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COMMENTARY: IL-4 AND IL-13 RECEPTORS AND SIGNALING

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Abstract

Interleukin (IL)-4 and IL-13 were discovered approximately 30 years ago and were immediately linked to allergy and atopic diseases. Since then, new roles for IL-4 and IL-13 and their receptors in normal gestation, fetal development, neurological function and in the pathogenesis of cancer and fibrosis have been appreciated. Studying IL-4/-13 and their receptors has revealed important clues about cytokine biology and lead to the development of numerous experimental therapeutics. Here we aim to highlight new discoveries and consolidate concepts in the field of IL-4 and IL-13 structure, receptor regulation, signaling and experimental therapeutics.

Keywords

interleukin-4; interleukin-13; signaling; regulation; receptor

Introduction

Interleukin (IL)-4 and IL-13 were among the first cytokines described in the early 1980s. The discovery and study of their structure, receptors and signaling laid the groundwork for much of our understanding of cytokine signaling networks, receptor complexing, immune function and powerful nature of cytokines in causing and ameliorating disease. IL-4 and IL-13 are structurally and functionally related cytokines not only involved in immune function but also in pregnancy, fetal development, mammary development and lactation, as well as in higher brain functions including memory and learning (1). Perhaps more widely known is the role of IL-4 and IL-13 in the pathogenesis of atopy, asthma, pulmonary fibrosis, and cancer.

IL-4 and IL-13 are short four α -helix bundle secreted glycoproteins with about 25% sequence similarity, encoded by adjacent genes that share several *cis*-acting transactivating

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regulatory regions. Although they share receptor subunits and signaling molecules, these two cytokines elicit overlapping but also unique biological responses. IL-4 and IL-13 are secreted by Th2-polarized T cells, granulocytes and monocytes/macrophages. In the last decade, new innate immune cell populations have been identified, including type 2 invariant natural killer T cells (iNKT2) and innate lymphoid type 2 cells (iLC2) in mice and CRTH2+ type 2 ILC in humans which secrete high levels of IL-13 but only secrete IL-4 under certain circumstances (2, 3). Every cell in the body has the potential to respond to IL-4 or IL-13, or both. In the context of the immune system, IL-4 and IL-13 trigger Th2 T cells differentiation, M2 macrophage polarization, MHCII expression, B cell and plasma cell differentiation and antibody isotype switch among others. Classically considered “anti-inflammatory” cytokines by virtue of their ability to inhibit type 1 inflammation (IFN- γ , IL-12, NO), IL-4 and IL-13 do not trigger immunological senescence, rather, they initiate potent type 2 inflammatory processes. IL-4 and IL-13 signal through cell surface receptor heterodimers composed of 3 possible subunits. These 3 subunits along with a myriad of cell signaling mediators, transcription factors and regulatory elements combine to mediate the critical physiological as well as pathophysiological manifestations of these cytokines. Herein we aim to highlight new discoveries in the field of IL-4 and IL-13 receptors and signaling as well as take a prospective look at the future of targeting this pathway to improve the human health.

Overview of IL-4/13 Receptor Assembly and Signaling

IL-4 signals through both the type I receptor composed of the IL-4R α and common gamma chain (γ C), and the type II receptor composed of the IL-4R α and IL-13R α 1. IL-4 binds IL-4R α in a species-specific manner with very high affinity ($K_D = 20\text{--}300$ pM), leading to the recruitment of either the common γ C or IL-13R α 1. The γ C and IL-13R α 1 subunits have lower, approximately equal affinity for the IL-4:IL-4R α complex ($K_D = 500$ nM). Therefore, the availability of each chain on the cell surface determines the signaling pathway activated within the responding cell (4, 5). While the IL-4R α and IL-13R α 1 chains are widely expressed at low levels on most cell types, the γ C chain is primarily expressed on hematopoietic immune cells. IL-4 binding the type I receptor complex phosphorylates JAK1/3, which in turn phosphorylates tyrosines within the IL-4R α cytoplasmic domain. These phospho-tyrosine residues create docking sites for STAT6 or/and IRS-2 (Figure 1). Although STAT3 and other STATs can also be activated in some cell types, STAT6 and IRS-2 are acknowledged as the primary pathways involved in IL-4/13 responsiveness. STAT6 tyrosine phosphorylation promotes phospho-STAT6 homodimerization, nuclear translocation and gene transcription. Tyrosine phosphorylation of IRS-2 leads to activation of PI3-K (6–8), AKT (9–12) and NF- κ B-driven (13–15) gene transcription, activation of the cell cycle, proliferation and survival (16, 17).

In addition to serving as a subunit of the type II IL-4 receptor, IL-13R α 1 is the ligand binding subunit for IL-13. IL-13 binds IL-13R α 1 with moderate affinity ($K_D = 30$ nM) relative to IL-4:IL-14R α interaction, leading to recruitment of the IL-4R α subunit. Binding of IL-13 to the type II receptor results in activation of JAK1 or JAK2/TYK2, STAT6, STAT3 and STAT1, STAT dimerization and nuclear translocation occurs, followed by activation of gene transcription (Figure 1). Since the binding affinity of IL-4 for the IL-4R α

is markedly higher than IL-13 for the IL-13R α 1, IL-4 and IL-13 may compete for availability of the ligand binding subunits in cells expressing the type II receptor. Consequently, the concentration of IL-4 vs. IL-13 in the extracellular milieu is an important determinant of signaling through the type II receptor in non-hematopoietic cells.

IL-13 also binds the cell surface IL-13R α 2 “non-signaling” subunit. IL-13R α 2 is believed to be an inhibitory subunit of the type II receptor by acting as a “decoy” receptor for IL-13 (18). However, there is mounting evidence this may not be the case and is addressed below. In addition to cell surface-associated receptor subunits, soluble forms of the IL-4 receptor chains exist (19–21). Soluble IL-13R α 2 has been identified in mice (22, 23) but not humans (24). The discovery of soluble receptors provided important cues for cytokine neutralization therapies being investigated in pre-clinical and clinical studies (25–27).

IL-4 and IL-13 receptor expression and regulation in health and disease

IL-4 and IL-13 receptor expression in cancer

The IL-4 and IL-13 receptor subunits are expressed at low levels under homeostatic conditions and are influenced by hormones, cellular/oxidative stress, infection and inflammation (28–32). In solid tumors, such as breast cancer, high IL-4R α expression is associated with increased cancer cell proliferation, epithelial invasion and more aggressive metastasis (33–35). Venmar *et al.* recently showed in human breast cancer, as well as in mouse models of metastatic breast cancer, that attenuating IL-4R α expression reduced tumor survival as well as metastatic potential by blunting AKT, ERK and mTOR signaling (36). These findings agree with the observations of others showing that phospho-STAT6, downstream of IL-4/-13 receptor engagement, regulates pro-metastatic behaviors, such as proliferation, migration and tissue invasion (37–39). In Hodgkin/Reed-Sternberg lymphoma (HL), transformed B-cells highly expressed IL-13R α 1, secreted IL-13 and displayed over-phosphorylation of STAT6 (40). In these cells, blocking IL-4R α and inhibiting STAT6 activation increased sensitivity to chemotherapeutics (40). In blood/bone marrow cancers such as HL, increased IL-13R α 1 (or IL-4R α) expression may serve as a biomarker for predicting how aggressive a cancer may be as well as provide a method to monitor effectiveness of cancer therapy.

