Targeting PI3K Isoforms and SHIP in the Immune System: New Therapeutics for Inflammation and Leukemia

Matthew D. Blunt and Stephen. G. Ward

Inflammatory Cell Biology Laboratory,
Department of Pharmacy and Pharmacology,
University of Bath,
Claverton Down,
Bath, BA2 7AY

Running Title Page:
Targeting PI3K signaling in the immune system

Corresponding author:
Prof. Stephen G. Ward,
Inflammatory Cell Biology Laboratory,
Department of Pharmacy and Pharmacology,
University of Bath,
Claverton Down,
Bath, BA2 7AY,
U.K.

E-mail address: S.G.Ward@bath.ac.uk
Tel: +44 (0) 1225 383641
Abstract:

PI3K is critical for the normal function of the immune system, however dysregulated PI3K mediated signalling has been linked to the development of many immune mediated pathologies. This review describes current progress in the development of isoform-specific PI3K inhibitors that hold promise for the treatment of hematopoietic malignancies as well as for inflammatory and autoimmune diseases. A SH2-domain containing inositol-5-phosphatase (SHIP) is a regulator of PI3K signalling, and is also discussed as a potential drug target for immunomodulation and the treatment of leukemia. Recent progress has been made in the development of small molecule compounds that potently and selectively modulate SHIP activity and hence provide a novel mechanism to alter PI3K mediated signalling.
**Introduction**

Activation of class 1 phosphoinositide 3-kinase (PI3K) occurs following ligation of a plethora of leukocyte receptors that are responsible for innate and adaptive immune responses [1-3]. Activation of PI3K results in the addition of a phosphate group to its lipid substrate PI(4,5)P$_2$, forming PI(3,4,5)P$_3$ in the cell surface membrane. PI(3,4,5)P$_3$ initiates a variety of signalling cascades by interaction with pleckstrin homology (PH) domain containing proteins, most notably the serine kinase Akt [4].

Class 1A isoforms (PI3K$\alpha$, PI3K$\beta$ and PI3K$\delta$) are activated by receptor tyrosine kinases, whilst the Class 1B isoform PI3K$\gamma$ is activated following GPCR ligation. However, these definitions are proving less rigid than previously thought. For example, PI3K$\beta$ has been shown to signal downstream of GPCR activation [5], whilst activation of the Class 1B isoform PI3K$\gamma$ has been reported by receptor tyrosine kinases and TLR/IL-1Rs [6]. PI3K$\gamma$ has also been reported to participate in T-cell-receptor signalling [7]. The PI3K$\alpha$ and PI3K$\beta$ isoforms appear to have a broad tissue distribution, while PI3K$\delta$ and PI3K$\gamma$ are primarily expressed in leukocytes. Studies using mice in which the genes encoding PI3K$\delta$ or PI3K$\gamma$ have been either altered to encode kinase-inactive mutants (e.g. PI3K$\delta^{D910A}$ mice) or deleted, have revealed that PI3K$\delta$ and PI3K$\gamma$ have non-redundant but often co-ordinated, functions in B cells, T cells, NK cell, neutrophils, mast cells and dendritic cells [1-3,8,9]. Indeed, when the immune system of these mice is challenged they exhibit severely defective responses to infection [1-3].
Development of PI3K inhibitors to target the immune system: lessons from PI3Kδ

The strong evidence of a role of dysregulated PI3Kγ and PI3Kδ in immune-mediated pathologies, has prompted intense efforts to develop selective inhibitors of these isoforms. The first compounds to block PI3K—the natural product wortmannin and Lilly’s Ly294002 were reported in the early 1990’s and served as useful experimental tools to research the PI3K pathway [2]. This is evidenced by nearly 6000 PUBMed hits for Ly294002 alone. However, the usefulness of wortmannin and Ly294002 both as research tools and potential drugs was tempered by their broad targeting of all PI3K isoforms and off-target effects [10]. However, the simple planar structure of Ly294002 (Figure 1) has aided the design of more selective pan-PI3K and isoform specific inhibitors. In particular, crystal structures of PI3Ks bound to Ly294002, have facilitated huge advances in the design of PI3K inhibitors which utilize the regions of the ATP binding pocket to achieve greater potency and selectivity as well as reduced toxicity [11-13]

