NUTS AND BOLTS

A comprehensive pathway map of IL-18-mediated signalling

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Abstract

Interleukin-18 (IL-18) is a member of the IL-1 family of cytokines and was initially described as an IFN-γ-inducing factor derived from anti-CD3-stimulated T-helper (Th)1 cells. IL-18 plays a significant role in the activation of hematopoietic cell types mediating both Th1 and Th2 responses and is the primary inducer of interferon- γ in these cells. The biological activity of IL-18 is mediated through its binding to the IL-18 receptor complex and activation of nuclear factor-κB (NF-κB), culminating in the production and release of several cytokines, chemokines, and cellular adhesion molecules. In certain cell types, IL-18 also activates mitogen-activated protein kinases (MAPKs) and phosphoinositide 3-kinase/ AKT serine/threonine kinase (PI3K/AKT) signaling modules leading to the production and release of proinflammatory cytokines. IL-18-mediated signaling acts as one of the vital components of the immunomodulatory cytokine networks involved in host defense, inflammation, and tissue regeneration. Albeit its biomedical importance, a comprehensive resource of IL-18 mediated signaling pathway is currently lacking. In this study, we report on the development of an integrated pathway map of IL-18/IL-18R signaling. The pathway map was developed through literature mining from published literature based on manual curation guidelines adapted from NetPath and includes information on 16 protein-protein interaction events, 38 enzyme-catalysis events, 12 protein translocation events, 26 activations/inhibition events, transcriptional regulators, 230 gene regulation events and 84 induced protein expression events. The IL-18 signaling pathway can be freely accessed through the WikiPathways database [\(https://www.wikipathways.org/index.php/Pathway:WP4754\)](https://www.wikipathways.org/index.php/Pathway:WP4754).

Keywords Inflammation · Proinflammatory cytokine · Post-translational modifications · Protein-protein interactions · Signaling pathways

Introduction

Interleukin-18 (IL-18), a member of the IL-1 family of cytokine, was initially termed as IFN-gamma inducing factor

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T. S. Keshava Prasad keshav@yenepoya.edu.in (Nakamura et al. [1989\)](#page-8-0) and was later defined as a proinflammatory cytokine with the ability to induce IFN γ (Dinarello 2001). The gene encoding $IL18$ is located on chromosomes 9 and 11 in mice and humans, respectively. It contains 7 exons

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with two distinct promoters on exon 1 and 2, including an interferon consensus sequence binding protein and a PU.1 binding sites (Nakanishi et al. [2001b\)](#page-8-0). IL18 encodes a 193 amino acid precursor, first synthesized as an inactive 24-kDa protein without a signal peptide, and is predominantly localized in the cytoplasm (Arend et al. [2008](#page-6-0); Carta et al. [2013](#page-6-0); Dinarello [2018\)](#page-6-0). The IL-18 precursor was primarily found to be expressed at high levels in Kupffer cells (Matsui et al. [1997;](#page-7-0) Tsutsui et al. [1997](#page-9-0)). Subsequent reports demonstrated that similar to other members of the IL-1 family such as IL-1 α and IL-33 but not IL-1β, IL-18 is constitutively expressed in most cell types including human peripheral blood mononuclear cells (PBMCs), macrophages, dendritic cells (DCs) (Chen et al. [2013\)](#page-6-0), osteoblasts, epithelial cells, chondrocytes, and epidermal keratinocytes (Gerdes et al. [2002](#page-7-0); Sanders and Mishra [2016](#page-8-0)) as well as in mouse peritoneal macrophages and spleen, thereby suggesting its vital pathophysiological role in health and disease. Additionally, the expression of a membrane-bound form of IL-18 in a subset of monocytes differentiated in vitro to macrophages by M-CSF has been reported (Bellora et al. [2012](#page-6-0)).

