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## 3-Azido-3-deoxy-glycopyranoside Derivatives as Scaffolds for the Synthesis of Carbohydrate-Based Universal Pharmacophore Mapping Libraries

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**Abstract**—The regiospecific syntheses of six monosaccharide scaffolds, 1–6, containing a carboxylic acid, an azido and a free hydroxyl group were accomplished through the utilization of a key intermediate, namely, methyl 3-azido-3-deoxy- $\beta$ -D-glucopyranoside (10). Scaffold 2 was also used in generating combinatorial libraries using solid-phase methodologies. © 2003 Elsevier Science Ltd. All rights reserved.

One aspect of combinatorial chemistry focuses on generating chemically diverse libraries suitable for broad screening. Our search for novel combinatorial platforms for preparing broad screening libraries lead us to investigate monosaccharides as versitile scaffolds. Conceptually, the use of a monosaccharide nucleus as a scaffold to display substituents in a stereochemically defined way was first pioneered by Hirschmann and coworkers in their effort to identify a mimetic of a known biologically active agent.<sup>1,2</sup>

We were particularly attracted to monosaccharides as combinatorial scaffolds because of the unique set of characteristics they possess. The monosaccharide system provides a rigid ring system with a defined three dimensional spatial arrangement of substituents. The high functional density of this nucleus provides multiple sites for chemical modification. However, the difficulty faced in trying to chemically differentiate each of the hydroxyl groups on any natural sugar nucleus presents a problem if one wants to use a sugar nucleus as a combinatorial scaffold.<sup>3</sup>

In designing a practical monosaccharide scaffold system we focused on building a system that would provide a classical three-point pharmacophoric recognition motif (Fig. 1).<sup>4,5</sup> Then to circumvent the chemoselectivity problem, we used an easily differentiated functional group triad that included a carboxylic acid, a free hydroxyl group and a masked amino group. The two remaining hydroxyl groups on a hexose nucleus would be converted to their methyl ethers. Methyl ethers were chosen because they provide permanent protecting groups that are stable to a wide variety of chemical transformations and we hoped that they would minimize the contribution of these sites to molecular recognition events maximizing the effect of the three point motif. To accommodate a carboxylic acid at secondary carbon sites, we chose to use a carboxymethyl unit. Although not isosteric with a C-6 uronic acid, the carboxymethyl group could be easily introduced by direct alkylation of a sugar hydroxyl group.

In our earlier report, our scaffolds incorporated a fluorenylmethoxycarbonyl (Fmoc) protected amino group as the masked amino functionality.<sup>4,5</sup> However, it became apparent to us that the Fmoc group lacked the chemical stability we required for doing a wide variety of chemical transformations. To circumvent this problem, we chose to replace the NHFmoc group with an azido group.

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OMe

Figure 1.

The scaffold design described above provides the potential for accessing a large diverse set of scaffolds that could be used to explore a wide range of threedimensional space. We envisioned that families of these scaffolds, would find utility in mapping pharmacophore recognition sites on target biomolecules and hence we refer to these libraries a carbohydrate-based universal pharmcophore mapping libraries. In this report we present the synthesis of a family of 3-azido containing monosaccharide scaffolds 1, 2, 3, 4, 5, and 6 and demonstrate the construction of a combinatorial library using scaffold 2.

