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## IL-6-mediated induction of MMP-9 is modulated by JAK-dependent IL-10 expression in macrophages<sup>1</sup>

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### Abstract

The mechanisms by which IL-6 contributes to the pathogenesis of chronic inflammatory diseases and cancer are not fully understood. We previously reported that cyclooxygenase-2 (Cox-2)-dependent PGE<sub>2</sub> synthesis regulates macrophage matrix metalloproteinase (MMP)-9 expression, an endopeptidase that participates in diverse pathologic processes. In these studies, we determined whether IL-6 regulates the Cox-2→PGE<sub>2</sub>→MMP-9 pathway in murine macrophages. IL-6 co-induced Cox-2 and microsomal prostaglandin E synthase-1 (mPGES-1), and inhibited the expression of 15-hydroxyprostaglandin dehydrogenase (15-PGDH), leading to increased levels of PGE<sub>2</sub>. In addition, IL-6 induced MMP-9 expression, suggesting that the observed proteinase expression was regulated by the synthesis of PGE<sub>2</sub>. However, inhibition of PGE<sub>2</sub> synthesis partially suppressed IL-6-mediated induction of MMP-9. In the canonical model of IL-6-induced signaling, JAK activation triggers STAT and MAPK<sup>erk1/2</sup>-signaling pathways. Therefore, the ability of structural diverse JAK inhibitors to block IL-6-induced MMP-9 expression was examined. Inhibition of JAK blocked IL-6 induced phosphorylation of STAT3, but failed to block the phosphorylation of MAPK<sup>erk1/2</sup>, and unexpectedly enhanced MMP-9 expression. In contrast, MEK-1 inhibition blocked IL-6 induced phosphorylation of MAPK<sup>erk1/2</sup> and MMP-9 expression without affecting the phosphorylation of STAT3. Thus, IL-6-induced MMP-9 expression is dependent on the activation of MAPK<sup>erk1/2</sup> and restrained by a JAK-dependent gene product. Utilizing pharmacologic and genetic approaches, JAK-dependent induction of IL-10 was identified as a potent feedback mechanism controlling IL-6 induced MMP-9 expression. Together, these data reveal that IL-6 induces MMP-9 expression in macrophages via Cox-2-dependent and -independent mechanisms, and identifies a potential mechanism linking IL-6 to the pathogenesis of chronic inflammatory diseases and cancer.

### Introduction

IL-6, a pleiotropic cytokine expressed by a variety of immune and non-immune cells, plays an important role in the recruitment and survival of neutrophils and macrophages, regulation

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of CD4 T cell effector functions, angiogenesis, bone and cartilage metabolism, lipid metabolism, and the expression of acute phase proteins (1,2). Circulating levels of IL-6 are elevated in patients with cancer (3) and several chronic inflammatory diseases including rheumatoid arthritis (4) and cardiovascular disease (5). In addition, adipose tissue is a major source of circulating IL-6, and levels are elevated in obese patients (6). Owing to its multiple roles in the pathogenesis of inflammatory diseases and cancer (1,2), IL-6 has emerged as a major target for therapeutic intervention (7,8).

IL-6-induced biological responses are mediated by the membrane bound IL-6 receptor (IL-6R $\alpha$ ; CD126) (7,9,10). The IL-6/IL-6R $\alpha$  complex engages transmembrane gp130 (IL-6R $\beta$ ; CD130), and the ternary complex dimerizes triggering the binding and phosphorylation of JAK, which then phosphorylates gp130 leading to the activation of STAT and MAPK signaling pathways. Despite the broad biologic activities of IL-6, surprisingly few cell types (e.g., monocyte/macrophages and hepatocytes) express membrane bound IL-6R $\alpha$ . In contrast, virtually all cells types express gp130, which can bind soluble IL-6/IL-6R $\alpha$  complexes (i.e. trans-signaling), thereby triggering STAT and MAPK signaling pathways.

Notwithstanding the significant progress in unraveling the many effects of IL-6 on immune and non-immune cells, the mechanisms by which IL-6 contributes to the pathogenesis of chronic inflammatory diseases and cancer is not fully understood. In this regard, evidence derived from mouse models indicates that MMP-9 (type IV collagenase; gelatinase B), a family member of Zn<sup>+2</sup>-dependent neutral endopeptidases, participates in the pathogenesis of arthritis (11), airway disease (12), cancer (13,14) and cardiovascular diseases (15-17). MMP-9 expression is low or absent in most normal tissues, and markedly elevated during inflammation, wound healing, and neoplasia (18-20). We and others have reported that macrophage MMP-9 expression is stimulated by PGE<sub>2</sub> (21-29). Increased Cox-2-dependent synthesis of PGH<sub>2</sub> and subsequent isomerization to PGE<sub>2</sub> by mPGES-1 (29-34), in combination with reduced catabolism by NAD<sup>+</sup>-dependent 15-PGDH (35,36), are largely responsible for elevated levels of PGE<sub>2</sub> associated with inflammation. Consequently, we determined whether IL-6 regulates the Cox-2→mPGES-1→PGE<sub>2</sub>→MMP-9 pathway in macrophages.

Results demonstrate that IL-6-induced MMP-9 expression in macrophages via Cox-2-dependent and independent pathways. Because IL-6 can activate both JAK/STAT and MAPK pathways, we explored their roles in regulating MMP-9 expression. Inhibition of MAPK<sup>erk1/2</sup> blocked IL-6-mediated induction of MMP-9. In contrast, inhibition of JAK led to a paradoxical increase in MMP-9 expression, which proved to be a consequence of diminished IL-10 levels. To the best of our knowledge, this is the first demonstration that IL-6 induces macrophage expression of MMP-9, which has been directly associated with the pathogenesis of chronic inflammatory diseases and cancer (18-20). In addition, these data suggest that JAK inhibitors have the potential to increase MMP-9 expression in macrophages via the inhibition of IL-10 expression.

## Materials and Methods

### Macrophages

Thioglycollate-elicited peritoneal macrophages were obtained from Swiss Webster mice by the method of Edelson and Cohn (37) as described previously (38). Mice were injected IP (3 ml/mouse) with 3% Brewer Thioglycollate Medium (DIFCO). Four days later, cells were harvested by lavage with cold Dulbecco's PBS. Peritoneal cells were recovered by centrifugation and resuspended in DMEM supplemented with 10% FBS, penicillin (100 U/ml), streptomycin (100  $\mu$ g/ml) and 4 mM glutamine, and plated into 6-well ( $2 \times 10^6$  cells/

well) or 12-well plates ( $6 \times 10^5$  cells/well). Cells were allowed to adhere for 4 h and then washed free of nonadherent cells. The murine macrophage cell line RAW264.7 (39) was obtained from American Type Culture Collection, and maintained as adherent cultures in DMEM-10% FBS. RAW264.7 cells were mechanically harvested and plated into 12-well plates ( $6 \times 10^5$  cells/well). For serum free conditions, macrophages were grown in DMEM supplemented with antibiotics, glutamine and 0.1% BSA (Sigma Aldrich) containing low levels of endotoxin ( $< 0.1$  ng/mg; LE-BSA). Cellgro® DMEM was obtained from Corning and heat inactivated FBS were obtained from Atlanta Biologicals. Antibiotics and glutamine were obtained from Gibco/Life Technologies. Macrophages were incubated with recombinant human IL-6 (Santa Cruz Biotechnology), recombinant mouse IL-10 (BioLegend), rat monoclonal anti-mouse IL-10 or non-immune rat IgG (R&D Systems) celecoxib (LC Laboratories), JAK Inhibitor 1 (Calbiochem/EMD Millipore), Tofacitinib (Selleckchem), or U0126 (Cayman Chemical). All animal studies described in this report have been reviewed and approved by the Weill Cornell Medical College Institutional Animal Care and Use Committee.

