TNF-α Signals Through PKCζ/NF-κB to Alter the Tight Junction Complex and Increase Retinal Endothelial Cell Permeability

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OBJECTIVE—Tumor necrosis factor- α (TNF- α) and interleukin-1 beta (IL-1 β) are elevated in the vitreous of diabetic patients and in retinas of diabetic rats associated with increased retinal vascular permeability. However, the molecular mechanisms underlying retinal vascular permeability induced by these cytokines are poorly understood. In this study, the effects of IL-1 β and TNF- α on retinal endothelial cell permeability were compared and the molecular mechanisms by which TNF- α increases cell permeability were elucidated.

RESEARCH DESIGN AND METHODS—Cytokine-induced retinal vascular permeability was measured in bovine retinal endothelial cells (BRECs) and rat retinas. Western blotting, quantitative real-time PCR, and immunocytochemistry were performed to determine tight junction protein expression and localization.

RESULTS—IL-1β and TNF-α increased BREC permeability, and TNF-α was more potent. TNF-α decreased the protein and mRNA content of the tight junction proteins ZO-1 and claudin-5 and altered the cellular localization of these tight junction proteins. Dexamethasone prevented TNF-α-induced cell permeability through glucocorticoid receptor transactivation and nuclear factor-kappaB (NF-κB) transrepression. Preventing NF-κB activation with an inhibitor κB kinase (IKK) chemical inhibitor or adenoviral overexpression of inhibitor κB alpha (IκBα) reduced TNF-α-stimulated permeability. Finally, inhibiting protein kinase C zeta (PKCζ) using both a peptide and a novel chemical inhibitor reduced NF-κB activation and completely prevented the alterations in the tight junction complex and cell permeability induced by TNF-α in cell culture and rat retinas.

CONCLUSIONS—These results suggest that PKC ζ may provide a specific therapeutic target for the prevention of vascular permeability in retinal diseases characterized by elevated TNF- α , including diabetic retinopathy. *Diabetes* **59:2872–2882, 2010**

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he cause of vision loss in diabetic retinopathy is complex and remains incompletely understood; however, changes in blood vessel permeability and macular edema are associated with loss of visual acuity (1-5), with center point thickness and fluorescein leakage combined with age accounting for 33% of the variation in visual acuity (5). A growing body of evidence suggests that diabetic retinopathy includes a neuroinflammatory component, with increased expression of cytokines, microglia activation, leukostasis, and vascular permeability (6-9). Increased levels of interleukin-1 beta (IL-1 β) and tumor necrosis factor- α (TNF- α) have been detected in the vitreous of diabetic patients with proliferative diabetic retinopathy (10,11) and in diabetic rat retinas (6,7,12). Moreover, intravitreal administration of IL-1β increases vascular permeability, associated with nuclear factor-kappaB (NF-κB) activation, increased leukocyte adhesion, and retinal capillary cell death (13,14). TNF-α also increases leukocyte adhesion to retinal endothelium (15) and blood-retinal barrier (BRB) permeability (16). The inhibition of TNF- α with etanercept, a soluble TNF-α receptor, inhibits NF-κB activation, leukostasis, and BRB breakdown in diabetic rat retinas (12). Altogether, these findings indicate that proinflammatory cytokines contribute to vascular permeability in diabetic retinopathy.

Changes in retinal vascular permeability may result from alterations of the tight junction complex. Tight junctions are composed of a combination of more than 40 proteins including the transmembrane proteins occludin, the claudin family, and the junction adhesion molecule (JAM) family, several peripheral membrane-associated proteins, including members of the zonula occludens (ZO) family, and several regulatory proteins (17). To date, evidence has been provided for the presence of occludin, claudin-5, ZO-1, and JAM-A in retinal vascular endothelium (18–21). Changes in occludin content, localization, and phosphorylation occur in response to vascular endothelial growth factor (VEGF) regulating endothelial permeability (18,19,22–24). Gene deletion studies have shown claudin-5 to be essential for blood-brain barrier function (25), and likely for the BRB as well. Recent studies demonstrated that ZO proteins are essential for the formation and organization of tight junction complex assembly (26,27). Therefore, changes in occludin, claudin-5, or ZO-1 likely contribute to alterations in endothelial permeability, in response to inflammatory cytokines.

Although several studies support the involvement of proinflammatory cytokines in BRB breakdown in diabetes, little attention has been given to the molecular mechanisms involved. Therefore, this study was designed to test the hypothesis that alterations to the tight junction protein complex confer endothelial permeability in response to TNF- α and to determine the signaling pathways underlying TNF- α -induced permeability.

TNF- α was shown to regulate permeability in an NF- κ B-dependent mechanism by downregulating both claudin-5 and ZO-1 expression. Further, protein kinase C zeta (PKC ζ) activity is essential for TNF- α -induced endothelial permeability, partly through NF- κ B activation, suggesting that targeting PKC ζ may provide a novel therapy to control retinal vascular permeability.

RESEARCH DESIGN AND METHODS

Materials used to carry out the experiments are described in the supplementary Materials and Methods in the online appendix available at http://diabetes.diabetes.journals.org/cgi/content/full/db09-1606/DC1.

Primary bovine retinal endothelial cell culture. Bovine retinal endothelial cells (BRECs) were isolated and grown as previously described (28), except medium was supplemented with 8 μ g/ml tylosin. For experimentation, BRECs were grown to confluence and then cell culture media was changed to MCDB-131 medium supplemented with 1% FBS, 0.01 ml/ml antibiotic/antimycotic, and 8 μ g/ml tylosin for one day and exposed to IL-1 β or TNF- α .

Measurement of BREC permeability. BREC monolayer permeability to 70 kDa rhodamine isothiocyanate dextran (Sigma-Aldrich, St. Louis, MO) was measured exactly as described previously (24). The average $P_{\rm o}$ for control conditions was 2.77×10^{-6} cm/s.

