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Review Article

Endogenous and Exogenous CD1-Binding Glycolipids

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In the same way that peptide antigens are presented by major histocompatibility complex (MHC) molecules, glycolipid antigens can also activate the immune response via binding to CD1 proteins on antigen-presenting cells (APCs) and stimulate CD1-restricted T cells. In humans, there are five members of the CD1 family, termed CD1a–e, of which CD1a–d are involved in glycolipid presentation at the cell surface, while CD1e is involved in the intracellular trafficking of glycolipid antigens. Both endogenous (self-derived) and exogenous (non-self-derived) glycolipids have been shown to bind to members of the CD1 family with varying degrees of specificity. In this paper we focus on the key glycolipids that bind to the different members of the CD1 family.

1. Introduction

Carbohydrates play an important role in various molecular recognition events. They are found on the surfaces of cells, bacteria, and viruses as glycoconjugates (such as glycoproteins and glycolipids), and it is therefore not surprising that their interaction with cells of the immune system forms an integral part of the body's immune response. The innate immune system operates largely via receptor-ligand interactions, and glycoconjugates are capable of inducing the strongest immune responses known. Classic examples include the landmark discovery of endotoxin by Pfeiffer in the late 19th century [1] and the discovery in 1928 by Landsteiner and Levine that blood group determinants were responsible for the blood cell rejection (also known as transfusion reaction) observed during blood transfusion with incompatible donors [2]. Only later was it revealed that endotoxin is in fact a glycolipid [3] and that the A, B, and O antigens responsible for transfusion reaction are glycolipids and glycoproteins found on the surface of red blood cells [4].

Recently, cell-mediated immunity to glycolipid antigens has been discovered, and a glycolipid presentation pathway which parallels that of peptide antigen presentation has emerged. The glycolipid pathway involves CD1 proteins, major histocompatibility complex- (MHC-) like molecules

that are capable of presenting glycolipids to a specialised subset of CD1-restricted T cells. Like MHC molecules, CD1 proteins are present on antigen-presenting cells (APCs), specialised immune cells that perform a crucial role in the immune system. After encountering an antigen, the APC sends messages to other immune cells, subsequently activating the full immune response. This paper will focus on CD1-binding glycolipids that are capable of activating and modulating the immune system via the activation of CD1-restricted T cells.

2. CD1 Molecules

The CD1 protein is a transmembrane glycoprotein consisting of a heavy chain with three extracellular domains ($\alpha 1$, $\alpha 2$, and $\alpha 3$) noncovalently linked to a β_2 -microglobulin (β_2m) chain (Figure 1) [5]. The nonpolar amino acids in the $\alpha 1$ and $\alpha 2$ chains form the hydrophobic pockets that can accommodate the long lipid tails of a glycolipid. The CD1 family contains five members, CD1a, CD1b, CD1c, CD1d, and CD1e, which, depending on their sequence homology, can be further classified into three groups: group 1 (CD1a, CD1b, and CD1c), group 2 (CD1d), and group 3 (CD1e). In a CD1-glycolipid complex (Figure 1), the polar carbohydrate group

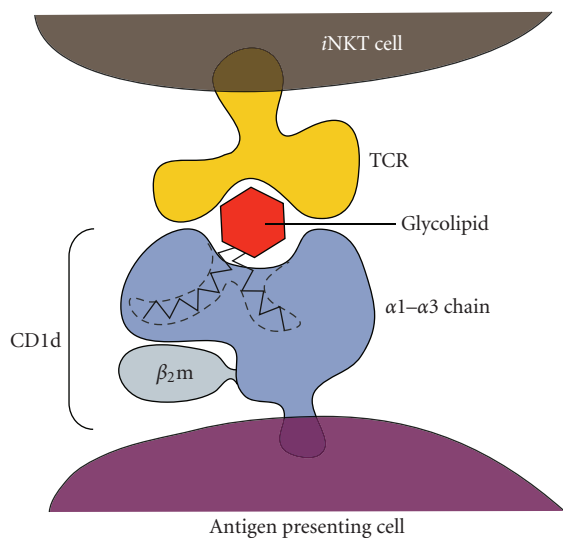


FIGURE 1: Representation of CD1-glycolipid-TCR interaction during glycolipid antigen presentation.

of the glycolipid protrudes from the CD1 binding groove, allowing it to be recognised by CD1-restricted T cells via the T cell receptor (TCR). The trimolecular interaction between CD1, glycolipid, and TCR initiates a cascade of intracellular signalling that activates the T cells to produce signalling molecules called cytokines. The type of immune response generated is dependent on the cytokine profile produced.

The exact effect that the glycolipid molecular structure has on the immune response is only known in a very general sense. Unlike MHCs, which specifically bind peptide antigens, CD1 molecules accommodate structurally diverse glycolipids. Extensive crystal structure analysis of CD1 molecules with bound ligands suggests that CD1 binding is less specific, which supports the observation that the length of a lipid tail does not greatly affect the immunomodulatory profile of a given glycolipid [6]. Instead, the highly specific interaction of the TCR with the CD1-glycolipid complex plays the principal role in dictating the outcome of the immune response [7, 8]. Other processes such as CD1 and glycolipid trafficking into the cell and glycolipid processing prior to TCR recognition have also been found to have a profound effect on the immune response [9–11]. The ability of glycolipids to modulate the immune response prompted researchers to develop specific CD1-binding glycolipids that can be used in the investigation and treatment of many diseases, including cancer, bacterial infections, and autoimmune diseases such as multiple sclerosis and systemic lupus.

3. Endogenous and Exogenous Glycolipids

Production of antigen-specific T cells requires positive selection in the thymus. For the glycolipid reactive CD1-restricted T cells, this means that endogenous glycolipids, originating from within the host organism, must be presented by CD1 proteins and interact with the TCRs of developing CD1-restricted T cells. Isoglobotrihexosylceramide (iGb3) is

the most studied endogenous glycolipid and has been suggested to be the pivotal ligand for the selection of CD1d-restricted T cells in the thymus [12], though further investigations are required to support this hypothesis [13]. Other endogenous glycolipids such as gangliosides and sulfatide, as well as phosphoglycerolipids and sphingomyelin, have also been found to stimulate group 1 CD1-restricted T cells. The exact role that endogenous lipids have in the activation of CD1-restricted T cells, however, remains unclear.

Of the exogenous glycolipids, numerous bacterial cell wall glycolipids have been shown to play an important role in the immune response. In particular, the discovery that mycolic acid, a nonpeptide lipid antigen from *Mycobacterium tuberculosis* (*M. Tb*), stimulates T cells via CD1 molecules paved the way towards the discovery of a host of glycolipids that activate the immune system in a similar manner [14]. Subsequently, other glycolipid cell wall components of *M. Tb*, such as glucose monomycolate (GMM) [14], phosphatidylinositolmannoside (PIM) [15], and lipoarabinomannan [16], have also been found to be immunogenic [17]. However, not all bacterial glycolipids are responsible for the activation of the immune system during infection. Interestingly, Mattner et al. discovered that the antigen responsible for activating CD1-restricted T cells in mice during an infection with Gram-negative, LPS-positive *Salmonella typhimurium*, was in fact the endogenous glycolipid, iGb3 [18].

