

Chemical Synthesis and Anti-Inflammatory Activity of Bikunin Associated Chondroitin Sulfate 24-mer

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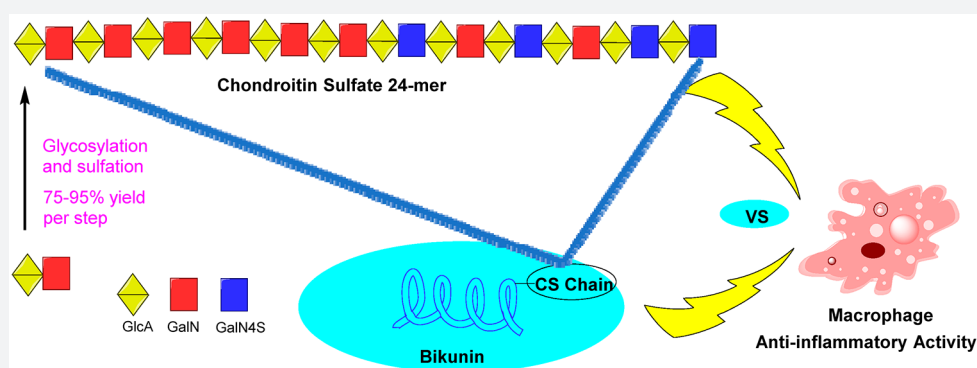
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ABSTRACT: Bikunin, a chondroitin sulfate (CS) proteoglycan clinically used to treat acute inflammation and sepsis, contains a CS chain with more than 20 monosaccharide units. To understand the function of the CS chain of bikunin, synthesis of long CS chains is needed. After exploring multiple glycosylation approaches and protective group chemistry, we report herein the successful generation of the longest CS chain to date (24-mer) in an excellent overall yield on a multi-mg scale. The anti-inflammatory activities of both bikunin and the synthetic 24-mer were determined, and the results demonstrate that both the glycan and the core protein are important for anti-inflammatory activities of bikunin by reducing macrophage production of proinflammatory cytokines.

INTRODUCTION

Bikunin, also known as the urinary trypsin inhibitor, is a glycoprotein in human plasma and urine.¹ Bikunin has many biological functions and is one of the main anti-inflammatory mediators.^{2–4} The levels of bikunin in plasma and urine can be significantly increased (up to 10-fold) for a range of conditions, including cancer and chronic inflammation.^{2,3,5,6} Clinically, bikunin has been used as a drug to treat acute inflammatory disorders such as pancreatitis, septic shock, and disseminated intravascular coagulation.

Bikunin is a proteoglycan consisting of a 147 amino acid residue core protein with a chondroitin sulfate (CS) chain attached to serine-10.^{7,8} Linhardt and co-workers have performed groundbreaking sequencing studies of bikunin CS chains using Fourier transform ion cyclotron resonance mass spectrometry.⁹ The CS chain of bikunin was determined to contain over 20 monosaccharide units with repeating glucuronic acid (GlcA) and *N*-acetyl galactosamine (GalNAc).^{9–11} Only 4-OH groups of GalNAc are partially sulfated in bikunin associated CS and 4–7 *O*-sulfate groups were found positioned toward the reducing end of the CS chain.^{3,9–13} Herein, we report the first synthesis of a CS chain 24-mer **1** from bikunin (Figure 1) and its anti-inflammatory activities.

The 24-mer is the longest CS chain synthesized to date. Biological studies showed that the CS chain is important for the anti-inflammatory effect of bikunin, while the core protein without the CS chain actually enhanced the levels of inflammatory cytokine secreted from macrophages treated with lipopolysaccharide (LPS).

Innovative technologies have been developed toward the synthesis of defined CS oligosaccharides,^{14,15} which include enzymatic synthesis,¹⁶ synthesis with building blocks derived from natural polysaccharides,^{17–20} automated solid phase synthesis,^{21,22} and small libraries of CS with varying sulfation patterns.^{23–28} The longest synthetic defined CS glycans produced to date were octasaccharides²³ and nonasaccharides.^{16,19} To assemble a long CS chain such as 24-mer **1**, new synthetic strategies need to be established.

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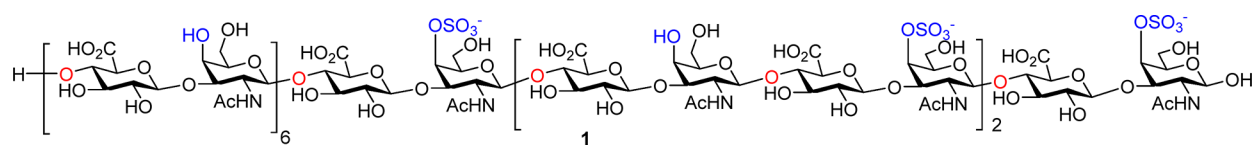
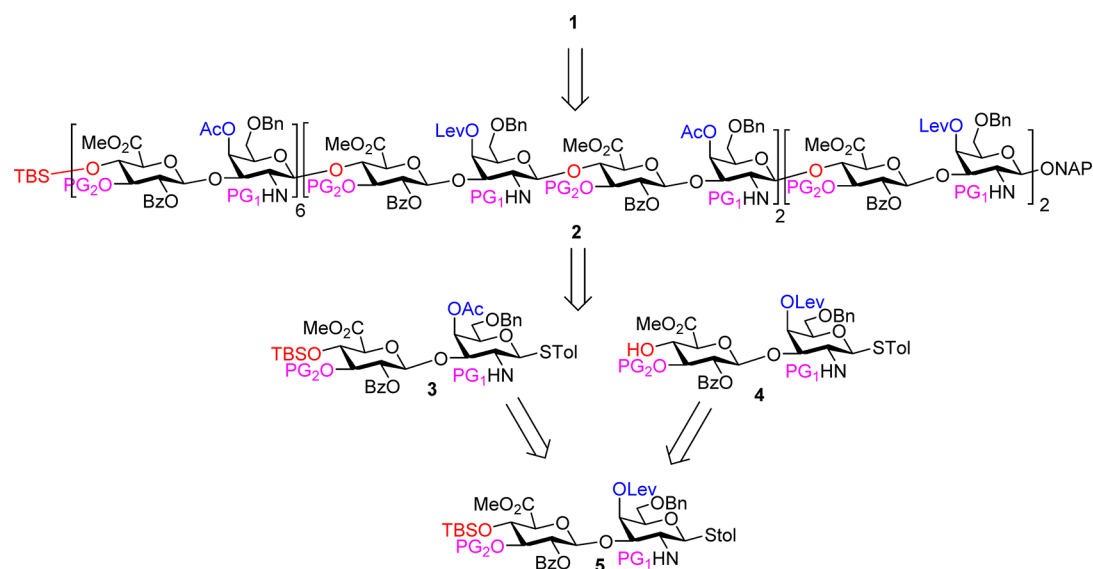