### Preparation of cell lysates

In experiments designed to monitor levels of phosphorylated and total STAT3 and MAPK<sup>erk1/2</sup>, cells were harvested in RIPA buffer (Sigma Aldrich) containing 1X protease inhibitor cocktail (Sigma Aldrich) and 1X Halt™ phosphatase inhibitor cocktail (Pierce), briefly sonicated and centrifuged ( $4,225 \times g$ ) for 10 min at 4°C. The supernatants were recovered, normalized for protein and mixed with SDS sample buffer with DTT and boiled for 5 min. Equal amounts of cell lysates were applied to gels based on protein content.

### Western blots

Cell lysates were electrophoresed in 8-16% polyacrylamide gels and proteins were transferred to a polyvinylidene fluoride (PVDF) membrane. The membrane was blocked in 5% BSA in TBST, washed in TBST and incubated 18 h in blocking buffer containing rabbit anti-mouse phospho-STAT3 (Tyr705) (1:2000; Cell Signaling Technology) or rabbit anti-human phospho-erk1/2 IgG (1:3000; Cell Signaling Technology). Membranes were washed 2X in TBST and incubated 1 h in blocking buffer containing affinity purified goat anti-rabbit IgG conjugated to HRP (1:3000; BioRad Laboratories). The membranes were washed 3X in TBST and bound HRP was visualized utilizing chemiluminescence (Pierce/Thermo Scientific). Membranes were then stripped in 0.1% SDS, 1.0% Tween 20 and 200 mM glycine, pH 2.2 10 min at RT, washed, blocked for 1 h in blocking buffer, and incubated with rabbit anti-mouse STAT3 (1:2000; Cell Signaling Technology) or mouse anti-erk1/2 (KLH-synthetic peptide) (1:3000; Cell Signaling Technology).

Macrophage conditioned media were electrophoresed in gradient gels and proteins were transferred to a PVDF membrane. The membrane was placed in 5% dry defatted milk in TBST for 1 h, washed in TBST and incubated 18 h in 3% blocking buffer containing rabbit anti-mouse MMP-9 IgG (1:2500; Abcam). Membranes were washed 2X in TBST and incubated 1 h in 3% blocking buffer containing goat anti-rabbit IgG conjugated to HRP (1:3000; BioRad Laboratories).

### IL-10 knockdowns

RAW264.7 macrophages were transfected with control siRNA (i.e. nonspecific, NS) or IL-10 siRNA (Santa Cruz Biotechnology) utilizing HiPerFect transfection reagent according to the manufacturer's protocol (Qiagen). Cells were aliquoted into a 12-well ( $3 \times 10^5$  cells/well) plate containing siRNA transfection complexes and incubated 30 h. PCR and immunoassay was utilized to determine the extent of IL-10 knockdown.

## Real-Time PCR analysis

RNA was prepared using Trizol reagent kits (Invitrogen). RNA (2  $\mu$ g) was reverse transcribed using Moloney murine leukemia virus reverse transcriptase (Invitrogen) and random hexamers. The resulting cDNA diluted 10-fold was used for amplification. For real-time PCR analysis, the reaction volume was 5  $\mu$ l and contained 1-2  $\mu$ l of the diluted cDNA, 1X Gene Expression Assay (Life Technologies), 1X PerfeCTa FastMix II, Rox (Quanta) using the StepOne instrument (Applied Biosystems). The following Applied Biosystems Gene Expression Assays were utilized: actin (mm00607939\_s1), Cox-2 (mm00478374\_m1), MMP-9 (mm00442991\_m1), mPGES-1 (mm00452105\_m1), IL-6 (mm00446190\_m1), 15-PGDH (mm00515121\_m1) and IL-10 (mm00439614\_m1). The comparative  $C_T$  ( $\Delta\Delta C_T$ ) method was used to determine the relative target quantity in samples. With the comparative  $C_T$  method, the StepOne software measures amplification of the target and endogenous control in the samples and in a reference sample. Measurements are normalized using the endogenous control and expressed as relative quantity (RQ).

## Determination of PGE<sub>2</sub> and IL-10 levels

The concentrations of PGE<sub>2</sub> and IL-10 in macrophage conditioned media were determined utilizing the PGE<sub>2</sub> EIA kit (monoclonal) (Cayman Chemical) and IL-10 Quantikine Immunoassay (R&D).

## Statistics

Relative quantities (RQ) of target gene mRNA and PGE<sub>2</sub> or IL-10 levels are expressed as the mean  $\pm$  SD of 3 independent samples from a single representative experiment. Each experiment was repeated 2-3 times. Where indicated, mean levels of mRNA, PGE<sub>2</sub> or IL-10 were compared utilizing t-test (e.g., comparing means of two groups) or single factor analysis of variance (e.g., comparing means of greater than 2 groups). Following the determination of significant differences between the means of a given experiment, subsequent pair-wise comparisons were performed utilizing Newman-Keuls multiple range testing (40).

## Results

### IL-6 stimulates Cox-2 and mPGES-1 expression, and inhibits 15-PGDH expression in macrophages

We determined whether IL-6 co-induces Cox-2 and mPGES-1 expression in macrophages, leading to increased levels of PGE<sub>2</sub>. Incubation with LPS, a potent inducer of Cox-2 and mPGES-1 in macrophages, served as a positive control (30,31,41). Levels of Cox-2 and mPGES-1 mRNA were determined utilizing quantitative PCR. Incubation of RAW264.7 macrophages with IL-6 induced Cox-2 and mPGES-1 expression in a dose- and time-dependent manner (Figures 1A and 1B). Since reduced catabolism of prostaglandins contributes to increased levels of PGE<sub>2</sub> in inflammatory foci, we also examined the effect of IL-6 on the expression of 15-PGDH, a key enzyme responsible for the catabolism of prostaglandins (35,36). In contrast to Cox-2 and mPGES-1 expression, IL-6 rapidly and potently inhibited 15-PGDH expression in macrophages (Figures 1A and 1B). Consistent with the IL-6-induced reciprocal changes in expression of genes directing the synthesis and degradation of PGE<sub>2</sub>, levels of PGE<sub>2</sub> in media recovered from RAW264.7 cells increased in a dose- and time-dependent manner (Figures 1A and 1B).

