Thrombin Inhibits Intercellular Calcium Wave Propagation in Corneal Endothelial Cells by Modulation of Hemichannels and Gap Junctions

Catbeleyne D’bondt,1 Raf Ponsaerts,1 Sangly P. Srinivas,2 Johan Vereecke,1 and Bernard Himpens1

PURPOSE. Thrombin, a serine protease, breaks down the barrier integrity of corneal endothelial cells by phosphorylation of the regulatory light chain of myosin II (myosin light chain; MLC), which induces contractility of the actin cytoskeleton. This study was undertaken to investigate the effect of thrombin on gap junctional (GJIC) and paracrine (PIC) intercellular communication in cultured bovine corneal endothelial cells (BCECs).

METHODS. An intercellular Ca2+ wave, a form of cell–cell communication, was elicited by applying a mechanical stimulus to a single cell in a confluent monolayer. Changes in [Ca2+]i were imaged by fluorescence microscopy with a fluorescent calcium indicator, and the images were used to calculate the area reached by the Ca2+ wave (active area). GJIC was assessed by fluorescence recovery after photobleaching (FRAP). Activity of hemichannels was assayed by lucifer yellow (LY) uptake and also by adenosine triphosphate (ATP) release by using the luciferin–luciferase technique.

RESULTS. RT-PCR showed transcripts for PAR-1 and -2 receptors, but not for PAR-4 receptors. Immunocytochemistry showed thrombin-sensitive PAR receptors as well as trypsin-sensitive PAR-2 receptors. Both thrombin and the selective PAR-1 agonist TRAP-6 reduced the active area of the Ca2+ wave. These agents also reduced the fluorescence recovery in FRAP experiments. The effect of thrombin on the Ca2+ wave was inhibited by a peptide antagonist of PAR-1, but not by a PAR-4 antagonist. Pretreatment with ML-7 (an MLCK inhibitor), Y-27632 (a Rho kinase inhibitor) or chelerythrine (a PKC inhibitor) prevented the effect of thrombin on the Ca2+ wave. Activation of PAR-1 did not affect the Ca2+ wave propagation in cells pretreated with Gap26, which blocks hemichannels. However, PAR-1 activation decreased the active area in cells pretreated with Gap27, which inhibits gap junctions. Thrombin abolished enhancement of the Ca2+ wave propagation by ARL-67156 (inhibitor of ecto-ATPases). The effect of the PAR-1 agonists on the Ca2+ wave was not detectable in cells pretreated with exogenous apyrase.

CONCLUSIONS. Thrombin inhibits intercellular Ca2+ wave propagation in BCECs. This effect is due to activation of PAR-1 receptors and involves MLC phosphorylation by MLCK, PKC- and Rho kinase–sensitive pathways. Thrombin mainly inhibits the ATP-mediated PIC pathway, and also reduces GJIC to a lesser extent. (Invest Ophthalmol Vis Sci. 2007;48:120–133)

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The corneal endothelium is a nonregenerative monolayer1 at the posterior surface of the cornea. Its main physiological role is maintenance of the transparency of the cornea, and its decomposition leads to loss of visual acuity and is a common indication for corneal transplantation.2 For maintaining corneal transparency, the endothelium carries out hydraulic control of the corneal stroma by bringing about an active fluid transport into the anterior chamber, so as to balance the fluid leak into the stroma.3,4 To sustain these functions in the absence of regeneration, the corneal endothelium must possess several defense mechanisms, to withstand extracellular stresses. Intercellular communication (IC), which promotes a coordinated response from the monolayer, possibly helps in maintaining the resilience of the endothelium against extracellular stresses, such as mechanical stress during intraocular surgery or exposure to inflammatory mediators during immune rejection or uveitis.5,6

Different groups6–16 including our own,17–19 have been investigating the mechanisms and roles of IC in the corneal endothelium and epithelium. As in many other nonexcitable cells, the corneal endothelial cells exhibit two distinct modes of IC:17,18 gap junctional IC (GJIC) and paracrine IC (PIC). In GJIC, a direct exchange of signaling molecules via gap junctions enables IC. Gap junctions are formed by the docking of two connexin (Cx) hemichannels (also called connexons) contributed by two apposing cells.20 In contrast to GJIC, PIC does not require cell–cell apposition, as it involves release of one or more diffusible signaling molecules that bring about IC by acting on the neighboring (NB) cells. A candidate paracrine factor in the propagation of intercellular Ca2+ waves that has been well investigated in several cell types,11,21,22 including corneal endothelium,15,16 is adenosine triphosphate (ATP). Furthermore, it has been demonstrated in many cell types, including in the corneal endothelium, that the nucleotide is released through hemichannels17,23,24 and that it evokes Ca2+ wave propagation through its action on purinergic receptors.18,19,21

As reported in previous studies, corneal endothelial cells express several connexin subtypes, including the widespread Cx43,8,10,13,19 Although we are beginning to understand the expression diversity of connexins and the mechanisms underlying PIC and GJIC, not much is known about factors that influence GJIC or PIC in corneal endothelial cells. In general, GJIC is influenced by gating and also by the number of gap junctional channels.20 The latter is influenced by (1) the level of expression of connexins; (2) the assembly of connexins into

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Connexons; (3) the trafficking of connexons, resulting in the formation of hemichannels in the plasma membrane; (4) formation of gap junction channels and their accretion into plaques; and (5) internalization and degradation of hemichannels and gap junction channels.\(^5\) Trafficking of connexons and the gating of gap junction channels are known to be influenced by several factors,\(^2\) including the connexin phosphorylation status.\(^2\) Gating of the gap junction channel is also affected by several physiological factors,\(^2\) such as intracellular acidification, \([\text{Ca}^{2+}],\) and transjunctional membrane voltage. Results in several recent studies have implied that the actin cytoskeleton has an active role in the regulation of GJIC.\(^2\) \(^{28,29}\) It has been shown, for example, that GJIC is affected by microinjection of cells with actin antibodies\(^3\) or exposure to agents that disrupt actin or the microtubule cytoskeleton.\(^2\) \(^{29,34}\)