When comparing endogenous and exogenous CD1-binding glycolipids, perhaps the most obvious structural difference is the linkage variation between the carbohydrate and the lipid backbone. Endogenous glycolipids such as iGb3 have β -linkages, while exogenous/pathogenic glycolipids (i.e., most bacterial glycolipids) typically have α -linkages. In this paper, endogenous and exogenous CD1-binding glycolipids will be described in the context of the CD1 isotype with which they interact. The origin/discovery, synthesis, and effect on the immune response of each glycolipid will also be discussed.

3.1. CD1a. The human CD1a molecule possesses two hydrophobic pockets (A' and F') that can bind aliphatic carbon chains of glycolipids via nonpolar Van der Waals interactions. The A' pocket of CD1a is narrow with only one entry point and hence has been proposed to be a “molecular ruler” that allows CD1a to bind C_{20} lipids or shorter [19]. The F' pocket of CD1a is also unique in that it lies close to the plane of TCR recognition and hence directly participates in antigen presentation [6].

3.1.1. Endogenous Glycolipids. Sulfatide, illustrated by 1 (Scheme 1), is a β -D-galactosylceramide sulfated at C-3 of D-galactose. Ceramides are long chain dihydroxy or trihydroxy bases which are linked to a fatty acid via an amide linkage at C-2. The sphingosine base, a dihydroxylated base with mono-unsaturation across C-4 and C-5, is commonly found in endogenous glycolipids and forms the lipid backbone of sulfatides. Sulfatide is a potent glycolipid produced by mammalian cells and is found mainly in the myelin sheath but is also present in small amounts in other tissues [20].

The ability of sulfatide to bind to all group 1 and group 2 CD1 proteins (CD1a–d) gives it the just reputation of being the most promiscuous CD1-binding glycolipid [20, 21]. Nonetheless, its binding affinity differs with each CD1 isotype, with the CD1a-sulfatide complex seeming to be the most stable [20]. Here the narrow A' pocket allows CD1a to selectively bind to the C₁₈ sphingosine backbone while the acyl lipid protrudes along the surface of the CD1a molecule and extends into the F' pocket. The 3-O-sulfogalactose protrudes from the CD1 pocket and interacts with the TCR of the T cell.

Elevated levels of sulfatide-reactive T cells in multiple sclerosis patients reveal that this endogenous glycolipid may be responsible, at least in part, for the pathogenesis of neurodegenerative diseases. The ability of the sulfatide glycolipid to stimulate CD1-restricted T cells has proven valuable in the understanding and treatment of autoimmune diseases of the central nervous system, particularly in multiple sclerosis.

With regard to their syntheses, several β -sulfatides with varying chain lengths and α -analogues of sulfatides have been reported by Compostella et al. and Franchini et al., respectively [22, 23]. The former employed a β -glycosylation of an azidosphingosine **2** with pivaloyl protected D-galactosyl trichloroacetimidate **3** as a key step (Scheme 1). After global deprotection of the coupled product **4**, the azide was reduced and the resulting amine derivatized with a fatty acid to obtain galactosyl ceramide **5**. Regioselective 3-sulfation of **5** then gave the desired sulfatide **1**, in excellent overall yield.

3.2. CD1b. The CD1b isotype, found in humans but not in mice, has a binding groove with four pockets: A', C', F', and T' [24]. The T' tunnel, which connects the A' and F' pockets, is unique to CD1b and accounts for the ability of the combined binding pockets to accommodate long alkyl chains (~C₆₈). The extra C' pocket, which is also only found in CD1b, contributes to the receptor's ability to host large bacterial antigens such as the sulfotrehalose and mycolyl antigens [25]. Similar to CD1a, nonpolar amino acids lining the interior of the binding pockets provide nonpolar Van der Waals interactions with the glycolipid lipid chains.

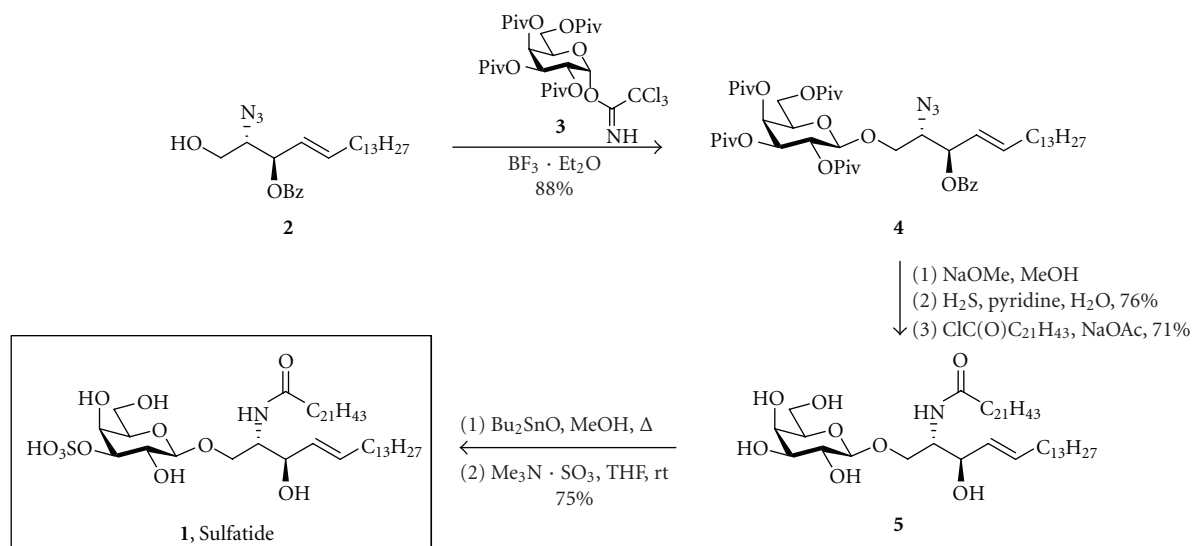
3.2.1. Endogenous Glycolipids. Gangliosides (e.g., GM1 (**6**) and GQ1b (**7**), Figure 2) are sialic acid containing glycosphingolipids and are constituents of the plasma membrane of various cells. Gangliosides are perhaps one of the largest classes of endogenous glycolipids, and autoreactivity of T cells or IgG autoantibodies to these gangliosides may be the cause of various immune-mediated diseases such as multiple sclerosis, rheumatoid arthritis, lupus erythematosus, and psoriasis [26]. In all gangliosides, the ceramide is linked via C-1 to a β -O-glucose moiety, which in turn is glycosylated at the 4-OH with a β -O-galactosyl residue. The structural diversity of gangliosides results from further glycosylation of this disaccharide.