To corroborate and extend these findings to primary macrophages, we examined the effect of IL-6 on the expression of Cox-2, mPGES-1 and 15-PGDH by macrophages recovered from thioglycollate-induced peritonitis (Figure 2). Following 18 h incubation with IL-6, levels of Cox-2 and mPGES-1 mRNA in elicited macrophages were markedly elevated, and

levels of 15-PGDH mRNA were reduced. Likewise, levels of PGE<sub>2</sub> in media recovered from elicited macrophages treated with IL-6 were increased dramatically.

In these studies, murine macrophages were incubated with recombinant human IL-6, which binds and activates the murine IL-6 receptor (42). To confirm that the observed alterations in Cox-2, mPGES-1 and 15-PGDH were due to IL-6 and not a potential contaminant, we pre-incubated human IL-6 with non-immune IgG or neutralizing anti-human IL-6 IgG prior to incubation with macrophages. Anti-IL-6 IgG blocked IL-6-induced alterations in gene expression in murine macrophages (data not shown). Taken together, these data demonstrate that IL-6 stimulates macrophage expression of Cox-2 and mPGES-1, and inhibits the expression of 15-PGDH, leading to increased levels of PGE<sub>2</sub>.

### **Cox-2-dependent PGE<sub>2</sub> synthesis plays a role in IL-6-mediated induction of MMP-9**

Macrophage MMP-9 expression is regulated by Cox-2-dependent synthesis of PGE<sub>2</sub> (26-29). Thus we examined whether the observed changes in expression of genes directing the synthesis and degradation of PGE<sub>2</sub> in response to IL-6 (Figures 1 and 2), were associated with changes in MMP-9 expression. Treatment of RAW264.7 macrophages with IL-6 resulted in a dose- and time-dependent increase in the levels of MMP-9 mRNA (Figures 3A and 3B). The dose-dependent increases in MMP-9 mRNA were mirrored by an accumulation of MMP-9 in cellular conditioned media (Figure 3A). Treatment with IL-6 similarly stimulated MMP-9 expression in elicited peritoneal macrophages (Figure 3C;  $p < 0.01$ ). IL-6 induced macrophage MMP-9 expression identifies a potential mechanism linking elevated IL-6 levels to the pathogenesis of chronic inflammatory diseases and cancer (18-20).

In contrast to earlier studies where MMP-1 and MMP-3 induced Cox-2 expression and increased PGE<sub>2</sub> synthesis preceded elevated MMP-9 expression (28), IL-6 induced MMP-9 expression parallels the expression of both Cox-2 and mPGES-1 (Figures 1 and 3). These findings raise the possibility that the increased synthesis of PGE<sub>2</sub> may not be solely responsible for the induction of MMP-9. However, since macrophage MMP-9 expression is stimulated by PGE<sub>2</sub> (26-29), IL-6-induced reciprocal changes in the expression of genes directing the synthesis and degradation of PGE<sub>2</sub> may contribute to MMP-9 expression. Therefore, we examined the effect of the selective Cox-2 inhibitor celecoxib on IL-6 induced PGE<sub>2</sub> levels and MMP-9 expression in macrophages. The levels of PGE<sub>2</sub> in conditioned media recovered from macrophages incubated with IL-6 (18 h) were increased >20-fold, as compared to control macrophages (Figure 4A). Pre-treatment with celecoxib completely inhibited IL-6-induced PGE<sub>2</sub> synthesis, but only led to a modest reduction in IL-6-mediated induction of MMP-9 mRNA (Figure 4B;  $p < 0.05$ ) and protein (Figure 4C). These data indicate that IL-6 stimulates MMP-9 expression by both Cox-2-dependent and -independent mechanisms.

### **IL-6-induced MMP-9 expression is MAPK<sup>erk1/2</sup>-dependent.**

In the canonical model of IL-6-induced signaling, binding of IL-6/IL-6R $\alpha$  to gp130 triggers the activation of the JAK and subsequent phosphorylation of several tyrosines within the cytoplasmic tail of gp130 creating binding sites for STAT and SHP2. The latter is believed to be critical for the activation of MAPK<sup>erk1/2</sup> (7,9,10). Thus, inhibition of JAK activity ought to block both STAT and MAPK<sup>erk1/2</sup>-signaling pathways. Therefore, we examined the effect of two selective inhibitors of JAK on IL-6-induced MMP-9 expression (Figure 5). Unexpectedly, pre-treatment of macrophages with JAK inhibitor 1 (1 – 100 nM) augmented IL-6-mediated induction of MMP-9 mRNA and MMP-9 protein in cellular conditioned media (Figure 5A). Likewise, pre-treatment with Tofacitinib (0.1 – 1  $\mu$ M), a structurally distinct JAK inhibitor, augmented IL-6-mediated induction of MMP-9 (Figure 5B). In

contrast, pre-treatment with the MEK1 inhibitor U0126 (1 - 10  $\mu$ M) led to a dose-dependent inhibition of IL-6 induced MMP-9 expression (Figure 5C). The failure of JAK inhibitors to block IL-6-induced MMP-9 expression was unexpected. Thus, we monitored their ability to inhibit the phosphorylation of STAT3 and MAPK<sup>erk1/2</sup>. As anticipated, pre-treatment with JAK Inhibitor 1 (100 nM) or Tofacitinib (1  $\mu$ M) completely blocked IL-6-induced phosphorylation of STAT3; whereas, IL-6-induced phosphorylation of MAPK<sup>erk1/2</sup> was unaffected (Figure 6C). Pre-treatment with U0126 blocked IL-6-induced phosphorylation of MAPK<sup>erk1/2</sup> (Figure 6B), but had no effect on IL-6-induced phosphorylation of STAT3 (Figure 6A). Although these data are unexpected based on the established model of IL-6-induced signaling, several investigators have reported that diverse JAK inhibitors block STAT activation without inhibiting MAPK<sup>erk1/2</sup> activation (43-47); however, the mechanism(s) underlying JAK-independent activation of MAPK<sup>erk1/2</sup> remains undefined at this time.

Although IL-6 is one of the most highly induced NF- $\kappa$ B-dependent cytokines (48), NF- $\kappa$ B is not generally reported to mediate IL-6-dependent signaling and gene expression (7,9,10). However, it has been reported that in addition to canonical JAK/STAT signaling, IL-6 induces the accumulation of unphosphorylated STAT3, which forms a complex with unphosphorylated NF- $\kappa$ B that can translocate into the nucleus (49). Therefore, we examined the effect of two cell permeable NF- $\kappa$ B activation inhibitors on IL-6-induced MMP-9 expression. Pre-incubation of RAW264.7 macrophages with several concentrations of NF- $\kappa$ B Activation Inhibitor I (CAS 545380-34-5) or II (CAS 749886-87-1) failed to block or attenuate IL-6-induced MMP-9 expression (data not shown). Taken together, our data demonstrate that IL-6-induced MMP-9 expression in macrophages is dependent on the activation of MAPK<sup>erk1/2</sup> and paradoxically enhanced by inhibition of JAK.