Because GJIC requires cell–cell apposition, it is very plausible that loss of intercellular tethering forces will induce a breakdown of the gap junctions, as reported widely for tight junctions.\(^35\) \(^{40}\) Recent studies show that the forces affecting intercellular junctions are influenced by contractility of the actin cytoskeleton,\(^36,39\) as they oppose the intercellular tethering forces established by the adherens junctions. The contractility of the actin cytoskeleton is induced by actomyosin interaction, which is under the control of phosphorylation of the regulatory light chain of myosin II (also called myosin light chain, MLC). The level of MLC phosphorylation is dynamically regulated by myosin light chain kinase (MLCK), PKA, PKC, and Rho kinase, secondary to activation of several G-protein-coupled receptors (GPCRs).\(^41,42\)

Much less is known about the regulation of hemichannels and PIC.\(^2\) \(^{24,25,43,45}\) In a recent study, the investigators suggested that ATP release through hemichannels is associated with depolymerization of the actin cytoskeleton.\(^45\) In the present study, the main objective was to investigate whether the altered MLC phosphorylation affects GJIC and PIC in corneal endothelial cells. Because the serine protease thrombin is well known to induce MLC phosphorylation and increase contractility of the actin cytoskeleton,\(^35,46,47\) we investigated the effects of this protease on GJIC and PIC in bovine corneal endothelial cells (BCECs) in the paradigm of intercellular propagation of the \([\text{Ca}^{2+}]\) wave elicited by a mechanical stimulus.\(^17,18\) Our results show that thrombin reduces \([\text{Ca}^{2+}]\) wave propagation, mainly by inhibiting a hemichannel-mediated ATP release-dependent PIC pathway and also by inhibiting GJIC, although to a lesser extent.

### Materials and Methods

#### Chemicals

Fluo-4 AM (F14217) and 6 carboxyfluorescein diacetate (C1362) were obtained from Invitrogen-Molecular Probes (Eugene, OR); chelerythrine (C-2932), ML7 (T-2764), thrombin (T-6468), apyrase VI (A6410), apyrase VII (A6355), ARL67156 (6-V,N-diethyl-β;γ-dibromomethylene-

#### Table 1. Gene-Specific Primer Pairs Used for RT-PCR of Bovine PAR-1, -2, and -4 and β-actin

<table>
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<th>Gene</th>
<th>Accession No.</th>
<th>Primer</th>
<th>Sequence (5' to 3')</th>
<th>Size (bp)</th>
</tr>
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<tr>
<td>PAR2</td>
<td>XM_604897</td>
<td>Forward</td>
<td>CACGTGAGTATTCTGTTG</td>
<td>360</td>
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<tr>
<td>PAR4</td>
<td>XM_585942</td>
<td>Forward</td>
<td>GTGTATAGCTGACAGATTGCT</td>
<td>248</td>
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<tr>
<td>β-Actin</td>
<td>NM_173979</td>
<td>Forward</td>
<td>GCTTTAGACCTGCTCTTCCT</td>
<td>178</td>
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</tbody>
</table>

#### RT-PCR Assay for Expression of PAR Receptors

Total RNA was extracted from BCECs (SV Total RNA Isolation System kit Z3100; Promega, Madison, WI). First-strand cDNA synthesis was performed (SuperScript II Reverse Transcriptase for RT-PCR, 11904-018; Invitrogen-Gibco) with random hexamers. To amplify cDNA for PAR receptors, PCR was performed for 35 cycles (94°C for 45 seconds; 55°C for 30 seconds; and 72°C for 90 seconds) in the final cycle) with Taq DNA polymerase (18038-042; Invitrogen-Gibco) and the gene-specific primers listed in Table 1. Amplified cDNA was visualized with ethidium bromide after performing DNA gel electrophoresis (2% agarose) in the presence of a 50-bp DNA ladder (G4521; Promega).

#### Expression of PAR Receptors by Immunocytochemistry

Confluent monolayers were fixed for 10 minutes with 4% paraformaldehyde in Dulbecco's PBS (14190-091; Invitrogen-Gibco). After fixation, the BCECs were incubated for 30 minutes in a freshly made PBS.
solution containing 3% BSA and 10% goat serum. They were rinsed with PBS after each treatment. After blocking, the cells were incubated overnight at 4°C with a rabbit polyclonal antibody directed against the thrombin receptor (ab13598; Abcam PLC, Cambridge, UK) or PAR-2 (ab13097; Abcam PLC). The secondary antibody was goat anti-rabbit IgG labeled with FITC (ab6717; Abcam PLC). The negative controls were obtained in identical conditions, but the cells were not exposed to the primary antibody. Finally, the cells were visualized with a confocal microscope (LSM510; Carl Zeiss Meditec, GmbH, Jena, Germany). The 488-nm line of the Argon laser was used for excitation. Emission was recorded via a long-pass filter at 505 nm.