Figure 1. Structure of bikunin CS chain (24-mer) 1.

Scheme 1. Retrosynthetic Design of 24-mer 1



RESULTS AND DISCUSSION

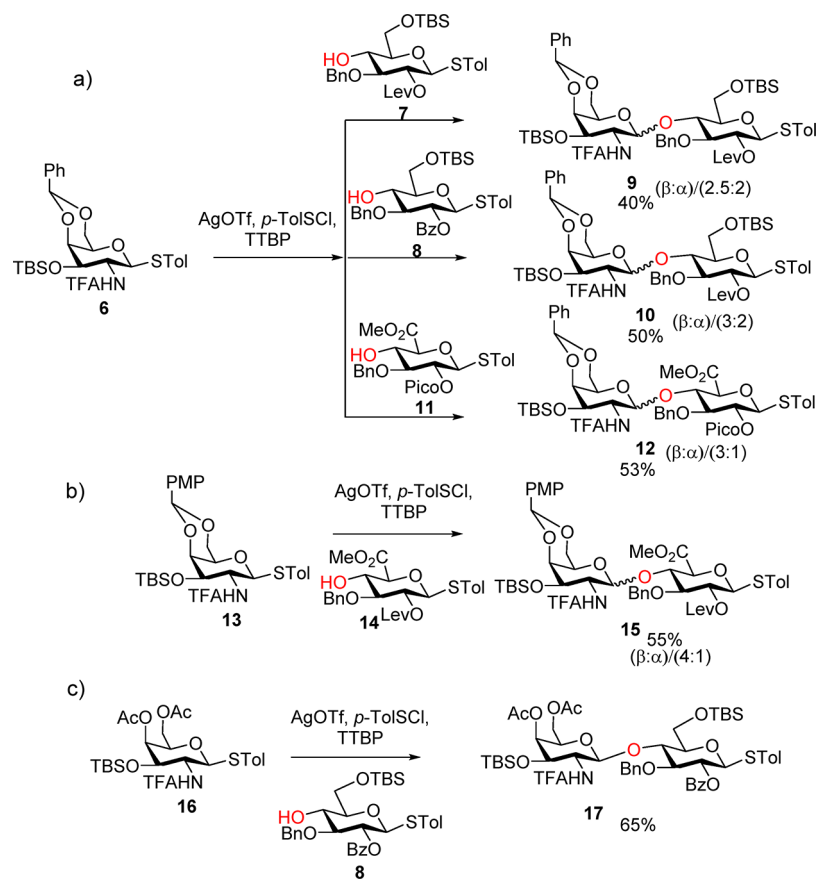
Synthesis of Fully Protected CS 24-mer. We envision that 24-mer 1 can be obtained from the fully protected precursor 24-mer 2, which in turn can be generated using disaccharide building blocks 3 and 4. The 4-OH groups of the reducing end galactosamine unit of both 3 and 4 are differentially protected for future selective sulfation. The disaccharides 3 and 4 can serve as both a glycosyl donor and an acceptor, which can be derived from the common precursor 5 (Scheme 1). A key in the synthesis is selection of the protective group on the amine, which should facilitate the formation of the desired 1,2-*trans* glycosides and be deprotected in high yields as there would be 12 such groups to be removed to generate the 24-mer 1. In addition, it was important to use the *t*-butyldimethylsilyl (TBS) group as the protective group on 4-OH of glucuronic acid (e.g., donor 3), as the presence of TBS helped enhance the solubility of the building blocks for high-yield glycosylation and the ease in purification.

We started our synthesis by testing trifluoroacetamide (TFA)^{27,29} protected galactosamine (GalN) monosaccharide as the donor due to the potential ease of TFA removal under a mild basic condition. However, the glycosylation between GalNTFA donor 6 and glucoside acceptor 7 or 8 gave disaccharides 9 and 10 as anomeric mixtures with little β -selectivity (β : α = 2.5:2, 3:2, respectively) (Scheme 2a). Switching the acceptor to the glucuronic acid derivative 11 or the donor to the more electron donating *p*-methoxybenzylidene protected GalNTFA 13 led to modest improvements (Schemes 2a,b). Interestingly, replacing the 4,6-benzylidene in the donor with di-4,6-*O*-acetates (donor 16) produced the desired β -linked disaccharide 17 as the only anomer isolated (Scheme 2c). These suggest that the 4,6-benzylidene moiety

likely restricted the conformational freedom of the pyranoside ring upon donor activation, rendering it difficult for TFA to assist the β -glycoside formation by neighboring group participation.