IL-4 responsiveness in the brain

Traditionally the brain is regarded as a site of immune privilege, separate from the rest of the immune system. As such, little attention has focused on investigating the role of the IL-4/-13 receptors in maintaining brain function and homeostasis. It has long been noted that T cell deficient mice have cognitive impairments and reduced adult neurogenesis, which is rescued by the adoptive transfer of Th2 polarized T cells (41–43). Aging in humans and mice is associated with decreased cognitive function that is inversely correlated with inflammation (44–46). This combined with the observations that neurological disorders such as multiple sclerosis (MS) (47, 48), Alzheimer’s disease (AD) (49, 50) and Parkinson’s disease (PD) (51) are associated with increased type 1 inflammation, suggests IL-4/-13 play an important regulatory role in the brain. Rodent models have revealed brain cells are not only sensitive to IL-4 through the expression of the IL-4 receptors, but can also produce IL-4, which is

critical for optimal cognitive function and memory formation (43). Two macrophage-like cell types, astrocytes and microglial cells, not only express the IL-4R α (but not the γ_c , suggesting type II receptor signaling) (52) but also produce IL-4 and possess an M2-like phenotype (53). These cells subsequently influence neuronal cells through the production of growth factors such as brain-derived neurotrophic factor (43), nerve growth factor (54, 55) and regulating pro-inflammatory mediators such as interferon- γ , IL-1 β , IL-6 and nitric oxide (54, 56). IL-4 induced M2 skewing of microglial cells alleviates some elements of AD pathology in rodent models and promotes amyloid- β phagocytosis (57, 58). IL-4 skewing of microglia is also critical to regulate neuro-inflammation in murine experimental autoimmune encephalitis (EAE), a model of human MS (53). Adoptive transfer of M2 polarized macrophages attenuated disease symptoms (59, 60). Evidence for the importance of IL-4 in balancing autoimmune neuro-inflammation and age-associated inflammation is strong. However most of the effects of IL-4 in the brain have been restricted to astrocytes and microglia. This view was recently challenged with the finding that following CNS injury, Th2-derived IL-4 bound to the IL-4R α on neurons. This binding not only prevented cell death but actually promoted neuronal re-growth, thus preserving cognitive function (61). This finding challenges those of other studies that suggest IL-4 in the brain solely functions to attenuate type 1 inflammation indirectly through astrocyte and glial cells. Rather it suggests IL-4 may act directly as a cyto-protective cytokine in neurons themselves. These findings are exciting and highlight several aspects of neuro-immunology that merit further investigation, particularly in the context of brain development, injury and stroke.

In contrast to evidence that the IL-4/-13 signaling axis positively influences brain function, increased expression of IL-4R α and IL-13R α 1 has been implicated in neurodegenerative disorders notably Parkinson's and Alzheimer's disease. While the exact mechanisms remain unclear, the increased expression of IL-4/13 receptor subunits is believed to cause inappropriate neuronal sensitivity to IL-4/-13 (62–65). Studies in mice revealed neurons co-express the dopaminergic receptor and the IL-13R α 1, the same neurons that are lost in the progression of Parkinson's disease (65). In this case, the IL-13R α 1 reportedly potentiated oxidative stress and cell death. It is important to note that many of these findings indicate that the IL-4/-13 receptors do not drive these disease processes alone (65). Another stressor, be it oxidative stress, a toxicant or inflammation, must also be present to drive pathology. These findings challenge our previous understanding of immune function in the brain and bring attention to a previously unappreciated role for IL-4 in maintaining homeostasis. Certainly IL-4 receptor subunits and IL-4 are necessary for normal brain development and maintaining an M2-reminicent environment, however, we are reminded that the brain exists in an exquisitely delicate balance. IL-4 signaling, while effective at attenuating type 1 inflammation, is none-the-less pro-inflammatory and aberrant receptor expression/signaling may in fact contribute to disease pathogenesis.

These new studies correlating specific IL-4/-13 receptor subunits in brain function, repair and disease in the absence of robust mechanistic studies, while very interesting, are still in their infancy. There are still many aspects of neurological IL-4 receptor expression, regulation and signaling that remain to be examined. Studying many of these aspects pose significant technical challenges and the use of animal models will be limiting. However,

advances in cell isolation techniques, *in vivo* imaging and the development of small molecules that penetrate the blood-brain barrier suggest renewed effort into understanding immune function within the brain and central nervous system could yield important insights into neurological diseases and elucidate novel therapeutic targets. Importantly findings highlight the current lack of understanding of the mechanisms that regulate receptor expression. Moreover, it is not yet clear if these changes are the driving force behind the disease or a downstream effect of other dysregulated pathways.

Molecular regulation of receptor subunit expression

Several stimuli that increase IL-4/-13 receptor subunit expression have been described. However, the mechanisms that regulate expression of each subunit and thus regulate responsiveness to IL-4/-13 remain largely undefined. A newly identified mechanism that regulates expression of IL-4R α under homeostatic conditions is the binding of Hepatocyte growth factor-regulated tyrosine kinase substrate (Hrs) to IL-4R α . Hrs binding to the cytoplasmic tail of surface-bound IL-4R α targets the complex to the late endosome, thereby limiting cell surface accumulation of the receptor (Figure 2A). This process is independent of ligand (cytokine) binding, ubiquitin-independent and appears to be a common regulatory mechanism for regulating homeostatic expression levels of other cytokine receptors including the IL-2R β (66, 67). Wei *et al.* reported a novel mechanism by which the IL-4R α expression is regulated by STIP1 homology and U Box containing protein (STUB)1 in response to cytokine stimulation (68). The authors showed that STUB1 interacts with the IL-4R α and facilitates IL-4R α ubiquitination and proteasomal degradation (Figure 2B). Importantly, they show that this is a cytokine-inducible event and a critical negative regulatory process in the development of allergic asthma. While they do not address whether STUB1 expression itself is induced by IL-4/-13 stimulation, it provides mechanistic insight into how cells attempt to regulate IL-4 signaling through the down regulation of the high affinity IL-4R α subunit. These findings highlight an important aspect of IL-4R regulation that has been largely overlooked. It has long been appreciated that MHC II expression is regulated by endosomal receptor internalization under steady-state as well as pathological conditions. However, this has not translated into the field of cytokine receptor regulation until this recent publication. This may reflect the technical challenges associated with low IL-4 receptor expression and the need to rely on cell culture systems for basic discovery.

Given these new discoveries in cancer and brain function, alongside those already described in asthma and atopy, we hope to see increased importance placed on understanding the dynamics and regulation of expression of the IL-4 receptor subunits between the cell surface and intracellular compartments. In-depth studies will provide meaningful insight to deciphering the mechanisms that regulate receptor subunit expression, localization of IL-4 receptor subunits and termination of signaling events in disease phenotypes.

IL-13R α 2 comes of age, more than just a cytokine decoy receptor

The IL-13R α 2 has greater than 35% homology to the IL-13R α 1 and binds IL-13 with extraordinarily high affinity [$<10^{-15}$ M, (18)], although does not bind IL-4. Primarily regarded as a “decoy” receptor for IL-13 because of its lack of cytoplasmic tail signaling motifs and high affinity for IL-13, the pathophysiological and therapeutic potential of this

