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Research paper

sp²-Iminosugar glycolipids as inhibitors of lipopolysaccharidemediated human dendritic cell activation in vitro and of acute inflammation in mice in vivo



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ABSTRACT

Glycolipid mimetics consisting of a bicyclic polyhydroxypiperidine-cyclic carbamate core and a pseudoanomeric hydrophobic tail, termed sp²-iminosugar glycolipids (sp²-IGLs), target microglia during neuroinflammatory processes. Here we have synthesized and investigated new variants of sp²-IGLs for their ability to suppress the activation of human monocyte-derived dendritic cells (DCs) by lipopoly-saccharide (LPS) signaling through Toll-like receptor 4. We report that the best lead was (1*R*)-1-dodecylsulfonyl-5*N*,6*O*-oxomethylidenenojirimycin (DSO₂-ONJ), able to inhibit LPS-induced TNF α production and maturation of DCs. Immunovisualization experiments, using a mannoside glycolipid conjugate (MGC) that also suppress LPS-mediated DC activation as control, evidenced a distinct mode of action for the sp²-IGLs: unlike MGCs, DSO₂-ONJ did not elicit internalization of the LPS co-receptor CD14 or induce its co-localization with the Toll-like receptor 4. In a mouse model of LPS-induced acute inflammatory interleukin-6. The ensemble of the data highlights sp²-IGLs as a promising new class of molecules against inflammation by interfering in Toll-like receptor intracellular signaling.

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1. Introduction

sp²-Iminosugars are an original family of sugar look-alikes possessing a pseudoamide-type nitrogen atom with a high sp²hybridation character at the position of the endocyclic oxygen [1–3], with the unique property of mimicking not only the structure and function of the natural counterparts, as classical iminosugars do, but also their chemistry [4]. Most significantly, sp²iminosugars can engage in glycosylation reactions, affording stable glycoconjugate analogs capable of agonizing or antagonizing the biological processes in which the natural counterparts are involved. Examples that illustrate the opportunities that sp²-iminosugar

https://doi.org/10.1016/j.ejmech.2019.02.078 0223-5234/© 2019 Elsevier Masson SAS. All rights reserved. conjugates offer in biology and medicine are the design of pharmacological chaperones targeting lysosomal glycosidases [5], the synthesis of multivalent constructs to explore multivalent enzyme inhibition [6] or the preparation of Tn antigen glycopeptide mimics [7].

Pursuing our interest in the potential of sp^2 -iminosugars as monosaccharide surrogates for neoglycoconjugate drug development, we recently turned our attention to glycolipids as immunoactive agents for clinical and mechanistic applications, a very active research area in medicinal chemistry. Studies report that glycolipids exert antiinflammatory and antiproliferative activity, as well as beneficial effects on lipid metabolism [8]. The chemical synthesis of glycolipid analogs is often challenging, however, especially when it comes to producing pure anomeric forms of the sugar head group. Remarkably, sp^2 -iminosugars benefit from an exacerbated anomeric effect, which warrants total α -stereoselectivity during *O*-, *S* or *N*- glycosylation steps and imparts

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chemical and enzymatic stability to the resulting glycosides. Indeed, some sp²-iminosugar glycolipids (sp²-IGLs) have been already prepared and found to exhibit a range of bioactivities including anti-proliferative [9], anti-metastatic [10,11], antiparasitic [12] and anti-neuroinflammatory activity [13,14]. The sp²-IGL representatives (1*R*,S*R*)1-dodecylsulfinyl-5*N*,60-oxomethylidenenojirimycin (R-DSO-ONJ) and (1R)-1-dodecylsulfonyl-5N.6O-oxomethylidenenojirimycin (DSO₂-ONI) were particularly efficient at inhibiting the lipopolysaccharide (LPS)-induced inflammatory signaling in mouse Bv.2 microglial cells by decreasing inducible nitric oxide synthase (iNOS) and pro-inflammatory cytokine production and simultaneously upregulating production of the anti-inflammatory cytokine IL-10. The latter compound was able to significantly reduce reactive gliosis in retinal explants of the diabetic *db/db* mice model that recapitulates the progression of diabetic retinopathy [14,15]. Although initially the protein target was hypothesized to be an endoplasmic reticulum α -glucosidase, the sp²-IGLs were found not to impact glycoprotein biosynthesis, which is consistent with their lack of toxicity in healthy cells [16]. Instead, they seemed to interfere with the mitogen activated protein kinase (MAPK) cascade, at the crossroad between immunity and cancer, offering new therapeutic opportunities.

Dendritic cells (DCs), present in most tissues and body liquids, are key players in the initiation and polarization of the immune response [17]. The investigation of key indicators of the inflammatory response in DCs upon treatment with sp²-IGLs was therefore mandatory to validate the above notion. These immune sentinels are highly sensitive to innate immune stimuli and mature into professional antigen-presenting cells producing inflammatory cytokines, such as TNFa, and upregulating the T cell co-stimulators CD83 and CD86 at the cell surface. Mature DCs are highly efficient in priming the T cell-mediated adaptive immune response [17]. A particularly efficient DC stimulator is LPS, an important toxin found in the outer membrane of Gram-negative bacteria involved in the development of septic shock, characterized by the hyperproduction of inflammatory cytokines and organ damage [18]. LPS-elicited acute inflammatory responses are mediated by the signaling receptor Toll-like receptor (TLR)-4. The signaling cascade is activated by the successive interaction of LPS with the lipopolysaccharide binding protein, the cluster differentiation antigen CD14, and the myeloid differentiation protein (MD-2) that binds LPS and presents it in a monomeric form to TLR-4 [19]. TLR-4 undergoes dimerization and recruits downstream adaptor proteins, which initiates an intracellular signaling cascade that leads to nuclear translocation of transcription factors and the biosynthesis of pro-inflammatory cytokines and interferons [20,21]. Therefore, therapeutic interventions to reduce TLR-4 signaling in response to LPS are of clinical importance.

With the objective to gather information on the mode of action of sp²-IGLs and on the structural features that determine their antiinflammatory properties, we have now generated a series of derivatives 1-8 (Fig. 1) that keep the dodecylsulfinyl or dodecylsulfonyl lipidic aglycone moiety but differ in the hydroxylation pattern (D-gluco, D-galacto or D-manno) of the sp²-iminosugar aglycone, and studied their capacity to inhibit the activation of human DCs by LPS. The sulfone derivative DSO₂-ONJ (compound **3**), having a hydroxylation profile of structural complementarity to α-D-glucopyranosides, emerged as the most active member in reducing TNF α production and downregulating the DC maturation markers CD83 and CD86. Interestingly, co-localization studies support that sp²-IGLs act by inhibition of intracellular signaling rather than by modifying LPS receptor cell-surface localization. This behavior is sharply different from that reported for other neoglycolipids such as the mannoside glycolipid conjugate (MGC) 9 (Fig. 1), which suppress TLR-4 signaling in human DCs by targeting



Fig. 1. Structures of sp^2 -IGL derivatives (1–8) and the mannoside glycolipid conjugate (MGC) 9 evaluated in this study.

early membrane steps of the CD14—TLR-4 signaling membrane complex [22–24]. In a mouse model of endotoxin shock, sp²-IGL **3** reduced the production of the inflammatory interleukin 6 (IL-6), supporting its therapeutic potential.

