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p38α senses environmental stress to control innate immune responses via mTOR

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Abstract

The mitogen-activated protein kinase p38a senses environmental stressors and orchestrates inflammatory and immunomodulatory reactions. However, the molecular mechanism how p38a controls immunomodulatory responses in myeloid cells remains elusive. We found that in monocytes and macrophages, p38a activated the mechanistic target of rapamycin (mTOR) pathway *in vitro* and *in vivo*. p38a signaling in myeloid immune cells promoted interleukin (IL)-10 but inhibited IL-12 expression via mTOR and blocked the differentiation of proinflammatory CD4+ T helper 1 cells. Cellular stress induced p38a-mediated mTOR activation that was independent of phosphoinositide 3-kinase (PI3K) but dependent on the kinase MK2 and on the inhibition of TSC1/TSC2 (tuberous sclerosis gene 1 and 2), a negative regulatory complex of mTOR signaling. Remarkably, p38a and PI3K concurrently modulated mTOR to balance IL-12 and IL-10 expression. Our data links p38a to mTOR signaling in myeloid immune cells that is decisive for tuning the immune response in dependence on the environmental milieu.

Introduction

Recognition of pathogen-associated molecular patterns (PAMPs) by innate immune receptors triggers inflammatory and immune responses involving several signaling molecules including the mitogen-activated protein kinases (MAPK) (1, 2). The MAPK p38a (also known as MAPK14) is one of four homologs of mammalian p38 and is essential in innate and adaptive immune signaling cascades (3). p38a is activated by diverse stimuli including Toll-like receptor (TLR) ligands, cytokines, and physicochemical stress signals such as ultraviolet (UV) irradiation, heat or osmotic shock, arsenite or anisomycin (4). p38a

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is ubiquitously expressed in most cell types and regulates diverse functions such as cell proliferation, differentiation, apoptosis, tissue repair, tumorigenesis, or inflammation (4, 5). For example, p38a was previously identified as activator of the proinflammatory cytokines interleukin (IL)-1 β , IL-6, tumor necrosis factor (TNF)- α , or cyclo-oxygenase 2 (6–8). Several kinases are activated by p38a such as mitogen- and stress-activated kinase (MSK) 1 and 2, MAPK-interacting serine/threonine-protein kinase (MNK) 1 and 2, or MAPK-activated protein kinase (MK) 2 and 3 that mediates TNF- α production (9, 10).

The finding that inhibiting p38 blocks lipopolysaccharide (LPS)-induced proinflammatory cytokine production (7) initiated the development of a wide range of p38 inhibitors for treatment of chronic inflammatory diseases such as rheumatoid arthritis (RA), psoriasis or Crohn's disease (11). SB203580 is a competitive inhibitor of p38a and p38 β by blocking ATP binding to the kinase, while BIRB796 is an allosteric inhibitor of p38a, p38 β , and p38 γ (12, 13). Remarkably, so far p38 inhibitors failed in clinical trials due to adverse and inflammatory side effects such as liver toxicity or skin rashes (11).

Recently, a more complex role of p38a has been reported (14–16). Expression of p38a in myeloid cells limits inflammation in an UV-induced irradiation model (14). These immunomodulatory effects of p38a may be mediated by the induction of the anti-inflammatory cytokine IL-10 (14, 15) and the inhibition of proinflammatory IL-12 (14, 16). However, the downstream pathway that controls coordinated IL-10 and IL-12 expression by p38a has remained elusive.

The classical insulin signaling pathway consisting of phosphoinositide 3-kinase (PI3K), Akt, and mechanistic target of rapamycin (mTOR) complex 1 (mTORC1) has recently emerged as key regulator of innate immune cell homeostasis (17-20). Stimulation of innate immune cells by TLR ligands activates the mTOR pathway and it is in fact a major pathway activated in LPS-stimulated macrophages based on phosphoproteomics (21). The function of PI3K-Akt-mTOR is cell type-specific, but it has been shown that inhibition of PI3K by wortmannin or mTOR by rapamycin in myeloid cells, such as human monocytes, macrophages or myeloid dendritic cells, enhances IL-12 production but blocks the release of IL-10 in vitro and in vivo (22-28). Tuberous sclerosis 2 (TSC2) is a tumor suppressor that is phosphorylated and inactivated by the protein kinase Akt, which itself is activated by PI3K (29). TSC2 forms a heterodimeric complex with TSC1 and negatively regulates mTOR (29). Conversely, knockdown of TSC2 in human monocytes or macrophages enhances IL-10 but inhibits IL-12 production (24, 30). In line, genetic inactivation of mTORC1 reduces IL-10 production in intestinal CD11c+CD11b+ dendritic cells (31). On the other hand, TSC1deficient macrophages show elevated production of TNF-a and IL-12p40 (32). Despite these observations, the precise regulatory units and upstream pathways controlling mTORdependent cytokine production are still unclear.