receptor chain has garnered attention in recent years. Structurally, the IL-13R α 2 does not contain a signaling tail and does not associate with the other known IL-4 receptor subunits. IL-13R α 2 is expressed at the surface of the cell before it is release in its soluble form as well as within the intracellular compartment (69). *In vivo* analyses report conflicting anti-asthmatic/inflammatory properties (70–72) and pro-asthmatic/inflammatory properties (73–76). The IL-13R α 2 is predominantly expressed on structural cells, such as epithelial cells, but has also been identified on fibroblasts (77) and in soluble form in the circulation in mice only (24, 78–80). Recently, it was reported that TNF α alone or in synergy with IL-17 increases expression of IL-13R α 2 on normal human lung fibroblasts (81, 82). In fibroblasts, IL-13R α 2 binds IL-13 but does not initiate the typical IL-13 signaling cascades. Rather, there are reports that the IL-13R α 2 regulates IL-4 signaling. Andrews *et al.* reported IL-13R α 2 attenuated IL-4 and IL-13 mediated STAT6 signaling in a human airway epithelial cell line, BEAS-2B (83). They further show that the cytoplasmic domain of the IL-13R α 2 is required to attenuate IL-4-mediated signaling but not IL-13-mediated signaling, which is universally blocked by overexpression of the IL-13R α 2. These findings support previous findings that IL-13R α 2 binds IL-13 with high affinity to block STAT6 signaling while only partially attenuating IL-4-driven STAT6 signaling (82, 84). Furthermore, they suggest that the IL-13R α 2 cytoplasmic domain may inhibit IL-4 signaling by preventing dimerization with a functional secondary chain or possibly by blocking recruitment of the IL13R α 1 (Figure 2C). While these studies strongly support the idea that the IL-13R α 2 does not signal to promote IL-4-driven inflammation, compelling studies from *in vivo* models suggest IL-13R α 2 is more than just a decoy receptor for IL-4/-13 signaling. Fichtner-Feigl *et al.* reported that the IL-13R α 2 is induced by TNF α resulting in increased IL-13 signaling through the IL-13R α 2 to promote AP1-driven TGF- β production in monocytes (Figure 2D) (85). This enhanced IL-13 signaling along with downstream TGF- β production promoted lung remodeling and fibrosis. These findings were further supported in a mouse model of chronic allograft rejection that demonstrates that IL-13R α 2 expression promotes TGF- β production by graft-infiltrating cells, pro-fibrotic changes in gene expression and collagen deposition in the heart (86). Both these publications conclude that since IL-13 expression was increased and increased expression of IL-13R α 2 co-localized with fibrosis and TGF- β , IL-13 was driving the fibrotic process. While this does indeed seem plausible, recent findings published by He *et al.* suggest this may not be the only mechanism by which IL-13R α 2 initiates signaling. The authors report IL-13R α 2 binds Chitinase 3-like 1 (Chi311, YKL-40) to promote MAPK, Akt/PKB, and Wnt/ β -catenin signaling leading to decreased oxidant injury and apoptosis/pyroptosis (Figure 2E) (87). They also noted activation of anti-bacterial defenses, inhibition of the inflammasome, stimulation of TGF- β production and increased lung metastasis in mouse models. They suggest that IL-13 can complex with Chi311 to bind IL-13R α 2. Taken together with the two studies discussed above, there is compelling evidence that the IL-13R α 2 may indeed have yet undefined binding partners and can act alone as well as synergistically with IL-13. This possibility may help us begin to understand the apparently contradictory findings for IL-13R α 2 in animal models and humans. Closer examination of the diseases, models and cytokine milieu may elucidate the function of the IL-13R α 2. It is not yet clear how the short 17 amino acid tail of IL-13R α 2 with no apparent signaling motifs is able to stimulate such profound, diverse biological effects. Several of these responses resemble the biological functions of IL-4/-13-stimulated

cell signaling, although not all of them can be attributed to IL-4/-13. Presumably these effects occur through differential association with other signaling receptors or signaling adaptor proteins, but this remains to be validated. These results are interesting nonetheless, in that they provide unique insight into the biological function of Chi311/YKL-40. Several studies suggest that YKL-40 is an inflammatory biomarker because it predicts severity of several diseases including asthma (88–91), cancer (92), neurodegenerative diseases (93) and glioblastoma multiforme (94–96), in spite of its largely unknown biological function. These developments finally help to reconcile the *in vivo* observations that IL-13R α 2 is pro-inflammatory with *in vitro* findings that in response to IL-13, the IL-13R α 2 has inhibitory activity. We can now appreciate that in the context of *in vivo*, inflammation, IL-13R α 2 is more than a simple decoy receptor and the aforementioned studies help to refine our understanding of the IL-13R α 2 in disease pathogenesis.

IL-4/-13 receptor polymorphisms in human disease

Polymorphisms in the promoters of the genes for *IL4*, *IL13* and the IL-4 receptor subunits have previously identified. Polymorphisms that both enhance and diminish the expression or activity of the gene products have been reported in various disease states. A single nucleotide polymorphism (SNP) in the *IL4* gene promoter, C590T has been shown to be associated increased *IL4* transcriptional activity and positively correlated with arthritis (97, 98) multiple sclerosis (MS) (99–101) and asthma (102, 103). Two SNPs in the *IL13* gene, C1055T in the promoter region and R130Q in exon 4 were reported to be associated with enhanced IL-13 biological activity and progression of pulmonary fibrosis (104), chronic obstructive pulmonary disease (COPD) (105) and asthma (106, 107). In addition, three SNPs in the *IL4RA* exons, I50V, S478P and Q551R, have been widely associated with hyper-IgE and atopy (102, 103, 108–110). SNPs in *IL4*, *IL13* and the receptor subunits are reportedly associated with glioma (111, 112). These polymorphisms highlight key features of IL-4/-13 receptor signaling that regulate disease susceptibility and severity, i.e. ligand binding, signaling and its regulation. Many functional consequences of these polymorphisms remain unexamined yet may reveal new therapeutic targets to diminish unwanted inflammation and identify unappreciated regulatory pathways. The I50V SNP in the extracellular domain of the IL-4R α has been highly correlated with many atopic diseases, allergies and asthma (106, 113). Several studies undertook a methodical examination of the functional consequences of this SNP on IL-4 ligand binding to the IL-4 receptor complexes and downstream signaling. Ford *et al.* demonstrated that this SNP caused not only prolonged IL-4 binding to the receptor but also prolonged STAT6 activation (113, 114). A transient increase in *CIS* mRNA expression, a negative regulator of cytokine signaling and member of the SOCS family, was noted in cells expressing the I50V form of the receptor in response to IL-4 stimulation (113). Enhanced STAT6 signaling further correlated with increased systemic IgE in patients with Graves Disease (114). These studies provide important insight into the effect of an SNP in the extracellular domain of the IL-4R α chain on intracellular signaling responses and biology. They also demonstrate that under normal “healthy” conditions, IL-4 signaling induced expression of its own negative regulator highlighting that regulatory mechanisms are in place to overcome a single polymorphism. Of course, these studies remind us that many diseases do not arise from a single change but rather represent one alteration in a combination of many changes. These kinds of investigations focused on the functional

consequences of SNPs provide insights into the molecular mechanisms and regulation of IL-4 signaling pathways, as well as identify altered regulatory mechanisms under disease conditions. Furthermore, studies that aim to dissect the precise mechanistic and biological consequence of the SNPs must be coordinated with large-scale correlative analyses in order to better define parameters of investigation and identify appropriate endpoints.

IL-4 and IL-13 also serve an important immunoregulatory role. In the context of T-cell-driven autoimmune inflammation such as arthritis, the IL-4R α plays an immunosuppressive role. Two major IL-4 receptor polymorphisms, the I50V and Q551R SNPs, have been linked with the susceptibility and severity of rheumatoid arthritis (RA) (115–118). Wallis *et al.* and Liepe *et al.* provided modest evidence that disease severity and IL-17 secretion is associated with the prevalence of the V allele (115, 119). Furthermore, they showed that IL-4 had little effect on modulating IL-17 production in patients with pre-existing RA or its secretion from already Th17 polarized cells. Their data suggest that the suppressive effect of IL-4 occurs at the level of T-cell priming and may not be able to influence a pre-existing Th17-phenotype due to inhibited STAT6 activation following CD4+ T cell differentiation. It must be noted, however, that IL-4R polymorphisms are equally represented in patient populations, suggesting dampened IL-4 signaling is only one of many factors that limit Th1-/Th17-driven disease. The efficacy or usefulness of IL-4 as an immunomodulator may reside therefore in its ability to limit *de novo* Th1/17 differentiation and help slow disease progression by limiting epitope spread/development of T cells against new epitopes. These ideas are further discussed below in the superkine section.