Animal model. Male Sprague-Dawley rats (Charles River Laboratories, Wilmington, MA) weighing 150-175 g were used to evaluate retinal vascular permeability and tight junction proteins localization in retinal blood vessels. Animals were housed under a 12-h light/dark cycle with free access to water and a standard rat chow. All experiments were conducted in accordance with the Association for Research in Vision and Ophthalmology (ARVO) Statement for the Use of Animals in Ophthalmic and Vision Research, and were approved and monitored by the Institutional Animal Care and Use Committee (IACUC) at the Penn State College of Medicine. Under anesthesia (66.7 mg/kg ketamine and 6.7 mg/kg xylazine), the animal received an intravitreal injection of TNF-α or vehicle (2.5 μl/eye) with a 5 μl Hamilton syringe (Hamilton Company, Reno, NV) through a puncture created by a 32-gauge needle. The animals were assessed for retinal permeability with Evans blue assay, 24 h after receiving the intravitreal injection of either PBS with 1% BSA, 10 ng TNF-α, 280 ng PKCζI-1, or TNF-α plus PKCζI-1. In a separate study, retinas were harvested 4 h after injection, immunolabeled for tight junction proteins, and analyzed by confocal microscopy.

Evans blue assay. Accumulation of the albumin binding dye Evans blue was used to assess changes in retinal vascular permeability. Evans blue dye accumulation in the retina was quantified using a published protocol (29), with normalization to blood plasma after 2-h circulation and expressed as microliters of plasma per gram of retina (dry weight) per hour of circulation. Assessment of cell viability. The LIVE viability assay (Molecular Probes, Invitrogen, Carlsbad, CA) was used for the assessment of cell viability according to manufacturer's instructions, as described in the supplementary Materials and Methods.

Caspase-3/7 activity assay. Caspase-3/7 activity was measured by the Apo-ONE Homogenous caspase-3/7 assay (Promega, Madison, WI), according to the manufacturer's instructions, as briefly described in the supplementary Materials and Methods.

Western blotting. Western blotting of cellular lysates was performed as described in the supplementary Materials and Methods.

RNA extraction and reverse transcription. RNA extraction and reverse transcription were performed as described in the supplementary Materials and Mathods.

Quantitative real-time PCR. Quantitative real-time PCR (qPCR) analysis was performed using the 7,900 HT Sequence Detection System in 384-well optical plates using TaqMan Universal PCR Master Mix Assay-on-Demand and Assay-by-Design primers and probes (Applied Biosystems) as previously described (30). Primer/probes used were as follows: claudin-5 Bt03288088_s1, occludin Bt03255225_m1, IL-8 Bt03211907_g1, and bovine ZO-1 specific primers 5'-AGAAAGATGTTTATCGTCGCATCGT-3' (forward), 5'-ATTCCT TCTCATATTCAAAATGGGTTCTGA-3' (reverse), and 6-carboxy-fluorescein (FAM) 5'-ACCCACATCGGATTCT-3' minor groove binding (MGB) probe. For each sample, qPCR were performed in triplicate and relative quantities were calculated using ABI SDS 2.0 RQ software and the $2^{-\Delta\Delta Ct}$ analysis method

(31), with glyceral dehyde-3-phosphate dehydrogenase (Bt03210919_g1) as the endogenous control.

Immunocytochemistry. Tight junction protein cellular localization was evaluated by immunocytochemistry, as previously described (32). Cells were incubated with a rat anti-ZO-1 monoclonal antibody culture supernatant (1:5), rabbit polyclonal anti-claudin-5 (1:500), or anti-occludin (1:300) antibodies overnight at 4°C. Primary antibodies were detected with Alexa Fluor 647conjugated anti-rat IgG (Molecular Probes, Invitrogen) or Cy3-conjugated anti-rabbit IgG secondary antibodies (Jackson Immunoresearch Laboratories, West Grove, PA). Coverslips were mounted onto slides using Aqua Poly/Mount (Aquamount; Polysciences, Warrington, PA) and analyzed in a Leica TCS SP2 AOBS confocal microscope. Ten confocal Z-stacks were collected over a depth of 2.56 µm and projected onto one image. ZO-1 and occludin localization in retinal vessels were assessed by immunohistochemistry in whole retinas, as described previously (33). The retinas were incubated with monoclonal anti-occludin (1:50) and polyclonal anti-ZO-1 (1:50) antibodies for three days at 4°C. Primary antibodies were detected with Cy2-conjugated anti-mouse IgG and Cy3-conjugated anti-rabbit IgG secondary antibodies (Jackson Immunoresearch Laboratories) for 24 h at 4°C. After incubation, retinas were washed, mounted in slides with Aqua Poly/Mount, and analyzed on a Leica TCS SP2 AOBS confocal microscope.

Adenovirus-mediated IkB α overexpression. IkB α was expressed using the AdEasy adenoviral vector system, as described (34). Subconfluent cells were transduced with adenovirus expressing IkB α (AdIkB α) or with an empty expression cassette as control (AdEmpty) at a multiplicity of infection of 20,000 for 6 h in MCDB complete medium. After incubation, cells were washed and MCDB media supplemented with 1% serum was added for a further 24 h. Successful viral infection was followed by fluorescence microscopy because the AdEasy vector contains a green fluorescent protein (GFP) cassette incorporated into the viral vector. More than 80% of the cells expressed detectable amounts of GFP after 30 h of adenoviral infection.

