



Published in final edited form as:

*Curr Signal Transduct Ther.* 2009 January 1; 4(1): 22–30. doi:10.2174/157436209787048711.

## Leptin signaling: A key pathway in immune responses

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### Abstract

Leptin is a hormone whose central role is to regulate endocrine functions and to control energy expenditure. After the discovery that leptin can also have pro-inflammatory effects, several studies have tried to address - at the molecular level - the pathways involved in leptin-induced modulation of the immune functions in normal and pathologic conditions. The signaling events influenced by leptin after its binding to the leptin receptor have been under scrutiny in the past few years, and considerable experimental work has elucidated the consequences of leptin effects on immune cells. This review examines the biochemistry, function and regulation of leptin signaling in view of possible intervention on this molecule for a better management and therapy of immune-mediated diseases.

### Introduction

The past few years of research on leptin have provided important insights into the intricate network that links metabolism and immune homeostasis. Leptin is a 16-kD hormone mainly produced by the adipose tissue that has the peculiarity of also acting as a cytokine (1).

As hormone, leptin helps monitoring body weight (as fat content) to adjust the metabolic rate, whereas as cytokine leptin can exert a prominent action on innate and adaptive immune responses (2,3). The signalling events that follow the binding of leptin to its receptors have been studied extensively and characterized at the biochemical and molecular levels in many systems and, more recently, in relation to immune responses. This review highlights the most recent advances in this field and discusses the newly envisioned possibilities of a modulation of leptin signalling for immunotherapeutic intervention.

### Leptin receptors

In addition to its hormonal activity, leptin effects on the immune system can be explained by the fact that leptin is a cytokine with a long-chain four-helical bundle typical of the type I cytokine family (4). Moreover, the pleiotropic biological effects of leptin can be explained by the wide distribution of leptin receptors on different types of cells, including those in extraneural tissues.

Structurally, leptin receptors (OB-R) belong to class I cytokine family receptors (which include the receptors of IL-2, IL-3, IL-4, IL-6, IL-7 and granulocyte colony-stimulating factor (GM-CSF) (5–11).

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Leptin receptors have characteristic extracellular motifs of four cysteine residues and contain the amino acid sequence WSXWS (Trp-Ser-Xaa-Trp-Ser) (12) and fibronectin type III domains (13,14). These receptors are produced as isoforms derived from alternative splicing of RNA transcripts of the *db* gene, and are designated OB-Ra, OB-Rb, OB-Rc, OB-Rd, OB-Re and OB-Rf (15). OB-Rb is expressed at high levels in the hypothalamus and, at lower levels, in many peripheral tissues (16,17). OB-Ra and OB-Rc are highly expressed in choroid plexus and microvessels, where they may play a role in leptin uptake or efflux from the cerebrospinal fluid, as well as in receptor-mediated transport of leptin across the blood-brain barrier (18).

All isoforms have a common extracellular domain of more than 800 amino acids, a transmembrane domain of 34 amino acids, and a variable intracellular domain characteristic of each isoform. Among the six isoforms, the main differences are in intracellular amino acids: 301 for OB-Rb, and 34, 32 and 40, respectively, for the short forms OB-Ra, OB-Rc and OB-Rd. The OB-Re isoform lacks the transmembrane and cytoplasmic parts, acting as a soluble receptor (19,6,15) which likely contributes to the regulation of plasma leptin levels by binding with high affinity free circulating leptin (20,21).

Another classification of OB-R isoforms is in three subclasses: short, long and secreted receptors. The long form of OB-R seems to be the only functional (signalling) isoform, while the lack of a full-length leads to the obese phenotype in *db/db* mice (22) – a phenotype rescued by neuronal transgenic expression of the long-form of OB-R (23).

The short and long isoforms share the extracellular and transmembrane domains as well as the same first 29 intracellular amino acids, diverging in sequence following alternative splicing of the 3' exons. Moreover, the OB-R extracellular domain has two cytokine-like receptor motifs and four fibronectin type III domains (5,8,9). Studies with mutant receptors have revealed that only the second putative binding domain mediates leptin binding and receptor activation, with an affinity in the nanomolar range (24).

Both long and short OB-R isoforms are capable to form homodimers in the absence of ligand (25–27) and the formation of dimers seems necessary for signalling (28). Considering that each OB-R binds one molecule of leptin, the result of dimerization is the formation of a tetrameric receptor/ligand complex (27) that induces a conformational change in the receptor structure which is critical for activation (24).