With the success of using 16 as the glycosyl donor, we examined the utility of disaccharide (Figure 2) donors such as 17 containing NHTFA protected galactosamine at the reducing end. Unfortunately, glycosylation of 17 with disaccharide acceptor 18 did not yield the desired tetrasaccharide, with the oxazoline 19 formed as the major side product in 55% yield. Changes of protective groups (donor 20), anomeric leaving group (21) on the donor, as well as the acceptor structure (22) did not lead to productive glycosylation either, with the donor mostly converted to the corresponding disaccharide oxazoline side product. The failures of disaccharide donors such as 20 and 17 are possibly because of their lower anomeric reactivities compared to monosaccharide 16 due to the electron withdrawing effect of the additional glycan ring³⁰ leading to enhanced stabilities of the oxazoline intermediates.

To overcome the challenges encountered with TFA protected disaccharide donors, we envision that the reactivities of building blocks can be enhanced by using less electron withdrawing protective groups such as the trichloroacetyl (TCA)^{14,17,18,26,31–33} and benzyl groups. Thus, glucoside donor 23 and TCA protected GalN acceptor 24 were designed and synthesized. Preactivation of donor 23 by *p*-TolSCl/AgOTf³⁴ followed by the addition of acceptor 24 produced the key disaccharide building block GlcA-GalN 25 in an excellent 88% yield (Scheme 3a). To analyze the ability of TCANH bearing disaccharide 25 as a glycosyl donor, glycosylation of 25 with 2-naphthol was carried out first, which, with subsequent TBS cleavage, afforded disaccharide 27 in 85% yield for the two steps (Scheme 3b). Furthermore, coupling between 25

Scheme 2. Challenges Encountered Using TFA Protected GalN Monosaccharide as Donors^a

^a(a) Stereochemical challenges in formation of disaccharides **9**, **10**, and **12** from donor **6**. (b) Modest stereoselectivity observed in the formation of disaccharide **15** from donor **13**. (c) Formation of **17** in high stereoselectivity from donor **16**.

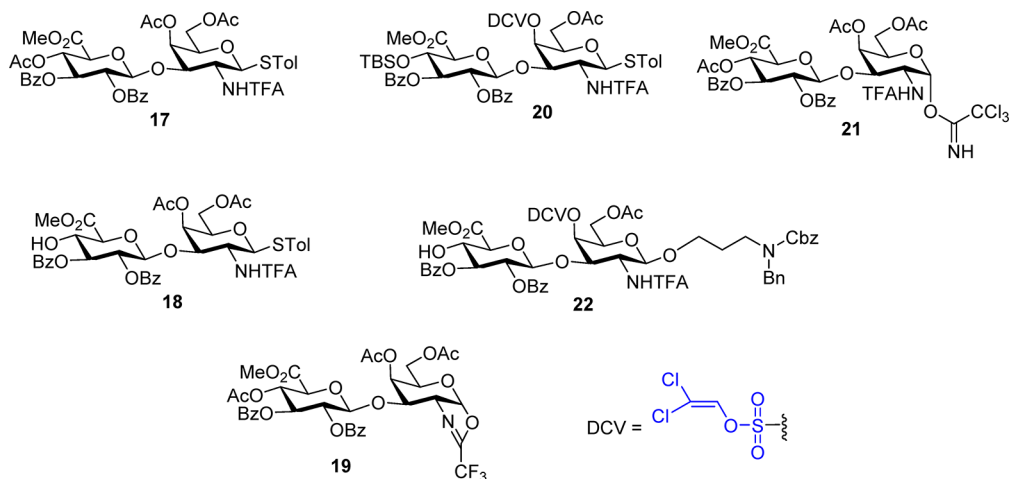


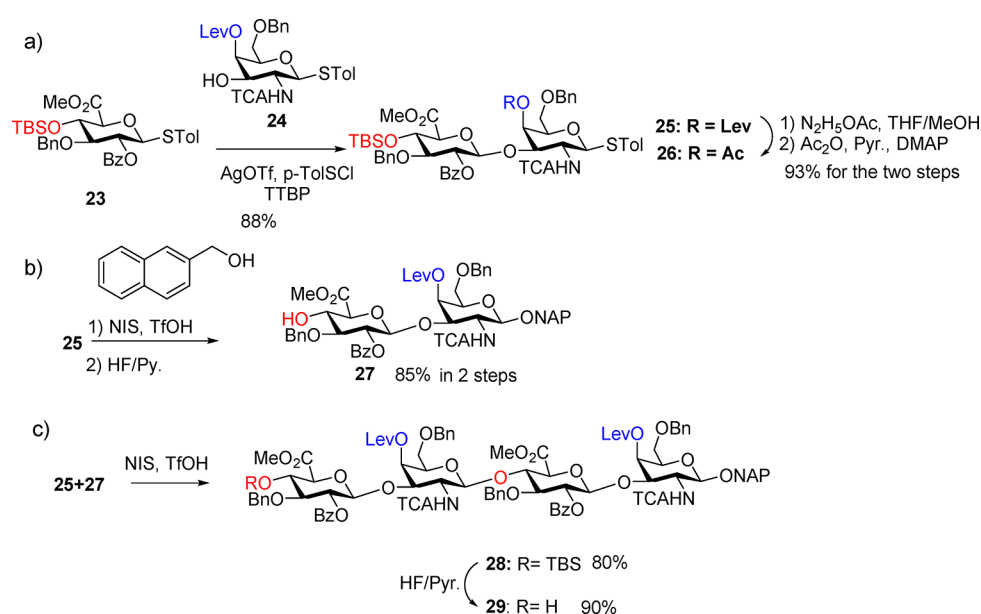
Figure 2. Structures of disaccharide building blocks examined containing NHTFA protected galactosamine.

and acceptor **27** generated the desired tetrasaccharide **28** in 80% yield (Scheme 3c), suggesting that disaccharide **25** is a suitable glycosyl donor for CS synthesis. Deprotection of the TBS group on **27** produced the tetrasaccharide **29** in an excellent 90% yield.