Compounds that selectively inhibit PI3Kγ have been identified, with a series of compounds designed by Merck Serono SA based on the thiazolidinedione scaffold. One of these, AS605240 (Figure 1), exhibited superior potency compared to related compounds, can be administered orally and has high cell membrane permeability [14]. However, the subsequent development of PI3Kγ inhibitors for the treatment of inflammatory disorders in humans has been largely disappointing. This is most likely due to the selectivity achieved for PI3Kγ over other class 1A PI3K isoforms, being insufficient to avoid off-target effects. In contrast, the development and implementation of inhibitors which target the PI3Kδ isoform has proven more successful. The discovery of the quinazolinone purine series, exemplified by the ICOS compound IC-87114 (Figure 1) demonstrated that the design of isoform selective PI3K inhibitors with at least 50-fold potency over other isoforms was possible to achieve [15].
In 2006 several members of ICOS Corporation formed a spin-out company, Calistoga Pharmaceuticals. Calistoga developed CAL-101, a PI3Kδ specific inhibitor that exhibits 40-300 fold selectivity over other PI3K isoforms. CAL-101 which was acquired by Gilead in February 2011 and recently renamed GS-1101) has shown success in clinical trials for treatment of B cell malignancies where it causes rapid lymph node shrinkage and lymphocytosis. CAL-101 displays a dual mechanism of action whereby it both decreases cell survival and reduces interactions that retain CLL cells in protective tissue microenvironments [16,17]. This inhibitor has therefore demonstrated an essential role for PI3Kδ in constitutive PI3K signaling that is required for the survival of malignant B cells. Oncogenic mutations of components of the PI3K signaling pathway are infrequent in B cell malignancies. A potential mechanism for PI3K activation in this setting is tonic antigen-independent B cell receptor (BCR) signaling that requires PI3Kδ for the transduction of proliferation and survival signals.

Inhibition of the PI3Kδ isoform for the treatment of inflammatory disorders is also being explored. Specifically, CAL-101 and CAL-263 have entered clinical trials for allergic rhinitis (Table 1). In addition, patents have been filed by several other companies (Amgen, Intellikine and Incyte) describing PI3Kδ inhibitors and the majority are based on the same basic pharmacophore identified by ICOS [18]. However, additional scaffolds have now been reported by several companies; almost all of these are with intended indications against B cell lymphomas [18].

The non-redundant and often coordinated roles of PI3Kδ and PI3Kγ in immune cell function described earlier, provide a rationale for targeting both isoforms simultaneously with a single compound. TargeGen described two diaminopteridine-diphenol-based compounds with good selectivity for PI3Kγ and PI3Kδ that showed early promise in animal models of myocardial ischemia as well as asthma and chronic obstructive pulmonary disease [19,20]. The TargeGen
compounds did not progress beyond phase I/II clinical trails. Infinity and Intellikine are currently in pre-clinical trials with IPI-145 (Table 1), the only PI3Kγ/δ inhibitor currently in development for the treatment of inflammatory disease [18].

There is an increasing appreciation of a role for PI3Kβ in the immune system including cooperation with PI3Kδ in the generation of reactive oxygen species (ROS) in neutrophils in response to fungal infection or immune complexes [21,22]. This provides a rationale for targeting PI3Kβ as well as PI3Kδ in the treatment of inflammatory disorders. Indeed, compounds with dual selectivity for PI3Kβ and PI3Kδ have been reported suggesting that this approach is feasible [13]. However, caution should be applied to the use of PI3Kβ inhibitors in inflammatory disorders due to the described role of PI3Kβ in thrombus formation and circulatory homeostasis [23].

**SHIP-1 as an alternative target for modulation of PI3K signaling.**

*Role of SHIP in the immune system*

The development of isoform specific PI3K inhibitors for inflammatory disease, particularly toward PI3Kγ, has been disappointing. This has lead to the search for alternative targets with which to modulate PI3K signalling specifically in the immune system. In this regard, attention has recently focussed on the lipid phosphatase SH2-domain containing inositol-5-phosphatase (SHIP), which de-phosphorylates PI(3,4,5)P₃ at the D5 position of the inositol ring to create PI(3,4)P₂. Multiple forms of SHIP have been reported, but the 145 kDa SHIP-1 (Figure 2) is a particularly ideal target for development of potential therapeutics for treating immune disorders. This is because its hematopoietic-restricted expression would limit the impact of SHIP-1 targeted drugs to the immune system. One would predict for example, that activators of SHIP-1 would lead to a reduction of cellular PI(3,4,5)P₃ levels and hence, mimic the effect
of PI3K inhibitors. SHIP-1 was initially considered as a negative regulator of PI3K signalling. However, the SHIP-1 product PI(3,4)P₂ is itself recognized by some PH domain-containing proteins [24]. Thus SHIP-1 can act as a molecular ‘switch’, re-directing PI3K signalling away from PI(3,4,5)P₃-dependent effector proteins, towards proteins which bind exclusively or partially to PI(3,4)P₂.

