

ARTICLE

Prostaglandin E₂ stimulates the epithelial sodium channel (ENaC) in cultured mouse cortical collecting duct cells in an autocrine manner

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Prostaglandin E₂ (PGE₂) is the most abundant prostanoid in the kidney, affecting a wide range of renal functions. Conflicting data have been reported regarding the effects of PGE₂ on tubular water and ion transport. The amiloride-sensitive epithelial sodium channel (ENaC) is rate limiting for transepithelial sodium transport in the aldosterone-sensitive distal nephron. The aim of the present study was to explore a potential role of PGE₂ in regulating ENaC in cortical collecting duct (CCD) cells. Short-circuit current (*I*_{SC}) measurements were performed using the murine mCCD_{cl1} cell line known to express characteristic properties of CCD principal cells and to be responsive to physiological concentrations of aldosterone and vasopressin. PGE₂ stimulated amiloride-sensitive *I*_{SC} via basolateral prostaglandin E receptors type 4 (EP₄) with an EC₅₀ of ~7.1 nM. The rapid stimulatory effect of PGE₂ on *I*_{SC} resembled that of vasopressin. A maximum response was reached within minutes, coinciding with an increased abundance of β-ENaC at the apical plasma membrane and elevated cytosolic cAMP levels. The effects of PGE₂ and vasopressin were nonadditive, indicating similar signaling cascades. Exposing mCCD_{cl1} cells to aldosterone caused a much slower (~2 h) increase of the amiloride-sensitive *I*_{SC}. Interestingly, the rapid effect of PGE₂ was preserved even after aldosterone stimulation. Furthermore, application of arachidonic acid also increased the amiloride-sensitive *I*_{SC} involving basolateral EP₄ receptors. Exposure to arachidonic acid resulted in elevated PGE₂ in the basolateral medium in a cyclooxygenase 1 (COX-1)-dependent manner. These data suggest that in the cortical collecting duct, locally produced and secreted PGE₂ can stimulate ENaC-mediated transepithelial sodium transport.

Introduction

In the kidney the amiloride-sensitive epithelial sodium channel (ENaC) is localized to the apical membrane of tubular epithelial cells lining the so-called aldosterone-sensitive distal nephron (ASDN). This comprises the late distal convoluted tubule, the connecting tubule, and the collecting duct (CD). In the ASDN, transepithelial sodium transport critically depends on the activity and abundance of ENaC. The precise regulation of ENaC is essential for the fine-tuning of urinary sodium excretion to match dietary sodium intake. Thus, ENaC regulation plays a key role for total body sodium balance and is critical for the long-term control of extracellular fluid volume and arterial blood pressure. ENaC is regulated by several hormones, including aldosterone and vasopressin, as well as by several local factors (Garty and Palmer, 1997; Loffing and Korbmaier, 2009; Bankir et al., 2010; Rossier, 2014; Kleyman et al., 2018).

Prostanoids are derivatives of arachidonic acid and are produced and secreted by many different cells. Prostanoids have

complex effects on renal function (Grantham and Orloff, 1968; Breyer and Breyer, 2000b) and contribute to the regulation of sodium and water excretion, renin secretion, renal blood flow, and glomerular filtration (Hao and Breyer, 2007). Because they are rapidly metabolized, these derivatives are thought to act within close proximity to the site of their synthesis in either an autocrine or paracrine manner (Fenton and Knepper, 2007). Prostaglandin E₂ (PGE₂), derived via cyclooxygenase (COX), is the most abundant prostanoid in the kidney. There are conflicting reports regarding the effects of PGE₂ on sodium and water transport within the collecting duct (Breyer and Breyer 2000a). Several studies suggest that in the renal medulla, PGE₂ reduces sodium absorption (Stokes and Kokko, 1977; Iino and Imai, 1978). In contrast, inhibition of prostaglandin synthesis has been reported to be associated with increased urinary sodium excretion in conscious dogs, probably due to diminished

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sodium reabsorption in the collecting duct (Kirschenbaum and Stein, 1976). Moreover, PGE₂ may be able to stimulate water and sodium absorption within the cortical CD (CCD) to maintain blood pressure in volume-contracted states (Hao and Breyer, 2008).

PGE₂ exerts its diverse effects by binding to four distinct G protein-coupled receptors: EP₁–EP₄ (Hao and Breyer, 2007). The expression pattern of these receptors determines local effects of PGE₂ (Breyer and Breyer, 2000b). EP₂ and EP₄ receptors are G_{as}-coupled receptors, and ligand binding to these receptors stimulates adenylyl cyclase (AC), raising cytosolic cAMP concentration. In addition, alternative EP₄ receptor pathways may play a role in mediating downstream effects (Fujino and Regan, 2006; Li et al., 2017). EP₃ receptor is coupled to G_{ai/o} and has an inhibitory effect on AC, thereby lowering cytosolic cAMP concentration. The EP₁ receptor is G_{aq/11} coupled and promotes signaling via inositol 1,4,5-trisphosphate (IP₃) and diacylglycerol resulting in elevated intracellular Ca²⁺ concentration and activation of protein kinase C (PKC; Narumiya et al., 1999; Breyer and Breyer, 2001).

Findings in animal models of nephrogenic diabetes insipidus indicate a likely role of prostaglandins in regulating renal water transport. In these animals EP₂ and EP₄ receptor activation alleviated urine-concentrating defects (Li et al., 2009; Olesen et al., 2011). Conversely, nephron-specific or collecting duct-specific knockout of EP₄ receptor in mice promoted urine-concentrating defects (Gao et al., 2015). Interestingly, in *in vitro* studies of the collecting duct activation of EP₂ and EP₄ receptors promoted apical targeting and phosphorylation of aquaporin-2 water channels, reminiscent of the effect of vasopressin (Olesen et al., 2011, 2016; Gao et al., 2015). Vasopressin-dependent coupling between amiloride-sensitive sodium transport and water flow has been demonstrated in a mouse CCD cell line (mCCD_{chl}; Gaeggeler et al., 2011) as well as in isolated perfused rat CCDs (Reif et al., 1986). Presently, it is not known whether a potential modulatory effect of PGE₂ on tubular water transport is associated with an effect of PGE₂ on ENaC-mediated sodium absorption in the CCD.

The aim of the present study was to investigate whether PGE₂ modifies ENaC-mediated transepithelial sodium transport in mCCD_{chl} cells. This cell line provides a highly differentiated and hormone-responsive model of CCD principal cells (Gaeggeler et al., 2005, 2011). It is well suited to study the regulation of electrogenic transepithelial ion transport by hormonal and local mediators (Edinger et al., 2014; Mansley et al., 2015, 2018, 2019). We sought to identify the prostaglandin receptors present in this model and explored the interplay between PGE₂ and two key hormones that promote salt and water reabsorption in the CCD, namely aldosterone and vasopressin. Finally, we investigated whether PGE₂ is synthesized and secreted by mCCD_{chl} cells.

Materials and methods

Cell culture

Mouse CCD cells (mCCD_{chl}) were kindly provided by Bernard Rossier (University of Lausanne, Lausanne, Switzerland) and cultured as described previously (Gaeggeler et al., 2005; Mansley et al., 2015, 2018). Cells were routinely passaged every 7 d (passage 25–34) and maintained in cell culture dishes at 37°C in a 5% CO₂ atmosphere in Dulbecco's modified Eagle's Medium

(DMEM)/Ham's F12 (1:1 vol/vol) medium supplemented with 2% FBS, 1 nM triiodothyronine, 60 nM sodium selenite, 10 ng·ml^{−1} epidermal growth factor, 5 μg·ml^{−1} human apotransferrin, 50 nM dexamethasone, 5 μg·ml^{−1} insulin, 100 U·ml^{−1} penicillin, and 100 μg·ml^{−1} streptomycin. For experimental procedures, cells were seeded onto Millicell-PCF culture plate inserts (EMD Millipore) with a membrane pore size of 0.4 μm and an effective surface area of either 0.6 cm² or 4.2 cm² and grown to form a polarized epithelial monolayer. At day 5 after seeding, the cell culture medium was replaced by a medium devoid of FBS, apotransferrin, and epidermal growth factor. Finally, 24 h before experiments, dexamethasone was removed from the medium.

Chemicals and solutions

DMEM/Ham's F12 (1:1 vol/vol) without phenol red was from Life Technologies and FBS from PAA and Bio&Sell. SC-560, AH23848, AH6809, GW627368X, and Sulprostone were from Cayman Chemical (Biozol); PGE₂ and indomethacin were from Enzo Life Sciences. TCS2510 was from Tocris. Lumiracoxib was kindly provided by Prof. Dr. K. Brune (Institute of Experimental and Clinical Pharmacology and Toxicology, Friedrich-Alexander University, Erlangen, Germany). All other drugs were ordered from Sigma-Aldrich.