An outstanding question is how myeloid immune cells adapt and coordinate their immune response to an infectious trigger towards the status of the environmental milieu; e.g. how to avoid detrimental tissue-destructive CD4+ Th1 responses under conditions of tissue repair. Moreover, it is imperative to explore the molecular signaling pathways that regulate p38a-mediated immune responses for a deeper understanding of the effects of p38 inhibitors for human health and disease. Therefore, we tested whether p38a is connected to PI3K-TSC2-mTOR signaling to regulate innate inflammatory responses. We found that TLR ligands or environmental stress activate the TSC2-mTOR pathway via p38a and MK2 to regulate the balance of IL-12 and IL-10. Importantly, p38a acts in parallel to PI3K to control the IL-12/IL-10 equilibrium in response to the environmental milieu.

Material and Methods

Reagents

LPS (*E. coli* 0111:B4), wortmannin, anisomycin, and rapamycin were from Sigma. SAC (PANSORBIN) and SD169 were from Calbiochem. BIRB796 was a kind gift of Sir Philip Cohen or purchased from Axon Medchem. SB203580 was from Tocris Bioscience and IFN- γ from R&D. Heat-killed cells of *Listeria monocytogenes* (L.m.) were prepared by incubating the viable log-phase bacterial suspension at 70°C for 1 h. For UV exposure, cell culture plates were placed on a 20×20 UV-transilluminator (MWG Biotech) and activated with UV light for 10 sec, 30 sec or 1 min.

Cell culture

Human peripheral blood mononuclear cells (PBMC) and peripheral human myeloid DCs were isolated as described (24). Monocytes were isolated from PBMCs by MACS using CD14 Microbeads (Miltenyi Biotec). RPMI 1640 supplemented with 2 mM L-glutamine, 100 µg/ml streptomycin, 100 U/ml penicillin (all from Gibco), and 10% FCS (Hyclone) was used as culture medium. Mouse embryonic fibroblasts (MEFs) were cultured in DMEM containing 4.5 g/L glucose, 2 mM L-glutamine, 100 µg/ml streptomycin, 100 U/ml penicillin, and 10% FCS. *Tsc2*^{+/+} *p53*^{-/-} and *Tsc2*^{-/-} *p53*^{-/-} as well as *Tsc1*^{+/+} and *Tsc1*^{-/-} MEFs were described previously (33, 34). p85a^{-/-} p85β^{-/-} MEFs were a kind gift of Lewis Cantley. p38a^{fl/fl} and p38a^{Δ M} mice were described previously (14). Bone marrow-derived macrophages (BMDMs) from mice were isolated and grown as described (35) and were replated one day prior to stimulation in full medium containing 2% FCS. *Mk2*^{-/-} immortalized murine macrophages stably reconstituted with MK2 or MK2^{K79R} were described previously (36).

Measurement of cytokine production

Cells were pretreated for 90 min with the indicated concentrations of SB203580, BIRB796, rapamycin, or wortmannin and then stimulated with 100 ng/ml LPS (+30 ng/ml IFN- γ as indicated), 75 µg/ml SAC, or 10⁷ L.m. in 48-well plates. Cell-free supernatants were collected after 22–24 h as indicated. Human and murine cytokines were determined by the Luminex bead system with beads from R&D Systems and Affymetrix and read on a Luminex 100 reader.

LPS injection

 $p38\alpha^{\Delta M}$ and $p38\alpha^{fl/fl}$ mice were housed and maintained at the Massachusetts General Hospital and Harvard Medical School. Mice were injected intraperitoneally with 30 µg/ mouse LPS. After 4 h, serum samples were taken and spleens were isolated. Cytokine levels in the sera were measured by Luminex. Homogenization of mouse tissue was performed by using the Precellys-ceramic kit 2.8 mm and the Precellys 24 tissue homogenizer (both from peQLab).

