



Review

From Lysosomal Storage Diseases to NKT Cell Activation and Back

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Abstract: Lysosomal storage diseases (LSDs) are inherited metabolic disorders characterized by the accumulation of different types of substrates in the lysosome. With a multisystemic involvement, LSDs often present a very broad clinical spectrum. In many LSDs, alterations of the immune system were described. Special emphasis was given to Natural Killer T (NKT) cells, a population of lipid-specific T cells that is activated by lipid antigens bound to CD1d (cluster of differentiation 1 d) molecules at the surface of antigen-presenting cells. These cells have important functions in cancer, infection, and autoimmunity and were altered in a variety of LSDs' mouse models. In some cases, the observed decrease was attributed to defects in either lipid antigen availability, trafficking, processing, or loading in CD1d. Here, we review the current knowledge about NKT cells in the context of LSDs, including the alterations detected, the proposed mechanisms to explain these defects, and the relevance of these findings for disease pathology. Furthermore, the effect of enzyme replacement therapy on NKT cells is also discussed.

Keywords: NKT cells; Lysosomal storage diseases; CD1d; lipids; lysosome

1. Introduction

The lysosome, designated as the recycling compartment of the cell, was initially described by Christian de Duve in 1955 [1]. It is a membrane-enclosed organelle, characterized by its acidic pH and the presence of a large number of hydrolases. Genetic defects in lysosomal hydrolases or in other proteins necessary for the degradation or transport of macromolecules in the lysosome lead to lysosomal storage diseases (LSDs). The main feature of LSDs is the accumulation of different types of molecules in the lysosome, leading to a disturbance in lysosomal homeostasis that has important implications in autophagy, protein degradation, and metabolic stress [2,3]. The most usual classification of LSDs is based on the type of material that is accumulated. LSDs are divided in sphingolipidoses (accumulation of sphingolipids), mucopolysaccharidoses (accumulation of glycosaminoglycans), mucolipidoses (accumulation of glycoproteins) [4]. The most common LSDs are sphingolipidoses, which are usually characterized by the accumulation of glycosphingolipids (GSLs): ceramide or sphingosine molecules modified by the addition of sugar head groups. GSLs have

been implicated in important immunological processes, such as T cell activation. More specifically, GSLs were shown to be antigenic for Natural Killer T (NKT) cells, a group of lipid-specific T lymphocytes with important functions in autoimmunity, infection, and cancer [5].

2. NKT Cells

NKT cells comprise a population of T lymphocytes with lipid-specific T cell receptors (TCRs). Peptide-specific T cells recognize antigens bound to Major Histocompatibility Complex (MHC) molecules at the surface of antigen presenting cells. Instead, NKT cells recognize lipid antigens that are bound to CD1d. CD1d stands for cluster of differentiation 1 d. In humans; CD1d molecules belong to a family of 5 MHC-class I like glycoproteins with hydrophobic grooves that have affinity for lipids. They are divided into three groups. Group I includes CD1a, CD1b, and CD1c isoforms. Group II includes CD1d, and group III is composed of CD1e. Group I and group II CD1 molecules present lipid antigens to lipid-specific T cells, while CD1e has a role in the loading of lipids in other CD1 molecules. Importantly, all these molecules traffic through the endo-lysosomal compartments and therefore are likely to be affected in LSDs. This review focuses on CD1d-restricted T cells, the NKT cells, the most studied lipid-specific T cells [6].

2.1. Classification and Characterization

Two different populations of NKT cells can be distinguished based on the TCR that they express (Table 1). Type I NKT cells, or invariant NKT (iNKT) cells, are characterized by the expression of a semi-invariant TCR composed of a $V\alpha 24J\alpha 18$ chain and a $V\beta 11$ chain in humans, or a $V\alpha 14J\alpha 18$ chain paired with a limited repertoire of $V\beta$ chains in mice [7–10].

Table 1. Main differences between iNKT and type II NKT cells. NKT, Natural Killer T; iNKT, invariant
NKT; TCR, T cell receptor; CD1d, cluster of differentiation 1 d.