Intracellularly, the OB-R does not have an intrinsic tyrosine kinase domain, therefore it binds cytoplasmic kinases - mainly Janus tyrosine kinase (Jak)2 (29).

Like other cytokine receptors, OB-R contains a highly conserved, proline-rich box1 (30) and two putative, less conserved, box2 motifs (29,31,32). Box1 and box2 motifs are considered important in recruiting and binding Jaks (33,34), yet recent studies showed that only box1 and surrounding amino acids (31–36) were essential for full Jak activation (32,35), being two conserved amino acids (Leu896, Phe897) crucial for signalling (35). However, although an intact box2 motif seems not required to activate Jaks (32,35), it is nonetheless important for full induction of the signal transducer and activator of transcription (STAT) signalling pathway (36).

## Jak/STATs

The finding of a homology of OB-R with other class I cytokine receptors has long suggested the possibility that leptin binding might mediate cytokine receptor-like signals, including the activation of Jaks and STATs (37–39).

The Jak/STAT pathway is mostly activated by interferons, interleukins or other cytokines whose receptors lack intrinsic kinase activity, and comprises a family of four non-receptor Jaks and seven 85–95 kDa transcription factors (STATs) that are regulated by phosphorylation on specific serine and tyrosine residues.

Of the four known members of the Jak family, Jak1, Jak2 and tyrosine kinase 2 (Tyk2) are widely expressed, while Jak3 is found only in cells of the haematopoietic system (40). Recent studies indicate that, under physiological conditions, only Jak2 is activated during OB-R signaling (32).

Activated Jaks transphosphorylate each other, as well as other tyrosine residues (Tyr985, Tyr1138 and Tyr 1077) of the OB-R (41,42), providing docking sites for downstream molecules such as STATs. At the end, OB-R signalling by leptin results in STAT3 binding (37,43–48), and activation of STAT1 (37,43), STAT5 (37,43,45) and STAT6 (43). Then, recruited STATs become tyrosine-phosphorylated by Jaks (14), with subsequent dissociation from the receptor and the formation of homo- or heterodimers. STAT dimers translocate into the nucleus and act as transcription factors by binding specific response elements in the promoter region of their target genes, such as sis-inducible-element (SIE), acute-phase-response-element (APRE) and GAS-like elements (14,37,43,).

Notably, the three intracellular tyrosine residues of OB-Rb exhibit different capabilities for downstream activation signalling. Tyr<sup>985</sup> is required for the activation of the Ras/Raf/ERK pathway. The phosphorylation of Tyr<sup>985</sup> creates a binding site for the C-terminal SH2 domain of the tyrosine phosphatase SHP-2, leading to the activation of the canonical p21<sup>Rasp</sup>/extracellular regulated kinase (ERK) signalling cascade through the recruitment of the SH2-domain-containing adapter protein, growth factor receptor binding-2 (Grb-2) (9). While either Tyr<sup>1077</sup> or Tyr<sup>1138</sup> are required for leptin-induced tyrosine phosphorylation of STAT5, Tyr<sup>1138</sup> is essential for the activation of STAT1 and STAT3 (49,50).

Also, Tyr<sup>1077</sup> appears as the only intracellular tyrosine residue sufficient to induce tyrosine phosphorylation of STAT5 and STAT5-driven reporter gene activity *in vitro* [50]. In *db/db* mice lacking the long form of the leptin receptor, impaired STAT signalling has been demonstrated (38), suggesting that *db* phenotypes are caused by a failure of STAT signalling. Confirming a role of STAT signalling in the control of body weight, leptin-deficient *ob/ob* mice show significantly lower STAT3 in the hypothalamic arcuate nucleus (51).

Leptin seems to exert its effects through the Jak/STAT pathway not only in the hypothalamus to control body weight, but also on immune cells. In blood mononuclear cells, leptin increases Jak2/3 and STAT3 phosphorylation, which promote proliferation and activation of T lymphocytes upon PHA-stimulation. On anti-CD3/CD28 stimulated T regulatory cells, leptin neutralization results in the phosphorylation of STAT3 associated with degradation of p27<sup>kip1</sup> (52).

## Suppressors of Cytokine Signalling (SOCS)

The Jak/STAT pathway of cytokine signalling is under the negative-feedback control of suppressors of cytokine signalling (SOCS) proteins (53,54). SOCS proteins are induced upon cytokine stimulation and attenuate signalling by various cytokine receptors, allowing possible cross-regulation among several cytokine systems.

