

Synthesis and Affinity Evaluation of a Small Library of Bidentate Cholera Toxin Ligands: Towards Nonhydrolyzable Ganglioside Mimics

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Abstract: A small library of nonhydrolyzable mimics of GM1 ganglioside, featuring galactose and sialic acid as pharmacophoric carbohydrate residues, was synthesized and tested. All compounds were synthesized from readily available precursors using high-performance reactions, including click chemistry protocols, and avoiding *O*-

glycosidic bonds. Some of the most active molecules also feature a point of further derivatization that can be used

Keywords: carbohydrates • cholera toxin • combinatorial chemistry • glycosides • weak affinity chromatography

for conjugation with polyvalent aglycons. Their affinity towards cholera toxin was assessed by weak affinity chromatography, which allowed a systematic evaluation and selection of the best candidates. Affinity could be enhanced up to one or two orders of magnitude over the affinity of the individual pharmacophoric sugar residues.

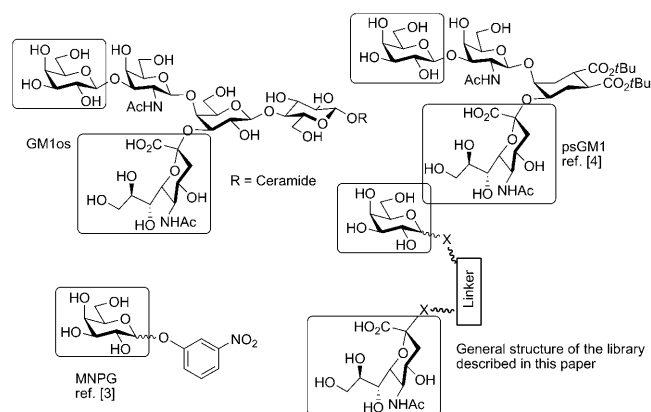
Introduction

Cholera toxin (CT) belongs to the AB₅ bacterial toxins family, named after the characteristic architecture comprising a single catalytically active component, A, and a nontoxic receptor-binding pentamer of B subunits.^[1] The B₅ pentamer is responsible for binding to GM1 ganglioside at the surface of host cells; the complete AB₅ holotoxin is required for the toxic effects. The CT family includes enterotoxins that are responsible for several disorders, from the relatively mild traveller's diarrhea (from *E. coli* heat-labile toxin, LT) to the much more serious cholera. Complexes formed between gangliosides and AB₅ toxins offer a paradigmatic model for studying the structural and thermodynamic basis

of protein-carbohydrate interactions. For medicinal chemistry, they may also provide key insights for the structure-based design of glycomimetic drug leads.^[2] A rational design of galactose-based ligands for CT and LT has been reported and recently reviewed.^[3] A series of CT ligands designed to mimic the three-dimensional structure of GM1 ganglioside has been reported^[4] and C-galactosides have been used as scaffolds to prepare a small library of nonhydrolyzable inhibitors of binding.^[5]

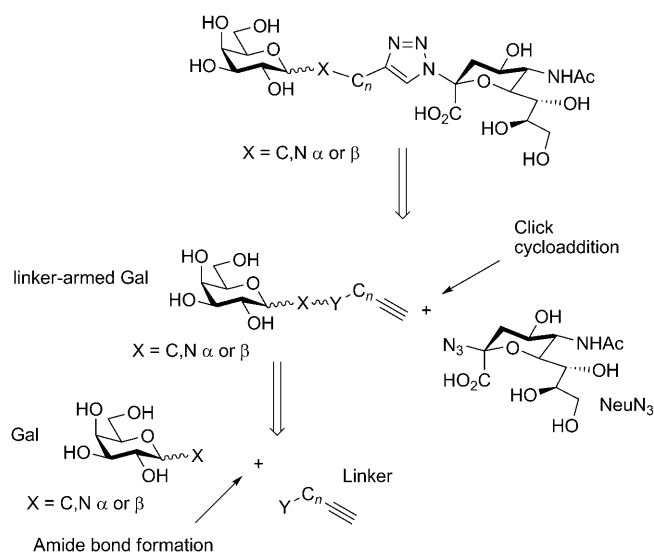
The B pentamer of CT (CTB) interacts with the soluble, monovalent oligosaccharide portion of GM1 (Galβ1-3GalNAcβ1-4(Neu5Acα2-3)Galβ1-4Glc, GM1os, Scheme 1) with a strong affinity (43 nM at room temperature, as measured by isothermal calorimetry (ITC)).^[6] The X-ray structure of the CTB-GM1os complex^[7] shows a “two-fingered grip” of the sugar on the toxin, created by a sialic acid “thumb” and a Galβ1-3GalNAc “forefinger”. The terminal galactose residue in the forefinger reaches a well-defined galactose binding pocket, which is lined by the indole side chain of Trp-88 and shielded from the solvent. The rest of the toxin binding site is shallow and exposed to solvent. The sialic acid (NeuAc) thumb interacts with a carboxylate binding region, which includes one highly conserved crystallographic water molecule. It has been shown by ITC that the individual “fingers” interact very weakly with the toxin (with dissociation constant in the high mM range),^[6] so the high binding affinity of the CTB-GM1os pair has been credited to structural preorganization of GM1os,^[8] which allows a near lock-and-key interaction with CTB.

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Scheme 1. GM1os, some known CT ligands (ps-GM1, MNPG), and general structure of the proposed library. The pharmacophoric sugar fragments are highlighted in the boxes.

