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1. Introduction

Sphingolipids belong to a major class of lipids that are ubiquitously synthesized by eukaryotic cells and are characterized by their sphingoid backbone and primary structural roles in membrane formation. Over the past two decades, sphingolipids have emerged as a source of important signalling molecules in addition to their structural functions (Merrill et al., 1997). In particular, sphingosine-1-phosphate (S1P), is a unique signalling molecule that has the ability to act as an intracellular second messenger, as well as an extracellular stimulus through specific G protein-coupled receptor (GPCRs) (Pyne and Pyne, 2000; Spiegel and Milstien, 2003; Van Brocklyn et al., 1998). S1P has been shown to mediate a variety of fundamental biological processes including cell proliferation, migration, invasion, angiogenesis, vascular maturation and lymphocyte trafficking. The intracellular level of S1P is tightly regulated by the equilibrium between its synthesis by sphingosine kinases (SphKs) and its degradation by S1P phosphatases (SPPs) and S1P lyase (SPL) (Saba and Hla, 2004). Activation of SphKs by a variety of agonists, including growth factors, cytokines and hormones, increases intracellular S1P. Once generated, S1P can either act intracellularly as a second messenger or be secreted out of the cell and act extracellularly by binding to and signalling through S1P receptors expressed on the surface of the same cell or nearby cells in autocrine and/or paracrine manners (Alvarez et al., 2007). Many of the biological effects of S1P are mediated by its binding to and activation of its specific receptors S1P1–S1P5 (Ishii et al., 2004). Alterations in S1P signalling, as well as in the enzymes involved in its synthesis and metabolism, have been observed in many types of pathological situations such as angiogenesis, metastasis, and autoimmunity.

The role played by S1P in the autoimmune disease rheumatoid arthritis (RA) has recently emerged. Activated SphK and elevated levels of S1P have been detected in the synovium and synovial fluids of patients with RA, respectively (Kamada et al., 2009; Kitano et al., 2006; Lai et al., 2008). In addition, exogenously applied S1P in cultured synovial fibroblasts from RA patients causes cell migration, production of cytokines/chemokines, expression of cyclooxygenase-2 and prostaglandins, as well as cell proliferation and survival (Kitano et al., 2006; Zhao et al., 2008). These results suggest a biological function for the SphK/S1P/S1P receptor (S1PR) axes in RA disease progression. In this review we summarize the current understanding of how S1P metabolism and signalling impact the biology of inflammation in
RA. We also discuss the potential therapeutic benefit of modulators of S1P metabolism and S1P receptors, including SphK/SPL inhibitors, FTY-720, S1P receptor antagonists, and anti-S1P monoclonal antibodies, in the treatment of RA.

2. S1P biosynthesis, metabolism and secretion

All cells are able to generate S1P during the normal physiologic metabolism of sphingolipids (Hla, 2004). Generally, systemic and local concentrations of S1P are tightly regulated by the balance between its synthesis and degradation via three enzyme families, including SphKs (SphK1 and SphK2), SPPs (SPP1 and SPP2), and SPL (Fig. 1). SphKs generate S1P through phosphorylation of its precursor, sphingosine, while SPPs reversibly convert S1P back to sphingosine by removing the phosphate (Xia et al., 2000). Irreversible degradation of S1P is carried out by a single enzyme SPL, which cleaves S1P into ethanolamine phosphate and hexadecenal, representing the last step in the sphingolipid degradation pathway (Xia et al., 2000).

![Sphingosine metabolism](image)

Fig. 1. S1P metabolism pathway. S1P homeostasis is tightly regulated by the balance between its synthesis and degradation via three enzyme families: (1) sphingosine kinases (SphK1 and SphK2), which generate S1P through phosphorylation of its precursor, sphingosine, (2) S1P phosphatases (SPP1 and SPP2), which reversibly convert S1P back to sphingosine, and (3) S1P lyase (SPL), which irreversibly degrades S1P to generate ethanolamine phosphate and hexadecenal.