Feature	iNKT Cells	Type II NKT Cells	
TCR	Semi-invariant; $V\alpha 24J\alpha 18 V\beta 11$ (humans) and $V\alpha 14J\alpha 18$ (mice)	Variable; αβ or γδ	
Preferred Antigens	α-linked monohexosylceramides	Phospholipids; β-linked glycosphingolipids	
Antigen Specificity	All cells recognize the same antigen	Different cells have different antigen specificities	
Identification	CD1d tetramers loaded with specific antigen; Antibodies against semi-invariant TCR CD1d tetramers load with specific antiger		
Whole Population Identified?	Yes	No	

On the contrary, type II NKT cells express variable TCRs. However, both mouse and human type II NKT cells present a bias towards some $V\alpha$ and $V\beta$ chains, suggesting that this population has some degree of oligoclonality [9,10]. These differences in TCR expression result in distinct antigen specificities (Table 1). While iNKT cells present a preference for α -linked monohexosylceramides, most known antigens for type II NKT cells are β -linked GSLs or phospholipids [11–16]. At the moment, there are no cell surface markers that allow for the identification of all type II NKT cells. However, iNKT cells can be easily recognized by the use of antibodies against a specific region of the semi-invariant TCR (6B11 antibody) or by CD1d tetramers loaded with the antigen α -galactosylceramide (α -GalCer) or analogues [17,18] (Table 1). The use of antibodies against only $V\alpha$ 24 or $V\beta$ 11 is not recommended for iNKT cell identification, as this will also detect T cells that are not restricted to CD1d [19].

2.1.1. iNKT Cells

iNKT cells are abundant in adipose tissue, and influence the development of obesity and metabolic disorder [20–26]. They are also frequent in the mouse liver and present, although at lower amounts, in the spleen, peripheral blood, lymph nodes, bone marrow, and thymus [19]. The development of iNKT cells in the thymus requires the interaction of thymocytes expressing the semi-invariant TCR with CD1d molecules loaded with self-antigen and expressed in other thymocytes [27]. Once outside the thymus, iNKT cells are able to respond very rapidly to both TCR-mediated and TCR-independent activation, producing large amounts of a broad array of both anti-inflammatory and pro-inflammatory cytokines [28]. In humans, iNKT cells can be divided into functionally different subsets, according to the cell surface expression of CD4 and CD8 molecules. iNKT cells that do not express CD4 (which include both CD8+ and CD8- cells) are characterized by the production of similar amounts of pro-inflammatory and anti-inflammatory cytokines [18,29–31]. Importantly, CD8 expression is associated with an increase in cytolytic activity by iNKT cells, as well as an increase in the production of pro-inflammatory cytokines when compared to anti-inflammatory cytokines [31]. On the other hand, cells expressing CD4 display lower cytolytic activity and tend to produce larger amounts of anti-inflammatory cytokines and lower quantities of pro-inflammatory cytokines [18,29-31]. The expression of the NK cell marker CD161 has also been associated with a functional subset of iNKT cells that produce IL (interleukin)-17 in response to pro-inflammatory conditions [32]. In mice, iNKT cells were shown to have plasticity. Under certain conditions, peripheral iNKT cells can acquire the ability to produce IL-17 and IL-9, despite being committed to other functions when they leave the thymus [33,34].

2.1.2. Type II NKT Cells

The identification of type II NKT cells is difficult to achieve due to the lack of specific surface markers and the diversity of CD1d-loaded lipids they recognize, thereby removing the possibility of using CD1d-tetramers to identify the entire type II population. Type II NKT cells can be identified by their CD1d-restriction and the absence of the invariant $V\alpha14/V\alpha24$ TCR α -chain. Type II NKT cells were initially described by comparing MHC-deficient mice (lacking conventional T cells) with MHC/CD1d double knockouts (lacking both conventional T cells and NKT cells). These studies identified a population of T cells in the MHC-deficient mice that did not recognize α -GalCer nor expressed the invariant TCR [35,36]. Later studies analyzing type II NKT cells were done by comparing mice lacking iNKT cells (J α 18-deficient mice) with mice lacking both iNKT cells and type II NKT cells (CD1d-deficient mice). These animal models have been very useful in defining the unique role of type II NKT cells in several pathological conditions including cancer, infection, and autoimmunity (the function of type II NKT cells has been extensively reviewed elsewhere [9,10]).