Transepithelial measurements

Experimental procedures were essentially the same as described previously (Bertog et al., 2008; Mansley et al., 2015, 2018). Briefly, transepithelial voltage (V_{te}) and resistance (R_{te}) were routinely checked using a commercially available epithelial volt-ohm meter and a set of two sticks "STX" electrodes (World Precision Instruments). On days 9–11, inserts with confluent mCCD_{chl} cell monolayers were transferred to Ussing chambers for continuous equivalent short-circuit current (I_{SC}) measurements using a CVC6 clamp device (Fiebig) as described previously (Bertog et al., 1999). Alternatively, modified Ussing chambers were used and kept in an incubator gassed with 5% CO₂, and the temperature was maintained at 37°C (Mansley et al., 2015, 2018). These miniaturized chambers were designed to minimize mechanical perturbations and to reduce the bath volumes in the apical and basolateral compartment to 0.35–0.6 ml and 0.55–1.0 ml, respectively. Both experimental approaches showed similar results in the transepithelial parameters investigated. R_{te} was evaluated every 2–30 s by measuring voltage deflections induced by 400-ms symmetrical square current pulses of ± 3–5 μA. Using R_{te} and open-circuit V_{te}, the equivalent I_{SC} was calculated according to Ohm's law. Conventionally, a lumen-negative V_{te} corresponds to a positive I_{SC} which may be due to electrogenic cation absorption, electrogenic anion secretion or a combination of both. After transfer into Ussing chambers, cells were allowed to equilibrate for 30 to 60 min before manipulations took place. At the end of each experiment, amiloride (10 μM) was applied to the apical compartment to determine the ENaC-mediated I_{SC} component.

Reverse transcription (RT) PCR

Total RNA was extracted from mCCD_{chl} cells following transepithelial measurements using NucleoSpin RNA Kit-XS (Macherey-Nagel)

Table 1. Primer pairs of target genes used for RT-PCR in this study

cDNA	Sense primer (5'→3')	Antisense primer (5'→3')
EP ₁	CGCAGGGTTCACGCACACGA	CACTGTGCCGGGAACACGC
EP ₂	AGGACTTCGATGGCAGAGGAG AC	CAGCCCCTTACACTTCTCCAATG AC
EP ₃	CCGGGCACGTGGTGCTTCAT	TAGCAGCAGATAAACCCAGG
EP ₄	TTCCGCTCGTGGTGCGAGTGT TC	GAGGTGGTGTCTGCTTGGGTCAG TC
COX-1	CCTCTTTCCAGGAGCTCACA	CGGGTAGAACTCTAAAGCATCG
COX-2	GGGAGTCTGGAACATTGTGAA	GCACATTGTAAGTAGGTGGAC TGT
mPGES1	AGCACACTGCTGGTCATCAA	CTCCACATCTGGGTCACTCC
mPGES2	GACCCTGTACCAGTACAAGAC	GAGGAGTCATTGAGCTGTTGC
cPGES	CGAATTTTGACCGTTTCTCTG	TGAATCATCATCTGCTCCATCT
β-Actin	TCACCCCACTGTGCCATCT AC	GAGTACTTGCGCTCAGGAGGAGC AC

according to the manufacturer's instructions. Lysates from two 0.6-cm² cell culture plate inserts were pooled to enhance the RNA concentration.

For prostanoid receptors (EP₁–EP₄), RT and PCR amplification were performed using an RNA Reverse Transcription System Kit (Promega). Specific primers (Table 1) for murine EP₁–EP₄ receptors and β-actin were designed as described previously (Arakawa et al., 1996) and synthesized by Invitrogen. PCR cycling conditions were 95°C for 2 min, followed by 32 repeats of 95°C for 30 s, 60°C for 45 s, and 72°C for 1 min. Final extension time was 4 min at 72°C.

For all other primers, RT was performed with 0.5 μg RNA using QuantiTect Reverse Transcription Kit (Qiagen) as per the manufacturer's protocol. Specific primers for the COX isoenzymes 1 and 2 (COX-1 and COX-2) and for the cytosolic PGE₂ synthase (cPGES) were designed using universal probe library system (Roche), whereas specific primers for the microsomal PGE synthases-1 and -2 (mPGES1 and, mPGES2) were designed as described previously (mPGES1: Soodvilai et al., 2009; mPGES2: Yang et al., 2006a). Primers were obtained from biomers.net. PCR was performed using 10 pM specific primers in PCR buffer (buffer Y; Peqlab). Samples were denatured for 5 min at 95°C, followed by 35 repeats of 94°C for 1 min, 60°C for 1 min, and 72°C for 1 min. Final extension time was 10 min at 72°C. All reactions were performed in a MJ-Research PTC-200 Peltier Thermo Cycler (Biozym). Amplified PCR products were separated by agarose gel electrophoresis (1.5% universal agarose; Bio&Sell) and stained by ethidium bromide. Bands of PCR products were extracted and sequenced (LGC Genomics). For sequence comparison we used the standard nucleotide BLAST software (National Center for Biotechnology Information, National Library of Medicine).

Biotinylation assay and Western blotting

To detect β-ENaC in Western blot experiments, a previously described custom-made antibody was used in a 1:2,000 dilution (Krueger et al., 2009; Nesterov et al., 2016; Mansley et al., 2018).

Horseradish peroxidase-coupled goat anti-rabbit antibodies were obtained from Santa Cruz Biotech and used in a dilution of 1:50,000.

Cell surface proteins were labeled using a biotinylation protocol similar to that previously described for surface biotinylation of lung epithelial cell monolayers (Woollhead and Baines, 2006). For biotinylation experiments, mCCD_{cl} cells were grown on permeable supports (Millicell-PCF inserts, membrane size 4.2 cm²; EMD Millipore) for 10 d. All biotinylation steps were performed at 4°C. Cells were chilled to 4°C by washing three times with ice-cold PBS containing 0.7 mM MgCl₂ and 0.5 mM CaCl₂ (PBS-CM). Cells were kept on ice, and biotinylation of the apical cell surface was achieved by adding borate buffer containing 85 mM NaCl, 5 mM KCl, and 15 mM Na₂B₄O₇, pH 9.0, containing 0.5 mg·ml⁻¹ EZ-link sulfo-NHS-SS-Biotin (Pierce). The basolateral compartment was exposed to PBS-CM + 10% FBS. Cells were kept on ice and rocked for 30 min. Cells were washed once with PBS-CM and then the reaction quenched by exposing both the apical and basolateral surface of cells to PBS-CM + 10% FBS for a further 30 min. Cells were washed twice with PBS-CM and then scraped into 200 μl lysis buffer (0.4% deoxycholic acid, 1% Triton X-100, 50 mM EGTA, and 10 mM Tris, pH 7.4) including a protease inhibitor cocktail (Roche). A small sample was taken to determine protein content by BCA assay. Biotinylated proteins were captured by exposing the sample to 50 μl ImmunoPure immobilized Neutravidin agarose beads (Pierce), which had been washed twice with PBS-CM and subsequently with lysis buffer. 250 μg sample was added to the washed beads, and tubes were incubated on a rotor overnight at 4°C. Samples were centrifuged for 2 min at 8,000 rpm (6,200 ×g) at 4°C, and the supernatant was collected separately for the detection of intracellular proteins. Neutravidin beads were washed and centrifuged four times and the biotinylated fraction was finally resuspended in 45 μl of 2× reducing SDS-PAGE sample buffer (Rotiload 1; Roth).

All protein samples were heated for 5 min at 95°C before loading on SDS gels. Proteins were separated by 10% SDS-PAGE, transferred to polyvinylidene difluoride membranes by semidry electroblotting, and probed with the indicated antibodies. Chemiluminescent signals were detected using Super Signal West Femto Chemiluminescent Substrate (Pierce).

Measurement of PGE₂ and cAMP concentrations

For PGE₂ measurements, an enzyme immunoassay was used according to the manufacturer's protocol (Prostaglandin E₂ Express EIA Kit—Monoclonal; Cayman Chemical Company). PGE₂ concentrations were determined in diluted basolateral cell culture medium after incubating mCCD_{cl} cells with arachidonic acid, COX inhibitors, or vehicle at 37°C for 10 min.

For cAMP measurements, mCCD_{cl} cells were lysed to release intracellular cAMP, which was quantified using an enzyme immunoassay according to the manufacturer's instructions (cAMP Biotrak competitive enzyme immunoassay system; GE Healthcare). Cytosolic cAMP concentrations were determined in cells exposed to PGE₂ or vehicle at 37°C for 10 min. Protein concentrations were determined with a bicinchoninic acid assay (BCA Protein Assay Kit; Thermo Scientific).