GM1 (**6**, Figure 2) is the most abundant ganglioside found in the human myelin sheath and consists of a ceramide with a core tetrasaccharide to which a single sialic acid residue is attached. GM1 binds to CD1b on the cell surface

at neutral pH and is recognised without internalisation and processing [27]. These CD1b-GM1 complexes can stimulate T cells to induce a T-helper 1- (Th1-) type response measured by the release of cytokines such as TNF- α and IFN- γ , which are important cytokines for the generation of a cytotoxic immune response [27]. Ganglioside-specific T cells can discriminate small differences in the carbohydrate portion of the glycolipid as revealed by a study whereby analogues of GM1 that lack the terminal galactose residue (GM2), sialic acid residue (asialo-GM1), or both were unable to stimulate production of IFN- γ by T cells [27]. This suggests that the GM1 pentasaccharide represents the minimum epitope for recognition by TCRs. The structure of the lipid tail and the fatty acid is also important for the specificity of CD1b binding, with stearic acid having the highest T cell stimulatory capacity [27]. GM1 binding to CD1b is highly reversible, and other ceramide-containing glycosphingolipids can displace GM1 in the CD1b complex. The first total synthesis of GM1 occurred in 1986 by Sugimoto et al. and took an impressive twelve linear steps [28]. From the retrosynthesis (Scheme 2), it can be seen that the main steps involved the coupling of the peracetylated pentasaccharide imidate **8** to sphingosine acceptor **9**, which was accomplished in 33% yield using BF₃·Et₂O as the activator. Donor **8** in turn was accessible via coupling of galactosyl galactosamine **10** and sialylated lactoside **11**.

More recently, the methyl glycoside of GM1 has been synthesised by Bhattacharya and Danishefsky along with the asialo GM1 analogue [29]. The study of the binding of GM1 and its analogues to CD1b and the subsequent activation of ganglioside-specific T cells will aid in the understanding of the autoimmune disease multiple sclerosis as it was found that gangliosides are present in increased frequencies in patients with this disease [30].

GQ1b (**7**, Figure 2) is another ganglioside that binds to CD1b and stimulates CD1-restricted T cells [31]. It was first isolated from the human brain in 1963 and later characterised in 1979 [32–34]. GQ1b consists of the core ganglioside tetrasaccharide to which four sialic acid residues are connected. Similar to other endogenous glycolipids, the ceramide contains the sphingosine backbone. GQ1b is most abundant in the mammalian central nervous system and participates in physiological activities such as toxin binding, modulation of protein phosphorylation, cell adhesion and growth, and apoptosis [35, 36]. GQ1b binds to CD1b molecules and, upon binding to the T cell TCR, induces the production of the Th1 cytokines IL-2 and IFN- γ while suppressing Th2 cytokine production [37]. While this suggests that perturbation of GQ1b levels may lead to a Th1/Th2 imbalance, elevated or suppressed levels of GQ1b have yet to be correlated with diseases resulting from this immunological imbalance [37]. Total synthesis of GQ1b has been accomplished by Ishida and coworkers in 1994 [31] and more recently by Imamura and colleagues in 2009 [35]. Ishida and coworkers used trimethylsilyloxyethyl (SE) protected octasaccharide **12** (Figure 3) to produce imidate donor **13**, which could be efficiently coupled to azide acceptor **2** (using TMSOTf and in 46% yield), *en route* to GQ1b.



SCHEME 1: Representative synthesis of a sulfatide.

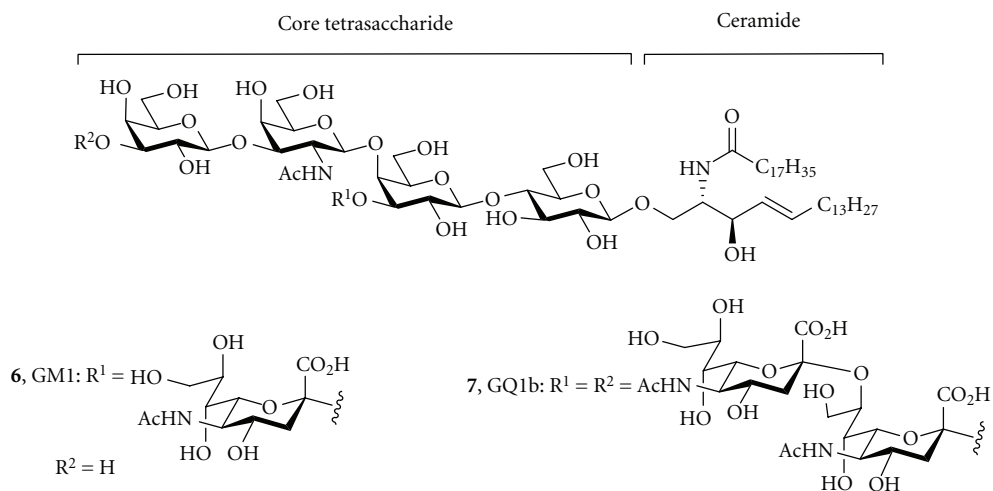


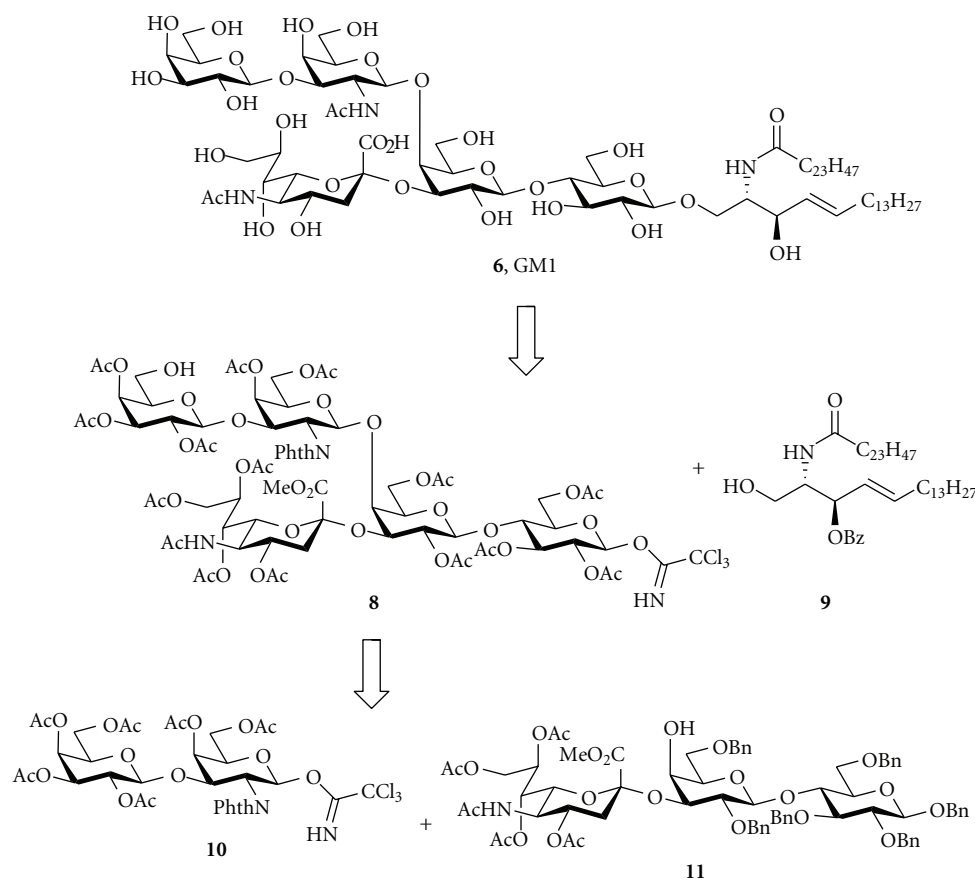
FIGURE 2: Gangliosides GM1 and GQ1b.

