

STUDIES ON MORPHOLINOSPHINGOLIPIDS: POTENT INHIBITORS OF GLUCOSYLCERAMIDE SYNTHASE

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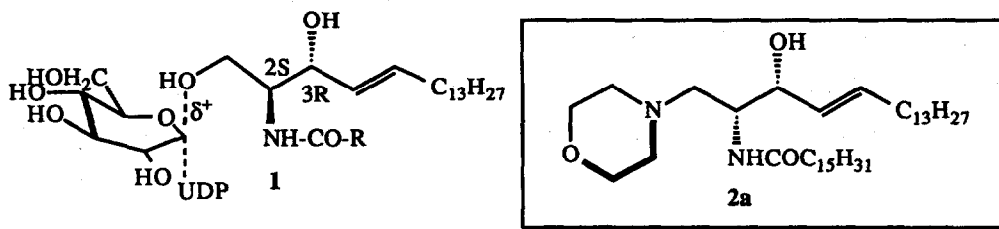
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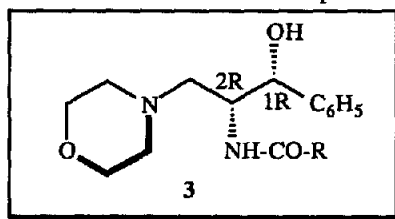
Abstract: Synthetic 1-morpholino-1-deoxyceramides were designed to inhibit glucosylceramide synthase. The most potent inhibitor **2a** possesses the unnatural R,R-configuration of D-threo-sphingosine.

Eucaryotic glycosphingolipids (GSLs) affect membrane physical properties, cell-cell and cell-matrix interactions, adhesiveness, cellular immune responses, and differentiation.¹ GSLs have also been implicated in cancer cell metabolism.² For example, malignant cells display marked abnormalities in their relative proportions of glycolipids,³ and produce GSLs with novel linkage and sugar specificities.⁴ Tumor cells also synthesize and shed excessive levels of GSLs which suppress lymphocyte responses in the host's immune system.



GSL biosynthesis begins with the coupling of UDP-glucose to C1 of an N-acylsphingosine (ceramide). The reaction is catalyzed by glucosylceramide synthase (GlcCer synthase; ceramide:UDP-glucose glucosyltransferase; EC 2.4.1.80), probably via a transition structure like **1**.⁵ The enzyme plays a pivotal role in GSL biosynthesis and represents a promising cancer chemotherapy target, since inhibitors can retard or arrest tumor growth.⁶ Here we report studies defining 1-morpholino-1-deoxyceramides such as **2a** as potent GlcCer synthase inhibitors.