The syntheses of the 3-azido scaffolds 1, 2, 3, 4, 5, and 6 began with construction of the common intermediate methyl 3-azido-3-deoxy- $\beta$ -D-glucopyranoside 10. Intermediate 10 would allow us to construct scaffold systems having a carboxylic acid unit or a free hydroxyl group in the 2-, 4- or 6-position of a glucose or galactose nucleus. For the synthesis of 10, we employed 1,2,4,6-tetra-Oacetyl-3-azido-3-deoxy-D-glucopyranose (7) which was prepared from 1,2-5,6-di-O-isopropylidene-a-D-allofuranose using a previously described procedure for the preparation of 3-azido-3-deoxy-D-mannose.<sup>6</sup> Treatment of 7 with Me<sub>3</sub>SiBr and BiBr<sub>3</sub><sup>7</sup> in CH<sub>2</sub>Cl<sub>2</sub> gave crude bromide 8. Treatment of 8 with MeOH– $CH_2Cl_2$  in the presence of Ag<sub>2</sub>O followed by de-O-acetylation with methanolic sodium methoxide afforded 10 in 87% yield after chromatographic purification on silica gel. The <sup>1</sup>H NMR spectrum [ $\delta$  4.25 (d, J = 7.5 Hz, H-1)] as well as <sup>13</sup>C NMR spectrum [δ 105.23 (C-1)] confirmed the formation of  $\beta$ -linkage at C-1 (Scheme 1).

The synthesis of scaffold 1 from intermediate 10, required that we oxidize the C-6 hydroxyl group to the corresponding carboxylic acid in the presence of the



**5** X = OMe, Y = OCH<sub>2</sub>CO<sub>2</sub>H, Z = CH<sub>2</sub>OH **6** X = OH, Y = OCH<sub>2</sub>CO<sub>2</sub>H, Z = CH<sub>2</sub>OMe

other functionality (Scheme 2). This was accomplished by first protecting the C-4 and C-6 hydroxyl groups as a 4,6-*O*-*p*-methoxybenzylidene cyclic acetal using anisaldehyde dimethyl acetal and *p*-toluenesulfonic acid monohydrate in DMF to give **11**. Acetylation of the free hydroxyl group followed by reductive ring opening<sup>8</sup> gave **13** in 96% yield.

With 13 providing the needed azido sugar with each of the hydroxyl groups differentiated we were able to proceed with completing the synthesis of scaffold 1. This was done by first methylating the C-4 hydroxyl group of 13 and removing the 6-*O*-*p*-methoxybenzyl group with DDQ.<sup>9</sup> Oxidation with Jones reagent<sup>10</sup> followed by de-*O*-acetylation produced compound 1 in 88% yield after silica gel column chromatography.

Scaffolds containing a carboxylate functionality attached to either the 2- or 4-position of the sugar nucleus required that we introduce a carboxymethyl substituent. In each case, we were able to accomplish the attachment of this carboxymethyl group by alkylation of a free hydroxyl group with methyl 2-bromo-acetate and Ag<sub>2</sub>O-KI. For the synthesis of scaffold 2 containing a carboxymethyl group at C-2, we began with compound 11 having a free hydroxyl at C-2. Alkylation with methyl 2-bromo-acetate gave 17 in 80% yield (Scheme 3). Conversion of the 4,6-benzylidene derivative 17 to the 4-O-methyl-6-hydroxy derivative 20 was accomplished as previously described for the preparation of compound 15. Hydrolysis of methyl ester in compound 20 provide scaffold 2 in 94% yield.

The synthesis of the 2,3,6-substituted scaffold 3 began with previously prepared intermediate 17 (Scheme 4); however, unlike in the preparation of scaffold 2, here we



Scheme 1. Reagents and conditions: (a) Me<sub>3</sub>SiBr, BiBr<sub>3</sub>, CH<sub>2</sub>Cl<sub>2</sub>, 4 h; (b) Ag<sub>2</sub>O, CaSO<sub>4</sub>, CH<sub>2</sub>Cl<sub>2</sub>, MeOH, 1 h; (c) MeOH, MeONa, 4 h.



Scheme 2. Reagents and conditions: (a)  $MeO-C_6H_4-CH(OMe)_2$ , p-TSA, DMF, 4 h; (b)  $C_5H_5N$ ,  $Ac_2O$ , 4 h; (c)  $NaBH_3CN$ , DMF, TFA, 3 Å molecular sieves, 48 h; (d)  $Ag_2O$ ,  $CH_3I$ , DMF, 16 h; (e) DDQ,  $CH_2Cl_2$  saturated with water, 2 h; (f) Jones reagent, acetone, 1.5 h; (g) MeOH-MeONa, 4 h.