SphK is represented by two different isoforms (SphK1 and SphK2). Although both SphK1 and SphK2 can phosphorylate sphingosine, SphK1 produces the majority of S1P in cells (Kohama et al., 1998). SphKs can be activated by a large variety of agonists, including growth factors (such as platelet-derived growth factor, vascular endothelial growth factor, epidermal growth factor and hepatocyte growth factor), cytokines (such as TNF-α), steroid hormones (such as estradiol), and GPCR ligands (such as lysophosphatidic acid (LPA) and S1P) (reviewed in (Alvarez et al., 2007; Spiegel and Milstien, 2003; Taha et al., 2006)). Following cellular activation by these agonists, SphK1 can be activated through: 1)
phosphorylation and translocation to the plasma membrane, 2) interaction with acidic phospholipids such as phosphatidic acid produced by phospholipase D (Delon et al., 2004), and 3) possible association with other proteins. All or parts of these mechanisms may be required for full activation of SphK1 (Takabe et al., 2008). Phosphorylation of SphK1 leads to its translocation to the plasma membrane where its substrate sphingosine is located, resulting in the production of S1P (Johnson et al., 2002; Pitson et al., 2005). S1P, in turn, activates specific S1P receptors present on the surface of the same cell or on nearby cells in autocrine and/or paracrine manners (Alvarez et al., 2007). The “inside-out” export of intracellularly generated S1P is crucial for many S1P functions (Takabe et al., 2008).

The mechanism by which S1P is exported from the inside to the outside of cells after synthesis is not fully understood. Several lines of evidence suggest the involvement of the ATP-binding cassette (ABC) family of transporters in S1P secretion (Kobayashi et al., 2006; Mitra et al., 2006; Sato et al., 2007). Mitra et al. (2006) revealed that the release of S1P from mast cells is regulated by ABCC1. Similarly, the ABCA1 transporter is critical for the release of S1P from astrocytes (Sato et al., 2007). In breast cancer the export of S1P mediated by ABCC1 and ABCG2 transporters is stimulated by estrogen receptor-α (Takabe et al., 2010).

Altogether, these studies suggest that members of the family of ABC transporters may be important for export of S1P out of cells.

3. S1P receptors and downstream signalling pathways

S1P evokes diverse biological functions by binding to its ubiquitously-expressed and specific cell surface receptors. So far, five S1P receptors, designated as S1P1/Edg1, S1P2/Edg5, S1P3/Edg3, S1P4/Edg6, and S1P5/Edg8, have been identified (An et al., 1997; Im et al., 2000; Lee et al., 1998; Van Brocklyn et al., 2000; Yamazaki et al., 2000). These receptors were initially named “Edg” receptors as they belong to the so-called endothelial differentiation gene (Edg) family; however “S1P receptors” is now preferred (Chun et al., 2002). S1P receptors are GPCRs that share high similarity with each other and with LPA receptors. S1P receptors exhibit variable tissue distribution, for example, S1P1, S1P2, and S1P3 are widely expressed in various tissues, whereas the expression of S1P4 and S1P5 is more restricted to cells of the immune system and nervous system, respectively (reviewed in (Sanchez and Hla, 2004)). Upon binding to S1P receptors, S1P activates downstream signalling pathways, leading to a variety of cellular responses such as proliferation, cell migration, actin cytoskeletal rearrangement, and adherens junction assembly (Kluk and Hla, 2002; Taha et al., 2004). Each S1P receptor is coupled to a specific heterotrimeric G protein (Gi/o, Gq, and G12/13), which, when activated, dissociates into its α and βγ subunits and transduces signals toward the downstream pathways. Particularly, S1P1 is coupled predominantly to Gi/0 through which it activates: 1) Rho kinase and tyrosine kinases, leading to cytoskeletal rearrangement (Garcia et al., 2001); 2) MAPK, leading to angiogenesis (Lee et al., 1999); and 3) Akt, leading to cell chemotaxis (Lee et al., 2001). On the other hand, S1P2 and S1P3 are linked predominantly to Gq and G12/13 through which they activate phospholipase C leading to Ca2+ mobilization (Ancellin and Hla, 1999) and positive or negative modulation of Rho and thus of cell migration (Sugimoto et al., 2003). More detailed information on the various signalling pathways turned on by S1P receptor activation is discussed in previous reviews (Choi et al., 2008; Rosen et al., 2009).

It is known that S1P can also act intracellularly to enhance cell proliferation and suppress apoptosis independently of S1P receptors (Rosenfeldt et al., 2001; Van Brocklyn et al., 1998). In plants and yeast, which do not express S1P receptors, for instance, intracellular S1P
regulates stomatal aperture (Coursol et al., 2005) or stress responses and survival (Jenkins and Hannun, 2001), respectively. Similarly, in mammalian cells intracellular S1P regulates calcium release independently of inositol trisphosphate formation and of S1P receptor activation (Blom et al., 2005). S1P also signals within the nucleus by binding to and inhibiting histone deacetylases HDAC1 and HDAC2, leading to the epigenetic regulation of gene expression (Hait et al., 2009). Further studies are needed, however, to reveal the intracellular second messenger functions of S1P.