2. Results

2.1. Design rational and synthesis

Compounds **1–8** were chosen for the initial cell toxicity and anti-inflammatory activity-screening, primarily because sp²-IGLs bearing sulfur anomeric functional groups and twelve-carbon aliphatic aglycones were previously identified as privileged structures in anti-cancer/anti-inflammatory tests in different settings [13,14,16]. The sulfinyl and sulfonyl derivatives, differing in their oxidation state, have been suggested [14] to emulate the phosphate group in anti-inflammatory phosphatidylinositol analogs (PIAs) and alkyl ether lipids (e.g., perifosine) for which a mechanism of action involving p38 MAPK binding has been evidenced [25]. Dodecyl-sulfinyl and -sulfonyl conjugates were therefore considered with sp²-iminosugar moieties that mimic D-gluco (1–3), D- galacto (**4**–**6**) and D-manno (**7** and **8**) glycolipids in order to assess the effect of the glycone configuration in the anti-inflammatory activity.

Compounds 1-3 were prepared following previously reported procedures [12.14]. The chemical syntheses of the new derivatives **4–8** follow a general scheme involving (i) steroselective α -S-glvcosidation of the corresponding peracetylated sp^2 -iminosugar (10 or **14**) with dodecanethiol and (ii) partial or full oxidation of the thioglycoside intermediate (11 or 15) with *m*-chloroperbezoic acid (m-CPBA) to afford the target sulfoxide and sulfone sp²-IGLs, respectively (Schemes 1 and 2). In the case of the galacto-configured sulfoxides, both the (S_S) and the (R_S) diastereomers **12** and **13** were formed in close to 1:1 ratio and could be separated by column chromatography before deacetylation to give **4** and **5**, respectively. Their absolute stereochemistry was assigned based on their ¹H NMR spectra. Notably, the H-5 resonance was deshielded by 0.67 ppm in **12** (δ 4.96 ppm) compared with **13** (δ 4.29 ppm). Literature data on alkyl α-D-glycosyl sulfoxides support then the (S_S) assignment for the first one, where H-5 and the sulfoxide oxygen would be located in close proximity in the more favourable exoanomeric-type conformation, that is, with C-2 in the ring and the exocyclic methylene carbon in anti-disposition [26]. The lower magnetic nonequivalence $(\Delta \delta)$ of the methylene protons vicinal to the chiral sulfur atom (SOCH₂) for 12 (57 Hz) as compared with 5 (81 Hz) is also in agreement with the NMR properties reported in the literature for (S_S) and the (R_S) alkyl α -D-glycosyl sulfoxides, respectively [27]. The partial oxidation of the *manno*-configured thioglycoside **15** afforded a major sulfoxide product (**16**) that was obtained in pure form after column chromatography and assigned the (S_S) configuration based on the $\Delta\delta$ value for the SOCH₂ protons (48 Hz) [27]. The (R_S) diastereomer was detected in the reaction mixture as a trace compound but could not be isolated in pure form. Finally, deacetylation of 16 by treatment with sodium methylate afforded the fully unprotected target sp²-IGL **7**.

2.2. Toxicity of sp²-IGLs on DCs

Human monocyte-derived dendritic cells (DCs) were differentiated from monocytes purified from the blood of different donors. The cytotoxicity of the sp²-IGL derivatives was evaluated by incubation of the individual components **1–8** at 25 and 50 μ M concentrations for 24 h, followed by exposure to 7-aminoactinomycin D, a DNA-intercalating fluorescent dye that only penetrates the cell upon cell death. As control, DMSO solvent was added. Assays were performed in parallel in the absence and presence of 100 ng mL⁻¹ LPS. In its absence, the mean cell viability did not fall below an 80%



Scheme 1. Synthesis of the D-galacto-configured sp²-IGLs 4–6.



Scheme 2. Synthesis of the D-manno-configured sp²-IGLs **7** and **8**.

threshold at 25 μ M or a 70% limit at 50 μ M upon treatment with the sp²-IGLs (Fig. 2a and b). However, in the presence of LPS, cell viability was reduced and fell below 70% for compounds **5** and **6** at 50 μ M (Fig. 2a and b). DMSO had no significant impact on cell viability irrespective of LPS. To be in a relatively safe window, we settled the dose limit to 50 μ M for future experimentation with the sp²-IGLs.

2.3. sp²-IGLs inhibit LPS activation of human DCs

We next examined the capacity of the selected sp²-IGLs representatives to inhibit the activation of DCs by LPS. By stimulating TLR-4-signaling, LPS is a potent activator of DCs leading to the production of $TNF\alpha$ and the upregulation of the T cell costimulators CD83 and CD86. A 24h exposure time to LPS in the absence or presence of 25 μ M and 50 μ M of sp²-IGLs was established to warrant significant expression of endogenous CD83 and CD86 [28,29]. LPS induced the production of TNFa, as determined by ELISA, but there was a significant reduction of $TNF\alpha$ by compound **3** at 25 μ M and by compounds **3**–**8** at 50 μ M (Fig. 3a and b). The expression of CD83 and CD86 on the cell surface was subsequently assessed by fluorescence-activated flow cytometry (FACS). At 25 µM, compounds 1 and 4 significantly reduced CD83 levels (Fig. 3c). As for CD86, only compound **3** was active at both $25 \,\mu\text{M}$ and 50 μ M (Fig. 3d). These results showed that most sp²-IGLs inhibited to some extent LPS-mediated DC maturation, compound **3** being the most active. Subsequent analysis of its TNFα production dose-dependent inhibition on LPS-activated DCs (Fig. 4a) allowed us to establish the dose-response curve for compound 3 and its IC_{50} at $20 \pm 10 \,\mu\text{M}$ (Fig. 4b). Its efficient downregulatory effect of CD86 expression was also confirmed (Fig. 4c), while there was a significant reduction of CD83 expression only at 50 µM (Fig. 4d). Taken together, the sp²-IGL **3** displayed a significant anti-LPS activity using human DCs as test cells.

2.4. sp²-IGL 3 remains active post-LPS exposure

We previously showed that the mannoside glycolipid (MGC) **9** counteracted LPS activation of DCs at similar concentrations as those now encountered for the sp²-IGL **3** [22]. In order to explore whether or not the two glycolipid mimics behave through similar mechanisms, a direct comparison of compound **3** with MGC **9** for their capacity to antagonize LPS in its stimulatory activity on DCs was conducted. It was observed that MGC **9** impaired TNF α



Fig. 2. Effect of sp²-IGLs on the viability of human DCs. The different sp²-IGL compounds were used at (a) 25 µM and (b) 50 µM. Solvent controls were performed with DMSO at 0.5 and 1%. Cell viability in the absence or presence of LPS (100 ng mL⁻¹) was measured after 24 h by 7-aminoactinomycin D (7-AAD) staining, followed by flow cytometry analysis. Each point corresponds to a different monocyte blood donor.