SHIP-1 is recruited to the surface membrane following ligation of a variety of receptors including, chemokine, antigen, co-stimulatory and cytokine receptors as well as IgG engagement [25]. SHIP-1 knockout mice have proven invaluable in identifying the crucial role of SHIP-1 in the regulation of mast cell degranulation, BCR signaling and autoantibody production, dendritic cell function and NK cell cytolytic function (Table 2). SHIP-1 also regulates TLR signaling [26,27,28], lymphocyte polarization and migration [29] and has a pivotal role in regulating the balance between pro-inflammatory and anti-inflammatory myeloid and lymphoid cells [30,31].

The Role of SHIP-1 in leukemia

Over-activation of PI3K-dependent signalling cascades is a common occurrence in many human cancers [32]. The lipid phosphatase PTEN which also negatively regulates PI(3,4,5)P₃ accumulation by de-phosphorylating the D3 position of the inositol ring, is a well characterized tumour suppressor gene [33]. Likewise, evidence for mutations of SHIP-1 have also been shown in acute lymphoblastic leukaemia [34] and in acute myeloid leukaemia [35]. SHIP-1 is targeted by miR-155 in B cells, with high levels of miR-155 and reduced SHIP-1 expression linked to the development of acute lymphoblastic leukaemia in mice [36]. miR-155 levels were also found to be significantly increased in human patients with diffuse large B cell lymphoma [37]. The loss of SHIP-1 has been shown to promote the development of erythroleukemia, with SHIP-1 identified as a target gene of the oncogene fli-1 [38]. The loss
of SHIP-1 has been shown to promote the development of erythroleukemia, with SHIP-1 identified as a target gene of the oncogene fli-1 [38]. The role of SHIP-1 as a tumour suppressor is also evident in the ability of SHIP-1 to restrict myeloid suppressor cells and regulatory T cells [30,39]. Therefore the loss of SHIP-1 expression/ function may lead to increased suppression of T-cell mediated anti-tumour immunity. Indeed, in murine pancreatic cancer SHIP-1 expression was shown to be reduced in splenocytes which also correlated with an increase in myeloid suppressor cell numbers [40]. Decreased SHIP-1 expression has also been shown in myelodysplastic syndrome progenitor cells, where over-expression of SHIP-1 inhibited myeloid leukemic growth [41].

The role of SHIP-1 in leukemia however, appears more complex than initially thought. For example, SHIP-1 was shown not to act as a tumour suppressor in myeloma cells [42]. The use of a small molecule SHIP-1 inhibitor demonstrated that catalytically active SHIP-1 is required for the survival of multiple myeloma cells [43] and that therefore, in certain cases, SHIP-1 actually supports cancer cell survival. This would be consistent with increased levels of the SHIP-1 enzymatic product PI(3,4)P₂ promoting Akt activation and survival/proliferation [44]. Indeed another group has shown that SHIP-1 inhibits CD95/Fas-mediated apoptosis of T cells [45].

**Pharmacological Targeting of SHIP**

**Allosteric SHIP-1 activators:**

The identification and description of an activator of SHIP-1, termed pelorol, was first described in 2005 and was derived from the marine invertebrate *Dactylospongia elegans* [46]. Aquinox Pharmaceuticals designed more potent synthetic analogies AQX-MN100 and AQX-016A (Figure 1) which along with PI(3,4)P₂ were shown to allosterically enhance catalytic
activity by binding to the C2 domain of SHIP-1 (Figure 2). AQX-016A and AQX-MN100 showed potent inhibition of immune cell activation *in vitro* and were anti-inflammatory *in vivo* using mouse models of endotoxemia and acute cutaneous anaphylaxis [47]. Intriguingly, these SHIP-1 activating compounds increased apoptosis of multiple myeloma cells *in vitro* and when used in combination with bortezomib (an established multiple myeloma treatment) proved more effective at inhibiting cancer cell proliferation than bortezomib alone [48]. Aquinox Pharmaceuticals are developing SHIP-1 activating compounds based on the structure of pelorol, to be used in inflammatory disorders. AQX-1125 is the most advanced and passed Phase 1 clinical trials in 2011, with Phase IIa clinical studies initiated in late 2011 for the treatment of mild and moderate asthma (Table 1).