This analysis suggests that functional mimics of GM1 may be designed by stringing together the two pharmacophoric sugar fragments, galactose and sialic acid, while trying to preserve the relative orientation adopted in GM1os. This concept was implemented in our initial work by generating pseudo-GM1 (psGM1, Scheme 1) structures that are close mimics of GM1os.^[4] Like other known inhibitors of CT binding, such as *meta*-nitrophenylgalactoside (MNPG, Scheme 1),^[3] this molecule contains enzymatically labile *O*-glycosidic linkages that limit their potential application in vivo. Furthermore, the synthetic methods used to connect the pharmacophoric sugar moieties in psGM1 are those of traditional carbohydrate chemistry, which are often laborious and low-yielding procedures. To develop metabolically stable and synthetically more accessible mimics of GM1os, we set our efforts towards the synthesis of a library of bi-functional compounds of the general formula shown in Scheme 1. The target molecules contain a galactose and a sialic acid moiety connected through a linker. By taking advantage of the terminal alkyne of the linker and the ready availability of the sialic acid azide **26**,^[9] a triazole was used to connect the NeuAc residue to the linker (Scheme 2) through a Cu-catalyzed (click) triazole formation reaction.^[10] The library design (Scheme 2) relied on combinatorial selection of the alkyne linkers and the galactose-mimicking fragments, with the following two constraints: 1) *O*-glycosides were excluded to stabilize the constructs against the activity of hydrolytic enzymes; 2) the functionalization of the galactose ring (X in Schemes 1 and 2) was chosen to allow a facile conjugation to the putative linkers. Selection of the linker and identification of the best bidentate ligand was achieved by synthesizing a small library of compounds and testing their CTB affinity by weak affinity chromatography (WAC). Herein we report our initial results.

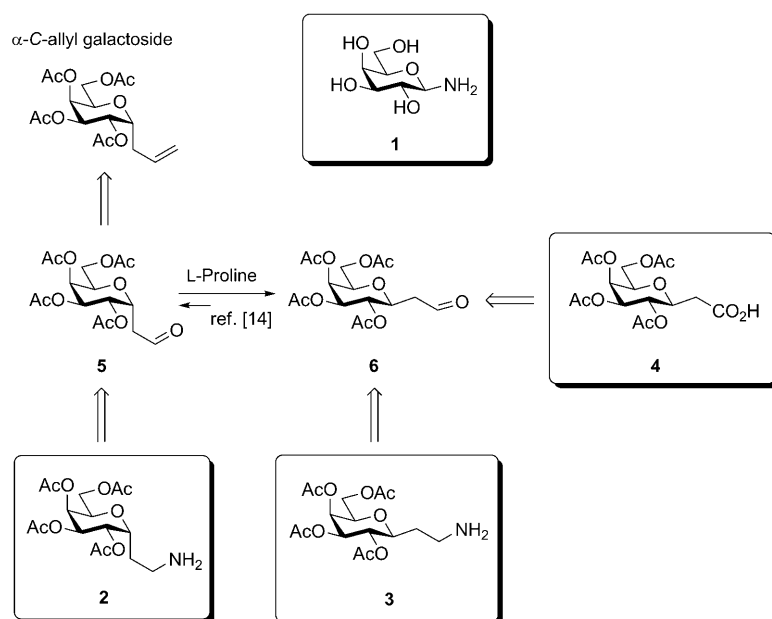


Scheme 2. General strategy for library assembly.

Results and Discussion

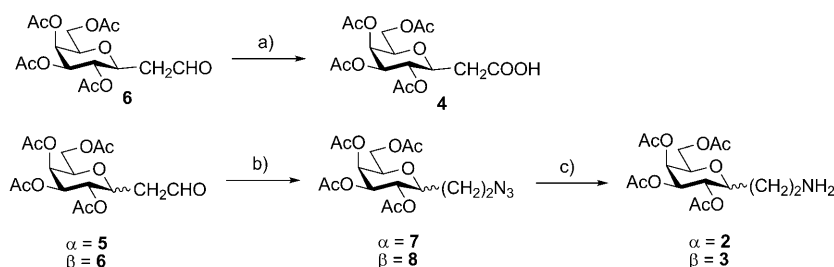
Synthesis of the library: A library of bidentate ligands was synthesized using linear alkyne linkers by following the strategy outlined in Scheme 2. Four functionalized Gal fragments were conjugated through an amide bond to linear alkyne linkers of variable length to obtain linker-armed galactose mimics. Incorporation of the sialic acid residue using click chemistry was planned as the final step of the synthesis, since Cu-catalyzed cycloaddition is known to tolerate a wide range of substrates, solvents, and reaction conditions.

The four functionalized Gal fragments (Scheme 3) used were: β -galactosyl amine **1**, α - or β -C-aminoethyl galactosides **2</**



Scheme 3. The functionalized Gal fragments used in this work.

yields (Scheme 4). Since direct reductive amination of α - and β -C-galactosyl aldehydes **5** and **6** to the corresponding amines **2** and **3** was complicated by an N-bisalkylation process and proceeded with poor yield, a two-step procedure via intermediate formation of azides **7** and **8** was used instead (Scheme 4).

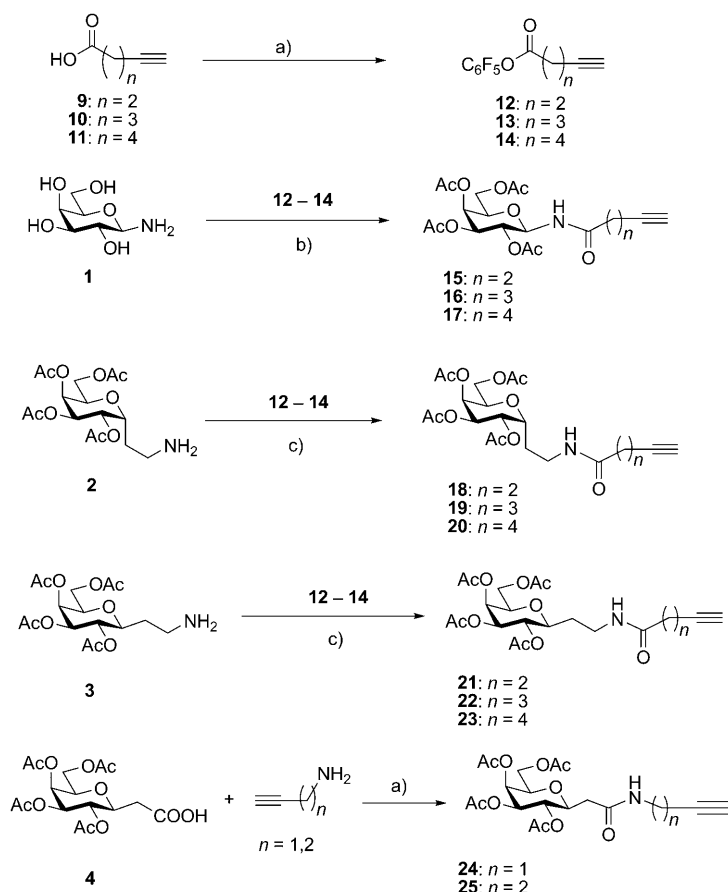
Scheme 4. Synthesis of **2–4**. Reagents and conditions: a) TEMPO/BAIB, 1:1 CH₃CN/H₂O; b) i) diphenylphosphoryl azide (DPPA), 120°C, microwave (MW); ii) NaBH₃(OAc); c) H₂, Pd/C.