**Fluorescence Recovery after Photobleaching**

We have described the FRAP (fluorescence recovery after photobleaching) protocol in BCECs previously. Briefly, cells were loaded with the Ca\(^{2+}\)-insensitive dye 6-carboxyfluorescein diacetate (10 μM) for 5 minutes at room temperature, and fluorescence recovery was measured with the confocal microscope. For recording fluorescence recovery, the dye was excited at 488 nm and its emission was recorded at 530 nm. A neutral-density filter was also used to minimize photobleaching. Before photobleaching, a polygon was drawn around a cell chosen for bleaching, and two images were obtained. The chosen cell was then exposed to 50 scans with the laser at 95% intensity, and fluorescence recovery was measured with the confocal microscope. For recording fluorescence recovery, the dye was excited at 488 nm and its emission was recorded at 530 nm. A neutral-density filter was also used to minimize photobleaching. Before photobleaching, a polygon was drawn around a cell chosen for bleaching, and two images were obtained. The chosen cell was then exposed to 50 scans with the laser at 95% intensity, and the recovery of fluorescence in the bleached cell was measured every 10 seconds over a period of 5 minutes. The decrease of fluorescence in a square region of interest widely distant from the bleached cell was measured as a reference for correction for bleaching due to the excitation light used for fluorescence detection while recording fluorescence recovery. After correction for background bleaching, recovery of fluorescence in the bleached cell at 5 minutes was compared with that of the prebleach scan, and the percentage recovery was calculated.

**Mechanical Stimulation for Inducing a Ca\(^{2+}\) Wave**

The protocol for point mechanical stimulation of a single cell has been described in our previous study. It consists of bringing about an acute deformation of the chosen cell by briefly touching less than 1% of its cell membrane with a glass micropipette (tip diameter <1 μm) coupled to a piezoelectric crystal (Piezo device P-280, Amplifier-E463; PI Polytech, Karlsruhe, Germany) mounted on a micromanipulator.

**Measurement of [Ca\(^{2+}\)]**

The Ca\(^{2+}\) wave propagation was assayed by imaging [Ca\(^{2+}\)], as described previously. Cells were loaded with the Ca\(^{2+}\)-sensitive dye, Fluo-4 AM (10 μM; Invitrogen-Molecular Probes) for 30 minutes at 37°C. The dye was excited at 488 nm, and its fluorescence emission was collected at 530 nm. Spatial changes in [Ca\(^{2+}\)] were measured with the confocal microscope (LSM510; Carl Zeiss Meditec, GmbH) equipped with a 10× objective (Air, 1.2 NA), but in experiments with ARL-67156, a 10× objective (Air, 0.3 NA) was used. Images were collected and stored on a personal computer. Polygonal regions of interest (ROIs) were drawn to define the borders of each cell. A cell (called the mechanically stimulated (MS) cell) was selected for point of mechanical stimulation. The NB cells immediately surrounding the MS cell were defined as NB cell layer 1 (NB1), the ones immediately surrounding the MS cell were defined as NB cell layer 2 (NB2), and so on. Fluorescence was averaged over the area of each ROI. Normalized fluorescence (NF) was then obtained by dividing the fluorescence by the average fluorescence before point mechanical stimulation. Intercellular propagation of the Ca\(^{2+}\) wave was characterized by maximum normalized fluorescence (NF), and percentage of responsive cells (%RC), as well as the total surface area of responsive cells (active area, AA) with NF ≥ 1.1.

Response to exogenously applied ATP after pretreatment with thrombin was also characterized by using Fluo-4 as just described, but...
the normalized fluorescence averaged over all cells in the field of view was used as a measure of \([\text{Ca}^{2+}]_i\).

### Measurement of ATP Release

ATP release, resulting from mechanical stimulation on the confocal microscope, was followed-up using the luciferin-luciferase bioluminescence protocol. One hundred microliters was sampled out from the 500 \(\mu\)l bathing solution covering the cells and taken to a custom-built photon-counting setup to measure the luminescence, as described previously. Briefly, photons emitted as a result of the oxidation of luciferin in the presence of ATP and O\(_2\) and catalyzed by luciferase were detected by a photon-counting photomultiplier tube (II7360-01; Hamamatsu Photonics, Hamamatsu City, Japan). Voltage pulses from the photomultiplier module were counted with a high-speed counter (PCI-6602; National Instruments, Austin, Texas, USA). Dark count of the photomultiplier tube was less than 80 counts/second.

### Lucifer Yellow Uptake Assay

Cells grown to confluence on chamber slides were incubated in a Ca\(^{2+}\)-rich PBS containing the drug of interest for 30 minutes. Cells were then exposed to PBS containing 2 mM EGTA and 2.5% Lucifer yellow (LY) for 5 minutes in the continued presence of the drug. After a brief wash with Ca\(^{2+}\)-containing PBS, LY fluorescence (\(\lambda_{ex} = 488 \text{ nm}; \lambda_{em} = 530 \text{ nm}\)) from cells was imaged with the confocal microscope (LSM510; Carl Zeiss Meditec, GmbH).

### Data Analysis

One-way ANOVA was used to compare the mean results of different treatments, whereas unpaired tests were used to compare the results of experiments with a single treatment and a single control (Prism 4.0 for Windows; GraphPad Software Inc., San Diego, CA). \(p < 0.05\) was considered statistically significant. Data are expressed as the mean ± SEM. \(N\) is the number of independent experiments (the number of MS cells), whereas \(n\) represents the total number of responsive cells.