In humans, type II NKT cells are very frequent in bone marrow and liver [37,38]. However, in previous studies, type II NKT cells were identified as being positive for CD3 and CD56 or CD161, and negative for V α 24J α 18 and V β 11 chains. Therefore, some non-CD1d-restricted T cells might have also been detected. Studies using CD1d tetramers loaded with specific lipids suggest that type II NKT cells outnumber iNKT cells in peripheral blood because frequencies of type II NKT cells specific for just one antigen are similar or greater for type II NKT than for iNKT cells [13,39,40]. The study of lipid-specific type II NKT cells with CD1d tetramers has also revealed other phenotypic and functional characteristics of type II NKT cells. CD1d tetramers loaded with sulfatide identified the majority of peripheral blood sulfatide-reactive NKT cells as $\gamma\delta$ T cells, which expressed the V δ 1 segment [40]. Another study using CD1d dimers loaded with lysophosphatidylcholine identified a group of $\alpha\beta$ CD3+ cells, negative for the V α 24J α 18 and V β 11 chains, in the peripheral blood of control subjects and multiple myeloma patients [39]. These cells can express CD4, CD8, or none of these molecules. Functionally, they produce several cytokines upon stimulation, including IL-13, IL-5, IL-8, and IFN- γ [39]. More recently, CD1d tetramers loaded with β -glucosylsphingosine and β -glucosylceramide identified a subset of type II NKT cells, designated NKT follicular-helper cells, that provide direct help to B cells [13].

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2.2. The Role of the Lysosome in NKT Cell Biology

Most of the studies that addressed the role of lysosomes in CD1d-mediated lipid antigen presentation focused only on iNKT cells. However, it is expected that some of the described mechanisms are also applicable for type II NKT cells.

There are mainly three lysosomal characteristics that contribute to the role of this organelle in iNKT cell activation: the content of hydrolytic enzymes capable of processing lipid antigens specific to iNKT cells; the content of lipid-transfer proteins (LTPs) that assist lipid loading in CD1 molecules; and the co-localization of CD1d molecules with lipid antigens in a low-pH compartment.

The processing of glycolipids by lysosomal hydrolases can lead both to the generation or inactivation of iNKT cell antigens. The role of lysosomal enzymes on the generation of lipid antigens was initially demonstrated for a synthetic lipid, $Gal(\alpha 1 \rightarrow 2) - \alpha$ -galactosylceramide, which upon galactose removal by α -galactosidase A (the enzyme deficient in Fabry disease), becomes an iNKT cell antigen [41]. However, it was suggested that naturally-occurring glycolipids, such as α -galactopyranosyl($1\rightarrow 2$)- α -glucosyl diacylglycerol from *Streptococcus pneumoniae*, might also need galactose removal to become antigenic [42]. Studies with Sandhoff disease mice showed that β -hexosaminidase activity was important for removing β -linked N-acetylgalactosamine from isoglobotetrahexosylceramide, yielding the iNKT cell antigen isoglobotrihexosylceramide [43]. Lysosomal phospholipase A2, responsible for the generation of lysophospholipids from phospholipids, was shown to be required for the normal presentation of endogenous antigens to iNKT cells [44,45]. However, the specific endogenous antigens processed by this enzyme were not identified [45]. Recently, lysosomal enzymes were also shown to be involved in the degradation of iNKT cell antigens through a sequential process that requires acid ceramidase (defective in Farber disease) to remove the acyl chain, and α -galactosidase A to remove the galactose residue [46].