4. Role of SphK/S1P/S1PR signalling in rheumatoid arthritis

4.1 Rheumatoid arthritis

Rheumatoid arthritis (RA) is an autoimmune disease characterized by chronic inflammation of the joint synovium, synovial hyperplasia and a massive infiltration of inflammatory cells into the affected joints, leading to the progressive destruction of cartilage and bone (Feldmann et al., 1996a). The inciting agent that triggers the development of RA is unknown; however, the disease is clearly an inflammatory process since the critical events in RA are largely coordinated by complex interplays of proinflammatory cytokines, chemokines, and matrix metalloproteinases produced by both synovial resident cells and infiltrating cells (Feldmann et al., 1996b; McInnes and Schett, 2007). These inflammatory mediators are involved in many pathological processes in RA. Increasing evidence demonstrates,

Fig. 2. S1P metabolism and signalling in RA synoviocytes. A range of agonists including growth factors, hormones, angiogenic factors and other stimuli activate SphKs through phosphorylation. S1P is then generated and exported out of the cells by ABC transporters to act on the same cell (autocrine) or on nearby cells (paracrine). S1P engages with its receptors (S1P1-3) to mediate a diverse array of signalling pathways, impacting fundamental biological processes that are integral to the pathogenesis of RA, such as cell proliferation and survival, cell migration, and inflammatory mediator secretion.
however, that the activation of RA synoviocytes can be maintained in the absence of inflammatory cytokines (Muller-Ladner et al., 1996), indicating the involvement of biologically active mediators other than inflammatory cytokines in RA disease progression. Emerging evidence suggests a role for S1P metabolism and signalling in various aspects of the pathogenesis of RA. Recent in vitro and in vivo studies evaluated the potential role of S1P synthesis, catabolism enzymes and S1P signalling in RA (Fig. 2).

4.2 S1P expression and SphK activity in RA synovium and synovial fluids

Several studies reveal high levels of S1P in RA synovial tissues and fluids (Kitano et al., 2006; Lai et al., 2008). Kitano et al. (2006) reported elevated levels of S1P in synovial fluids of RA patients (1078 pM/ml) in comparison to those of osteoarthritis (OA) patients (765 pM/ml). The S1P content in RA synovial fluids was significantly higher than those in serum (400 pM/ml) or plasma (100 pM/ml) from normal people. Lai et al. (2008) also measured the amount of S1P by a competitive ELISA and detected up to 17 μM S1P in RA synovial fluids, which was more than five fold higher than that found in OA synovial fluids (3 μM). The increased level of S1P could be responsible for the recruitment and the infiltration of immune cells into the synovium (see section 4.2.2). Pi et al. (2006) also showed that peripheral blood B lymphoblastoid cell lines derived from patients with RA exhibit a high level of S1P synthesis. SphK expression and activity were also demonstrated in RA. Increased SphK1 expression and activity were found in RA B lymphoblastoid cell lines, which were implicated as the underlying mechanism of impaired Fas-mediated death signalling in RA (Pi et al., 2006). More recently, SphK2 was shown to be expressed in rheumatoid synovial fibroblasts in vivo and in vitro and associated with the up-regulation of S1P production (Kamada et al., 2009). Surprisingly, FTY-720 (2-amino-2-[2-(4-octylphenyl) ethyl] propane-1, 3-diol hydrochloride), a sphingosine analog that is phosphorylated to the active metabolite FTY-720-phosphate (FTY-720-P) by SphK2 (Brinkmann et al., 2002; Mandala et al., 2002), induces apoptosis of synovial fibroblasts. This effect could be due, at least in part, to FTY-720-P-mediated degradation of specific S1P receptors (Matloubian et al., 2004). Corroborating this observation, we reported that S1P protects synovial fibroblasts from ongoing apoptosis (Zhao et al., 2008). Using the mouse model of collagen-induced arthritis (CIA), Lai et al. (2009) highlighted distinct roles for SphK1 and SphK2 in regulating cell growth and survival. The in vivo administration of SphK1 siRNA reduced inflammation whereas mice treated with SphK2 siRNA resulted in a more aggressive disease and greater secretion of proinflammatory cytokines (IL-6, TNF-α and IFN-α) by immune cells. How and why SphK1 and SphK2 exert distinct opposing roles in the regulation of inflammatory arthritis remains unclear and may be related to the location of S1P production (Maceyka et al., 2005). The role of SphK2 in chronic inflammatory diseases remains ambiguous since the selective inhibitor of SphK2 ABC249640 has been reported to reduce bone and cartilage degradation in the mouse models of adjuvant-induced arthritis (AIA) and of CIA (Fitzpatrick et al., 2011). The SphK1 signalling pathway is activated in RA (Limaye et al., 2009). In the mouse model of CIA, administration of a non-specific pharmacological inhibitor of SphKs, N,N-dimethylsphingosine (DMS), and a siRNA approach to knockdown the SphK1 isoform, markedly suppressed joint pathologies such as adjacent cartilage and bone erosion, synovial hyperplasia, and infiltration of inflammatory cells into the joint compartment (Lai et al., 2008). Moreover, SphK1 deficiency in hTNF-α transgenic mice that develop arthritis at an early age was related to less synovial inflammation and bone erosion (Baker et al., 2010). To
our knowledge the effect of specific inhibitors of SphK1, such as BML-258 (Paugh et al., 2008), has not been tested in animal models of arthritis.