For the synthesis of the linker-armed Gal fragments from amines **1–3**, three commercially available, linear, terminal alkynoic acids **9–11** (from C₅ to C₇) were converted into the corresponding pentafluorophenyl esters **12–14** and coupled with either freshly prepared crude galactosylamine **1**, or aminoethyl galactosides **2** and **3** (Scheme 5). Attempts to apply a direct Staudinger acylation procedure by treating pentafluorophenyl esters **12–14** with azides **7** or **8**^[17] did not lead to any significant improvement in overall yields.

Ligands with shorter linker chains were prepared by coupling the β -galactosyl acid **4**, with either propargylamine or

butynylamine to afford alkynes **24** and **25**, respectively (Scheme 5).

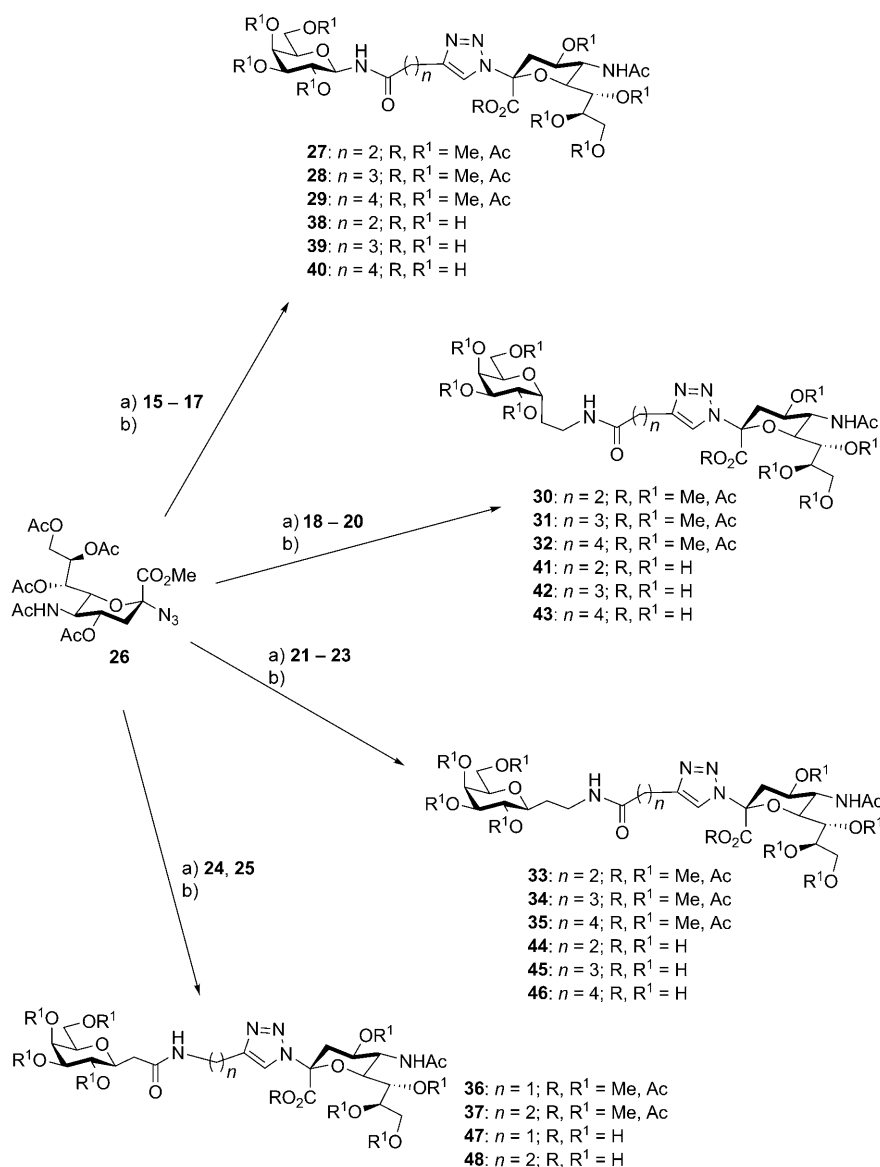
Finally, Cu-catalyzed (click) cycloaddition of alkynes **15–25** with fully



Scheme 5. Synthesis of linker-armed Gal fragments **15–25**. Reagents and conditions: a) $\text{C}_6\text{F}_5\text{OH}$, dicyclohexylcarbodiimide (DCC), THF; b) i) Et_3N , DMF; ii) $\text{Ac}_2\text{O}/\text{Py}$; c) Et_3N , THF.

(trifluoroacetic acid (TFA)), and treated with either benzoyl chloride or the pentafluorophenyl ester of phenylacetic acid to afford, after deprotection, amides **56** and **57** as a pair of inseparable isomers. Alternatively, acylation of the free amine with either phenylisocyanate or benzylisocyanate, followed by deacetylation, yielded the two ureas **58** and **59**.