Inside cells, lipids usually travel in membranes or in association with proteins. Lysosomal LTPs facilitate loading in CD1d molecules, either by promoting lipid removal from membranes or transfer between proteins. It has been shown that saposins, Niemann-Pick type C2 (NPC2) protein, GM2 activator protein, and CD1e can assist lipid binding to CD1d [47–53]. Saposins are important for endogenous and exogenous lipid removal and loading into mouse and human CD1d, both in the steady-state and during infection [47–50]. They perform lipid exchange, meaning that saposins are not capable of removing lipids from CD1d if they are not being replaced by another lipid. GM2 activator protein is able to remove mouse CD1d bound-lipids [48] and NPC2 protein can load lipids into mouse CD1d molecules [51]. Whether this is true for human cells has never been addressed. Finally, CD1e, which is present in human but not in mouse cells, promotes the loading and unloading of lipids on CD1d and on group I CD1 molecules [52,53]. This suggests that CD1e tunes the CD1d-mediated immune response by selecting ligands to load on CD1d molecules, consequently influencing presentation to both iNKT and type II NKT cells [53].

The lysosome is also a meeting point for CD1d molecules and lipid antigens. This is especially important for mouse CD1d molecules, which are known to localize mainly to the lysosome. In contrast, CD1d in humans is mostly targeted to late endosomes rather than lysosomes. Either way, both compartments carry lipids with long saturated tails, which may match the CD1d binding groove [27]. The loading of lipids in CD1d molecules is also facilitated by the low pH of the lysosome, which allows relaxation of the CD1d structure and consequently enables binding and dissociation of lipids [54]. Interestingly, in mice, type II NKT cells (identified as CD1d-restricted T cells that do not express $V\alpha14$ or NK1.1) do not depend on lysosomal localization of CD1d for activation [55]. Studies with tail-truncated mouse or human CD1d (not able to reach lysosomes or late endocytic compartments) showed how CD1d trafficking is important for lipid antigen presentation and iNKT cell development [50,56,57].

More recently, autophagy, a process regulated by the lysosome, was shown to be important for the development of iNKT cells [58,59], once again reinforcing the importance of the lysosome for NKT cell biology.

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3. Lysosomal Storage and NKT Cells

3.1. NKT Cells in LSDs

The study of NKT cells in LSDs was centered on iNKT cells for a long time. Table 2 summarizes the previous studies analyzing NKT cells in animal models of LSDs. The first report describing iNKT cell alterations in LSDs was published in 2001 [41]. In this paper, preliminary data suggested that iNKT cells were reduced by approximately 50% in Fabry disease mice when compared to wild-type mice [41]. A few years later, a reduction in iNKT cell number was also described in a mouse model of Sandhoff disease [43]. Initially, these defects were thought to be disease-specific. However, in the following years, other papers were published describing iNKT cell alterations in GM1 gangliosidosis [60,61], Niemann-Pick type C1 (NPC1) [61,62], NPC2 [51,60], or Tay-Sachs [61] diseases, and confirming the alterations in Fabry [61,63-65] and Sandhoff diseases [61]. It was believed that lysosomal dysfunction per se, independently of the type of substrate stored, had a negative impact on the number of iNKT cells. However, a few years later, a report described normal iNKT cell numbers in mouse models of metachromatic leukodystrophy, mucopolysaccharidosis type I, and Krabbe disease [66]. These results raised the hypothesis that the iNKT cell defects encountered in LSDs were not merely caused by lysosomal storage, but related to the accumulation/absence of a specific group of lipids/enzymes. Recently, a decrease in the frequency of iNKT cells in a mouse model of Gaucher disease was reported [13]. In Table 3, studies on NKT in LSDs patients are listed. Earlier studies in Gaucher and Fabry disease patient, analyzed "iNKT-like" cells by the use of an antibody against V α 24 TCR α -chain, and no differences were observed between patients and controls [67–69]. Subsequent studies determined the frequency of iNKT cells in the peripheral blood of Fabry, Gaucher, and NPC disease patients, and no alterations in iNKT cell frequency were detected when compared with control subjects [13,70,71]. This can be attributed to the different trafficking pathways between human and mouse CD1d. While mouse CD1d is mainly found in the lysosomes, human CD1d is often found in late endosomes [14,72]. Thus, it is reasonable to suggest that mouse iNKT cells may be more affected by lysosomal dysfunction. Another possibility is related to the analysis of distinct tissues. In mouse models of LSDs, iNKT cells were analyzed in the thymus, spleen, and liver, whereas in patients, iNKT cells were collected from the blood. Blood contains a small number of iNKT cells and it was previously shown that the frequency of iNKT cells in blood do not correlate with the percentage of thymic iNKT cells [73]. Despite the absence of alterations in the frequency of iNKT cells in LSDs patients, the analysis of iNKT cell subsets revealed disease-specific alterations. While the iNKT cell population from Fabry disease patients has a decreased percentage of cells expressing CD4 with a concomitant increase of CD4 – iNKT cells, iNKT cell subsets are unaffected in NPC patients [70,71]. Curiously, mouse models of both Fabry and NPC diseases display a reduction in iNKT cells expressing CD4 [60,64]. In the case of Fabry disease, in both mice and patients, these phenotypic alterations were accompanied by functional alterations accessory to a pro-inflammatory status that has been described in these patients [71,74]. Thus, iNKT cells might play a role in vascular inflammation leading to endothelial cell dysfunction, a feature of Fabry disease.