Targeting IL-4 and IL-13 receptors to treat disease

IL-4 triggers cell signaling, first by binding with high affinity to the IL-4R α subunit, which causes a conformational change, followed by association with one of two low-affinity receptor subunits either the common γ C or the IL-13R α 1 chain (120). IL-4 engagement of the type I IL-4 receptor induces stronger tyrosine phosphorylation of IRS-2, whereas IL-4 or IL-13 binding the type II receptor results in weak tyrosine phosphorylation of IRS-2. This signaling pattern correlates with the degree of target gene expression and maturation/differentiation in macrophages. These differential associations likely account for the recruitment of different JAKs and STAT/IRS pathways triggered and the different biological consequences downstream. To better understand how cytokine binding to the receptor elicited differential IRS-2 signaling responses, the cytoplasmic domain of the type I receptor γ C was swapped with the type II receptor IL-13R α 1 cytoplasmic domain and *vice versa* (9). Interestingly, the enhanced phospho-IRS-2 signaling responses could not be transferred to the type II IL-4 receptor simply by replacing the IL-13R α 1 cytoplasmic domain with that of the γ C chain. The reverse “tail swap” also held true: type I IL-4 receptor responses could not be diminished by the presence of the IL-13R α 1 cytoplasmic domain in place of the γ C tail. These results suggested that signaling responses are dictated, for the most part, by the interaction of the cytokines IL-4 and IL-13 with the extracellular and transmembrane domains of the receptor. These results have fortuitous implications for therapies directed at modulating IL-4/-13-mediated responses. Blocking or augmenting agents of downstream signaling can be directed at the cytokine:receptor complex outside the cell and they would not necessarily need to be cell-penetrative. Taken together with the success of cytokine

neutralization in animal models, these findings fostered the idea that manipulating cytokine structure may result in different biological outcomes (121). Moreover, the specificity of the cytokine:receptor binding also presents a unique opportunity to develop novel therapeutics that would effectively target a single receptor and block cytokine signaling while avoiding the complications associated with antibody neutralizing strategies.

IL-4 antagonists

Anti-cytokine therapy in the form of antibodies directed against the soluble cytokine (122–127), cytokine receptor decoys (25, 128, 129) and antibodies direct against the receptor (129, 130) have been widely studied and shown success in the clinic. Anti-cytokine therapies targeting soluble IL-4 and IL-13 have been less successful in treating human atopy/asthma, possibly because of their overlapping receptor binding properties. Several IL-4 mutants (or “muteins”) have been generated by various genetic/chemical modification strategies, including some currently in pre-clinical and clinical trials. Duppatla *et al.* demonstrate that chemical modifications to human IL-4 at residue 121 allowed for binding to the high affinity IL-4 α chain but blocks the recruitment of either the γ C or IL-13R α , preventing IL-4R signaling in Jurkat T-cells (131). Pitrakinra (AER-001, BAY-16-9996) has been well studied in several clinical trials (129, 130). In asthmatics, pitrakinra reduced exacerbations as well as the need for rescue inhaler although these effects were predominantly observed in patients with the specific SNPs rs3024585, rs3024622, and rs4787956 (132, 133). In spite of blocking wildtype IL-4/-13 binding by up to 99%, IL-4 antagonists have not been able to block the biological activity of endogenous IL-4/-13 completely. This may be due to incomplete receptor saturation, the ability of a small number of receptors to assemble an alternative IL-4R complex configuration that leads to signaling or may reflect the very small amount of endogenous cytokine necessary to initiate a signaling response. In clinical studies, several safety aspects need to be considered including the effect of exogenous inhibitory cytokine on endogenous cytokine levels, immune function and cognitive function based on the emerging data as to the role of IL-4 in the brain. In addition, it will be important to sample a wide array of tissue in order to fully ascertain the consequences of inhibiting the IL-4/-13 signaling pathways.

IL-4 superkines

Cytokine modification can also enhance the biological properties of IL-4/-13. Junttila *et al.* generated modified human IL-4 that activated the IL-4R with greater affinity than endogenous cytokine and specifically activated the type I IL-4R or the type II IL-4R by increasing the affinity for the γ C or the IL-13R α 1 chain respectively (4). The authors demonstrated that these superkines stimulate more robust and prolonged IL-4-induced STAT6 signaling than endogenous IL-4. In addition, a superkine with high affinity for γ C only induced changes in gene transcription consistent with the type I receptor signaling, while the superkine with high affinity for the IL-13R α 1 induced a type II receptor gene profile. These findings have important implications as a therapeutic agent. These superkines would have the potential to specifically target immune cells to modulate Th1 and/or Th17-driven inflammation with out influencing structural cells or off-target effects. IL-17-producing CD4+ T cells and innate $\gamma\delta$ T cells have been associated with extreme, corticosteroid resistant forms of asthma. To date, very few therapeutic approaches have been

effective in regulating IL-17 responses. Since IL-4 is known to be a potent negative regulator of CD4⁺ Th1/Th17 (134) and IL-17-producing $\gamma\delta$, a therapeutic IL-4 superkine may be an effective way to dampen unchecked Th1/17-driven inflammation in autoimmune diseases and asthma. Since human T and B cells predominantly express γC and not IL-13R α 1, a superkine that specifically binds γC to enhance the Th2-differentiation program, without influencing non-structural cells expressing the IL-13R α 1, may be an effective way to modulate inappropriate inflammation. Such an approach may also influence the differentiation of monocytes/macrophages as well as dendritic cells. However, *ex vivo* experiments suggest that this may not be the case. Surprisingly, human PBMC-derived dendritic cells (DCs) exposed to super-IL-4, which preferentially recruits the γC , failed to mature, compared to DCs treated with native IL-4 (4). These observations indicate that human blood-derived DC specifically used the type II IL-4R to differentiate in response to IL-4, whereas human monocytes utilized both the γC (type I receptor) and the IL-13R α 1 (type II receptor) (4). These results suggest that a highly specific superkine may provide an important adjunct therapy specifically targeted to affect the function of relevant cell types in different disease. Ideally, combining the technologies of IL-4 antagonists and superkines could yield a highly specific therapy that could effectively target a single receptor subunit to block IL-4 signaling.

IL-4 and IL-13 signaling in health and disease

New insights into the regulation of IL-4 and IL-13 signaling

IL-4 and IL-13 take advantage of numerous signaling molecules to trigger activation and gene transcription namely 2 main pathways; the STAT6 pathway and the IRS pathway. STAT6 signaling in asthma is well characterized. In mouse models of allergic airway disease, a model of asthma, STAT6 phosphorylation is critical for Th2 differentiation (135), eosinophil migration (136, 137), mucus production and airway hyper-reactivity (138). More recently, IL-13 derived from NKT cells promotes ulcerative colitis through the activation of STAT6 (139). Rosen *et al.* describe a STAT6-dependent role for IL-13 signaling in the expression of claudin-2, a pore-forming protein that promotes epithelial barrier dysfunction. Furthermore, IL-13-driven STAT6 signaling promotes the production of IL-33 and thymic stromal lymphopoietin (TSLP), cytokines which promote Th2 differentiation and amplify intestinal inflammation (139).