Luciferase reporter assay. A 293T/NFκB-luc stable reporter cell line (Panomics-Affymetrix, Fremont, CA) was used to evaluate NF-κB transcription factor activity. These cells maintain a chromosomal integrated luciferase reporter gene regulated by multiple copies of the NF-κB response element. After treatments, cells were harvested and lysed with Passive Lysis Buffer (Promega), and the luciferase activity was measured using the Dual-Luciferase Reporter Assay system (Promega) according to manufacturer's instructions. Luciferase activity was normalized to the total protein content of each sample.

Statistical analysis. Results are expressed as mean \pm SEM. Data were analyzed using Student t test or one-way ANOVA followed by Dunnett or Bonferroni post hoc test. A value of P < 0.05 was considered significant. Prism 4.0 (GraphPad Software, San Diego, CA) was used for all statistical analysis.

RESULTS

IL-1β and TNF-α increase BREC permeability. BRECs grown on transwell filters were exposed to increasing concentrations and various times of IL-1 β and TNF- α , and the monolayer permeability was evaluated. IL-1\beta significantly increased cell permeability at a concentration of 10 ng/ml $(220.3 \pm 34.2\% \text{ of control}; \text{ Fig. 1A})$, and this effect was statistically significant after 24 h of treatment (Fig. 1B). Similarly, TNF-α increased cell permeability in a concentration- and time-dependent manner. However, TNF-α was maximally effective at a concentration of 5 ng/ml (365.3 ± 47.8% of control, Fig. 1C) and at a shorter time point than IL-1β (6 h, Fig. 1D). To ensure that the increase in permeability induced by the cytokines was not due to cell death, the effect of IL-1 β and TNF- α on BREC viability was evaluated using IL-1 β at 10 ng/ml and TNF- α at 5 ng/ml. IL-1 β had no effect on cell viability. TNF-α-induced a significant decrease in cell viability after 24 h of treatment (70.9 \pm 2.6% of control), but no effect was observed after 6 h of exposure (Fig. 1E). The effect of IL-1 β and TNF- α on caspase-3/7 activity, a marker of apoptosis, was also evaluated. IL-1β had no effect on caspase-3/7 activity, whereas TNF- α induced a significant increase in caspase-3/7 activity, but again only after 24 h of exposure (Fig. 1F). Therefore, the increase in cell permeability induced by TNF- α after 6 h of exposure was not due to cell death. Because the effect of TNF- α was more robust than that of IL-1β, the molecular mechanisms under-

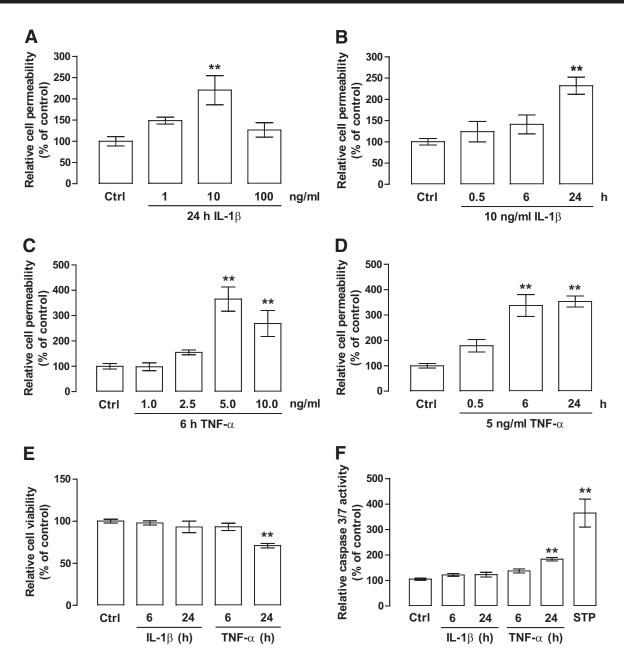
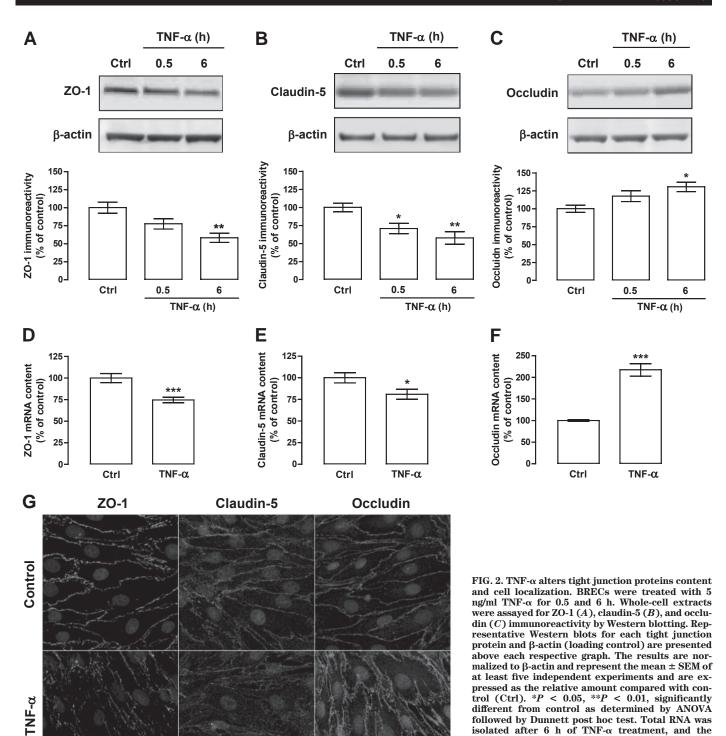


FIG. 1. IL-1 β and TNF- α increase retinal endothelial cell permeability. A-D: BRECs were grown to confluence on transwell filters and then exposed to IL-1 β or TNF- α in a concentration- and time-dependent manner. The monolayer permeability to 70 kDa dextran was measured as described in Research design and methods. A: Cells were treated with 1, 10, and 100 ng/ml IL-1 β for 24 h. B: Cells were treated with 10 ng/ml IL-1 β for 0.5, 6, and 24 h. C: Cells were treated with 1, 2.5, 5, and 10 ng/ml TNF- α for 6 h. D: Cells were treated with 5 ng/ml TNF- α for 0.5, 6, and 24 h. E: Relative cell viability was measured by calcein AM cleavage by live cells. F: Retinal endothelial cell apoptosis was evaluated by caspase-3/7 activation. As a positive control of apoptosis induction, cells were treated with 100 nmol/l staurosporine (STP) for 6 h. The results represent the mean \pm SEM of at least four independent experiments and are expressed relative to control (Ctrl). **P < 0.01, significantly different from control as determined by ANOVA followed by Dunnett post hoc test.

lying TNF- α -induced retinal endothelial cell permeability were further investigated.

