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## PKC-epsilon and TLR4 synergistically regulate resistin-mediated inflammation in human macrophages

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### Abstract

**Background and aims**—Resistin has been associated with atherosclerotic inflammation and cardiovascular complications. We and others have previously shown that PKC-epsilon (PKC $\epsilon$ ) is involved in resistin-induced smooth muscle cell (VSMC) dysfunction at a high pathological concentration. This study aimed to evaluate the role and potential pathways of resistin at a physiological concentration, in atherosclerosis-related inflammation.

**Methods**—Plasma from patients with atherosclerosis was analyzed for resistin concentration. Patients were divided into tertiles based on resistin levels and cytokines were compared between tertiles. Macrophages were then treated with resistin in the presence or absence of PKC $\epsilon$  inhibitor and/or TLR4 blocking-antibody, and their inflammatory state was evaluated with ELISA, RT-PCR, immunocytochemistry, and Western blot.

**Results**—We observed significant associations between plasma resistin levels and TNF- $\alpha$ , IL-6, IL-12, MIP-1 $\alpha$ , MIP-1 $\beta$ , and CD40L. Our *in vitro* analyses revealed that resistin activated PKC $\epsilon$  via TLR4. This was followed by NF- $\kappa$ B activation and induction of a pro-inflammatory phenotype in macrophages, significantly upregulating CD40, downregulating CD206 and stimulating gene expression and secretion of the inflammatory cytokines, for which we found association in our plasma analysis. Resistin also induced persistent TRAM and CD40L upregulation up to 36 hours after resistin treatment. PKC $\epsilon$  and TLR4 inhibitors suppressed gene expression to levels similar to control, especially when used in combination.

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#### Conflict of interest

The authors declared they do not have anything to disclose regarding conflict of interest with respect to this manuscript.

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**Conclusions**—Resistin, at a physiological concentration, exacerbates the inflammatory response of macrophages. PKCε is a key upstream mediator in resistin-induced inflammation that may interact synergistically with TLR4 to promote NF-κB activation, while TRAM is an important signal. PKCε and TRAM may represent novel molecular targets for resistin-associated chronic atherosclerotic inflammation.

### Keywords

resistin; macrophage; cytokine; atherosclerosis; protein kinase C epsilon; toll-like receptor 4

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### Introduction

Resistin, an adipokine with a cysteine-rich C terminus, was originally thought to link obesity and diabetes due to its ability to ‘resist’ insulin action in murine adipocytes.<sup>1</sup> However, further investigations revealed key differences between mouse and human resistin. In humans, resistin is mainly produced by monocytes and macrophages, primary inflammatory mediators in atherosclerosis.<sup>2, 3</sup> Unlike in mice, human resistin is a 12.5 kDa peptide whose gene resides on chromosome 19, and it has different promoter regions, suggesting different regulatory mechanisms and tissue distribution.<sup>4</sup> Thus, the specific function and underlying mechanisms of resistin action in human pathology are still debated.

Clinical observations indicate that resistin has important roles in cardiovascular disease and atherosclerotic inflammation. Recently, the Multi-Ethnic Study of Atherosclerosis (MESA), with over 1900 subjects, found a strong association between higher resistin levels and incidence of cardiovascular disease (CVD) and heart failure.<sup>5</sup> Resistin has also been associated with coronary artery disease and linked to increased risk of CVD for diabetic patients<sup>6</sup> and cardiovascular events in patients with different comorbidities.<sup>7–10</sup> These clinical studies support resistin roles in atherosclerosis and its associated complications. However, the mechanistic insights have not yet been elucidated.

Human plasma resistin level was found to be positively correlated with the levels of tumor necrosis factor-α receptor-2 (TNFα-R2) and interleukin 6 (IL-6) in non-diabetic individuals.<sup>11</sup> Moreover, several *in vitro* studies have investigated resistin’s atherogenic properties. Human recombinant resistin has been shown to increase adhesion molecules expression in endothelial cells, promoting monocyte attachment.<sup>12, 13</sup> We and others have demonstrated that resistin promotes vascular smooth muscle cell migration, proliferation, and dedifferentiation, and we reported that protein kinase C epsilon (PKCε) mediates such processes.<sup>14–16</sup> Resistin can also induce IL-6 and TNF-α productions in peripheral blood mononuclear cells (PBMCs)<sup>17</sup> and promote inflammation in macrophages.<sup>18</sup> Tarkowski et al. have proposed that toll like receptor 4 (TLR4) may serve as a binding protein for resistin.<sup>19</sup> However, the specific pathways triggered by resistin to promote its inflammatory properties in macrophages remain uncertain. Additionally, the levels of resistin being used in these studies ranged from 25 ng/ml to 100 μg/ml, which are vastly different from physiological (clinically relevant) and pathological levels reported in patients with atherosclerosis and cardiovascular disease.<sup>5, 8, 11</sup> This study aimed to elucidate the role and mechanistic pathways of resistin in atherosclerosis-related inflammation at a clinically

relevant concentration. Hence, using a highly translational approach, we first evaluated patients with atherosclerosis for resistin levels and the relationship of resistin with other inflammatory cytokines in our patient population. We then chose a resistin concentration consistent with our cohort findings and explored the molecular mechanisms triggering resistin-mediated inflammation in macrophages. We focused on the role of upstream modulators, specifically PKC $\epsilon$ , its interaction with TLR4, and in TLR4-related downstream pathways in macrophages.

## Materials and methods

### Human plasma analysis

This study was approved by the Stanford University Investigational Review Board and the Palo Alto VA Research and Development Committee (IRB 23476), and it conforms to the principles outlined in the Declaration of Helsinki. Written informed consent was obtained from each patient included in the study. Plasma samples were collected from 99 patients undergoing carotid surgery interventions following an established protocol. Samples were stored at  $-80^{\circ}\text{C}$  and analyzed with a Luminex magnetic bead-based assay for circulating cytokine levels at the Stanford University Human Immune Monitoring Center.