At present, evidence for roles of SPPs and SPL in RA is limited. However, up-regulation of SPP2 was detected in samples of skin lesions from patients with psoriasis, a chronic inflammatory skin disease (Mechtcheriakova et al., 2007). Interestingly, elevated mRNA expression of SPP1 and SPL was observed in RA synoviocytes compared to non-arthritic synoviocytes (Zhao et al., unpublished data).

4.3 S1P receptor expression in RA synovial fibroblasts

In RA, the abnormal growth and erosive activity of synovial fibroblasts suggest that these cells are important contributors to chronic inflammation. During RA disease progression, synovial fibroblasts become hyperplastic and closely interact with infiltrated immune cells to form the aggressive pannus tissue that eventually promotes joint destruction (reviewed in (Feldmann et al., 1996a)). Synovial fibroblasts have been reported to express three of five known S1P receptors, S1P1-3 (Kitano et al., 2006; Nochi et al., 2008; Zhao et al., 2008). Expression of S1P1 in RA synovial tissues was significantly higher compared to that of OA synovial tissues (Kitano et al., 2006). Moreover, pretreatment of synovial fibroblasts with TNF-α, the well-recognized critical cytokine in RA, resulted in up-regulation of S1P3 receptor expression in synovial fibroblasts, which likely contributes to the synergistic production of inflammatory cytokines/chemokines upon subsequent exposure to S1P (Zhao et al., 2008).

4.4 S1P signalling in RA pathophysiology

4.4.1 S1P signalling in migration of synovial fibroblasts and osteoclasts

As constituents of the synovial pannus in RA, synovial fibroblasts have long been considered as key players in the aggressive destruction of cartilage and bone (Shiozawa et al., 1983). Using an in vitro wound-closing assay, we previously reported that S1P could induce synoviocyte migration (Zhao et al., 2008). S1P-induced cell migration is mediated through the activation of extracellular signal-regulated kinase-1 and -2 (ERK1/2), as well as p38 mitogen-activated protein kinase (MAPK) and of Rho kinase. S1P-driven motility of synoviocytes was mainly via S1P1 and S1P3 receptor activation since the effect of S1P was mimicked by the S1P1 receptor agonist (SEW2871) but abolished by S1P1/3 (VPC23019) and S1P3 (CAY10444) receptor antagonists.

S1P has also been shown to induce chemotaxis and regulate migration of osteoclasts and their precursors in vitro and in vivo (Ishii et al., 2009). Cells bearing the properties of osteoclast precursors express S1P1 receptors and migrate along an S1P gradient in vitro. The S1P1 agonist, SEW2871, stimulated motility of osteoclast precursors-containing monocytoid populations in vitro. In addition, osteoclast/monocyte lineage-specific conditional S1P1 knockout mice show osteoporotic changes due to increased osteoclast attachment to the bone surface, suggesting a crucial role of the S1P-S1P1 receptor axis in regulating the migratory behaviour of osteoclast precursors and bone mineral homeostasis.