Intrinsic factors, such as JAK/STAT inhibitors of cytokine signaling as well as extrinsic factors such as immunoregulatory cytokines, all participate to limit IL-4/-13 signaling. Intrinsic regulatory mechanisms target signal transduction molecules within the cell to limit their activation and prevent subsequent signaling. Src homology region 2 domain-containing phosphatase-1 (SHP1, also known as Tyrosine-protein phosphatase non-receptor type 6 or PTPN6), a protein tyrosine phosphatase, is a negative regulator of several signaling pathways by promoting the dephosphorylation of signaling molecules. Signaling studies revealed STAT6 is dephosphorylated by Shp1 and the IL-4R α SH2 binding domain of the ITIM is required for efficient Stat6 phosphorylation (Figure 3A) (140). Mice deficient in Shp1 (motheaten and viable motheaten mice) develop spontaneous Th2-like disease and are highly susceptible to allergic airway disease marked by hyper-activation of Th2 cells, hyper-

Ig production and lung tissue damage (141–144). *In vivo* studies indicate Shp1 regulates the activation and proliferation of T cell function, macrophage, epithelial cell responsiveness, mucus production and B cell development (141, 145–150). Conflicting reports have made it difficult to decipher how SHP1 deficiency regulates Th2-like disease that *in vitro* systems have not been able to clarify these issues. Conditional knockouts were recently generated to help address the cell-type specific role of Shp1 in regulating allergic disease. Zhou *et al.* reported that Shp1 regulates the development of mast cells in a murine model of anaphylaxis but reported no role for T and B cells or macrophages (151). Previous studies suggested that in T cells, Shp1 regulated TCR:MHC affinity, proliferation and Treg polarization, however, these reports were contradictory [reviewed in (152)]. To clarify the role of Shp1 in CD4+ T cells, Johnson *et al.* generated a conditional CD4+ T cell Shp1 knockout mouse. These mice did not develop the characteristics *motheaten* inflammation seen in whole body knockouts. In contrast to studies carried out in whole body knockout mice and *in vitro*, the authors report normal thymocyte development and normal TCR signaling in these mice, however, more activated CD4⁺CD44^{hi} T cells could be detected in the periphery (153). These cells were highly responsive to IL-4 such that neutralizing IL-4 attenuated CD4+ T cell activation. Furthermore, using bone marrow chimera mice, they show that autocrine IL-4 is a potent activator of these autoreactive T cells. It is important to note that these CD4+ T cells developed an activated phenotype in the absence of a specific antigen, suggesting they are autoreactive. Phenotypic analysis revealed increased IL-4 production from these cells and enhanced serum IgE levels, supporting the idea that Shp1 negatively regulates Th2 polarization and T cell helper functions, possibly through the production of IL-4. Furthermore, they show that blocking IL-4 *in vivo* attenuates CD4+ T cell activation, suggesting IL-4 signaling contributes to the diffuse T cell activation in these mice (153). Although neither of these recent papers address the signaling processes reported *in vivo*, we can extrapolate our understanding of the function of Shp1 in regulating STAT6 phosphorylation to appreciate that these events are likely IL-4-driven. In humans, decreased SHP1 expression is linked to some lymphoma and leukemia as well as MS (152, 154, 155). Taken together, we can begin to appreciate that targeting STATs with small molecules such as FDA-approved statins [reviewed in (156)] may be an effective way to modulate IL-4-driven eosinophil/mast cell recruitment and differentiation, T cell activation, B cell activation and differentiation as well as hyper-Ig syndromes.

IRS-2 is also a critical mediator of IL-4 and IL-13 signaling. IL-4 induces strong phosphorylation of IRS-2 whereas IL-13 only weakly induces IL-13 phosphorylation (157). This level of phosphorylation correlates with the strength of M2 gene induction in macrophages and may account for some of the different physiological effects of these two cytokines *in vivo*. It has been reported that insulin signaling through IRS-2 leads to downregulation of the IRS-2 receptor (158), presumably through the induction of a negative regulator. Indeed both suppressor of cytokine signaling (SOCS)1 and SOCS3 have been shown to negatively regulate IRS-2 expression through the ubiquitination and proteasomal degradation leading to insulin insensitivity (159, 160). In the context of IL-4/-13 signaling, the role of the SOCS proteins is less well characterized. SOCS1 and SOCS3 have been implicated in macrophage polarization although studies of SOCS1 have been particularly difficult to interpret. SOCS1, but not SOCS3 is induced by IL-4 and was associated with an

M2 phenotype (161). In contrast, SOCS3 was associated with an M1 phenotype, yet SOCS1 reportedly plays a role in negatively regulating M1 polarization as well (161–163). These findings have yet to define the molecular mechanisms that would account for observations in macrophage polarization and relate these observations to human disease. In line with the understanding that SOCS1 regulates M2 polarization and IRS-2 signaling, we have also found a role for SOCS1 but not SOCS3 regulation of IRS-2 in human monocytes (McCormick and Heller, manuscript in preparation). The importance of IRS expression regulation has important implications in the field of IL-4-driven cancer proliferation and metastasis. Porter *et al.* examined the expression of signaling adaptor proteins IRS-1 and IRS-2 in human breast cancer tissues (164). They found a correlation between high IRS-2 expression and invasive ductal carcinomas, whereas, high expression of IRS-1 correlated with poorly differentiated, less invasive cancers. Moreover and IRS-1 correlated with sensitivity to chemotherapy. Since IL-4 potently activates IRS-2 to trigger transcriptional activation, suppress apoptosis and actively promote differentiation, increased IRS-2 expression may begin to account for the exquisite sensitivity of breast cancer to IL-4/-13-driven metastasis. Furthermore, given the importance of IL-4/-13 activated tumor infiltrating macrophages with aggressive cancers, these observations suggest a possible link between macrophage IRS-2 expression and IL-4 sensitivity in tumor progression. Most importantly, the authors have identified not only a therapeutic target to dampen IL-4 signaling but also a possible prognostic marker. Investigation into strategies that modulate IL-4 signaling through the regulation of the expression of IL-4 receptor subunits or the activation of downstream STAT6 and IRS-2 signaling is warranted. Such approaches would be promising as a combinational therapy alongside standard therapies to combat aggressive tumors and metastasis.

mTOR signaling: a master regulator of IL-4-induced signal transduction

Early studies in the 1990s had uncovered that IL-4 stimulation lead to activation of PI3K through IRS-2 (165, 166) and to subsequent activation of AKT (167, 168). Other studies demonstrated a role for the IL-4/STAT6 pathway in regulating peripheral nutrient metabolism through immune cells resident in metabolic tissues such as liver, fat and skeletal muscle (168). Metabolic responses are coordinated inside the cell by the serine/threonine kinase, mTOR, acting as a sensor of glucose and amino acid availability (169). As background, mTOR is the catalytic protein that nucleates two different complexes: mTOR complex 1 (TORC1) and mTOR complex 2 (TORC2, (170)). TORC1 and TORC2 are made up of multiple proteins, with mTOR being common to both, although they localize to different regions within the cell (Figure 2C). The small GTPase, Ras homolog enriched in brain (Rheb), activates TORC1 when it binds GTP. Activation of the TORC1 kinase by Rheb is controlled by tuberous sclerosis complex 1 (TSC-1) and TSC-2. The Tsc-1/-2 complex, in turn, is phosphorylated by AKT, inhibiting its GTPase activating protein (GAP) activity and allowing GTP-bound Rheb to remain associated with TORC1. Activated TORC1 phosphorylates multiple substrates including p70 S6-kinase and 4E-BP1. The TORC1 regulates multiple cellular processes including protein synthesis, lipid biogenesis, inhibition of autophagy and other processes (169). TORC1 is sensitive to inhibition by rapamycin mediated by FKBP12. Typically, mTORC2 is not rapamycin-sensitive but long-term rapamycin treatment in some cell types inhibits this complex (171). The upstream

activation of TORC2 is less well understood but thought to occur through PIP3 (172). TORC2 fully activates AKT by phosphorylation of serine 473 (173, 174). Thus, AKT is both an activator of TORC1 and a substrate of TORC2.