### **Tofacitinib inhibition of IL-6-induced IL-10 enhances MMP-9 expression**

The observation that IL-6-induced MMP-9 expression in macrophages was enhanced by JAK inhibition suggested that a JAK-dependent gene product serves to limit IL-6-induced MMP-9 in macrophages. In this regard, the pro-inflammatory properties of LPS are limited, in part, by the JAK-dependent induction of IL-10, a potent anti-inflammatory cytokine (46), which is also reported to inhibit MMP-9 expression (50-53). Therefore, we examined whether IL-6 stimulated IL-10 expression in macrophages. As seen in Figure 7A and B, levels of IL-10 mRNA in macrophages and IL-10 in conditioned media were increased, in a dose-dependent manner. Likewise, IL-6 induced IL-10 expression in a time-dependent manner (Figure 7C and D). When compared to control cells, levels of IL-10 mRNA are increased > 60-fold in as little as 2 h, and >1000 fold at 18 h. Levels of IL-10 in the conditioned media recovered from cells incubated with IL-6 were first elevated at 4 h (124  $\pm$  90 pg/ml; mean  $\pm$  SD, n=3) and increased to 2031  $\pm$  65 at 18 h. Importantly, IL-6-induced IL-10 expression was inhibited, in a dose-dependent manner, by JAK Inhibitor 1 or Tofacitinib (Figure 7E and F).

Next, we determined the effect of exogenous IL-10 on IL-6-induced MMP-9 expression. When macrophages were pre-treated (1 h) with as little as 1 ng/ml IL-10, subsequent exposure to IL-6 failed to induced MMP-9 expression (Figure 8A). The IL-10-dependent inhibition of MMP-9 expression was blocked, when macrophages were pre-incubated with neutralizing anti-IL-10 receptor IgG (Figure 8B). These data clearly demonstrate that pre-exposure of macrophages to IL-10 potently inhibits the ability of IL-6 to induce MMP-9 expression. In addition, IL-10 effectively turned off previously up-regulated MMP-9 expression. As seen in Figure 8C, levels of MMP-9 mRNA in macrophages incubated with IL-6 for 18 hours was increased ~3-fold; whereas, levels of MMP-9 mRNA in cells incubated with IL-6 for six hours followed by the addition of IL-10 were similar to control macrophages at 18 hours. Likewise, levels of MMP-9 in conditioned media of cells

incubated 6 hours with IL-6 prior to the addition of IL-10 were reduced >50%, as compared macrophages treated with IL-6 alone (Figure 8C). Together, these data demonstrate that exogenously added IL-10 is a potent inhibitor of IL-6 induced MMP-9 expression by macrophages.

To directly test whether endogenously synthesized IL-10 inhibits IL-6-induced MMP-9 expression, macrophages were treated with IL-6 in the presence of anti-mouse IL-10 IgG or normal IgG. Levels of MMP-9 mRNA in macrophages incubated with IL-6 were increased 5.8 fold over control cells (Figure 9A;  $P < 0.05$ ). In the presence of anti-IL-10, levels of MMP-9 mRNA increased to 14-fold ( $p < 0.01$ ). Likewise, IL-6-induced MMP-9 levels in macrophage conditioned media were increased in the presence of anti-IL-10 IgG (Figure 9A). To corroborate these findings, macrophages were transfected with either nonspecific (NS)-siRNA or IL-10 siRNA, and treated with IL-6 for 18 h. Levels of IL-10 in macrophage conditioned media were significantly ( $p < 0.01$ ) reduced in IL-6 treated cells transfected with IL-10 siRNA (Figure 9B). As predicted, levels of IL-6-induced MMP-9 mRNA were significantly ( $p < 0.01$ ) increased in cells transfected with IL-10 siRNA. Taken together, these data demonstrate that IL-6 induced IL-10 serves to limit MMP-9 expression, and administration of JAK inhibitors, which selectively block IL-10 expression, led to increased MMP-9 expression.

## Discussion

In the present study, we have examined the regulatory role of IL-6 on macrophage MMP-9 expression, which participates in diverse physiologic and pathologic processes. Similar to other MMP family members, the degradation of extracellular matrix components is generally thought to be the primary role of MMP-9. However, in addition, MMP-9 modifies the activities of cytokines and chemokines, growth factors, and proteinase inhibitors (18-20). In these studies, we demonstrate that IL-6 rapidly and robustly stimulated an increase in MMP-9 mRNA levels in RAW264.7 macrophages and elicited peritoneal macrophages, which led to an accumulation of MMP-9 in their conditioned media. These data reveal that IL-6, recognized as a critical mediator of inflammation, induces macrophage expression of a metalloproteinase that is directly associated with the pathogenesis of chronic inflammatory diseases and cancer (18-20).

We and others have reported that macrophage MMP-9 expression is stimulated by PGE<sub>2</sub> (21-29). PGE<sub>2</sub>-dependent proteinase expression is dependent on the EP4 prostanoid receptor and subsequent MAPK<sup>erk1/2</sup> activation (27,28). Since it is well established that elevated levels of PGE<sub>2</sub> associated with inflammation depends on the coupling of Cox-2 and mPGES-1 expression (30-33,41,54-59), as well as reduced catabolism by 15-PGDH (35,36), we examined whether IL-6 regulates the expression of these genes in macrophages. Results demonstrate that IL-6 co-induced Cox-2 and mPGES-1 expression, and inhibited the expression of 15-PGDH. Consistent with these changes in gene expression, levels of PGE<sub>2</sub> in macrophage conditioned media were increased. These data suggested a targetable pathway (Cox-2→mPGES-1→PGE<sub>2</sub>) by which IL-6 induced MMP-9 could be inhibited. However, pre-treatment of macrophages with celecoxib, which completely blocked IL-6-induced PGE<sub>2</sub> synthesis, resulted in a partial reduction in MMP-9 mRNA and protein levels.

