J* = 5.5, 11.3 Hz, H-5a), 3.90 (2H, t, *J* = 9.0 Hz, H-2), 3.709-3.647 (2H, m, H-4), 3.50 (2H, t, *J* = 9.4 Hz, H-3), 3.47 (2H, t, *J* = 10.9 Hz, H-5b; ¹³C NMR (CD₃OD): 144.8, 138.3, 128.4, 127.4, 127.3, 123.1, 89.0, 77.4, 72.7, 72.0, 69.5, 68.7, 62.8; ESI-MS: Calcd for C₂₄H₃₃N₆O₁₀: 565.2258 [M+H]⁺, found 565.1189. Xyloside **15**: ¹H NMR (CD₃OD): δ 8.14 (2H, s, triazolyl H), 7.34 (4H, s), 5.51 (2, d, J = 9.4 Hz, H-1), 4.65 (4H, s), 4.58 (4H, s), 4.01 (2H, dd, J = 5.1, 11.1 Hz, H-5a), 3.88 (2H, t, J = 9.4 Hz, H-2), 3.71-3.64 (2H, m, H-4), 3.49 (2H, t, J = 9.4 Hz, H-3), 3.46 (2H, t, J = 10.9 Hz, H-5b); ESI-MS: Calcd for C₂₄H₃₂N₆O₁₀Na: 587.2078 [M+Na]⁺, found 587.1150.

Xyloside **16**: ¹H NMR (CD₃OD): δ 7.91 (2H, s, triazolyl H), 7.22 (1H,s, Ar-H), 7.21 (2H, d, *J* = 7.0 Hz, Ar-H), 7.29 (1H, t, *J* = 7.0 Hz, Ar-H), 5.46 (2H, d, *J* = 9.0 Hz, H-1), 4.51 (4H, s), 3.99 (2H, dd, *J* = 5.5, 11.3 Hz, H-5a), 3.86 (2H, t, *J* = 9.0 Hz, H-2), 3.74 (4H, t, *J* = 6.6 Hz), 3.67-3.63 (2H, m), 3.49 (2H, t, *J* = 9.0 Hz, H-3), 3.45 (4H, t, *J* = 10.9 Hz, H-5b), 3.00 (4H, t, *J* = 6.6 Hz,); ESI-MS: Calcd for C₂₆H₃₆N₆O₁₀Na: 615.2391 [M+Na]⁺, found 615.1550.

Xyloside **17**: ¹H NMR (CD₃OD): δ 7.83 (2H, s, triazolyl H), 7.34-7.25 (4H, m), 5.44 (2H, d, *J* = 9.0 Hz, H-1), 4.51 (4H, s), 3.99 (2H, dd, *J* = 5.5, 11.3 Hz, H-5a), 3.83 (2H, t, *J* = 9.0 Hz, H-2), 3.70-3.64 (2H, m), 3.54-3.42 (8H, m), 2.81 (4H, t, *J* = 7.8 Hz), 1.99-1.92 (4H, m); ESI-MS: Calcd for C₂₈H₄₀N₆O₁₀Na: 643.2704 [M+Na]⁺, found 643.2550. Xyloside **18**: ¹H NMR (CD₃OD): δ 8.33 (2H, s, triazolyl H), 7.81 (2H, d, *J* = 8.6 Hz, Ar-H), 7.35 (2H, t, *J* = 7.8 Hz, Ar-H), 7.10 (2H, d, *J* = 7.8 Hz, Ar-H), 5.55 (2H, d, *J* = 9.0 Hz, H-1), 5.35 (4H, s), 4.02 (2H, dd, *J* = 5.5, 11.3 Hz, H-5a), 3.93 (2H, t, *J* = 9.0 Hz, H-2), 3.74-3.65 (2H, m, H-4), 3.50 (2H, t, *J* = 9.0 Hz, H-3), 3.48 (2H, t, *J* = 11.3 Hz, H-5b); ¹³C NMR (CD₃OD): δ 155.2, 145.1, 128.1, 126.3, 124.6, 115.9, 107.2, 90.3, 78.6, 73.9, 70.7, 69.9, 62.7; ESI-MS: Calcd for C₂₆H₂₉N₆O₁₀: 585.1945 [M-H]⁻, found 585.0000. Xyloside **19**: ¹H NMR (CD₃OD): δ 8.32 (2H, s, triazolyl H), 7.73 (2H, d, *J* = 9.0 Hz, Ar-H), 7.39 (2H, s, Ar-H), 7.18 (2H, d, *J* = 9.0 Hz, Ar-H), 5.55 (2H, d, *J* = 9.0 Hz, Ar-H), 7.39 (2H, s, Ar-H), 7.18 (2H, d, *J* = 9.0 Hz, Ar-H), 5.55 (2H, d, *J* = 9.0 Hz, Ar-H), 7.39 (2H, s, Ar-H), 7.18 (2H, d, *J* = 9.0 Hz, Ar-H), 5.55 (2H, d, *J* = 9.0 Hz, H-1), 5.29