4.4.2 S1P signalling in the secretion of cytokines/chemokines and lipid mediators

In the inflamed RA synovium, large amounts of pro-inflammatory cytokines and chemokines may contribute directly to cartilage and bone erosion by promoting matrix metalloproteinase (MMP) production and suppressing chondrocyte/osteoclast functions (Feldmann et al., 1996b). Previous studies from our group and others demonstrated that exogenously applied S1P could stimulate synovial fibroblasts to release various...
inflammatory mediators, including cytokines, chemokines, and prostaglandin E2 (PGE2) (Kitano et al., 2006; Zhao et al., 2008). S1P administration strongly stimulated the secretion of IL-8 (interleukin-8), IL-6, MCP-1 (monocyte chemotactic protein-1), and RANTES (regulated on activation normal T cells expressed and secreted) via S1P2 and S1P3 receptors. The signalling pathways involved in S1P-mediated cytokine/chemokine secretion were dependent on ERK1/2, p38 MAPK, and Rho kinase activation. These cytokines/chemokines may subsequently increase the recruitment of inflammatory cells such as neutrophils into the synovium, as chemokines such as IL-8 exhibit selective chemotactic activity for neutrophils, whereas MCP-1, MIP-1α (macrophage inflammatory protein-1α), MIP-1β (macrophage inflammatory protein-1β), and RANTES primarily attract monocytes (Koch, 2005). S1P may therefore contribute to the regulation and maintenance of the inflammatory response in RA, in part through stimulation of multiple cytokine/chemokine secretion by synovial fibroblasts. SphK/S1P signalling was also implicated in cell-contact-mediated proinflammatory cytokine/chemokine secretion in RA synovium without additional exogenous stimulation. Lai et al. (2008) used peripheral blood mononuclear cells from RA patients in cell-cell contact experiments. They discovered that activated peripheral T lymphocytes from RA patients induced substantial production of TNF-α, IL-1β, IL-6, MCP-1 and MMP-9 by autologous peripheral monocytes. Importantly, treatment with a potent SphK inhibitor, DMS, significantly suppressed production of these cytokines. The results suggest the importance of SphK/S1P signalling in cell-contact-mediated inflammatory mediators, which is relevant to RA pathology.

Regarding PGE2, S1P was reported to stimulate COX-2 expression and super-production of PGE2 by RA synovial fibroblasts (Kitano et al., 2006; Nochi et al., 2008). PGE2 is an autocrine lipid mediator derived from arachidonic acid metabolism by cyclooxygenases (COX-1 or COX-2) (Ghosh et al., 2004). High levels of PGE2 were detected in synovial fluids of RA patients (Hidaka et al., 2001; Lettesjo et al., 1998). PGE2 has pro-inflammatory properties and contributes to the pathogenesis of arthritis, since PGE2 stimulates angiogenesis in the rheumatoid synovium (Ben-Av et al., 1995) and triggers bone resorption by osteoclasts (Mino et al., 1998). Thus, S1P may aggravate synovial hyperplasia, inflammation and angiogenesis through the induction of COX-2 and PGE2 in RA synovial tissues. Indeed, the COX-2-PGE2 axis has been suggested to play an important role in arthritic diseases by stimulating inflammation, angiogenesis, and osteoclastic bone destruction (Martel-Pelletier et al., 2003).

4.4.3 S1P signalling in the proliferation and survival of RA synovial fibroblasts, B lymphoblastoid cells and chondrocytes

Abnormal proliferation and resistance to apoptosis of synovial fibroblasts have been suggested to contribute directly to the hyperplasia of the rheumatoid synovium (Osple et al., 2004). Indeed, S1P appears capable of increasing cell survival and inhibiting apoptosis of various cell types, including B lymphoblastoid cells derived from RA patients (Pi et al., 2006). We previously demonstrated that S1P, through S1P1, protected synovial fibroblasts from apoptosis (Zhao et al., 2008). In our study we were not able to demonstrate a significant effect of S1P on RA synoviocyte proliferation, although Kitano et al. (2006) reported a small stimulatory effect of S1P on cell proliferation. Nonetheless these studies suggest that S1P can induce the proliferation of synovial fibroblasts, which may contribute to synovial hyperplasia (Knedla et al., 2007). S1P was also reported to aggravate arthritis by inducing chondrocyte proliferation through the stimulation of COX-2 and PGE2 production and via the activation of ERK (Kim et al., 2006; Masuko et al., 2007).
4.4.4 Synergistic action of S1P and TNF-α on synovial fibroblast functions