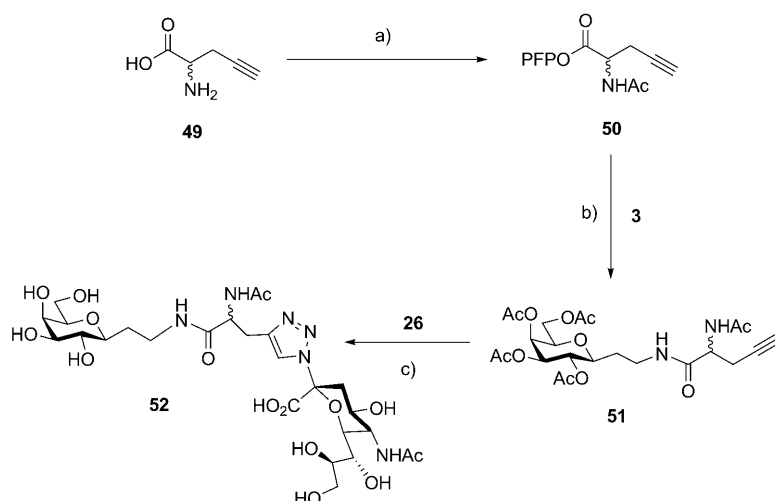
Ligand ranking by weak affinity chromatography (WAC):



Scheme 6. Synthesis of the bidentate adducts **38–48**. Reagents and conditions: a) CuSO₄/Na ascorbate; b) NaOH, MeOH/H₂O.

order of magnitude higher than the revelation threshold. These values may be assumed to result from simultaneous interaction of the two sugar fragments with the toxin, which apparently is optimally allowed in this series by the framework of ligand **44** (C₅ linker chain from β-Gal-C framework).

As mentioned above, further characterization of the interaction of **44** with CTB by fluorescence spectroscopy was precluded by the interference of the triazole moiety, which absorbs in the same range as CTB Trp-88, and, therefore, filters the excitant radiation. However, saturation transfer difference (STD) NMR spectroscopy^[25] experiments allowed the observation of binding events between CT and ligand **44**. In STD experiments, irradiation of the protein is followed by transfer of magnetization to the ligand



Scheme 7. Reagents and conditions: a) i) Ac_2O ; ii) $\text{C}_6\text{F}_5\text{OH}$, DCC, THF; b) Et_3N , THF; c) i) CuSO_4 /sodium ascorbate; ii) NaOH , $\text{MeOH}/\text{H}_2\text{O}$.

The loss of affinity upon branching of the linker suggests that the presence of the NHAc substituent disfavors the conformation required to allow simultaneous contact of the two pharmacophoric sugars with the toxin. However, this unfavorable factor could be offset by additional interactions between the linker side chain and the toxin binding site. Since the CTB binding site is known to possess a lipophilic patch near the sialic acid side-chain binding region,^[27] the contribution of more lipophilic side chains was explored by testing ligands **56–59** (Table 1, entries 15–21), which showed similar pH dependence to the parent compound **44**. As can be seen from Table 1, the affinity of most structures improved significantly compared with the *N*-acetyl derivative

52, to approach the potency of the underivatized ligand **44**. Among the compounds tested, one of the epimers of phenylacetamide (**57**) and one of the phenylureas (**58**) were found to be the most active ligands, with K_d values of 1.2 and 0.8 mM, respectively. When the two *R* and *S* isomers of **58** were synthesized separately, starting from enantiomerically pure *D*- or *L*-propargyl glycine, WAC analysis clearly showed that the *R* isomer of **58** had higher affinity than the corresponding *S* isomer (Table 1). This, in fact, had been suggested by docking models obtained by using Glide,^[28] which showed that the *R* epimer of the side-chain stereocenter should direct the *N*-substituent towards the lipophilic region of the binding site (Figure 3).

Conclusion

A modular approach to a library of new glycomimetic CT ligands has been developed. The cornerstone of this approach consists of tethering carbohydrate epitopes that are known to interact with the CTB binding site, galactose, and sialic acid, to a properly designed linker through nonhydrolyzable, non-*O*-glycosidic bonds. The Gal epitope was introduced as one of four simple *C*- or *N*-galactosides. The NeuAc residue

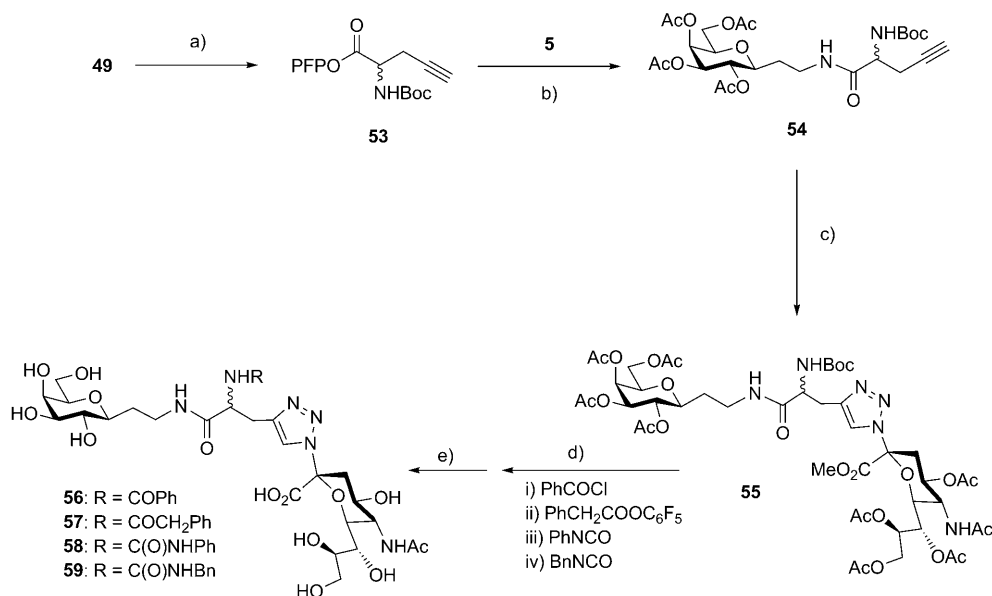


Table 1. Retention factor (k') and dissociation constant (K_d) of CT ligands at 23 °C determined with WAC at pH 6 and 7.