T cell differentiation

Monocytes were incubated with medium, 200 nM BIRB796, 2 μ M SB203580 or 100 nM rapamycin for 90 min and stimulated with 100 ng/ml LPS for 24 h. The cells were then washed with PBS and incubated with allogeneic T lymphocytes at a ratio of 1:1 in 24-well plates in RPMI complete medium. After one week, IFN- γ production was determined in cell-free supernatants by Luminex. The primed cells were further activated for 5 h with 50 ng/ml phorbol-12-myristat-13-acetat (PMA) and 200 ng/ml ionomycin (both from Sigma) in the presence of 10 μ g/ml brefeldin A (Sigma) for the last 3 h. Afterwards, cells were stained

with FITC–labeled anti-IFN- γ , PE-labeled anti-IL-4, and APC-labeled anti-CD4 (all BD Bioscience) and analyzed by flow cytometry.

Analysis of signal transduction events

Monocytes, BMDMs, or 70% confluent MEFs starved overnight were treated and stimulated as indicated. Extract preparation and Western Blotting was done as described (24). Antibodies were p-p70S6K (Thr389), p70S6K, p-4E-BP1 (Thr37/46), p-p38 (Thr180/Tyr182), p-S6 (Ser 240/244), p-Akt (Ser473), p-MK2 (Thr334), GAPDH, S6-ribosomal protein, p38 MAPK, p38a MAPK, p38β MAPK, p38δ MAPK, Tuberin/TSC2 (all Cell Signaling Technology), p-Erk (Tyr204), IkBa and p38 (Santa Cruz Biotechnology).

Quantitative RT-PCR

RNA from human monocytes or murine immortalized macrophages was extracted in TRIzol (Invitrogen). cDNA was generated by Superscript II (Invitrogen). mRNA levels were determined by TaqMan Gene Expression Assays (Applied Biosystems) on a StepOnePlusTM Real-Time PCR System and normalized to ubiquitin.

Transfection of MEFs

Tsc2–/– MEFs in 6-well plates at 20–40% confluency were transfected in DMEM without antibiotics with 1µg pcDNA3-HA-TSC2 WT, pcDNA3-HA-TSC2 S1210A, or empty vector) with Lipofectamine 2000 Reagent (Invitrogen) for 36h and afterwards starved for 12 h in DMEM without antibiotics and FCS before stimulation.

Immunoflourescence microscopy

Cells were applied to 8-well PermanoxTM chamber slides (Lab-Tek Chamber Slide System), fixed with 4% paraformaldehyde, quenched with 100mM glycine, permeabilized with methanol, blocked with 1% BSA, and stained with p-S6 antibody, p-MK2 antibody or isotype control overnight at 4° C. Cells were stained with AlexaFluor488-labeled goat anti-rabbit IgG (Invitrogen) followed by nuclear tracking using 0.1 μ g/ml Hoechst-33342 (Invitrogen) and mounted in Vectashield® mounting medium.

Statistics

Results are expressed as means \pm SEM. Student's t test was used to detect statistical significance.

Results

p38α modulates IL-12 and IL-10 production in mice and men

To evaluate the role of p38a in the myeloid immune system, we generated bone marrowderived macrophages (BMDM) from mice with a deletion of p38a in cells expressing the lysozyme M gene (p38a^{Δ M}) and stimulated these cells with LPS, *Staphylococcus aureus* (SAC), or heat-killed *Listeria monocytogenes* (L.m.) (Fig. 1A). Deficiency of p38a enhanced IL-12p40 and IL-12p70 expression while the anti-inflammatory cytokine IL-10 was blocked compared to controls carrying homozygously the floxed p38a gene (p38a^{fl/fl}) (Fig. 1A). Other cytokines such as IL-1 β , IL-23, or TNF-a were not significantly altered (data not shown). Injection of LPS into p38a^{Δ M} mice similarly deviated the production of IL-12 and IL-10 *in vivo* (Fig. 1B). Next, we characterized the precise function of p38a, which is the most highly expressed p38 isoform in human monocytes. We found that inhibition of p38 with different concentrations of SB203580 or BIRB796 as well as with the mTOR inhibitor rapamycin strongly increased the production of IL-12p40 and IL-12p70 but blocked secretion of IL-10 after stimulation with LPS, SAC, or L.m. (Fig. 1,C–F). Notably, SB203580 did not augment IL-12p40 production in L.m.-stimulated monocytes. Enhanced IL-12p40 but reduced IL-10 expression was also observed in SB203580- or BIRB796- treated peripheral human myeloid dendritic cells stimulated with LPS, SAC or L.m. (data not shown).