3.2.2. *Exogenous Glycolipids.* Glucose-monomycolate (GMM, **14**, Figure 4) is the 6-O-mycolate ester of D-glucose and is found in bacterial species including *Mycobacteria*, *Rhodococcus*, and *Nocardia*. Mycolic acids themselves are α -alkyl, β -hydroxy fatty acids that are present in most mycobacterium species and also in related taxa. Mycolic acids from different species of bacteria may vary to give rise to α -alkyl-, keto-, methoxy-, or epoxy-mycolic acids. The GMM (**14**) illustrated contains the α -mycolic acid backbone. It should also be noted that mycolic acids were the first lipid antigens revealed to activate CD1-restricted T cells via binding to CD1b molecules [14].

GMM is a potent ligand for CD1-restricted T cells when presented by CD1 molecules [7]. For example, studies have shown that the CD1b-restricted T-cell recognition of GMM is highly specific for the glucose moiety since naturally occurring mycolate esters of glycerol and arabinose are not

recognised by CD1-restricted T cells. In addition, the *R,R* configuration of the hydroxy acid and the position of the linkage of the mycolate to the glucose unit are also important for CD1 binding [7]. Structural and mass-spectroscopy analysis has shown that high-affinity binding also depends on the exact structure of the acyl side chains of the antigen [38]. Crystal structure studies of the CD1b-GMM complex indicate that both acyl chains are buried in the antigen binding groove of CD1b, thus leaving the glucose unit exposed to the surface for recognition by TCRs. This explains the observation that one TCR can recognise GMMs with varying chain lengths derived from different bacterial sources [39].

Diacylated trehalose sulphates (Acyl₂SGL, **15**, Figure 5) are lipid antigens produced by virulent *M. Tb* and consist of a trehalose core that is acylated at the 2-position by palmitic (or stearic) acid and acylated at the 3-position by



SCHEME 2: Synthesis of GM1.

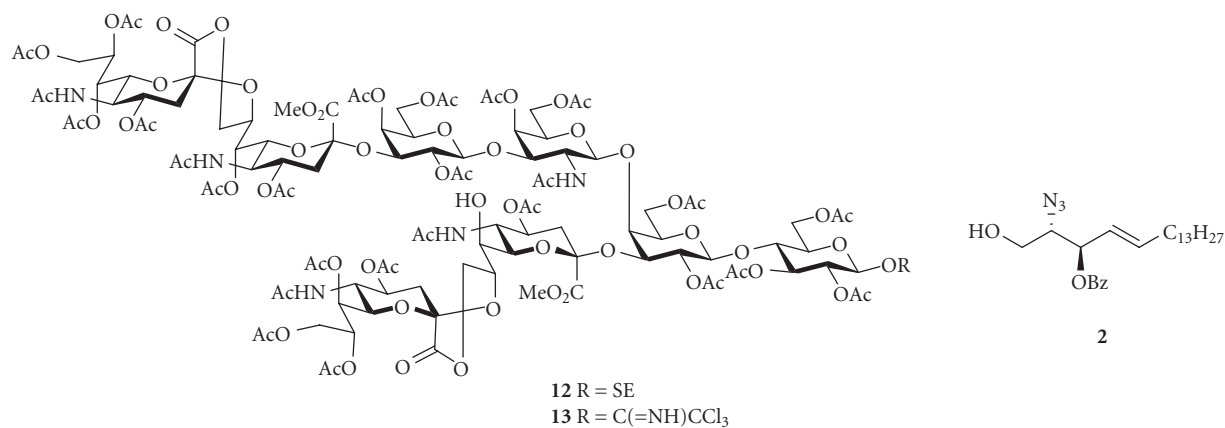


FIGURE 3: Building blocks for the synthesis of GQ1b.

hydroxyphthioceranoic acid [40]. While the methyl branches were assigned to be of the *L*-series, the stereochemistry of the hydroxy-substituted carbon has not been assigned [41].

Sulfoglycolipids bind to human CD1b, stimulate CD1b-restricted T cells to release the cytotoxic cytokine IFN- γ , and recognise *M. Tb*-infected cells in a host, leading to killing of intracellular bacteria [40]. The influence that the lipid chains have on T cell activation in relation to the length, position, and structure of the fatty acid residues has been studied,

and it was observed that the presence and stereochemistry of a multimethyl-branched fatty acid appendage on the 3-position in Acyl₂SGL were crucial for T cell stimulation [42]. Naturally occurring Acyl₂SGL, illustrated in 15, was seen to be most potent in T-cell activation as revealed in a study by Guiard and coworkers, who synthesised a range of sulfoglycolipid analogues in which the hydroxyphthioceranoic acid has been replaced by less complex acids (Scheme 3) [42]. The synthesis started with the selective acylation of the

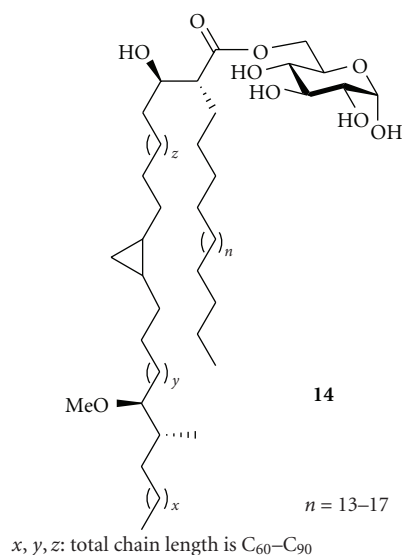


FIGURE 4: Representative glucose monomycolate (GMM).

readily available 4,6:4',6'-di-*O*-benzylidene protected α,α' -trehalose **16**, which gave the 2-*O*-acyl product **17** without the need for stannylene chemistry or elaborate protecting group manipulations. Subsequent protection of the diol in **17** with dichloro-tetraisopropylidisiloxane gave alcohol **18** which was then acylated at the 3-position with various long chain branched fatty acids. Finally, the siloxane protection in **19** was removed using TBAF and the resulting diol **20** selectively sulphated at the 2'-position to gain sulphate **21**. Final deprotection using catalytic acid provided the SGL analogues (e.g., **22**). Although none of the synthetic compounds were as potent as the natural sulfoglycolipid **15**, some of the sulfoglycolipid analogues showed promising T-lymphocyte-activation properties [42]. It was later shown that the presence of T cells that can recognise Acyl₂SGL, in patients with active or latent tuberculosis, demonstrates the potential of sulfoglycolipids as subunit vaccine against tuberculosis [43].