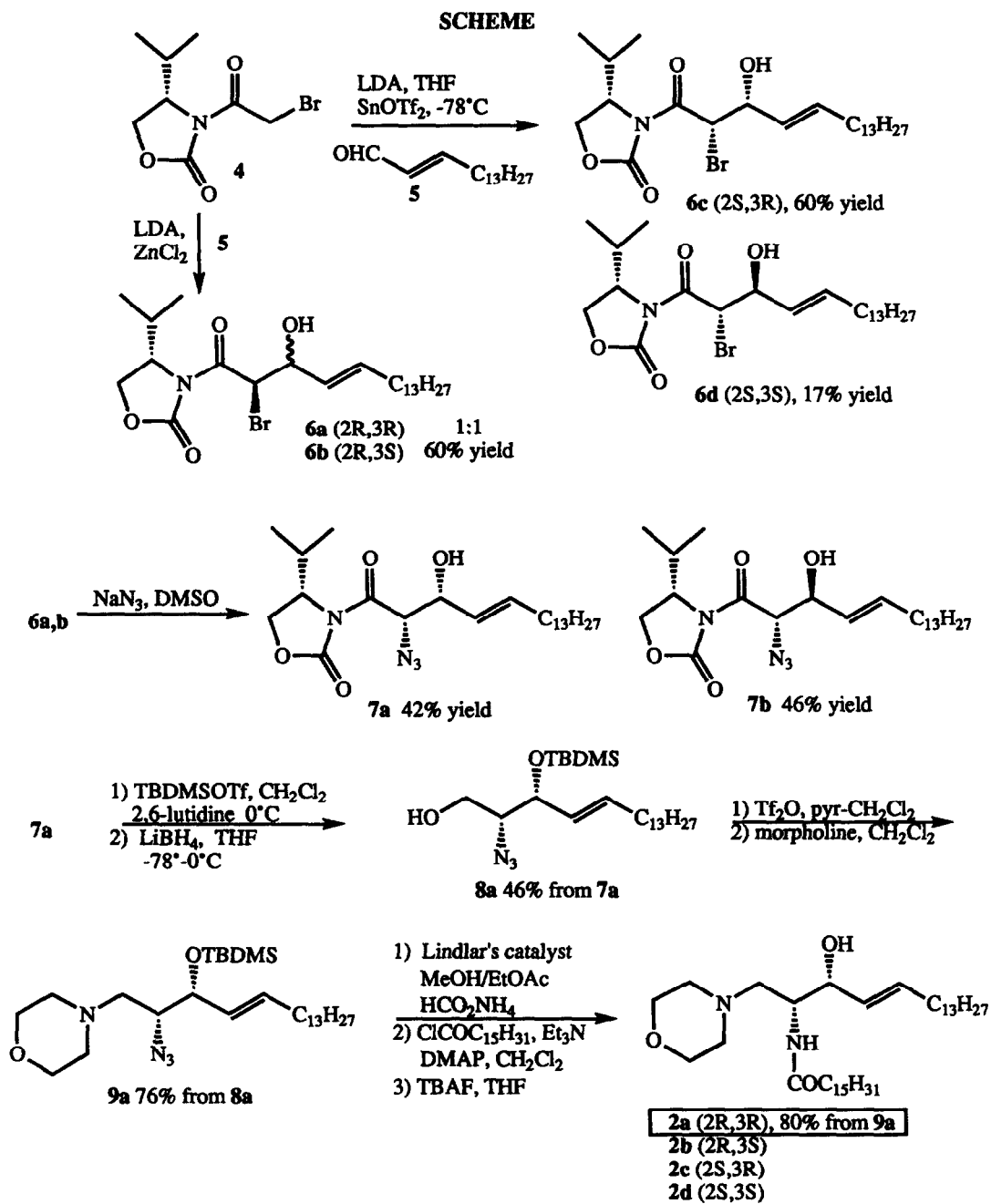
Inhibitor design was based on the observation that *D-threo*-1-phenyl-2-decanoylamino-3-morpholino-1-propanol (PDMP) **3** is an active competitive inhibitor of GlcCer synthase.⁷ Presuming that the morpholine ring mimics the cationic charge of the GlcCer synthase transition state **1**, the *syn*-stereochemistry in **3** suggests that the bioactive (1*R*,2*R*)-PDMP stereoisomer is at variance with the *D-erythro*- or *anti*-configuration of naturally-occurring GSLs shown in (2*S*,3*R*)-**1**. This stereochemical difference was a central concern in our plan to design more potent, sphingosine-based GlcCer synthase inhibitors containing glucopyranose analogs having ring conformations related to **1**.⁸ To probe this stereochemical issue, we developed syntheses of all four isomeric 1-morpholino-1-deoxyceramides **2a-d**, as shown in the Scheme.



Our synthetic approach was based on the method of Evans *et al.* for the enantioselective aldol condensation of oxazolidinone **4** with unsaturated aldehyde **5**.⁹ According to Abdel-Magid *et al.*,¹⁰ the observed stereochemistry of condensation is influenced by chelation to the oxazolidinone oxygen in the transition state. In fact, condensation of **5** with the tin(II) enolate of **4** (prepared as shown in the Scheme) gave mostly **6c**, whereas the Zn(II) enolate of **4** gave an inseparable mixture of **6a** and **6b**. Both reactions gave minor amounts of the (2*S*,3*S*) isomer **6d**, which could be separated by flash column chromatography.

The mixture of bromohydrins **6a** and **6b** could not be resolved, but was separable after conversion to the corresponding azides **7a** and **7b**, following the procedure of Nicolaou *et al.*¹¹ Alcohol **7a** was then protected as its *t*-butyldimethylsilyl ether, and the chiral auxiliary was reductively removed to afford sphingol ether **8a**. The primary alcohol in **8a** was next activated as its triflate, and displacement with morpholine afforded aminoether **9a**. Reduction of the azide group in **9a** was effected by transfer hydrogenation using ammonium formate as the hydrogen source. Subsequent N-acylation and desilylation led to the final product **2a** in good overall yield.¹² This constitutes the first use of metal-dependent aldol condensations in sphingosine synthesis, and the first synthesis of 1-azaceramides. Diastereomers **2b-d** were synthesized in like fashion from aldol adducts **6b-d**.

Morpholinoceramides **2a-d** were evaluated as inhibitors of GlcCer synthase from Madin-Darby canine kidney cell homogenates, using octanoyl sphingosine as glucose acceptor (thermostatted ultrasonic bath, triplicate assays).¹³ Diastereomer **2a** was clearly the most powerful inhibitor of GlcCer synthase (73% inhibition at 5 μ M), and was significantly more potent than PDMP **3** (16-20% inhibition at 5 μ M) or isomers **2b-d** (5-20% at 5 μ M).



These findings set the stage for the refinement of GlcCer synthase transition structure mimics, including bi-substrate analogs and other active inhibitors based on the corresponding glucoamidrazone and glucoamidoxime derivatives⁸ of sphingosine.

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