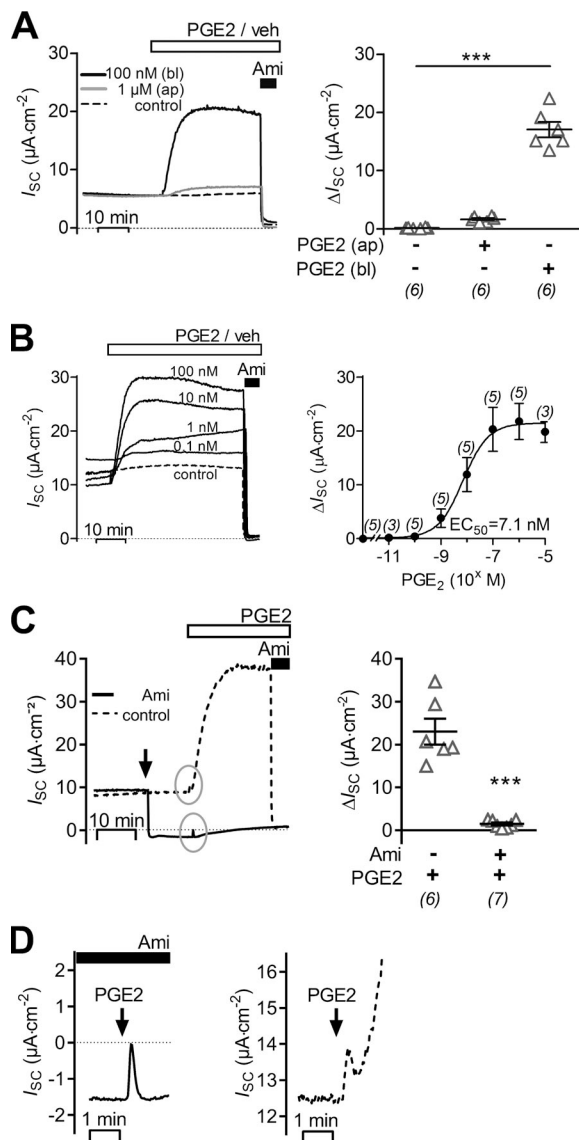


Figure 1. Basolateral PGE₂ stimulates amiloride-sensitive I_{sc} in mCCD_{cl1} cells. (A–C) Representative traces of continuous I_{sc} recordings from mCCD_{cl1} cells are shown in the left panels and data from similar experiments are summarized in the corresponding right panels. During the time periods indicated by the open horizontal bar, PGE₂ or vehicle (veh) was present. Apical application of 10 μ M amiloride (Ami) is indicated by a black horizontal bar. (A) I_{sc} recordings with basolateral PGE₂ (bl; 100 nM), apical PGE₂ (ap; 1 μ M) or vehicle (control) are represented by a black, gray, or dashed line, respectively. (B) I_{sc} recordings are shown with basolateral application of PGE₂ (black lines) in concentrations from 0.1–100 nM or with vehicle (dashed line). A nonlinear fit for the I_{sc} response upon PGE₂ application in concentrations of 10 pM to 10 μ M was used to estimate EC_{50} . (C) Basolateral PGE₂ (100 nM) was applied in the absence (control; dashed line) or presence of 10 μ M amiloride (black line). Arrow indicates time point of apical addition of amiloride or vehicle (control). In the control PGE₂ recording, apical amiloride (10 μ M) was added at the end of the experiment. To summarize data from different experiments as ΔI_{sc} values, the individual I_{sc} values were corrected by subtracting the corresponding baseline I_{sc} value. Gray circles highlight the initial phases of the I_{sc} responses to PGE₂. Summary data (right panels) are presented as individual data points and/or mean values \pm SEM. Numbers of experiments are given in parenthesis. Statistical analysis was determined using one-way ANOVA with Tukey's multiple comparison test or Student's t test where appropriate; ***, $P < 0.001$. (D) The initial phases of the responses to basolateral PGE₂ encircled in C are shown enlarged using

All samples were measured as duplicates.

Data analysis

Data were analyzed using PRISM 5.04 for Windows (GraphPad Software). Summarized data are presented as mean values \pm SE (SEM). Multiple comparisons were subjected to one-way ANOVA followed by ad hoc post-tests as specified in the figure legends; otherwise, Student's t tests were used. P values < 0.05 were required to reject the null hypothesis; *, **, and *** represent P values < 0.05 , 0.01 , and 0.001 , respectively, and "ns" represents P values ≥ 0.05 . Numbers in parentheses in the figures signify the number of samples studied.

Results

Basolateral application of PGE₂ stimulates ENaC-mediated transepithelial sodium transport in mCCD_{cl1} cells

Basolateral application of 100 nM PGE₂ caused a sustained increase of I_{sc} (Fig. 1A), which rose from a baseline value of $6.7 \pm 0.7 \mu$ A \cdot cm⁻² to a maximal value of $23.7 \pm 2.1 \mu$ A \cdot cm⁻² ($n = 6$, $P < 0.001$) within 5–10 min. This I_{sc} increase of $17.0 \pm 1.3 \mu$ A \cdot cm⁻² was associated with a decrease in R_{te} from 6.49 ± 0.41 to 2.61 ± 0.18 k Ω \cdot cm² ($P < 0.001$). In contrast, in matched vehicle-treated control cells I_{sc} and R_{te} remained stable ($6.3 \pm 0.9 \mu$ A \cdot cm⁻² and 6.40 ± 0.54 k Ω \cdot cm² versus $6.4 \pm 0.9 \mu$ A \cdot cm⁻² and 6.34 ± 0.54 k Ω \cdot cm²; $n = 6$). Importantly, apical application of PGE₂ in a concentration of up to 1 μ M had negligible effects on I_{sc} (Fig. 1A). This indicates that the stimulatory effect of PGE₂ is mediated by a basolateral receptor. Apical application of amiloride (10 μ M) at the end of the experiments almost completely inhibited baseline I_{sc} in vehicle-treated cells and the stimulated I_{sc} in PGE₂-treated cells. The inhibitory effect of amiloride on I_{sc} was accompanied by a 1.7-fold R_{te} increase in vehicle-treated cells (from 6.34 ± 0.54 k Ω \cdot cm² to 10.56 ± 0.45 k Ω \cdot cm², $n = 6$; $P < 0.001$) and a 2.6-fold R_{te} increase in PGE₂-treated cells (from 2.61 ± 0.18 k Ω \cdot cm² to 6.69 ± 0.18 k Ω \cdot cm², $n = 6$; $P < 0.001$). Thus, the stimulatory effect of basolateral PGE₂ on I_{sc} can be attributed to an increase in ENaC-mediated transepithelial sodium transport most likely due to an activation of ENaC activity. The latter conclusion is supported by the associated decrease in R_{te} most likely reflecting an increased sodium conductance of the apical cell membrane. As illustrated in Fig. 1B, the stimulatory effect of basolateral PGE₂ was concentration dependent, with a half-maximal effective concentration (EC_{50}) of ~ 7.1 nM. Furthermore, the PGE₂-induced increase in I_{sc} was prevented when apical application of amiloride preceded addition of PGE₂ (Fig. 1C), which confirms that the stimulatory effect of PGE₂ is dependent on ENaC function. Interestingly, a small and transient I_{sc} peak response to PGE₂ was consistently observed in the presence of amiloride (Fig. 1C, left panel; and Fig. 1D, left panel). It was not an experimental artifact, because it was not observed in vehicle-treated controls. This amiloride-insensitive I_{sc} component is most likely due to a transient chloride secretory response mediated by Ca²⁺-

expanded scales. Arrows indicate the time point of PGE₂ application in the continuous presence (black bar, left panel) or absence (right panel) of amiloride.

activated chloride channels (Sandrasagra et al., 2004; Mansley et al., 2015). A close inspection of the initial phase of the I_{SC} response to PGE_2 revealed that a similar initial I_{SC} peak was also detectable in the absence of amiloride (Fig. 1 D, right panel). However, this I_{SC} peak response was highly variable and in most recordings was at least partially concealed by the overlapping rapid onset of the much larger stimulatory effect of PGE_2 on the amiloride-sensitive ENaC-mediated I_{SC} . Therefore, it was not feasible to study this peak response systematically. Taken together, these data indicate that in mCCD_{cl1} cells basolateral PGE_2 predominantly stimulates ENaC-mediated transepithelial sodium transport most likely by increasing apical ENaC activity.

PGE_2 increases the abundance of β -ENaC at the plasma membrane

An increase in the activity of ENaC can be caused by increasing the open probability of the channel (P_o), by increasing the number of channels at the cell surface (N), or by a combination of both factors. Using a biotinylation assay and Western blot analysis with antibodies directed against mouse β -ENaC, an increase (~ 2.5 times) of β -ENaC protein was detected at the apical surface of cells treated with 100 nM PGE_2 on the basolateral side compared with vehicle-treated cells (Fig. 2, A and B). In the corresponding cytosolic fractions, a difference in the abundance of β -ENaC was not detected (data not shown). These data indicate that the stimulatory effect of PGE_2 on ENaC can be attributed at least in part to an increased channel expression at the apical plasma membrane of mCCD_{cl1} cells.

PGE_2 mediates its effects via basolateral EP_4 receptors

To identify the prostanoid receptor responsible for the stimulatory effect of PGE_2 on ENaC activity in mCCD_{cl1} cells, a pharmacological approach with known receptor agonists and antagonists was used. Basolateral addition of an EP_2 receptor antagonist (AH6809; 10 μ M), which also shows some affinity to murine EP_1 receptor (Kiriya et al., 1997), or of sulprostone (100 nM), an agonist of EP_1 and EP_3 receptors, had no effect on baseline I_{SC} and did not alter the stimulatory response to PGE_2 (Fig. 3). In contrast, the EP_4 receptor antagonists AH23848 (100 μ M) or GW627368X (2 μ M) largely diminished or nearly abolished the PGE_2 -mediated stimulatory response, respectively (Fig. 4 A). The small transient peak response to PGE_2 appeared to be preserved in the presence of the EP_4 receptor antagonist GW627368X (Fig. 4 A, middle panel). Basolateral application of TCS2510 (100 nM), an EP_4 receptor agonist, stimulated the amiloride-sensitive current in a similar manner as PGE_2 (Fig. 4 B). These data indicate that PGE_2 stimulates ENaC activity via a basolateral $G_{\alpha s}$ -coupled EP_4 receptor in mCCD_{cl1} cells.