To elongate the CS backbone toward the CS 24-mer, the 4-*O* acetate bearing disaccharide **26** (Scheme 3a) was obtained as a second disaccharide building block. Glycosylation between donor **26** and tetrasaccharide acceptor **29** proceeded smoothly

to give hexasaccharide **30** in 77% yield (Scheme 4). TBS deprotection of **30** followed by glycosylation with levulinyl ester bearing donor **25** gave octasaccharide **32** in an excellent yield (90%) (Scheme 4). The TBS deprotection and glycosylation with donor **26** and **25** were repeated iteratively, producing deca-saccharide **34** in 75% yield, dodeca-saccharide **36** in 77% yield, tetradeca-saccharide **38** in 89% yield, hexadeca-saccharide **40** in 82% yield, octadeca-saccharide **42** in 91% yield, icosasaccharide **44** in 86% yield, docosa-sacchar-

Scheme 3. (a) Synthesis of Disaccharide Donors 25 and 26, (b) Synthesis of Acceptor 27, and (c) Synthesis of Tetrasaccharide Acceptor 29



ide 46 in 95% yield, and tetracosasaccharide 48 in 90% yield (Scheme 4). This glycosylation approach was readily scalable, generating the protected 24-mer 48 on a 200 mg scale. The TBS group in tetracosasaccharide 48 was deprotected leading to tetracosasaccharide 49, which, with its free OH at the nonreducing end, could serve as an acceptor for further chain elongation if necessary.

DEPROTECTION OF THE 24-MER 49

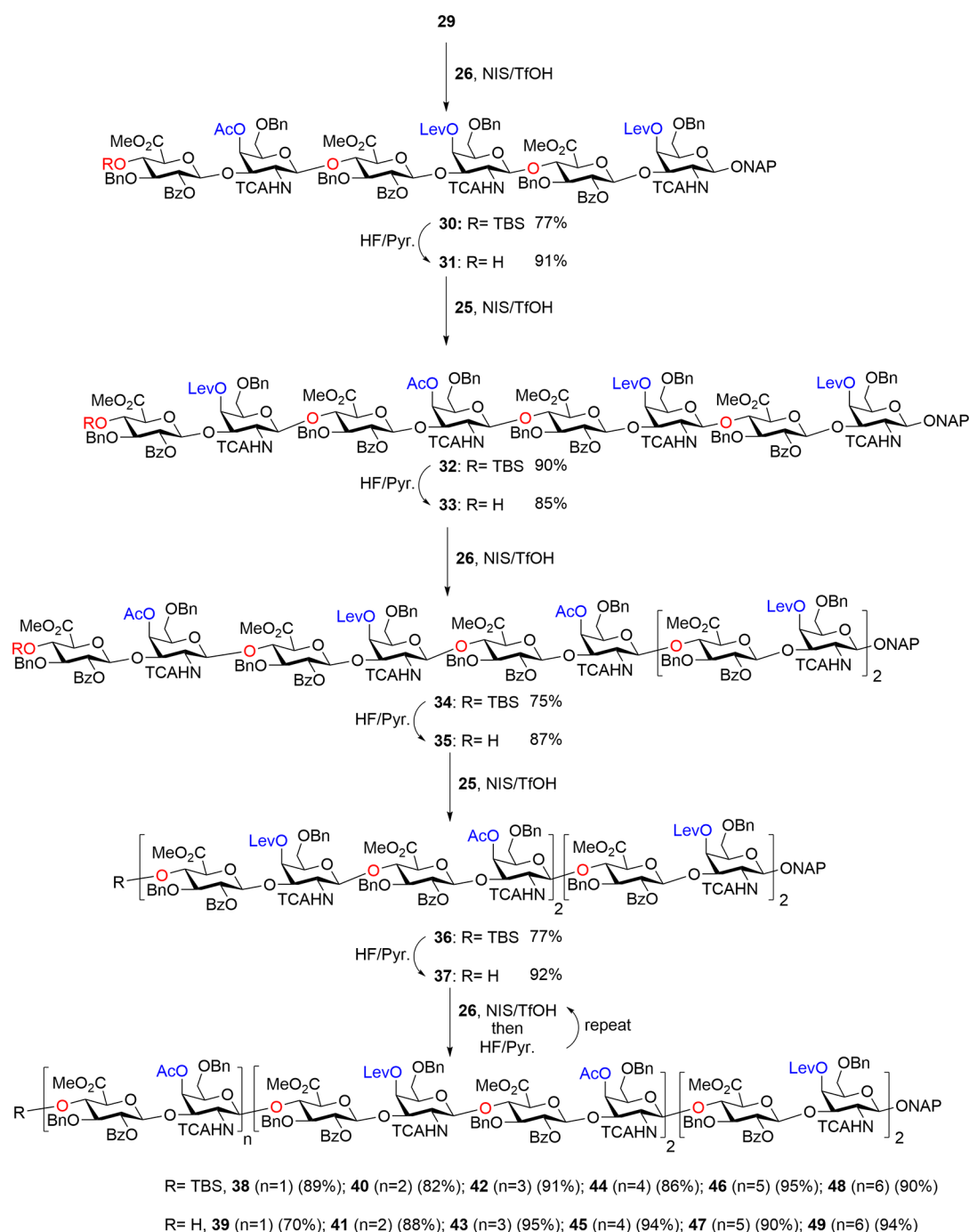
With the fully protected 24-mer 49 in hand, we went forward to deprotection and sulfation (Scheme 5). 49 was acetylated with Ac₂O to give tetracosasaccharide 50. Selective cleavage of the Lev groups was performed by treatment of 50 with hydrazine acetate to afford the tetraol 51 in 91% yield, which was subjected for sulfation using sulfur trioxide–triethylamine complex (SO₃–NEt₃) in dry DMF to provide the sulfated 24-mer 52 in 85% yield.