Insomuch that celecoxib was partially effective, we explored alternative therapeutic approaches to inhibit IL-6 induced MMP-9 expression. In this regard, the IL-6 binding to the IL-6 receptor complex (IL-6R $\alpha$ /gp130) triggers phosphorylation of JAK, and subsequent activation of STAT and SHP2 (60,61). Phosphorylated STAT dimerizes and translocates to the nucleus to regulate STAT-dependent genes; whereas, the phosphorylation of SHP2 plays an important role in the activation of MAPK<sup>erk1/2</sup> and MAPK-dependent gene expression

(7,9). Thus, we examined the ability JAK inhibitors to block MMP-9 expression. The inhibition of JAK predictably blocked IL-6 induced activation of STAT3, but failed to block the activation of MAPK<sup>erk1/2</sup>, and unexpectedly enhanced MMP-9 expression. In contrast, inhibition of MEK-1 blocked IL-6 induced phosphorylation of MAPK<sup>erk1/2</sup> and MMP-9 expression without affecting the phosphorylation of STAT3. Thus, IL-6-induced MMP-9 expression is dependent on the activation of MAPK<sup>erk1/2</sup> and restrained by a JAK-dependent gene product. Although these data were unexpected based on the established model of IL-6-induced signaling, several investigators have reported that diverse JAK inhibitors, including the highly specific Tofacitinib and Ruxolitinib, effectively block STAT activation without inhibiting MAPK<sup>erk1/2</sup> activation (43-47). Although the mechanism(s) underlying JAK-independent activation of MAPK<sup>erk1/2</sup> remains undefined at this time, the binding of Src kinase family member HCK to gp130, distal to the SHP2 docking site, reportedly plays a role in IL-6-induced MAPK<sup>erk1/2</sup> activation (62).

Utilizing pharmacologic and genetic approaches, we identified IL-10 as the IL-6 induced JAK-dependent suppressor of MMP-9 expression in macrophages. First, the robust induction of IL-10 expression by IL-6 was completely blocked by JAK Inhibitor 1 and Tofacitinib. Second, as little as 1 ng/ml exogenous IL-10 completely blocked IL-6 induced MMP-9 expression. Third, IL-6-induced MMP-9 expression in macrophages was significantly increased, when cells were incubated in the presence of neutralizing anti-IL-10 IgG. Fourth, IL-6-induced MMP-9 expression was significantly increased in macrophages in which IL-10 expression was knocked down via transfection with IL-10 siRNA. Thus, JAK-dependent induction of IL-10 is an important feedback mechanism to control IL-6 induced MMP-9 expression in macrophages. These data are consistent with the recognized role IL-10 plays in limiting the inflammatory response (63). For example, LPS-induced IL-10 inhibits expression of pro-inflammatory cytokines TNF $\alpha$ , IL-6 and IL-12 via activation of the JAK/STAT3 pathway (46). Treatment of cells with the selective JAK inhibitors resulted in enhanced production of LPS-induced inflammatory mediators, which was due to inhibition of IL-10 expression and signaling. Likewise, LPS-induced cytokine production is enhanced by macrophages recovered from IL-10 null mice (46). Taken together, JAK inhibitors have the potential to paradoxically increase expression of inflammatory mediators under defined circumstances by inhibiting IL-10 expression.

In the present studies, we report that IL-6 induced macrophage expression of MMP-9 via Cox-2-dependent and -independent pathways, and required the activation of MAPK<sup>erk1/2</sup>. Inasmuch that MMP-9 plays a causal role in the pathogenesis of chronic inflammatory diseases and cancer, these data reveal a direct mechanism by which elevated IL-6 can contribute to these diseases. Current strategies to block IL-6 induced inflammation utilize anti-IL-6R $\alpha$  (i.e. Tocilizumab), soluble gp130 and small molecule inhibitors of JAK (i.e. Ruxolitinib and Tofacitinib)(7). As reported here, the use of JAK inhibitors appears to have the unintended consequence of blocking IL-6-induced IL-10 expression resulting in increased MMP-9 expression in macrophages.

### Abbreviations used in this paper

<b>ApoE</b>	apolipoprotein E
<b>CVD</b>	cardiovascular disease
<b>Cox-2</b>	cyclooxygenase-2
<b>EIA</b>	enzyme immunoassay
<b>IL-6R<math>\alpha</math></b>	IL-6 receptor alpha chain



<b>IP</b>	intraperitoneal
<b>LE</b>	low levels of endotoxin
<b>MMP</b>	matrix metalloproteinase
<b>mPGES</b>	microsomal prostaglandin synthase
<b>NS</b>	nonspecific
<b>PGDH</b>	hydroxyprostaglandin dehydrogenase
<b>PVDF</b>	polyvinylidene fluoride
<b>RIPA</b>	radioimmunoprecipitation assay
<b>RQ</b>	relative quantity
<b>siRNA</b>	small interfering RNAI

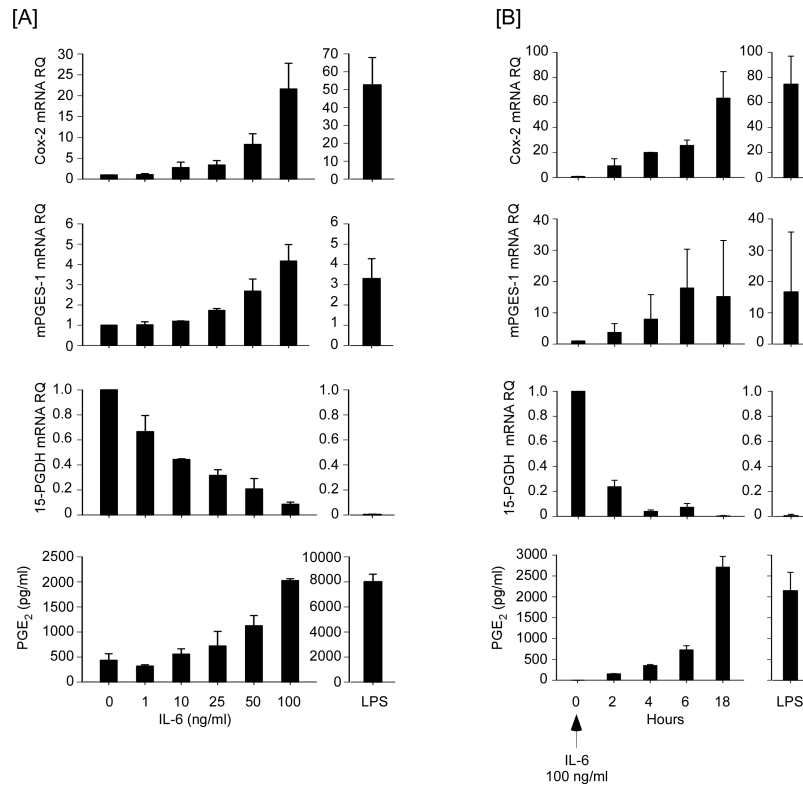
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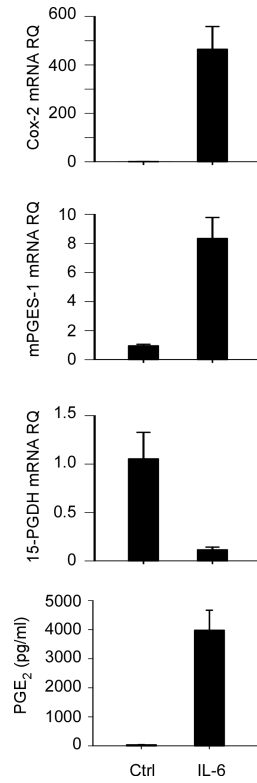
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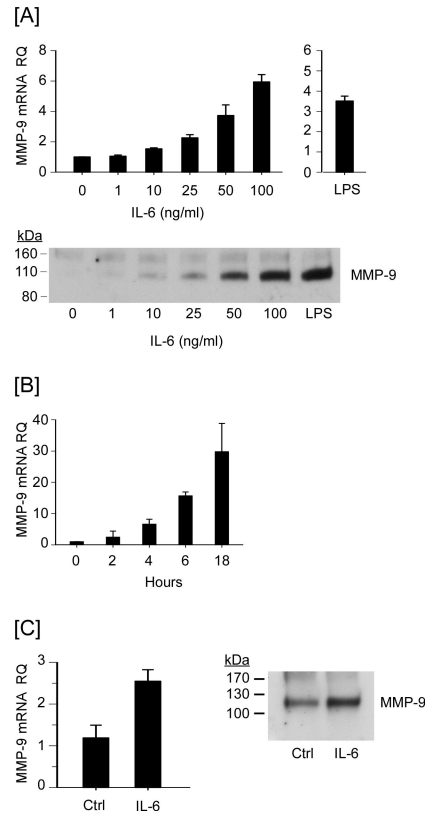
**Figure 1.**