Entry	Ligand	pH 6		pH 7	
		k'	K_d [mM]	k'	K_d [mM]
1	MNPG ^[a]	1.28	1.3	1.58	1.1 ^[b]
2	38	< 0.05	> 30	< 0.05	> 30
3	39	< 0.05	> 30	< 0.05	> 30
4	40	< 0.05	> 30	< 0.05	> 30
5	41	0.17	9.9	0.09	19
6	42	0.38	4.4	0.19	9.1
7	43	0.09	20	0.05	31
8	44	2.17	0.8	1.25	1.3
9	45	0.76	2.2	0.37	4.6
10	46	0.25	6.6	0.14	12
11	47	< 0.05	> 30	< 0.05	> 30
12	48	< 0.05	> 30	< 0.05	> 30
13	(<i>R</i>)- 52 ^[c]	0.70	2.4	0.47	3.6
14	(<i>S</i>)- 52 ^[c]	0.07	26	< 0.05	> 30
15	(<i>R/S</i>)- 56 ^[c]	1.10	1.5	0.71	2.4
16	(<i>R</i>)- 57 ^[c]	1.35	1.2	0.86	2.0
17	(<i>S</i>)- 57 ^[c]	1.19	1.4	0.86	2.0
18	(<i>R</i>)- 58 ^[c]	2.			

of the ligand, thus increasing its preorganization and, ultimately, its affinity for CT.

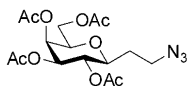
All compounds reported herein could be synthesized from precursors that are readily available on a large scale by using high-performance reactions, including click chemistry protocols. Previously known artificial CT ligands basically fall into two classes: the MNPG derivatives introduced by the group of Fan^[3] and the structural mimics of the GM1os reported by our group (Scheme 1).^[4] Both types contain enzymatically labile *O*-glycosidic linkages. Several of the non-hydrolyzable ligands described herein demonstrated reasonably high CT affinities that were similar to or higher than the affinity of MNPG.^[20,24] They are therefore among the most active monovalent CT ligands that do not closely reproduce the GM1 structure. Some of the most active molecules identified herein also feature a point of further derivatization that can be used for conjugation with polyvalent aglycons. Even though there was either no, or negligible, affinity gained by linker derivatization, this approach opens the way to the creation of polyvalent constructs that may be used to block the toxin in a therapeutically relevant context.^[18]

Interesting findings, such as pH dependence of the binding affinity of the ligands and the structural preference of the CTB binding site towards one of the two diastereomeric forms of the ligand, were assessed with WAC, which will be useful for further development of inhibitors of CT binding.

Experimental Section

General: WAC was performed essentially as described previously.^[20] In short, recombinant CTB (SBL vaccines, Stockholm, Sweden) was immobilized with reductive amination to aldehyde-derivatized Nucleosil silica (10 μ m, 300 Å; Macherey–Nagel, Düren, Germany) and packed into a 50 $\times$ 2.1 mm column. The number of binding sites on the column was determined with frontal chromatography to be 261 nmol at pH 7. All chromatography experiments were performed on an Agilent 1100 series HPLC system (Agilent Technologies, Santa Clara, CA, USA); the mobile phase was 0.15 M sodium chloride with 10 mM sodium phosphate (adjusted to pH 5.5, 6.0, 6.5, or 7.0 as indicated in the text); the flow rate was 0.1 mL min⁻¹ and the temperature was 23 °C; the injection volume was 5 μ L and the concentration of each sample was 20 μ g mL⁻¹. Detection was performed at 220 nm and the detector signal was collected with an Agilent ChemStation chromatography system. The retention factor (k') was calculated from the retention time (T_r) and the mobile phase hold up time, also called the void time (T_m), according to the equation: $k' = (T_r - T_m)/T_m$. The void time of the column was determined to be 1.55 min. Dissociation constants (K_d) of the ligands were calculated from the following equation: $K_d = B_{tot}/(k' \cdot V_m)$ in which B_{tot} is the number of binding sites in the column and V_m is the void

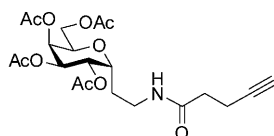
while stirring. The mixture was then subjected to microwave irradiation for 20 min at 120 °C (dynamic control) at 200 W power. The solvent was evaporated and the mixture was purified by flash chromatography (hexane/EtOAc) to afford the azide **8** (560 mg, 77.3 %). $[\alpha]_D^{25} = +1.8$ ($c = 1$ in CHCl_3); $^1\text{H NMR}$ (400 MHz, CDCl_3): $\delta = 5.41$ (dd, $J_1 = 1$, $J_2 = 3$ Hz, 1H; H-4), 5.07 (t, $J = 10$ Hz, 1H; H-2), 5.01 (dd, $J_1 = 3$, $J_2 = 10$ Hz, 1H; H-3), 4.16–4.46 (m, 2H; H-6), 3.86 (brt, $J = 7$ Hz, 1H; H-5), 3.69 (td, $J_1 = 4$, $J_2 = 9$ Hz, 1H; H-1), 3.41 (dt, $J = 6$ Hz, 2H; CH_2N_3), 1.92–2.15 (m, 12H; $4 \times \text{CH}_3\text{CO}$), 1.74–1.82 ppm (m, 2H; $\text{CH}_2\text{CH}_2\text{N}_3$); $^{13}\text{C NMR}$: $\delta = 170.5$, 170.2, 170.1, 169.9 ($4 \times \text{CH}_3\text{CO}$), 75.0 (C-1), 74.3 (C-5), 72.0 (C-3), 69.2 (C-2), 67.7 (C-4), 61.7 (C-6), 47.1 (CH_2N_3), 31.0 ($\text{CH}_2\text{CH}_2\text{N}_3$), 20.8, 20.7, 20.6 ppm ($4 \times \text{CH}_3\text{CO}$); MS (ESI): m/z : 423.9 [$M^+ + \text{Na}$].



Compound 2: Acetic acid (500 μL) and a catalytic amount of Pd/C was added to a stirred solution of the azide **7** (636 mg, 1.6 mmol) in a mixture of MeOH/ H_2O (5:1, 30 mL). The reaction vessel was filled with hydrogen and the reaction mixture was vigorously stirred at RT for ≈ 2 h until the reduction was complete. The reaction mixture was filtered through Celite and the resulting solution was evaporated and dried in vacuo to afford amine **2** as the acetate salt (834 mg, 100 %). $[\alpha]_D^{25} = +25.4$ ($c = 1$ in MeOH); $^1\text{H NMR}$ (400 MHz, CD_3Cl_3): $\delta = 5.39$ (t, $J_1 = J_2 = 3$ Hz, 1H; H-4), 5.25 (dd, $J_1 = 3$, $J_2 = 8$ Hz, 1H; H-3), 5.19 (dd

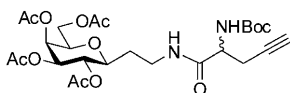
(acetate salt; 0.5 M) in THF. The reaction mixture was stirred at RT overnight, then the solvent was evaporated, and resulting solid was purified by flash chromatography (hexane/EtOAc).