The analysis of type II NKT cells is complicated by the lack of specific markers that identify these cells. Nevertheless, a recent study used CD1d tetramers to identify glucosylceramide and glucosylsphingosine-specific type II NKT cells in Gaucher disease mice and patients [13]. They found that glucosylsphingosine-specific type II NKT cells were increased in both mice and patients when compared to control subjects [13]. Importantly, in Gaucher disease patients, the frequency of these glucosylsphingosine-specific type II NKT cells is positively correlated with the activity of chitotriosidase in serum, an established biomarker of disease severity [13,75]. This, together with the fact that these cells strongly communicate with B cells, suggests a possible implication of type II NKT cells in the pathology of Gaucher disease, and more specifically in the development of B cell malignancies.

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Disease	Protein Defect	iNKT Cells	Type II NKT Cells	iNKT Cell Subsets
Fabry	α-galactosidase A	↓ [41,61,63–65]	ND	↓ CD4+ iNKT cells [64]
NPC1	NPC1	↓ [61,62]	ND	ND
NPC2	NPC2	↓ [51,60]	ND	↓ CD4+ iNKT cells [60]
Sandhoff	β-hexosaminidase A/B β subunit	↓ [43,61]	ND	ND
Tay-Sachs	β-hexosaminidase A $α$ subunit	↓[61]	ND	ND
GM1 gangliosidosis	β-galactosidase	↓ [60,61]	ND	ND
Metachromatic leukodystrophy	Arylsulphatase A	= [66]	ND	ND
MPS I	α-L-iduronidase	= [66]	ND	ND
Krabbe	β-galactosylceramidase	= [66]	ND	ND
Gaucher	β-glucosidase (Glucocerebrosidase)	↓ [13]	↑ (glucosylsphingosine-specific) [13]	ND

Table 2. NKT cells in Lysosomal Storage Diseases mouse models.

^{=,} normal; \uparrow , increase; \downarrow , reduction; ND, not done; MPS I, Mucopolysaccharidosis type I; NPC, Niemann-Pick type C.

Disease	Protein Defect	iNKT Cells	Type II NKT Cells	iNKT Cell Subsets
Gaucher	β-glucosidase (Glucocerebrosidase)	= [13]	↑ (glucosylsphingosine-specific) [13]	ND
Fabry	α-galactosidase A	= [71]	ND	↓ CD4+ iNKT cells [71]
NPC	NPC1	= [70]	ND	No alterations in CD4+ iNKT cells [70]

Table 3. NKT cells in Lysosomal Storage Diseases (LSDs) patients.