The surface expression of the costimulatory molecule CD86, important for T cell activation and priming, was enhanced upon p38 blockade in human monocytes after stimulation with LPS, SAC or L.m. (data not shown). In line, we found that inhibition of p38 or mTOR in monocytes stimulated with LPS strongly enhanced the production of IFN- γ (Fig. 1G) and the differentiation of CD4+ Th1 cells (Fig. 1H) in an allogeneic T cell activation model. In summary, these data show that p38a differentially regulates the expression of IL-12 and IL-10 in human monocytes and DCs as well as in BMDM *in vitro* and *in vivo* in response to microbial insult.

p38 activation stimulates mTOR signaling

The concurrent regulation of IL-12/IL-10 by p38 and mTOR inhibitors indicated that these molecules might be connected. Therefore, we explored whether inhibition of p38 or mTOR might mutually influence the other kinase. Rapamycin did not modulate the phosphorylation of p38 or its downstream kinase MK2 after stimulation of human monocytes with LPS but blocked the phosphorylation of the mTOR substrates p70S6K and 4EBP1 (Fig. 2A). In contrast, inhibition of p38 with either SB203580 or BIRB796 blocked MK2 activation as expected, but also decreased the phosphorylation of 4EBP1 and p70S6K as well as S6 suggesting that p38 also activates mTOR signaling. Total levels of the investigated proteins were not altered by treatment with the inhibitors (data not shown). Interestingly, phosphorylation of Akt at Ser473 was blocked while activation of Erk was enhanced with BIRB796 and SB203580 (Fig. 2, A and C). However, inhibition of Erk did not inhibit mTOR and did not modulate the production of IL-12 or IL-10 (data not shown). Degradation of $I\kappa B$ -a was not influenced by either p38 or mTOR (Fig. 2A). Activation of p38 by the two environmental stress signals anisomycin or UV also activated mTOR signaling in human monocytes in a p38-dependend manner (Fig. 2, B and C). SD169, another reported inhibitor of p38a, was without effect in monocytes (Fig. 2, A and C). Next we tested whether hyperactivation of p38 with anisomycin in the presence of LPS or SAC could influence cytokine expression. Indeed, anisomycin reduced IL-12p40 mRNA levels but increased IL-10 mRNA levels in LPS- or SAC-stimulated macrophages (Fig. 2, D and E). Remarkably, hyperactivation of p38 with anisomycin further increased mTOR activity in LPS-activated monocytes (Fig. 2F). These results indicate that p38 activates the mTOR pathway and thereby regulates the expression of IL-12 and IL-10.

p38α deletion blocks mTOR activation in vitro and in vivo via MK2

To extend our results genetically, we examined the mTOR signaling pathway in $p38a^{\Delta M}$ BMDM. Deletion of p38a in macrophages abolished MK2 activation and also strongly diminished the phosphorylation of p70S6K, S6, and 4EBP1 after stimulation with LPS, UV, anisomycin, or L.m. (Fig. 3A and data not shown). Erk activation was only observed after LPS or L.m. treatment and not modified in $p38a^{\Delta M}$ BMDM (Fig. 3A). Moreover, activation of p70S6K and 4EBP1 was blocked in spleens from $p38a^{\Delta M}$ but not $p38a^{f1/f1}$ mice that were challenged with LPS (Fig. 3B) demonstrating that p38a activates mTOR signaling *in vitro* and *in vivo*. Next we analyzed whether MK2 may mediate the effect of p38a on mTOR signaling. Therefore, we used macrophages expressing a catalytic-dead mutant of MK2 (K79R) or its wild-type control. Strikingly, activation of mTOR after stimulation with LPS or anisomycin was severely compromised in the K79R mutant compared to wild-type MK2 (Fig. 3C). Moreover, the K79R macrophages showed absent IL-10 expression after LPS or SAC stimulation, while IL-12p40 was strongly increased (Fig. 3, D and E). These data indicate that the kinase activity of MK2 transmits the p38a signal to stimulate mTOR signaling and to regulate the production of IL-12 and IL-10.