### RESULTS

#### Effects of Thrombin on Ca\(^{2+}\) Wave Propagation

In control conditions, the MS cell showed a transient \([\text{Ca}^{2+}]_i\) increase that originated at the point of stimulation and spread out to the NB cells in a wavelike manner, as shown by the fluorescence images in Figure 1A. Ca\(^{2+}\) transients were observed up to approximately four to six cell layers away from the MS cell. The line graph (Fig. 1A, right) shows the time course of the Ca\(^{2+}\) transients (represented as NF values) in the MS cell and in the NB cell layers one to five. The figure also shows that the normalized fluorescence decreased, whereas the time delay for the onset of the rise in \([\text{Ca}^{2+}]_i\) was lengthened, with increasing distance from the MS cell. Cells subjected to mechanical stimulation after incubation with thrombin (2 U/ml) for 5 minutes, also show intercellular propagation of the Ca\(^{2+}\) wave, but the spread of the wave is limited to only approximately two to four cell layers (Fig. 1B). A quantitative summary of similar experiments is provided in Figure 2 and Table 2. The normalized fluorescence and the percentage of responsive cells (%RC) decreases as a function of the distance of the cell layer from the MS cell. In the presence of thrombin, this spatial decline is accelerated. The AA, a measure of efficacy of the Ca\(^{2+}\) wave, was \(51,400 \pm 1,300 \mu m^2 (N = 140)\) under control conditions and was reduced by \(\approx 70\%\) to \(15,000 \pm 1,000 \mu m^2 (N = 140)\) in the presence of thrombin in contemporaneous experiments (Fig. 2C). Therefore, the results in Figures 1 and 2 demonstrate that thrombin significantly inhibits the intercellular propagation of Ca\(^{2+}\) waves in BCECs.

Because the effect of thrombin on the AA after mechanical stimulation could be due to several factors other than propagation of the Ca\(^{2+}\) wave (e.g., thrombin could have emptied Ca\(^{2+}\) stores or have an effect on ATP-mediated IP3 release), we also measured the Ca\(^{2+}\) rise in response to exogenous ATP (10 \(\mu\)M) in cells pretreated with thrombin and in untreated cells. The peak increase in \([\text{Ca}^{2+}]_i\), in thrombin-treated cells on exposure to ATP was not significantly different from that in untreated cells (\(N = 25, P < 0.05\)), suggesting that the effect of thrombin is not due to an effect on P2 receptor-mediated signaling.

#### Identity of Receptors Involved in the Effects of Thrombin

It is known that thrombin activates GPCRs—more specifically, proteinase-activated receptors (PAR family of receptors). To date, four PAR receptors have been described, and most cell types express multiple PAR receptors. To find the identity

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**Figure 2.** Quantification of the spread of the Ca\(^{2+}\) wave in control conditions and in the presence of thrombin. (A) Average NF in the MS cell and in NB cell layers 1 to 5 in control conditions and in the presence of thrombin (2 U/ml for 5 minutes). (B) Percentage of responsive cells (%RC) in MS and NB1 to NB5 in control conditions and in the presence of thrombin. (C) AA in the presence of thrombin versus control conditions. Data represent average values from 140 contemporaneous experiments. *\(P < 0.001\) for thrombin versus control.
of PAR receptors expressed in BCECs, we performed RT-PCR and immunocytochemistry. RT-PCR demonstrated expression of mRNA for PAR-1 and -2 in BCECs, but not PAR-4 (Fig. 3A). Immunocytochemistry was performed to confirm the expression using an antithrombin receptor antibody as well as a PAR-2–specific antibody. As shown in Figure 3, both thrombin-sensitive receptors (Fig. 3C) and PAR-2 receptors (Fig. 3D) are very well expressed on the cell surface. In consistence with the expression of PAR-1 and -2 receptors and their coupling to $G_{\alpha/11}$, acute exposure to thrombin as well as trypsin (1 $M$) produced a transient increase in $[Ca^{2+}]_{i}$ (NF $= 2.08 \pm 0.06$; $N = 25$; NF $= 2.43 \pm 0.12$; $N = 10$, respectively).

We also investigated the identity of the PAR receptors involved in the inhibitory effect of thrombin on the intercellular propagation of the $Ca^{2+}$ wave, using PAR agonists and antagonists (Fig. 4). Pretreatment with TRAP-6 (thrombin receptor-activator peptide-6 with the sequence SFLLRN; 10 $\mu M$ for 30 minutes), a selective PAR-1 activating peptide, also reduced the AA from 62,000 $\pm 2,800 \mu m^2$ ($N = 50$) in control conditions to 26,000 $\pm 3,200 \mu m^2$ ($P < 0.001; N = 50$; Fig. 4A). In accordance with this PAR-1–mediated effect, thrombin failed to induce reduction in the AA in the presence of a PAR-1 antagonist (a mimetic peptide with the sequence 3-Mpr-FCha-Cha-RKPNDK-NH$_2$; 10 $\mu M$ for 30 minutes; $P < 0.001; N = 20$; Fig. 4B). We also investigated possible involvement of PAR-4 receptors using the indazole derivative YD-3, a selective PAR-4 antagonist. On pretreatment with YD-3 (50 $\mu M$ for 30 minutes), thrombin caused a significant reduction of the AA ($P < 0.001; N = 20$) similar to the reduction caused by thrombin in the absence of the antagonist. Therefore, we conclude that PAR-4 receptors do not contribute to the observed effect of thrombin on the $Ca^{2+}$ wave propagation.

### Table 2. Average Maximum NF, Percentage Responsive Cells, and AA in the MS and NB Layers during Mechanical Stimulation in Control Conditions and after Treatment with Thrombin

<table>
<thead>
<tr>
<th></th>
<th>MS</th>
<th>NB1</th>
<th>NB2</th>
<th>NB3</th>
<th>NB4</th>
<th>NB5</th>
<th>AA ($\mu m^2$)</th>
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<tr>
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<td></td>
<td></td>
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<td>51,366 $\pm$ 1,294</td>
</tr>
<tr>
<td>NF $\pm$ SEM</td>
<td>2.61 $\pm$ 0.09</td>
<td>3.20 $\pm$ 0.04</td>
<td>2.64 $\pm$ 0.03</td>
<td>2.11 $\pm$ 0.03</td>
<td>1.79 $\pm$ 0.03</td>
<td>1.54 $\pm$ 0.03</td>
<td></td>
</tr>
<tr>
<td>% RC</td>
<td>100</td>
<td>99</td>
<td>94</td>
<td>76</td>
<td>46</td>
<td>33</td>
<td></td>
</tr>
<tr>
<td>n</td>
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<td>340</td>
<td>654</td>
<td>834</td>
<td>646</td>
<td>423</td>
<td></td>
</tr>
<tr>
<td>Thrombin</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td>15,013 $\pm$ 993</td>
</tr>
<tr>
<td>NF $\pm$ SEM</td>
<td>2.27 $\pm$ 0.09*</td>
<td>2.20 $\pm$ 0.04*</td>
<td>1.85 $\pm$ 0.03*</td>
<td>1.55 $\pm$ 0.03*</td>
<td>1.37 $\pm$ 0.02*</td>
<td>0.00 $\pm$ 0.00*</td>
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<tr>
<td>% RC</td>
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<td>568</td>
<td>298</td>
<td>174</td>
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Data were collected during mechanical stimulation in control conditions and in cells treated with thrombin (2 U/mL for 5 min).