Compound 18: The reaction of α -aminoethylgalactoside **2** (30 mg, 0.069 mmol, *N*-acetate form) with pentafluorophenyl ester **12** (22 mg, 0.083 mmol) afforded alkyne **18** (25 mg, 0.055 mmol, 80 %). ^1H NMR (400 MHz, CDCl_3): δ =6.18 (brt, J =6.5 Hz, 1 H; NH), 5.38 (t, J =3 Hz, 1 H; H-4), 5.05–5.15 (m, 2 H; H-2, H-3), 4.50 (dd, J_1 =8.3, J_2 =11 Hz, 1 H; H-6b), 4.20 (dt, J_1 =3, J_2 =11 Hz, 1 H; H-5), 4.05–4.10 (m, 1 H; H-1), 3.95 (dd, J_1 =11, J_2 =5 Hz, 1 H; H-6a), 3.38–3.50 (m, 1 H; $\text{CH}_2\text{bNC(O)}$), 3.08–3.18 (m, 1 H; $\text{CH}_2\text{aNC(O)}$), 2.45–2.55 (m, 2 H; $\text{HNC(O)CH}_2\text{CH}_2\text{CCH}$), 2.40–2.50 (m, 2 H; $\text{HNC(O)CH}_2\text{CH}_2\text{CCH}$), 1.95–2.07 (m, 13 H; $\text{C}\equiv\text{CH}$, $4\times\text{CH}_3\text{CO}$), 1.75–1.95 ppm (m, 2 H; $\text{CH}_2\text{CH}_2\text{NC(O)}$).



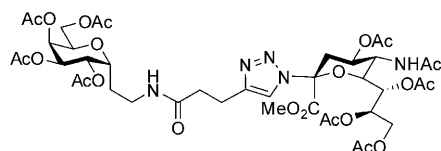
Compound 19: The reaction of α -aminoethylgalactoside **2** (32 mg, 0.073 mmol, *N*-acetate form) with pentafluorophenyl ester **13** (31 mg, 0.111 mmol) afforded alkyne **19** (27 mg, 0.058 mmol, 78 %). ^1H NMR (400 MHz, CDCl_3): δ =6.10 (brt, J =6.5 Hz, 1 H; NH), 5.38 (t, J =3 Hz, 1 H; H-4), 5.10–5.20 (m, 2 H; H-2, H-3), 4.57 (dd, J_1 =8.3, J_2 =11 Hz, 1 H; H-6b), 4.23 (dt, J_1 =3, J_2 =11 Hz, 1 H; H-5), 4.10–4.15 (m, 1 H; H-1), 4.00 (dd, J_1 =11, J_2

Compound 54: The reaction of the pentafluorophenyl ester **53** (234 mg, 0.619 mmol) with β -aminoethylgalactoside **3** (*N*-acetate salt, 224.5 mg, 0.516 mmol), afforded alkyne **54** (178 mg, 60%). ^1H NMR (400 MHz, CDCl_3): δ =6.59 (s, 0.5H; CH_2NH), 6.53 (t, J =6 Hz, 0.5H; CH_2NH), 5.42 (brs, 1H; H-4), 5.28 (s, 1H; NHBOc), 5.08 (t, J_1 =10 Hz, 1H; H-2), 4.99 (dd, J_1 =3, J_2 =10 Hz, 1H; H-3), 4.20–4.31 (m, 1H; CHNH), 4.06–4.18 (m, 2H; H-6), 3.86 (qd, J_1 =1, J_2 =6 Hz, 1H; H-5), 3.44–3.64 (m, 2H; H-1, $\text{CH}_2\text{CH}_2\text{NH}$), 3.25–3.40 (m, 1H; $\text{CH}_2\text{CH}_2\text{NH}$), 2.71–2.83 (m, 1H; $\text{CH}_2\text{C}\equiv\text{CH}$), 2.55–2.68 (m, 1H; $\text{CH}_2\text{C}\equiv\text{CH}$), 1.94–2.18 (m, 13H; $4\times\text{CH}_3\text{CO}$, $\text{C}\equiv\text{CH}$), 1.61–1.92 ppm (m, 2H; $\text{CH}_2\text{CH}_2\text{NH}$); ^{13}C NMR: δ =170.6–170.0 ($4\times\text{CH}_3\text{CO}$, CH_2NHCO), 77.4 (C-1), 74.7 (C-5), 71.7 (C-3), 69.1 (C-2), 67.9 (C-4), 62.0 (C-6), 53.2 (CHNH), 36.6 (CH_2NH), 31.0 ($\text{CH}_2\text{CH}_2\text{NH}$), 28.4 (*t*Bu), 22.8 ($\text{CH}_2\text{C}\equiv\text{CH}$), 20.7–20.9 ppm ($4\times\text{CH}_3\text{CO}$).



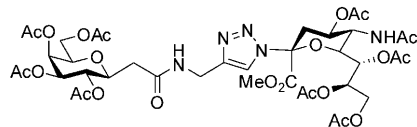
Compound 24: The pentafluorophenyl ester of β -caboxymethylgalactoside **4**

Compound 30: The reaction of alkyne **18** (23 mg, 0.050 mmol) with **26** following the general procedure described above afforded **30** (31 mg, 0.032 mmol, 63 %). ¹H NMR (400 MHz, CDCl₃): δ = 7.71 (s, 1H; H-Triaz), 6.45 (br s, 1H; NH), 5.28–5.40 (m, 3H; H-4Gal, H-7Neu, H-8Neu), 5.25 (d, *J* = 9.8 Hz, 1H; AcNH), 5.00–5.17 (m, 3H; H-4Neu, H-2Gal, H-3Gal), 4.46–4.50 (m, 1H; H-6bGal), 3.90–4.30 (m, 7H; H-9Neu, H-5Neu, H-6Neu, H-1Gal, H-5Gal, H-6aGal), 3.71 (s, 3H; COOCH₃), 3.28–3.46 (m, 2H; CH_{2b}NC(O), H-3_{eq}Neu), 3.24–3.06 (m, 1H; CH_{2a}NC(O)), 2.49–2.71 (m, 5H; H-3_{ax}Neu, CH₂CH₂C(O)NH), 1.80–2.15 (m, 27H; 9 × CH₃CO), 1.60–1.75 (m, 1H; CH_{2b}CH₂NC(O)), 1.50–1.60 ppm (m, 1H; CH_{2a}CH₂NC(O)).