The stimulatory effect of PGE_2 on ENaC is similar to that of vasopressin and forskolin and is associated with an increase in intracellular cAMP

It has previously been reported that in mCCD_{cl1} cells, ENaC-mediated transepithelial sodium transport can be stimulated by vasopressin via a basolateral V2 receptor (Gaeggeler et al., 2011). Therefore, experiments were performed to compare the

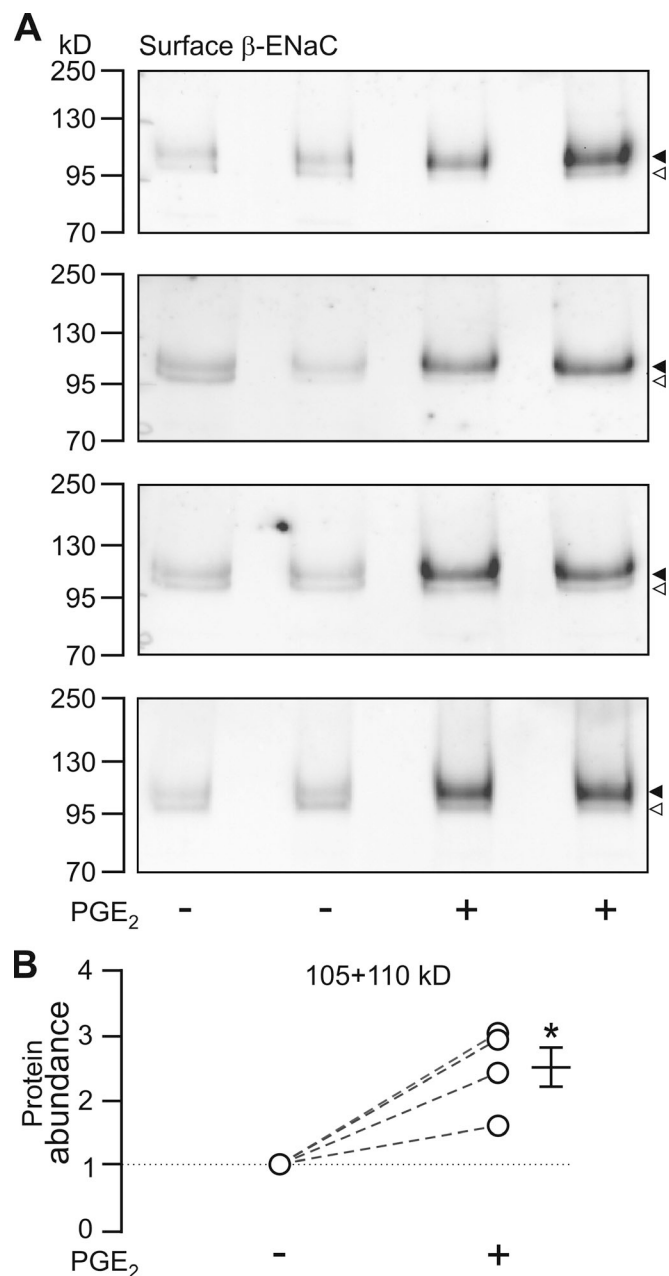


Figure 2. PGE_2 elevates the abundance of β -ENaC at the cell surface.

(A) β -ENaC was detected by Western blot analysis using biotinylated cell surface proteins isolated from mCCD_{cl1} cells exposed on the basolateral side for 30 min to 100 nM PGE_2 (+) or vehicle (–). The four blots represent data from four individual experiments using two filters of mCCD_{cl1} cells per experimental group. Arrowheads to the right of the blots indicate band sizes of ~ 110 kD (filled) and ~ 105 kD (open) as expected for glycosylated and nonglycosylated β -ENaC. (B) The blots shown in A were analyzed by densitometry, and the signals of the lower and upper bands of each lane were combined (105 + 110 kD). In each experiment, two matched mCCD_{cl1} samples were used per group (\pm PGE_2) and an average value was determined for each group. For each experiment, the average value from the PGE_2 -treated cells was normalized to the average value of the nontreated cells (corresponding values are connected by a dashed line). The mean \pm SEM value represents the average change in protein abundance compared with vehicle treatment. Statistical analysis was performed using Student's *t* test; *, $P < 0.05$ versus solvent vehicle controls.

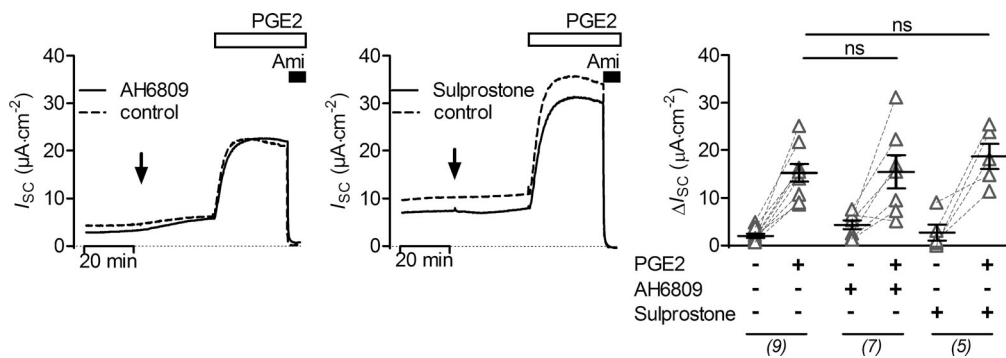


Figure 3. EP₁, EP₂, and EP₃ receptors are not involved in mediating the stimulatory effect of PGE₂ on ENaC. In the left and middle panels, representative I_{sc} recordings are shown. At the time point indicated by an arrow, cells were exposed to 10 μ M AH6809 (EP₂ receptor antagonist; left panel, solid line), 100 nM sulprostone (EP₁/EP₃ receptor agonist; middle panel, solid line), or vehicle in matched control recordings (control, dashed line). Approximately 30 min later, all cells were exposed to 100 nM basolateral PGE₂ and subsequently to apical amiloride (Ami; 10 μ M) as indicated. Summary data (right panel) are presented as ΔI_{sc} values, which were determined by subtracting the corresponding baseline I_{sc} from the I_{sc} reached in the presence (+) or absence (–) of PGE₂, AH6809, and sulprostone as indicated. Data are presented as individual values and their mean \pm SEM; paired data are connected by dashed lines. Numbers of experiments are given in parenthesis. ns, $P > 0.05$, one-way ANOVA with Tukey's multiple comparison test.

effect of vasopressin with that of PGE₂ and to investigate a possible interdependence of these effects. First, the stimulatory effect of vasopressin was confirmed (Fig. 5, A and B). Basolateral application of 25 pM vasopressin increased I_{sc} to a similar extent as 100 nM basolateral PGE₂ in matched experiments. Interestingly, exposure to PGE₂ after prestimulation with vasopressin had no additional stimulatory effect on I_{sc} (Fig. 5 A). Similarly, when cells were initially exposed to PGE₂, subsequent application of vasopressin failed to stimulate I_{sc} further (Fig. 5 B). These findings suggest that the effects of PGE₂ and vasopressin involve the same signaling pathway, i.e., cAMP/PKA. To

confirm this, further experiments were performed with forskolin, a known activator of AC. Forskolin (10 μ M) increased I_{sc} in a similar manner as vasopressin and PGE₂ (Fig. 5 C). After exposure to forskolin, application of PGE₂ (100 nM) had no additional stimulatory effect. To provide evidence that PGE₂ causes an increase in intracellular cAMP, an enzyme immunoassay was used. Indeed, in PGE₂-treated mCCD_{cl1} cells cytosolic cAMP concentration was elevated to 137.6 ± 47.2 fmol/ μ g whole-cell protein compared with 9.5 ± 3.8 fmol/ μ g whole-cell protein in matched vehicle-treated control cells ($n = 3$).