**SHIP-1 inhibitors**

A novel small molecule selective inhibitor of SHIP-1, termed 3 α-aminocholestane (3AC, Figure 1) has also recently been identified using high throughput screening, although the site of action is currently unclear [43]. Consistent with observations from SHIP-1 deficient mice, treatment of mice with 3AC led to increased numbers of myeloid suppressor cells and reduced ability of peripheral lymphoid tissues to prime myeloid-associated responses and protected against Graft-versus-host disease [43]. The inhibition of SHIP-1 using pharmacological compounds may therefore offer the potential to aid transplant acceptance in patients undergoing transplant surgery. 3AC also increased levels of granulocytes, red blood cells, neutrophils and platelets in mice and could therefore, have potential to improve blood cell number in patients with myelodysplastic syndrome or myelosuppressive infection.

Remarkably, SHIP-1 inhibition using 3AC induced the apoptosis of human AML cell lines which is consistent with SHIP-1 being anti-apoptotic under some circumstances [43]. Further studies showed that 3AC inhibited multiple myeloma cell growth in a tumour xenograft
model in mice [49]. Since both substrate [PI(3,4,5)P₃] and product [PI(3,4)P₂] of SHIP-1 have been shown to influence Akt activation and cell survival, this may explain in part, why both activators and inhibitors of SHIP-1 have shown efficacy against leukemic cells [50].

**SHIP-2 inhibitors**

SHIP-2 shares 35% homology with SHIP-1, and is thought to be involved in type-2 diabetes and obesity [51]. Its ubiquitous expression means that it is also expressed in leukocytes and one concern is that inhibition of SHIP-1 will be compensated for by SHIP-2. In recent years progress has been made in the development of small molecule compounds which inhibit the catalytic activity of SHIP-2 [52]. Another group has identified a novel biphenyl 2,3’4,5’,6-pentakisphosphate [BiPh(2,3’,4,5’,6)P₅], which showed potent inhibition of SHIP-2 catalytic activity [53]. This biphenyl compound is however, not cell permeable and does not possess drug-like properties. Nevertheless, the crystal structure of the phosphatase domain of SHIP2 bound to BiPh(2,3’,4,5’,6)P₅ has recently been reported. Molecular dynamics simulations suggest that when BiPh(2,3’,4,5’,6)P₅ binds to SHIP2, a flexible loop folds over and encloses the ligand [54]. Compounds targeting such a closed conformation might therefore deliver SHIP2-specific drugs. Recently, pan-SHIP-1/2 inhibitors have been reported to kill multiple myeloma cells [49]. The identification of both pan–SHIP and SHIP-2 inhibitors therefore provides a useful tool in the identification of the less well-characterized role of SHIP-2 in immune cell function.

**Summary**

The β, γ and δ isoforms of PI3K have important non-redundant roles in multiple cells of the immune system. Consequently alterations of the PI3K signaling pathway can lead to inflammatory and autoimmune disorders as well as leukemia. This, together with growing appreciation of the crystal structure of the catalytic isoforms which help define the structure-
activity rules for obtaining selectivity, will spur the continued design and development of improved PI3K inhibitors. These offer opportunities to manipulate the PI3K signaling network in immune cells for inflammation and transplantation as well as cancer. The latter may include non-leukemic cancers, given the up-regulation of PI3Kγ and PI3Kδ in some forms of non-immune cell malignancies [32,55,56]. One concern would be that it might be difficult to avoid effects on the immune system that could impair the endogenous anti-tumor response.