3.3. CD1c. To date, there is no high-resolution structural data for CD1c, but computational studies of the protein in complex with a *M. Tb* glycolipid, mannosyl- β -1-phosphomycoketide (MPM), indicate that the enlarged F' pocket can accommodate bacterial polyketides with multiply branched unsaturated alkyl tails [44]. From the structural model of CD1c with phosphomycoketide, it was hypothesised that the A' pocket does not contribute much to the specificity of the CD1c binding domain as it is similar to the A' pocket of CD1b and CD1d. Moreover, CD1c is set apart from other members of the CD1 family due to its unusual ability to present glycolipids that only have a single lipid tail.

3.3.1. Endogenous Glycolipids. Mannosyl- β -1-phosphodolichol (MPD, **23**, Figure 6) belongs to the family of glycosyl-1-phosphopolyprenols that are found in all cells [45]. They function as sugar donors in the glycan biosynthesis pathways

of eukaryotes and bacteria and are important for cell-wall assembly in prokaryotes. The length, saturation, phosphorylation, and glycosylation of the isoprenoid backbones vary in different organisms. Multicellular organisms have the longest (C_{90-100}) dolichols, while fungi and protozoa have shorter chain lengths (C_{70-90} and C_{50-65} , resp.) [46, 47].

MPD consists of a mannoside β -linked to a phosphopolyprenoid lipid via a phosphate group. Except for the proximal isoprene, the lipid contains unsaturated isoprene units linked in *trans* and *cis* configurations. The ability of CD1c, but not other CD1 proteins, to accommodate methyl branching within a single lipid indicates the specificity of CD1c for this branched motif [45].

Through the testing of various isolated and semisynthetic MPD analogues, a correlation between dolichol chain length and T-cell response has been observed. MPDs with a chain length similar to those found in *Mycobacteria* (C_{30-35}) gave the strongest T-cell responses, while no activation was observed for lipids with longer chains (C_{55-95}) [45]. This is probably because long lipid chains resemble the endogenously expressed MPD, which should not generate any immune response in the absence of infection and illustrates the capability of CD1c to distinguish between self-like and foreign-like glycolipids.

3.3.2. Exogenous Glycolipids. Mannosyl- β -1-phosphomycoketides (MPMs, **24**, Figure 7) are potent mycobacterial antigens recently isolated from *Mycobacterium avium* (**24a**) and *Mycobacterium tuberculosis* (**24b**) [44]. These phospholipids consist of a mannosyl- β -phosphate moiety similar to that in the endogenous MPD (**23**) but possess an unusual saturated isoprenoid backbone. The antigenicity of the MPM is largely attributed to the saturation of the lipid chain and the shorter length of the backbone (C_{30}) as compared to their endogenous dolichol counterparts [48]. Like MPDs, the methyl branches (*S*-configuration) of this mannosyl- β -1-phosphomycoketide help retain the lipid in the F' pocket, thus allowing for T-cell recognition and activation [49]. The synthesis of MPM **24a** was first reported by Crich and Dudkin in 2002 [50], and, more recently, MPM **24b** and analogues thereof were synthesised by Van Summeren and coworkers [51]. Crich and Dudkin (Scheme 4(a)) used a strategy whereby triflate donor **25** was coupled to the isoprenoid phosphate **26** to give mannosyl phosphate **27** which could be deprotected in a single Birch reduction to provide MPM **24a**. Van Summeren and coworkers, however, phosphorylated tetraacetylmannose **28** to give phosphate triester **29** (Scheme 4(b)). After deprotection of **29**, phosphate **30** was coupled to isoprenoid alcohol **31** to give phosphodiester **32**. Final deacetylation then gave MPM **24b**. These syntheses not only confirmed the structural identification of these naturally occurring MPMs lipids but also assist in the development of a subunit vaccine against tuberculosis [52].

3.4. CD1d. CD1d is perhaps the most widely studied CD1 protein due to its structural homology across mice and humans [6] and its importance in cancer immunotherapy.

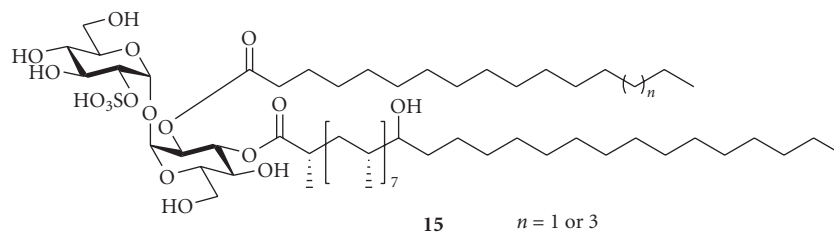
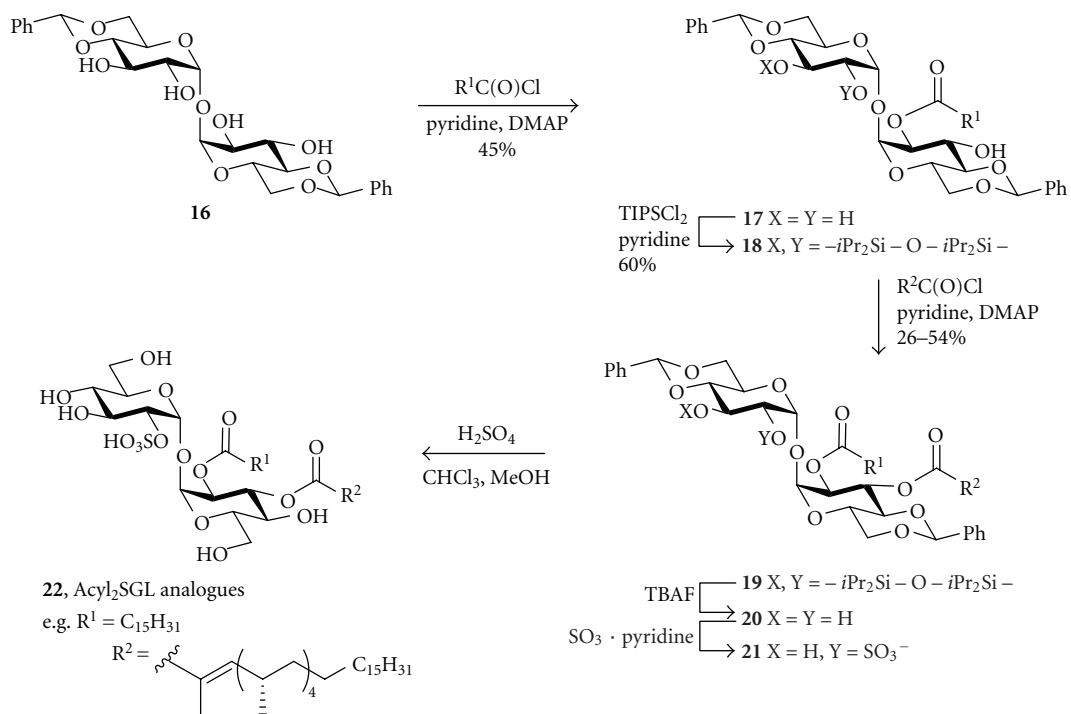
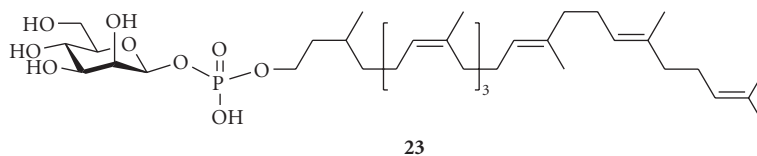
FIGURE 5: Diacylated trehalose sulphate (Acyl₂SGL).SCHEME 3: Synthesis of Acyl₂SGL analogues.