Fig. 3. sp²-IGLs are anti-inflammatory for LPS-activated DCs. Cells were stimulated with LPS (100 ng mL⁻¹) in the absence or presence of sp²-IGLs for 24 h (a, 25 μ M; b, 50 μ M) Effect on the pro-inflammatory cytokine TNF- α released in the media. (c, d) Effect on the relative expression of CD83 and CD86 cell surface maturation markers. Non-treated cells (n.t.; control) or treated with the solvent (DMSO) are indicated. Data correspond to at least 3 cell donors/independent assays. Bars represent means \pm SEM.

production more strongly than compound **3** when added at the same time as LPS or 0.5 h after LPS exposure (Fig. 5a and b).

During the first hour of LPS stimulation of macrophages there is an initial delay in the production of biologically active TNF α [30]. This is followed by a rapid increase in TNF α accumulation, reaching a peak by 3 h, followed by a rapid decline, and reaching a plateau in TNF α accumulation in the supernatant by 6 h [31]. Thus, it was of interest to examine sp²-IGL **3** versus MGC **9** regulation of TNF α production during this time frame. When the compounds were added 2 h, 4 h or 6 h after LPS, it was apparent that while inhibition by compound **3** remained essentially constant, there was a clear loss of MGC **9** activity (Fig. 5a and b). Analysis of CD83 and CD86 markers confirmed these properties (SI, Fig. S1). In summation, MGC **9** was found to be a more potent anti-inflammatory agent when added conjointly with LPS, but compound **3** displayed a more efficient post-LPS inhibitory activity.

2.5. The mode of operation of sp^2 -IGLs and **MGC 9** are different

The above findings suggest that the mode of action differs



Fig. 4. Anti-inflammatory effects of sp²-IGL **3** in DCs. Cells were stimulated with LPS (100 ng mL⁻¹) in the absence or presence of **3**, as in Fig. 3. After 24 h, analyses were performed for (a) TNF-α released in the media, (b) IC₅₀ on TNF-α production, and (c, d) cell surface maturation markers CD86 and CD83 relative expression. Non-treated cells (n.t.) are indicated. Data correspond to at least 3 cell donors/independent assays. Bars represent means ± SEM.



Fig. 5. Delayed addition of compound **3** conserves inhibitory activity. DCs were left untreated (n.t.) or treated with LPS, either alone or with (a) compound **3** or (b) MGC **9**, added at the indicated times. TNF-*α* production was analyzed after 24 h. Data correspond to the mean (±SD) of n > 3 independent assays (each dot corresponds to an independent assay).

between the MGC and sp²-IGL LPS-signaling inhibitor families. We have previously shown that MGC **9** at 100 μ M elicited CD14 internalization from the cell surface and triggered the co-localization of CD14 with the TLR-4, the two LPS co-receptors, thereby preventing their availability to interact with the LPS at the cell membrane leading to the inflammatory response [22]. To investigate whether sp²-IGLs affected receptor co-localization, DCs were left untreated or exposed for 30 min to 100 μ M of MGC **9** or 25 μ M and 50 μ M of the *D*-gluco-configured sp²-IGLs **1**, **2** and **3**, followed by flow cytometry analysis of CD14 cell surface expression. As expected, MGC **9** internalized CD14 from the cell surface, but none of the sp²-IGLs reduced the CD14 concentration at the cell-surface (Fig. 6a). To assess CD14—TLR-4 co-localization and the potential effect of variations in the monosaccharide mimic configurational pattern, the cells were next labeled with anti-CD14 and anti-TLR-4 antibodies

and treated with the more active *D*-gluco compounds, namely the sulfoxide **1** and the sulfone **3**, or the **3**-epimers **6** and **8**, and visualized by confocal microscopy as described.¹⁸ In comparison to untreated cells, MGC **9** provoked the co-localization of the two LPS co-receptors. However, none of the sp²-IGLs, irrespective of the glycone configuration (**1** and **3** for *D*-gluco, **6** for *D*-galacto and **8** for *D*-manno) induced CD14—TLR-4 proximity (Fig. 6b). These data show that sp²-IGLs have no impact on cellular LPS-receptor co-localization and support the notion that their inhibitory mechanism is different from that of MGCs.

2.6. sp^2 -Iminosugar glycolipid **3** inhibits acute inflammation in mice

We next tested if sp²-IGL **3** can function as an anti-inflammatory



Fig. 6. Effect of sp²-IGLs on DCs membrane proteins. (a) Cell surface expression of CD14 analyzed by flow cytometry, after 30 min treatment with MGC **9** (100 μM) or compounds **1**, **2**, **3** (25 and 50 μM; light and deep blue dots, respectively). (b) Pearson coefficient of colocalization of CD14 and TLR-4 in permeabilized DCs after 30 min treatment with MGC **9** (100 μM) or compounds **1**, **3**, **6**, **8** (50 μM). Each dot corresponds to a field of 15–20 cells visualized by confocal microscopy. Data correspond to two independent assays. Dots above the grey line (area noted as yes) indicate an increased tendency for colocalization. (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)

agent in a mouse model of LPS-induced inflammation. For practical reasons, the response of the proinflammatory interleukin IL-6, which reaches much higher concentrations in serum after LPS administrations as compared with TNF α or other cytokines/chemo-kines, was determined in this assay [32]. First, toxicity was assessed by comparing the weight change of 2 female and 2 male C57BL/6J mice after repeated intraperitoneal administration of sp²-IGL **3** or control solvent for 4 successive days. There was no weight loss in either female or male mice, demonstrating good tolerance of the compound at this dose (Fig. 7a and b). Next, mice received intraperitoneally either a single dose of 50 mg/kg of sp²-IGL **3** or solvent control 1 h prior to a single peritoneal injection of 0.5 mg/kg of LPS. Two hours later, the mice were bled and serum prepared for IL-6

measures by ELISA. As expected [33], the proinflammatory cytokine IL-6 rose to high levels in the blood in response to LPS. However, the administration of sp^2 -IGL **3** led to a significant reduction in IL-6 levels. These findings unequivocally show that sp^2 -IGL **3** reduces the inflammatory response to LPS stimulation in a mouse model.