TNF- α alters tight junction protein content and localization. To determine the effect of TNF- α on expression of specific tight junction proteins, BRECs were exposed to 5 ng/ml TNF- α for 0.5 and 6 h and the protein contents of ZO-1, claudin-5, and occludin were determined by Western blotting. TNF- α significantly decreased ZO-1 content (58.2 ± 6.5% of control) after 6 h of exposure (Fig. 2A). Claudin-5 content was rapidly reduced after 0.5 h of treatment (70.8 ± 7.0% of control, Fig. 2B), and 6 h of TNF- α exposure further downregulated claudin-5 content

 $(57.6\pm8.7\% \text{ of control})$. In contrast, TNF- α increased occludin content $(130.6\pm6.6\% \text{ of control})$; Fig. 2C). To determine whether alterations in protein content were due to changes in mRNA expression, total mRNA content was evaluated by qPCR 6 h after TNF- α exposure. TNF- α significantly decreased ZO-1 $(74.6\pm3.3\% \text{ of control})$; Fig. 2D) and claudin-5 mRNA content $(80.9\pm5.7\% \text{ of control})$; Fig. 2E) but induced a twofold increase in occludin mRNA (Fig. 2F). To investigate whether TNF- α alters the tight junction complex at the cell membrane, the cellular localization of the tight junction proteins were evaluated by immunocytochemistry and confocal microscopy. In con-



cantly different from control as determined by Student t test. G: Cells were immunolabeled for ZO-1, claudin-5, and occludin 6 h after TNF- α treatment, and 10 confocal Z-stacks were taken through 2.56 μ m and projected into one image. Arrows indicate continuous staining at cell borders. These results are representative of four independent experiments. Scale bar, 25 μ m.

trol conditions, ZO-1, claudin-5, and occludin immunoreactivity appeared as a near continuous staining at the cell border (Fig. 2G and supplementary Fig. 1, available in an online appendix). Upon TNF- α treatment, a loss of both ZO-1 and claudin-5 immunostaining was observed, leading to a fragmented border staining, although the

effect on ZO-1 was more pronounced. Also, a diffuse cytoplasmic distribution of claudin-5 and occludin was observed in TNF- α -treated cells. After TNF- α treatment, occludin staining increased, and it was irregularly distributed at the cell border (Fig. 2G and supplementary Fig. 1).

isolated after 6 h of TNF- α treatment, and the transcript levels of ZO-1 (D), claudin-5 (E), and occludin (F) were analyzed by qPCR. Glyceraldehyde-3-phosphate dehydrogenase was used as an endogenous control. Results represent the mean \pm SEM of eight independent experiments and are expressed as the relative amount compared with control conditions. *P < 0.05, ***P < 0.001, signifi-

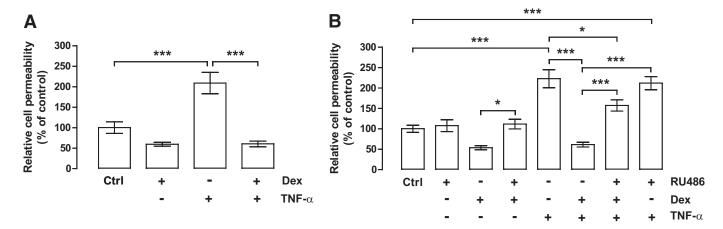


FIG. 3. Dexamethasone prevents TNF- α -induced cell permeability through transactivation of the glucocorticoid receptor. A: BRECs were grown to confluence on transwell filters and treated with 50 ng/ml dexamethasone (Dex) 18 h before TNF- α treatment (5 ng/ml, 6 h). B: Cells were treated with 5 μ mol/l RU486 1 h before Dex treatment. The monolayer permeability to 70 kDa dextran was measured as described in RESEARCH DESIGN AND METHODS. The results represent the mean \pm SEM of at least seven independent experiments and are expressed relative to control (Ctrl). *P < 0.05, ***P < 0.001, significantly different as determined by ANOVA followed by Bonferroni post hoc test.