The next challenging task is the unprecedented conversion of 12 NHTCAs to acetamides. The Zn–Cu couple reduction method^{33,35} was tested first on 50, which led to incomplete conversions and glycan decompositions. Next, a basic condition was applied to remove the TCA. 52 was first treated with LiOH (1 M) and H₂O₂³¹ for deprotection of methyl esters and a multitude of acyl groups. However, this reaction was slow with only partial removal of TCA. Similar phenomena were reported as in the synthesis of hyaluronan decasaccharide; deprotection of the five TCA moieties took 3 weeks.³⁶ Addition of NaOMe to the reaction mixture of 52 with heating at 50 °C did not give the desired fully deprotected product. Increasing the basicity of the reaction led to cleavage of the glycan chain. After extensive screening, we discovered that treatment of the reaction mixture after NaOMe with concentrated ammonia (2 days at 55 °C) could cleanly remove all 12 TCA protective groups as monitored by mass spectrometry (see the SI, page S186). The resulting free amines were selectively *N*-acetylated to afford 53 in 60% yield over 4 steps from 50. Finally, hydrogenolysis of 53 using Pd(OH)₂/C gave the desired 24-mer 1 in 58% yield (Scheme 5).

CS Chain Is Important for the Anti-Inflammatory Activities of Bikunin.

With the homogeneous 24-mer CS 1 in hand, we investigated its anti-inflammatory effects using a macrophage/lipopolysaccharide (LPS) assay.^{37,38} Found on the outer membrane of Gram negative bacteria, LPS can activate macrophages to release a large amount of proinflammatory cytokines such as tumor necrosis factor- α (TNF- α), which may lead to septic shock, a potentially fatal medical condition resulting from infections.³⁹ To better understand the structural requirement of bikunin on its inflammatory responses, bikunin was treated with chondroitinase ABC to cleave off the endogenous CS chain producing the core protein lacking the CS chain.⁴⁰ In addition, bikunin was treated with actinase E to completely digest the core protein backbone generating the bikunin associated CS chain, which mainly consists of CS-A⁹ as confirmed by our NMR and MS analysis. The abilities of macrophages to respond to LPS upon incubation with bikunin, core protein, synthetic CS 24-mer 1, commercially available CS-A polysaccharide, and bikunin CS were determined by quantifying the amounts of TNF- α released from macrophages using an enzyme linked immunosorbent assay (ELISA) (Figure 3). Consistent with its clinical application, bikunin was able to reduce the amounts of TNF- α from macrophages induced by LPS in a dose dependent manner (Figure 3).^{38,41} At 5 and 50 μ M of bikunin, the levels of TNF- α were reduced to 65% and 39% of those from macrophages not receiving bikunin. Interestingly, under the same condition, macrophages treated with the core protein without CS produced more TNF- α , suggesting that the core protein alone actually exacerbated macrophage inflammations. In contrast to the core protein, commercial CS-A polysaccharide and bikunin CS were able to reduce TNF- α levels to 80% at 5 μ M concentration. The synthetic CS 24-mer 1 exhibited improved anti-inflammatory effects compared to both CS-A polysaccharide and bikunin CS. At a low dose (5 μ M) of CS 24-mer 1, the TNF- α level from treated macrophages decreased to 60% of the control, similar to that by bikunin. The further increase of the doses of CS-24-mer 1 did not lead to significant differences in reduction of TNF- α

Scheme 4. Synthesis of Protected 24-mer 49



levels (Figure 3). Our results suggest that the CS chain could modify the function of bikunin core protein, leading to anti-inflammatory activities against macrophages.

A possible reason for the absence of dose dependent responses in the anti-inflammatory activities from CS polysaccharide or the synthetic CS 24-mer is that CS and bikunin may have different aggregation states. To test this, we measured hydrodynamic diameters of CS and bikunin in solution by dynamic light scattering (DLS), and both CS and bikunin showed small size decreases at higher concentrations, possibly due to increased repulsive interactions between polymer chains (Figure S2). Furthermore, little UV-vis absorbance at 600 nm was observed with aqueous solutions

of CS or bikunin at all concentrations. These results suggest that CS and bikunin are soluble and not aggregating at concentrations studied. As solubility and aggregation are not likely the main factor, the differential dose dependent responses may be due to different cellular receptors of CS from that of bikunin for cellular response or transport, which will require further investigation.

CONCLUSION

In conclusion, due to the structural heterogeneity of CS polysaccharide from nature,^{2,9,42} it is important to develop synthetic methodologies to access well-defined CS chains. While multiple innovative syntheses of CS oligosaccharides up

Scheme 5. Deprotection and Sulfation of Protected 24-mer 55 Successfully Generated CS 24-mer 1