Members of the SOCS family, which contain an SH2 domain, are induced by a variety of cytokines, acting as negative regulators of signalling by binding to phosphorylated Jak proteins or by direct interaction with tyrosine phosphorylated receptors. The family of SOCS proteins consists of 8 members: cytokine inducible Src-like homology 2 (SH-2) protein (CIS) and

SOCS1 through SOCS7. Structurally, SOCS proteins are characterized by a central SH-2 domain, an N-terminal preSH-2 domain, with an ESS (extended SH-2 subdomain) region and in some cases a kinase inhibitory region (KIR) domain, which abolishes the kinase activity of the Jaks, and a more conserved C-terminal SOCS-box (55)- a key mediator of proteasomal degradation (by linking ubiquitin to the substrate).

Only SOCS1 and 3 carry a KIR domain in their N-terminal region involved in the inhibition of the Jak activity. They both inhibit Ob-R signalling, using different mechanisms. SOCS1 directly interacts with the kinase domain of Jak2 by targeting the phosphotyrosine Y<sup>1007</sup> in the Jak2 activation loop (56,57). The KIR domain association with the catalytic groove of Jak2 suggests that it might act as a pseudosubstrate mimicking the activation loop that regulates the access to the catalytic groove (it could obstruct the ATP binding pocket and hinder accessibility for substrates) (56,57). Differently from SOCS1, SOCS3 has only weak affinity for Jak2. It is thought to inhibit kinase activity through its KIR domain after the binding through its SH-2 domain with phosphotyrosine motifs in the receptor in the proximity of the Jaks (58).

Interestingly, leptin can induce SOCS3 expression (59–63) and the Tyr<sup>985</sup> of OB-Rb is a high-affinity binding site for SOCS3 (19,41,59,62,64). Endogenous SOCS3 expression inhibits tyrosine phosphorylation of OB-R, thus providing an important feedback mechanism for receptor signalling at the transcriptional level (62). In this context, the participation of SOCS3 in the negative-feedback mechanism of leptin signalling has been proposed to underlie the development of leptin resistance in relation to the hyperleptinaemia observed in the context of the majority of obesity cases (41).

Other SOCS proteins, in addition to the negative regulatory effects, can have positive effects on cytokine signalling (65,66). For example, SOCS2 can interfere with other SOCS proteins on several cytokine receptors, including OB-R signalling (65,67–69). SOCS2 can impair the inhibitory effect of SOCS1 or SOCS3 on leptin-induced signalling (69), and it can interfere with the association of CIS to the OB-R membrane proximal tyrosine (70). Also SOCS6 and SOCS7 can interact with the OB-R. SOCS7 may be implicated in OB-R signalling termination (71), i.e. by inhibiting STAT3 activation and/or by interacting with activated STAT3 to prevent translocation to the nucleus (72).

## PTP 1B

PTP1B (protein tyrosine phosphatase 1B) is another negative regulator of leptin signalling (and also insulin signalling (73,74) both *in vivo* and *in vitro* via dephosphorylation of Jak2 (75–77). Under physiological conditions, the effects of PTP1B are likely to be exerted via central and peripheral actions. Mice lacking PTP1B are resistant to developing diet-induced obesity and do not exhibit hyperphagia despite a clear hypoleptinemia (75). It appears that PTP1B-mediated hypophosphorylation of Jak2 can abrogate leptin-dependent induction of the STAT3 and MAPK-inducible SOCS3 and c-fos genes, respectively (76).

## Src-like homology 2 (SH2) domain containing protein tyrosine phosphatase (SHP-2)

Kinases are typically considered “dominant” regulators of intracellular signalling. However, in the past few years there has been growing attention on the study of phosphatases, which have opposite effects to kinases.

SH-2 domain-containing phosphatase-2 (SHP-2) is a constitutively expressed tyrosine phosphatase that regulates leptin signalling (78). SHP-2 is involved in the dephosphorylation of Jaks *in vitro* and down-modulates Jak2 and STAT3 activation *in vivo* (79).

SHP-2 carries two tandem SH-2 domains followed by a tyrosine phosphatase catalytic domain (80). When one SH-2 domain interacts with a tyrosine phosphorylated ligand, a conformational change occurs and brings this phosphatase to activation of OB-R at position Y<sup>985</sup> (indicated by the fact that in 293T cells SHP-2 binding to OB-Rb is prevented by the Y<sup>985</sup> mutation) (80).