p38 signals to mTOR via TSC1/TSC2 to regulate IL-12 and IL-10

TSC1 and TSC2 act as dimer to inhibit activation of mTORC1 (29). To investigate whether p38-MK2 may mediate mTORC1 activation via TSC1/TSC2, we made use of murine embryonic fibroblasts (MEF) deficient in either TSC1 or TSC2. Treatment with anisomycin or UV stimulated p38 and mTOR signaling in $Tsc1^{+/+}$ as well as in $Tsc2^{+/+}$ cells and inhibition of p38 by either SB203580 or BIRB796 strongly reduced mTOR activation as shown by diminished phosphorylation of p70S6K or S6 (Fig. 4, A–D). Strikingly, inhibition of p38 did not block mTOR signaling in cells deficient of TSC1 or TSC2 (Fig. 4, A–D). Rapamycin, which acts downstream of TSC1/TSC2, still inhibited mTOR in these cells (Fig. 4, A–D). These results suggest that the TSC1/TSC2 complex is a critical signaling nodule that senses p38 activity to regulate mTOR activation. Interestingly, serum, a well known mTOR activator, did not stimulate p38 activation and inhibition of p38 did not influence mTOR activation induced by serum (Fig. 4E). Previously, it has been shown that MK2 phosphorylates serine 1210 (Ser¹²¹⁰) in TSC2 (37). However, it remained to be verified whether p38 activation regulates TSC2 activity via Ser¹²¹⁰ phosphorylation by MK2. To explore this possibility, we transfected $Tsc2^{-/-}$ cells with plasmids encoding either wild-type TSC2 or a S1210A TSC2 mutant and observed that wild-type but not mutant TSC2 restored the ability of anisomycin to activate p70S6K (Fig. 4F). Moreover, BIRB796 inhibited p70S6K activation upon p38 stimulation in cells transfected with wild-type but not mutant TSC2 (Fig. 4F). Overall, these results demonstrate that activation of p38 inhibits the TSC1/ TSC2 complex allowing mTOR activation potentially via MK2-dependent phosphorylation of TSC2 at Ser¹²¹⁰.

p38α and PI3K independently regulate mTOR

Classical activation of mTOR signaling by growth factors or TLR ligands is dependent on PI3K (17, 38). Our results so far established that p38a activates mTOR via MK2 involving a Ser¹²¹⁰-dependent regulation of TSC2 to modulate IL-12/IL-10 signaling. To further delineate the relative requirements of PI3K and p38a for stimulating mTOR, we inhibited p38 in the presence or absence of wortmannin, a specific covalent PI3K inhibitor. Treatment with either wortmannin or BIRB796 considerably diminished phosphorylation of S6 in human LPS- or anisomycin-activated monocytes (Fig. 5, A and B). The combination of both inhibitors led to a near complete abolishment of S6 phosphorylation (Fig. 5, A and B). Likewise, mTOR was blocked more efficiently in BMDM from $p38a^{\Delta M}$ than from $p38a^{fl/fl}$ mice after wortmannin treatment (Fig. 5, C and D). Immunofluorescence analysis confirmed that wortmannin inhibited the phosphorylation of S6 more completely in $p38a^{\Delta M}$ than in p38a^{fl/fl} BMDM (Fig. 5E) suggesting that PI3K and p38 cooperatively stimulate mTOR signaling. Next, we analyzed MEFs deficient in p85a/p85β, the regulatory subunits of PI3K, to assess the importance of p38 versus PI3K for maximum activation of mTOR (Fig. 5F). Phosphorylation of p70S6K and 4E-BP1 induced by LPS, anisomycin or UV exposure was severely reduced but still detectable in p85^{-/-} MEFs compared to their wild-type counterparts (Fig. 5F). However, concurrent inhibition of p38 by BIRB796 in p85^{-/-} MEFs further inhibited activation of mTOR (Fig. 5F). Together, these results strongly suggest that full activation of mTOR in myeloid immune cells is dependent on p38 and PI3K.

p38 and PI3K coordinately control expression of IL-12 and IL-10

Finally, we wanted to elucidate the relative contribution of p38a and PI3K for the production of IL-12 and IL-10. BMDM showed maximum production of IL-12p70 or IL-12p40 when PI3K was inhibited regardless of whether p38a was present or not (Fig. 6A and data not shown). This indicates that PI3K activation is dominant after LPS stimulation

and that p38a modulates IL-12p40 within this context. In contrast, IL-10 production seems to more thoroughly depend on p38a as deletion of p38a already diminished IL-10 production of BMDM to a level that could not be further inhibited by PI3K inhibition (Fig. 6A). Similar results were obtained in human monocytes (Fig. 6B). In these cells, p38 was also dominant for IL-10 production, whereas PI3K and p38 both regulated Il-12p40 (Fig. 6B). In conclusion, these results suggest that PI3K and p38a coordinately modulate mTOR signaling to regulate the expression of IL-12 and IL-10 in myeloid immune cells.