TNF-α is central to RA pathogenesis as the pro-inflammatory cytokine not only stimulates the production of other inflammatory mediators but also directly triggers the activation of synovial cells and osteoclasts leading to the irreversible damage of soft tissues and bone (Olsen and Stein, 2004). Several studies have suggested that S1P and TNF-α might synergize in their regulation of synovial fibroblast functions. On the one hand, the expression of S1P3 receptor was enhanced by TNF-α, leading to amplified secretion of cytokines/chemokines, including IL-6, IL-8, MCP-1, and RANTES (Zhao et al., 2008). On the other hand, S1P enhanced TNF-α-mediated COX-2 expression and production of PGE_2 by RA synoviocytes (Kitano et al., 2006). Thus, the elevated levels of TNF-α seen in RA synovium may increase S1P synthesis via activation of SphKs and make synovial fibroblasts more responsive to S1P possibly through up-regulation of S1P receptor expression. The synergy between TNF-α and S1P may eventually exacerbate the clinical manifestations of RA, including enhanced synovial tissue invasion by aggressive fibroblasts (due to enhanced cell mobility), synovial hyperplasia (due to proliferation and survival of synoviocytes), and exacerbated inflammation (due to over production of inflammatory mediators).

5. Potential S1P-targeted therapy for RA

Despite great efforts devoted to the development of new therapeutic targets for RA, currently used drugs have limitations in their use for the treatment of RA. As discussed in the above sections, S1P appears to be an important modulator of many aspects of the pathogenesis of RA. Manipulation of endogenous local and systemic amounts of bioactive S1P by inhibiting metabolic enzymes such as SphKs and SPL, by applying S1P-blocking agents such as FTY-720, VPC23019 and JTE-013 or anti-S1P antibodies is thus a promising approach for the treatment of RA. Studies targeting S1P metabolism and signalling for the treatment of RA are summarized in Table 1.

5.1 Targeting S1P metabolism

5.1.1 Reducing S1P formation with sphingosine kinase inhibitors

Blockade of SphK1 activity in the mouse model of CIA significantly suppressed articular inflammation and joint destruction, reduced disease severity, and down-regulated proinflammatory cytokine production and inflammatory cell recruitment into the synovium (Lai et al., 2009). Decreasing S1P production by inhibiting SphK activity is, therefore, a promising therapeutic option in chronic inflammatory arthritis. Indeed, the inhibition of S1P synthesis by blocking SphKs activity has proven useful as an anti-inflammation strategy in cancer therapy in vitro and in animal models (Billich et al., 2005; Bonhoure et al., 2006; French et al., 2006; Gamble et al., 2006; Leroux et al., 2007).

5.1.2 Downregulation of S1P receptor activity with S1P lyase inhibitors

SPL catalyzes the irreversible degradation of intracellular S1P to hexadecenal and phosphoethanolamine. It is the major and constitutively active S1P-degrading enzyme in cells and tissues (Fig. 1). As a result, S1P concentrations in tissues are maintained at very low levels. Inhibition of SPL leads to the accumulation of S1P in various tissues (Schwab et al., 2005). Treatment with SPL inhibitor results in accumulation of S1P in lymphoid tissues.
Inhibition of SphK activity

<table>
<thead>
<tr>
<th>Approach and reagent</th>
<th>Mechanism of action</th>
<th>Target</th>
<th>Experimental system</th>
<th>Reference</th>
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<tbody>
<tr>
<td>N,N-dimethylsphingosine (DMS, SphK inhibitor)</td>
<td>Reduces S1P formation</td>
<td>SphK1 and SphK2</td>
<td>In vivo</td>
<td>Lai et al., 2008</td>
</tr>
<tr>
<td>ABC249640 (SphK2 inhibitor)</td>
<td>Reduces S1P formation</td>
<td>SphK2</td>
<td>In vivo</td>
<td>Fitzpatrick et al., 2011</td>
</tr>
<tr>
<td>siSphK1</td>
<td>Reduces S1P formation</td>
<td>SphK1</td>
<td>In vivo</td>
<td>Lai et al., 2008</td>
</tr>
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</table>

Inhibition of SPL activity

| SPL inhibitor | Downregulates S1PR activity | SPL | Clinical trial | Bagdanoff et al., 2010 |

Sequestration of S1P

| SIP specific antibody | Depletes extracellular SIP | SIP | In vivo | Visentin et al., 2006 |

