

Stereochemical analysis of D-galacto-sulfoxides using (*S*)- α -methoxyphenylacetic acid

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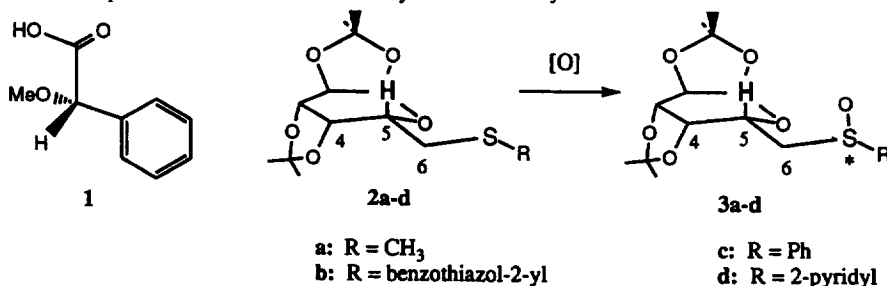
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Abstract: (*S*)- α -Methoxyphenylacetic acid (MPAA) was used as an NMR shift reagent to predict the absolute configuration of some sulfoxide-containing sugars. The correctness of this assignment was confirmed by X-ray crystallography. © 1997 Elsevier Science Ltd

We have recently reported the use of (*S*)-(+)- α -methoxyphenylacetic acid (MPAA) **1** as a chiral NMR shift reagent to determine the enantiomeric purity and absolute configuration of a number of acyclic sulfoxides.^{2a} In this paper, we demonstrate that our methodology can be used to analyze the stereochemistry of some epimeric D-galacto-sulfoxides **3a–d** which are available by oxidation³ of the corresponding sulfides **2a–d**. The determination of the absolute configuration at the sulfinyl center of **3a–d** is of current interest since these compounds are valuable synthetic intermediates whose further elaboration can depend on the stereochemistry of the sulfinyl function.⁴



We began our stereochemical analysis by recording the ¹H (400 MHz) and ¹³C (100.6 MHz) NMR spectrum of the two simplest epimeric sulfoxides **3a1:3a2** (2:3 mixture) in CDCl₃ and examining the effect of adding (*S*)-(+)-MPAA (3 equiv.) on the signals assigned to the α -sulfinyl methyl group. The ¹H resonances (singlets) of both epimers (2.701 ppm, **3a1**; 2.642 ppm, **3a2**) moved downfield due to H-bonding of the carboxyl group to the sulfinyl oxygen but the resonance belonging to **3a2** shifted to a lesser extent (0.017 ppm, **3a1**; 0.012 ppm, **3a2**). The corresponding ¹³C signals (37.662 ppm, **3a1**; 39.736 ppm, **3a2**) moved upfield as anticipated^{2b} but the resonances due to **3a2** were shifted to a greater extent (–1.133 ppm, **3a1**; –1.207 ppm, **3a2**). These results can be rationalized in terms of the typical shielding effects predicted by our Pirkle-type binding model^{2a} and on this basis one would assign the (*R*)-configuration to **3a1** and the (*S*)-configuration to **3a2** (Figure 1).

Support for these configurational assignments was obtained by examining the effect of (*S*)-MPAA addition on the ¹H resonances due to the α -sulfinyl methylene group of the two epimers, **3a1** and **3a2** in combination with conformational analysis. As shown in Figure 2, addition of (*S*)-MPAA (3 equiv.) caused the upfield half of the C-6 AB quartet (H_B) of **3a2** to shift downfield significantly (0.18 ppm)

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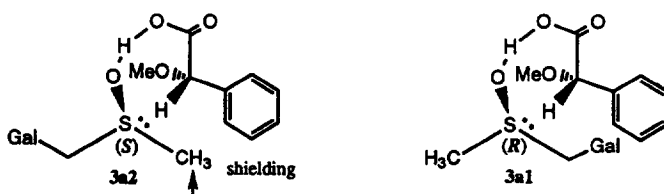


Figure 1. Binding model for the interaction of (*S*)-MPAA with the two epimers of **3a**.

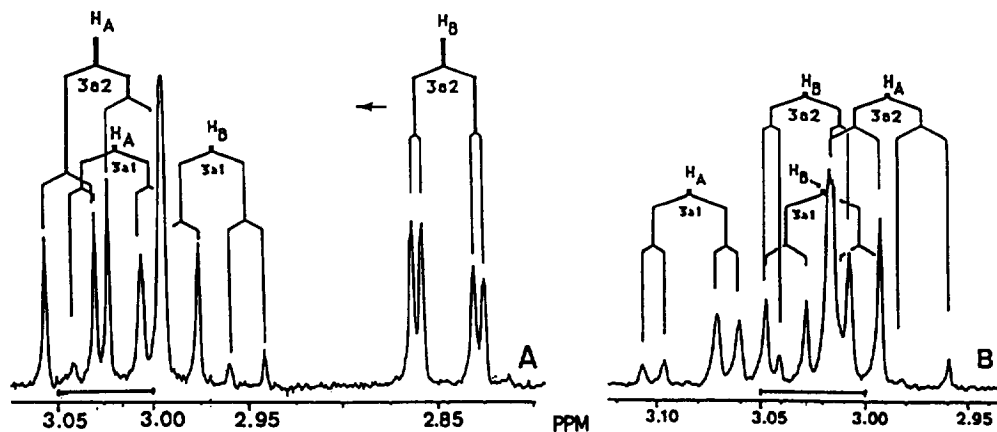


Figure 2. ^1H NMR (400 MHz) resonances due to the α -sulfinyl methylene group of **3a1** and **3a2** (A) before and (B) after addition of (*S*)-(+)-MPAA (3 equiv.).

