

Parallel Synthesis of Glycomimetic Libraries: Targeting a C-Type Lectin

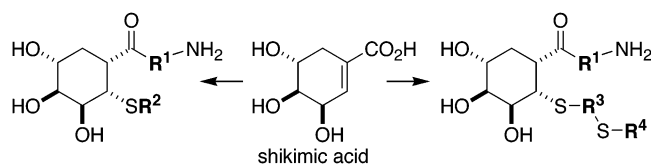
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ABSTRACT



We have developed methods for the parallel synthesis of two libraries of non-carbohydrate-based analogues of mannose on a solid support. The natural product shikimic acid was used as a key building block. The ability of the compounds to block the binding of the C-type lectin MBP-A to a mannosylated surface was assessed in a high-throughput assay. Ten library members with inhibitory activities equivalent to that of α -methyl mannopyranoside were identified.

Carbohydrate–protein interactions are critical in physiological and pathological processes.¹ Consequently, synthetic glycomimetics and oligosaccharides designed to inhibit these interactions have been pursued as potential therapeutic agents.² Effective inhibitors have been identified for some carbohydrate–protein binding events, but most of these are derived from carbohydrate scaffolds.³ The potential lability of synthetic carbohydrates toward *in vivo* glycosidases has prompted the search for stable carbohydrate analogues (glycomimetics) with potent biological activities.

We sought to develop non-carbohydrate-based glycomimetics targeted toward the C-type lectin family. The C-type lectins are a group of Ca²⁺-dependent carbohydrate-binding proteins, many of which are found in mammals.⁴ Human C-type lectins include the mannose-binding proteins (MBPs),

the selectins (E-, L- and P-selectin), and DC-SIGN.⁵ These proteins are all involved in immune system regulation.⁶ Studies of ligand-bound MBP-A, MBP-C, E-selectin, and P-selectin by X-ray crystallography have revealed the importance of a vicinal axial–equatorial–equatorial display of hydroxyl groups on the carbohydrate ligand (Figure 1A).⁷ We sought a core structure that was capable of providing a collection of diverse ligands and that could give rise to

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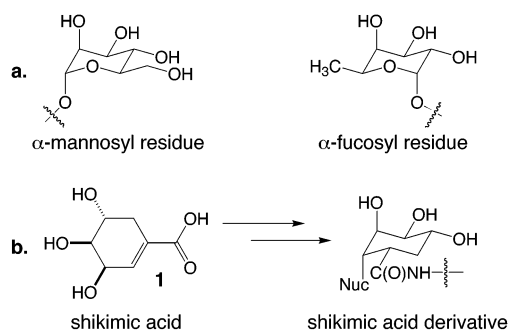


Figure 1. (a) Mannose and fucose residues participate in binding to C-type lectins via their axial–equatorial–equatorial vicinal hydroxyl group. (b) Transformation of shikimic acid affords products with the desired hydroxyl group arrangement.

ligands that target a range of lectins within the C-type lectin superfamily. A key criterion for our scaffold was that it should allow for the streamlined synthesis of libraries.⁸

We chose to build our library from the carbocycle shikimic acid. Shikimic acid possesses three hydroxyl groups that are displayed in a *pseudoaxial–pseudoequatorial–pseudoequatorial* configuration (Figure 1B). We envisioned that glycomimetics could be generated by the stereoselective conjugate addition of nucleophiles. Diversification could be achieved by using different thiol building blocks and by elaboration of the carboxylic acid. We chose to use thiolates as building blocks because they are excellent nucleophiles, and the corresponding conjugate addition reactions should proceed under mild conditions. With proper stereochemical control, conjugate addition could provide the requisite *axial–equatorial–equatorial* display of vicinal hydroxyl groups (Figure 1B).⁹ Multivalent derivatives of shikimic acid itself have been used previously to inhibit a protein–carbohydrate interaction.¹⁰ Because of the mode of C-type lectin ligand binding, we surmised that the proposed transformations of shikimic acid would yield more potent ligands.

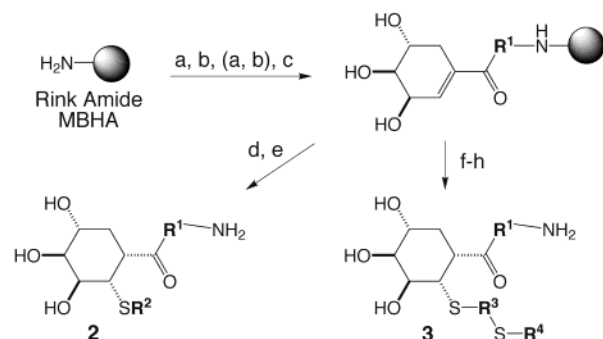
To facilitate preparation of glycomimetic libraries, two related solid-phase, parallel syntheses were envisioned (Scheme 1). The key step in each strategy, the conjugate addition of a thiolate to a shikimic acid derivative, was tested in solution. At issue was the stereochemical outcome of the conjugate addition reaction. Specifically, we anticipated that

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Scheme 1. Solid-Supported Synthesis of Libraries 1 and 2^a



^a Conditions: (a) Fmoc-(R¹)-OH, HBTU, DIEA, DMF; (b) piperidine, DMF; (c) shikimic acid, DIC, C₆F₅OH, HOBT, DMF; (d) R²SH, KO^t-Bu, *t*-BuOH, DMF; (e) TFA, Et₃SiH, H₂O, HSCH₂CH₂SH; (f) HS-R³-SH, KO^t-Bu, *t*-BuOH, DMF; (g) R⁴-Br, KI, Et₃N, DMF; (h) TFA, H₂O.

the incoming nucleophile would approach the unsaturated carbonyl compound on the *si* face opposite the allylic hydroxyl group. The stereochemical outcome from protonation of the incipient enolate, however, was more difficult to predict. We found that when the addition was conducted using a shikimic acid derivative with free hydroxyl groups, the desired isomer was obtained. Under the optimized conditions, thiolate addition occurs from the *si* face, followed by *axial* protonation to afford the desired isomer (Figure 2). With this knowledge, we set about to develop solid-phase routes to the target compounds.