IL-18 similar to IL1-β, is synthesized as an inactive precursor form and cleaved to its active form (18KDa) by caspase-1 in response to inflammasome activation mediated by pathogen-associated molecular patterns (PAMPs) or damage-associated molecular patterns (DAMPs) recognition (Fabbi et al. [2015;](#page-6-0) Jacobs and Damania [2012;](#page-7-0) Zitvogel et al. [2012\)](#page-9-0). The active form of IL-18 is released primarily from macrophages and dendritic cells. Additionally, caspase-1 independent mechanisms of IL-18 processing have been reported. Importantly, caspase-8 mediated maturation and release of IL-18 in myeloid cells have been demonstrated as Fasdependent but independent of RIP3 or inflammasomes (Bossaller et al. [2012](#page-6-0); Tsutsui et al. [1999](#page-9-0)). Furthermore, proIL-18, similar to IL-1 α and IL-33, is also released from dying cells and is likely processed extracellularly by neutrophil proteases such as neutrophil-derived proteinase 3 (Sugawara et al. [2001\)](#page-8-0), Granzyme B produced mainly by NK cells and cytotoxic T lymphocytes (Omoto et al. [2010\)](#page-8-0), or by mast cell chymase (Omoto et al. [2006](#page-8-0)).

Signaling by IL-18 is mediated through IL-18 receptor, a heterodimeric complex consisting of the ligand-binding chain termed as IL-18Rα, and the co-receptor chain or the signaltransducing chain termed as IL-18Rβ (Boraschi and Tagliabue [2013;](#page-6-0) Kim et al. [2001](#page-7-0)) belonging to the IL-1 receptor family (Lee et al. 2004). The alpha chain (IL-18R α) was initially reported as an orphan receptor termed as the IL-1R-related protein (IL-1Rrp), which was later identified and demonstrated to have increased binding capacity to IL-18, leading to the activation of NF-κB-driven luciferase reporter gene (Torigoe et al. [1997\)](#page-8-0). The co-receptor or the beta chain (IL-18Rβ) has a low binding affinity to IL-18 alone but demonstrates a higher affinity when IL-18 is coupled to IL-18R α chain (Born et al. [1998\)](#page-6-0) resulting in the

initiation of downstream signaling and induction of inflammatory mediators (Bossaller et al. [2012](#page-6-0)). X-ray crystallography studies reveal similarities in the 3D-structure of IL-18 and its coreceptor with the IL-1β receptor complex (Tsutsumi et al. [2014\)](#page-9-0). IL-18R is mainly expressed in hematopoietic cells such as CD4 + NKT cells, mast cells, basophils, T-cells with the highest expression observed in NK cells driving its differentiation and activation (Nakamura et al. [2000\)](#page-8-0). The expression of IL18R on Th1 cells and B cells is mainly driven by IL-12 (Yoshimoto et al. [1998\)](#page-9-0) and in human NK and T cells by IL-12 and IFN α (Sareneva et al. [2000\).](#page-8-0) Recent studies also indicate the expression of IL18R in non-immune cells, such as nerve cells and epithelial cells (Nakanishi [2018\)](#page-8-0).