Scheme 3. Reagents and conditions: (a) Ag<sub>2</sub>O, BrCH<sub>2</sub>COOMe, KI, DMF, 16 h; (b) NaBH<sub>3</sub>CN, DMF, TFA, 3 Å molecular sieves, 48 h; (c) Ag<sub>2</sub>O, CH<sub>3</sub>I, DMF, 16 h; (d) DDQ, CH<sub>2</sub>Cl<sub>2</sub> saturated with water, 2 h; (e) Dioxane, 2N NaOH, 1 h.



Scheme 4. Reagents and conditions: (a) NaBH<sub>3</sub>CN, Me<sub>3</sub>SiCl, 3 Å molecular sieves, CH<sub>3</sub>CN, 4 h; (b) 21, Ag<sub>2</sub>O, CH<sub>3</sub>I, DMF, 16 h; (c) DDQ, CH<sub>2</sub>Cl<sub>2</sub> saturated with water, 2 h; (d) dioxane, 2N NaOH, 1 h.

needed to achieve acetal ring opening that would favor *p*-methoxybenzyl protection on the C-4 hydroxyl. This was accomplished by treating compound **17** with Me<sub>3</sub>SiCl–NaBH<sub>3</sub>CN<sup>11</sup> in acetonitrile. These conditions provide the 6-*O*-*p*-methoxybenzyl derivative **18** and 4-*O*-*p*-methoxybenzyl derivative **21** in 1:2 ratio. The C-6 hydroxyl group was then methylated using Ag<sub>2</sub>O and methyl iodide and the *p*-methoxybenzyl group was removed under oxidative conditions. Hydrolysis of the methyl ester gave scaffold **3** in 57% overall yield.

For the preparation of scaffold 4, we needed to differentiate the C-2, C-4, and C-6 hydroxyl groups as we had done in the preparation of scaffold 1. Therefore, we again used intermediate 11 and employed the regioselective *p*-methoxybenzylidene ring opening protocol (Scheme 5). Starting with 11, treatment with CH<sub>3</sub>I-NaH in DMF gave the C-2 methyl ether 23 in 95% yield. Acetal ring opening and attachment of the carboxymethyl group gave 25. Removal of the C-6 hydroxyl protecting group and ester hydrolysis gave scaffold 4 in 38% overall yield.

Synthesis of galactose scaffolds **5** and **6** from the glucopyranosyl intermediate **18** required inverting the stereochemistry at C-4. This inversion was accomplished through intermediacy of the C-4 triflate **27** which was obtained from **18** on treatment with triflic anhydride in dichloromethane (Scheme 6). Nucleophilic displacement of the C-4 triflate with tetrabutyl-ammonium benzoate in toluene gave the axial benzoate **28** in 86% yield. Removal of the *p*-methoxybenzyl protecting group gave **29**. We next wanted to methylate the C-6 hydroxyl group; however, simultaneous transfer of the benzoate group from the C-4–C-6 hydroxyl group and methylation of the C-4 hydroxyl occurred on exposure



Scheme 5. Reagents and conditions: (a) NaH, DMF, CH<sub>3</sub>I, 1 h; (b) NaBH<sub>3</sub>CN, DMF, TFA, 3 Å molecular sieves, 48 h; (c) Ag<sub>2</sub>O, BrCH<sub>2</sub>COOMe, KI, DMF, 4 h; (d) DDQ, CH<sub>2</sub>Cl<sub>2</sub> saturated with water, 2 h; (e) Dioxane, 2N NaOH,1 h.