### Monocytes isolation and culture

PBMCs were collected from buffy coats of healthy control male donors (Stanford Blood Center), age  $>20$  using the Ficoll-Paque density gradient method. Then, cluster of differentiation 14 (CD14)-positive monocytes, were isolated using negative magnetic sorting (Stemcell Technologies). Population purity was assessed with flow cytometry by conjugating a sample of sorted cells to CD14-FITC antibody (Invitrogen). Cells were differentiated to macrophages during a 7-day culture in RPMI media supplemented with 10% FBS and 100 ng/mL of macrophage colony stimulating factor (PeproTech). Effective macrophage differentiation was confirmed by immunocytochemistry as described below. Differentiated macrophages were then treated with resistin.

### *In vitro* cell treatments

Human recombinant resistin (PeproTech) was used to treat cells. To choose a treatment concentration, we first evaluated resistin's ability to polarize human macrophages by testing concentrations representative of our patient population (2, 5, and 10 ng/ml, Figure 2A). Based on results from these studies, we chose 10 ng/ml for further *in vitro* experiments. For all experiments, except for time-specific evaluations, cells were treated for 18 hours based on literature for macrophage polarization and phenotype-priming studies.<sup>20–22</sup> The PKC $\epsilon$ -specific inhibitor  $\epsilon\text{V1-2}$  (from Dr. Mochly-Rosen Laboratory, Stanford University) and an anti-toll like receptor 4 (TLR4) antibody (Abcam)<sup>23</sup> were used for inhibitor and receptor-blocking studies, respectively.  $\epsilon\text{V1-2}$  was used at 1  $\mu\text{M}$  concentration;<sup>24</sup> anti-TLR4 at 0.5  $\mu\text{g/ml}$ ;<sup>23</sup> with a 30 min pre-treatment prior to the addition of resistin. The TLR4 inhibitor TAK-242 (EMD Millipore) was also used for confirmatory studies at 2  $\mu\text{M}$  concentration, with a 30 min pre-treatment time.

### Real-time polymerase chain reaction (RT-PCR)

Total RNA from macrophages was isolated using TRIzol according to standard protocols. SYBR green PCR master mix was used for real time PCR. Human primers are listed in Supplementary Table 1.<sup>25–29</sup> RT-PCR was performed in a Mastercycler RT-PCR detection system (Eppendorf, Westbury, NY). The relative level of target (respective cytokine or key pathway modulators) gene in each group was normalized against internal housekeeping gene  $\beta$ -actin using the calculation formula of  $2^{-CT(\beta\text{-actin-target})}$ . The target gene levels in drug treated groups were further normalized against the control group.

### Macrophage cytokine measurement

Conditioned media from macrophage cultures were analyzed using standard ELISA manufacturer's protocols (Millipore, Invitrogen, and Thermo Scientific). Our analysis focused on macrophage pro-inflammatory cytokines that were found elevated in our patient population, were shown upregulated gene expression after resistin treatment, and have been proposed to play key pro-inflammatory roles in atherosclerosis. These included resistin, TNF- $\alpha$ , IL-6 and CD40L.

### Fluorescent immunocytochemistry

Phenotypic signatures were evaluated with fluorescent immunocytochemistry. Briefly, cells were differentiated to macrophages as described above. After respective treatments, coverslips with macrophages were fixed, washed, and incubated with primary and fluorescence-conjugated secondary antibodies according to manufacturer's recommendations and as previously described.<sup>30</sup> Cells were visualized with laser scanning confocal microscope (ZEISS Confocal LSM-710) using DAPI staining as the focal reference point. Co-expression CD68 and CD11b was used as pan-macrophage marker, accounting for successful Macrophage differentiation. For the M1 subtype, we evaluated expression of cell surface markers CD40 and CD80; and for the M2 subtype, expression of CD163 and Mannose Receptor (CD206). At least 5 representative images (covering different areas of the coverslip) were recorded per treatment group. Mean fluorescence intensity (MFI) was measured using ImageJ and ZEN 2010 6.0 software packages, and relative MFI for each maker was compared among groups.

### PKC activity

Total cell lysates were collected and the quantification of PKC $\epsilon$ -specific activity was carried out with a PKC activity kit (Enzo Life sciences) according to the manufacturer's instructions. Cell samples were assayed in the presence and absence of eV1–2 (1  $\mu$ M) and the difference in values is used to calculate PKC $\epsilon$ -specific activity. The assay was quantified with a spectrophotometric microplate reader (iMark, Bio-Rad) at a dual wavelength of 450/595 nm. Data is presented as amount of active PKC $\epsilon$  ng/ $\mu$ g protein.

### Western blot

Cell lysates (10  $\mu$ g of protein) separated on a 4–20% polyacrylamide gel for electrophoresis were transferred onto nitrocellulose membranes using Bio-Rad Mini-Trans-Blot system. The membranes were blocked with 5% BSA in TBS with Tween-20 (TBST, 0.1% Tween-20) at

room temperature. The blocked membrane was incubated with primary antibodies overnight at 4°C and with secondary antibody for 1 hour at RT. The immunoreactive bands were detected using a BIO-RAD chemiluminescence system, and the bands were captured and intensity quantified by with BIO-RAD ChemiDoc XRS+ camera and BIO-RAD Image Lab software respectively.