Blocking S1PR

| FTY-720 (sphingosine analogue) | Down-regulates S1PRs and renders cells unresponsive to S1P | S1P1, S1P3, S1P4 and S1P5 | In vivo | Matsuura et al., 2000; Tsunemi et al.; Wang et al., 2007 |
| VPC23019 (S1P1/3 antagonist), CAY10444 (S1P3 antagonist) and JTE-013 (S1P2 antagonist) | Block SIP binding to S1PR | S1P1-3 | In vitro | Kitano et al., 2006; Zhao et al., 2008 |

Table 1. Targeting the SphK/S1P/S1PR signalling pathway in RA.

and induces premature internalization of the exit-signal-sensing S1P1 receptor on lymphocytes (Schwab et al., 2005). S1P1 receptor internalization renders lymphocytes unresponsive to S1P, preventing their egress from the thymus and lymph nodes (Lo et al., 2005). One physiological outcome of this systemic redistribution of lymphocytes is potent immunosuppression, which offers new opportunities for developing immunoregulatory agents to treat autoimmune and inflammatory diseases (Gardell et al., 2006; Huwiler and Pfeilschifter, 2008; Mandala et al., 2002; Rosen et al., 2003; Zhang and Schluesener, 2007). Indeed, SPL-deficient mice showed resistance to various inflammatory and autoimmune challenges (Bagdanoff et al., 2009; Bandhuvula et al., 2007; Vogel et al., 2009). The evaluation of a synthetic SPL inhibitor, LX2931, is currently underway in phase II clinical trials in RA patients (Bagdanoff et al., 2010).
5.1.3 Depletion of extracellular S1P with S1P antibody
Specific antibodies against S1P have been developed and are currently being tested in clinical studies for the treatment of cancer, fibrosis, inflammation, macular degeneration, diabetic retinopathy, glaucoma, and other diseases (Graler and Goetzl, 2004). The antibodies are thought to bind S1P and reduce the extracellular pool of bioactive S1P (Sabbadini, 2006). Indeed, a preclinical study using blocking S1P antibodies to prevent tumor progression was recently reported (Visentin et al., 2006). In this hallmark study, a specific monoclonal antibody recognizing S1P was administered to mice harboring human cancer xenografts. The intervention reduced, and in some cases completely eliminated, tumor formation and accompanying tumor angiogenesis. The results suggest that antibody-mediated inhibition of S1P signalling may be developed as a strategy for inhibiting pannus formation and angiogenesis in RA.

5.2 Targeting S1P receptors
S1P and its receptors are powerful regulators of various critical functions of RA synoviocytes. Targeting these cells with specific S1P receptor agonists or antagonists can modulate specific inflammatory responses locally and systemically by altering cell migration, cytokine and chemokine secretion, and proliferation/survival. RP-001, a potent S1P1 receptor agonist with properties similar to those reported for FTY-720 on S1P1 receptor degradation and lymphocyte sequestration from the blood, was recently characterized (Cahalan et al., 2011). The short half-life of RP-001 in vivo, however, disqualifies this compound for clinical use.

5.2.1 S1P receptor agonists, FTY-720
The immunosuppressant drug FTY-720 (fingolimod) was originally described as a sphingosine analogue that is phosphorylated in vivo by SphK2 to a S1P agonist for all S1P receptors except S1P2 (Brinkmann et al., 2002; Mandala et al., 2002). Further studies revealed that FTY-720-P not only activates S1P receptors, it also down-regulates them and consequently renders cells unresponsive to S1P (Matloubian et al., 2004). FTY-720 has been suggested to play a major role in autoimmune disease, such as multiple sclerosis [reviewed in (Nicholas et al., 2011)]. Multiple sclerosis is a common neurological disability, in which autoreactive T-cells migrate across the blood–brain barrier and attack myelin sheaths, leading to demyelination and axonal damage. FTY-720 deprives thymocytes and lymphocytes by downregulating S1P receptors and interfering with S1P signalling necessary for their egress from secondary lymphoid tissues (Cyster, 2005; Graler and Goetzl, 2004; Kappos et al., 2006; Matloubian et al., 2004). The results of a Phase II clinical trial evaluating the efficacy and safety of FTY-720 for treating relapsing multiple sclerosis showed that the annualized relapse rate of the FTY-720 group was significantly decreased (Kappos et al., 2006). FTY-720 was recently FDA-approved for the treatment of multiple sclerosis (Strader et al., 2011).