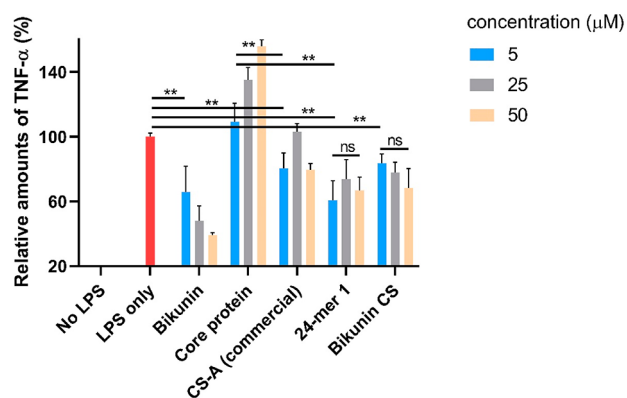
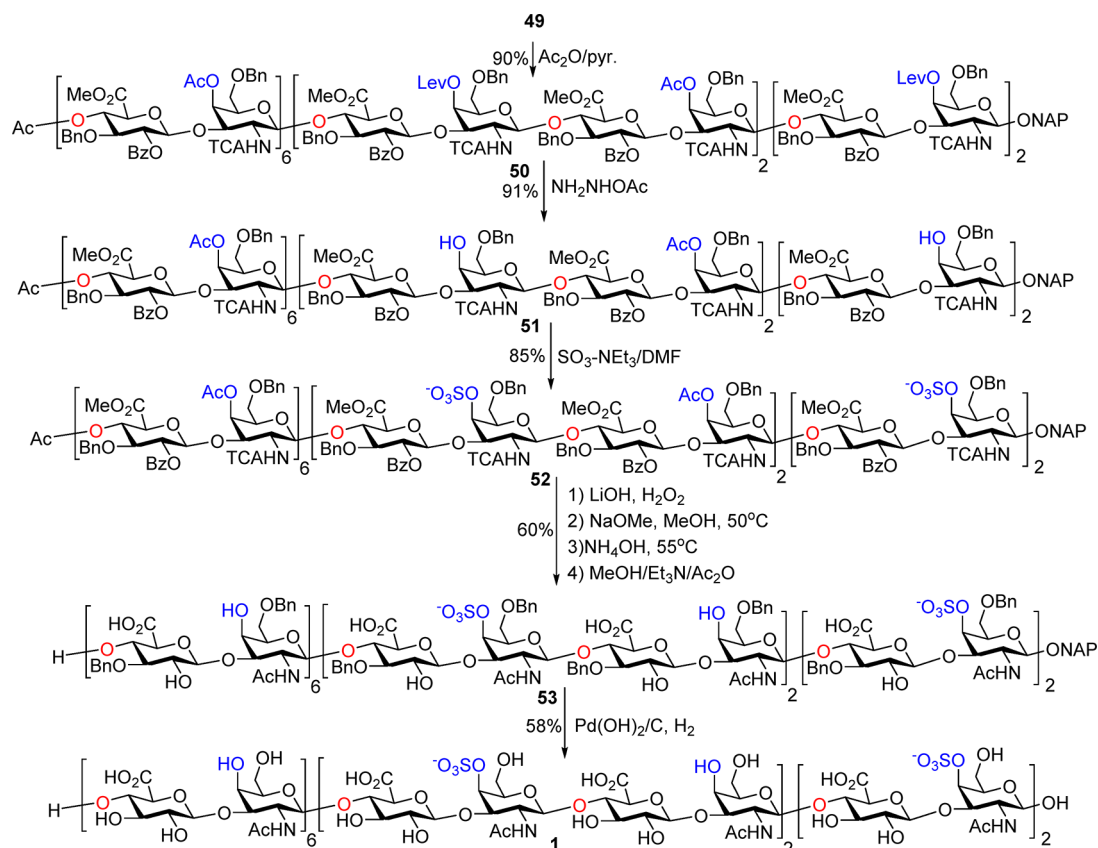


Figure 3. Amounts of TNF- α secreted by macrophage raw 264.7 cells upon treatment with LPS (100 ng/mL) in the presence of bikunin, bikunin core protein, CS-A polysaccharide, bikunin CS, and the synthetic CS 24-mer 1 at 5, 25, and 50 μ M concentrations, respectively. The amounts of TNF- α secreted by macrophages were quantified by ELISA and normalized against the levels of TNF- α from cells treated with LPS only. Each experiment was performed at least three times with the mean values presented. The error bars represent standard deviations of the measurements. Statistical analyses were performed through a two-tailed *t* test using GraphPad Prism software. ** *p* < 0.01. ns: no significant differences.

to nonasaccharides have been reported, the need to study CS proteoglycans such as bikunin requires longer CS chains. Herein, we report that by judicious design of building blocks and protective groups, fully protected CS 24-mer could be prepared via chemical glycosylation in a high overall yield on a 200 mg scale. Deprotection and sulfation conditions have been

established to remove the multitude of protective groups and install the requisite *O*-sulfates producing the CS 24-mer 1, which is the longest CS chain synthesized to date.

As bikunin is an approved drug to treat inflammatory conditions, the abilities of the CS 24-mer 1 to reduce the inflammatory effects of macrophages were studied and compared with bikunin glycoprotein and the corresponding core protein backbone. Interestingly, the core protein of bikunin lacking the endogenous CS chain was found to stimulate the production of TNF- α , a powerful proinflammatory cytokine, from macrophages treated with LPS. In contrast, CS 24-mer 1 exhibited anti-inflammatory activities by reducing the levels of TNF- α from macrophages, and the level of reduction was similar to that obtained using the full bikunin glycoprotein at 5 μ M concentrations of bikunin. The similar activities observed of synthetic CS 24-mer compared to bikunin CS suggest the synthetic 24-mer can recapitulate the function of native CS polysaccharides. The ability of obtaining long CS chains opens up avenues to investigate biological functions of CS with its native length in glycoproteins. Studies are underway to synthesize such glycopeptides and glycoproteins and investigate their multifaceted biological functions.

■ ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acscentsci.9b01199>.

Detailed experimental procedures, characterization data of all the synthesized compounds and intermediates, NMR spectra (^1H , ^{13}C , COSY, HSQC, HMBC), and MS (ESI-MS or MALDI-MS) spectra (PDF)

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Notes

The authors declare no competing financial interest.

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■ REFERENCES

- (1) Proksch, G. J.; Routh, J. I. The purification of the trypsin inhibitor from human pregnancy urine. *Transl. Res.* **1972**, *79*, 491–499.
- (2) Puglia, M. J.; Valdes, R.; Jortani, S. A. Bikunin (urinary trypsin inhibitor): structure, biological relevance, and measurement. *Adv. Clin. Chem.* **2007**, *44*, 223–245.
- (3) Fries, E.; Blom, A. M. Bikunin — not just a plasma proteinase inhibitor. *Int. J. Biochem. Cell Biol.* **2000**, *32*, 125–137.
- (4) Maehara, K.; Kanayama, N.; Halim, A.; Elmaradny, E.; Oda, T.; Fujita, M.; Terao, T. Down-regulation of interleukin-8 gene expression in HL60 cell line by human kunitz-type trypsin inhibitor. *Biochem. Biophys. Res. Commun.* **1995**, *206*, 927–934.
- (5) Lin, S. D.; Endo, R.; Kuroda, H.; Kondo, K.; Miura, Y.; Takikawa, Y.; Kato, A.; Suzuki, K. Plasma and urine levels of urinary trypsin inhibitor in patients with chronic liver diseases and hepatocellular carcinoma. *J. Gastroenterol. Hepatol.* **2004**, *19*, 327–332.