### Statistical analysis

All *in vitro* experiments were performed independently at least four times in duplicate. Results are expressed as the mean +SEM. Statistical analyses were performed with THE GraphPad Prism 6 (GraphPad Software, Inc., La Jolla, CA) software package. Data sets were tested for parametric or non-parametric analysis, and the appropriate statistical test was selected. Dose-dependent data were compared by one-factor analysis of variance (ANOVA) followed by Dunnett's test. Data comparing inhibitors effect, from PKC $\epsilon$  inhibitor and TLR4-blocking experiments, on gene expression, protein expression, and resistin secretion, were analyzed using two-factor ANOVA followed by Tukey's test. To compare differences between tertiles in cytokine levels and comorbidity distributions, categorical variables were analyzed with the Chi-square test and continuous variables with ANOVA, followed by Bonferroni correction for the multiple-comparisons test. Statistical significance was considered if the *p*-value was <0.05.

## Results

### Clinical characteristics and medical comorbidities in patients with varying levels of resistin

Arterial blood plasma from 99 male patients who underwent carotid interventions with a mean age of 69.3 years were analyzed. Medical comorbidities included hypertension (94%), diabetes (42%), CAD (49%), and obesity (42%), and most patients had a history of smoking (83%). Subjects were divided into tertiles (n=33 each) based on plasma resistin concentration (mean, range): low (2257.5 pg/ml, 1031.6 pg/ml – 3416.3 pg/ml), medium (4225.2 pg/ml, 3420.6 pg/ml – 4874.2 pg/ml), and high (7285.5 pg/ml, 4968.4 – 13266.5 pg/ml). The distribution of comorbidities, pre-operative symptoms, and medications was similar within each tertile (Table 1).

### High resistin tertile exhibits higher levels of inflammatory cytokines/chemokines

There was no significant difference in various growth factors (vascular endothelial growth factor A, macrophage colony-stimulating factor, and granulocyte-macrophage colony-stimulating factor), neurotrophic factors (brain-derived neurotrophic factor and beta-nerve growth factor), or adhesion molecules (intercellular adhesion molecule 1 and vascular cell adhesion protein 1) between the tertiles. However, significantly higher levels of the pro-inflammatory cytokines/chemokines were observed in the high resistin tertile compared to the low tertile cohort including tumor necrosis factor alpha (TNF- $\alpha$ ; *p*<0.001; Fig. 1A), interleukin-6 (IL-6; *p*<0.01; Fig. 1B), interleukin-12 subunit p40 (IL-12p40; *p*<0.01; Fig. 1C), macrophage inflammatory protein 1-alpha (MIP-1 $\alpha$ ; *p*<0.05; Fig. 1D), macrophage inflammatory protein 1-beta (MIP-1 $\beta$ ; *p*<0.05; Fig. 1E), and soluble CD40 ligand (CD40L; *p*<0.05; Fig. 1F). Similarly, significantly higher levels of the aforementioned pro-

inflammatory cytokines were observed in the high compared to the medium tertile: IL-6,  $p < 0.05$ ; IL-12p40,  $p < 0.01$ ; MIP-1 $\beta$ ,  $p < 0.05$ ; and CD40L,  $p < 0.05$ .

### Resistin promotes a pro-inflammatory phenotype in macrophages

We examined the direct effects of physiological levels of resistin on macrophage-mediated inflammation *in vitro*. To evaluate if resistin changes the relative polarization state of differentiated macrophages we used double markers to discern between the two end-of-the-spectrum macrophage states: M1 and M2. The M1 phenotype is typically associated with a pro-inflammatory state in macrophages, whereas the M2 phenotype is attributed to tissue repair activities. CD40 and CD80 have been observed to be upregulated in M1.<sup>31, 32</sup> On the other hand, the mannose receptor (CD206) and CD163, a scavenger receptor, are generally attributed to the M2 phenotype.<sup>33, 34</sup> First, macrophages were treated with resistin at a various resistin levels, representative of our patient population (2, 5, and 10 ng/ml) and examined for CD40 expression (Fig. 2A, top panel and Supplementary Fig. 1A). We observed that resistin dose dependently upregulated CD40 expression, and results at 10 ng/ml were verified with immunofluorescence (Fig. 2A, bottom panel). We therefore decided to use 10 ng/ml of resistin for our remaining experimental studies, which is within the high resistin tertile range (5–13 ng/ml) of our patient population. In addition, increased sections of key pro-inflammatory cytokines in macrophages including resistin itself (Fig. 2C), TNF- $\alpha$  (Fig. 2D), CD40L (Fig. 2E), and IL-6 (Fig. 2F) were observed following resistin treatment, while CD206 expression was downregulated as evidenced by Western blot results (Fig. 2B, top panel and Supplementary Fig. 1B) and immunofluorescence (Fig. 2B, bottom panel). However, not significant change in CD80 (Supplementary Fig. 1C) or CD163 (Supplementary Fig. 1D) was observed.

### Resistin activates PKC $\epsilon$ via TLR4 through a resistin-induced positive feedback loop

Resistin significantly increased PKC $\epsilon$  activity as early as 30 min post treatment, and PKC $\epsilon$  remained active throughout the time points investigated and peaking at 6 hours (Fig. 2G). As PKC $\epsilon$  was significantly active at 30 min, we performed our inhibitory studies at this time point. We observed that blocking TLR4 or inhibiting PKC $\epsilon$  with  $\epsilon$ V1–2 both prevented resistin-induced PKC activation (Fig. 2H). In addition, resistin-mediated CD40 upregulation was significantly inhibited by  $\epsilon$ V1–2 and by blocking TLR4 (Fig. 2I), reiterating the TLR4–PKC $\epsilon$  connection in resistin-induced inflammation. Moreover, resistin treatment further increased resistin gene expression and protein secretion (Supplementary Fig. 2A and B), which was significantly higher than the treatment dose, suggesting a positive feedback loop. The increase in gene expression was time-dependent, peaking at 18 hours. Inhibition with  $\epsilon$ V1–2 and anti-TLR4 blocking mitigated gene upregulation and secretion of resistin (Supplementary Fig. 2A and B).