(4H, s), 4.01 (2H, dd, J = 5.5, 11.3 Hz, H-5a), 3.91 (2H, t, J = 9.4 Hz, H-2), 3.70-3.64 (2H, m, H-4), 3.49 (2H, t, J = 9.4 Hz, H-3), 3.48 (2H, t, J = 10.94 Hz, H-5b); ¹³C NMR (CD₃OD): δ 156.4, 145.1, 131.4, 129.6, 124.7, 120.2, 108.6, 90.2, 78.7, 73.9, 70.7, 69.9, 62.4; ESI-MS: Calcd for C₂₆H₂₉N₆O₁₀: 585.1945 [M-H]⁻, found 585.0667. Xyloside **20**: ¹H NMR (CD₃OD): δ 8.22 (2H, s, triazolyl H), 5.73 (2H, s), 5.55 (2H, d, J =9.4 Hz, H-1), 4.60 (4H, s), 4.12 (4H, s), 4.01 (2H, dd, J = 5.5, 11.3 Hz, H-5a), 3.91 (2H, t, J = 9.4 Hz, H-2), 3.67-3.53 (2H, m, H-4), 3.51-3.44 (4H, m, H-3, H-5b); ESI-MS: Calcd C₂₀H₃₀N₆O₁₀Na: 537.202 [M+Na]⁺, found 537.151 Table S1. Sulfation density analysis of xyloside-primed GAG chains. GAG chains primed by cluster-xylosides in pgsA-745 cell lines at 100 μ M were analyzed using anion exchange HPLC; the bound GAG chains were eluted with a linear gradient of 0.2 M to 1 M NaCl over 80 minutes at a flow rate of 1 ml/min and elution times indicate charge density of primed GAG chains.

Xylosides	Elution times of primed GAG chains (min)
Mono-xyloside 1	44.5
Mono-xyloside 2	40.8
Bis-xyloside 3b	49.8
Bis-xyloside 4	50.2
Bis-xyloside 5	50.8
Bis-xyloside 6	51.5
Bis-xyloside 7b	50.6
Tris-xyloside 8a	46.0
Tris-xyloside 9	47.2
Tetrakis-xyloside 10a	48.9
Tetrakis-xyloside 11a	49.7

Figure Legends

Figure S1. Comparison of GAG priming activity of protected bis-xylosides and deprotected bis-xylosides. The priming ability of protected bis-xylosides (**3a**, **7a**) and deprotected bis-xylosides (**3b**, **7b**) were examined using xylosyl transferase deficient CHO cells (pgsA-745). 400,000 cells were seeded in wells of 6-well plates and treated with xyloside (**3a**, **3b**, **7a** or **7b**) at 100 μ M and 250 μ M concentrations in the presence of 100 μ Ci of *D*-[6-³H]-glucosamine. The medium was removed from the well at 24 hours, GAG chains were purified and quantified as described in the method section.