The difficulties of developing PI3Kγ inhibitors with sufficient selectivity over PI3K isoforms has in part, led to the exploitation of endogenous and leukocyte-restricted regulators of PI3K signaling, namely the lipid phosphatase SHIP-1. Small molecule regulators of this protein have shown early promise in inflammatory, transplantation and cancer settings, and are currently in phase I clinical trials to evaluate the safety, tolerability and pharmacokinetics (Table 1). However, the targeting of SHIP-1 (particularly with inhibitors), is not without its problems. For example, SHIP-1 deficiency leads to a number of pathologies including fibrotic lung disease [57], osteoporosis [58] and the development of Crohn’s disease-like ileitis [59]. Another point to take into account when pharmacologically targeting SHIP-1, is the fact that SHIP-1 has important non-catalytic functions [60]. Therefore inhibition of the catalytic site of SHIP-1 may not be sufficient to inhibit SHIP-1 mediated modulation of PI3K activity in all cases.

References:


** This study refutes current dogma that PI3Kgamma is activated only by GPCRs by demonstrating that myeloid cell PI3Kgamma is unexpectedly activated by tyrosine kinases and Toll-like/IL1-receptors. Whereas GPCRs activate PI3Kgamma in a Ras/p101-dependent manner, receptor tyrosine kinase and Toll-like/IL1 receptors directly activate PI3Kgamma in a Ras/p87-dependent manner. These studies reveal that PI3Kgamma is a single convergent point controlling tumor inflammation and progression.


** Describes the first crystal structures of the catalytic subunit of PI3Kδ. Revealing the structure of this enzyme provides exciting new insights into the behavior of the PI(3)K enzymes, and particularly into the reasons for isoform selectivity of small-molecule inhibitors. This study provides the first detailed structural insights into the active site of a class IA PI3K occupied by non-covalently bound inhibitors and suggests mechanisms to increase the potency of inhibitors without sacrificing isoform selectivity and also how to optimize solubility, pharmacokinetics/metabolism and pharmacodynamic behavior.


** Using tumor cell lines and primary patient samples representing multiple B-cell malignancies, this study demonstrated that constitutive PI3K pathway activation is PI3Kδ-dependent. CAL-101 blocked constitutive PI3K signaling, resulting in decreased phosphorylation of Akt and other downstream effectors, an increase in poly(ADP-ribose) polymerase and caspase cleavage and an induction of apoptosis. These effects were been observed across a broad range of immature and mature B-cell malignancies, thereby providing an impetus for subsequent clinical evaluation of CAL-101.

** This study reveals that PI3Kδ inhibition by CAL-101 disrupts crosstalk between the CLL and its microenvironment in several ways: (i) CLL chemokine receptor function and signaling are modulated by CAL-101, causing diminished leukemia cell chemotaxis and migration beneath marrow stromal cells; (ii) CAL-101 impairs CLL cell viability, both by disrupting BCR signaling but also by antagonizing support from nurse-like cells and by interrupting paracrine secretion of chemokines by CLL cells; (iii) CAL-101 also reduced the exaggerated production of other chemokines and cytokines that occurred when CLL cells
were co-cultured with nurse-like cells.


** This exciting study provides the first evidence that PI3Kβ plays an essential, non-redundant role in the efficient activation of mouse neutrophils by IgG-containing immune complexes. LTB4 is produced in response to FcγR ligation and, following its release, signals in an autocrine or paracrine manner through its GPCR BLT1 to enhance immune complex-induced ROS production. Intriguingly, this signalling loop in response to FcγR ligation was not abolished in PI3Kγ-deficient neutrophils, despite the known role for PI3Kγ downstream of BLT1. In fact, the response to co-stimulation with LTB4 and immune complexes was largely dependent on PI3Kβ. The unique ability of PI3Kβ to mediate signalling downstream of both tyrosine kinase-linked receptors and GPCRs may allow this isoform to integrate signals from FcγRs and BLT1.


*SHIP has a key role in regulating CD4 T cell differentiation via its capacity to promote Th17 and limit Treg development. In vitro and in vivo analyses of SHIP null mice revealed that in the absence of SHIP, T cells have an enhanced capacity to develop into Tregs and a parallel decrease in Th17 cell development. These data suggest that modulation of the PI3K pathway is central to T cell-mediated immune regulation and suggest that pharmacological manipulation of SHIP activity may represent a new strategy for manipulating the balance between Tregs and Th17 cells.*


*Demonstrates a role for SHIP as a repressor of Th2 skewing. This occurs, at least in part, by SHIP inhibiting IL-3– and IgE-induced IL-4 production from basophils. SHIP may therefore be a novel regulator of type 2 immune responses via its repression of basophil activation.*


33. Hollander MC, Blumenthal GM, Dennis PA: **PTEN loss in the continuum of common cancers, rare syndromes and mouse models.** *Nat Rev Cancer* 2011, **11**:289-301.