### PKC $\epsilon$ mediates resistin-induced upregulation of pro-inflammatory cytokine genes in macrophages

Resistin significantly increased gene expressions of CD40L, IL-12p40, IL-6, TNF- $\alpha$ , MIP-1 $\alpha$ , MIP-1 $\beta$  after 3 hours of treatment at 10 ng/ml. The trend persisted at 6 hours and up to 18 hours for CD40L, IL-12p40, IL6, and TNF-1 $\alpha$  (Fig. 3A–C). 36 hours after treatment, CD40L remained significantly elevated in the resistin-treated group (Fig. 3D).

Our research group has shown that PKC $\epsilon$  mediates resistin-induced vascular smooth muscle cell-migration,<sup>24</sup> but its involvement in resistin-triggered macrophage activation had not been previously investigated. Thus, we used  $\epsilon$ V1–2, a PKC $\epsilon$  specific inhibitor, to further study PKC $\epsilon$  regulatory roles in macrophages. We observed that  $\epsilon$ V1–2 markedly inhibited resistin-induced cytokine gene upregulations (Fig. 3). In addition, we tested the involvement of TLR4, since it has been suggested as a receptor for resistin in mononuclear cells,<sup>19</sup> and we observed similar inhibitory trends as  $\epsilon$ V1–2 (Fig. 3). The relative suppression of resistin-induced upregulation of inflammatory cytokines varied for  $\epsilon$ V1–2 and TLR4-blocking, but it appeared to be more effective at the early time point of 3 hours. However, a combined inhibition of PKC $\epsilon$  and TLR4 proved more effective, bringing cytokine/chemokine expression to levels similar to control.

### Resistin activates TLR4-mediated downstream pathways in macrophages

We observed significant increases in TLR4 and PKC $\epsilon$  gene expressions at 3 hours (Fig. 4A), 6 hours (Fig. 4B), and 18 hours (Fig. 4C) following resistin treatment. Significant upregulations of TRIF-related adaptor molecule (TRAM) and myeloid differentiation primary response gene 88 (MYD88), well-known TLR4 downstream pathways, were also observed up to 18 hours; however, only TRAM upregulation was statistically significant after 36 hours of treatment (Fig. 4D). The PKC $\epsilon$  inhibitor and TLR4 antibody suppressed resistin-induced gene upregulation of TLR4, PKC $\epsilon$ , TRAM, and MYD88 at early time points, but they were less effective at later time points. Similarly to gene expression of cytokines, when both TLR4 was blocked and PKC $\epsilon$  inhibited, gene expression returned to levels similar to control. The nuclear factor kappa-light-chain-enhancer of activated B cells (NF- $\kappa$ B) subunits showed a trend for upregulated expression in the early time points (3 and 6 hours), but only the NF- $\kappa$ B1 subunit was significant at 6 hours. In addition, we evaluated the protein expression of TLR4, PKC $\epsilon$ , TRAM, and MYD88 after 18 hours of resistin treatment and observed similar trends (Supplementary Fig. 3). Finally, we tested NF- $\kappa$ B activation by evaluating I $\kappa$ B $\alpha$  phosphorylation in lysates from resistin-treated macrophages. Resistin induced increased I $\kappa$ B $\alpha$  phosphorylation, an indicator NF- $\kappa$ B activation (Supplementary Fig. 4A). When TLR4 or PKC $\epsilon$  were inhibited in macrophages prior to resistin treatment, I $\kappa$ B $\alpha$  phosphorylation was prevented (Supplementary Fig. 4B), suggesting that TLR4 and PKC $\epsilon$  mediate resistin-induced NF- $\kappa$ B activation.

### Discussion

The specific roles of resistin in cardiovascular disease and atherosclerosis have long been debated. Although it has been suggested that resistin's inflammatory properties promote its pathological consequences, the mechanisms and pathways remain uncertain. Furthermore, studies performed with super-pathological concentrations of resistin, which are far beyond the observed clinically relevant physiological ranges, have produced unclear results.<sup>18</sup> To our knowledge, this is the first systematic study to evaluate the effects of resistin that are representative of a physiological level in patients with atherosclerosis. We validated the inflammatory property of resistin and identified several novel mechanistic interactions.

Reilly et al. were the first to propose that resistin serves as an atherosclerotic inflammatory marker after finding a positive association between plasma levels of resistin and TNF- $\alpha$  and IL-6 in the context of coronary atherosclerosis.<sup>13</sup> Others have linked serum resistin to inflammatory activation in hypertension,<sup>35</sup> metabolic syndrome,<sup>36</sup> and rheumatoid arthritis.<sup>37</sup> Comparing patients with similar comorbidities and demographics, we too observed significant associations between plasma resistin and TNF- $\alpha$  and IL-6. In addition, we identified several novel associations between resistin and IL-12, MIP-1 $\alpha$ , MIP-1 $\beta$ , and CD40L, cytokines that are typically secreted by macrophages in their pro-inflammatory phenotype.<sup>38–40</sup>

Our observations in our patient cohort prompted a more detailed *in vitro* investigation into the role of resistin in the vasculature, its inflammatory stimuli, and the pathways involved. We utilized our clinical results to guide the selection of a clinically relevant resistin concentration for our *in vitro* studies and to select the markers and pathways to be investigated. Using a stepwise approach, we first evaluated macrophage phenotype changes, where we observed that resistin significantly induced CD40 upregulation and CD206 downregulation, suggesting pro-inflammatory transformation of macrophages.<sup>41</sup> In addition, we also observed that resistin increased secretion of TNF- $\alpha$ , CD40L, and IL-6 and gene expression of CD40L, IL-12p40, IL-6, TNF- $\alpha$ , MIP-1 $\alpha$ , and MIP-1 $\beta$ . These are well known, pro-inflammatory cytokines and markers, which further validate a pro-inflammatory macrophage transformation induced by resistin at high physiological levels. Although not significant, we observed a trend for downregulation for IL-10, an M2-related cytokine. These comprehensive evaluations prove, for the first time, that resistin has an independent inflammatory property that promotes macrophage pro-inflammatory polarization, inducing an inflammatory cascade of cytokines and chemokines. We also show for the first time that resistin induced further resistin expression and secretion in macrophages, possibly via a positive feedback loop. However, we cannot eliminate the possible contribution of other cytokines produced by resistin-treated macrophages to promote resistin expression and the overall inflammatory state of macrophages. Likewise, Shyu et al. have previously suggested that TNF- $\alpha$  induces resistin expression.<sup>42</sup>