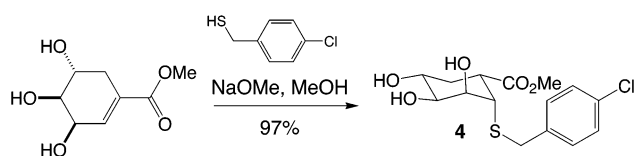


Figure 2. Desired stereochemical outcome was obtained from thiolate conjugate addition to shikimic acid methyl ester. Protonation of the incipient enolate on the other face affords a product that would exist in a conformer in which the hydroxyl groups are improperly oriented for C-type lectin binding.

In both of the synthetic routes developed, Rink amide 4-methylbenzhydrylamine polystyrene resin was utilized as the synthetic support. It is stable toward a variety of reaction conditions and delivers compounds with terminal amide groups.¹¹ To the immobilized amine was coupled either one or two commercially available amino acids via standard methods. Shikimic acid, which could be obtained via fermentation¹² or isolated from star anise, was then coupled to the free N-terminal amine.¹³ Conveniently, shikimic acid could be added without protection of its secondary hydroxyl

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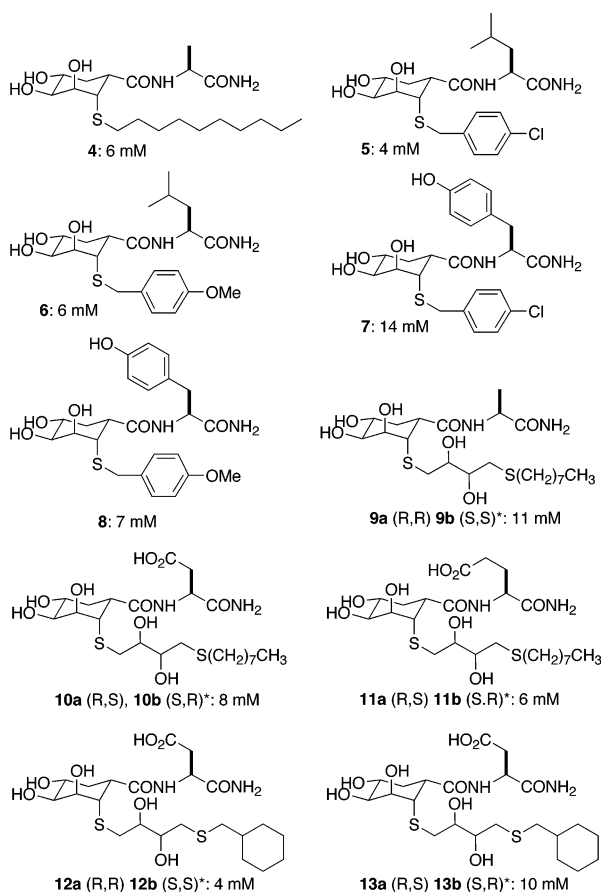


Figure 5. Compounds with activity in the MBP-A bead elution assay from Libraries 1 and 2. The reference IC_{50} for α -MeMan in this assay is 10 mM. An asterisk (*) indicates where compounds synthesized with DTT or DTE afforded diastereomers that were not separated.

to that of α -MeMan for MBP-A (Figure 5). Shikimic acid alone was inactive in the assay even at a concentration of 100 mM. These results indicate that members of the designed library have the key structural features required for interaction with the C-type lectins.

Analysis of the data reveals several trends. The amino acid substituent of the active ligands varies, suggesting that this

position has only a minor influence on binding. The charged amino acid side chains in the active ligands, however, are all anionic; thus, they may engage in a Coulombic interaction or hydrogen bond. With regard to the thiol substituent, all of the active ligands possess a hydrophobic group at their terminus. In both libraries, none of the compounds bearing an anionic functional group at this position were found to have activity. The identification of MBP-A ligands that possess hydrophobic groups at this position suggests the presence of a hydrophobic pocket near the carbohydrate binding site. Significantly, ligands that bind as well as the carbohydrates that bind MBP-A have been identified from these libraries of modest size. These results indicate that shikimic acid is useful in glycomimetic synthesis.

The glycomimetics described here are effective inhibitors of a C-type lectin. Shikimic acid is a valuable building block because its carboxylic acid group and conjugated alkene allow for the incorporation of diversifying elements. These elements may be used to distinguish between members of the C-type lectin family. Thus, we anticipate that the synthetic strategy outlined here may serve as a general method for producing selective inhibitors of C-type lectins.

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Supporting Information Available: Experimental procedures for the synthesis of library members, description and synthesis of thiol- and alkyl bromide library building blocks, and NMR spectra for select compounds. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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