3. Discussion

With the aim of exploring alternative therapeutic approaches against inflammation, here we provide a proof-of-principle study, which lays the foundation for novel strategies based on the use of sp²-iminosugar glycolipids to modulate the intracellular events



Fig. 7. Compound **3** is active in a mouse model of acute LPS-induced inflammation. (a, b) The weight curves of male and female C57BL/6J mice (relative to their weight on day 0) treated with (a) control PBS/DMSO or (b) compound **3** (30 mg/kg) administered i.p. on day 0, 1, 2, 3, shows the absence of toxic effect of **3**. (c) Mice were treated with solvent (PBS/DMSO 2:1) or compound **3** (in PBS/DMSO; 50 mg/kg) administered i.p., 1 h before LPS (0.5 mg/kg) challenge. Histograms show the IL-6 production in serum 2 h after LPS injection. Bars are mean values \pm SD of one typical experiment out of 2. * p \leq 0.05.

following an external inflammatory stimulus. The ensemble of data on the expression of LPS-elicited human dendritic cell maturation markers upon treatment with the synthesized sp²-iminosugar glycolipids evidenced the critical effect of the nature of the functional group linking the glycone portion and the lipid chain in the biological activity. The efficiency of the inflammatory response inhibition is further modulated by the configurational pattern of the sugar-like motif. Thus, sp²-IGLs with a hydroxylation profile analogous to that of α -D-glucose were more active than the corresponding epimers with α -D-galactose or α -D-mannose configuration. Nevertheless, the changes in the relative orientation of the hydroxyl groups in the sugar-like portion did not affect drastically their potential as anti-inflammatory agents. Sulfone 3 was identified as the most active sp²-IGL in the series, exhibiting low toxicity in vitro up to 50 μ M and the capability to inhibit TNF α production in a dose-dependent manner, with an IC₅₀ of 20 µM. Time-course addition experiments and determination of the CD14 and the TLR-4 LPS-receptor localization revealed marked differences between the sp^2 -IGLs and a control glycolipid, MGC **9**: whereas the latter acts at the cell membrane level, preventing binding of LPS to the CD14 and TLR-4 co-receptors, the sp²-IGLs do not; therefore, they interfere with TLR-4 intracellular signaling. This is consistent with the reported data suggesting that compound 3 interacts with p38a MAPK in microglia [14]. The promise of small molecule kinase inhibitors for patient benefit in the context of autoimmune and inflammatory processes is well recognized [34]. Finally, our results provide evidence for the efficacy of compound **3** in an *in vivo* model of LPS-mediated inflammation and incite further work on the potential of sp²-IGLs in anti-inflammatory therapies.

4. Experimental

4.1. General methods

Reagents and solvents were purchased from commercial sources and used without further purification. Optical rotations were measured with a JASCO P-2000 polarimeter, using a sodium lamp $(\lambda = 589 \text{ nm})$ at 22 °C in 1 cm or 1 dm tubes. NMR experiments were performed at 300 (75.5), 400 (100.6) and 500 (125.7) MHz for ¹H (¹³C, respectively). 2-D COSY and HMQC experiments were carried out to assist on signal assignment. In the FABMS spectra, the primary beam consisted of Xe atoms with maximun energy of 8 keV. The samples were dissolved in *m*-nitrobenzyl alcohol or thioglycerol as the matrices and the positive ions were separated and accelerated over a potential of 7 keV. NaI was added as cationizing agent. For ESI mass spectra, 0.1 pm sample concentrations were used, the mobile phase consisting of 50% aq MeCN at 0.1 mL min⁻¹. Thin-layer chromatography was performed on precoated TLC plates, silica gel 30F-245, with visualization by UV light and by carring with 10% H₂SO₄ or 0.2% w/v cerium (IV) sulphate-5% ammonium molybdate in 2 MH₂SO₄ or 0.1% ninhydrin in EtOH. Column chromatography was performed on Chromagel (silice 60 AC.C 70-200 µm). Deacetylation reactions were carried out by using Zemplén procedure, e.i., addition of NaOMe (0.1 equiv/Ac mol) in MeOH at room temperature, followed by neutralization with solid CO₂, evaporation of the solvent and purification by column chromatography. All compounds were purified to \geq 95% purity as determined by elemental microanalysis results obtained on a CHNSTruSpect Micro elemental analyzer (Instituto de Investigaciones Químicas de Sevilla, Spain) from vacuum-dried samples. The analytical results for C, H, N, and S were within ± 0.4 of the theoretical values. Compounds 1-8 were examined for known classes of assay interference compounds using the freely available research tool for ligand discovery ZINC 15 (http://zinc15.docking. org) [35].²⁴ Starting materials **10** and **14** [36] and compounds **1–3** were synthesized following reported procedures [12]. Mannoside glycolipid conjugate (MGC) **9** was synthesized by the company Roowin (Riom, France). The synthesis, physico-chemical properties and cytotoxicity were previously reported [22–24].

4.2. Synthesis

4.2.1. (1R)-2,3,4-Tri-O-acetyl-1-dodecylthio-5N,6O-oxomethylidenegalactonojirimycin (11)

To a stirred solution of (1R)-1,2,3,4-tetra-O-acetyl-5N,6O-oxomethylidenegalactonojirimycin (10, 0.29 mmol) in anhydrous DCM (4 mL) at 0 °C, BF₃.OEt₂ (134 µL, 1.09 mmol) and dodecylthiol (0.61 mmol) were added under Ar atmosphere. The mixture was stirred for 30 min (TLC monitoring), diluted with DCM (50 mL) and washed with water (10 mL), aq NaHCO₃ (10 mL) and brine (10 mL), dried (MgSO₄) and concentrated under reduced pressure. Purification by column chromatography (1:3 EtOAc-cyclohexane) yielded **11** (118 mg, 79%). R_f 0.68 (1:1 EtOAc-cyclohexane). [α]_D +161.7 $(c \ 1.0 \ in \ DCM)$.¹H NMR (500 MHz, CDCl₃): $\delta 5.78 \ (d, 1 \ H, J_{1,2} = 5.1 \ Hz,$ H-1), 5.43 (t, 1 H, J_{3,4} = J_{4,5} = 1.6 Hz, H-4), 5.26–5.16 (m, 2 H, H-2, H-3), 4.43–4.34 (m, 2 H, H-5, H-6a), 4.04 (dd, 1 H, J_{6a,6b} = 13.5 Hz, $J_{5,6b} = 10.0$ Hz, H-6b), 2.57 (ddd, 1 H, ${}^{2}J_{H,H} = 12.7$ Hz, ${}^{3}J_{H,H} = 8.3$ Hz, ${}^{3}J_{H,H} = 6.0$ Hz, SCH₂), 2.43 (ddd, 1 H, SCH₂), 2.17–2.00 (3 s, 9 H, *Me*CO), 1.70–1.20 (m, 20 H, CH₂), 0.87 (t, 3 H, ${}^{3}J_{H,H} = 7.0$ Hz, CH₃). ${}^{13}C$ NMR (125.7 MHz, CDCl₃): δ 170.4-169.8 (MeCO), 155.7 (CO), 68.9 (C-3), 68.4 (C-4), 67.2 (C-2), 62.8 (C-6), 58.1 (C-1), 50.4 (C-5), 32.0-22.8 (CH₂), 30.4 (SCH₂), 20.8-20.7 (MeCO), 14.2 (CH₃). ESIMS: m/z 538.5 [M + Na]⁺. HRFABMS Calcd for C₂₅H₄₁NO₈SNa [M + Na]⁺ 538.2451, found 538.2444.