Our previous studies have shown that PKC $\epsilon$  mediates resistin-induced VSMC migration, proliferation, and dedifferentiation.<sup>16, 24</sup> In this study, we observed that PKC $\epsilon$  is a key mediator of resistin signaling for macrophage activation, suggesting that inhibiting PKC $\epsilon$  has anti-inflammatory properties. Our study is in line with previous findings proposing that TLR4 participates in resistin-associated activity in macrophages.<sup>19</sup> TLR4 signaling and interaction between CD40-CD40L have both been independently attributed to atherosclerotic plaque development and atherosclerosis-mediated inflammation.<sup>25, 43–45</sup> Hoebe et al. have demonstrated that binding of lipopolysaccharides to TLR4 induced upregulation of co-stimulatory signals, including CD40 in macrophages via TRAM,<sup>46</sup> and Lundberg et al. recently reported that TRAM signaling is crucial in TLR4-mediated atherosclerotic inflammation.<sup>47</sup> Unique to this study, we observed persistent TRAM and CD40L upregulation up to 36 hours after resistin treatment, suggesting that TRAM-related downstream activation and CD40L expression mediate chronic cellular effects in macrophages. We suspect that the positive feedback mechanism triggered by resistin may

contribute to continuous pro-inflammatory stimulation leading to the to100-fold increase in TRAM and up to 1400-fold increase in CD40L.

Studies have shown that PKC $\epsilon$  is phosphorylated by TLR4 when signaling through MyD88 and TRAM.<sup>48, 49</sup> Our results suggested that resistin may act through similar pathways to trigger pro-inflammatory properties in macrophages. First, we showed that it increased PKC $\epsilon$  activity significantly as early as 30 min after treatment, and that this effect was prevented when TLR4 was blocked, indicating a sequential nature in PKC $\epsilon$  activation. Second, we observed that resistin induced increased gene and protein expression of these downstream molecules (TRAM and MYD88) and that inhibition of PKC $\epsilon$ /TLR4 blocks such effects. Finally, although resistin only moderately increased gene expression of NF- $\kappa$ B at early hours, it significantly triggered I $\kappa$ B $\alpha$  phosphorylation, an indicator of NF- $\kappa$ B activation, within an hour of treatment. I $\kappa$ B $\alpha$  holds NF- $\kappa$ B in the cytoplasm, but its phosphorylation releases NF- $\kappa$ B to translocate to the nucleus. NF- $\kappa$ B is a common transcription factor for TLR4-mediated inflammatory processes. Its activation indicates inflammatory transformation of macrophages. Overall, our findings suggest that resistin interacts with TLR4, activates PKC $\epsilon$  within 30 min, and then elicits NF- $\kappa$ B activation to promote inflammation in macrophages, which further induces increased expression of pro-inflammatory cytokines and molecules.

There were a few unexpected observations in our study. We observed that TLR4 blocking and PKC $\epsilon$  inhibition were more effective at early time points, and that each alone only partially blocked gene upregulations of TRAM and CD40L. When both TLR4 and PKC $\epsilon$  were inhibited, expressions of TRAM and CD40L returned to control levels. These findings suggest that TLR4 and PKC $\epsilon$  synergistically propagate resistin-mediated gene expression of inflammatory signals in macrophages. Finally, although we focused on inflammatory cascades in this study, we cannot ignore the possible connection between reactive oxygen species (ROS) production and inflammatory transformation of macrophages. We recently showed that resistin increased ROS production, NADPH oxidase (Nox) activity, and inflammatory cytokine secretion in VSMCs.<sup>16</sup> We speculate that ROS production and Nox activity may also contribute to this pathway, and play important roles in the overall inflammatory cascades that are activated by resistin.

In conclusion, our study highlights the mechanistic insight in resistin-mediated atherosclerotic inflammation. Our findings, for the first time, suggest that PKC $\epsilon$  is a key mediator for resistin-induced signaling and activation in macrophages. We also offer valid evidence that resistin at a physiological concentration has the potential to exacerbate the inflammatory response of macrophages, increasing cytokines and CD40/CD40L expression. Additionally, our findings allude to TRAM as an important downstream signaling molecule to induce NF- $\kappa$ B activation. Targeting resistin-mediated inflammation and its downstream signaling pathways may present a novel strategy in atherosclerosis prevention and management.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

## Acknowledgments

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## Abbreviations

<b>M<math>\Phi</math></b>	macrophages
<b>PKC<math>\epsilon</math></b>	protein kinase C epsilon
<b>TLR4</b>	toll-like receptor 4
<b>TRAM</b>	TRIF-related adaptor molecule
<b>MyD88</b>	myeloid differentiation primary response gene 88
<b>NF-<math>\kappa</math>B</b>	nuclear factor kappa-light-chain-enhancer of activated B cells
<b>CVD</b>	cardiovascular disease