- (6) Matsuzaki, H.; Kobayashi, H.; Yagy, T.; Wakahara, K.; Kondo, T.; Kurita, N.; Sekino, H.; Inagaki, K.; Suzuki, M.; Kanayama, N.; Terao, T. Plasma bikunin as a favorable prognostic factor in ovarian cancer. *J. Clin. Oncol.* **2005**, *23*, 1463–1472.

- (7) Zhuo, L.; Hascall, V. C.; Kimata, K. Inter-alpha-trypsin inhibitor, a covalent protein-glycosaminoglycan-protein complex. *J. Biol. Chem.* **2004**, *279*, 38079–38082.

- (8) Xu, Y.; Carr, P. D.; Guss, J. M.; Ollis, D. L. The crystal structure of bikunin from the inter- α -inhibitor complex: a serine protease inhibitor with two Kunitz domains. *J. Mol. Biol.* **1998**, *276*, 955–966.

- (9) Ly, M.; Leach, F. E., III; Laremore, T. N.; Toida, T.; Amster, I. J.; Linhardt, R. J. The proteoglycan bikunin has a defined sequence. *Nat. Chem. Biol.* **2011**, *7*, 827–833.

- (10) Enghild, J. J.; Thøgersen, I. B.; Cheng, F.; Fransson, L.-Å.; Roepstorff, P.; Rahbek-Nielsen, H. Organization of the inter- α -inhibitor heavy chains on the chondroitin sulfate originating from ser10 of bikunin: posttranslational modification of I α I-derived bikunin. *Biochemistry* **1999**, *38*, 11804–11813.

- (11) Capon, C.; Mizon, C.; Lemoine, J.; Rodié-Talbert, P.; Mizon, J. In acute inflammation, the chondroitin-4 sulphate carried by bikunin is not only longer; it is also undersulphated. *Biochimie* **2003**, *85*, 101–107.

- (12) Toyoda, H.; Kobayashi, S.; Sakamoto, S.; Toida, T.; Imanari, T. Structural analysis of a low-sulfated chondroitin sulfate chain in human urinary trypsin inhibitor. *Biol. Pharm. Bull.* **1993**, *16*, 945–947.

- (13) Yamada, S.; Oyama, M.; Kinugasa, H.; Nakagawa, T.; Kawasaki, T.; Nagasawa, S.; Khoo, K.-H.; Morris, H. R.; Dell, A.; Sugahara, K. The sulphated carbohydrate-protein linkage region isolated from chondroitin 4-sulphate chains of inter- α -trypsin inhibitor in human plasma. *Glycobiology* **1995**, *5*, 335–341.

- (14) Vibert, A.; Jacquinet, J. C.; Lopin-Bon, C. Recent advances in the chemical and enzymatic chondroitin sulfate synthesis. *J. Carbohydr. Chem.* **2011**, *30*, 393–414.

- (15) Ramadan, S.; Yang, W.; Huang, X. Chapter 8. Synthesis of chondroitin sulfate oligosaccharides and chondroitin sulfate glycopeptides. *Synthetic Glycomes: The Royal Society of Chemistry* **2019**, 172–206 and references cited therein.

- (16) Li, J.; Su, G.; Liu, J. Enzymatic synthesis of homogeneous chondroitin sulfate oligosaccharides. *Angew. Chem., Int. Ed.* **2017**, *56*, 11784–11787.

- (17) Vibert, A.; Lopin-Bon, C.; Jacquinet, J.-C. From polymer to size-defined oligomers: a step economy process for the efficient and stereocontrolled construction of chondroitin oligosaccharides and biotinylated conjugates thereof: part 1. *Chem. - Eur. J.* **2009**, *15*, 9561–9578.

- (18) Lopin, C.; Jacquinet, J.-C. From polymer to size-defined oligomers: an expeditious route for the preparation of chondroitin oligosaccharides. *Angew. Chem., Int. Ed.* **2006**, *45*, 2574–2578.

- (19) Zhang, X.; Liu, H.; Lin, L.; Yao, W.; Zhao, J.; Wu, M.; Li, Z. Synthesis of fucosylated chondroitin sulfate nonasaccharide as a novel anticoagulant targeting intrinsic factor xase complex. *Angew. Chem., Int. Ed.* **2018**, *57*, 12880–12885.

- (20) Chng, Y. S.; Tristan, G.; Yip, G. W.; Lam, Y. Protecting-group-free synthesis of chondroitin 6-sulfate disaccharide and tetrasaccharide. *Org. Lett.* **2019**, *21*, 4559–4562.

- (21) Eller, S.; Collot, M.; Yin, J.; Hahm, H. S.; Seeberger, P. H. Automated solid-phase synthesis of chondroitin sulfate glycosaminoglycans. *Angew. Chem., Int. Ed.* **2013**, *52*, S858–S861.

- (22) Liang, C. F.; Hahm, H. S.; Seeberger, P. H. Automated synthesis of chondroitin sulfate oligosaccharides. *Methods Mol. Biol.* **2015**, *1229*, 3–10.

- (23) Tamura, J.; Nakada, Y.; Taniguchi, K.; Yamane, M. Synthesis of chondroitin sulfate E octasaccharide in a repeating region involving an acetamide auxiliary. *Carbohydr. Res.* **2008**, *343*, 39–47.

- (24) Matsushita, K.; Nakata, T.; Takeda-Okuda, N.; Nakanaka, S.; Kitagawa, H.; Tamura, J. Synthesis of chondroitin sulfate CC and DD tetrasaccharides and interactions with 2H6 and LY111. *Bioorg. Med. Chem.* **2018**, *26*, 1016–1025.