On binding to its receptor complex, IL-18 activates multiple signaling pathways by first recruiting MyD88, probably mediated by TRAM (Ohnishi et al. [2012](#page-8-0)), followed by IRAK4 (Suzuki et al. [2003\)](#page-8-0) and IRAK1/2 forming a molecular assembly called Myddosome (Tsutsumi et al. [2014](#page-9-0)). This complex recruits TRAF6 eventually activating NF-κB and mitogen-activated protein kinase (MAPK) pathways (Adachi et al. [1998](#page-5-0); Cao et al. [1996\)](#page-6-0). IL-18-mediated signaling is responsible for the induction of various inflammatory factors involved in both innate and adaptive immune responses. In an IL-12 or IL-15 dependent setting, IL-18 induces Th1-mediated immune responses by activating NK cells and Th1 cells, essential for host defense against intracellular pathogen infection through IFNγ production (Chaix et al. [2008](#page-6-0)). Additionally, IL-18 also induces the production of TNF and FasL involved in growth, survival, and apoptosis (Tsutsui et al. [1999;](#page-9-0) Zhang et al. [2011](#page-9-0)). However, independent of synergistic signals, IL-18 potentially induces IL-4 and IL-13 production in T cells, NK cells, mast cells, and basophils driving a Th2 response (Nakanishi et al. [2001a;](#page-8-0) Yoshimoto et al. [1999\)](#page-9-0). Other mediators induced by IL-18 include inducible nitric oxide; cyclooxygenase (Cox-2); proinflammatory cytokines IL-1β, IL-6; chemokines IL-8, MCP-1, and MIP-1 α ; intracellular adhesion molecule ICAM-1; growth factor GM-CSF (Dinarello [1999;](#page-6-0) Kohka et al. [1998](#page-7-0); Nakanishi et al. [2001b](#page-8-0); Olee et al. [1999\)](#page-8-0). IL-18 also induces the production of cytokines such as IL-12, IL-2 (Okamura et al. [1998;](#page-8-0) Takeuchi et al. [1997\)](#page-8-0) which was observed in experimental murine dengue models, where the synergetic effect of IL-12 and IL-18 are considered as critical mediators in dengue host defense (Fagundes et al. [2011\)](#page-6-0). Given the pleiotropic functions of IL-18, regulation of IL-18 activity is vital to prevent an aberrant immune response. The cell activation mediated by IL-18 is mainly regulated by IL-18 binding protein (IL-18BP) (Dinarello et al. [2013](#page-6-0)) which binds to IL-18 at a higher affinity than IL-18R α (Novick et al. [1999\)](#page-8-0) and thereby suppresses its Th-1 response in physiological conditions (Kim et al. [2000;](#page-7-0) McInnes et al. [2000](#page-7-0)). Recent evidence suggests IL-37, an anti-inflammatory cytokine belonging to the IL1 family, acts as a negative regulator by binding to IL-18R α with low affinity, resulting in the loss of recruitment of IL-18R β and downstream signaling (Nold-Petry et al. [2015](#page-8-0)).

The role of IL-18 in host defense mechanisms against intracellular pathogens has been well documented. Increasing evidence also suggests its vital role in the maintenance of metabolic homeostasis (Lindegaard et al. [2013;](#page-7-0) Netea et al. [2006;](#page-8-0) Zorrilla et al. [2007\)](#page-9-0) and protection against the development of metabolic syndrome (De Nardo and Latz [2011](#page-6-0); Murphy et al. [2016\)](#page-8-0). Elevated signaling by IL-18 has been implicated in a number of pathological conditions such as several acute and chronic inflammatory diseases, including COPD, asthma, atopic dermatitis, rheumatoid arthritis, lupus erythematosus, atherosclerosis, graftversus-host disease, renal diseases and hepatitis (Briend et al. [2017;](#page-6-0) Lee et al. [2015](#page-7-0); Liew et al. [2003](#page-7-0); Sharma et al. [2009](#page-8-0); Xu et al. [2017](#page-9-0)). Interestingly, the role of IL-18 in ameliorating cancer metastasis has been increasingly recognized (Dupaul-Chicoine et al. [2015](#page-6-0); Smyth et al. [2004\)](#page-8-0).

Despite the importance of IL-18-mediated signaling, detailed documentation of IL-18 mediated signaling events is currently lacking. Though the signaling pathway map of IL-18 is available in KEGG and Reactome pathway database, it currently provides only a generic view of the signaling modules regulated by IL-18. In the current study, we developed a resource of signaling events mediated by IL-18/IL-18R in similar lines to earlier reports on comprehensive signaling maps (Bhat et al. [2019;](#page-6-0) Pinto et al. [2018;](#page-8-0) Subbannayya et al. [2014\)](#page-8-0). Our resource contains the complete cascade of IL-18 signaling, including 319 molecules and 402 reactions stimulated by IL-18 through manual annotation of the data mined from the publicly available literature. The signaling pathway map is made available through WikiPathways [\(https://www.](https://www.wikipathways.org/index.php/Pathway:WP4754) [wikipathways.org/index.php/Pathway:WP4754](https://www.wikipathways.org/index.php/Pathway:WP4754)).