Although the PI3K-AKT-TOR pathway was described as activated downstream of IL-4 in T cells (175), how canonical IL-4 signaling pathways and nutrient/metabolic responses were integrated was not known. New insights into the role of the TOR complexes and TORC-activated molecules in activation and regulation of IL-4 signaling have been recently uncovered in a variety of immune cell types. The different downstream pathways and functions of mTORC1- and mTORC2-activated signaling following IL-4 stimulation has been an emerging focus of research in recent years.

The importance of mTOR-activated pathways in response to IL-4 leading to polarization of CD4⁺ T-cells was uncovered recently. Deletion of Rictor the mTORC2 adaptor, in CD4⁺ T cells leads to a deficit of Th2 polarization and IL-4 production (176, 177). However, the signaling pathways downstream of mTORC2 activation that promote Th2-polarization were not known. Heikamp *et al.* demonstrated that serum- and glucocorticoid-regulated kinase 1 (SGK1) is a critical regulator of Th2-cell lineage commitment downstream of mTORC2 (178). Under Th2 polarizing conditions (IL-4, α -IL-12 and α -IFN- γ), they found that SGK1 deletion in CD4⁺ T cells resulted in diminished Th2 cytokine production. These SGK1-deficient cells secreted much more IFN- γ , even without polarization, than wildtype CD4⁺ T cells under the same conditions. This led to suppressed allergic responses in the *in vivo* model of allergic lung inflammation. The molecular mechanism underlying changes in T cell polarization in Rictor-deficient cells was due to the lack of SGK-mediated inhibition of the E3 ubiquitin ligase, Nedd4-2. This resulted in enhanced ubiquitin-mediated degradation of JunB, which is required for Th2-cell development.

Another important IL-4-induced mTOR-mediated response is the differentiation of macrophages into the M2 macrophage phenotype *in vitro*. New data have demonstrated the essential role of the TOR pathway in regulating IL-4-polarization of macrophages. Byles *et al.* used bone marrow macrophages from myeloid-specific Tsc1-deficient mice to highlight the role of TORC1 as a negative feedback regulator of IL-4-activated AKT signaling in macrophages (179). When Tsc1, the negative regulator of TORC1, is deleted, constitutive TORC1 activity leads to suppression of AKT signaling and deficient M2 polarization of macrophages. Validation of this study by Zhu *et al.* with similarly *Tsc1-LyzMcre*-floxed animals reinforced the finding that Tsc1 deletion in macrophages resulted in diminished AKT phosphorylation and M2 polarization following IL-4 stimulation (180). The authors further demonstrated that C/EBP β , a critical transcription factor for the M2 fate, was also decreased in the nuclei from IL-4-stimulated macrophages derived from the Tsc1-deleted mice. Furthermore, the importance of the TORC1 complex as a regulator of M2 polarization in human macrophages was described by Mercurio *et al.* (181). Rapamycin treatment of human monocyte-derived macrophages polarized with IL-4 resulted in a decrease in the number of M2 macrophages and increased apoptosis. On the other hand, M1 polarization markers were enhanced. These *in vitro* findings were also recapitulated in monocyte-derived macrophages from patients treated with rapamycin prior to islet transplantation. The authors concluded that the mTOR pathway was essential for M2 macrophage survival although no

deeper insight into the molecular mechanism in human macrophages was provided. Further investigation will be required to reconcile the apparent opposite conclusions made from mouse and human studies described above.

Taken together, these papers point to mTORC1 and 2, or TORC1 and 2-activated pathway(s), as canonical regulators of IL-4 and IL-13 signaling pathways contributing to Th2 and M2-type inflammation. Therefore, manipulation of TOR activity with rapamycin as a therapeutic in the setting of IL-4-driven Th2-mediated disease was investigated in several new publications. Mushaben *et al.* (182) repeated early studies by Fredricksson *et al.* (183) comparing the effects of rapamycin treatment during the induction phase of allergic lung inflammation versus established disease. Mushaben and colleagues described the importance of mTORC1-mediated signaling in promoting the early phases of allergic disease using the house dust mite (HDM)-induced allergic inflammation model in mice. Rapamycin had little effect on the later, effector phase of established disease. Therefore, the utility of rapamycin or mTORC1 inhibition as a therapeutic is somewhat tempered by these *in vivo* findings. There was no effect on AHR in the most recent study (182). In fact, AHR and airway inflammation became worse when rapamycin was given to animals with established allergic lung inflammation (183). Uncovering more details of the precise molecular underpinnings of mTORC1 or mTORC2 regulation to subvert pathogenic IL-4 signaling in allergic inflammation will aid in more effective manipulation of these pathways for therapeutic benefit.

Concluding remarks

IL-4 and IL-13 are important cytokines for cell differentiation, promoting allergic and anti-helminth inflammation, proliferation and cancer. In addition, these cytokines regulate type 1- and type 17-driven T cell inflammation, but should they not be confused as anti-inflammatory cytokines. Both IL-4 and IL-13 have important functions and drive equally important and potentially pathogenic type 2 inflammation. Ubiquitous expression of the type I and II receptors on immune cells and the type II IL-4 receptor on structural cells means that IL-4 and IL-13 have the potential to affect every cell in the body. IL-4 receptor subunit expression is tightly regulated but can be modified by genetic, environmental and inflammatory stimuli. Insights into the mechanisms that regulate receptor sensitivity, density and signaling are important for identifying appropriate therapeutic targets in several diseases. In many diseases, aberrant signaling causes or certainly amplifies disease progression. Understanding the regulation of IL-4/-13 signaling is increasingly important in order to identify and develop therapies that augment the action of these pathways. The discovery of mutein cytokines have precipitated the development of antagonists as well as high affinity superkines. Superkines with modified affinity for the secondary chain could show promise in modulating activation/signaling and downstream responses in targeted cell types. These modified cytokines along with the refinement of cytotoxins could be exploited to diminish or even reverse disease severity in a variety of diseases, notably, cancer, neurological disorders, asthma and allergies.

Abbreviations

AP-1	Activator protein-1
CNS	central nervous system
DEPTOR	DEP-domain-containing mTORinteracting protein
FDA	US Food and Drug Administration
FoxO	forkhead box protein O
Hsp70	heat shock protein 70
Hrs	hepatocyte growth factor-regulated tyrosine substrate
Ig	immunoglobulin
ITIM	immunoreceptor tyrosine-based inhibition motif
IL	interleukin
JAK	Janus Kinase
IRS	insulin receptor substrate
IFN	interferon
mLST8	mammalian lethal with Sec13 protein 8
MHC	major histocompatibility complex
mSIN1	mammalian stress-activated protein kinaseinteracting protein 1
NO	nitric oxide
PI3K	phosphatidylinositol-4,5-bisphosphate 3-kinase
PIP3	phosphatidylinositol (3,4,5)-trisphosphate
PDK1	3-phosphoinositide dependent protein kinase-1
PRAS40	proline-rich Akt substrate of 40 kDa
PROTOR	protein observed with RICTOR
RICTOR	rapamycin-insensitive companion of TOR
RHEB	Ras homolog enriched in brain
TSC1	tuberous sclerosis complex 1
Raptor	regulatory-associated protein of mTOR
SGK1	serum and glucocorticoid-inducible kinase 1
STAT	Signal transducer and activator of transcription
STUB1	STIP1 homology and U-Box containing protein 1
SOCS1	suppressor of cytokine signaling 1

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Highlights

- IL-4 and IL-13 receptors are expressed on every cell in the body and expression is altered during disease
- New mechanisms that regulate receptor expression have been identified yet highlight the lack of understanding of cytokine receptors in disease processes
- IL-4/-13 and their receptors play a greater role in brain homeostasis than previously appreciated
- Therapeutics that specifically target a single receptor subunit may hold promise in ameliorating treatment-refractory diseases
- New insights into regulation of IL-4/-13 signaling suggest new therapeutic approaches are on the horizon