The role of SHP-2 in OB-R signalling has been matter of debate, and initially suggested as inhibitor of OB-R signalling (mutation of the Y<sup>986</sup> position in the human OB-R leads to increased STAT3 signalling) (81). It is known that SHP-2 strongly activates the MAPK pathway and the SHP-2 induction of the ERK pathway can enhance leptin signalling, switching towards the MAPK signalling. ERK activation occurs predominantly through SHP-2 recruitment at tyrosine Y<sup>985</sup> via its C-terminal SH-2 domain. SHP-2 can be phosphorylated by Jak2 to form a docking site for the adaptor protein Grb2, leading to the activation of the ERK signalling cascade (41). Alternatively, ERK can be activated directly by Jak2, but still requires the intervention of SHP-2 (19).

### Cyclic AMP and PDE

Cyclic AMP activates protein kinase A (PKA) and plays a pivotal role in the crosstalk between many signalling systems (82,83). Metabolism of cAMP, and cGMP, is controlled by a family of PDE enzymes (84).

Leptin decreases cAMP levels in pancreatic  $\beta$ -cells via the activation of PDE3B (85), and the inhibition of GLP-1-stimulated insulin secretion (85) is one of the most evident physiological consequence of the response to leptin.

### 5'-AMP-activated protein Kinase (AMPK)

Leptin stimulates fatty acid oxidation to exert protective effects against lipotoxicity in the non-adipose tissues (86). Leptin selectively activates the  $\alpha$ 2 catalytic subunit of AMPK in skeletal muscle, which stimulates fatty-acid oxidation by blocking the effect of ACC (acetyl-CoA carboxylase) (87). AMPK represents a heterotrimeric enzyme that functions as a 'fuel gauge' to monitor cellular energy status (88,89). AMPK regulates food intake by responding to hormonal and nutrient signals in the hypothalamus (88). Activation of AMPK represents a signal to shut down anabolic pathways and to promote catabolic processes in response to a decrease in the ATP/AMP ratio by phosphorylating key enzymes of intermediary metabolism. In parallel with AMPK activation, leptin suppresses ACC activity, thereby stimulating  $\beta$ -oxidation in muscle by disinhibiting carnitine palmitoyltransferase 1 (CTP1). The direct activation of AMPK by leptin can explain, at least in part, the findings that leptin increases both *in vitro* and *in vivo* glucose uptake and metabolism.

### Phosphatidylinositol (PI) 3-kinase (PI3K)

PI 3-kinase activity is regulated by a wide spectrum of ligands, in particular growth factors such as insulin (90). The binding of PI3K regulatory subunit to tyrosine-phosphorylated proteins induces a conformational change allowing the activation of its catalytic subunit and consequent full activation of PI3K. This event leads this kinase to add a phosphate to the 3' position of the inositol ring of phosphatidylinositols, allowing switching of protein kinase-dependent cascades to lipid-dependent signalling cascades (90). PI 3-kinase products typically stimulate protein kinases such as Akt, also called protein kinase B (PKB) and protein kinase C (PKC) (90).

Leptin can act through some of the components of the insulin signalling cascade, as most insulin-dependent actions involve the activation of PI3k (91). The binding of insulin to its

receptor recruits several IRSs (insulin receptor substrates) that are tyrosine-phosphorylated by the intrinsic kinase activity of the receptor. This suggests a potential crosstalk between insulin and leptin signalling pathways (although the magnitude of PI3k stimulation in response to leptin is less than with insulin) (92).

The stimulation of PI3K leads to activation of PtdIns(3,4,5) $P_3$ -dependent serine/threonine kinases such as PDK1 (phosphoinositide-dependent kinase 1), which can activate Akt, a key serine/threonine kinase in subsequent downstream signalling. Although the contribution of insulin versus leptin-induced PI3K stimulation to signalling has to be better clarified, the current findings suggest that the PI3K/PDE3B/cAMP pathway interacting with the Jak2/STAT3 cascade represents a critical component of leptin signalling (53).

### Protein kinase B (PKB or Akt)

PKB is a serine/threonine kinase that plays an important role in many cellular processes including cell survival and carbohydrate metabolism (93). The binding of its PH domain to D3-phosphorylated phosphoinositides leads to PKB activation and subsequent phosphorylation on T<sup>308</sup> and S<sup>473</sup> residues (93).