Methodology

Literature survey and catalog of signaling events

An extensive literature search was carried out in PubMed using key terms "IL-18" OR "IL18" OR "Interleukin-18" OR "IL-18 induced" AND "Signaling" OR "Pathway" NOT 'Review.' The downstream signaling cascade under the influence of IL-18 was screened, and only those molecular events regulated/influenced by IL-18 were considered for further documentation as per NetPath criteria (Kandasamy et al. [2010\)](#page-7-0). From the screened research articles, information on IL-18 induced protein-protein interactions (PPIs), post-translational modifications (PTMs), gene regulation (upregulated or downregulated in response to IL-18 stimulation), protein activation/inhibition and protein translocation events were systematically annotated based on NetPath criteria. The information manually annotated includes the experimental conditions- in vivo or in vitro, cell line model, protein site/domain involved in PPI, transcription regulators (if present). Additional information about post-translational modifications, including the residue and site of modification, has been

included. The curated data were subjected to quality control by both internal and external review process.

Development of signaling pathway map

The signaling pathway map of IL-18-mediated signaling events has been developed using PathVisio software (van Iersel et al. [2008\)](#page-9-0). The final IL-18 mediated signaling pathway was visualized based on NetPath criteria and implemented in WikiPathways ([https://www.wikipathways.org/index.php/](https://www.wikipathways.org/index.php/WikiPathways) [WikiPathways](https://www.wikipathways.org/index.php/WikiPathways)). The reactions activated/induced by IL-18 are arranged in a topological order starting from the ligandreceptor interaction to signaling modulators, transcription factors, and transcriptionally regulated genes. Pathway modules such as MAPK signaling, PI3K/AKT signaling, which are regulated by IL-18, have been depicted in the pathway map.

Results

The development of the IL-18/IL-18R signaling pathway map involved screening of a total of 1748 research articles from PubMed until July 2019 using keywords as described in the methods section. Of these, 292 research articles had information related to IL-18 induced signaling as compared with an unstimulated condition in humans or mammals. From the manually screened articles, a total of 406 IL-18 mediated signaling events arbitrated by 324 molecules were curated (Supplementary Data S1-S7). These events included 16 protein-protein interaction events, 38 enzyme-catalysis events, 12 protein translocation, 26 activations/inhibition 230 gene regulation, and 84 induced protein expression events were manually annotated based on modified NetPath criteria (Kandasamy et al. [2010\)](#page-7-0). The PPIs included both 'binary' and 'complex' (multimeric) associations. In the case of enzyme-substrate reactions, depending on the information available on the upstream enzyme, the reactions were classified as "direct" or "induced" reactions. Additionally, the site and residue of PTMs were documented depending on the available literature evidence. In all, information was available for 38 PTM events. All the events that have been annotated include a link to the corresponding references.

The pathway map was drawn in PathVisio. All curated events were included in the map. Pathway topology was determined based on known signaling modules, by comparison with existing pathway resources, and comparing with IL-1 signaling (Fig. [1](#page-3-0)).

Summary of IL-18 mediated signal transduction

IL-18 is an important inflammatory cytokine that initiates signaling by binding to its receptor-binding protein IL-18 alpha chain (IL-18R α) and co-receptor (IL-18R β), forming a ternary complex (Dinarello et al. [2013](#page-6-0)). The central role of IL-18R α

Fig. 1 Schematic representation of IL-18 signaling pathway. Schematic representation of IL-18 induced signaling reactions. The signaling pathway map represents molecules involved in ligand-receptor interactions and IL-18 induced downstream molecular events including molecular

association, catalysis, translocation, and gene regulation events. Information regarding the post-translational modification site and the residue is also shown in the pathway.

was demonstrated by Hoshino K et al. using IL-18Rα- deficient mice, where they observed a lack of Th1 response upon IL-18 stimulation. Other studies also reported inhibition of c-Jun Nterminal kinase and NF-κB activation in Th1 cells from IL-