The effects of FTY-720 have been examined in several animal models of arthritis (Matsuura et al., 2000; Tsunemi et al., 2010; Wang et al., 2007). In the study of Wang et al. (2007), CIA rats were treated daily with FTY-720 for 28 days and the arthritis index was measured. Radiological analysis revealed that FTY-720-treated CIA rats had less joint damage in comparison to untreated CIA rats. Moreover, while histological assessment showed that CIA rats suffered from inflammatory cell infiltration and synovial hyperplasia in their joints,
FTY-720 treatment clearly reduced these pathological parameters. Similarly, Matsuura et al. (2000) compared FTY-720 with two other anti-rheumatic compounds, mizoribine and prednisolone, in the rat models of CIA and AIA. FTY-720 completely suppressed the increase in hind paw volume and bone destruction to normal control levels by inhibiting leukocyte accumulation in the arthritis joint. Moreover, FTY-720 was shown to possess anti-arthritic activity with a wider margin of safety in AIA and CIA models as compared to mizoribine and prednisolone. A recent study by Tsunmi et al. (2010) revealed that FTY-720 administration suppressed the progression of laminarin-induced arthritis in SKG mice. FTY-720 treatment decreased IL-6 and TNF-α expression by synovial fibroblasts, diminished the number of inflammatory cells migrating into the joints, and suppressed bone destruction.

5.2.2 S1P receptor antagonists
Blocking S1P receptor activity by using S1P receptor antagonists may lead to promising therapeutic strategies for patients suffering from RA. Despite the lack of more comprehensive in vivo data, considerable progress has been made in the identification of the S1P receptors regulating synovial fibroblast migration, production of cytokines/chemokines and PGE₂, proliferation, and survival. Several compounds targeting S1P receptors have also been used to decipher the biological roles of S1P receptors. The S1P1/3 antagonist VPC23019 and S1P3 antagonist CAY10444 blocked S1P-mediated synoviocyte migration and cytokine secretion, whereas the S1P2 antagonist JTE-013 attenuated S1P-mediated cytokine secretion (Kitano et al., 2006; Zhao et al., 2008). The pharmacokinetics, bioavailability and metabolic characteristics of these S1P antagonists are essential to advance in vivo studies and for therapeutic intervention. Indeed, pharmacological approaches have been developed to block the action of S1P in the context of cancer progression (reviewed in Peyruchaud, 2009).

6. Conclusion
In summary, SphKs and S1P/S1PR signalling appear to play essential roles in modulating RA pathogenesis. The SphK1 pathway is activated and likely plays a pro-inflammatory role in mouse models of inflammatory arthritis. It is fascinating that the blockade of SphK1 activity results in the simultaneous reduction of several inflammatory responses such as pro-inflammatory cytokines and inflammatory cell infiltration into the synovium. Excessive S1P and enhanced S1P receptor expression are detected in the synovium of RA patients. S1P and signalling through S1P receptors induce expression of inflammatory cytokines and suppress apoptosis of B lymphoblastoid cells and fibroblast-like synoviocytes. Although the possible mechanism by which S1P exerts its activity in RA remains to be fully characterized, further understanding of S1P metabolism and S1P receptor expression by synovial tissues represents an exploitable objective for the development of novel chemotherapeutic agents in the treatment of RA.

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8. References


Bagdanoff, J. T., et al., 2010. Inhibition of sphingosine 1-phosphate lyase for the treatment of rheumatoid arthritis: discovery of (E)-1-(4-((1R,2S,3R)-1,2,3,4-tetrahydroxybutyl)-1H-imidazol-2-yl)ethanone oxime (LX2931) and (1R,2S,3R)-1-(2-(isoxazol-3-yl)-1H-imidazol-4-yl)butane-1,2,3,4-tetraol (LX2932). J Med Chem. 53, 8650-62.


Ospelt, C., et al., 2004. Synovial activation in rheumatoid arthritis. Front Biosci. 9, 2323-34.


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The purpose of this book is to provide up-to-date, interesting, and thought-provoking perspectives on various aspects of research into current and potential treatments for rheumatoid arthritis (RA). This book features 17 chapters, with contributions from numerous countries (e.g. UK, USA, Canada, Japan, Sweden, Turkey, Bosnia and Herzegovina, Slovakia), including chapters from internationally recognized leaders in rheumatology research. It is anticipated that Rheumatoid Arthritis - Treatment will provide both a useful reference and source of potential areas of investigation for research scientists working in the field of RA and other inflammatory arthropathies.

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