Discussion

Monocytes, macrophages and DCs are emerging therapeutic targets in cardiovascular, malignant, and autoimmune disorders. Addressing gaps in knowledge about the function of these cells in response to environmental stimuli is required to understand the *in vivo* responses to therapies that target these cells. Stringent control of MAPK signaling is critical for balancing pro- versus anti-inflammatory signaling to enable efficient pathogen killing but also to limit detrimental tissue pathology. In that regard p38 is one of the most studied drug targets for anti-inflammatory therapy. This kinase directly or indirectly regulates many transcription factors and therefore participates in the gene induction of cytokines and other inflammatory molecules. p38 is also important in the post-transcriptional regulation of gene expression during inflammation (39). Animal studies have shown that p38 inhibitors are efficacious in several disease models, including inflammation, arthritis and other joint diseases, septic shock and myocardial injury (11). However, translation into the clinic has been difficult either due to lack of efficiency or the appearance of adverse effects including inflammation such as skin rashes.

p38a is activated in diverse cell types by a wide array of stress stimuli including genotoxic agents, PAMPs, proinflammatory cytokines, heat or osmotic shock, oxygen partial pressure or chemical insults (arsenite and anisomycin) (40). The simple view of an entirely proinflammatory kinase promoting the expression of TNF-a and IL-1 β shifted to a more complex role in recent years demonstrating that p38a controls IL-12 and IL-10 expression (14–16). However, the downstream pathway that regulates these immunomodulatory cytokines remained unknown.

Here we have demonstrated that p38a utilizes the TSC2/mTOR signaling pathway to control the balance of IL-12 and IL-10. Our data suggest that TLR ligands or stress stimuli lead to an activation of p38a that in turn activates its downstream kinase MK2. The kinase activity of MK2 most likely phosphorylates Serine 1210 of TSC2 leading to inactivation of the TSC1/TSC2 complex and in turn activation of the mTOR pathway (Fig. 6C). Activation of mTOR then promotes IL-10 production, while reducing IL-12 expression. Our work indicates that p38a-mediated mTOR activation occurs in parallel to the well-known PI3K pathway that activates mTOR in response to TLR signals (17, 19). Hence, both pathways concurrently control mTOR activation to precisely allow the expression of pro- and antiinflammatory cytokines in response to environmental stress. We are unaware of a report demonstrating that two stimuli additively regulate the activation of mTOR via the TSC complex. We suggest that p38a-mediated mTOR activation in addition to the PI3K pathway represents a tuning mechanism to regulate immunomodulatory cytokines to adapt the immune response to the environmental milieu. This is supported by the observation that hyperactivation of p38a by anisomycin can modulate IL-12 and IL-10 expression on top of a TLR signal (Fig. 2D). The p38/MK2 axis is required after excessive tissue damage to induce tissue repair (41, 42). In such situations p38a may promote mTOR activation in resident and recruited macrophages to reduce IL-12 and augment IL-10 production that limits the generation of a proinflammatory CD4+ Th1 response that would further exaggerate tissue damage (43).

A link from p38ß to mTOR has been described in Drosophila that occurs via a TSC2independent mechanism (44). In line, p38ß was recently shown to phosphorylate the essential mTORC1 binding protein Raptor and to participate in arsenite-induced mTOR activation in fibroblasts (45). In contrast, in the same cell p38ß can also inhibit mTOR upon energy starvation via phosphorylation and inactivation of Ras homologue enriched in brain (Rheb), a key component of the mTORC1 pathway (46). We now show that p38a via MK2 promotes mTOR activation dependent on the TSC1/TSC2 complex in myeloid immune cells. Indeed, MK2 was shown to phosphorylate TSC2 at Ser^{1210} in fibroblasts (37). In addition, MK2 was also described to phosphorylate Akt at Ser⁴⁷³ in neutrophils (47), in line with the inhibition of Akt Ser⁴⁷³ by the p38 inhibitors in our cells (Fig. 2A and Fig. 6). We have previously shown that mTOR regulates NF-*k*B and STAT3 signaling (24). Interestingly, also p38a and MK2 are required for STAT3 activation and IL-10 production (14, 36). These effects are likely to be indirectly mediated by p38 and mTOR and the precise downstream pathways how mTOR regulates IL-12 vs. IL-10 needs further investigation. In this study we focused on delineating the mechanism of p38-depdendent mTOR regulation in myeloid immune cells.