Figure S2. Comparison of the priming ability of GAG chains by bis-xylosides at various concentrations. The priming ability of the bis-xylosides (3b and 7b)were examined using xylosyl transferase deficient CHO cells (pgsA-745). 400,000 cells were seeded in wells of 6-well plates and treated with bis-xylosides at 100 μ M, 300 μ M, 600 μ M and 1 mM concentrations in the presence of 100 μ Ci of *D*-[6-³H]-glucosamine. The medium was removed from the well at 24 hours, GAG chains were purified and quantified as described in the method section.

Figure S3. Comparison of GAG priming activity of protected cluster-xylosides and deprotected cluster-xylosides. The priming ability of protected and deprotected tris and tetrakis-xylosides was examined using xylosyl transferase deficient CHO cells (pgsA-745). 400,000 cells were seeded in wells of 6-well plates and treated with cluster xylosides at 100 μ M in the presence of 100 μ Ci of *D*-[6-³H]-glucosamine. The medium was removed from the well at 24 hours, GAG chains were purified and quantified as described in the method section.

Figure S4. Priming activity of bis-xylosides in the pgs A-745 cell line at 1 mM. CHO cells were treated with bis- xylosides at 1 mM in the presence of ³H-GlcNH₂ (100 μ Ci) as described in the method section. The GAG chains were purified by anion exchange chromatography and quantitated using a liquid scintillation counter. The results were average of two independent experiments that were carried out in duplicate.

Figure S5. Calibration of size exclusion column with polystyrene sulfonate

standards. Polystyrene sulfonate standards of various molecular weights "I (3600 Da), II (6530 Da), III (14900 Da), IV (32900 Da), V (63900 Da), VI (152000)" were analyzed by size exclusion chromatography. 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 cluster-xylosides were compared to the calibration curve to determine the molecular weight of primed GAG chains.

Figure S6. Effect of various hydrophobic moieties on the DEAE elution profiles of

GAG chains. GAG chains primed by various cluster-xylosides eluted from an analytical anion-exchange column (7.5 mm x 7.5 cm) using linear gradient of 1 M NaCl solution. GAG samplers were diluted five-fold prior to analysis by DEAE-HPLC. The variations in the elution profiles and migration times indicate differences in both sulfation pattern and extent of sulfation of the primed GAG chains. The elution profiles are representative of at least two independent experiments.

Figure S7. Structural analysis of GAG chains primed by bis-xyloside (3b) at 100 μ M. PgsA-745 cells were incubated with the bis-xyloside (3b) for 24 hours. The primed GAG chains were then purified as described in the method section. The HS/CS composition of the primed GAG chains was determined by digesting the GAG chains

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with heparitinase I/II/III or chondroitinase ABC enzymes. The reaction mixture was incubated at 37 °C for 2 hours, the solution was then loaded on to two tandem G3000 SWXL columns (7.8 mm x 30 cm) and analyzed with the aid of an inline radiometric detector using phosphate buffer as an eluent. The HS/CS composition was determined based on the percentage area of undigested and digested GAG peak. **A**: The elution profile of GAG chains without heparitinase I/ II/ III or chondroitinase ABC treatment. **B**: The elution profile of GAG chains after treatment with heparitinase I/ II/ III enzyme. **C**: The elution profile of GAG chains after treatment with chondroitinase ABC enzyme. **D**: The elution profile of GAG chains after treatment with heparitinase I/ II/ III and chondroitinase ABC enzyme.

Figure S8. GAG priming activity was determined by additional bis-xylosides. A: Priming activity of bis-xylosides (12 and 13). B: priming analysis of bis-xylosides (14, 15, 16 and 17). C: Priming activity of bis-xylosides (18 and 19). The priming ability of bis-xylosides were examined using xylosyl transferase deficient CHO cells (pgsA-745). 400,000 cells were seeded in wells of 6-well plates and treated with bis-xylosides at 100 μ M in the presence of 100 μ Ci of *D*-[6-³H]-glucosamine. The medium was removed from the well at 24 hours, GAG chains were purified and quantified as described in the method section.