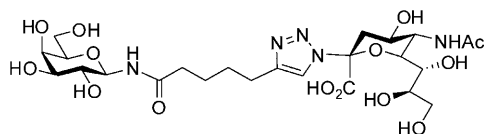
Compound 31: The reaction of alkyne **19** (27 mg, 0.058 mmol) with **26** following the general procedure described above afforded **31** (51 mg, 0.052 mmol, 90 %). ¹H NMR (400 MHz, CDCl₃): δ = 7.70 (s, 1H; H-Triaz), 6.30 (br s, 1H; NH), 5.36–5.48 (m, 3H; H-4Gal, H-7Neu, H-8Neu), 5.33 (d, *J* = 9.0 Hz, 1H; AcNH), 5.11–5.23 (m, 3H; H-4Neu, H-2Gal, H-3Gal), 4.45 (t, *J* = 7.5 Hz, 1H; H-6bGal), 4.23–4.37 (m, 3H; H-9Neu, H-5Gal), 4.00–4.19 (m, 4H; H-5Neu, H-6Neu, H-1Gal, H-6aGal), 3.70 (s, 3H; COOCH₃), 3.39–3.53 (m, 2H; CH_{2b}NC(O), H-3_{eq}Neu), 3.12–3.30 (m, 1H; CH_{2a}NC(O)), 2.68–2.90 (m, 2H; CH₂CH₂CH₂C(O)NH), 2.68 (t, *J* = 12 Hz, 1H; H-3_{ax}Neu), 2.28 (t, *J* = 7 Hz, 2H; CH₂CH₂C(O

Triaz, 6.80 (br s, 1H; NHCH_2), 5.31–5.51 (m, 3H; H-4Gal, H-8Neu, H-7Neu), 4.96–5.20 (m, 3H; H-4Neu, H-2Gal, H-3Gal), 4.47–4.67 (m, 2H; H-6Neu, H-9aNeu), 4.21–4.36 (m, 2H; H-5Neu, H-9bNeu), 3.90–4.17 (m, 6H; H-1Gal, H-5Gal, H-6Gal, NHCH_2 -Triaz), 3.78 (s, 3H; CH_3OCONeu), 3.40 (dd, $J_1=4$, $J_2=13$ Hz, H-3_{eq}Neu), 2.65 (t, $J=12$ Hz, 1H; H-3_{ax}Neu), 2.48 (brs, 2H; Gal- CH_2), 1.80–2.20 ppm (m, 27H; $8\times\text{CH}_3\text{CO}$, CH_3CONH).



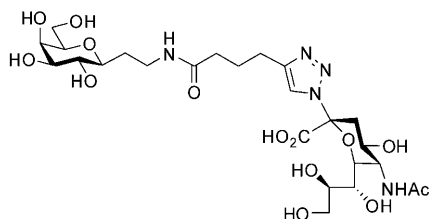
Compound 37: The reaction of alkyne **25** (13.1 mg, 0.0297 mmol) with **26** following the general procedure described above afforded **37** (8.0 mg, 0.0132 mmol, 45%). ^1H NMR (400 MHz, CDCl_3): $\delta=7.77$ (s, 1H; CH-Triaz), 6.79 (t, $J=6$ Hz, 1H; NHCH_2), 5.40–5.48 (m, 2H; H-4Gal, H-8Neu), 5.37 (dd, $J_1=2$, $J_2=9$ Hz, 1H, H-7Neu), 5.18 (dt, $J_1=4$, $J_2=12$ Hz, 1H; H-4Neu), 5.10 (t, $J=10$ Hz, 1H; H-2Gal), 5.05 (dd, $J_1=3$, $J_2=10$ Hz, 1H; H-3Gal), 4.28–4.35 (m, 2H; H-6Neu, H-9aNeu), 4.03–4.15 (m, 4H; H-6Gal, H-5Neu, H-9bNeu), 3.91–3.99 (m, 2H; H-1Gal, H-5Gal), 3.80 (s, 3H; CH_3OCONeu), 3.55–3.72 (m, 2H; CHNH), 3.43 (dd, $J_1=4$, $J_2=13$ Hz, H-3_{eq}Neu), 2.93 (t, $J=6$ Hz, 2H; CH CH_2 -Triaz), 2.66 (dd, $J_1=12$, $J_2=13$ Hz, 1H;

400 MHz): δ = 177.1, 175.2 (C=O), 121.4 (CH-triaz), 79.0 (C-1Gal), 76.7, 74.3, 73.1, 71.1, 70.7, 68.9, 68.7, 67.8, 67.6, 66.8 (C-2Gal, C-3Gal, C-4Gal, C-5Gal, C-4Neu, C-5Neu, C-6Neu, C-7Neu, C-8Neu), 62.7, 61.1 (C-6Gal, C-9Neu), 51.6 (C-5Neu), 39.4 (C-3Neu), 34.6 ((CH₂)₃CH₂C(O)NH), 25.0 (CH₂(CH₂)₂CH₂C(O)NH), 24.2 (CH₂(CH₂)₃C(O)NH), 21.1 ppm (NHCOCH₃); HRMS (FT-ICR, ESI): m/z calcd for C₂₄H₃₉N₅O₁₄ [M-H]⁻: 620.241530 found: 620.24059.