* $P < 0.05$ versus control.
potential contribution of other thrombin-sensitive receptors or proteolytic effects on proteins other than PAR-1, we investigated the Ca\(^{2+}\) wave propagation after treating the cells with a combination of the PAR-1 and -4 antagonists. Under these conditions, thrombin failed to produce a decrease in the AA (Fig. 4B), excluding a significant contribution of other receptors to the inhibitory effect of thrombin on the Ca\(^{2+}\) wave. Exposure to the PAR-2 agonist trypsin (1 \(\mu\)M for 5 minutes) had no significant influence on Ca\(^{2+}\) wave propagation (AA 55,100 \(\pm\) 1,700 \(\mu\)m\(^2\) in the presence of trypsin versus 59,400 \(\pm\) 4,500 \(\mu\)m\(^2\) in the control (\(N = 25\)). Based on these observations, we conclude that the effect of thrombin on the Ca\(^{2+}\) wave propagation in BCECs is mediated through PAR-1 receptors.

**Signal Transduction Underlying the Effects of Thrombin**

It has been demonstrated that exposure of BCECs to thrombin increases MLC phosphorylation, which enhances contractility of the actin cytoskeleton and breaks down the barrier integrity. MLC phosphorylation is driven by MLCK, whereas its dephosphorylation is brought about by myosin light chain phosphatase (MLCP). MLCK can be activated by thrombin, because PAR-1 is coupled to G\(_{\alphaq/11}\), which results in Ca\(^{2+}\) release and hence Ca\(^{2+}\)-calmodulin binding to the kinase. PAR-1 receptors also activate PKC via G\(_{\alphaq/11}\), whereas they activate Rho kinase via G\(_{\alpha12/13}\); these two pathways contribute toward inactivation of MLCP. The transduction of the thrombin effect on MLC phosphorylation can thus occur via two pathways—namely, inhibition of the MLCP or stimulation of the MLCK. We, therefore, performed experiments to investigate whether thrombin exerts its effect on the Ca\(^{2+}\) wave propagation through the same transduction cascade that decreases the barrier integrity.

Pretreatment with MLCK inhibitor ML-7 (10 \(\mu\)M for 30 minutes) significantly suppressed the effect of thrombin or TRAP-6 on the propagation of the Ca\(^{2+}\) wave (\(P < 0.001; N = 25\); Fig. 5A). Pretreatment with the Rho kinase inhibitor Y-27632 (10 \(\mu\)M for 30 minutes) also significantly suppressed the effect of thrombin on the propagation of the Ca\(^{2+}\) wave (\(P < 0.05; N = 25\); Fig. 5C). The AA in the presence of a combination of Y-27632 plus chelerythrine, or in the presence of a combination of ML-7 + Y-27632 + chelerythrine, was also not significantly affected by thrombin (\(N = 25\); Fig. 5C). The absence of a significant effect of thrombin in the presence of the combination of the three protein kinase inhibitors provides evidence that the major pathways for the effect of thrombin on the AA are via MLCK, Rho kinase, and PKC.

**Effects of Connexin Mimetic Peptides.** In our previous studies, we used connexin mimetic peptides (Gap27 and Gap26) to distinguish relative contributions of GJIC and PIC to intercellular Ca\(^{2+}\) wave propagation after mechanical stimulation. We used the same peptides and followed identical treatment protocols to explore the effect of thrombin on the two mechanisms. Gap27 was shown to decrease significantly the propagation of the Ca\(^{2+}\) wave in BCECs via inhibition of gap junction channels. As shown in Figure 7A, pretreatment with Gap27 (300 \(\mu\)M for 30 minutes) led to a reduction in the AA (\(P < 0.001; N = 40\)). Exposure to thrombin led to a further decrease (\(P < 0.05; N = 40\); Fig. 7A). This observation suggests...
that thrombin has an effect on wave propagation that is insensitive to Gap27. The presence of an inactive control peptide did not influence the effect of thrombin. We obtained similar findings when TRAP-6 was used instead of thrombin (Fig. 7B). These results provide additional evidence that thrombin and TRAP-6 exert their major effects on \( \text{Ca}^{2+} \) wave propagation through inhibition of PIC.

The connexin-mimetic peptide Gap26, which has been demonstrated to block connexin hemichannels,\(^{17,19}\) significantly reduces the propagation of the \( \text{Ca}^{2+} \) wave in BCECs.\(^{17}\) As shown in Figure 7A, pretreatment with Gap26 (300 \( \mu \text{M} \) for 30 minutes) reduced the AA. Exposure to thrombin after pretreatment of the cells with Gap26 did not cause a further reduction of the AA. Similar effects were obtained using TRAP-6 (Fig. 7B). These findings provide evidence that thrombin exerts an effect on PIC that is due to inhibition of connexin hemichannels.