Rapamycin is currently evaluated as vaccine adjuvant, especially because of its ability to enhance memory CD8+ T cell responses. In addition, previous work also established that rapamycin exerts immunostimulatory effects via the innate immune system that may contribute to the adjuvant properties of rapamycin (23–25, 48, 49). However, its immunosuppressive activity will likely prevent the inclusion of rapamycin in widely-distributed vaccines. Our data suggest that inhibition of p38 might be similarly effective as adjuvant strategy where strong Th1 responses are desired and, moreover, it might avoid the potent immunosuppressive effects on the T cell compartment as p38 is regarded dispensable for T cell function (50). Indeed, in a mouse model of *Leishmania major* infection, vaccination with SB203580 was protective by inducing efficient Th1 immunity (16).

In summary, we have identified and characterized a pathway from p38a to mTOR via MK2 and TSC1/TSC2 in myeloid immune cells that tunes the immune response according to environmental input signals.

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FIGURE 1.

p38a modulates IL-12 and IL-10 in mice and men. (A) p38a^{fl/fl} and p38a^{ΔM} bone marrowderived macrophages (BMDM) were stimulated with medium (-), LPS, SAC or Listeria monocytogenes (L.m.) for 24 h. IL-12p40, IL-12p70, and IL-10 in the supernatants were determined by Luminex. Data are shown as means \pm SEM for five mice. (B) Serum levels of IL-12p40, IL-12p70, and IL-10 were determined in p38a^{fl/fl} and p38a^{ΔM} mice injected with LPS for 4 h (means ± SEM for 3 mice). (C-F) Human monocytes were treated with medium (-), BIRB796 (BIRB; 100 or 200 nM), SB203580 (SB; 200 nM or 2 µM) or rapamycin (Rap; 100 nM) and then stimulated with (C) LPS, (D) SAC, (E) L.m. or (F) LPS and IFN- γ for 22h. Cytokines in cell-free supernatants were measured by Luminex (means \pm SEM of at least 3 donors). (G-H) Human monocytes were treated as indicated, washed and added to allogeneic T cells for one week. (G) IFN- γ of cell-free supernatants was determined by Luminex. Data represent means \pm SEM for three independent experiments. (H) Primed T cells were activated for 5 h with PMA/Ionomycin in the presence of Brefeldin A. Intracellular cytokine staining for IL-4 and IFN- γ in CD-4 T cells is illustrated. One representative experiment out of three is shown. *, P < 0.05 compared with the respective controls.



FIGURE 2.

p38 activation stimulates mTOR signaling. (A–C) Human monocytes were incubated with medium (–), BIRB796 (200 nM), SB203580 (2 μ M), SD169 (200 nM) or rapamycin (Rap; 100 nM) and stimulated with (A) LPS (100 ng/ml), (B) anisomycin (50 ng/ml) or (C) UV (30 sec) for the indicated times. Cell lysates were analyzed by immunoblotting. Representatives of three independent experiments are shown. (**D**,**E**) Murine macrophages were stimulated with (D) LPS or (E) SAC in the presence or absence of anisomycin (50 ng/ml) for 1 h. IL-12p40 and IL-10 mRNA levels were measured by RT-PCR. Levels were normalized to ubiquitin and are shown relative to the (D) LPS- or (E) SAC-stimulated samples, which were set to 1 (means ± SEM; n=3). *, P < 0.05. (F) Monocytes were incubated with LPS (10 ng/ml) or anisomycin (10 ng/ml) for 30 min as indicated. Cell lysates were analyzed by immunoblotting.