3.2. Proposed Mechanisms Behind NKT Cell Alterations

Although a clear relationship between lysosomal dysfunction and alterations in the iNKT cell compartment was established more than 10 years ago, the elucidation of the mechanisms behind these alterations continues until today. Importantly, all explanations are based on events occurring at the antigen presenting cell, which lead to an alteration in the amount of CD1d molecules bound to endogenous antigens at the cell surface and consequently to an impaired iNKT cell selection or activation (Figure 1).

The first explanation was proposed together with the first description of a reduction in iNKT cell numbers in Fabry disease [41]. In this study, α -galactosidase A was shown to be necessary for the processing of a synthetic glycolipid, which upon galactose removal gains antigenic activity [41]. Even though they could not exclude other mechanisms, the lack of correct antigen processing was proposed as the mechanism behind iNKT cell reduction. By analyzing lipid antigen presentation in the Sandhoff mouse model, β-hexosaminidase was identified as being essential for the generation of isoglobotriosylceramide (iGb3), an endogenous iNKT cells antigen [43]. Therefore, in the absence of this enzyme, there is no available antigen for thymic selection of iNKT cells, leading to a reduction in their numbers. Later, a putative accumulation of iGb3 in the Fabry disease mouse model has been suggested to be the link to the overstimulation and decreased number of iNKT cells in these mice [63]. However, the accumulation of iGb3 was not addressed directly. The presence of iGb3 in the thymus was controversial, and in 2012 it was shown that, in Fabry disease mice, the storage of globotriaosylceramide (Gb3) is responsible for the reduction of both iNKT cell numbers and antigen presentation [65]. Indeed, it is now known that different antigens can select iNKT cells [46,76]. Sagiv and colleagues proposed an explanation for the iNKT cell reduction observed in NPC1-deficient mouse model [62]. They demonstrated that the abnormal lipid trafficking existent in NPC1 cells

^{=,} normal; ↑, increase; ↓, reduction; ND, not done; NPC, Niemann-Pick type C.

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prevented the co-localization of iNKT cell antigens with CD1d molecules, thus preventing the loading of antigens onto CD1d molecules [62]. Soon after, the NPC2 protein, defective in NPC2 disease, was shown to be fundamental for the loading of lipid antigens in CD1d, working as a LTP. This justified the strong reduction of iNKT cells observed in NPC2-deficient mice [51,60]. At this point, the proposed mechanisms behind iNKT cell alterations were related to the lack of antigen presentation and consequent defective selection of iNKT cells in the thymus. This idea was then challenged by the discovery that the enzyme α -galactosidase A, defective in Fabry disease, has a role in the degradation of endogenous antigens [46], suggesting that the reduction in iNKT cell numbers could be related not to a lack of selection, but instead, due to the death caused by an overstimulation potentiated by an excess of endogenous antigen. This notion was further supported by the identification of signals of chronic activation in residual iNKT cells from Fabry disease mice [63]. It is known that several GSLs have the capacity to bind to CD1d molecules [14], however, the impact of this binding in NKT cell activation is not clear. One of the explanations proposed in 2006 by Gadola and co-workers was that the high amounts of lipids accumulating in the lysosomes could easily out-compete the endogenous antigens for CD1d binding, thus leading to iNKT cell defects [61]. This hypothesis was recently proven for the GSL Gb3, which accumulates in high amounts in Fabry disease. Gb3 was identified as a CD1d ligand capable of inhibiting iNKT cell activation by competing with endogenous antigens for CD1d binding [77].