FIGURE 6: Mannosyl-β-1-phosphodolichol (MPD).

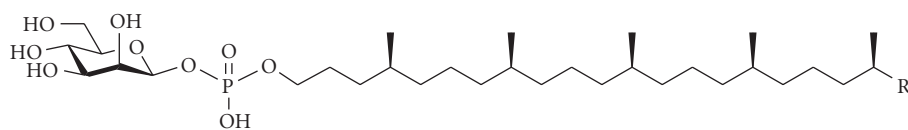
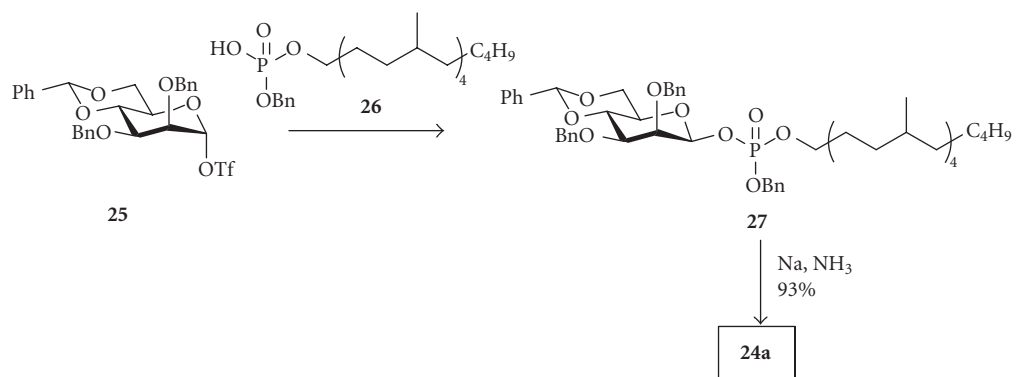
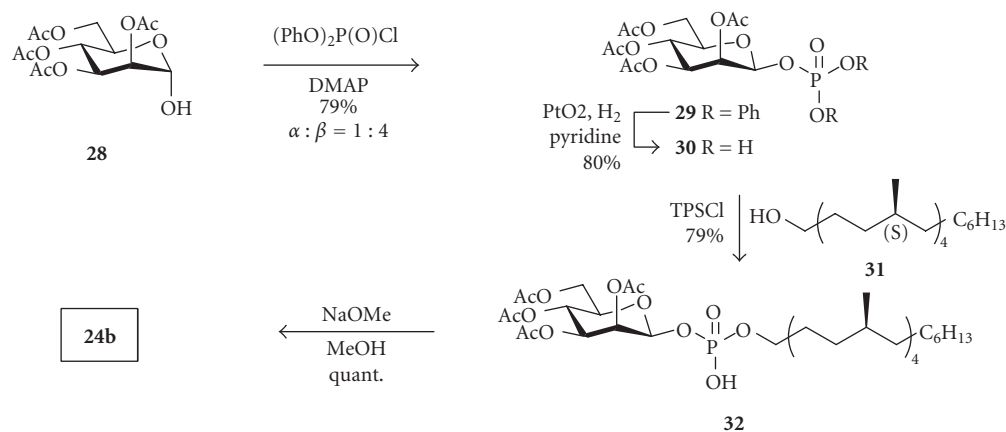


FIGURE 7: Mannosyl-β-1-phosphomycoketide (MPM).



(a)



(b)

SCHEME 4: Syntheses of MPMs.

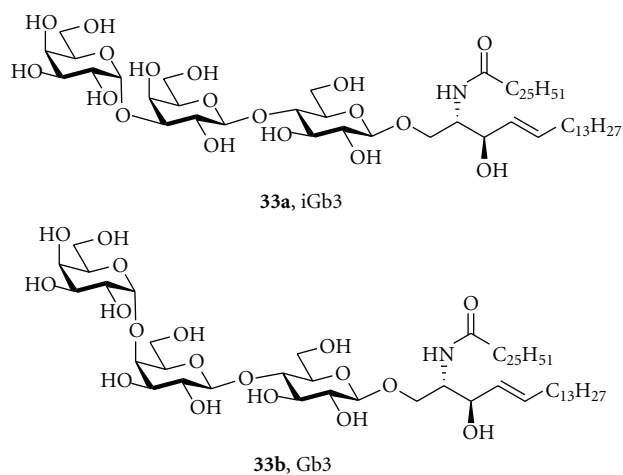
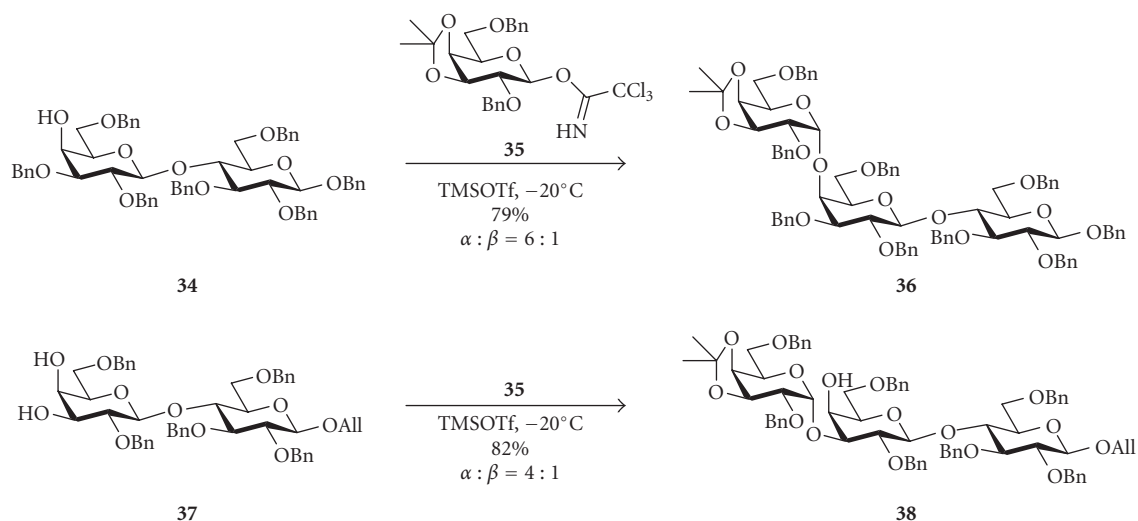


FIGURE 8: iGb3 and Gb3.