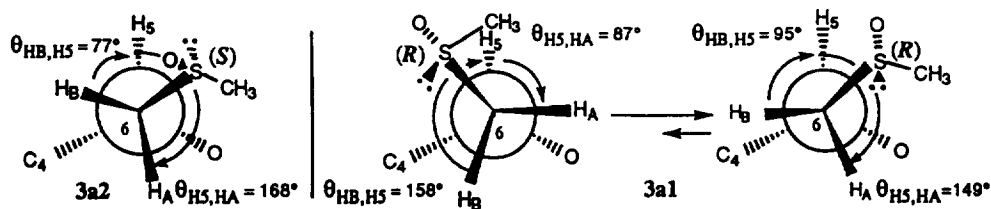


Figure 3. Newman Projections of energy-minimized conformations calculated for **3a1** and **3a2**.

while the corresponding signals due to H_A of **3a2** and the entire AB quartet for **3a1** are only slightly affected.

In order to interpret this data, energy-minimized⁵ conformation(s) for the side-chain of each epimer were calculated. As shown in Figure 3, a single preferred conformation was identified for **3a2** while a pair of conformations, separated in energy by 0.6 kcal/mole was found for **3a1**. The calculated⁶ values for $^3\text{J}_{5,6}$ of each epimer are in qualitative agreement with the observed values for $^3\text{J}_{5,6}$ obtained from the ^1H NMR spectrum displayed in Figure 2A: (calculated $^3\text{J}_{5,6}$ **3a2**: 9.4 Hz (H_A), 0.7 Hz (H_B) and observed $^3\text{J}_{5,6}$ **3a2**: 10.6 Hz (H_A), 2.1 Hz (H_B); calculated $^3\text{J}_{5,6}$ **3a1**: 5.4 Hz (H_A), 2.6 Hz (H_B) and observed $^3\text{J}_{5,6}$ **3a1**: 7.2 Hz (H_A), 4.8 Hz (H_B)).

Assuming that the conformations of the side-chain do not change upon addition of (*S*)-MPAA,⁷ the dramatically different effect of this shift reagent on the ^1H chemical shifts of the α -sulfinyl methylene hydrogens of each epimer can now be explained. That is, H-bonding of the sulfanyl oxygen in **3a2** strongly deshields⁸ the proximal hydrogen, namely H_B . In contrast, the sulfanyl oxygen of **3a1** is pointing away from H_A and H_B in both conformations and thus the ^1H signals due to both hydrogens are affected only very slightly and to a similar extent (Figure 2).

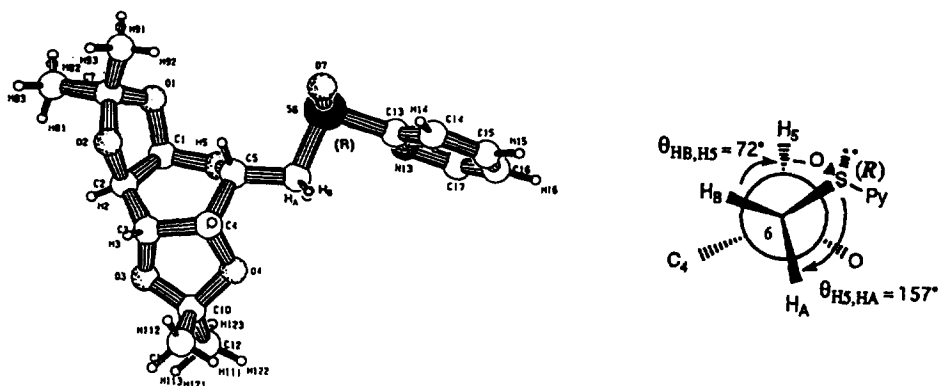


Figure 4. X-ray structure of the more polar epimer of **3d** and a Newman projection along the C-5,6 bond.

A similar dichotomy in sulfinyl side-chain conformations was detected for the two epimeric 2-pyridyl sulfoxides, **3d**. These compounds could be separated by silica gel column chromatography and were examined separately by ^1H NMR with the aid of (*S*)-MPAA. The more polar epimer of **3d** possesses disparate $^3J_{5,6}$ values (10.5 Hz (H_A), 2.5 Hz (H_B)=*anti* conformation) and upon addition of the chiral shift reagent, a substantial downfield shift of 0.12 ppm was observed for the upfield half of the C-6 AB quartet (H_B) while the downfield half (H_A) was essentially unaffected. In contrast, the less polar epimer of **3d** exhibits more similar $^3J_{5,6}$ values (4.6 Hz, 7.8 Hz), and both halves of the C-6 AB quartet move downfield very slightly (ca. 0.02 ppm) upon complexation as anticipated. These phenomena clearly link the two sets of epimers, **3a** and **3d**, and on this basis one would assign the (*R*)-configuration (priority change) to the sulfinyl group of the more polar epimer of **3d** and the (*S*)-configuration to its diastereomer. The former compound was subsequently crystallized and our configurational assignments were validated by X-ray crystallography (Figure 4). (Full details of the X-ray work will be published elsewhere (*Journal of Carbohydrate Chemistry*)). It is interesting to note that the conformation of the pendant chain shown in this X-ray structure is similar to that calculated for **3a2** as shown in Figure 3.

Stereochemical analyses of other related compounds in this series are planned.

Acknowledgements

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