(25) Miyachi, K.; Wakao, M.; Suda, Y. Syntheses of chondroitin sulfate tetrasaccharide structures containing 4,6-disulfate patterns and analysis of their interaction with glycosaminoglycan-binding protein. *Bioorg. Med. Chem. Lett.* **2015**, *25*, 1552–1555.

(26) Gama, C. I.; Tully, S. E.; Sotogaku, N.; Clark, P. M.; Rawat, M.; Vaidehi, N.; Goddard, W. A., 3rd; Nishi, A.; Hsieh-Wilson, L. C. Sulfation patterns of glycosaminoglycans encode molecular recognition and activity. *Nat. Chem. Biol.* **2006**, *2*, 467–473.

(27) Macchione, G.; Maza, S.; Kayser, M. M.; de Paz, J. L.; Nieto, P. M. Synthesis of chondroitin sulfate oligosaccharides using N-(tetrachlorophthaloyl)- and N-(trifluoroacetyl)galactosamine building blocks. *Eur. J. Org. Chem.* **2014**, *2014*, 3868–3884.

(28) Yang, S.; Liu, Q.; Zhang, G.; Zhang, X.; Zhao, Z.; Lei, P. An approach to synthesize chondroitin sulfate-E (CS-E) oligosaccharide precursors. *J. Org. Chem.* **2018**, *83*, 5897–5908.

(29) Maza, S.; Mar Kayser, M.; Macchione, G.; López-Prados, J.; Angulo, J.; de Paz, J. L.; Nieto, P. M. Synthesis of chondroitin/dermatan sulfate-like oligosaccharides and evaluation of their protein affinity by fluorescence polarization. *Org. Biomol. Chem.* **2013**, *11*, 3510–3525.

(30) Koeller, K. M.; Wong, C.-H. Synthesis of complex carbohydrates and glycoconjugates: enzyme-based and programmable one-pot strategies. *Chem. Rev.* **2000**, *100*, 4465–4493.

(31) Tully, S. E.; Mabon, R.; Gama, C. I.; Tsai, S. M.; Liu, X.; Hsieh-Wilson, L. C. A chondroitin sulfate small molecule that stimulates neuronal growth. *J. Am. Chem. Soc.* **2004**, *126*, 7736–7737.

(32) Coutant, C.; Jacquinet, J. C. 2-Deoxy-2-trichloroacetamido-D-glucopyranose derivatives in oligosaccharide synthesis - from hyaluronic-acid to chondroitin 4-sulfate trisaccharides. *J. Chem. Soc., Perkin Trans. 1* **1995**, 1573–1581.

(33) Ramadan, S.; Yang, W.; Zhang, Z.; Huang, X. Synthesis of chondroitin sulfate A bearing syndecan-1 glycopeptide. *Org. Lett.* **2017**, *19*, 4838–4841.

(34) Huang, X.; Huang, L.; Wang, H.; Ye, X.-S. Iterative one-pot synthesis of oligosaccharides. *Angew. Chem., Int. Ed.* **2004**, *43*, 5221–5224.

(35) Vibert, A.; Lopin-Bon, C.; Jacquinet, J.-C. Efficient alternative for the reduction of N-trichloroacetyl groups in synthetic chondroitin oligosaccharide intermediates. *Tetrahedron Lett.* **2010**, *51*, 1867–1869.

(36) Lu, X.; Kamat, M.; Huang, L.; Huang, X. Chemical synthesis of a hyaluronic acid decasaccharide. *J. Org. Chem.* **2009**, *74*, 7608–7617.

(37) Wakahara, K.; Kobayashi, H.; Yagyu, T.; Matsuzaki, H.; Kondo, T.; Kurita, N.; Sekino, H.; Inagaki, K.; Suzuki, M.; Kanayama, N.; Terao, T. Bikunin suppresses lipopolysaccharide-induced lethality through down-regulation of tumor necrosis factor- α and interleukin-1 β in macrophages. *J. Infect. Dis.* **2005**, *191*, 930–938.

(38) Matsuzaki, H.; Kobayashi, H.; Yagyu, T.; Wakahara, K.; Kondo, T.; Kurita, N.; Sekino, H.; Inagaki, K.; Suzuki, M.; Kanayama, N.; Terao, T. Bikunin inhibits lipopolysaccharide-induced tumor necrosis factor α induction in macrophages. *Clin. Diagn. Lab. Immunol.* **2004**, *11*, 1140–1147.

(39) Annane, D.; Bellissant, E.; Cavaillon, J.-M. Septic shock. *Lancet* **2005**, *365*, 63–78.

(40) Lord, M. S.; Day, A. J.; Youssef, P.; Zhuo, L.; Watanabe, H.; Caterson, B.; Whitelock, J. M. Sulfation of the bikunin chondroitin sulfate chain determines heavy chain-hyaluronan complex formation. *J. Biol. Chem.* **2013**, *288*, 22930–22941.

(41) Kanayama, S.; Yamada, Y.; Onogi, A.; Shigetomi, H.; Ueda, S.; Tsuji, Y.; Haruta, S.; Kawaguchi, R.; Yoshida, S.; Sakata, M.; Sado, T.; Kitanaka, T.; Oi, H.; Yagyu, T.; Kobayashi, H. Bikunin suppresses expression of pro-inflammatory cytokines induced by lipopolysaccharide in neutrophils. *J. Endotoxin Res.* **2007**, *13*, 369–376.

(42) Nadanaka, S.; Clement, A.; Masayama, K.; Faissner, A.; Sugahara, K. Characteristic hexasaccharide sequences in octasaccharides derived from shark cartilage chondroitin sulfate D with a neurite outgrowth promoting activity. *J. Biol. Chem.* **1998**, *273*, 3296–3307.