The heavy α chains form the hydrophobic binding pockets A' and F' that are able to accommodate lipid tails of glycolipids,

as illustrated by the crystal structure of CD1d bound to α -galactosyl ceramide [53]. An extensive hydrogen bonding network holds the sugar head group in place for recognition by the T cell receptor.

3.4.1. Endogenous Glycolipids. Isoglobotrihexosylceramide (iGb3, **33**, Figure 8) is a triglycosylceramide with a lactose residue β -linked to a ceramide with a sphingosine backbone and a terminal galactose residue, bound via an $\alpha(1-3)$ linkage to the nonreducing end of the lactosylceramide. iGb3 can be isolated from cat intestines but has also been chemically [54] and enzymatically [10] synthesised. Interestingly, iGb3 has been shown to play an important role in the clearing of bacterial infections [12]. This was illustrated in a study whereby mice, lacking in an enzyme that converts the biological precursor iGb4 to iGb3, were found to be deficient in CD1d-restricted T cells and subsequently incapable of clearing infection by *Salmonella typhimurium*. Also of note is that the globoside Gb3, which contains an $\alpha(1-4)$ -linked terminal galactose instead of the $\alpha(1-3)$ linkage in iGb3, does not stimulate CD1-restricted T cells [55]. This indicates that



SCHEME 5: Synthesis of globoside trisaccharides.

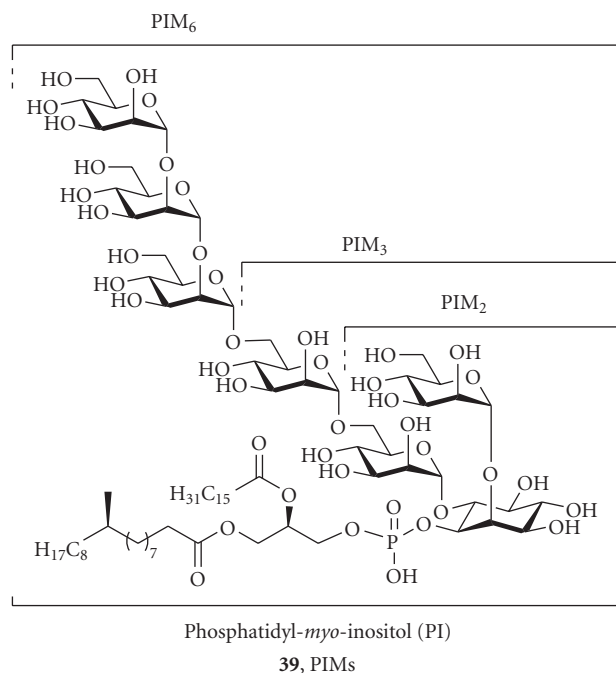
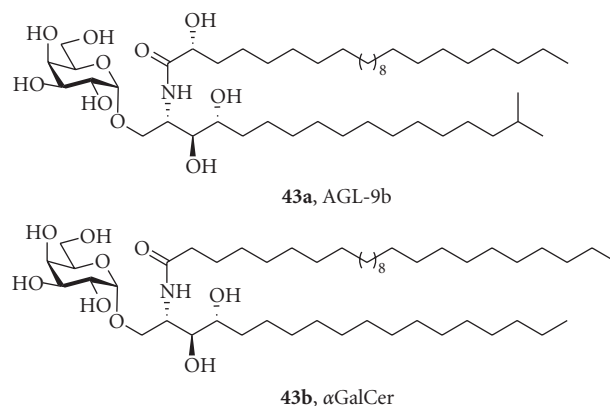


FIGURE 9: Phosphatidylinositolmannosides (PIMs).

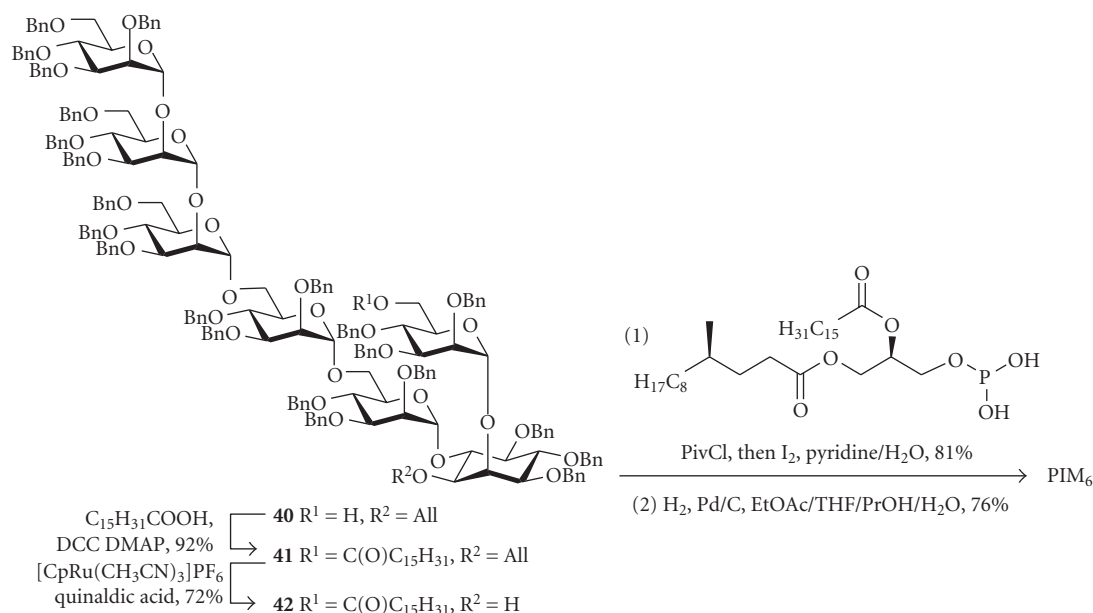
there is a high level of specificity for the interaction between CD1d-glycolipid complexes with the TCRs of CD1-restricted T cells.