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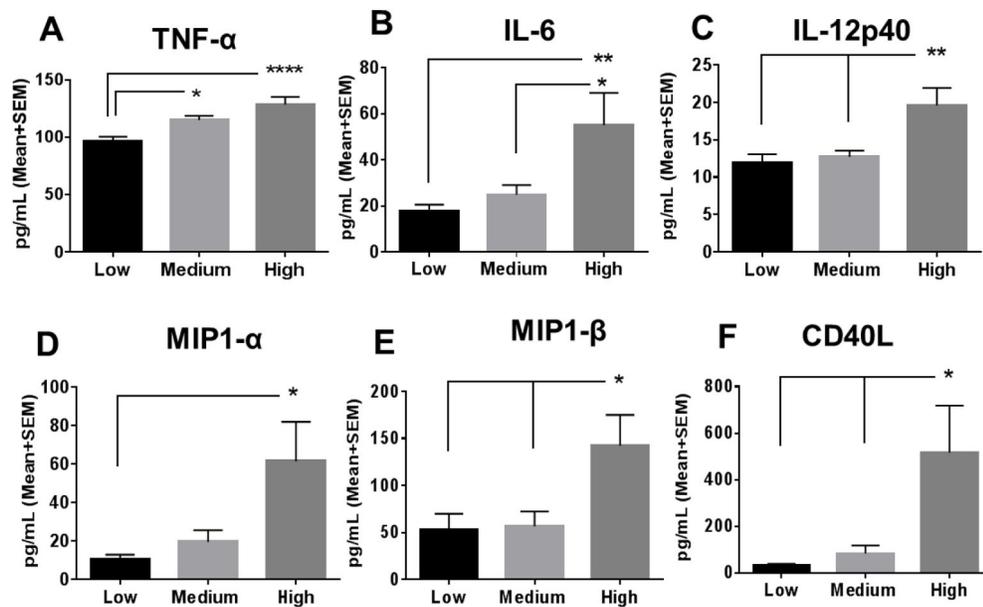
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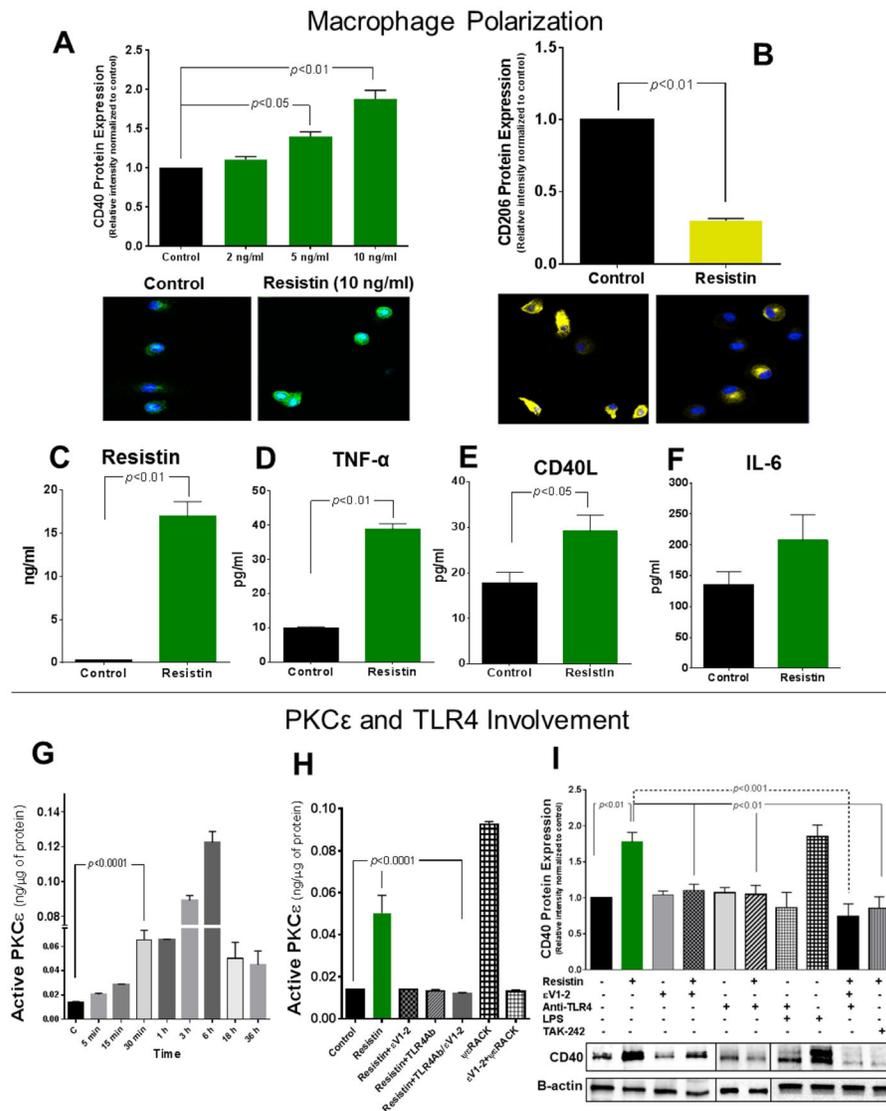
### Highlights

- High plasma resistin levels are associated with high levels of inflammatory cytokines in patients with atherosclerosis.
- Resistin induces a pro-inflammatory phenotype in macrophages via a TLR4 and PKC $\epsilon$ .
- Resistin stimulates NF- $\kappa$ B activity, significantly upregulating CD40, downregulating CD206 and stimulating gene/protein expression of inflammatory molecules and cytokines.
- Resistin treatment further increased resistin gene expression and protein secretion, suggesting a positive feedback loop.
- TRAM and CD40L are persistently upregulated up to 36 hours after resistin treatment.