Figure S9. Disaccharide profiles of xyloside-primed CS chains. GAG chains, which were primed by cluster-xylosides (3b, 8a and 10a) at 100 μ M, were treated with chondroitinase ABC and the resulting disaccharides were analyzed by SAX-HPLC. I: Δ UA-GalNAc; and II: Δ UA-GalNAc4S

Figure S10. Periodate oxidation-alkaline elimination of GAG chains. GAG chains, which were primed by mono-xyloside 2 and bis-xyloside 7 at 100 μ M, were subjected to periodate oxidation-alkaline elimination as described in the method section. **A**: The elution profiles of GAG chains, primed by mono-xyloside 2, before (gray trace) and after (black trace) treatments are shown. **B**: The elution profiles of GAG chains primed by bis-xyloside 7, before (gray trace) and after (black trace) treatments are shown. This data showed that only GAG chains primed by bis-xylosides can be broken by oxidation-elimination treatments whereas GAG chains primed by mono-xylosides cannot be cleaved and thus, demonstrated the bidirectional GAG priming ability of bis-xylosides.

Figure S11. **Ozonolysis of GAG chains**. GAG chains, which were primed by bisxyloside **20** at 100 μ M, were subjected to ozonolysis. A: Schematic representation that shows the action of ozone on double bond. Bis-xyloside **20** that carries a double bond is susceptible to ozone treatment resulting in the cleavage of two GAG chains that are assembled on the same scaffold. B: The elution profiles of GAG chains primed by bisxyloside **20** before (black trace) and after ozonolysis (gray trace) are shown and the data shown here demonstrate the bidirectional priming ability of bis-xyloside **20**.

