

# Synthesis of alkylated sugar amino acids: conformationally restricted L-Xaa-L-Ser/Thr mimics†

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Two synthetic strategies for the generation of  $\delta$ -substituted pyranoid sugar amino acids (SAAs) are evaluated. The first employs chiral nonracemic *tert*-butane sulfinamides as key reagents. Regardless of the stereochemistry of the applied sulfinamide, the product formed has a stereochemistry resembling that of a D amino acid at C7. Direct Grignard reaction on formyl-tetra-*O*-benzyl- $\beta$ -D-*C*-glucopyranoside in the second strategy and subsequent Mitsunobu inversion, yields the L,L-dipeptide isomers.

## Introduction

As part of our ongoing research on artificial peptide-like materials, we recently reported the synthesis of glucopyranose-based sugar amino acids (SAAs) **1** (Fig. 1).<sup>1</sup> In dipeptide isosteres **1**, the stereocentre at C2 has the *S*-configuration, thereby resembling the  $\alpha$ -carbon in L-serine or L-threonine.<sup>2</sup> The secondary amine at the N-terminus (C7) in **1** resembles the side-chain at the  $\alpha$ -carbon of amino acids other than glycine. By tuning the nature of the R<sup>1</sup> group, functionalities corresponding to specific amino acid side-chains can be incorporated into the SAAs.<sup>3</sup> This feature distinguishes compounds **1** from SAAs reported in the literature,<sup>4</sup> of which the large majority are primary amines.<sup>5</sup> As a whole, SAAs **1** can be viewed as conformationally constrained H-Xaa-Ser/Thr-OH mimics. The configuration of C7 corresponds to that of the  $\alpha$ -carbons in D-amino acids, rather than the proteinogenic L-amino acids. In contrast to stereochemical control over the C-terminal portion, which originates from selection of the carbohydrate template, controlling the configuration of the newly introduced stereocentre at C7 stems from asymmetric organic synthesis. The work presented here concerns adaptation of the synthetic strategy we applied to prepare SAAs **1** to provide C7-*S* configured SAA building blocks **2**.

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Protected SAAs **1** were previously prepared using the stereoselective alkylation of *R*-*tert*-butanesulfinamide **4**, which was obtained by the condensation of formyl tetra-*O*-benzyl- $\beta$ -D-*C*-glucopyranoside **3** with *R*-*tert*-butanesulfinyl amide<sup>7</sup> (Scheme 1). Alkylation of compound **4**, subsequent acid-mediated hydrolysis of the *R*-*tert*-butanesulfonyl group and instalment of the Fmoc protective group gave compound **7**, which could be transformed into carboxylate **8** by selective acidolysis of the primary benzyl ether, ester hydrolysis and oxidation. The alkylation of sulfinamide **4** proceeds in good diastereoselectivity. For instance, reaction of **4** with 3 equivalents of MeMgBr in methylene chloride at  $-78$  °C gave *R*-methyl adduct **5** in 20-fold excess over the other diastereoisomer **6**. Similar results were obtained by using toluene as a solvent. Performing the alkylation in THF resulted in a drop in diastereoselectivity (**5**–**6** = 13 : 1).

Either the *R*-*tert*-butanesulfinyl chiral auxiliary or the chiral carbohydrate template, or a combination of both may be responsible for the observed diastereoselectivity. Would the first be true, then L,L-dipeptide isostere SAAs **2** would be directly accessible by following the same synthetic scheme, but employing *S*-*tert*-butanesulfinamide as the chiral auxiliary. In order to investigate this possibility, we prepared *S*-*tert*-butanesulfinamide **9**, the diastereoisomer of **4** with respect to the chirality at the sulfur atom. Treatment of *S*-*tert*-butanesulfinamide **9** with 3 equivalents of MeMgBr (CH<sub>2</sub>Cl<sub>2</sub>,  $-78$  °C) gave a diastereoisomeric ratio for **10**–**11** of 13 : 1, as monitored by inverse gated <sup>13</sup>C NMR measurements<sup>8</sup> on the crude Grignard products (Scheme 1). The absolute stereochemistry of **10** was unambiguously established by acidic removal of the *S*-*tert*-butanesulfonyl group, giving, after Fmoc-protection, a compound that in all spectroscopical