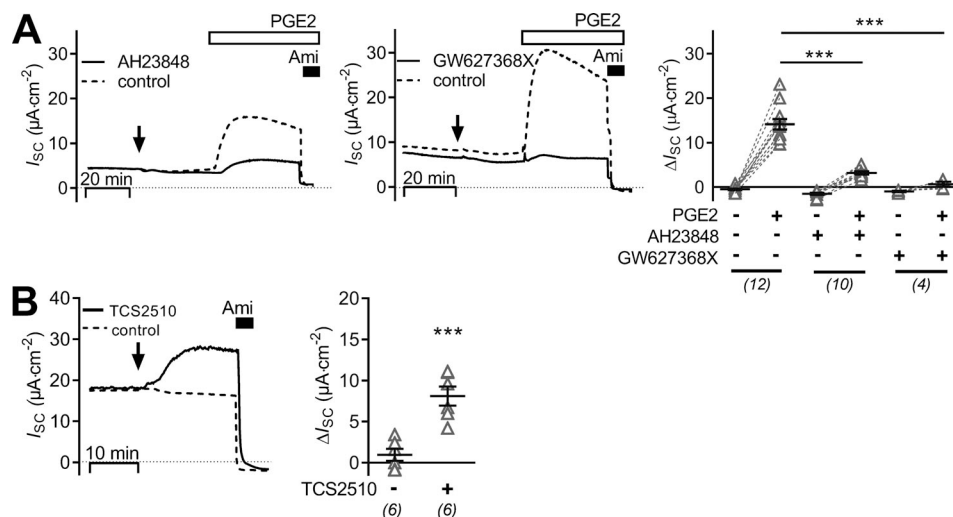


Figure 4. PGE₂ stimulates ENaC activity via basolateral EP₄ receptors. (A) The left and middle panel show representative I_{sc} recordings in which vehicle (control; dashed lines) or EP₄ receptor antagonists (100 μ M AH23848 or 2 μ M GW627368X; solid lines) were added basolaterally at the time point indicated by an arrow. Approximately 30 min later, all cells were exposed to 100 nM basolateral PGE₂ and subsequently to apical amiloride (Ami; 10 μ M) as indicated. Summary data (right panel) are presented as ΔI_{sc} values, which were determined by subtracting the corresponding baseline I_{sc} from the I_{sc} reached in the presence (+) or absence (–) of PGE₂, AH23848, and GW627368X as indicated. (B) Representative I_{sc} recordings (left panel) from cells exposed basolaterally to the EP₄ receptor agonist TCS2510 (solid line; 100 nM) or vehicle (control, dashed line) at the time point indicated by the arrow. Apical amiloride (10 μ M) was applied as indicated. Summary data (right panel) are presented as ΔI_{sc} values determined by subtracting the corresponding baseline I_{sc} from the I_{sc} reached after treating cells with TCS2510 (+) or vehicle (–). Data are presented as individual values and their mean \pm SEM; paired data are connected by dashed lines. Numbers of experiments are given in parentheses. ***, $P < 0.001$, one-way ANOVA with Tukey's multiple comparison test.

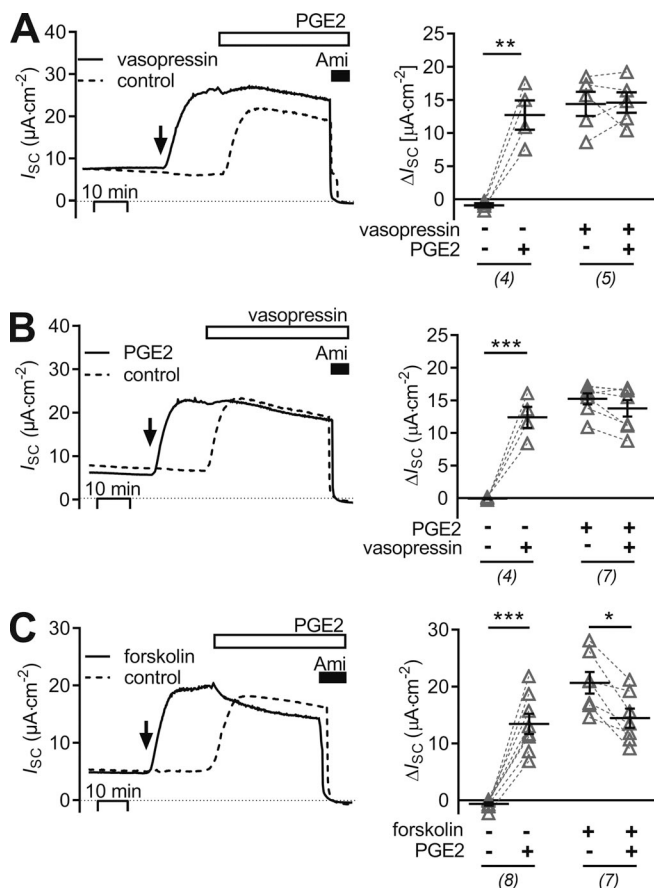


Figure 5. The stimulatory effect of PGE₂ is similar and not additive to those of vasopressin and forskolin. (A–C) Representative traces of continuous I_{sc} recordings from mCCD_{cl1} cells are shown in the left panels, and data from similar experiments are summarized in the corresponding right panels. (A) At the time point indicated by an arrow, cells were exposed to 25 pM basolateral vasopressin (solid line) or vehicle in matched control recordings (control, dashed line). About 20 min later all cells were exposed to 100 nM basolateral PGE₂ and subsequently to apical amiloride (Ami; 10 μ M) as indicated. (B) At the time point indicated by an arrow, cells were exposed to 100 nM basolateral PGE₂ (solid line) or vehicle in matched control recordings (control, dashed line). Approximately 20 min later, all cells were exposed to 25 pM basolateral vasopressin and subsequently to apical amiloride (10 μ M) as indicated. (C) At the time point indicated by an arrow, cells were exposed to 10 μ M forskolin (solid line) to stimulate adenylyl cyclase or to vehicle in matched control recordings (control, dashed line). Approximately 20 min later, all cells were exposed to 100 nM basolateral PGE₂ and subsequently to apical amiloride (10 μ M) as indicated. Summary data (right panel) are presented as ΔI_{sc} values, which were determined by subtracting the corresponding baseline I_{sc} from the I_{sc} reached in the presence (+) or absence (–) of PGE₂, vasopressin, and forskolin as indicated. Data are presented as individual values and their mean \pm SEM. Paired data are connected by dashed lines. Numbers of experiments are given in parentheses. *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.001$, paired Student's t test.

The stimulatory effect of PGE₂ on ENaC activity is preserved in cells pretreated with aldosterone

As shown previously (Gaeggeler et al., 2005; Mansley et al., 2018), mCCD_{cl1} cells treated with a physiological concentration of aldosterone (3 nM) responded with a sustained increase of I_{sc} from $8.7 \pm 0.9 \mu A \cdot cm^{-2}$ to $17.9 \pm 1.6 \mu A \cdot cm^{-2}$ ($n = 9$, $P < 0.001$; Fig. 6) reaching a new plateau within ~ 2 h. Basolateral application

of PGE₂ (100 nM) at the plateau of the aldosterone response caused a further rapid (within 5–10 min) increase in I_{sc} by $16.8 \pm 1.1 \mu A \cdot cm^{-2}$. This stimulatory effect of PGE₂ in aldosterone-treated cells was similar to that in matched vehicle-treated control cells, averaging $17.3 \pm 1.5 \mu A \cdot cm^{-2}$ ($n = 11$; Fig. 6). These results indicate that the stimulatory effect of PGE₂ on ENaC is independent of the signaling pathways mediating the stimulatory effect of aldosterone.

PGE₂ is generated and secreted by mCCD_{cl1} cells exposed to the precursor arachidonic acid and stimulates ENaC activity in an autocrine manner

The finding that exogenously applied PGE₂ stimulates ENaC activity in mCCD_{cl1} cells raised the question whether endogenously generated prostanoids may elicit a similar response. To promote the synthesis of endogenous PGE₂, its precursor arachidonic acid was applied at a concentration of 50 μ M. At this concentration, arachidonic acid is not rate limiting for the biosynthesis of PGE₂ (Bonvalet et al., 1987). Apical application of arachidonic acid increased I_{sc} from $9.8 \pm 1.5 \mu A \cdot cm^{-2}$ to $25.1 \pm 2.4 \mu A \cdot cm^{-2}$ ($n = 8$, $P > 0.001$; Fig. 7 A, dashed line). The stimulated I_{sc} was inhibited by amiloride, which indicates that the stimulatory effect of arachidonic acid is due to increased ENaC-mediated transepithelial sodium transport. The stimulatory effect of arachidonic acid was similar to that of PGE₂, and subsequent application of PGE₂ (100 nM) to the basolateral bath did not increase I_{sc} further ($\Delta I_{sc} = -1.5 \pm 0.6 \mu A \cdot cm^{-2}$). Importantly, treating mCCD_{cl1} cells with the COX-1/2 inhibitor indomethacin (50 μ M) completely prevented the stimulatory response to arachidonic acid ($\Delta I_{sc} = 0.3 \pm 0.2 \mu A \cdot cm^{-2}$, $n = 8$; Fig. 7 A, solid line). In contrast, the stimulatory effect of subsequently applied PGE₂ was fully preserved ($\Delta I_{sc} = 10.7 \pm 1.3 \mu A \cdot cm^{-2}$, $n = 8$). Thus, mCCD_{cl1} cells are capable of synthesizing and releasing a COX-derived prostanoid, most likely PGE₂, which can stimulate ENaC activity in an autocrine manner, preventing a further stimulation by exogenous PGE₂. This hypothesis is further supported by the finding that the stimulatory effect of arachidonic acid was largely reduced in the presence of the EP₄ receptor antagonists AH23848 (100 μ M) and GW627368X (2 μ M) in the basolateral solution (Fig. 7 B). In mCCD_{cl1} cells, we did not observe an inhibitory effect of arachidonic acid on ENaC-mediated I_{sc} . In contrast, ENaC inhibition via cytochrome P450 epoxygenase-dependent pathways has been reported in microdissected rat CCD (Wei et al., 2004) and mpkCCD cells (Pavlov et al., 2011). The lack of an inhibitory effect in mCCD_{cl1} cells cannot be attributed to an insufficient concentration of arachidonic acid used in our experiments because in rat CCD, the concentration required to inhibit ENaC activity by 50% was $\sim 2 \mu$ M.