Cell treatment with leptin induces Akt phosphorylation (94), although - as for PI-3K - the magnitude of this effect is small and much less than that produced by insulin.

Other studies suggest that the ability of leptin to stimulate Akt may depend on high intracellular cAMP levels, and this mediator might inhibit a phosphatase resulting in leptin-induced increase Akt phosphorylation (53).

### Protein kinase C (PKC)

Leptin has been shown to have both stimulatory and inhibitory effects on PKC, a serine/threonine kinase implicated, like PKB, in a wide range of cellular effects (95). Leptin seems to decrease insulin release from pancreatic islets of *ob/ob* mice in response to PKC stimulation (96), leptin action in pancreatic islets may involve inhibition of the PKC-regulated component of the phospholipase C (PLC)-PKC signalling system (that normally elicits insulin secretion) (97).

### MAPK

ERK members of the MAPK family are serine/threonine kinases of 44 and 42 kDa activated by a wide range of stimuli, including leptin, and are components of the Ras/MAPK signalling cascade (98).

The MAPK pathway can be stimulated by either the long or the short isoform of OB-R (at lesser extent by the former) (19,41) - an observation that supports the idea that the distal portion of the OB-R may not be essential for activation of MAPK.

It is possible that leptin can stimulate the MAPK pathway in two different ways: via tyrosine phosphorylation of Jak2 receptor-associated activation, or independently of receptor phosphorylation.

In any case, Tyr<sup>985</sup> of the long OB-R isoform has an important role in leptin-induced ERK activation (41). As a result of leptin administration, Tyr<sup>985</sup> becomes phosphorylated by recruited Jaks (mainly Jak2 and Jak1), and provides a docking site for SHP-2. After binding to that specific tyrosine residue, SHP-2 is phosphorylated at the C-terminus. This phosphorylated form, together with its adapter molecule Grb-2, activates downstream signalling, leading to the activation of the p21<sup>Ras</sup>/ERK signalling cascade (41). Subsequent

steps lead to the activation of ras and raf molecules, followed by the activation of MEK1 (99). The final consequences of ERK activation by leptin are many and include induction of specific target genes expression, such as c-fos or egr-1, a zinc-finger transcription factor that influences the initiation of growth and differentiation (100,101).

The activation of the MAPK signalling cascade by leptin has been observed both *in vitro* (30,41) and *in vivo*, both centrally and peripherally (101,102,103).

In monocytes, leptin induces expression and secretion of the IL-1 receptor antagonist (IL-1Ra) using the MAPK pathway that activates NF- $\kappa$ B (104). Also, the expansion of T regulatory cells following anti-leptin neutralization is mediated by the induction of ERK1/2 phosphorylation (52).

Leptin has also been shown to induce apoptosis through the MAPK pathway in precursor cells of the osteoblastic lineage. In this case, ERK1/2 activates cytosolic phospholipase A2 (cPLA2) that leads to cytochrome c release and caspase-3 and -9 activation, which coordinate cell death (105).

Finally, a wide range of stimuli, including cytokines, heat shock, osmotic stress, and ultraviolet light are can activate another member of the MAP kinase family, p38 MAPK (106). Treatment of human mononuclear cells with leptin increases p38 MAPK phosphorylation (107). However, a study in L6 muscle cells suggests that leptin does not alter p38 MAP kinase phosphorylation *per se* but rather reduces insulin-stimulated p38 MAP kinase phosphorylation (108).

Finally, leptin shares with other cytokines, growth factors and stressors, the ability to activate the stress-activated protein kinase c-Jun N-terminal kinase (JNK). For example, leptin enhances tumour necrosis factor (TNF)- $\alpha$  production via p38 and JNK MAPK (109). Among the possible downstream targets of leptin-induced activation of p38 and JNK MAPK pathways, the regulation of the transcription factor NF- $\kappa$ B appears important for the transcriptional regulation of pro-inflammatory cytokines such as TNF $\alpha$  and IL-1 $\beta$ .

## Leptin signalling in immune cells

OB-R lacks an intrinsic tyrosine kinase activity, and requires activation of receptor-associated kinases of the Jaks (38), which initiate downstream signalling and activate STATs (37) which dimerize and translocate to the nucleus, where specific gene responses are elicited (110).