Figure S1

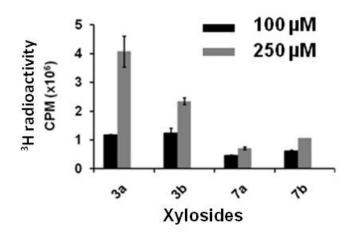


Figure S2

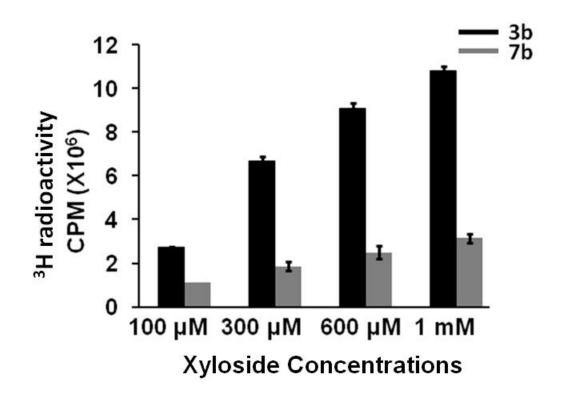


Figure S3

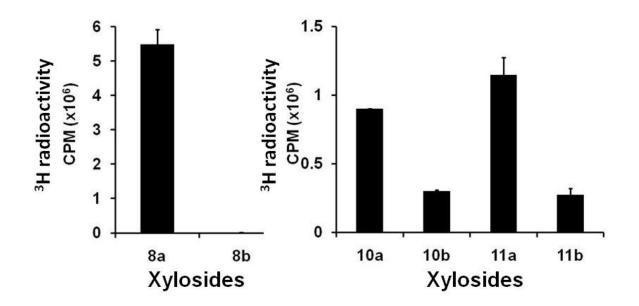


Figure S4

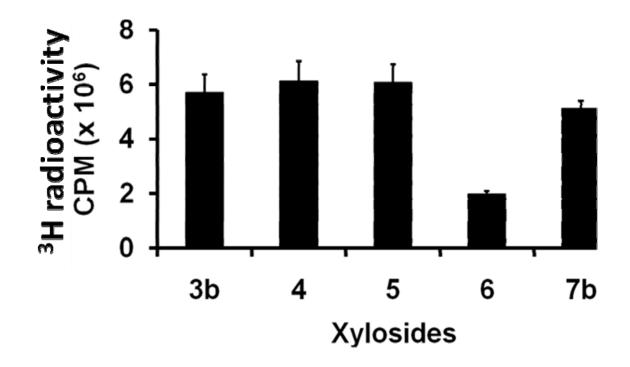


Figure S5

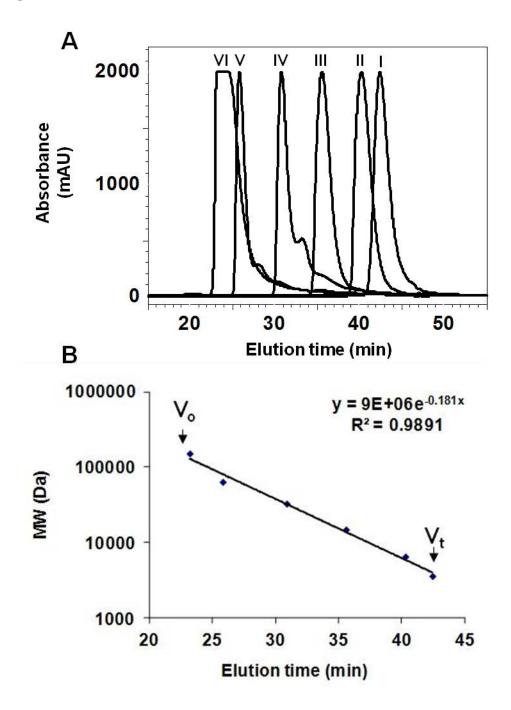
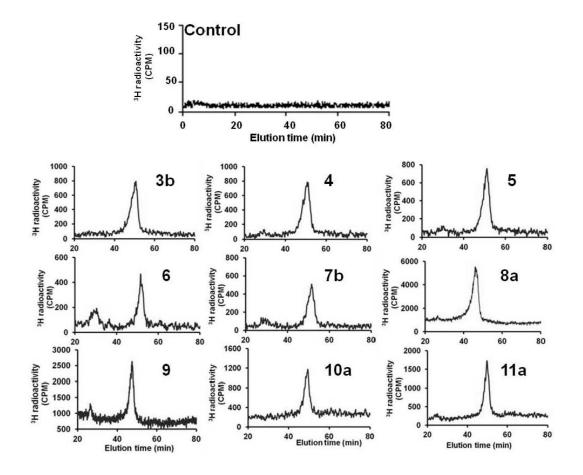