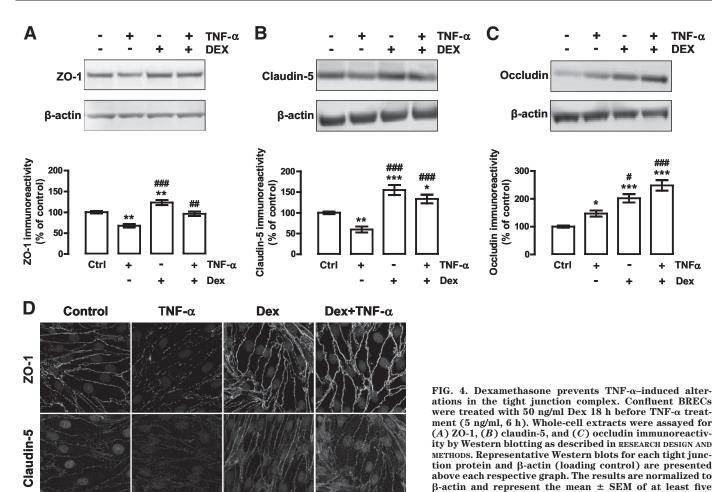
Dexamethasone prevents TNF-α-induced BREC per**meability.** The effect of the glucocorticoid dexamethasone on TNF-α-induced endothelial permeability was determined. BRECs were treated with 50 ng/ml dexamethasone 18 h before TNF- α treatment. The increase in cell permeability induced by TNF- α was completely prevented by dexamethasone (Fig. 3A). To investigate if the effect of dexamethasone was dependent on transcriptional activation by the glucocorticoid receptor, BRECs were pretreated with 5 µmol/l RU486, a glucocorticoid receptor antagonist that inhibits the transactivation function of this receptor (35). A 1-h pretreatment of RU486 before the addition of dexamethasone significantly reduced the protective effect of dexamethasone on TNF-α-induced cell permeability (Fig. 3B). RU486 alone had no effect on permeability and did not alter the TNF- α response but did prevent the dexamethasone reduction in permeability as previously reported (36). These data show that dexamethasone's protective effect is partially due to glucocorticoid receptor transactivation but also suggest that transrepression of TNF-α-responsive transcription factors also contributes to the inhibition of TNF-α-induced cell permeability.

Dexamethasone prevents TNF-α-induced alterations in tight junction proteins. The effect of dexamethasone on TNF-α-induced changes in tight junction protein content and cellular localization was also evaluated. Dexamethasone alone significantly increased ZO-1, claudin-5, and occludin protein content (Fig. 4). TNF-α decreased ZO-1 and claudin-5 protein content and these effects were prevented by dexamethasone pretreatment (Fig. 4A and B), whereas dexamethasone and TNF-α treatment yielded an additive threefold increase in occludin protein content (Fig. 4C). Dexamethasone increased ZO-1, claudin-5, and occludin staining at the cell border and prevented the TNF-α-induced fragmentation of these tight junction proteins (Fig. 4D and supplementary Fig. 1).

NF-κB inhibition reduces TNF- α -induced cell permeability. The involvement of NF-κB activation in TNF- α -induced endothelial permeability was investigated. BRECs were exposed to 1 μ mol/l IKK VII, an IκB kinase complex inhibitor, 30 min before TNF- α addition. As a control for inhibitor IKK VII effectiveness, TNF- α -induced IκB α phosphorylation by IKK was evaluated. Figure 5A shows that

IκBα phosphorylation was blocked by IKK VII. The TNF- α -induced increase in permeability (275.0 \pm 46.7% of control) was significantly reduced by IKK VII (160.5 ± 21.7% of control; Fig. 5B). The effect of adenovirusmediated overexpression of $I\kappa B\alpha$ on cell permeability was also evaluated. Western blotting was used to confirm the adenovirus-mediated expression of GFP and IκBα 30 h after adenoviral infection. GFP expression was similar in both AdEmpty- and AdIκBα-transduced cells, while IκBα was heavily expressed in AdIκBα-transduced cells compared with AdEmpty-transduced cells (Fig. 5C). The ability of IκBα overexpression to inhibit NF-κB activation was evaluated by examining the TNF- α -induced expression of IL-8 mRNA, a transcriptional target of NF-κB. In both nontransduced and AdEmpty-transduced cells, IL-8 mRNA levels were significantly increased (by 18- and 16-fold, respectively) after 2 h of TNF- α stimulation (Fig. 5D). This increase in IL-8 expression was significantly reduced (by 41%) in AdIκBα-transduced cells, demonstrating that adenovirus-mediated overexpression of IκBα effectively blocked NF-κB activation after TNF-α treatment. Subsequently, the effect of $I\kappa B\alpha$ overexpression on $TNF-\alpha$ induced cell permeability was determined. In AdEmptytransduced cells, TNF- α induced a significant increase in cell permeability (273.6 \pm 71.0% of control; Fig. 5E), similar to nontransduced cells (Fig. 5E). IkB α overexpression reduced TNF- α -induced permeability to 157.5 \pm 5.7% of control. These data suggest that NF-kB activation is necessary for a substantial part of the TNF- α -induced changes in BREC permeability.

PKCζI-1 inhibits NF-κB activation induced by TNF-α. PKCζ has a critical role in the activation of the NF-κB pathway (37). To determine whether PKCζ modulates NF-κB activation in retinal endothelial cells induced by TNF-α, the effect of PKCζI-1, a novel PKCζ inhibitor with little or no inhibitory activity on PKCβ or PKCδ (manuscript in preparation), on TNF-α-induced IL-8 mRNA expression, was evaluated. BRECs were treated with 10 μmol/l PKCζI-1 30 min before treatment of cells with TNF-α for 2 h. TNF-α induced a significant 14-fold increase in IL-8 transcripts, which was significantly reduced to 6-fold by PKCζI-1 (Fig. 6A). The effect of PKCζI-1 on TNF-α-induced NF-κB reporter activity was investigated in a 293/NF-κB/luc stable reporter cell line. Cells were



treated with 10 or 50 μ mol/l PKC ζ I-1 30 min before the addition of TNF- α for 6 h, and luciferase activity was determined in whole-cell lysates. The results show that PKC ζ I-1 significantly decreased TNF- α -induced NF- κ B-responsive luciferase expression in these cells (Fig. 6B). These observations suggest that pharmacological inhibition of PKC ζ reduces NF- κ B activation in response to TNF- α .