IL-6 induces Cox-2 and mPGES-1 expression, and inhibits 15-PGDH expression in RAW264.7 macrophages. [A] Macrophages were incubated in DMEM-0.1% LE-BSA containing 0 - 100 ng/ml IL-6 or 10 ng/ml LPS. Following 18 h incubation, total RNA and conditioned media were recovered. [B] Macrophages were incubated 0 – 18h in DMEM-0.1% LE-BSA containing 100 ng/ml IL-6 or 10 ng/ml LPS for 18 h. The relative quantity (RQ) of Cox-2, mPGES-1 and 15-PGDH mRNAs were determined by quantitative RT-PCR as described in Materials and Methods. Levels of PGE<sub>2</sub> in conditioned media were determined utilizing ELISA. Target gene mRNA and PGE<sub>2</sub> levels are expressed as the mean  $\pm$  SD of 3 independent samples from a representative experiment.



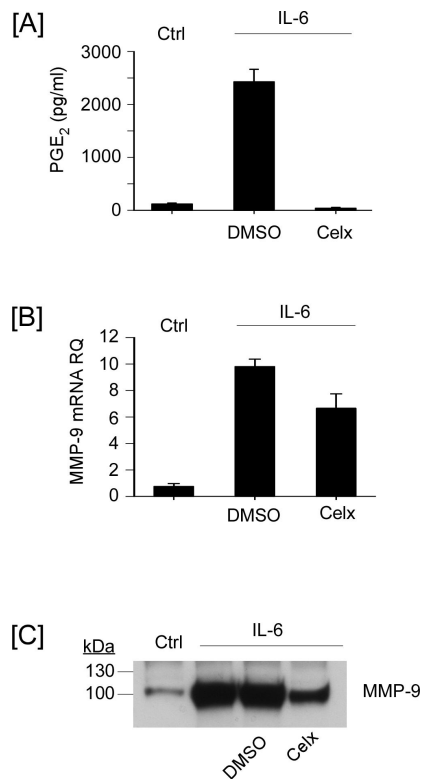
**Figure 2.**

IL-6 induces Cox-2 and mPGES-1, and inhibits 15-PGDH expression in thioglycollate-elicited peritoneal macrophages. Elicited macrophages were incubated in DMEM-0.1% LE-BSA containing 100 ng/ml IL-6 for 18 h and total RNA were recovered. Levels of Cox-2, mPGES-1 and 15-PGDH mRNAs were determined by quantitative RT-PCR, and levels of PGE<sub>2</sub> in conditioned media were determined utilizing ELISA. Target gene mRNA and PGE<sub>2</sub> levels are expressed as the mean  $\pm$  SD of 3 independent samples from a representative experiment.



**Figure 3.**

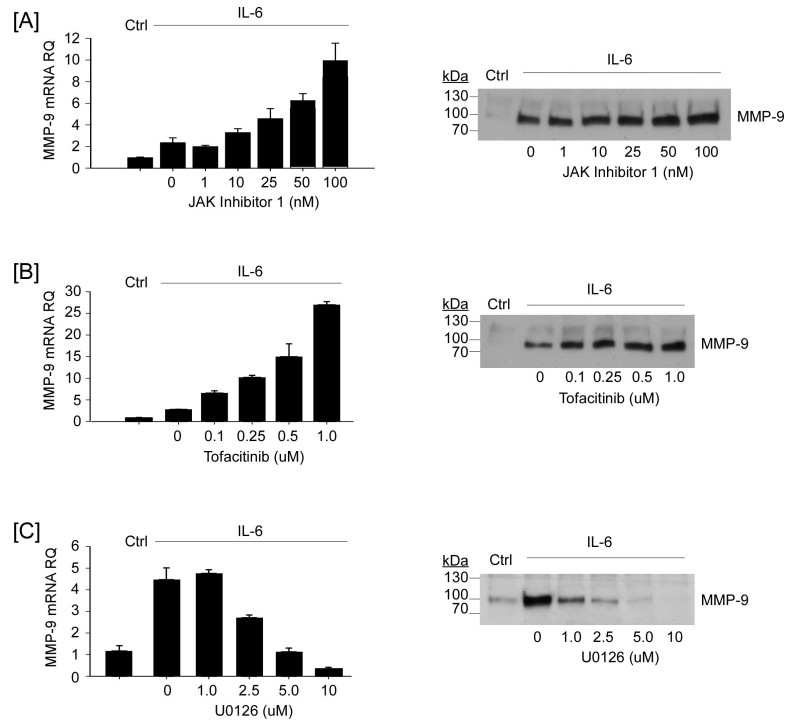
IL-6 induces MMP-9 expression in macrophages. [A] RAW264.7 macrophages were incubated 18 h in DMEM-0.1% LE-BSA containing 0 - 100 ng/ml IL-6 or 10 ng/ml LPS. [B] Macrophages were incubated 0 – 18h in DMEM-0.1% LE-BSA containing 100 ng/ml IL-6. [C] Thioglycollate-elicited macrophages were incubated 18 h in DMEM-0.1% LE-BSA containing 100 ng/ml IL-6. Following the indicated incubation, total RNA and conditioned media were recovered. Levels of MMP-9 mRNA and MMP-9 in conditioned media were determined utilizing quantitative RT-PCR and Western blot, respectively. MMP-9 mRNA levels are expressed as the mean  $\pm$  SD of 3 independent samples from a representative experiment.