4.2.2. (1R,SS)- and (1R,SR)-2,3,4-Tri-O-acetyl-1-dodecylsulfinyl-5N,6O-oxomethylidenegalactonojirimycin (12 and 13)

To a solution of **11** (131 mg, 0.23 mmol) in DCM (6 mL), *m*CPBA (70%, 40 mg, 0.23 mmol) was added at 0 °C. The reaction mixture was stirred for 10 min, diluted with DCM (50 mL), washed with aqueous NaHCO₃ (10 mL), brine (10 mL), dried (MgSO₄) and concentrated under reduced pressure. The resulting crude was purified by column chromatography (1:2 EtOAc-cyclohexane) to afford the pure diastereomers **12** and **13**.

Yield of **12**: 44 mg (35%). R_f 0.53 (3:2 EtOAc-cyclohexane). [α]_D +68.4 (c 0.8 in DCM). ¹H NMR (300 MHz, CDCl₃): δ 5.72 (dd, 1 H, J_{2,3} = 10.7 Hz, J_{3,4} = 2.6 Hz, H-3), 5.49 (t, 1 H, J_{4,5} = 2.6 Hz, H-4), 5.45 (dd, 1 H, J_{1,2} = 7.5 Hz, H-2), 5.05 (d, 1 H, H-1), 4.96 (ddd, 1 H, J_{5,6a} = 9.2 Hz, J_{5,6b} = 3.8 Hz, H-5), 4.50 (t, 1 H, J_{6a,6b} = 9.2 Hz, H-6a), 4.07 (dd, 1 H, H-6b), 2.84 (ddd, 1 H, ²J_{H,H} = 13.0 Hz, ³J_{H,H} = 8.1 Hz, ³J_{H,H} = 6.8 Hz, SOCH₂), 2.65 (ddd, 1 H, SOCH₂), 2.17–2.01 (2 s, 9 H, MeCO), 1.78–1.20 (m, 20 H, CH₂), 0.87 (t, 3 H, ³J_{H,H} = 6.6 Hz, CH₃). ¹³C NMR (75.5 MHz, CDCl₃): δ 170.3–169.5 (MeCO), 157.3 (CO), 68.9–68.7 (C-3, C-4), 67.0–66.8 (C-1, C-2), 63.9 (C-6), 54.2 (C-5), 49.3 (SOCH₂), 32.0–22.8 (CH₂), 20.7 (MeCO), 14.2 (CH₃). ESIMS: m/z 554.3 [M + Na]⁺. Anal. Calcd for C₂₅H₄₁NO₉S: C, 56.48; H, 7.77; N, 2.63; S, 6.03. Found: C, 56.63; H, 7.90; N, 2.44; S, 5.79.

Yield of **13**: 39 mg (32%). R_f 0.47 (3:2 EtOAc-cyclohexane). [α]_D +75.4 (*c* 1.0 in DCM). ¹H NMR (300 MHz, CDCl₃): δ 5.65 (dd, 1 H, $J_{2,3} = 10.8$ Hz, $J_{3,4} = 2.6$ Hz, H-3), 5.58–5.48 (m, 2 H, H-2, H-4), 5.23 (d, 1 H, $J_{1,2} = 6.2$ Hz, H-1), 4.42 (t, 1 H, $J_{5,6a} = J_{6a,6b} = 9.0$ Hz, H-6a), 4.29 (ddd, 1 H, $J_{5,6b} = 4.1$ Hz, $J_{4,5} = 1.6$ Hz, H-5), 4.07 (dd, 1 H, H-6b), 2.98 (ddd, 1 H, $^2J_{H,H} = 13.0$ Hz, $^3J_{H,H} = 9.3$ Hz, $^3J_{H,H} = 5.3$ Hz, SOCH₂), 2.71 (m, 1 H, SOCH₂), 2.18–2.02 (3 s, 9 H, MeCO), 1.90–1.18 (m, 20 H, CH₂), 0.87 (t, 3 H, $^3J_{H,H} = 6.6$ Hz, CH₃). ¹³C NMR (75.5 MHz, CDCl₃): δ 170.3–169.5 (MeCO), 157.3 (CO), 68.9 (C-3), 68.7 (C-4), 67.0 (C-1), 66.8 (C-2), 63.9 (C-6), 52.2 (C-5), 49.3 (SOCH₂), 32.0–22.7 (CH₂), 20.7 (*Me*CO), 14.2 (CH₃). ESIMS: *m*/*z* 554.3 [M + Na]⁺. Anal. Calcd for C₂₅H₄₁NO₉S: C, 56.48; H, 7.77; N, 2.63; S, 6.03. Found: C, 56.55; H, 7.63; N, 2.31; S, 5.67.

4.2.3. (1R,SS)-1-Dodecylsulfinyl-5N,6O-

oxomethylidenegalactonojirimycin (4)

Compound **4** was obtained from **12** (30 mg, 0.06 mmol) by conventional *O*-deacetylation (see General Methods) and purification by column chromatography (4:1 EtOAc-cyclohexane \rightarrow EtOAc). Yield: 24 mg (98%). R_f 0.09 (EtOAc). [α]_D +64.1 (*c* 1.0 in MeOH). ¹H NMR (300 MHz, CD₃OD): δ 4.51 (t, 1 H, $J_{6a,6b} = J_{5,6a} = 8.6$ Hz, H-6a), 4.42 (dd, 1 H, $J_{5,6b} = 5.4$ Hz, H-6b), 4.35 (ddd, 1 H, $J_{4,5} = 2.0$ Hz, H-5), 4.29 (dd, 1 H, $J_{2,3} = 9.9$ Hz, $J_{1,2} = 6.8$ Hz, H-2), 3.99 (dd, 1 H, $J_{3,4} = 2.5$ Hz, H-3), 3.88 (t, 1 H, H-4), 3.00–2.82 (m, 2 H, SOCH₂), 1.84–1.73 (m, 2 H, SOCH₂CH₂), 1.58–1.21 (m, 18 H, CH₂), 0.90 (t, 3 H, ³ $_{JH,H} = 6.5$ Hz, CH₃). ¹³C NMR (75.5 MHz, CD₃OD): δ 160.2 (CO), 72.8–72.4 (C-1, C-3), 70.1 (C-4), 67.7 (C-2), 65.5 (C-6), 57.3 (C-5), 51.9 (SOCH₂), 33.0–23.7 (CH₂), 14.4 (CH₃). ESIMS: *m/z* 428.2 [M + Na]⁺. Anal. Calcd for C₁₉H₃₅NO₆S: C, 56.27; H, 8.70; N, 3.45; S, 7.91. Found: C, 56.16; H, 8.65; N, 3.13; S, 7.60.