Qiu and Schmidt provided convenient routes for the synthesis of globosides (Scheme 5) [54]. Here, glycosylation of lactose acceptor **34** with imidate **35** provided the isogloboside trisaccharide **36**, while regioselective glycosylation of lactose acceptor **37** with the same donor provided the isomeric trisaccharide **38**. The total syntheses of iGb3 and Gb3 were finalised by transforming **36** and **38** into their respective imidates, coupling to a sphingosine acceptor (cf. **2**, Scheme 1) and deprotection/acylation.

FIGURE 10: Agelasphin-9b (AGL-9b) and α -galactosyl ceramide (α GalCer).

Sulfatide (cf. **1**, Scheme 1). As previously mentioned, sulfatide binds to all members of the CD1 family. In the case of CD1d, Shamshiev and coworkers found that the presence of the 3-O-sulfate group as well as the β -anomeric linkage was crucial for the activation of CD1d-restricted T cells [20].

3.4.2. Exogenous Glycolipids. Phosphatidylinositolmannosides (PIMs, **39**, Figure 9) were first isolated from *M. Tb* in 1930 [56] and shown to be a natural antigen for CD1d-restricted T cells in 2004 [15]. *M. Tb* contains phosphatidyl-*myo*-inositol (PI) and PIM₁–PIM₆, the most abundant being PIM₂ and PIM₆ (Figure 9). In 1963, Ballou and colleagues characterised PIMs and deduced that, in PIM₁ the inositol residue of PI is mannosylated at the C-2 position, while, in PIM₂, PIM₁ is further mannosylated at the C-6 position of the inositol moiety. Further α (1–6) mannosylation of PIM₂ gives rise to PIM₃ and PIM₄, and the higher PIM₅ and PIM₆ motifs, which are made by consecutive α (1–2) mannosylation of the PIM₄ precursor [57].

SCHEME 6: Synthesis of PIM₆.

PIMs stimulate CD1-restricted T cells via CD1d binding and the subsequent triggering of antigen-specific IFN- γ and cell-mediated cytotoxicity [15]. The structural requirements for PIMs to bind to CD1d are the possession of two fatty acyl chains and a polar head group [58]. The increased complexity of the carbohydrate moiety does not affect CD1d binding with mycobacterial PI behaving in the same way as PIM₂ and PIM₆ [58]. Interestingly, the CD1e isoform has been shown to activate lysosomal mannosidases, which break down larger PIM structures, such as PIM₆, into PIM₂ that is then presented by CD1d [59]. PIM ether analogues are also effective in enhancing IL-12 production by immature bovine dendritic cells and thus have the potential to be used as an adjuvant [60]. The first total synthesis of the more complex PIM₆ was reported in 2006 by Liu and coworkers [61]. Pseudoheptasaccharide **40** (Scheme 6) was first acylated and ester **41** subsequently allyl-deprotected to give alcohol **42**. Alcohol **42** was then phosphorylated, oxidised, and deprotected to obtain the triacyl PIM₆.

α -Galactosyl ceramides (α GalCer, **43**, Figure 10) were first isolated from the marine sponge *Agelas mauritianus* in 1993 [62]. Of these, agelasphin-9b (AGL-9b, **43a**) and the structurally optimised synthetic α -galactosyl ceramide (KRN7000, **43b**), now widely known as α GalCer, were found to possess potent antitumour properties. The finding that these molecules formed antigens in T cell-mediated immunology led to a burst of interest in a class of CD1d-restricted T cells known as invariant natural killer T (*i*NKT) cells [63]. α GalCer consists of a galactosyl moiety α -linked to a ceramide containing a phytosphingosine base and a C₂₆ lipid connected via an amide bond. The phytosphingosine base differs from the sphingosine base found in the endogenous sulfatide glycolipids in that it has an additional hydroxyl at C-4 of the fully saturated lipid backbone.

The antitumour activities of the isolated **43a** and synthetic derivative **43b** are comparable; however, as **43b** has an extended acyl chain (extra two carbons) and lacks the hydroxyl group on the acyl chain and the methyl branch on the phytosphingosine backbone, the chemical synthesis of **43b** is more straightforward and is thus a better drug candidate.

α GalCer has been reported to have potential in the treatment of several diseases including cancer, malaria, type 1 diabetes, and multiple sclerosis [64]. The mechanism by which α GalCer exerts its therapeutic effect only became known with the discovery that it binds to CD1d and activates *i*NKT cells [65]. α GalCer is the first agonist found to activate the CD1d-restricted *i*NKT cells that express an invariant TCR α chain (V α 24J α 18 in humans and V α 14J α 18 in mice). The first chemical synthesis of α GalCer was reported by Morita et al. [63] in 1995 and subsequently optimised by others in representative syntheses such the use of thioglycoside donor using NIS/TfOH as the promoter system [66] and trichloroacetimidate galactosyl donor using boron trifluoride etherate as the activating agent [67] in the key glycosylation step. Much effort is currently being made to develop derivatives of α GalCer [68] with improved antitumour activity, particularly within the context of cancer immunotherapy.

3.5. CD1e. Unlike CD1a–d, which are expressed on the cell surface of antigen presenting cells, CD1e is primarily found in intracellular compartments (i.e., Golgi apparatus, endosome, and lysosome) and is believed to be crucial for glycolipid processing and loading onto CD1 molecules [59, 69]. In this context, CD1e has been shown to process PIM₆ to the smaller PIM₂ motif for presentation by CD1d [59].

TABLE 1: Ligands of CD1 molecules.

CD1 isoform	Endogenous ligand	References	Exogenous ligand	References
CD1a	Sulfatide	[20, 22, 23]	—	—
CD1b	GM1	[27–29]	GMM	[14, 38, 39]
	GQ1b	[31, 35, 37]	Acyl ₂ SGL	[40–43]
CD1c	MPD	[45–47]	MPM	[48–52]
CD1d	iGb3	[10, 54]	PIMs	[15, 56–61]
	Sulfatide	[20–23]	α -GalCer	[62–68]
CD1e	Involved in glycolipid antigen presentation			[59, 69]

4. Concluding Remarks

Like their protein counterparts, glycolipids can regulate the immune system in a very specific manner via binding to the CD1 family of proteins and subsequent activation of CD1-restricted T cells. Glycolipids that bind to CD1 include both endogenous glycolipids, found within the host, and exogenous glycolipids from foreign materials or pathogens (Table 1). The five members of the CD1 family (CD1a–e) show binding specificities to individual or families of glycolipids, and as such, the resulting immune response to each CD1-glycolipid complex can be used for therapies ranging from cancer vaccine adjuvants to the treatment of autoimmune diseases. As more complex biochemical tools are developed, it is almost certain that the role of CD1-binding glycolipids in the immune system becomes better understood and opens the gateway to more specific therapies and enhanced preventative vaccination protocols.

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