The presence of Gap27 and Gap26, did not affect the activity of thrombin as measured by thrombin time. Based on the thrombin recognition and cleavage motif\(^{52,55}\) and on the amino acid sequences of Gap27 and Gap26, proteolysis of the peptides is not expected.

### Effects of Stimulated or Inhibited Extracellular ATP Hydrolysis.

Because PIC in the endothelium is largely through ATP release,\(^{17,18}\) inhibition of ectonucleotidases, known to be expressed in the endothelium,\(^{18,48,49}\) results in strong enhancement of the wave propagation, as has been demonstrated previously in BCECs.\(^{17,18}\) As shown in Figure 8A, exposure to the ectonucleotidase inhibitor ARL67156 (100 \( \mu \text{M} \) for 30 minutes) increased the AA from 55,500 ± 3,700 to 258,300 ± 2,300 \( \mu \text{m}^2 \) \( \left( P < 0.001; N = 20 \right) \). Furthermore, the same figure shows that this large AA was reduced to 33,500 ± 8,000 \( \mu \text{m}^2 \) \( \left( P < 0.001; N = 20 \right) \) in the presence of thrombin. Thus, the very pronounced enhancement of the \( \text{Ca}^{2+} \) wave produced by
ARL-67156 was almost completely inhibited in the presence of thrombin (reduction of AA by 87%). These experiments demonstrate the influence of thrombin on ATP-mediated PIC. Specifically, these findings suggest that the effect of thrombin is due to a block of ATP release.

We therefore investigated the effect of thrombin after treatment of the cells with the nucleotidases apyrase VI (5 U/mL) and apyrase VII (5 U/mL), to inhibit PIC via hydrolysis of ATP and adenosine diphosphate (ADP). The propagation of the Ca\(^{2+}\) wave is markedly reduced in the presence of these nucleotidases.\(^{17,18}\) In the presence of the nucleotidases, neither thrombin nor TRAP-6 had a significant effect on the AA of BCECs pretreated with Gap26 or flufenamic acid in a condition where PIC was almost completely inhibited using ARL-67156. Under these conditions, thrombin failed to reduce the AA in cells pretreated with Gap26 (Fig. 9A; column 4 versus column 3). Furthermore, this AA was also not significantly different from the one obtained without pretreatment with Gap26 (Fig. 9A; column 4 versus column 2). These results demonstrate that thrombin exerts its effect via a Gap26-sensitive pathway, indicating the involvement of connexin hemichannels. Similar results were obtained with flufenamic acid (50 \(\mu\)M for 30 minutes; \(N = 35\)), which is also known to block hemichannels in BCECs\(^{17}\) and other cells, such as astrocytes,\(^{24}\) although the drug is not selective, because it is also known to inhibit Cl\(^{-}\) channels\(^{24}\) (Fig. 9B). These results indicate that pathways other than connexin hemichannels do not contribute to the effect of thrombin on the Ca\(^{2+}\) wave propagation.

Dye Uptake Experiments. The sensitivity of the wave propagation to Gap26 demonstrated in our experiments, and similar findings reported previously,\(^{17,50}\) indicated the involvement of hemichannels in ATP release. As shown in previous studies,\(^{17}\) connexin hemichannels are significantly permeable to the hydrophilic dye LY in Ca\(^{2+}\)-free solutions containing EGTA (2 mM). To investigate whether thrombin inhibits hemichannel-mediated PIC, we examined the effects of thrombin (2 U/mL) and TRAP-6 (10 \(\mu\)M; pretreatment for 30 minutes) on LY uptake. As shown in Figure 10A, exposure to PAR-1 agonist (thrombin or TRAP-6, \(N = 10\)) under Ca\(^{2+}\)-free medium (containing 2 mM EGTA and LY) led to complete inhibition of LY uptake. Furthermore, exposure to drugs which inhibit MLC phosphorylation overcame the effect of thrombin. Thus, when cells were pretreated with ML-7 (10 \(\mu\)M for 30 minutes, \(N = 4\)), Y-27632 (10 \(\mu\)M for 30 minutes, \(N = 3\)), or chelerythrine (5 \(\mu\)M for 30 minutes, \(N = 3\)), LY uptake was similar to control conditions (Figs. 10B, 10C, and 10D, respectively). These results indicate that PAR-1 activation blocked LY uptake rapidly, and therefore, we suggest that thrombin inhibits connexin hemichannels and hence the PIC pathway.

Effects on ATP Release. To study whether PAR-1 agonists inhibit ATP release on mechanical stimulation in BCECs, we measured extracellular ATP levels by the luciferin-luciferase technique. Thrombin (2 U/mL for 5 minutes) markedly reduced ATP release on mechanical stimulation (median reduction, 69%; \(N = 55\)). The reduction by thrombin was only 9% after pretreatment with the PAR-1 antagonist (10 \(\mu\)M for 30 minutes; \(N = 3\)), 23% in the presence of Y-27632 (10 \(\mu\)M for 30 minutes; \(N = 5\)), and 20% in the presence of chelerythrine (5 \(\mu\)M for 30 minutes; \(N = 5\)). The PAR-4 antagonist YD-3 (50 \(\mu\)M for 30 minutes; \(N = 3\)) did not influence the effect of thrombin on ATP release. Similar to thrombin, TRAP-6 (10 \(\mu\)M for 30 minutes) caused a marked inhibition of ATP release on mechanical stimulation (median reduction of 50%; \(N = 16\)).