18Rα−/− mice (Hoshino et al. [1999;](#page-7-0) Yoshimoto et al. [1998\)](#page-9-0). Following the complex formation, the classical Myddosome complex (MYD88/ IRAK/TRAF6) is recruited (Kojima et al. [1998](#page-7-0)), and the recruitment of MYD88 likely involves TRAM, a sorting adaptor known for its role in TLR signaling (Ohnishi et al. [2012](#page-8-0)). It is presumed that similar to IL-1/IL1R signaling, IL-18 mediated TRAF6 ubiquitination of IκBα kinase (CHUK) results in its degradation and concomitant activation and release of NFκB that translocates to the nucleus to induce transcription of inflammatory genes, including proinflammatory cytokines, chemokines, and adhesion molecules (Chandrasekar et al. [2004](#page-6-0), [2006b](#page-6-0), [2008;](#page-6-0) Doffinger et al. [2001](#page-6-0); Finotto et al. [2004](#page-7-0); Lee et al. [2004](#page-7-0); Leyfer et al. [2004](#page-7-0); Reddy et al. [2010,](#page-8-0) [2011](#page-8-0); Zabalgoitia et al. [2008](#page-9-0)). IL-18 also activates the MAPK cascade to induce STAT3 activation, IFNγ production and resultant cytotoxic activity in NK cells (Kalina et al. [2000\)](#page-7-0). Additionally, IL-18 increases the expression of nitric oxide synthase (iNOS) in leukocytes via IL-18R/ p38-MAPK phosphorylation (Jablonska et al. [2008\)](#page-7-0).

IL-18 also activates phosphatidylinositol-kinase (PI-3K)/AKT/ mammalian target of rapamycin (mTOR) pathway to regulate pathogenic Th17 cell differentiation and expression of Bcl-xL and Bcl2 (Deason et al. [2018](#page-6-0); El-Darawish et al. [2018](#page-6-0)). Additionally, Akt phosphorylation is also induced in several hematopoietic and epithelial cells such as neutrophils, basophils, macrophages, HCF, HCMEC, VSMC, SMCs, and mouse aortic smooth muscle cells (ASMC) (Chandrasekar et al. [2005,](#page-6-0) [2006b,](#page-6-0) [2008](#page-6-0); Finotto et al. [2004](#page-7-0); Kroeger et al. [2009;](#page-7-0) Morel et al. [2002](#page-8-0); Reddy et al. [2008,](#page-8-0) [2010,](#page-8-0) [2011;](#page-8-0) Venkatesan et al. [2009;](#page-9-0) Yoo et al. [2005\)](#page-9-0). In rheumatoid arthritis synovial tissue fibroblasts, IL-18 induced the cell surface expression of VCAM1 through two distinct signaling mechanisms involving direct activation of Src by IL-18 which in turn activates transcription factor AP-1 through Ras-Raf1 induced ERK1/2 activation. Independent of Src, IL-18 also activated the PI3K-AKT module to induce VCAM1 (Morel et al. [2002\)](#page-8-0). Crosstalk between GRP78, phospho-Akt, NF-κB, and XIAP is necessary for the aggressive growth of cancer via CASP3 in normal human neonatal foreskin epidermal keratinocytes (Hosotani et al. [2008](#page-7-0)). IL-18 also induces JNK/Sp1 signaling and MMP-9 expression in part via EMMPRIN/ BSG and through MAPK mediated AP-1 and NFkB activation (Chandrasekar et al. [2006a;](#page-6-0) Reddy et al. [2010\)](#page-8-0).

In addition to the activation and inhibition of signaling events in different cell types, the downstream signaling proteins that are induced by IL-18 also varies with cell types. IL-18 induces the activation and nuclear translocation of cytosolic NFκB1 proteins in normal human epidermal melanocytes (NHEM), cortical neurons, myelomonocytic cells, cardiac microvascular endothelial cells (EC) which results in the production and release of MIP1-alpha and -beta, MIP2-alpha and -

beta, and IL-8 (Leyfer et al. [2004;](#page-7-0) Zabalgoitia et al. [2008;](#page-9-0) Zhou et al. [2013](#page-9-0)). In chondrocytes, embryonic stem (ES) cells, human cardiac microvascular endothelial cells, and human alveolar basal epithelial cells, IL-18 induces the expression of IL-6 and IL-8. In murine peritoneal macrophages, IL-18 produces TNF α , IL-6, IL-1 α , and IL-1 β , which leads to joint inflammation resulting in cartilage destruction (Dai et al. [2004](#page-6-0); Hosotani et al. [2008;](#page-7-0) Volin and Koch [2011](#page-9-0)). IL-18 stimulation also induces IL23A, IL17A, and MCP-1 expression in macrophage and microglia (Chandrasekar et al. [2004;](#page-6-0) Cheung et al. [2005](#page-6-0); Olee et al. [1999\)](#page-8-0).