Figure S6





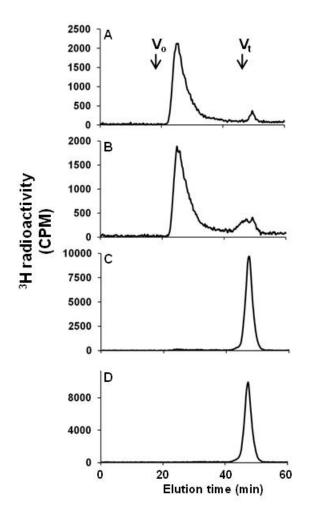


Figure S8

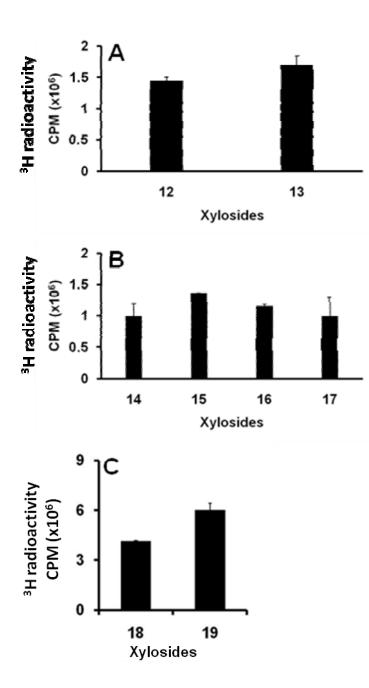
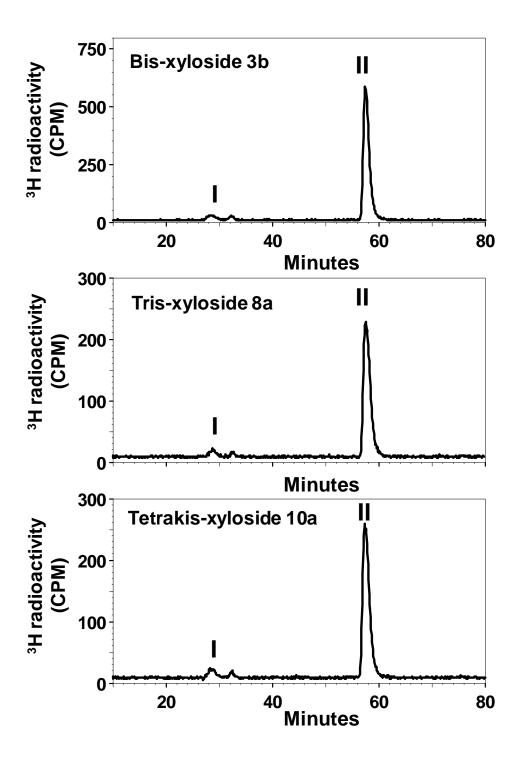
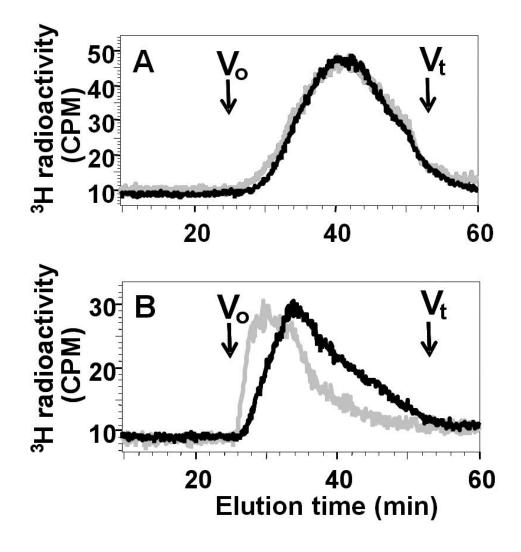
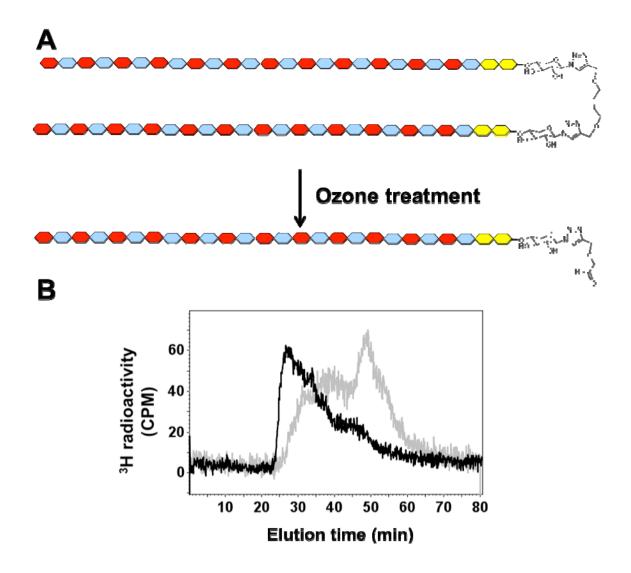


Figure S9







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