In additional experiments cell culture medium was collected from the basolateral bath of cells treated with apical arachidonic acid (donor cells). The collected medium was transferred to the basolateral compartment of nontreated mCCD_{cl1} cells (receiver cells) in a 1:1 (vol/vol) ratio. This resulted in a significant stimulation of I_{sc} in the receiver cells ($\Delta I_{sc} = 5.6 \pm 1.4 \mu A \cdot cm^{-2}$, $n = 7$). Subsequent application of PGE₂ had an additional but less pronounced stimulatory effect on I_{sc} . The stimulatory effect of medium from donor cells was preserved when COX-1/2 activity

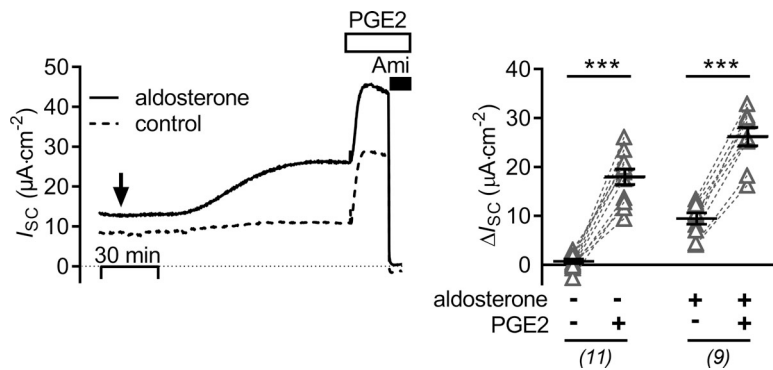


Figure 6. The stimulatory effect of PGE₂ upon ENaC activity is preserved in cells pretreated with aldosterone. The left panel shows representative I_{sc} recordings in which vehicle (control; dashed line) or aldosterone (3 nM; solid line) were added bilaterally at the time point indicated by an arrow. Approximately 2 h later, cells were exposed to 100 nM basolateral PGE₂ and subsequently to apical amiloride (Ami; 10 μ M) as indicated. Summary data (right panel) are presented as ΔI_{sc} values, which were determined by subtracting the corresponding baseline I_{sc} from the I_{sc} reached in the presence (+) or absence (–) of aldosterone and PGE₂ as indicated. Data are presented as individual values and their mean \pm SEM. Paired data are connected by dashed lines. Numbers of experiments are given in parentheses. ***, $P < 0.001$, paired Student's t test.

in the receiver cells was inhibited by indomethacin (ΔI_{sc} $5.4 \pm 1.3 \mu A \cdot cm^{-2}$, $n = 7$; Fig. 7 C). This rules out the possibility that the stimulatory effect of the transferred medium was mediated by a contamination with arachidonic acid. Finally, the stimulatory effect of the transferred medium on ENaC activity was largely abolished when the receiver cells were pretreated with the EP₄ receptor antagonist GW627368X in the basolateral bath (ΔI_{sc} = of $1.4 \pm 0.5 \mu A \cdot cm^{-2}$). In contrast, medium transfer increased I_{sc} by $12.4 \pm 1.3 \mu A \cdot cm^{-2}$ in nontreated matched control cells ($n = 4$, $P < 0.001$; Fig. 7 D).

Stimulation of ENaC in mCCD_{cl} cells by arachidonic acid requires COX-1 activity

Similar to treating mCCD_{cl} cells with the nonselective COX-1/2 inhibitor indomethacin (Fig. 7 A), treating cells with the selective COX-1 inhibitor SC-560 (0.14 μ M; Fig. 8 A) largely prevented the stimulatory effect of arachidonic acid on ENaC activity. The arachidonic acid-induced ΔI_{sc} was reduced to $4.4 \pm 0.6 \mu A \cdot cm^{-2}$ ($n = 6$) compared with the effect of arachidonic acid in matched controls with a ΔI_{sc} of $17.1 \pm 2.7 \mu A \cdot cm^{-2}$ ($n = 7$, $P < 0.05$). In contrast, lumiracoxib (2 μ M; Fig. 8 B), a highly selective COX-2 inhibitor, did not prevent the stimulatory effect of arachidonic acid. Indeed, in cells treated with lumiracoxib, the effect of arachidonic acid was preserved with a ΔI_{sc} of $15.7 \pm 2.7 \mu A \cdot cm^{-2}$ ($n = 6$), which was not different from the ΔI_{sc} of $18.9 \pm 2.7 \mu A \cdot cm^{-2}$ ($n = 6$) elicited by arachidonic acid in vehicle-treated control cells. These data indicate that COX-1 is the dominant isoenzyme involved in PGE₂ generation in mCCD_{cl} cells. To confirm synthesis and basolateral secretion of PGE₂, its concentration was measured in the basolateral culture medium of mCCD_{cl} cells. The concentration of PGE₂ in the basolateral medium of mCCD_{cl} cells exposed to apical arachidonic acid was significantly higher (16.3 ± 3.8 nmol·l⁻¹, $n = 6$) than that of vehicle-treated control cells (0.3 ± 0.2 nmol·l⁻¹, $n = 8$, $P < 0.01$; Fig. 8 C). The arachidonic acid-dependent increase in basolateral PGE₂ concentration was abolished by the COX-1 inhibitor SC-560 (0.3 ± 0.1 nmol·l⁻¹, $n = 7$, $P < 0.001$), but not by the COX-2 inhibitor lumiracoxib (7.4 ± 5.4 nmol·l⁻¹, $n = 7$, $P > 0.05$), which is consistent with the I_{sc} data reported above.

Transcripts for EP receptors and for enzymes of the PGE₂ biosynthetic pathway were detected in mCCD_{cl} cells

RT-PCR experiments revealed the presence of transcripts for EP₄ and EP₁ receptors in mCCD_{cl} cells. In contrast, transcripts for

EP₂ or EP₃ receptors were not detected (Fig. 9 A). Moreover, transcripts for several enzymes (COX-1, COX-2, mPGES2, and cPGES) involved in the biochemical pathway of PGE₂ synthesis were detected not including mPGES1 (Fig. 9 B). Targets were confirmed by extracting and sequencing the PCR bands of interest.

Discussion

The present study provides evidence that basolateral, but not apical application of PGE₂ stimulates ENaC-dependent transepithelial sodium transport in mCCD_{cl} cells. Moreover, it demonstrates that this stimulatory effect is mediated by EP₄ receptor signaling and is due to increased ENaC activity resulting, at least in part, from enhanced channel expression at the apical cell surface. Finally, experiments with the PGE₂ precursor arachidonic acid indicate that mCCD_{cl} cells can synthesize and release PGE₂ to stimulate transepithelial sodium transport in an autocrine manner.

The sustained stimulatory effect of PGE₂ on ENaC-mediated I_{sc} in mCCD_{cl} cells is reminiscent of similar effects previously observed in *Xenopus laevis* A6 (Kokko et al., 1997; Matsumoto et al., 1997) and canine MDCK renal epithelial cells (Wegmann and Nüssing, 2003). This challenges the view that PGE₂ mainly inhibits sodium absorption in the collecting duct. Interestingly, an inhibitory effect of PGE₂ on sodium absorption was not observed in isolated perfused rat CCD (Chen et al., 1991). Moreover, the inhibitory effects of PGE₂ on Na⁺ transport in microdissected rabbit tubules (Stokes and Kokko, 1977; Iino and Imai, 1978) and primary cultures of rabbit principal CCD cells (Ling et al., 1992) were short-term, occurring within a few minutes of basolateral PGE₂ application. In contrast, the stimulatory effect observed in mCCD_{cl} cells in the present study reached a maximum within ~10 min and was sustained. In this context, it is of interest that a biphasic action of PGE₂ has been reported in *Xenopus* A6 renal epithelial cells with an acute inhibition (3–6 min) and a delayed stimulation (10–50 min) of ENaC by basolateral PGE₂ (Kokko et al., 1994, 1997; Worrell et al., 2001). The acute inhibition is probably caused by an increase in intracellular Ca²⁺ signaling. In contrast, the delayed stimulatory effect is due to an increase in intracellular cAMP leading to an increase in the number of channels expressed in the apical membrane (Kokko et al., 1994).