**Fig. 1. Levels of inflammatory cytokines in patients with clinical atherosclerosis**

Subjects were divided in tertiles (n=33 each) based on resistin levels: low resistin (Low; mean=2257.5 pg/ml), medium resistin (Medium; mean =4225.2 pg/ml), and high resistin (High; mean=7285.5 pg/ml), and inflammatory cytokines were compared between tertiles. Significantly higher levels of (A) tumor necrosis factor alpha (TNF- $\alpha$ ), (B) interleukin-6 (IL-6), (C) interleukin-12 subunit p40 (IL-12p40), (D) macrophage inflammatory protein 1- $\alpha$  (MIP-1 $\alpha$ ), (E) macrophage inflammatory protein 1- $\beta$  (MIP-1 $\beta$ ), and (F) CD40 ligand (CD40L) were observed in the High tertile as compared to the Low tertile. Data are shown as mean + SEM, \* $p$  < 0.05; \*\* $p$  < 0.01; and \*\*\*\* $p$  < 0.0001.



**Fig. 2. Resistin induces a pro-inflammatory macrophage phenotype, activates PKC $\epsilon$  via TLR4, and initiates a positive feedback loop in macrophages**

(A) Macrophages treated with resistin at various physiological levels, representative of our patient population, showed significant upregulation of CD40 expression after 18 hours of treatment, as demonstrated by Western blot (top panel) and immunofluorescence (bottom panel). (B) Resistin promoted decreased expression of CD206 at 10 ng/ml as analyzed by Western blot (top panel) and immunofluorescence (bottom panel). Resistin increased pro-inflammatory cytokine secretion of resistin (C), TNF- $\alpha$  (D), CD40L (E), and IL-6 (F), after 18 hours of 10 ng/ml treatment. (G) Specific PKC $\epsilon$  activity was measured at various time points after 10 ng/ml of resistin treatment, and PKC $\epsilon$  was significantly activated after 30 min of treatment. (H) Inhibition of TLR4 prevented PKC $\epsilon$  activation (30 min time point).  $\psi$ RACK, a specific PKC $\epsilon$  activator, was used as a positive control at 1  $\mu$ M concentration. (I) PKC $\epsilon$  inhibitor and TLR4-blocking prevent resistin-induced CD40 upregulation; LPS (100 ng/ml) was used as a positive control; and TAK-242, a TLR4 inhibitor, at 2  $\mu$ M, was

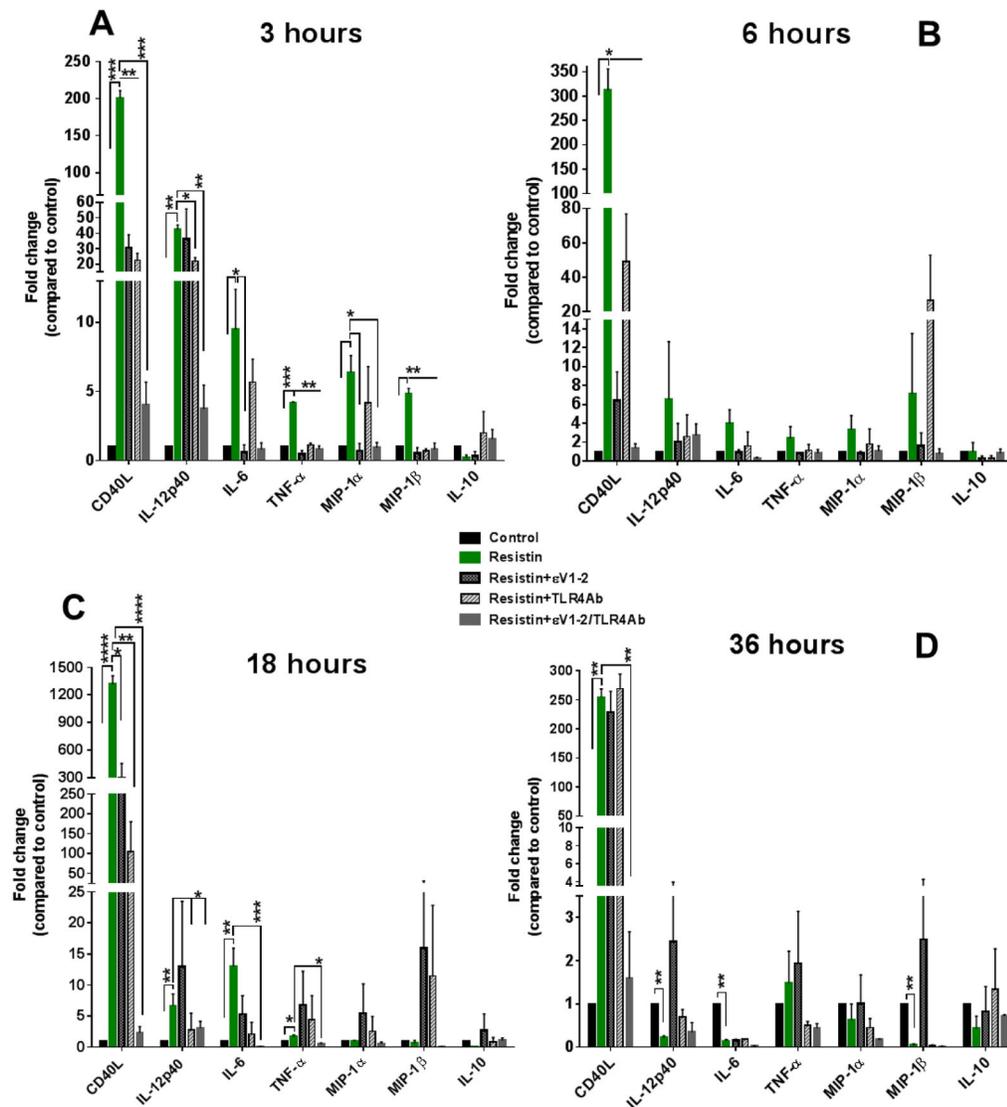
used to confirm TLR4 involvement in resistin-induced upregulation. Data are shown as mean + SEM of at least 4–12 independent experiments analyzed by one-way or two-way ANOVA.