Leptin stimulates and promotes the proliferation of human peripheral blood mononuclear cells (PBMC) (4). The presence of OB-R on monocytes and lymphocytes has been shown in mice (2,111) and confirmed in human peripheral blood T lymphocytes (both CD4 and CD8) (112).

The Jak-STAT signalling pathway triggered by leptin stimulation has been studied in human PBMC (47). It was shown that Jak2/3 are activated by tyrosine phosphorylation, and this effect is transient. Both Jak isoforms are physically associated with OB-R, as indicated by the co-immunoprecipitation of OB-R with Jak2 or Jak3. A preassociation of Jak2 with OB-R has also been described (38,40).

In PBMC, leptin stimulation induces tyrosine phosphorylation and translocation to the nucleus of STAT3 molecules (47,110,113), in addition to the phosphorylation of the STAT3 associated RNA binding protein Sam68 (39,40,114,115) (a tyrosine phosphorylated adaptor protein in T cell receptor activation that is associated with the SH2 and SH3 domains of Src and other signalling molecules, such as Grb2, PLC- $\gamma$ -1, and PI3K) (116-119). Since Sam68 is an RNA-binding protein and tyrosine phosphorylation is known to modulate RNA binding activity, the

data implicate possible regulation of RNA as a component of tyrosine kinases signalling pathways (117–119), i.e. through modification of the mRNA stability and/or translation.

This PI3K pathway has also been explored in PBMC responses to human leptin. PI3K activity associated with tyrosine phosphorylated proteins is increased more than 3-fold after leptin stimulation (120–122). Leptin treatment, similarly to insulin response, induced tyrosine phosphorylation of Sam68 and IRS-1, which associated with p85 (123,124), the regulatory subunit of PI3K via the SH2 domain, recruiting and leading to stimulation of PI3K activity (122). It is not yet known whether Tyr-phosphorylated Sam68 contributes to the increase in PI3K activity along with IRS-1, or whether it is only working as a docking protein.

Recently, leptin was shown to inhibit apoptosis in the thymus through an IRS-1/PI3K-dependent and Jak2-independent pathway (125). Treatment with leptin reduced thymic apoptosis, an effect that was not inhibited by the JAK inhibitor AG490 but that was inhibited by the PI3K inhibitor LY294002 and by antisense oligonucleotides to IRS-1 (125).

Moreover, the activation of MAPK by leptin in PBMC has also been assessed. Both ERK-1 and ERK-2 were found phosphorylated in a dose-dependent fashion in PBMC after incubation with human leptin (122). It was also found that leptin could induce sustained phosphorylation of p38 MAPK in human PBMCs (126), and the phosphorylation of the ribosomal protein S6 - the only protein in the large 40S subunit that has been shown to be phosphorylated in response to growth factors and mitogens (126). One route of leptin-induced S6 phosphorylation in human PBMCs is via MEK and p42/p44 MAPK (19,127–129), which activate MAPK-dependent S6 Kinase p90 RSK and S6. The other route seems to be mediated via activation of p70 S6 kinase, since it has been shown that leptin phosphorylates p70 S6 kinase at Thr<sup>389</sup>, the mammalian target of rapamycin (mTOR) target residue in the linker domain - an event that plays a central role in p70 activation (130). Accordingly, pre-treatment of cells with rapamycin (inhibitor of mTOR) abolished this phosphorylation and substantially reduced S6 phosphorylation (130). Strikingly, the MEK inhibitor PD98059 not only inhibited p90 RSK phosphorylation, as expected, but also abolished p70 S6 Kinase and S6 phosphorylation, suggesting an essential role of MEK activation in a full induction of p70 S6 kinase activity in human PBMC (131, 132).

In CD4<sup>+</sup>CD25<sup>-</sup> T cells, leptin induced strong STAT3 phosphorylation, while stimulation of CD4<sup>+</sup>CD25<sup>+</sup> T cells was not associated with a marked increase of phosphorylated STAT3 (52). SOCS3, a negative regulator of cytokine signaling, was activated by leptin blockade in CD4<sup>+</sup>CD25<sup>+</sup> T cells, in which the stimulation with anti-CD3/28 induced phosphorylation of ERK1/2 and subsequent cell proliferation (52). In the same subset of cells, the cyclin-dependent kinase inhibitor p27 (p27<sup>kip1</sup>), a molecule involved in the control of cell cycle and T cell energy) was elevated before and after anti-CD3/28 stimulation, and leptin neutralization induced degradation of this molecule, partly explaining the reversal of the anergic state and proliferation of these cells.