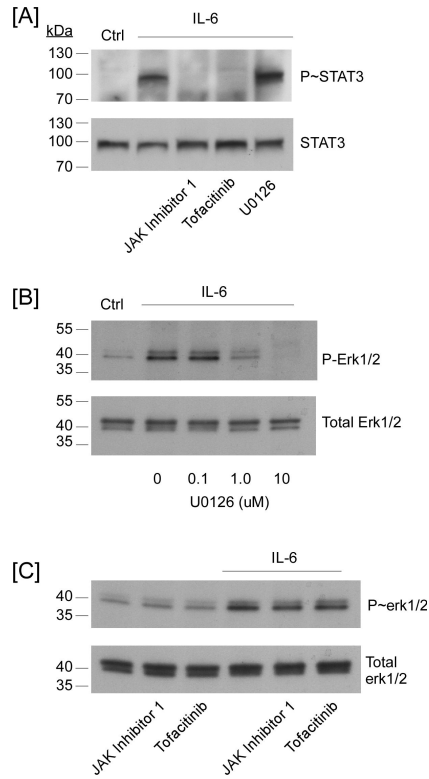
**Figure 4.**

Cox-2-dependent PGE<sub>2</sub> synthesis contributes to IL-6-induced MMP-9 expression. [A] RAW264.7 macrophages were pre-incubated in DMEM-0.1% LE-BSA containing DMSO (vehicle control) or 5  $\mu$ M celecoxib for 30 min, following which 100 ng/ml IL-6 was added. Following 18 h incubation, conditioned media were recovered for determination of [A] PGE<sub>2</sub> and [C] MMP-9 levels utilizing ELISA and Western blot, respectively. [B] Total RNA were recovered and levels of MMP-9 mRNA were determined utilizing quantitative RT-PCR. Levels of PGE<sub>2</sub> and MMP-9 mRNA are expressed as the mean  $\pm$  SD of 3 independent samples from a representative experiment.



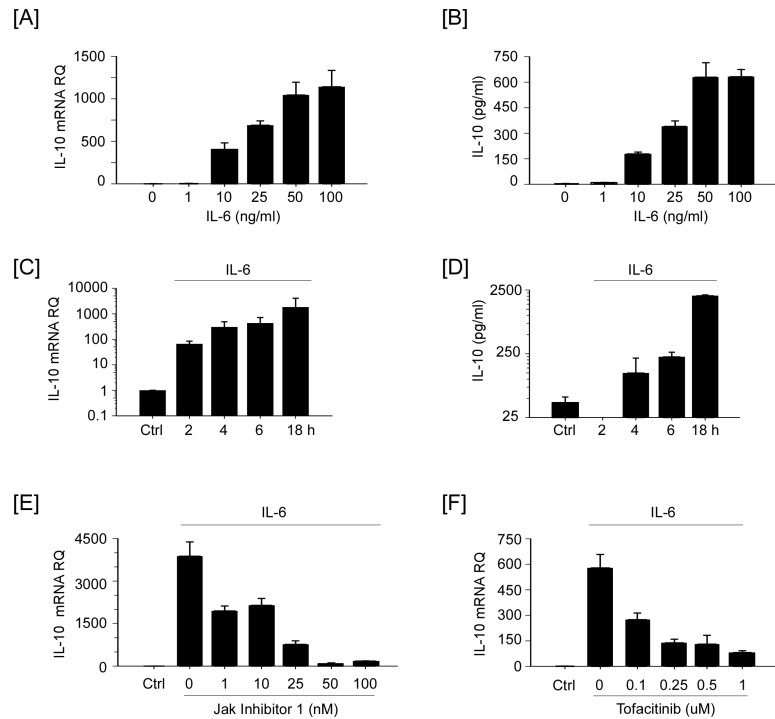


**Figure 5.** Opposing effects of JAK and MEK-1 inhibitors on IL-6-dependent MMP-9 expression. RAW264.7 macrophages were pre-incubated 30 min in DMEM-0.1% LE-BSA containing [A] 0-100 nM JAK Inhibitor 1, [B] 0-1.0  $\mu$ M Tofacitinib or [C] 0-10  $\mu$ M U0126, followed by the addition of 100 ng/ml IL-6. Following 18 h incubation, total RNA and conditioned media were recovered. Levels of MMP-9 mRNA and MMP-9 in conditioned media were determined utilizing quantitative RT-PCR and Western blot, respectively. MMP-9 mRNA levels are expressed as the mean  $\pm$  SD of 3 independent samples from a representative experiment.



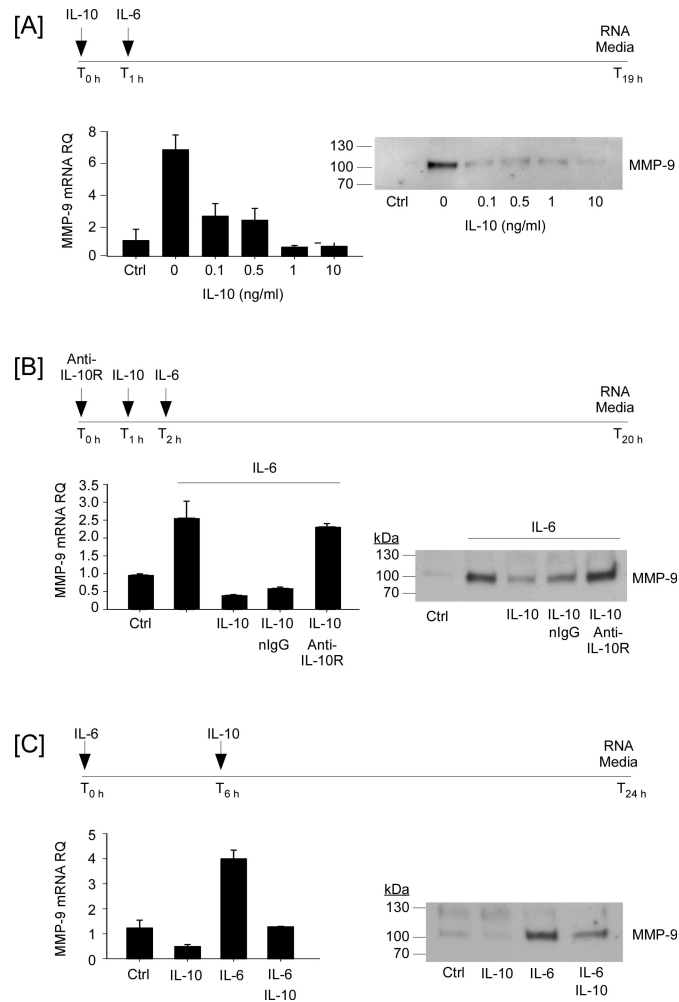
**Figure 6.**

JAK inhibitors block IL-6-induced phosphorylation of STAT3 in macrophages without affecting phosphorylation of MAPK<sup>erk1/2</sup>. [A] RAW264.7 macrophages were pre-incubated 30 min in DMEM-0.1% LE-BSA containing 100 nM JAK Inhibitor 1, 1.0 μM Tofacitinib or 10 μM U0126, followed by the addition of 100 ng/ml IL-6. Following 90 min incubation, cell lysates were recovered, and levels of phosphorylated and total STAT3 were determined utilizing Western blot. Macrophages were pre-incubated 30 min in DMEM-0.1% LE-BSA containing [B] 0-10 μM U0126, [C] 100 nM JAK Inhibitor 1 or 1.0 μM Tofacitinib followed by the addition of 100 ng/ml IL-6. Following 90 min incubation, cell lysates were recovered, and levels of phosphorylated and total MAPK<sup>erk1/2</sup> were determined utilizing Western blot. Data are representative of two separate experiments.