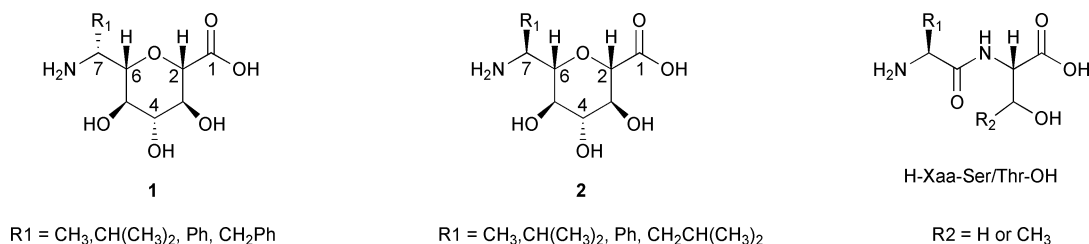
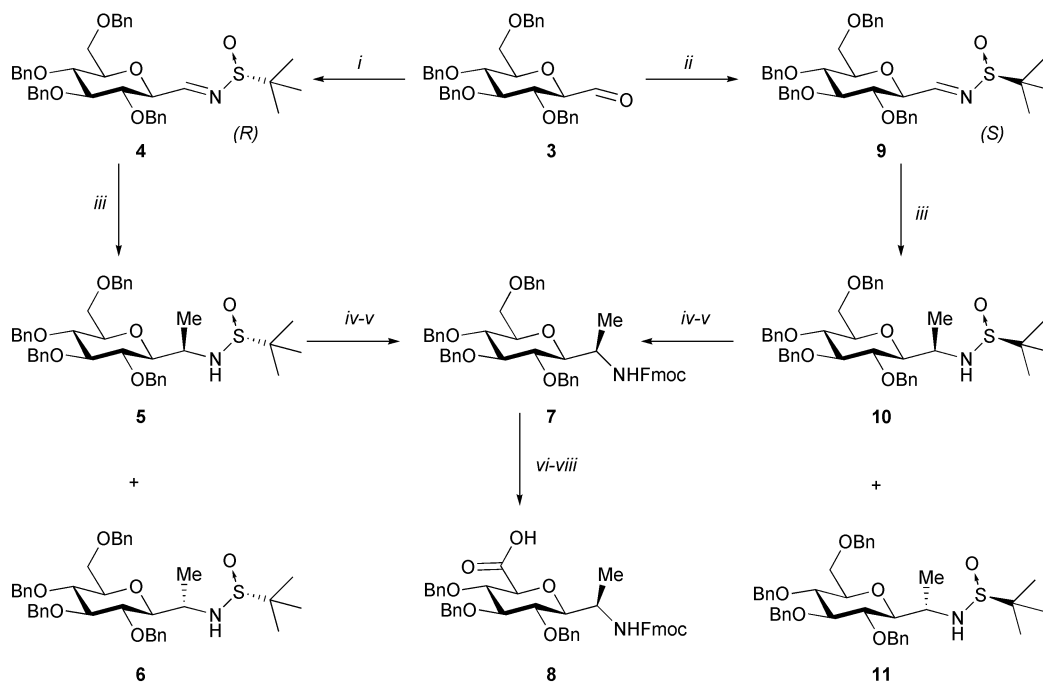


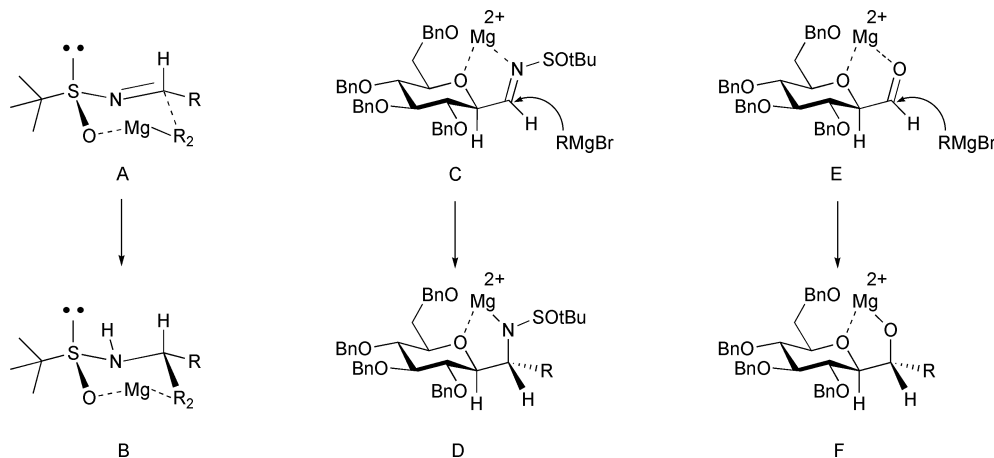
Fig. 1 Alkylated SAAs as Xaa-Ser/Thr mimics.



**Scheme 1** Reagents and conditions: (i) *R*-*tert*-butanesulfonamide,  $\text{Ti}(\text{OiPr})_4$ ,  $\text{CH}_2\text{Cl}_2$ , 70%, (ii) *S*-*tert*-butanesulfonamide,  $\text{Ti}(\text{OiPr})_4$ ,  $\text{CH}_2\text{Cl}_2$ , 72%, (iii)  $\text{MeMgBr}$ ,  $\text{CH}_2\text{Cl}_2$ ,  $-78^\circ\text{C}$ , (iv)  $\text{HCl}$ ,  $\text{MeOH}$ , (v)  $\text{FmocOSu}$ ,  $\text{DIPEA}$ , dioxane,  $\text{CH}_2\text{Cl}_2$ , (from **4**: 71%, from **9**: 75%, three steps), (vi)  $\text{ZnCl}_2$ ,  $\text{HOAc}$ ,  $\text{Ac}_2\text{O}$ , (vii)  $\text{HCl}$ ,  $\text{MeOH}$ , (viii)  $\text{TEMPO}$ ,  $\text{BAIB}$ ,  $\text{CH}_2\text{Cl}_2$ ,  $\text{H}_2\text{O}$  (64%, two steps).