Scheme 6. Reagents and conditions: (a) Triflic anhydride, pyridine,  $CH_2Cl_2$ , 0 °C, 2 h; (b) Tetrabutylammonium benzoate, toluene, 1 h; (c) DDQ,  $CH_2Cl_2$  saturated with water, 2 h; (d)  $Ag_2O$ ,  $CH_3I$ , DMF, 16 h; (e) Dioxane, 2N NaOH, 1 h.



Scheme 7. Reagents and conditions: (a) Triflic anhydride, pyridine,  $CH_2Cl_2$ , 0 °C, 2 h; NaNO<sub>2</sub>, DMF, 4 h; (b) 95% aq. TFA,  $CH_2Cl_2$ , 2 h; (c) Trimethyloxonium tetrafluoroborate, 2,6-di-*tert*-butyl-4-methyl-pyridine,  $CH_2Cl_2$ , 16 h; (d) Dioxane, 2 N NaOH, 1 h.

to  $Ag_2O/CH_3I$  in DMF providing compound **30** in 94% yield. With compound **30** in hand, base hydrolysis gave scaffold **5** in 93% yield (Scheme 6).

To access scaffold **6**, triflate **27** was treated with sodium nitrite<sup>12</sup> in DMF providing the desired C-4 axial hydroxyl group (Scheme 7). Subsequent removal of the C-6 *p*-methoxybenzyl group with TFA–CH<sub>2</sub>Cl<sub>2</sub> provided diol **31** in 40% yield after silica gel column chromato-

graphy. Selective methylation of **31** with trimethyloxonium tetrafluoroborate-2,6-di*tert* butyl-4-methylpyridine<sup>13</sup> gave the 6-*O*-methyl derivative **32**. The hydrolysis of methyl ester provided **6** in 87% yield.

To demonstrate the use of a 3-azido scaffold in the construction of a library of substituted monosaccharides we chose scaffold **2**. Using Rink amide resin<sup>14</sup> as the solid support, a blocked amino acid,



Scheme 8. Reagents and conditions: (a) (i) 20% piperidine, DMF, 0.5 h; (ii) Fmoc histidine (trt), HATU, DIEPA, DMF, 16 h; (b) 20% piperidine, DMF, 0.5 h; (c) 2, HATU, DIEPA, DMF, 16 h; (d) isocyanate, THF, Et<sub>3</sub>N, 6–16 h or isocyanate, DMF, CuCl; (e) Me<sub>3</sub>P, THF/EtOH/H<sub>2</sub>O(4/4/1); (f) carboxylic acid, HATU, DIEPA, DMF, 16 h; (g) 20% TFA, CH<sub>2</sub>Cl<sub>2</sub>, 2% Et<sub>3</sub>SiH.



## Figure 2.

Fmoc-histidine (Trt), was first coupled to the support (Scheme 8). Removal of the Fmoc protecting group was then followed by coupling of scaffold 2 to the free amino group by amide bond formation to the C-2 carboxymethyl group of 2. The resin-bound scaffold was then treated with a series of isocyanates in the presence of CuCl in DMF to generate the corresponding C-6 carbamates. Reduction of the C-3 azido group with trimethylphosphine generated the desired amines which were then reacted with a series of carboxylic acids to provide the corresponding amides.

A library of 48 compounds was prepared using the directed sorting mix-and-split synthesis method<sup>15</sup> (Fig. 2).

All 48 compounds were analyzed by LC/MS using evaporative light-scattering detection (ELSD). The yield of each product was determined from the ELSD trace using standard curves and product identity was determined by electrospray mass spectal analysis. The average yield for the six-step sequence was 55%. One of the library products was further purified by preparative HPLC and its structure was confirmed by NMR spectroscopy and LC/MS analysis to be the desired compound **40**.<sup>16</sup>



In conclusion, we believe that carbohydrate scaffolds are biologically relevant molecular platforms that can be used in combinatorial library strategies to identify unique ligands for a wide variety of biomolecular drug targets. To exploit this potential, we have developed a series of 3-azido glycopyranosides.

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