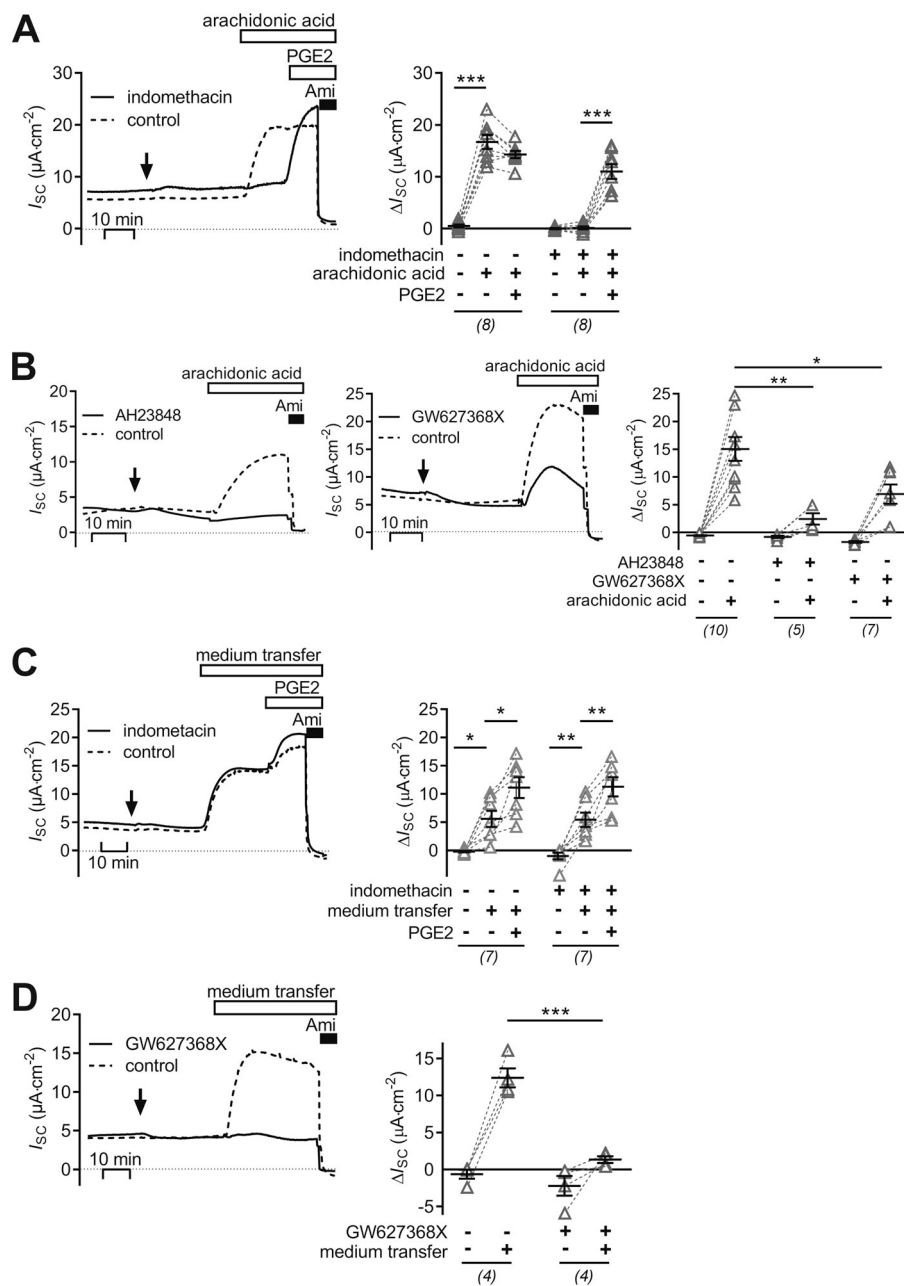


Figure 7. Stimulation of ENaC activity in mCCD_{cl1} cells by arachidonic acid is mediated via basolateral EP₄ receptors and constitutively active COX. The left panel in A and the left and middle panels in B show representative traces of continuous I_{sc} recordings. 50 μM indomethacin (A) or 100 μM AH23848 or 2 μM GW627368X (B; solid lines) was added basolaterally at the time point indicated by an arrow. Dashed lines indicate matched control recordings treated with the respective vehicle (control). About 30 min later cells were exposed to 50 μM apical arachidonic acid and in A followed by basolateral 100 nM PGE₂. Finally, apical 10 μM amiloride was added as indicated. Summary data (right panel) are presented as ΔI_{sc} values, which were determined by subtracting the corresponding baseline I_{sc} from the I_{sc} reached in the presence (+) or absence (-) of indomethacin, arachidonic acid, PGE₂, AH23848, and GW627368X as indicated. The left panels in C and D show representative traces of continuous I_{sc} recordings in which vehicle (control; dashed lines), 50 μM indomethacin, or 2 μM GW627368X (solid lines) was added basolaterally at the time point indicated by an arrow. Approximately 30 min later, medium from the basolateral compartment of a second set of cells (donor cells; not depicted) which were treated with apical 50 μM arachidonic acid was transferred to the basolateral compartment (indicated as medium transfer) in a 1:1 (vol/vol) ratio. This was followed in C by basolateral 100 nM PGE₂. Finally, as indicated, 10 μM amiloride was applied to all cells. Summary data (right panel) are presented as ΔI_{sc} values, which were determined by subtracting the corresponding baseline I_{sc} from the I_{sc} reached in the presence (+) or absence (-) of indomethacin, GW627368X, PGE₂, and medium transfer as indicated. Summary data are presented as individual values and their mean \pm SEM. Paired data are connected by dashed lines. Numbers of experiments are given in parentheses. ***, $P < 0.001$; **, $P < 0.01$; *, $P < 0.05$, one-way ANOVA with Tukey's multiple comparison test or unpaired Student's t test, where appropriate.

This latter interpretation is in good agreement with our observation in mCCD_{cl1} cells.

In a previous study, our group reported that PGE₂ stimulates Cl⁻ secretion in murine M-1 CCD cells (Sandrasagra et al., 2004). In this latter study, no stimulatory effect of PGE₂ on the amiloride-sensitive I_{sc} component was observed. A possible explanation for this discrepancy is that in M-1 cells, ENaC-dependent I_{sc} is usually maximally stimulated under baseline conditions, which makes it difficult to detect any additional stimulatory effects on ENaC. A prestimulation of ENaC by deoxycorticosterone acetate in the majority of experiments may be a reason why no stimulatory effect of 10–100 nM PGE₂ was detected in isolated rabbit collecting tubules (Stokes and Kokko, 1977; Iino and Imai, 1978). Moreover, in our previous study (Sandrasagra et al., 2004), we focused on the short-term effects

of PGE₂ and may have overlooked a delayed stimulatory response. In mCCD_{cl1} cells the sustained I_{sc} increase in response to PGE₂ is clearly due to a stimulatory effect on ENaC. This is evidenced by the finding that the response could not be elicited in the presence of amiloride and that the PGE₂-stimulated I_{sc} was completely inhibited by amiloride. In contrast, a minor rapid and transient I_{sc} peak preceding the sustained stimulatory response to PGE₂ was preserved in the presence of amiloride or in the presence of the EP₄ receptor antagonist GW627368X. This suggests that the transient peak response is not mediated by EP₄ and reflects electrogenic Cl⁻ secretion reminiscent of the Cl⁻ secretory response observed in M-1 cells (Sandrasagra et al., 2004). Thus, under certain conditions, mCCD_{cl1} cells, like M-1 cells and mouse inner medullary collecting duct cells (Rajagopal et al., 2014), may have the ability to secrete Cl⁻ in response to

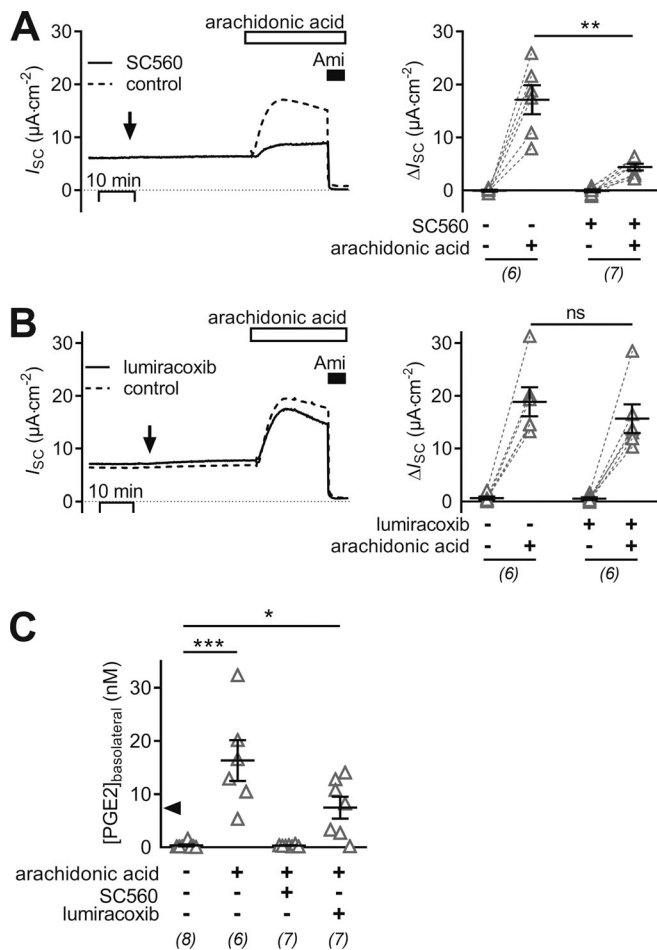


Figure 8. Stimulation of ENaC in mCCD_{cl1} cells by arachidonic acid requires COX-1 activity. (A and B) Representative traces of continuous I_{sc} recordings from mCCD_{cl1} cells are shown in the left panels, and data from similar experiments are summarized in the corresponding right panels. At the time point indicated by an arrow, cells were exposed to 0.14 μM SC560 (COX-1 selective inhibitor, solid line) or vehicle in matched control recordings (A; control, dashed line) or 2 μM lumiracoxib (COX-2 selective inhibitor, solid line) or vehicle in matched control recordings (B; control, dashed line). Approximately 30 min later, all cells were exposed to 50 μM apical arachidonic acid and subsequently to apical amiloride (Ami; 10 μM) as indicated. Summary data (right panel) are presented as ΔI_{sc} values, which were determined by subtracting the corresponding baseline I_{sc} from the I_{sc} reached in the presence (+) or absence (-) of SC560, lumiracoxib, and arachidonic acid as indicated. (C) Concentration of PGE₂ measured in basolateral cell culture medium collected from cells in the presence (+) or absence (-) of 50 μM arachidonic acid, 0.14 μM SC560 and 2 μM lumiracoxib as indicated. The solid arrowhead indicates the EC₅₀ value for the I_{sc} response upon PGE₂ as determined in Fig. 1. Data are presented as individual values and their mean \pm SEM. Numbers of experiments are given in parentheses. ***, $P < 0.001$; **, $P < 0.01$; *, $P < 0.05$; ns, $P \geq 0.05$, one-way ANOVA with Dunn's multiple comparison test.