aspects was identical to the previously synthesized **7** ( $\text{R}^1 = \text{CH}_3$ ).<sup>1</sup> From this it follows that the minor product **11** is the *S*-methyl diastereomer of **10** with respect to the newly formed stereocentre. Apparently, *re*-side addition is favored irrespective of the nature of the chiral auxiliary on the imine nitrogen. Changing the solvent system from  $\text{CH}_2\text{Cl}_2$  to THF resulted in a slightly more favored *si*-side addition, and **10** and **11** were formed in equal amounts. This result is the best we obtained in favor of the desired diastereoisomer **11** and we conclude that at least in this system chiral sulfinylimides are not useful intermediates in the construction of L,L-dipeptide isomers.

At the moment we do not have a satisfying model that explains the different product ratios we observe, but it seems likely that the chelation model proposed by Ellman and co-workers in their explanation<sup>7f</sup> of chirality transfer (**A** to **B**, Fig. 2) is counterbalanced by competing chelation of magnesium ions to hetero-atoms on the carbohydrate template. This chelation (for instance **C**, leading to **D**) may occur irrespective of the configuration on the sulfur atom. Whether this reasoning is valid or not, it does present an obvious strategy towards the desired L,L-dipeptide isomers. When one assumes that a formyl-*C*-glycoside chelates just as the sulfinyl imides do with magnesium and that



**Fig. 2** Transition states for Grignard reactions (**A** → **B**: sulfinyl imines in general, **C** → **D**: glycosyl sulfonamide through intramolecular chelation, **E** → **F**: glycosyl aldehyde through intramolecular chelation).



## Crystallography

Compound **12d** was crystallized from mixture of diethyl ether and light petroleum at room temperature as parallelepiped blocks.† A crystal of approximately 0.6 × 0.35 × 0.15 mm was cut from a larger one and analyzed on the Kappa CCD at 293 K using MoK $\alpha_1$  radiation. The full sphere was collected up to  $\theta = 27.5^\circ$ . The data collection and processing were done using the Scalepack software.<sup>13,14</sup> The structure was solved by the direct method and refined using the SHELX97 package.<sup>15</sup> All of the non-H atoms were refined anisotropically while all H atoms were found on the Fourier Difference map and refined isotropically. The absolute configuration of the molecule was deduced knowing the absolute configurations of the four stereocentres at C2, C3, C4 and C5 as occurring in D-glucose, and the Flack parameters.<sup>16</sup>

## Acknowledgements

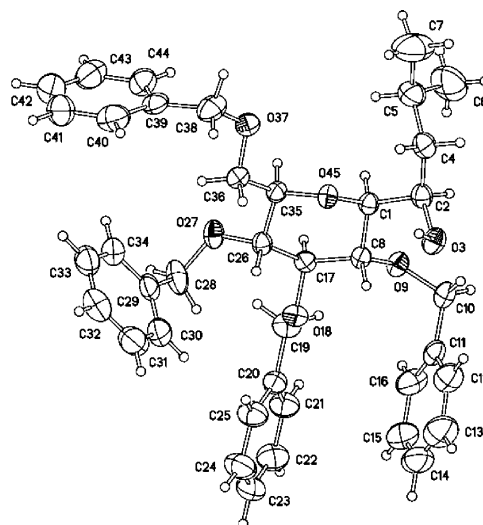
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Empirical formula	C <sub>39</sub> H <sub>46</sub> O <sub>6</sub>
Formula weight	610.76
T[K]	293(2)
$\lambda$ [Å]	0.71073
Crystal system	Orthorhombic
Space group	P 2 <sub>1</sub> 2 <sub>1</sub> 2 <sub>1</sub>
Unit cell dimensions	
a [Å]	6.0520(2)
b [Å]	23.7560(5)
c [Å]	23.9440(5)
V [Å <sup>3</sup> ]	3442.5(2)
Z	4
D <sub>m</sub> [g/cm <sup>3</sup> ]	1.178
Absorption coefficient mm <sup>-1</sup>	0.078
F(000)	1312
Crystal size [mm <sup>3</sup> ]	0.6 × 0.3 × 0.15
$\theta$ range	3 → 27.5
Reflections collected	17437
Independent reflections	7673 [R <sub>int</sub> = 0.0411]
Data / restraints / parameters	7673 / 0 / 566
S	1.025
R [ $>2\sigma(I)$ ]	R1 = 0.0465, wR2 = 0.0858
R indices (all data)	R1 = 0.0747, wR2 = 0.0965
Absolute structure parameter	-0.4(7)

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