FIGURE 3.

p38a activates mTOR via MK2. (**A**) p38a^{*fl/fl*} and p38a^{ΔM} BMDM were stimulated with medium (–), LPS (100 ng/ml), anisomycin (100 ng/ml), UV (1 min) or L.m. (10⁷ cells). Cell lysates were analyzed by immunoblotting. Representatives of three independent experiments are shown. (**B**) p38a^{*fl/fl*} and p38a^{ΔM} mice were injected with LPS for 4 h. Isolated and homogenized spleens were analyzed by immunoblotting. Results for two mice per genotype are shown. Similar results were obtained in an independent experiment (data not shown). (**C**) MK2^{-/-} macrophages reconstituted with either MK2^{K79R} or wild-type (WT) MK2 were stimulated with medium, LPS (100 ng/ml) or anisomycin (100 ng/ml) for 1 h and whole-cell lysates were analyzed by immunoblotting. Representatives of three independent experiments are shown. (**D**,**E**) MK2 WT and MK2^{K79R} macrophages were stimulated with medium (–), LPS (100 ng/ml) or SAC (75 µg/ml) for 2 h. mRNA levels of IL-12p40 and IL-10 were measured by RT-PCR. Levels were normalized to ubiquitin and are shown relative to the (**D**) LPS- or (**E**) SAC-treated MK2 WT samples, which were set to 1 (means ± SEM; n=3). *, P < 0.05.



FIGURE 4.

p38 signals to mTOR via TSC1/TSC2. (**A**, **B**) *Tsc1*^{+/+} and *Tsc1*^{-/-} MEFs or (**C**-**E**) *Tsc2*^{+/+} and *Tsc2*^{-/-} MEFs were starved overnight and treated with medium (–), BIRB796 (200 nM), SB203580 (2 μ M) or rapamycin (Rap; 100 nM) for 90 min. Afterwards, MEFs were stimulated with (A, C) anisomycin (100 ng/ml), (B, D) UV (1 min) or (E) 10% serum for 60 min. Whole-cell lysates were analyzed by immunoblotting. (**F**) *Tsc2*-/- MEFs were transfected with empty vector (e.v.), HA-tagged wild-type TSC2 (WT), or HA-tagged TSC2 S1210A and then treated with medium, BIRB796, and anisomycin as indicated. Immunoblotting was performed with the indicated antibodies. Long exposure revealed the phosophorylation of the p85S6K (p85) isoform in addition the the p70S6K (p70) isoform. Nonspecific bands (NB). Densitometric analysis was performed on the intensity of p-p70S6K. One representive of three different experiments is shown.



FIGURE 5.

p38a and PI3K independently regulate mTOR. (**A**, **B**) Human monocytes were incubated with medium (–), BIRB796 (200 nM) and/or wortmannin (WM; 100 nM) for 90 min and stimulated with (A) LPS (100 ng/ml) or (B) anisomycin (100 ng/ml). Cell lysates were analyzed by immunoblotting. (**C**–**E**) p38a^{fl/fl} and p38a^{Δ M} BMDM were treated with medium (–), BIRB796 (200 nM), SB203580 (2 μ M) and/or wortmannin (WM; 100 nM) as indicated and stimulated with (C, E) LPS (100 ng/ml) or (D) anisomycin (100 ng/ml) for 1 h. (C, D) Cell lysates were analyzed by immunoblotting. (**E**) Phosphorylation of S6 and MK-2 was analyzed by immunofluorescence. One representative experiment of three is shown. (**F**) p85^{+/+} and p85^{-/-} MEFs were treated with medium (–) or BIRB796 (200 nM) followed by stimulation with LPS (100 ng/ml), anisomycin (100 ng/ml) or UV (10 sec) for 60 min. Cell lysates were analyzed by immunoblotting.



FIGURE 6.

p38a and PI3K coordinately regulate the IL-12/IL-10 balance. (A) p38a^{fl/fl} and p38a^{ΔM} BMDM were treated with wortmannin (WM; 100nM) and stimulated with medium (–), SAC (75 µg/ml), or LPS (100 ng/ml) for 24 h. Cytokine levels in cell-free supernatants were determined by Luminex (means ± SEM; n=5). *, P < 0.05; n.s., non significant. (B) Human monocytes were treated with wortmannin (WM; 100 nM) or BIRB796 (200 nM) as indicated and stimulated with LPS (100 ng/ml) for 24 h. IL-12p40 and IL-10 were determined in the supernatants by Luminex (means ± SEM; n=3). *, P < 0.05 (C) Model of mTOR-regulated production of IL-12/IL-10 via PI3K and p38 on the level of TSC1/TSC2.