4.2.4. (1R,SR)-1-Dodecylsulfinyl-5N,60oxomethylidenegalactonojirimycin (5)

Compound **5** was obtained from **13** (21 mg, 0.04 mmol) by conventional *O*-deacetylation (see General Methods) and final purification by column chromatography (4:1 EtOAc-cyclohexane \rightarrow EtOAc). Yield: 15 mg (96%). R_f 0.09 (EtOAc). [α]_D +65.0 (*c* 1.0 in MeOH). ¹H NMR (300 MHz, CD₃OD): δ 4.96 (d, 1 H, $J_{1,2} = 6.5$ Hz, H-1), 4.50–4.40 (m, 2 H, H-6a, H-6b), 4.37 (dd, 1 H, $J_{2,3} = 10.0$ Hz, H-2), 4.16 (ddd, 1 H, $J_{5,6a} = 8.7$ Hz, $J_{5,6b} = 5.2$ Hz, $J_{4,5} = 2.0$ Hz, H-5), 4.05 (dd, 1 H, $J_{3,4} = 2.5$ Hz, H-3), 3.90 (t, 1 H, H-4), 3.12 (ddd, 1 H, $^2J_{H,H} = 13.0$ Hz, $^3J_{H,H} = 8.7$ Hz, $^3J_{H,H} = 5.8$ Hz, SOCH₂), 2.98 (ddd, 1 H, SOCH₂), 1.87–1.70 (m, 2 H, SO₂CH₂CH₂), 1.60–1.21 (m, 18 H, CH₂), 0.90 (t, 3 H, $^3J_{H,H} = 6.8$ Hz, CH₃). ¹³C NMR (75.5 MHz, CD₃OD): δ 159.2 (CO), 73.1–72.3 (C-1, C-3), 70.0 (C-4), 69.5 (C-2), 65.3 (C-6), 56.5 (C-5), 50.8 (SOCH₂), 33.1–23.7 (CH₂), 14.4 (CH₃). ESIMS: *m/z* 428.2 [M + Na]⁺. Anal. Calcd for C₁₉H₃₅NO₆S: C, 56.27; H, 8.70; N, 3.45; S, 7.91. Found: C, 56.30; H, 8.90; N, 3.17; S, 7.75.

4.2.5. (1R)-1-Dodecylsulfonyl-5N,6O-

oxomethylidenegalactonojirimycin (6)

To a solution of 11 (51 mg, 0.09 mmol) in DCM (3 mL), mCPBA (70%, 31 mg, 0.18 mmol) was added at 0 °C. The reaction mixture was stirred for 20 min, diluted with DCM (50 mL), washed with aqueous NaHCO3 (10 mL), brine (10 mL), dried (MgSO4), concentrated under reduced pressure. Further purification by column chromatography (4:1 EtOAc-cyclohexane \rightarrow EtOAc) and conventional O-deacetylation (see General Methods) afforded 6. Yield: 18 mg (47% global yield). R_f 0.20 (EtOAc). [α]_D +23.2 (*c* 1.0 in MeOH). ¹H NMR (300 MHz, CD₃OD): δ 5.16 (d, 1 H, $J_{1,2}$ = 6.0 Hz, H-1), 4.50 (t, 1 H, $J_{6a,6b} = J_{5,6a} = 8.5$ Hz, H-6a), 4.42 (dd, 1 H, $J_{5,6b} = 5.2$ Hz, H-6b), 4.31 (ddd, 1 H, *J*_{4,5} = 1.7 Hz, H-5), 4.28–4.24 (m, 2 H, H-2, H-3), 3.88 (t, 1 H, J_{3,4} = 1.7 Hz, H-4), 3.39–3.34 (m, 2 H, SO₂CH₂), 1.84 (m, 2 H, ${}^{3}J_{H,H} = 7.2 \text{ Hz}, \text{ SO}_{2}\text{CH}_{2}\text{CH}_{2}$), 1.48–1.29 (m, 18 H, CH₂), 0.90 (t, 3 H, ${}^{3}J_{H,H} = 6.7$ Hz, CH₃). 13 C NMR (75.5 MHz, CD₃OD): δ 159.2 (CO), 72.3 (C-1), 70.7 (C-2), 70.2 (C-4), 67.8 (C-3), 65.4 (C-6), 56.3 (C-5, CH₂SO₂), 33.0–22.5 (CH₂), 14.4 (CH₃). ESIMS: *m*/*z* 444.2 [M + Na]⁺. Anal. Calcd for C₁₉H₃₅NO₇S: C, 54.14; H, 8.37; N, 3.32; S, 7.61. Found: C, 53.90; H, 8.18; N, 3.11; S, 7.28.

4.2.6. (1R)-2,3,4-Tri-O-acetyl-1-dodecylthio-5N,6O-oxomethylidenemannonojirimycin (15)

To a stirred solution of (1R)-1,2,3,4-tetra-O-acetyl-5N,6O-oxomethylidenemannojirimycin (**14**, 0.29 mmol) in anhydrous DCM (4 mL) at 0 °C, BF₃.OEt₂ (134 µL, 1.09 mmol) and dodecylthiol (0.61 mmol) were added under Ar atmosphere. The mixture was stirred for 30 min (TLC monitoring), diluted with DCM (50 mL) and washed with water (10 mL), aq NaHCO₃ (10 mL) and brine (10 mL), dried (MgSO₄) concentrated under reduced pressure and subjected to column chromatography (1:4 \rightarrow 1:2 EtOAc-cyclohexane) to give **15.** Yield: 101 mg (68%). R_f 0.72 (1:1 EtOAc-cyclohexane). [α]_D +18.2 (*c* 1.0 in DCM). ¹H NMR (300 MHz, CDCl₃): δ 5.33–5.28 (m, 2 H, H-2, H-3), 5.22 (t, 1 H, J_{3,4} = J_{4,5} = 9.9 Hz, H-4), 5.15 (d, 1 H, J_{1,2} = 2.0 Hz, H-1), 4.41 (dd, 1 H, J_{6a,6b} = 9.1 Hz, J_{5,6a} = 8.2 Hz, H-6a), 4.32 (dd, 1 H, J_{5,6b} = 5.1 Hz, H-6b), 4.08 (ddd, 1 H, H-5), 2.73–2.51 (m, 2 H, CH₂S), 2.10–1.99 (3 s, 9 H, MeCO), 1.67–1.51 (m, 2 H, CH₂CH₂S), 1.41–1.17 (m, 18 H, CH₂), 0.86 (t, 3 H, ³J_{H,H} = 6.6 Hz, CH₃).¹³C NMR (75.5 MHz, CDCl₃): δ 170.3–169.6 (MeCO), 156.3 (CO), 70.8, 69.1 (C-2, C-3), 69.5 (C-4), 66.1 (C-6), 58.3 (C-1), 52.8 (C-5), 31.9–28.7 (CH₂), 20.8–20.6 (MeCO), 14.2 (CH₃). ESIMS: *m*/z 538.3 [M + Na]⁺. Anal. Calcd for C₂₅H₄₁NO₈S: C, 58.23; H, 8.01; N, 2.72; S, 6.22. Found: C, 58.32; H, 8.20; N, 2.60; S, 6.04.