TNF-α-induced cell permeability requires PKCζ. PKCζ activation can occur downstream of the PI_3K pathway (38), which has been shown to be activated by TNF-α in vascular endothelial cells (39). Therefore, the effect of $PI_3K/PKCζ$ pathway inhibition on retinal endothelial cell permeability was investigated. BRECs were treated with 2 μmol/1 LY294002, a PI_3K inhibitor, 30 min before the treatment with TNF-α for 6 h. The increase in cell permeability induced by TNF-α was significantly inhibited by 56% in the presence of LY294002 (Fig. 7A). Next, BRECs were exposed to 10 μmol/1 PKCζI-1 or the myristoylated pseudosubstrate inhibitor of PKCζ (PKCζp; 250 nmol/1) 30 min prior to TNF-α addition. Both PKCζI-1 and PKCζp completely suppressed the increase in cell permeability induced by TNF-α (Fig. 7B and C). Because conventional

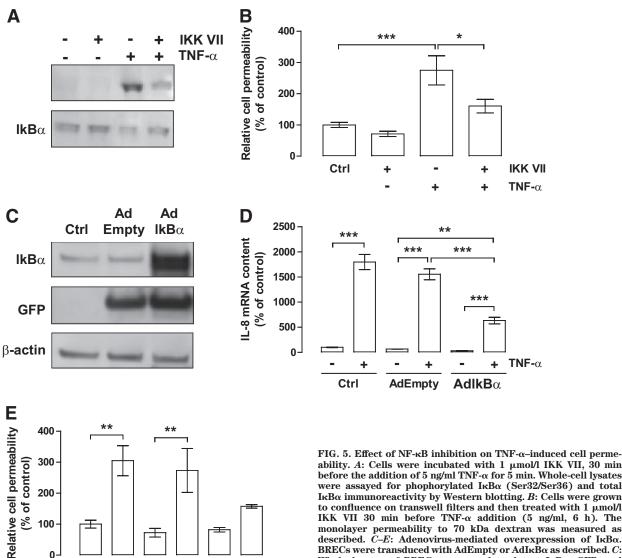
PKC isoforms contribute to the VEGF-induced permeability in retinal endothelial cells (23), the contributions of conventional and novel PKC isoforms for TNF-αinduced cell permeability were also evaluated. BRECs were treated with 5 μmol/l BIM I (inhibitor of conventional and novel PKC isoforms) or with 30 nmol/l PKCB inhibitor 30 min before TNF-α treatment. These inhibitors had no effect on TNF-α-induced cell permeability (supplementary Fig. 2A and B, available in an online appendix). These results suggest that PKCζ, but not conventional and novel PKC isoforms, mediate TNF-αinduced retinal endothelial cell permeability. The effect of PKCζ inhibition on tight junction protein content in TNF-α-treated BRECs was also evaluated. The protein content of ZO-1, claudin-5, and occludin was not affected by treatment with PKC I-1. However, the decrease in ZO-1 and claudin-5 protein levels induced by TNF- α was effectively prevented by PKC ζ I-1 (Fig. 7E and F). Surprisingly, in the presence of PKC ζ I-1, TNF- α still caused an increase in occludin protein content, which actually was greater than the effect of TNF- α alone (Fig. 7G).

periments. Scale bar, 25 µm.

independent experiments and are expressed as the relative amount compared with control. *P < 0.05, **P < 0.01, ***P < 0.001, significantly different from control; #P < 0.001

0.05, ##P < 0.01, ###P < 0.001, significantly different from TNF- α , as determined by ANOVA followed by Bonferroni post hoc test. D: Cells were immunolabeled for ZO-1, claudin-5, and occludin and 10 confocal Z-stacks were taken through 2.56 μ m and projected into one image. These results are representative of four independent ex-

Occludin



ability, A: Cells were incubated with 1 µmol/l IKK VII, 30 min before the addition of 5 ng/ml TNF-α for 5 min. Whole-cell lysates were assayed for phophorylated IκBα (Ser32/Ser36) and total IκBα immunoreactivity by Western blotting. B: Cells were grown to confluence on transwell filters and then treated with 1 µmol/l IKK VII 30 min before TNF-α addition (5 ng/ml, 6 h). The monolayer permeability to 70 kDa dextran was measured as described. \hat{C} -E: Adenovirus-mediated overexpression of IkB α . BRECs were transduced with AdEmpty or AdI κ B α as described. C: Whole lysates of BRECs were used to detect IκBα, GFP, and β-actin (loading control) by Western blotting. D: After 28 h of adenovirus transduction, cells were exposed to 5 ng/ml TNF-α for 2 h. Total RNA was isolated, and the transcript levels of IL-8 were analyzed by qPCR. E: Cells were grown to confluence on transwell

filters and after 24 h of adenovirus transduction cells were treated with 5 ng/ml TNF- α for 6 h. The monolayer permeability to 70 kDa dextran was measured. The results represent the mean \pm SEM of at least three independent experiments and are expressed relative to control (Ctrl). *P < 0.05, **P < 0.01, ***P < 0.001, significantly different as determined by ANOVA followed by Bonferroni post hoc test.

AdlkB α

TNF-α

PKCζI-1 prevents TNF-α-induced BRB permeability in vivo. The effect of PKCζI-1 on TNF-α-induced BRB permeability was evaluated in vivo by the Evans blue assay. The injection of TNF- α in the vitreous induced the accumulation of Evans blue $(20.30 \pm 2.46 \,\mu l \, plasma/g/h)$ as compared with PBS-injected animals (10.25 \pm 2.48 μ l plasma/g/h; Fig. 8A). PKCζI-1 alone had no effect on the accumulation of Evans blue but completely prevented TNF- α -induced accumulation of the dye. To determine the effect of TNF-α injection and PKCζ inhibition on retinal vascular tight junction organization, retina whole flat mounts were immunolabeled for ZO-1 and occludin proteins after injection of the cytokine or cytokine with PKCζI-1. In PBS-injected rat eyes, ZO-1 and occludin immunoreactivities were intense and localized at the junctions of the cell membranes of endothelial cells in retinal vessels. In TNF-α-injected eyes, changes in ZO-1 immunostaining were particularly evident, which became markedly reduced and intermittently absent from the cell

Ctrl

AdEmpty

borders. These alterations were prevented by cotreatment with PKC I-1 (Fig. 8B). Consistent with the results obtained with BRECs in culture, the occludin content was not reduced with TNF- α but displayed regions with disorganized cell-border labeling that were reversed by PKCζI-1 treatment.