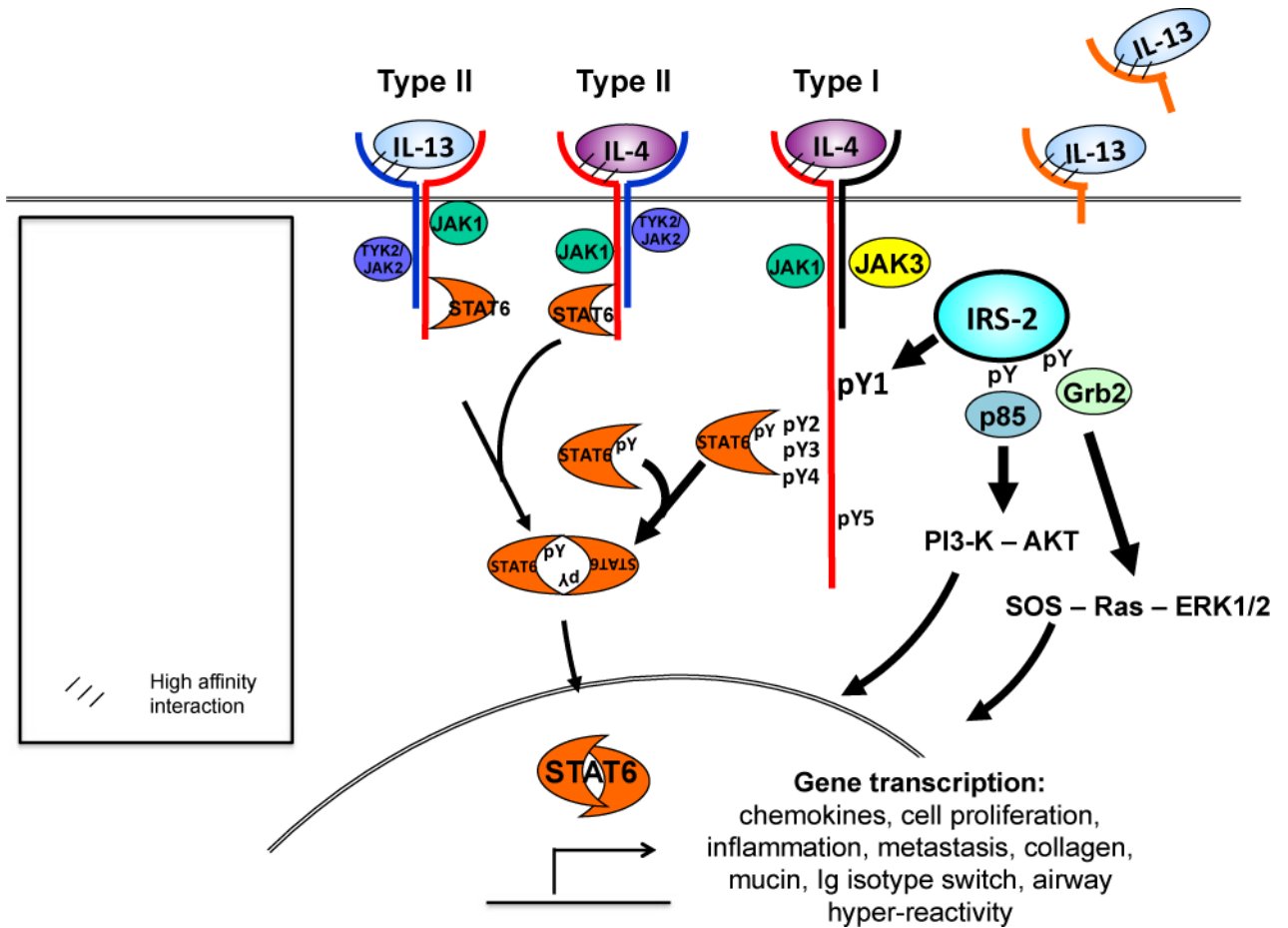


Figure 1. IL-4 and IL-13 receptor structure

IL-4 signals through two possible receptor complexes composed of a heterodimer of the IL-4R α (140 kDa) and γ c chain (60 kDa); type I receptor or the IL-4R α and IL-13R α 1 (65–70 kDa) chain; type II receptor. IL-4 binds IL-4R α with high affinity, triggering dimerization with the secondary signaling chain. IL-13 binds IL-13R α 1 which complexes with the IL-4R α , forming the type II receptor. Signaling through the type I receptor leads to activation JAKs and downstream signaling adaptor molecules STAT6 and IRS-2, whereas signaling through the type II receptor predominantly activates STAT6. IL-4 signaling also activates PI3-K and AKT. IL-13 also binds cell-surface and soluble forms of the IL-13R α 2, this so-called inhibitory subunit binds IL-13 with greater affinity than IL-13R α 1 and acts as a “cytokine sink”.