**Discussion**

Intercellular Ca\(^{2+}\) wave propagation in response to a point mechanical stimulus is a distinct paradigm for investigating IC.\(^{28,59–61}\) In corneal endothelial cells, the intercellular wave propagation results from a wave of Ca\(^{2+}\) entry and subsequent release of ATP at the leading edge of the cell, followed by a wave of Ca\(^{2+}\) entry and ATP release in the trailing edges of the adjacent cells. This wave of Ca\(^{2+}\) entry is followed by a local release of ATP only at the leading edge of the cell, and this wave of ATP release is followed by a wave of Ca\(^{2+}\) entry only at the trailing edges of the adjacent cells. This pattern of intercellular wave propagation is not observed in all cell types, and it is not clear what factors determine the pattern of intercellular wave propagation.}

**Figure 6.** Gap junctional communication analysis by FRAP. Cells were loaded with carboxyfluorescein. Recovery of the fluorescence after photobleaching (corrected for background bleaching) of a single cell plotted as function of time after bleaching. (A) BCECs pretreated with thrombin versus control conditions. Three minutes after bleaching, a recovery of 68% ± 0.97% was noted in control conditions whereas, in the presence of thrombin, 58% ± 1.29% of the fluorescence was recovered (\(P < 0.001, N = 190\)). (B-D) Effects of thrombin in the presence or absence of ML-7 (10 \(\mu\)M; \(N = 70\)), Y27632 (10 \(\mu\)M; \(N = 70\)) or chelerythrine (5 \(\mu\)M; \(N = 25\)) applied for 30 minutes. \(P < 0.001\) for each combination of inhibitor plus thrombin versus thrombin alone.
propagation is dominated by PIC. \textsuperscript{17,19} mediated by ATP release through hemichannels. \textsuperscript{17,19} In this study, we investigated whether altered MLC phosphorylation influences GJIC and PIC in BCECs. Since thrombin has been shown to produce well-characterized and reproducible acute effects in terms of MLC phosphorylation, \textsuperscript{46,48,49} we examined its effect on GJIC and PIC in conjunction with a known set of agents that inhibit GJIC and PIC. \textsuperscript{17,18} The main findings of the present study are that thrombin brings about a very robust inhibition of the Ca\textsuperscript{2+} wave propagation through mechanisms linked to MLC phosphorylation and that the dominant effect of the protease is through inhibition of the ATP-dependent PIC.

**Effects of Thrombin: Dependence on PAR-1 Receptors**

To unravel the signaling pathway underlying inhibition of the Ca\textsuperscript{2+} wave highlighted in Figures 1 and 2, we explored the identity of the thrombin receptors that cause the effect. Experiments with PAR agonists (thrombin and TRAP-6) and antagonists\textsuperscript{56,57} provide clear evidence that the effect of thrombin is mediated through PAR-1 receptors (Figs. 3, 4). Independent experiments demonstrated that cells pretreated with thrombin showed an increase in [Ca\textsuperscript{2+}], on exposure to exogenous ATP that was not significantly different from that in untreated cells. This suggests that thrombin had no effect on the P2 receptors and their signaling pathway involved in the Ca\textsuperscript{2+} wave propagation, or on the intracellular Ca\textsuperscript{2+} stores. Moreover, the effect of TRAP-6 excluded that the inhibitory effect of thrombin on the wave propagation is mediated by proteolytic activity on cell surface molecules other than PAR-1 (Fig. 4A).

**Effects of Thrombin on GJIC and PIC: Dependence on MLC Phosphorylation**

As shown in our previous studies, \textsuperscript{46,48,49} thrombin induces MLC phosphorylation, leading to a breakdown of barrier integrity in BCECs. The pathways involved are illustrated in Figure 11. Our observation that the effect of thrombin on Ca\textsuperscript{2+} wave propagation is suppressed by Y-27632, chelerythrine and ML-7 (shown in Fig. 5), is an indication of an involvement of MLC phosphorylation in intercellular communication. Specifically, as shown in Figure 11, Y-27632, chelerythrine and ML-7 inhibit Rho kinase, PKC, and MLCK, respectively, each of which affect MLC phosphorylation.

**Effect on GJIC.** FRAP protocol\textsuperscript{19} is quantitative and hence we used it to assess the effect of thrombin on GJIC. As shown...
also inhibited by thrombin, and this effect was also over-

FIGURE 9. Effect of Gap26 and flufenamic acid on reduction of AA induced by thrombin in the presence of ARL-67156. Cells were treated with thrombin (2 U/mL for 5 minutes) after incubation with ARL-67156 (ARL, 100 μM for 30 minutes) and with the combination of ARL with either Gap26 (300 μM for 30 minutes) or flufenamic acid (FFA; 50 μM for 30 minutes). (A) AA in the absence of PAR-1 agonist (control). (A) AA in the presence of ARL and Gap26 and thrombin is not significantly different from that in the presence of ARL and Gap26 alone (N = 40). (B) Similar results were obtained with FFA (N = 35). *P < 0.001 for comparison between AA in the presence versus absence of the PAR-1 agonist in contemporaneous experiments.

in Figure 6, in the FRAP experiments thrombin inhibited GJIC, although the reduction was limited (~15%). Furthermore, in the presence of the selective protein kinase inhibitors, the reduction in the fluorescence recovery was suppressed. This indicates that GJIC is inhibited in response to MLC phosphorylation.

Effect on PIC. The large reduction of the AA in response to Gap26 (known to block connexin hemichannels; Fig. 7) or to apyrases (Fig. 8B) clearly highlights that PIC, mediated by ATP release through hemichannels, dominates the wave propagation in BCECs. When the hydrolysis of the released ATP is prevented using ARL-67156, the Ca2+ wave propagation is significantly enhanced. As shown in Figure 8A, this increase of the AA is almost completely abolished by thrombin, implying that thrombin has a major inhibitory effect on PIC.