DISCUSSION

Elevated levels of proinflammatory cytokines IL-1\beta and TNF-α have been detected in the vitreous of diabetic patients with retinopathy (10,11) and in diabetic rat retinas correlated with increased vascular permeability (12,40). However, the mechanisms underlying the effects of IL- 1β and TNF- α on retinal microvascular barrier have not been addressed. The results presented herein demonstrate that IL-1 β and TNF- α increase retinal endothelial cell permeability and that TNF- α acts through PKC ζ /NF- κ B to reduce the expression and alter the distribution of the tight

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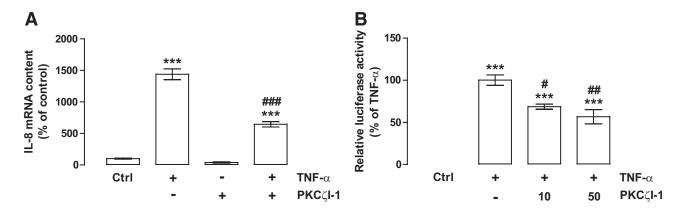


FIG. 6. Inhibition of PKC ζ inhibits NF- κ B activation by TNF- α . A: BRECs were treated with 10 μ mol/l PKC ζ I-1, a PKC ζ inhibitor, 30 min before the addition of 5 ng/ml TNF- α for 2 h. Total RNA was isolated, and the transcript levels of IL-8 were analyzed by qPCR. The results represent the mean \pm SEM of six independent experiments and are expressed relative to control (Ctrl). B: 293T-NF-kB-luc cells, with a κ B-dependent luciferase reporter gene, were treated with 10 or 50 μ mol/l PKC ζ I-1 30 min prior to the addition of TNF- α for 6 h. Cells were harvested, and luciferase activity was determined in whole-cell lysates. The results represent the mean \pm SEM of four independent experiments and are expressed relative to TNF- α . ***P < 0.001, significantly different from control; *P < 0.05, *P < 0.01, *P < 0.001, significantly different from TNF- α , as determined by ANOVA followed by Bonferroni post hoc test.

junction proteins claudin-5 and ZO-1. Moreover, glucocorticoid treatment completely prevented the TNF- α -induced increase in retinal endothelial cell permeability through both transactivation of the glucocorticoid receptor and transrepression of the NF- κ B signaling pathway. Pharmacological inhibition of PKC ζ reduced NF- κ B activation and

prevented TNF- α -induced retinal endothelial cell permeability both in vitro and in vivo.

The increase in retinal vascular permeability is associated with changes in the expression or distribution of the tight junction proteins. The content of occludin and ZO-1 decreases in the retinas of diabetic animals (18,33,41). In

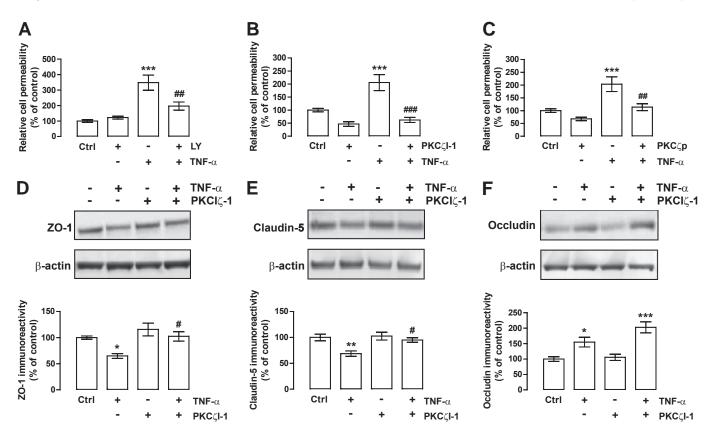


FIG. 7. TNF- α increases cell permeability in a PKC ζ -dependent manner. BRECs were grown to confluence on transwell filters and then treated with (A) 2 µmol/l LY294002 (LY), (B) 10 µmol/l PKC ζ I-1, and (C) 250 nmol/l PKC ζ pseudosubstrate inhibitor (PKC ζ p). The monolayer permeability to 70 kDa dextran was measured as described. Results represent the mean \pm SEM of at least five experiments and are expressed relative to Ctrl. D-F: PKC ζ I-1 prevents tight junction complex disruption induced by TNF- α . Whole-cell extracts were assayed for (D) ZO-1, (E) claudin-5, and (F) occludin immunoreactivity by Western blotting. Representative Western blots for each tight junction protein and β -actin (loading control) are presented above each respective graph. The results are normalized to β -actin and represent the mean \pm SEM of at least eight independent experiments and are expressed as the relative amount compared with control (Ctrl). All inhibitors were added to the cell culture medium 30 min before TNF- α addition (5 ng/ml, 6 h). *P < 0.05, **P < 0.01, ***P < 0.001, significantly different from control; #P < 0.05, ##P < 0.01, significantly different form TNF- α , as determined by ANOVA followed by Bonferroni post hoc test.