4.2.7. (1R,SS)-2,3,4-Tri-O-acetyl-1-dodecylsulfinyl-5N,6Ooxomethylidenemannonojirimycin (**16**)

To a solution of 15 (131 mg, 0.23 mmol) in DCM (6 mL), mCPBA (70%, 40 mg, 0.23 mmol) was added at 0 °C. The reaction mixture was stirred for 10 min, diluted with DCM (50 mL), washed with aqueous NaHCO3 (10 mL), brine (10 mL), dried (MgSO4) and concentrated under reduced pressure. The resulting crude was purified by column chromatography (Et₂O) to give 16. Yield: 48 mg (39%). R_f 0.59 (Et₂O). [a]_D - 4.1 (c 1.0 in DCM). ¹H NMR (300 MHz, CDCl₃): δ 5.73 (t, 1 H, $J_{1,2} = J_{2,3} = 2.5$ Hz, H-2), 5.61 (dd, 1 H, $J_{3,4} = 9.0$ Hz, H-3), 5.27 (t, 1 H, $J_{4,5} = 9.0$ Hz, H-4), 4.62 (d, 1 H, H-1), 4.60–4.50 (m, 1 H, H-5), 4.52 (t, 1 H, $J_{6a,6b} = J_{5,6a} = 8.3$ Hz, H-6a), 4.38 (dd, 1 H, J_{5.6b} = 3.7 Hz, H-6b), 2.95 (m, 1 H, SOCH₂), 2.79 (m, 1 H, SOCH₂), 2.16-2.01 (3 s, 9 H, MeCO), 1.79-1.18 (m, 20 H, CH₂), 0.87 (t, 3 H, ${}^{3}I_{HH} = 6.9$ Hz, CH₃). ${}^{13}C$ NMR (75.5 MHz, CDCl₃): δ 170.3–169.4 (MeCO), 157.7 (CO), 69.6 (C-2), 69.0 (C-3), 68.8 (C-4), 68.4 (C-1), 66.9 (C-6), 55.8 (C-5), 50.8 (SOCH₂), 31.9-22.4 (CH₂), 20.7-20.6 (MeCO), 14.1 (CH₃). ESIMS: m/z 554.5 [M + Na]⁺. Anal. Calcd for C₂₅H₄₁NO₉S: C, 56.48; H, 7.77; N, 2.63; S, 6.03. Found: C, 56.13; H, 7.44; N, 2.29; S, 5.76.

4.2.8. (1R,SS)-1-Dodecylsulfinyl-5N,60-

oxomethylidenemannonojirimycin (7)

Compound **7** was obtained from **16** (20 mg, 0.04 mmol) by conventional O-deacetylation. Column chromatography (5:1 EtOAc-MeOH). Yield: 14 mg (86%). R_f 0.52 (5:1 EtOAc-MeOH). [α]_D +23.8 (*c* 0.6 in MeOH). ¹H NMR (300 MHz, CD₃OD): δ 4.67 (d, 1 H, $J_{1,2} = 2.0$ Hz, H-1), 4.63 (t, 1 H, $J_{6a,6b} = J_{5,6a} = 8.7$ Hz, H-6a), 4.38 (t, 1 H, $J_{2,3} = 2.0$ Hz, H-2), 4.34 (dd, 1 H, $J_{2,3} = 2.0$ Hz, H-6b), 4.11 (ddd, 1 H, $J_{4,5} = 9.3$ Hz, H-5), 3.90 (dd, 1 H, $J_{2,3} = 2.0$ Hz, $J_{3,4} = 9.3$ Hz, H-3), 3.75 (t, 1 H, H-4), 2.98–2.81 (m, 2 H, SOCH₂), 1.82–1.71 (m, 2 H, SOCH₂CH₂), 1.52–1.24 (m, 18 H, CH₂), 0.90 (t, 3 H, ³ $J_{H,H} = 6.7$ Hz, CH₃). ¹³C NMR (75.5 MHz, CD₃OD): δ 160.5 (CO), 73.6–73.0 (C-1, C-3), 71.9–71.2 (C-2, C-4), 68.7 (C-6), 58.9 (C-5), 51.1 (SOCH₂), 33.1–23.6 (CH₂), 14.5 (CH₃). ESIMS: *m/z* 428.3 [M + Na]⁺. HRFABMS Calcd for C₁₉H₃₅NO₆SNa [M + Na]⁺ 428.2077, found 428.2081.

4.2.9. (1R)-2,3,4-Tri-O-acetyl-1-dodecylsulfonyl-5N,6Ooxomethylidenemannojirimycin (17)

To a solution of **15** (51 mg, 0.09 mmol) in DCM (3 mL), *m*CPBA (70%, 31 mg, 0.18 mmol) was added at 0 °C. The reaction mixture was stirred for 20 min, diluted with DCM (50 mL), washed with aqueous NaHCO₃ (10 mL), brine (10 mL), dried (MgSO₄) and concentrated under reduced pressure. Final purification by column chromatography (1:3 EtOAc-cyclohexane) afforded **17**. Yield: 33 mg (71%). R_f 0.66 (1:1 EtOAc-cyclohexane). [α]_D -6.5 (*c* 1.0 in DCM). ¹H NMR (300 MHz, CDCl₃): δ 5.93 (t, 1 H, $J_{1,2} = J_{2,3} = 2.6$ Hz, H-2), 5.54 (dd, 1 H, $J_{3,4} = 9.0$ Hz, H-3), 5.21 (t, 1 H, $J_{4,5} = 9.0$ Hz, H-4), 5.06 (d, 1 H, H-1), 4.56–4.49 (m, 1 H, H-6a), 4.43–4.37 (m, 2 H, H-5, H-6b), 3.05 (m, 2 H, SO₂CH₂), 2.13–2.02 (3 s, 9 H, MeCO), 1.95–1.78 (m, 2 H, SO₂CH₂CH₂), 1.46–1.19 (m, 18 H, CH₂), 0.87 (t, 3 H, ³ $J_{HH} = 6.7$ Hz,

CH₃). ¹³C NMR (75.5 MHz, CDCl₃): δ 170.4–169.2 (MeCO), 156.6 (CO), 69.4 (C-1), 69.0 (C-4), 68.9 (C-3), 67.2 (C-6), 65.0 (C-2), 54.1 (C-5), 52.2 (CH₂SO₂), 32.0–21.7 (CH₂), 20.8–20.6 (*Me*CO), 14.2 (CH₃). ESIMS: *m*/*z* 570.4 [M + Na]⁺. Anal. Calcd for C₂₅H₄₁NO₁₀S: C, 54.83; H, 7.55; N, 2.56; S, 5.85. Found: C, 55.01; H, 7.69; N, 2.29; S, 5.62.