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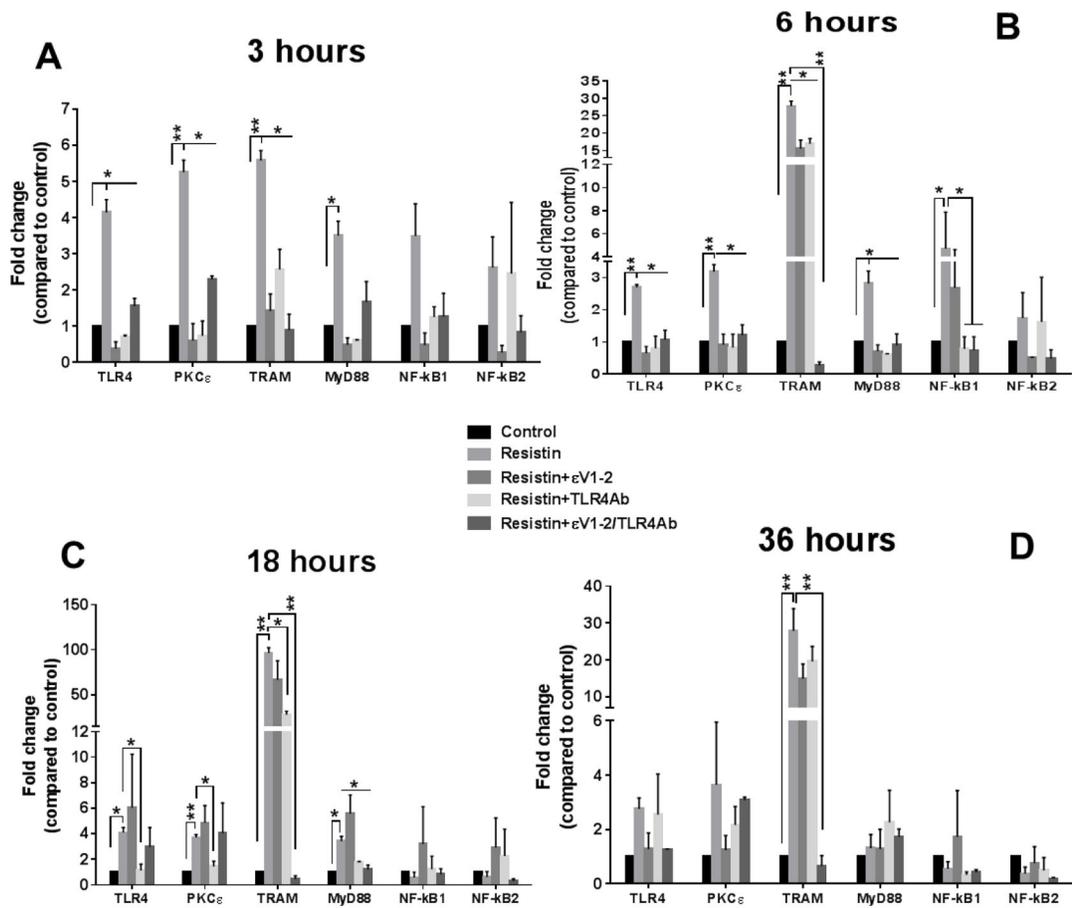
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**Fig. 3. PKC $\epsilon$  and TLR4 synergistically promote resistin-induced upregulation of pro-inflammatory cytokine genes in macrophages**

Resistin upregulated gene expression of inflammatory cytokines, which was examined after 3 hours (A), 6 hours (B), 18 hours (C), and 36 hours (D) of resistin treatment at 10 ng/ml. Inhibition of TLR4 or PKC $\epsilon$  partially suppressed resistin-induced upregulation (A and C), but this was more effective when  $\epsilon$ V1-2/TLR4-blocking were used in combination. Resistin-treated cells were compared to control and to PKC $\epsilon$ -inhibited/TLR4-blocked cells for the different cytokines. Data are shown as mean + SEM. \* $p$ <0.05; \*\* $p$ <0.01; \*\*\* $p$ <0.001; and \*\*\*\* $p$ <0.0001 by one-way or two-way ANOVA.



**Fig. 4. Resistin significantly upregulates TLR4 and PKCε gene expression as well as several downstream pro-inflammatory pathways in macrophages**

Resistin increased gene expression of TLR4, PKCε and key mediators of downstream inflammatory pathways after 3 hours (A), 6 hours (B), 18 hours (C), and 36 hours (D) of resistin treatment at 10 ng/ml. TRAM expression remained statistically significant after 36 hours of treatment (D). Inhibition of TLR4/PKCε prevented resistin-induced increased gene expression (A–D). Resistin-treated cells were compared to control and to PKCε-inhibited/TLR4-blocked cells for the different molecules/transcription factors. Data are shown as mean + SEM. \* $p < 0.05$  and \*\* $p < 0.01$  by one-way or two-way ANOVA.

**Table 1**

Distribution of comorbidities and risk factors in resistin tertiles.

Variables	1st tertile, n =33 (Low)	2nd tertile, n=33 (Medium)	3rd tertile, n=33 (High)	p-value
Age, average yrs (range)	68 (54–84)	70 (59–87)	68 (55–91)	0.24
Procedure (CAS/CEA)	16/17	22/11	18/15	0.32
Risk factors and comorbidities				
History of smoking, n	27	29	26	0.61
Diabetes, n	10	16	16	0.23
Obesity, n	9	16	17	0.09
CAD, n	18	12	19	0.18
Hypertension, n	29	31	33	0.20
Preoperative symptoms				
Symptomatic, n	16	18	13	0.46
Prior stroke, n	7	5	8	0.64
Key medications				
Antiplatelets, n	24	24	20	0.47
Satins, n	28	28	30	0.70
Anticoagulants, n	0	4	2	0.12

CAS, carotid artery stenting; CEA, carotid endarterectomy; CAD, coronary artery disease.