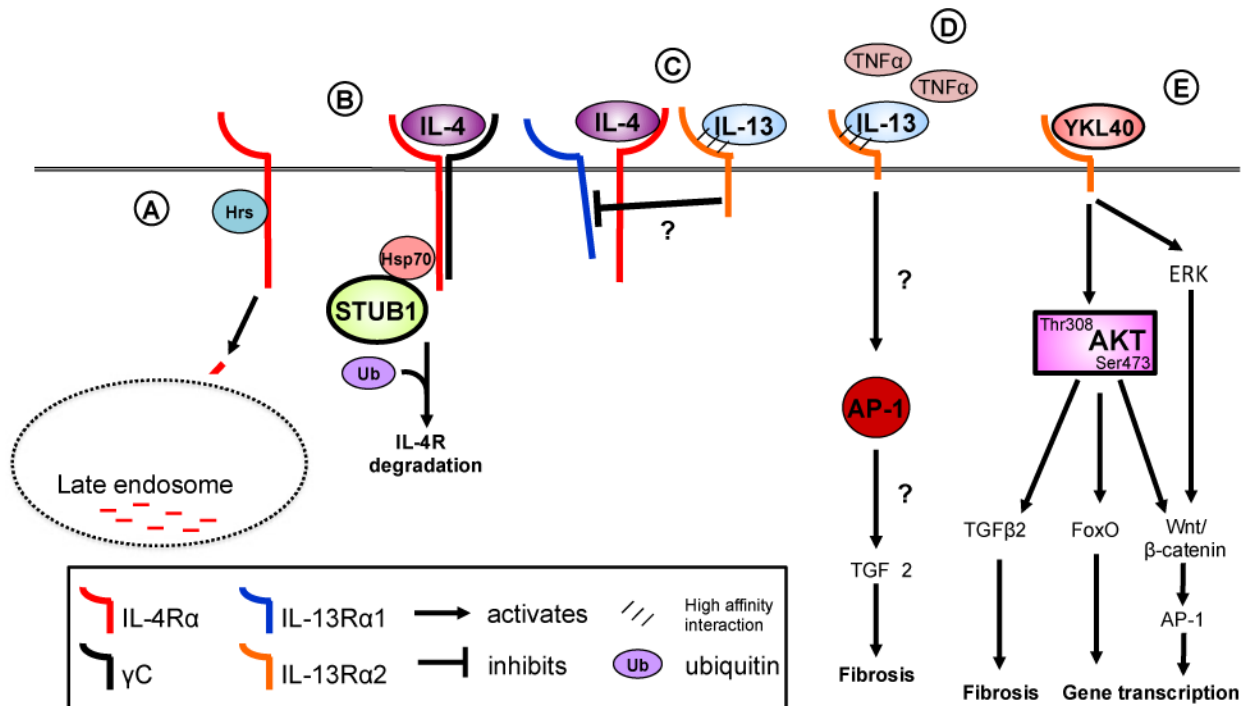


Figure 2. Regulation of signaling receptors

IL-4R α expression is regulated in the steady state by Hepatocyte growth factor-regulated tyrosine substrate (Hrs) (A). Hrs binding targets the IL-4R α to the late endosome for degradation. Downregulation of IL-4R α occurs through STUB1 binding the cytoplasmic domain of the receptor through Hsp70 (B). Subsequent ubiquitination and degradation the IL-4R α reduces cell surface expression and cellular responsiveness to IL-4. The IL-13R α 2 acts as a negative regulator of IL-4- but not IL-13-induced signaling through the type II IL-4 receptor (C). The IL-13R α 2 acts as a negative regulator of IL-4- but not IL-13-induced signaling through the type II IL-4 receptor although the precise mechanisms of this inhibition are not understood. IL-13R α 2 expression is induced by TNF α in fibroblasts and leads to AP-1 driven pro-fibrotic reprogramming (D). YKL40 binds binds to the IL-13R α 2 chain leading to serine 473 phosphorylation in addition to triggering ERK signaling (E). AKT phosphorylation following binding of YKL40 to IL-13R α 2 activates a myriad of signaling pathways including FoxO1/3 and the Wnt/b-catenin pathway. Hepatocyte growth factor-regulated tyrosine substrate, Hrs: STIP1 homology and U-Box containing protein 1, STUB1; heat shock protein 70, Hsp70; forkhead box protein O, FoxO; transforming growth factor- β , TGF- β ; activator protein-1, AP-1.

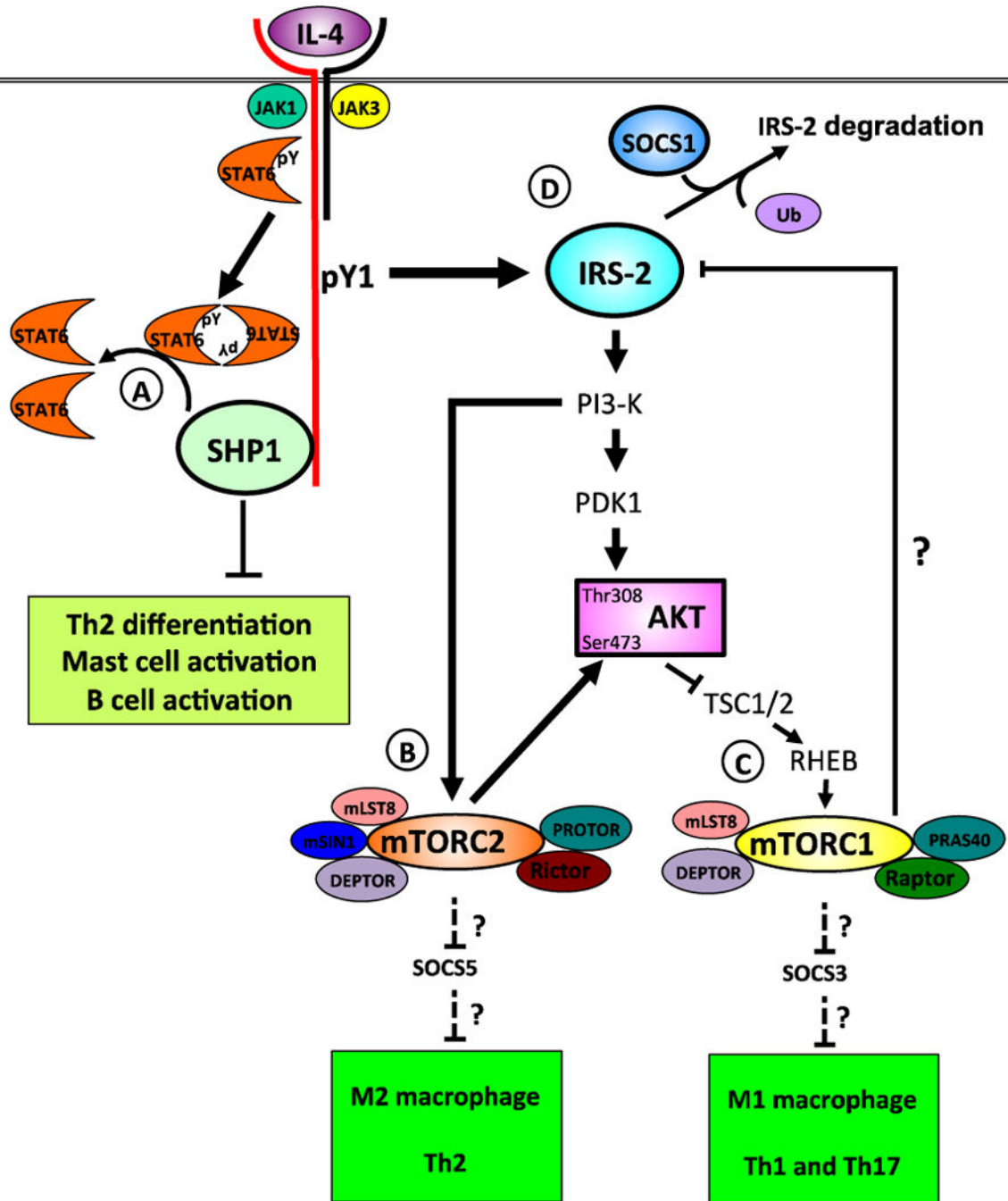


Figure 3. Regulation of signaling in response to IL-4 and IL-13

Engagement of the type I IL-4 receptor by IL-4 brings about tyrosine phosphorylation of the associated JAKs, key tyrosine residues (pY1-5) in the cytoplasmic domain in IL-4Ra, STAT6 and IRS-2. STAT6 is dephosphorylated by SHP1, a protein tyrosine phosphatase (A). De-phosphorylation of STAT6 is critical to limit autoimmune and Th2 allergic inflammation, mast cell, B cell and epithelial cell activation. IRS-2 binds the regulatory p85 subunit of PI3-K, activating the enzyme to generate PIP3. In turn, PIP3 activates PDK1 and mTORC2. The proteins comprising the two mTOR complexes 1 and 2 are shown (B). The proteins comprising the two mTOR complexes 1 and 2 are shown (B).

TORC2 kinase phosphorylates AKT on serine 473 and to SGK1 in CD4+ T cells, leading to Th2 polarization. AKT inactivation of the Tsc1/2 complex prevents hydrolysis of GTP bound to Rheb. GTP-bound Rheb activates mTORC1, triggering a negative feedback loop via multiple pathways on IL-4-activated IRS-2/PI3-K/AKT signaling (C). The composition of the mTORC1 complex is shown. The mTORC1 complex also regulates macrophage polarization, as Raptor-deficient macrophages have enhanced M1 and M2 responses. The mTORC1 and mTORC2 complexes also regulate by unknown mechanisms, the polarization of T cells [adapted from (170)]. An additional negative regulatory pathway triggered by IL-4 is the induction of the SOCS proteins. SOCS1 binds IRS-2 and targets it for ubiquitin-mediated degradation (D). Phosphotyrosine, pY; phosphatidylinositol-4,5-bisphosphate 3-kinase, PI3-K; phosphatidylinositol (3,4,5)-trisphosphate, PIP3; 3-phosphoinositide dependent protein kinase-1, PDK1; mTOR complex 2, mTORC2; serum and glucocorticoid-inducible kinase 1, SGK1; Ras homolog enriched in brain, RHEB; tuberous sclerosis complex 1, TSC1; regulatory-associated protein of mTOR, Raptor; mammalian lethal with Sec13 protein 8, mLST8; proline-rich Akt substrate of 40 kDa, PRAS40; DEP-domain-containing mTOR-interacting protein, DEPTOR; rapamycin-insensitive companion of TOR, RICTOR; mammalian stress-activated protein kinase-interacting protein 1, mSIN1; protein observed with RICTOR, PROTOR; suppressor of cytokine signaling 1, SOCS1.