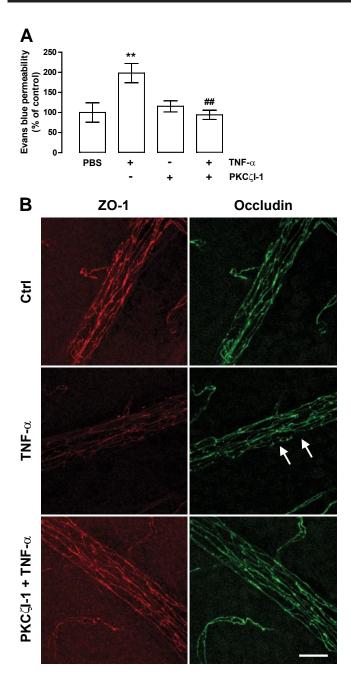


FIG. 8. PKCζI-1 prevents TNF- α -induced retinal vascular permeability in vivo. Animals' eyes were injected with PBS with 0.1% BSA, TNF- α (10 ng); PKCζI-1 (280 ng); or with both PKCζI-1 and TNF- α . A: Evans blue leakage was evaluated 24 h after intravitreous injections. The results represent the mean \pm SEM (n=7–8 animals per group) and are expressed relative to control (Ctrl; PBS-injected eyes). **P < 0.01, significantly different from control; ##P < 0.01, significantly different from TNF- α , as determined by ANOVA followed by Bonferroni post hoc test. B: PKCζI-1 prevents the alterations in tight junction proteins induced by TNF- α in vivo. Whole retinas were immunolabeled for ZO-1 and occludin 4 h after injection. Images were obtained on a Leica TCS SP2 AOBS confocal microscope and are presented as a maximum projection. Arrows indicate loss and/or discontinuous cell border staining. Scale bar, 25 μ m. (A high-quality digital representation of this figure is available in the online issue.)

BRECs, TNF- α downregulated ZO-1 and claudin-5 expression and decreased the junctional localization of both proteins, which was associated with an increase in cell permeability. These studies are consistent with previous publications demonstrating reduced claudin-5 gene expression after TNF- α treatment in brain capillary endothe-

lial cells (42,43). However, in these same studies, a decrease in occludin was observed after TNF- α treatment, which contrast the increase in occludin observed in the present study. Further, TNF- α reduced occludin promoter activity in a human intestinal cell line (44). The cause for this difference is unclear but may relate to the use of cell lines versus primary culture or differences in epithelial and endothelial cell types. Regardless, TNF- α consistently reduces both claudin-5 and ZO-1 expression, two tight junction proteins essential for barrier properties.

The effect of TNF- α and VEGF on retinal endothelial permeability is distinct. Occludin becomes phosphorylated on multiple sites after VEGF treatment in a conventional PKC-dependent manner (23) and Ser490 has recently been identified as a phosphorylation site necessary for VEGF-induced permeability (24,45). TNF- α did not lead to an increase in occludin phosphorylation on Ser490 (data not shown) but rather led to an increase in occludin expression. Further, inhibition of conventional or novel PKC isoforms did not prevent TNF- α -induced permeability. These data demonstrate that VEGF and TNF- α alter retinal endothelial barrier properties by distinct molecular mechanisms.

The data presented here suggest that glucocorticoid receptor activation inhibits TNF-α-induced cell permeability through both transactivation and transrepression mechanisms. Inhibition of IKK and adenovirus-mediated overexpression of IκBα blocked the increase in BREC permeability induced by TNF- α treatment. Together, these data demonstrate that TNF-α-induced retinal endothelial cell permeability is mediated, at least in part, by NF-kB. The mechanisms by which NF-kB may regulate cell permeability and the tight junction complex are largely unknown. NF-κB putative binding sites and several E-box sequences were identified within the claudin-5 promoter sequence (43). Recent studies demonstrated that NF-κB induces the expression of Snail and Slug transcription factors (46,47), which repress E-cadherin, occludin, and claudin family members gene expression by binding to specific E-box sequences during epithelial-mesenchymal transition (48–50) These reports suggest the possibility that a similar regulation by transcriptional repressors might also play a role in claudin-5 and ZO-1 expression in retinal endothelial cells.

PKC ζ has been shown to contribute a critical and selective role in the regulation of NF-κB. In PKC ζ -deficient mice, NF-κB transcriptional activity as well as the phosphorylation of p65 in response to TNF-α is severely impaired (37). In endothelial cells, the transcriptional activity of NF-κB is dependent on the phosphorylation of the p65 subunit by PKC ζ (51). The observations that TNF-α-induced increase in IL-8 expression and NF-κB-dependent luciferase reporter expression are inhibited by PKC ζ I-1 further support the hypothesis that PKC ζ is important for NF-κB transcriptional activity.

In endothelial cells, PKC ζ has been shown to be activated by TNF- α (51–53). The mechanism of activation of PKC ζ remains to be fully clarified, but it has been shown to be an important downstream target of PI₃K (38). TNF- α stimulates PI₃K in endothelial cells through endothelial/epithelial tyrosine kinase induced activation of VEGFR2 (39). Interestingly, this is only a partial activation of VEGFR2 that fails to activate the classical PKC isoforms necessary for VEGF-stimulated permeability (23,54). PI₃K inhibition and inhibition of its downstream target PKC ζ (38) blocked TNF- α -induced cell permeability, whereas

classical PKC inhibitors failed. Targeting PKC ζ reduced NF- κ B activation and most likely inhibited additional signaling pathways that contribute to the regulation of the tight junction complex and endothelial permeability induced by TNF- α . PKC ζ I-1 is also effective at blocking VEGF-induced permeability (manuscript in preparation). Then, although VEGF and TNF- α alter the tight junction complex by distinct mechanisms, inhibition of PKC ζ is a common target for blocking both VEGF- and TNF- α -induced cell permeability. Therefore, targeting PKC ζ may provide a specific therapeutic option for the prevention of vascular permeability in retinal diseases with elevated TNF- α and VEGF, such as diabetic retinopathy.

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