The significant effect of thrombin on PIC led us to explore further the effects of MLC phosphorylation on the activity of hemichannels. We first examined whether hemichannel-mediated ATP release in response to mechanical stimulation could be blocked by agents that affect MLC phosphorylation. The ATP release was inhibited by thrombin, and moreover, this effect was suppressed by the kinase inhibitors (ML-7, chelerythrine, and Y-27632). Our second approach was to confirm whether hemichannels are influenced by MLC phosphorylation, based on LY uptake in a Ca2+-free medium, which occurs through hemichannels. As shown in Figure 10, LY uptake is also inhibited by thrombin, and this effect was also over-

come by the kinase inhibitors. Thus, PIC, through hemichannel-mediated ATP release, is inhibited by MLC phosphorylation. Cotrina et al.29 also have implicated actin cytoskeleton in the propagation of MS Ca2+ waves in astrocytes. However, in contrast to our findings in BCECs, they observed inhibition of the intercellular Ca2+ wave propagation in response to ML-7.

The above observations led us to conclude that the effect of thrombin on the Ca2+ wave propagation is largely through inhibition of PIC.

Effects of Thrombin: Role of Cortical Actin

Gap junctions are located in close proximity to the tight and adherens junctions, and share common linker proteins that bind to the actin cytoskeleton.28–30 The adherens junctions, by promoting cell–cell adhesion, stabilize the interactions of transmembrane proteins of the tight junction complex (i.e., claudin, occludin, and junctional adhesion molecule) that are essential to occlude the paracellular space.30–32 This tethering force brought on by the adherens junctions is opposed by the centripetal forces produced by an increase in contractility of the cortical actin cytoskeleton.35–37 Thus, agents that raise contractility of the actin cytoskeleton are known to breakdown the barrier integrity of cellular monolayers. Recent studies on the corneal endothelium also reflect these findings.36–39 When BCECs were exposed to thrombin, a significant increase in the paracellular permeability was noticed concomitant with a disruption of the cortical actin.36

These observations suggest that agents that can increase MLC phosphorylation, could disrupt gap junctions and thereby affect GJIC, because in nonmuscle cells, the primary mechanism of increasing contractility of the actin cytoskeleton is through an increase in MLC phosphorylation.40–44 Therefore, our findings suggest an involvement of increased contractility of the actin cytoskeleton in the inhibition of gap junctions by thrombin (Fig. 11). In agreement with the above hypothesis, our results show a significant, although small, inhibition of GJIC in response to increased MLC phosphorylation.

Although much less is known about the effects of an altered actin cytoskeleton on hemichannels, it is possible that connexon trafficking, internalization, and gating of hemichannels could be influenced by the cortical actin cytoskeleton through altered scaffolding, or altered signaling via direct protein–protein interactions or downstream transduction pathways.

Potential Effects of MLC Phosphorylation on Cx43 and ZO-1 Interactions

As noted in Figure 11, a key mechanism toward MLC phosphorylation that is upregulated by thrombin is the activation of the RhoA–Rho kinase axis.45–47 Thus, an inhibition of the Ca2+ wave secondary to the thrombin-induced MLC phosphorylation and its prevention by the Rho kinase inhibitor Y-27632 (Figs. 5, 6) suggest an involvement of the RhoA-Rho kinase axis in GJIC and PIC. In line with this conclusion, Anderson et al.9 have reported that inhibition of Rho kinase in corneal epithelial cells enhances dye coupling. Rouach et al.66 demonstrated RhoA-dependent inhibition of GJIC by SIP (sphingosine-1-phosphate), based on dye coupling and electrophysiologic experiments.

In other studies on the effect of thrombin and other GPCR agonists, inhibition of GJIC by tyrosine phosphorylation of Cx43 has been shown.68–70 Specifically, the nonreceptor tyrosine kinase src was implicated in acute reduction of GJIC, and this effect was shown to be independent of the RhoA-Rho

FIGURE 10. The bar graph shows the increased LY uptake in the presence of thrombin and ARL (ARL, 100 μM; 15%) and in the presence of thrombin and Gap26 (300 μM; 15%). As shown in Figure 10, LY uptake is also inhibited by thrombin, and this effect was also over-

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kinase pathway. However, it has been shown that this effect of Src disrupts the interaction between ZO-1 and Cx43, reducing the stability of Cx43 at the cell surface. Furthermore, a recent study showed that MLCK-mediated MLC phosphorylation brings about a structural separation of ZO-1 and occludin from the actin cytoskeleton. In this context, Cx43, which is known to interact with ZO-1 in several cell types, is also expressed in corneal endothelial cells, and such an interaction is implicated in connexin trafficking and internalization and accretion at gap junctions. However, taking into account the finding that thrombin inhibits Ca²⁺ wave propagation, FRAP and dye uptake within minutes (Fig. 10), we speculate that thrombin-induced redistribution of ZO-1, secondary to MLC phosphorylation, can also affect gating of the Cx43-dependent hemichannels and gap junctions.

**Physiological Significance in the Corneal Endothelium**

Our observations highlight the influence of MLC phosphorylation and contractility of the actin cytoskeleton on IC in the corneal endothelium. Although this is partly reminiscent of the role of the actin cytoskeleton in the loss of barrier integrity, the block of hemichannels involved in the PIC suggests the importance of the disposition of cortical
actin in the function of the endothelium in addition to the role of the tethering forces. In situations of inflammatory stress, which usually result in MLC phosphorylation, blockage of the hemichannels can be thought of as being a defense mechanism. Thus, such a blockade would limit IC and prevent potential bystander effects, through limiting the release of signaling molecules. In addition, it is plausible that hemichannel blockage can prevent disturbances of intracellular homeostasis by solute entry/loss through the channels. Furthermore, the influence of the cortical actin cytoskeleton organization and/or contractility on hemichannels suggests their altered activity in response to hypoxia and aging, since, under these conditions, the corneal endothelial cells show significant alterations in the actin cytoskeleton, as suggested by polymegathism and polymorphism.

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References