Ligand 41: Deprotection of **30** (23 mg, 0.024 mmol) following the general procedure described above afforded ligand **41** (14 mg, 0.023 mmol, 95 %). [α]_D = +40.9 (c = 0.2 in MeOH); ¹H NMR (D₂O, 400 MHz): δ = 7.87 (s, 1H; H-triaz), 3.95–4.00 (m, 14H; H-4Neu, H-5Neu, H-6Neu, H-7Neu, H-8Neu, 2×H-9Neu, H-1Gal, H-2Gal, H-3Gal, H-4Gal, H-5Gal, 2×H-6Gal), 3.03–3.15 (m, 3H; H-3_{eq}Neu, CH₂NHC(O)), 2.87 (t, J = 7.1 Hz, 2H; CH₂CH₂C(O)NH), 2.44 (t, J = 7.1 Hz, 2H; CH₂CH₂C(O)NH), 2.08 (m, 1H; H-3_{ax}Neu), 1.90 (s, 1H; NHC(O)CH₃), 1.50–1.75 ppm (m, 2H; CH₂CH₂NHC(O)); ¹³C NMR (D₂O, 400 MHz): δ = 177.4, 174.8 (C=O), 120.0 (CH

(CH₂NHC(O)), 33.7 (CH₂CH₂C(O)NH), 29.3 (CH₂CH₂NHC(O)), 23.8 (CH₂CH₂CH₂), 22.7 (CH₂CH₂C(O)NH), 20.9 ppm (NHCOCH₃); HRMS (FT-ICR, ESI): *m/z* calcd for C₂₅H₄₁N₅O₁₄: 634.257180 [M-H]⁻; found: 634.25636.



Ligand 46: Deprotection of **35** (13 mg, 0.013 mmol) following the general procedure described above afforded ligand **46** (8 mg, 0.012 mmol, 95%). [α]_D = +27.4 (*c* = 0.11 in MeOH); ¹H NMR (D₂O, 400 MHz): δ = 7.81 (s, 1H; H-triaz), 3.31–3.90 (m, 12H; H-4Neu, H-5Neu, H-6Neu, H-7Neu, H-8Neu, 2×H-9Neu, H-3Gal, H-4Gal, H-5Gal, 2×H-6Gal), 3.27 (t, *J* = 9.4 Hz, 1H; H-2Gal), 3.00–3.23 (m, 4H; CH₂NHC(O), H-3_{eq}Neu, H-1Gal), 2.57 (t, *J* = 9.4 Hz, 1H; CH₂CH₂C(O)NH), 2.03–2.15 (m, 3H; CH₂CH₂C(O)NH, H-3_{ax}Neu), 1.83–1.96 (m, 4H; NHC(O)CH₃, CH_{2b}CH₂NH), 1.37–1.57 ppm (m, 5H; (CH₂)₂, CH_{2a}CH₂NH); ¹³C NMR (D₂O, 400 MHz): δ = 175.5, 173.8, 169.7 (C=O), 146.8 (C-triaz), 119.5 (CH-triaz), 89.5 (C-2Neu), 77.4, 72.9, 72.8, 70.2, 68.0, 66.9, 66.8 (C-3Gal, C-4Gal, C-5Gal, C-4Neu, C-6Neu, C-7Neu, C-8Neu), 76.2 (C-1Gal), 69.6 (C-2Gal), 61.6, 60.2 (C-6Gal, C-9Neu), 50.4 (C-

evaporated, and the crude product was purified by flash chromatography ($\text{CH}_2\text{Cl}_2/\text{acetone}$ gradient). Deprotection of the product (9 mg, 0.0082 mmol) afforded ligand **56** as a 1:1 mixture of diastereomers (6.0 mg, 0.0081 mmol, 58 %). ^1H NMR (400 MHz, D_2O): δ = 8.13 (s, 1H; H-triaz), 7.77 (dd, J = 4.3, 5.9 Hz, 2H; Ph), 7.67–7.73 (m, 1H; Ph), 7.60 (t, J = 7.5 Hz, 2H; Ph), 4.85 > (brt, J = 6.5 Hz, 1H; $\text{CH}(\text{NHBz})\text{CH}_2$), 3.48–4.00 (m, 12H; H-4Neu, H-5Neu, H-6Neu, H-7Neu, H-8Neu, 2 $\times$ H-9Neu, H-3Gal, H-4Gal, H-5Gal, 2 $\times$ H-6Gal), 3.43 (m, 1H; H-2Gal), 3.17–3.38 (m, 5H; $\text{CH}(\text{NHBz})\text{CH}_2$, $\text{CH}_2\text{CH}_2\text{NHC}(\text{O})$, H-3_{ax}Neu), 2.20–2.30 (m, 1H; H-3_{eq}Neu), 1.85–2.07 (m, 4H; CH_3CO , $\text{CH}_2\text{CH}_2\text{NHC}(\text{O})$), 1.50–1.64 ppm (m, 1H; $\text{CH}_2\text{CH}_2\text{NHC}(\text{O})$); ^{13}C NMR (400 MHz, D_2O): δ = 132.6, 128.8, 127.8 (Ph), 122.1 (C-triaz), 78.7, 77.5, 74.1, 71.2, 69.7, 68.7, 67.7 (C-3Gal, C-4Gal, C-5Gal, C-6Gal, C-4Neu, C-6Neu, C-7Neu, C-8Neu), 77.5 (C-1Gal), 70.7 (C-2Gal), 62.4, 61.2 (C-6Gal, C-9Neu), 54.1 ($\text{CH}(\text{NHAc})\text{CH}_2$), 51.3 (C-5Neu), 39.5 (C-3Neu), 30.1 ($\text{CH}_2\text{CH}_2\text{NHC}(\text{O})$), 27.0 ($\text{CH}_2\text{CH}_2\text{NHC}(\text{O})$), 21.8 ppm (CH_3CO); HRMS (FT-ICR, ESI): m/z calcd for $\text{C}_{31}\text{H}_{43}\text{N}_6\text{O}_{15}$: 739.27919 [$M-\text{H}$] $^-$; found: 739.27851

(NHCOCH₃); HRMS (FT-ICR, ESI): *m/z* calcd for C₃₂H₄₆N₇O₁₅: 768.30574 [*M*–H][–]; found: 768.30668.

Acknowledgements

Support by COST-D34 and Consorzio Interuniversitario Nazionale Metodologie e Processi Innovativi di Sintesi (CINMPIS) is gratefully acknowledged. We thank Maurizio Benaglia for a kind gift of PEG-supported proline and Francesca Vasile for performing the STD titration experiments. P.C. was supported by a Marie Curie International fellowship within the 6th EU Framework Programme (MIF1-CT-2006-040158).

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