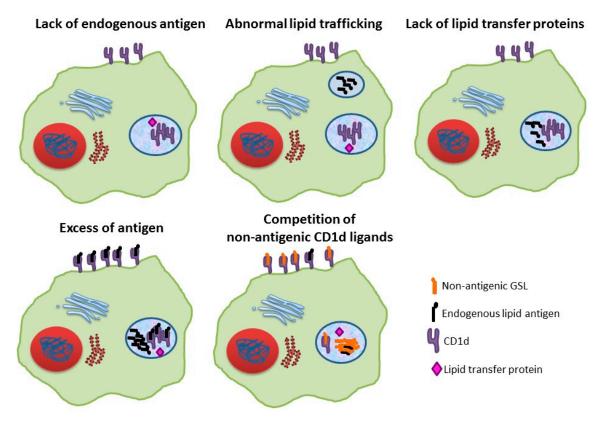


Figure 1. Proposed mechanisms, occurring in the antigen-presenting cell, that can explain invariant Natural Killer T (iNKT) cell defects in lysosomal storage diseases (LSDs). GSL, glycosphingolipid.

3.3. The Effect of Enzyme Replacement Therapy on NKT Cells

Enzyme replacement therapy (ERT) is available for some LSDs, namely Gaucher disease, Fabry disease, Pompe disease, mucopolysaccharidosis (MPS)-I, MPS-II, MPS-Iva, and MPS VI [78]. ERT consists of the infusion of a recombinant enzyme that is targeted to the lysosomes, thus recovering the function of the defective enzyme. For some LSDs, this treatment option successfully ameliorates disease pathology. Due to the properties of the blood-brain barrier, ERT is not applicable to treat

pathologies of the central nervous system that are characteristic to some LSDs. In Gaucher disease, the first disease for which ERT was available, ERT is very successful, correcting hepatosplenomegaly and thrombocytopenia and reducing bone complications [79,80]. In Fabry disease, clinical trials have shown an improvement in pain and in cardiac disease [81]. However, little is known about the effect of ERT on NKT cells.

In the Fabry disease mouse model, ERT was shown to partially prevent the decrease in the number of splenic iNKT cells [64]. However, it was not capable of correcting the defect already existent at the beginning of the ERT [64]. When the number of iNKT cells was compared between patients receiving ERT and not receiving ERT, no significant alterations were found [69]. Similarly, a longitudinal analysis starting four months after the beginning of the ERT and along a total of 24 months (with a four month periodicity) revealed no significant alterations in the frequency of iNKT cells or iNKT CD4/CD8 subsets [71]. In Gaucher disease patients, the levels of CD4+ and CD8+ T cells expressing the TCR chain $V\alpha 24$ —a population that includes, but is not limited to, iNKT cells—were similar between untreated and treated patients, suggesting that ERT does not have an influence in iNKT cell frequency or phenotype [67]. Contrarily, ERT seems to successfully reduce the increase of glucosylsphingosine-specific type II NKT cells observed in Gaucher disease patients [13]. This might be related to different antigen presentation requirements between iNKT and type II NKT cells. The type II NKT cells analyzed in this study are specific to glucosylsphingosine. Therefore, it is expected that by reducing the amount of antigen available through ERT, the glucosylsphingosine-specific type II NKT cells would decrease. iNKT cells, on the contrary, can recognize different antigens, and the defects observed in LSDs may be explained by distinct mechanisms that are not be overcome by ERT. Additional studies, targeting the characterization of iNKT cells in LSDs before and at different times of ERT and analyzing the effect of ERT on the different proposed mechanisms behind iNKT cell alterations, would be important in order to clarify this subject.

4. Conclusions

A tight relationship between NKT cells and LSDs is now established. The lysosomal impairment observed in LSDs is sufficient to cause alterations in this subset of T cells. However, at this point, the contribution of NKT cell defects to LSDs pathology is not completely understood. Nevertheless, NKT cells are known to play a pivotal role in autoimmunity, infections, and cancer, conditions for which some patients with LSDs present increased risk [82–84]. It is important to continue studying NKT cells in the context of LSDs. Ideally, these studies should be performed in human samples because of the differences in NKT cell biology between humans and mice. Another important task would be to correlate NKT cell alterations with clinical parameters in patients in order to identify new biomarkers. Finally, the study of NKT cells in LSDs is crucial to unveil the importance of the lysosome and of different lipids in NKT cell biology, thereby opening new ways to modulate NKT cell responses.

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