Regulation of IL-18 expression and activity occurs at several levels. Several repressors at the genomic and posttranscriptional levels have been identified, including B cell lymphoma 6 protein (Bcl6) that has been demonstrated as one of the critical repressors of the IL18 gene (Takeda et al. [2003](#page-8-0)). miRNAs such as miR-346 in synovial cells (Alsaleh et al. [2009](#page-5-0)), miR-197 in hepatitis B and chronic liver failure (Chen et al. [2013\)](#page-6-0), miR-134 in adultonset Still's disease (Liao et al. [2017](#page-7-0)) have shown to lower IL-18 expression and secretion. The biological activity of soluble IL-18 is mainly mediated by IL-18 binding protein (IL-18BP) (Dinarello et al. [2013\)](#page-6-0) which binds to IL-18 at a higher affinity than IL-18Rα (Novick et al. [1999](#page-8-0)) and thereby suppressing its Th-1 response in physiological conditions (Kim et al. [2000;](#page-7-0) McInnes et al. [2000](#page-7-0)). A feedback response by IL-18BP to elevated IL-18 helps regulate untoward IFN-γ signaling, thus reducing the damage resulting from excessive "free" IL-18. IL-18BP also acts as a carrier protein circulating with IL-18 and is required for the normal NK cell function (Harms et al. [2017\)](#page-7-0). The severity of the disease is associated with an imbalance of both IL-18 and IL-18BP, with elevated levels of free IL-18 associated with the risk of development of disorders such as autoimmune disorders, schizophrenia, sepsis, among others (Michels et al. [2015](#page-7-0)). Several studies have demonstrated considerable changes in pathology when administered with IL-18BP in experimental murine models of arthritis, colitis, endotoxic shock, ischemia-reperfusion injury and type 1 diabetes (Banda et al. [2003;](#page-6-0) Colafrancesco et al. [2012](#page-6-0); Faggioni et al. [2001](#page-6-0); He et al. [2008](#page-7-0)). A recent study also demonstrated the association of IL-18BP deficiency with the development of exacerbated colitis and arrested maturation of goblet cells (Nowarski et al. [2015\)](#page-8-0). Considering the vital role of IL18BP in IL18 mediated signaling, further studies are required for the better understanding of the mechanism of action of IL-18BP in the IL-18 signaling pathway. The signaling pathway map (Fig. [1\)](#page-3-0) provides a complete picture of the signaling events mediated by IL-18, and we hope that this will enable researchers to obtain insights into the signaling cross-talk across the various signaling modules regulated by IL-18.