** A high throughput screening strategy identified a SHIP1 selective inhibitor that is capable of increasing myeloid immunoregulatory cell numbers and function in vivo, impairing the ability of peripheral lymphoid tissues to prime allogeneic T cell responses, enhancing blood cell production in myelosuppressed hosts and promoting apoptosis of blood cancer cells.

44. Manning BD, Cantley LC: AKT/PKB signaling: navigating downstream. *Cell* 2007, **129**:1261-1274


** The authors demonstrate that a small molecule allosteric activator (AQX-MN100) of SHIP is sufficient to prevent growth and induce cytotoxicity of multiple myeloma cell lines, while having no significant effects on non-hematopoietic cells lacking SHIP. AQX-MN100 also augments the effects of the established agents dexamethasone and bortezomib. These results provide the basis for the further study of small molecule SHIP activators to improve multiple myeloma patient outcomes.


* The authors describe a novel inhibitor of SHIP-2 catalytic activity. This compound provides an extremely useful tool for investigating the role of SHIP-2 in immune cell function and the potential of targeting SHIP-2 for therapeutic purposes.


* Reports the structure of the phosphatase domain of human SHIP2 in complex with BiPh(2,3′,4,5′,6)P5 at 2.1 Å resolution. Together with results of molecular dynamics simulations, this work suggests a rationale for the design of inhibitors of SHIP2.


  * Identifies a mechanism for the regulation of PI3K by SHIP1 that occurs by the docking of SHIP1 at an immunoreceptor ITAM, which directly prevented the recruitment of PI3K to the signaling complex. This mechanism may be particularly important for regulation of tonic ITAM signals or to facilitate the generation of inhibitory signals that are needed to prevent excessive innate immune responses.


Figure Legends:

**Figure 1**: Chemical Structures of PI3K inhibitors and SHIP-targeting compounds.

**Figure 2. Structure of SHIP-1**
SHIP-1 possesses a centrally located 5’ phosphatase catalytic domain. A SH2 domain at the N-terminus and NPXY motifs at the C-terminus [25]. SHIP also has a C2 domain adjacent to the catalytic domain which, when bound to PI(3,4)P2, acts to allosterically enhance the catalytic activity of SHIP [47].
<table>
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<tr>
<th>Compound</th>
<th>Company</th>
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<th>Phase</th>
<th>Status</th>
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Table 2. Impact of SHIP-1 Gene Targeting on Leukocytes

<table>
<thead>
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<th>Cell Type</th>
<th>Phenotype of SHIP-1 KO</th>
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<tr>
<td>Mast cell</td>
<td>Degranulation with IgE stimulation alone [61]</td>
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<tr>
<td>T cell</td>
<td>Increased regulatory T cell differentiation, decreased Th17 development [30]</td>
</tr>
<tr>
<td></td>
<td>Enhanced Th1 differentiation and CD8 cytotoxic activity. Decreased Th2 differentiation [62]</td>
</tr>
<tr>
<td>Basophils</td>
<td>SHIP(^{-}) mice show increased Th2 skewing due to increased IL-4 secretion from basophils [31])</td>
</tr>
<tr>
<td>B cell</td>
<td>Btk membrane association increased. Hyper-responsive to cross-linking of BCR [63]</td>
</tr>
<tr>
<td></td>
<td>Loss of anergy, production of auto-antibodies [64]</td>
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<tr>
<td>Myeloid cell</td>
<td>Increased myeloid suppressor cell numbers [39]</td>
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<tr>
<td></td>
<td>Increased M2 macrophage skewing (indirect mechanism via increased IL-4 secretion from basophils) [65]</td>
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<tr>
<td></td>
<td>Increased ratio of PI(3,4,5)P(_3) to PI(3,4)P(_2) on phagosomal membrane.</td>
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<td></td>
<td>Decreased early NADPH oxidative activity in phagosomes [66]</td>
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<tr>
<td>Dendritic cell</td>
<td>Enhanced survival and proliferation, but impaired maturation [67]</td>
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<td></td>
<td>Reduced nitric oxide production; SHIP null DC’s suppress T cell proliferation [68]</td>
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<tr>
<td>Natural Killer cells</td>
<td>Deficient receptor repertoire. Defective IFNγ secretion. Increase in peripheral number. Defective cytolytic function. [69].</td>
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