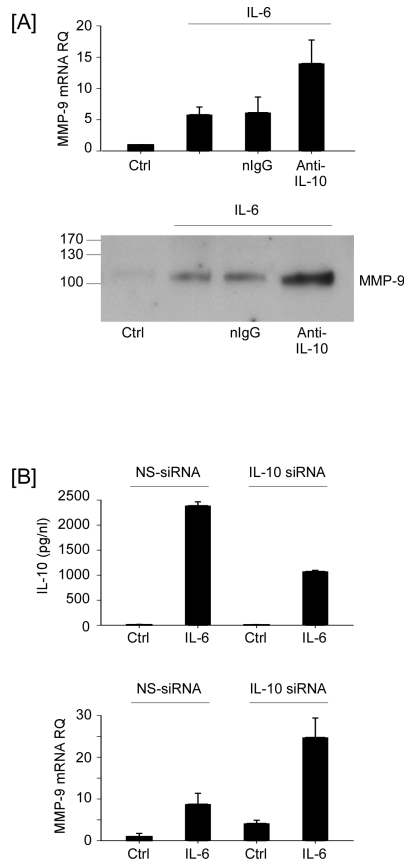


**Figure 7.**

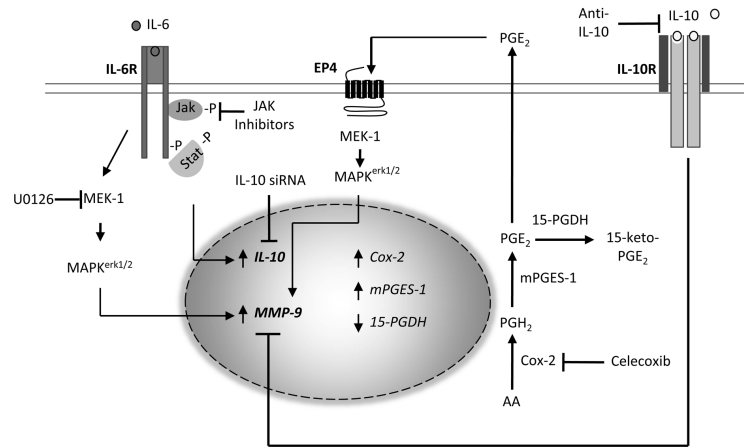
JAK inhibitors block IL-6 induced IL-10 expression in macrophages. [A and B] RAW264.7 macrophages were incubated in DMEM-0.1% LE-BSA containing 0-100 ng/ml IL-6 for 18 h. [C and D] Macrophages were incubated for 18 h in media alone (Ctrl) or 2 - 18 h in media containing 100 ng/ml IL-6. Following the recovery of total RNA and conditioned media, levels of IL-10 mRNA and IL-10 in conditioned media were determined utilizing quantitative RT-PCR and ELISA, respectively. Macrophages were pre-incubated 30 min in DMEM-0.1% LE-BSA containing [E] 0-100 nM JAK Inhibitor 1 or [F] 0-1.0  $\mu$ M Tofacitinib, following which 100 ng/ml IL-6 was added and cells incubated for 18 h. Following the recovery of total RNA, levels of IL-10 mRNA were determined. IL-10 mRNA levels and IL-10 in conditioned media are expressed as the mean  $\pm$  SD of 3 independent samples of a representative experiment.

**Figure 8.**

IL-10 potently suppresses IL-6-induced MMP-9 expression in macrophages. [A] RAW264.7 macrophages were pre-incubated 1 h in DMEM-0.1% LE-BSA containing 0-100 ng/ml recombinant mouse IL-10, following which 100 ng/ml IL-6 was added, and incubated 18 h. [B] Macrophages were pre-incubated 1 h in DMEM-0.1% LE-BSA containing non-immune (n) IgG or anti-IL-10 receptor (R), followed by 1h incubation with 10 ng/ml IL-10, and an additional 18 h incubation with 100 ng/ml IL-6. [C] Macrophages were incubated DMEM-0.1% LE-BSA containing 100 ng/ml IL-6 for 6 h, following which 10 ng/ml IL-10 was added and incubated for 18 h. Following the recovery of total RNA and conditioned media, levels of MMP-9 mRNA and MMP-9 in conditioned media were determined utilizing quantitative RT-PCR and Western blot, respectively. Levels of MMP-9 mRNA are expressed as the mean  $\pm$  SD of 3 independent samples of a representative experiment.



**Figure 9.** Inhibition of IL-10 activity or expression enhances IL-6-induced MMP-9 expression in macrophages. [A] RAW264.7 macrophages were incubated in DMEM-0.1% LE-BSA alone (Ctrl) or media supplemented with 100 ng/ml IL-6, IL-6 and 2 ug/ml anti-murine IL-10 IgG, or IL-6 and normal IgG for 18 h. Following the recovery of total RNA, levels of MMP-9 mRNA were determined utilizing quantitative RT-PCR and levels of MMP-9 in conditioned media were determined by Western blot. [B] Macrophages transfected with nonspecific (NS) or IL-10 siRNA received DMEM-0.1% LE-BSA (Ctrl) or media containing 100 ng/ml IL-6, and were incubated 18 h. Total RNA was recovered and levels of MMP-9 mRNA were determined utilizing quantitative RT-PCR, and levels of IL-10 in conditioned media were determined utilizing ELISA. Data are the mean  $\pm$  SD of 3 independent samples of a representative experiment.



**Figure 10.**

IL-6-mediated induction of MMP-9 is modulated by JAK/STAT-dependent IL-10 expression in macrophages. IL-6, heretofore recognized as a marker of CVD, induces macrophage expression of MMP-9, which is directly associated with the pathogenesis of vascular diseases. IL-6 stimulation of MMP-9 expression was dependent on the activation of MAPK<sup>erk1/2</sup>, and partially dependent on the synthesis of PGE<sub>2</sub>, which we have previously shown to bind EP4 and activate MAPK<sup>erk1/2</sup>. IL-6 also induced IL-10 expression, which potently suppressed MMP-9 expression. Small molecule inhibitors of JAK blocked IL-6-induced IL-10 expression, and augmented MMP-9 expression. Likewise, inhibition of IL-10 expression utilizing siRNA or engagement of the IL-10R utilizing anti-IL-10 led to enhanced MMP-9 expression.