In macrophages, ERK-mediated phosphorylation of Ser<sup>727</sup> was found required for full stimulation of STAT3 by leptin (133). Macrophages express high levels of OB-Rb, and leptin stimulates STAT3 phosphorylation on both Tyr<sup>705</sup> and Ser<sup>727</sup> (133). While the MEK-1 inhibitor PD98059 had no effect on leptin-stimulated phosphorylation of STAT3 Tyr<sup>705</sup>, it greatly attenuated leptin's effects on STAT3 Ser<sup>727</sup> phosphorylation (133). Leptin-induced ERK activation in macrophages showed a biphasic pattern, with an initial reduction in ERK phosphorylation and a subsequent increased phosphorylation of ERK (133) paralleled by phosphorylation of Ser<sup>727</sup> and STAT3 DNA binding activity.

In human neutrophils, leptin was found to have activity as chemoattractant devoid of secretagogue properties but capable of inhibiting chemotaxis to classical neutrophilic



chemoattractants (134). This effect was dependent on the activation of intracellular kinases involved in F-actin polymerization and neutrophil locomotion (134). Indeed, p38 MAPK and Src kinase, but not ERK, were activated by short-term incubation with leptin (135). Moreover, the p38 MAPK inhibitor SB203580 and Src kinase inhibitor PP1, but not the MEK inhibitor PD98059, blocked neutrophils chemotaxis toward leptin (ref.).

In neutrophils, the ability of leptin to delay apoptosis seemed secondary to effects on the PI3K and MAPK-dependent pathways (136). Additionally, leptin delayed the cleavage of Bid and Bax, the mitochondrial release of cytochrome c and mitochondria-derived activator of caspase, as well as the activation of both caspase-8 and caspase-3 in neutrophils (136). The use of the specific inhibitors SB203580 and LY294002 (respectively a selective inhibitor of p38 MAPK and PI3K) blocked the anti-death effects of leptin, suggesting the key role of these pathways in transducing leptin-mediated antiapoptotic signals in neutrophils (136). The same effects of leptin mediated by PI3K and MAPK were observed on eosinophils (137).

It is possible that leptin signaling can also play a role in dendritic cell (DC) development and function. Optimal level of STAT3 activation induced by leptin are critically important for normal DC differentiation (138), and PI3K/Akt, p38 MAPK and NF- $\kappa$ B signalling pathways seem to have a central role in the survival of DC. In immature and LPS-induced matured DC in *db/db* mice, markedly reduced levels of active Akt and active PDK1 were observed (139). In line with those findings, active PTEN (a major negative regulator of PI3K/Akt signaling pathway) was also found up-regulated (139). Finally, both immature and LPS-matured *db/db* DC had decreased level of c-Raf phosphorylation at the inhibitory binding site (ser 259) (139), and *db/db* DC had decreased level of active p38 and phosphorylated I $\kappa$ B- $\alpha$  and high level of I $\kappa$ B- $\alpha$  (139). Taken together, these findings suggest that OB-R deficiency can impair signal transduction - including the Akt, MAPK and NF- $\kappa$ B pathways - in DC, affecting their survival and maturation.

## Conclusions

Leptin is a multifunctional hormone cytokine that is involved in processes as disparate yet intimate as metabolism and immune response. Because of its broad action, signalling pathways triggered by the OB-R can affect intracellular transduction pathways in different types of cells and/or promote cross talking among different cells.

Although many effects of leptin have been elucidated in recent times, the details of the signalling events that govern the response to leptin need further investigation to understand how the different pathways downstream of leptin are ultimately integrated. It will be also worthwhile to focus in the future on how leptin signalling integrates with the intracellular cascades activated by other factors in the immune cells, to have a better view of leptin's actions - ultimately allowing new strategies of therapeutic targeting.