4.2.10. (1R)-1-Dodecylsulfonyl-5N,6Ooxomethylidenemannonojirimycin (8)

Compound **8** was obtained by conventional *O*-deacetylation (see General Methods) of **17**. Column chromatography (4:1 EtOAc-cyclohexane \rightarrow EtOAc). Yield: 12 mg (64%). R_f 0.20 (EtOAc). [α]_D +23.2 (*c* 1.0 in MeOH). ¹H NMR (300 MHz, CD₃OD): δ 5.03 (d, 1 H, $J_{1,2} = 1.6$ Hz, H-1), 4.65–4.59 (m, 2 H, H-2, H-6a), 4.36 (dd, 1 H, $J_{6a,6b} = 9.0$ Hz, $J_{5,6b} = 5.0$ Hz, H-6b), 4.08 (td, 1 H, $J_{4,5} = J_{5,6a} = 9.0$ Hz, H-5), 3.86 (dd, 1 H, $J_{3,4} = 9.5$ Hz, $J_{2,3} = 3.1$ Hz, H-3), 3.73 (t, 1 H, H-4), 3.25–3.06 (m, 2 H, SO₂CH₂), 1.89–1.73 (m, 2 H, SO₂CH₂CH₂), 1.47–1.26 (m, 18 H, CH₂), 0.89 (t, 3 H, ³ $J_{H,H} = 6.7$ Hz, CH₃). ¹³C NMR (75.5 MHz, CD₃OD): δ 159.5 (CO), 74.0 (C-1), 73.0 (C-3), 70.7 (C-4), 68.6 (C-6), 66.9 (C-2), 57.0 (C-5), 52.0 (CH₂SO₂), 32.0–22.7 (CH₂), 14.4 (CH₃). ESIMS: *m/z* 444.3 [M + Na]⁺. Anal. Calcd for C₁₉H₃₅NO₇S: C, 54.14; H, 8.37; N, 3.32; S, 7.61. Found: C, 53.88; H, 8.12; N, 3.04; S, 7.32.

4.3. Cell culture

Human primary cells were cultured at 37 °C and 5% CO₂ in RPMI 1640 supplemented with gentamycin and 10% (v/v) heatinactivated fetal calf serum (complete medium). Human monocytes were purified from buffy coats by successive Ficoll and Percoll gradients. Monocyte-derived DCs were differentiated from monocytes by 5 days of culture with GM-CSF (50 ng ml⁻¹; ImmunoTools) and IL-4 (10 ng ml⁻¹; ImmunoTools), as described.¹⁹ Proper differentiation was characterized by low CD14 and high CD1a and DC-SIGN expression levels.

4.4. Flow cytometry

The cell phenotype was analyzed using the following antibodies from BD Bioscience: HLA-DR-FITC (Tü39 or L243), DC-SIGN/CD209-PerCP-Cy5.5 (DCN46), CD1a-APC (HI149), CD86-FITC (FUN-1), CD83-APC (HB15e), and CD14-PE (MEM-15, ImmunoTools). Data were acquired on a FACSCalibur (Becton-Dickinson) or Gallios (Beckman-Coulter) after exclusion of dead cells by Sytox Red (Molecular Probes, Invitrogen) or 7-AAD (BD-Pharmingen) DNA intercalating dyes. Data were analyzed using the Cell Quest Pro software (BD Bioscience) or FlowJo (Treestar).

4.5. Cell activation

DCs (10^5 cells/well) were cultured in complete medium ($200 \,\mu$ L) and incubated in the presence of Ultra pure E. *coli* K12 lipopoly-saccharide (LPS; $100 \,ng \,mL^{-1}$). Compounds were usually added at the same time as LPS, or, where indicated, 2, 4 or 6 h after LPS addition.

4.6. Enzyme-linked ImmunoabSorbent Assay (ELISA)

After 24 h of stimulation with LPS, cell supernatants were collected and stored at -20 °C. Levels of TNF- α released in the media, or in the case of *in vivo* assays, levels of IL-6 in the mouse sera, were measured by ELISA with the corresponding BD OptEIA Set (BD-Pharmingen).

4.7. Effect on cell surface proteins

DCs (10^5 cells) in complete medium (200μ L) were plated in a 96-well plate. They were left untreated or treated with sp²-IGLs (25 and 50μ M) or conjugate MGC **9** (100μ M) for 30 min at 37 °C. They were washed, kept on ice, labeled with anti-CD14-PE, and analyzed by flow cytometry.

4.8. Co-localization analysis of CD14 and TLR-4 by confocal microscopy

Assays were performed according to the previously described procedure [22]. Briefly, DCs (10^5 cells per chamber) were cultured on poly-L-Lysine-coated, 8-chambers slides (Lab-Tek, Nunc) in complete medium ($300 \,\mu$ L). Cells were left untreated or treated with sp²-IGLs ($50 \,\mu$ M) or MGC **9** ($100 \,\mu$ M) for 30 min. Cells were washed, fixed, permeabilized and labeled with anti-TLR-4 polyclonal goat antibody (R&D Systems AF1478, $5 \,\mu$ g mL⁻¹) followed by AF488-donkey anti-goat (Molecular Probes, Invitrogen), and anti-CD14-APC (ImmunoTools). DAPI was used as nuclear counterstaining. Slides were mounted using Fluoromount (Dako). Images were acquired on a Zeiss LSM 780 confocal microscope with GaAsP detector and Zen acquisition software. Images were processed using the ImageJ software.

4.9. In vivo assays in C57BL/6J mice

All animals received humane care in compliance with the guidelines formulated by the French Ministry of Agriculture and of Higher Education and Research, and all procedures were reviewed by the Regional Ethical Committee for Animal Experimentation of the University of Strasbourg. sp²-IGL **3** was prepared in sterile solvent (7.25 mg mL⁻¹) by solubilization in DMSO (10.5 mg in 1 mL) followed by addition of H_2O (0.5 mL) and PBS x10 (150 μ L). LPS (0.5 mg/kg; strain O111.B4; Sigma-Aldrich) was prepared in sterile PBS. To evaluate the in vivo toxicity of the compound, 8-week old male and female mice received intraperitoneal (i.p.) administrations of 200 μ l of sp²-IGL **3** (30 mg/kg) or of control DMSO/PBS on day 0, 1, 2, 3. Evaluation of their well-being and weight were performed every day up to day 12. To assess the compounds for their capacity to block LPS-induced IL-6 production, C57BL/6J male 8week old mice received a single intraperitoneal (i.p.) administration of sp²-IGL **3** (50 mg/kg) 1 h before LPS (0.5 mg/kg) challenge. Control animals received equivalent volumes of solvent (PBS or PBS/DMSO 2:1) (200 μ L) by the intraperitoneal route. Blood (150 µL) was taken 2 h after LPS injection. Serum was prepared and IL-6 measured by ELISA (see above).

Statistical Analysis. Statistical analysis was performed using paired Student's t-test. Data were considered significantly different when p was less than 0.05.

Notes

The authors declare no competing financial interest.

Abbreviations used

DC, dendritic cell; DMSO, dimethyl sulfoxide; DCM, dichloromethane; *m*CPBA, *m*-chloroperoxybenzoic acid; sp²-IGL, sp²-iminosugar glycolipid; *R*-DSO-ONJ, (1*R*)-1-dodecylsulfinyl-5*N*,60oxomethylidenenojirimycin; DSO₂-ONJ, (1*R*)-1-dodecylsulfonyl-5*N*,60-oxomethylidenenojirimycin; LPS, lipopolysaccharide; MGC, mannoside glycolipid conjugate; IL-10, interleukin-10; IL-6, interleukin-6; TNF- α , Tumor Necrosis Factor- α . MAPK, mitogen activated protein kinase.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.ejmech.2019.02.078.

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