IL-18 signaling in disease

IL-18 and its receptor (IL-18R) are closely involved in regulating both adaptive and innate immune responses. With increasing evidence suggesting a crucial role of inflammasomes in regulating IL-1β and IL-18 secretion, several genetic disorders associated with the components of inflammasomes may likely result in their increased levels in circulation, and thereby resulting in autoinflammatory syndromes such as cryopyrin-associated periodic syndrome (CAPS) (Hoffman et al. [2001\)](#page-7-0), macrophage-activating syndrome (MAS) (Canna et al. [2014;](#page-6-0) Girard-Guyonvarc'h et al. [2018;](#page-7-0) Yasin et al. [2019\)](#page-9-0), adult-onset Still's disease (AOSD) (Girard et al. [2016\)](#page-7-0) and several other autoinflammatory and autoimmune diseases including inflammatory bowel disease, schizophrenia and sarcoidosis (McInnes et al. [2000](#page-7-0); Zhang et al. [2016](#page-9-0)). Additionally, genetic polymorphisms in IL18 or IL18R also contribute to allergic reactions, metabolic and autoimmune disorders. Several studies have demonstrated the presence of IL-18 and its receptor subunit in the neurons confirming its ability to cross the blood-brain barrier (Alboni et al. 2010). Studies have also tried to demonstrate its role in neurophysiological and neuro-pathological diseases (Andoh et al. [2008;](#page-6-0) Culhane et al. [1998](#page-6-0); Wang et al. [2006;](#page-9-0) Wheeler et al. [2000\)](#page-9-0). The level of IL-18 and IL-18R subunits can be induced and regulated in the central nervous system (CNS) as demonstrated in hippocampus of mouse by increased level of IL-18 and IL-18R α during kainic acid (KA)-induced excitotoxicity and marked increase of IL-18 in mouse microglia during hypoxic-ischemic brain injury (Hedtjarn et al. [2002](#page-7-0); Jeon et al. [2008](#page-7-0); Miyoshi et al. [2008](#page-7-0)). Studies have reported increased level of IL-18 in autoimmune neurodegenerative diseases such as multiple sclerosis (MS), Alzheimer's disease (AD) and autoimmune encephalomyelitis (EAE) confirming its pivotal role in the pathogenesis of the disease (Huang et al. [2004;](#page-7-0) Jander and Stoll [1998](#page-7-0); Motta et al. [2007;](#page-8-0) Nicoletti et al. [2001;](#page-8-0) Ojala et al. [2009;](#page-8-0) Wildbaum et al. [1998](#page-9-0)). It has also been reported in mediating angiogenesis and vascular remodeling in inflammageing (Fahey and Doyle [2019;](#page-7-0) Rodriguez-Menocal et al. [2014](#page-8-0)). Induced IL-18 and IL-18R α play a critical role in the pathogenesis of cigarette smoke-induced pulmonary emphysema and inflammation (Kang et al. [2007\)](#page-7-0). IL-18 has been associated with inflammatory disorders and numerous reactions in human body such as atherosclerosis, atopic eczema, COPD, maintenance of homeostasis, development of autoimmune disease along with the significant role in the prevention of infectious disease such as tuberculosis(Akdis et al. 2016; Blankenberg et al. [2002](#page-6-0); Briend et al. [2017](#page-6-0); Lee et al. [2015;](#page-7-0) Terada et al. [2006](#page-8-0); Wawrocki et al. [2016](#page-9-0); Wawrocki et al. [2019](#page-9-0)). Hence, the neutralization of IL-18 antibody can be utilized in the preclinical models of inflammation, metastasis, and tissue injury (Lauw et al. [2002](#page-7-0); Nakajima and Owen [2012](#page-8-0); Yu et al. [2002](#page-9-0)). The

inhibition of IL-18 in autoimmune diseases such as Crohn's disease and psoriasis could act as an attractive therapeutic regimen as IFNγ production decreases with the reduced IL-18 (Dinarello et al. [2013;](#page-6-0) Joosten et al. [2000\)](#page-7-0).

Conclusions

IL-18, a member of the IL-1 family, is reportedly involved in the activation of hematopoietic cell types and plays a crucial role in the regulation of autoimmune diseases and cancers. The importance IL-18 mediated signaling pathway and the absence of signaling resources led us to drive manual curation efforts from published literature and catalog signaling events upon IL-18 stimulation. We anticipate that the detailed pathway map of IL-18 and the compendium of IL-18 signaling events emanating from this study will serve as a useful resource for researchers. Besides expanding understanding of IL-18 signaling in both normal physiology and disease, the availability of this pathway resource on the WikiPathways resource will enable researchers to use this signaling pathway map to carry out pathway analysis of high-throughput omics data obtained from various platforms.

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Compliance with ethical standards

Conflict of interest The authors report no conflicts of interest.

Abbreviations IL-18, Interleukin-18; IFN γ , Interferon gamma; IL-1 β , Interleukin-1beta; M-CSF, Macrophage colony-stimulating factor; NF-κB, Nuclear factor-κB; PPI, Protein-protein interaction; IL-18BP, Interleukin-18 binding protein

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