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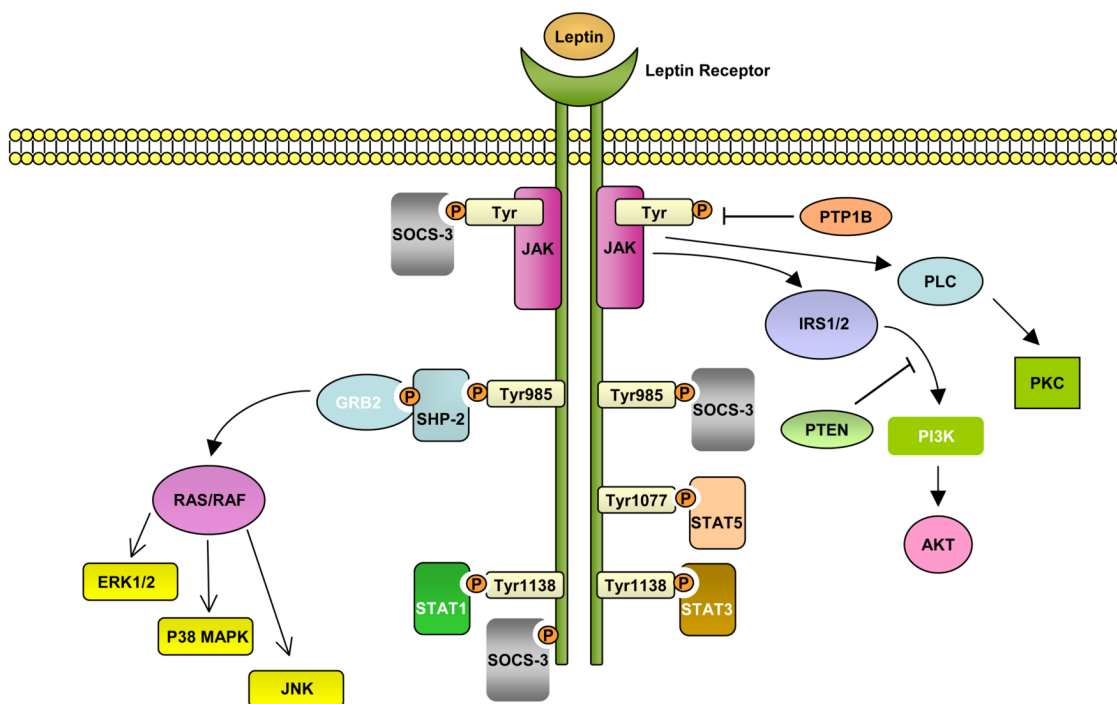
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### 1. Figure Schematic diagram of the pathways influenced by leptin

After leptin binds to the long isoform of the leptin receptor (OB-Rb), Jak2 is activated at the box 1 motif, resulting in the autophosphorylation of tyrosine residues and phosphorylation of tyrosines that provide docking sites for signalling proteins containing src homology 2 (SH2) domains. The autophosphorylated Jak2 at the box 1 motif can phosphorylate insulin receptor substrate1/2 (IRS1/2) that leads to activation of phosphatidylinositol 3-kinase (PI3K)/Akt pathway. Akt can regulate a wide range of targets including FOXO1 and NF- $\kappa$ B. Activation of NF- $\kappa$ B after leptin binding has been shown to induce Bcl-2 and Bcl-XL expressions. Leptin binding to OB-Rb can also activate the phospholipase C (PLC) for stimulation of c-jun N-terminal protein kinase (JNK) via protein kinase C (PKC).

Both Tyr<sup>1077</sup> and Tyr<sup>1138</sup> bind to STAT5, whereas only Tyr<sup>1138</sup> recruits STAT1 and STAT3. STAT3 proteins form dimers and translocate to the nucleus to induce expression of genes such as c-fos, c-jun, egr-1, activator protein-1 (AP-1) and suppressors of cytokine signaling 3 (SOCS3). SOCS3 negatively regulates signal transduction by leptin by binding to phosphorylated tyrosines on the receptor, to inhibit the binding of STAT proteins and the SH2 domain-containing phosphatase 2 (SHP2). SHP2 activates the mitogen-activated protein kinase (MAPK) pathways including extracellular signal-regulated kinase (ERK1/2), p38 MAPK and p42/44 MAPK through an interaction with the adaptor protein growth factor receptor-bound protein 2 (GRB2), to induce cytokine and chemokine expression in immune cells. SOCS2 binds to Tyr<sup>1077</sup> and might interfere with STAT5 binding. After stimulation with leptin, Src associated in mitosis protein 68 (Sam68) can form a complex with activated STAT3, leading to its dissociation from RNA. Sam68 can also be directly activated by Jak2 to phosphorylate IRS1/2 for Akt activation.