PGE₂. However, in mCCD_{cl1} cells, the predominant effect of PGE₂ is a substantial and sustained stimulation of ENaC-dependent Na⁺ absorption. Importantly, the estimated EC₅₀ of ~7.1 nM for this stimulatory PGE₂ effect is within a range that may be relevant in vivo (Bonvalet et al., 1987).

The present study provides pharmacological evidence that the stimulatory effect of PGE₂ on ENaC is mediated by EP₄ receptors in the basolateral membrane of mCCD_{cl1} cells. This

conclusion is further supported by the finding that PGE₂ caused an increase in intracellular cAMP as expected for a G_{as}-coupled receptor. In agreement with the functional data, transcripts for EP₄ receptor were detected in mCCD_{cl1} cells consistent with previous reports of EP₄ receptor expression in the collecting duct (Jensen et al., 2001; Hao and Breyer, 2008). In contrast, transcripts for EP₂ or EP₃ receptors were not detected in mCCD_{cl1} cells. The absence of EP₂ receptor transcripts is plausible, because renal EP₂ receptor expression is thought to be limited to interstitial cells and the vasculature (Li et al., 2017). The lack of EP₃ receptor expression in mCCD_{cl1} cells was unexpected and may not reflect the situation in the native collecting duct, where EP₃ receptor is expressed, but its cellular localization and precise physiological role is less clear (Hao and Breyer, 2008). Of interest is the detection of EP₁ receptor transcripts in mCCD_{cl1} cells consistent with the reported expression of EP₁ receptor in the collecting duct (Guan et al., 1998) and M-1 cells (Sandrasagra et al., 2004). EP₁ receptor is a G_{aq/11}-coupled receptor, and its activation leads to a rise in cytosolic Ca²⁺ via the IP₃/diacylglycerol pathway (Narumiya et al., 1999; Breyer and Breyer, 2000b). An EP₁-mediated increase in intracellular calcium is thought to be the mechanism by which PGE₂ inhibits sodium absorption in rabbit and mouse CCD (Hébert et al., 1991; Guan et al., 1998; Nasrallah et al., 2018). Thus, whether PGE₂ inhibits or stimulates sodium absorption in the collecting duct may depend on the relative expression of EP₁ versus EP₄ receptors which may vary in different parts of the collecting duct according to physiological needs. It has been speculated that in M-1 cells the initial Cl⁻ secretory response to PGE₂ is mediated through EP₁ receptor activation and subsequent stimulation of Ca²⁺-activated chloride channels (Sandrasagra et al., 2004). This may also be the mechanism underlying the small ENaC-independent transient I_{sc} response to PGE₂ occasionally observed in mCCD_{cl1} cells. Presently, it is unclear whether Cl⁻ secretion is physiologically relevant in CCD cells or becomes relevant under pathophysiological conditions like in polycystic kidney disease, where it is thought to contribute to cyst enlargement (Liu et al., 2012; Blanco and Wallace, 2013). In contrast, it is well established that ENaC function in the CCD and its appropriate regulation play a key role in maintaining sodium homeostasis. Thus, the reported stimulatory effect of PGE₂ on ENaC-mediated transepithelial sodium transport is potentially important under certain physiological and pathophysiological conditions.

PGE₂ is the most abundant prostanoid detected in the kidney, but the physiologically relevant sites for the synthesis of PGE₂ remain to be defined (Hao and Breyer, 2008). The present study provides evidence that mCCD_{cl1} cells can synthesize and secrete PGE₂. Indeed, apical exposure to the PGE₂ precursor arachidonic acid caused substantial secretion of PGE₂, which reached a concentration of ~16 nM in the basolateral culture medium of treated cells compared with ~0.3 nM in the basolateral medium of control cells. From the PGE₂ concentration reached in the basolateral medium of treated cells it can be estimated that confluent mCCD_{cl1} cells, covering one tissue culture insert, secrete ~4 μg PGE₂ over a 30-min period. This corresponds to a secretion rate of ~79 fg PGE₂/ng cellular protein per 30 min. It is conceivable that this secretion rate may be achieved in vivo,

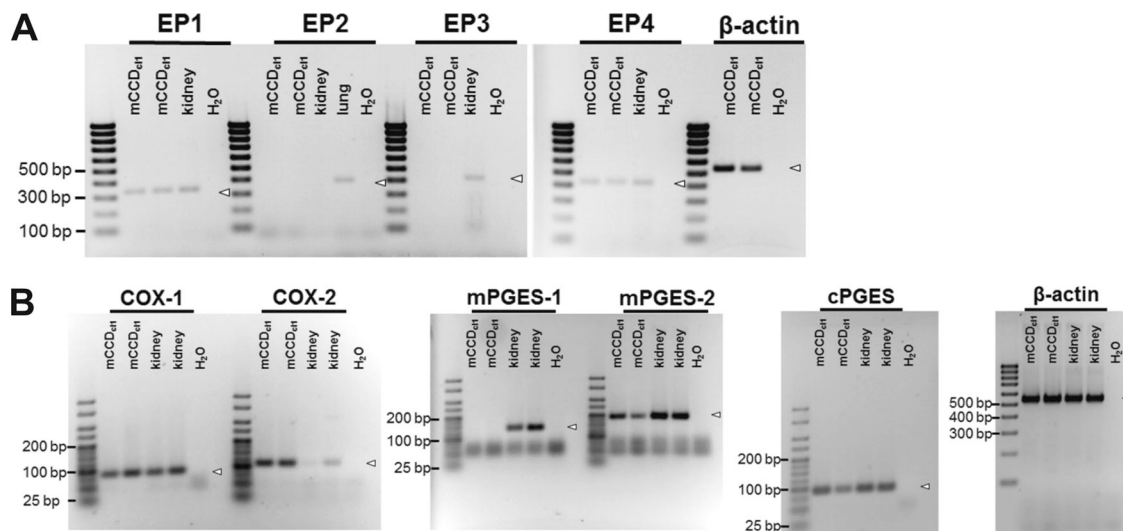


Figure 9. Transcripts for EP receptors and for enzymes of the PGE₂ biosynthetic pathway were detected in mCCD_{cl1} cells. Specific primers were used to detect the transcript expression of the prostanoid receptors EP₁–EP₄ (A), as well as key enzymes involved in PGE₂ synthesis, namely COX-1, COX-2, mPGES1, mPGES2, and cPGES (B). Transcript expression was determined in RNA extracted from mCCD_{cl1} cells, as well as from mouse cortical kidney and lung homogenates. H₂O and primers recognizing β-actin served as no-template and loading control, respectively. Expected individual transcript sizes are indicated by open arrowheads.

because synthesis rates ranging from ~800 to >8,000 fg·ng protein⁻¹·30 min⁻¹ have been reported for PGE₂ synthesis in the collecting duct (Bonvalet et al., 1987; Liu et al., 2014). In light of the EC₅₀ value of ~7.1 nM, the concentration of PGE₂ of ~16 nM explains the robust stimulatory effect observed in medium transfer experiments when receiver cells were exposed to a 1:1 dilution of the basolateral medium from donor cells. It also explains why the stimulatory effect was submaximal and why subsequent exposure to 100 nM PGE₂ further stimulated I_{SC} in these experiments (Fig. 7 C). In contrast, in mCCD_{cl1} cells exposed to arachidonic acid on the apical side subsequent exposure to 100 nM PGE₂ on the basolateral side had no additional stimulatory effect (Fig. 7 A). This is plausible, because the local PGE₂ concentration reached within the vicinity of the basolateral membrane of mCCD_{cl1} cells exposed to apical arachidonic acid is probably much higher than the ~16 nM measured in the bulk medium. Thus, autocrine secretion of PGE₂ may well be sufficient to achieve a maximal EP₄ receptor-mediated stimulatory effect on the amiloride-sensitive I_{SC} (Fig. 7).

In the renal cortex, COX-1 and downstream prostaglandin synthases are constitutively expressed, while basal expression of COX-2 is low (Yang et al., 1998; Murakami et al., 2002; Tanikawa et al., 2002; Hao and Breyer, 2008). This study provides evidence that transcripts for COX-1, mPGES2, and cPGES, but not mPGES1, are expressed in mCCD_{cl1} cells. Once arachidonic acid is processed by COX-1, the biosynthesis of PGE₂ can be achieved by mPGES2 or cPGES. The stimulatory effect of arachidonic acid on ENaC-dependent I_{SC} was abolished in mCCD_{cl1} cells treated with the nonselective COX-1/2 inhibitor indomethacin or the selective COX-1 inhibitor SC560. In contrast, the selective COX-2 inhibitor lumiracoxib did not prevent the stimulatory effect of arachidonic acid. These findings indicate that COX-1 plays a major role in PGE₂ synthesis in mCCD_{cl1} cells, which is consistent

with the finding that COX-1 is the primary isoform in native CCD (Vitzthum et al., 2002) with a preferential localization in CCD principal cells (Câmpeanu et al., 2003).

cPGES transcripts are ubiquitously expressed in mouse epithelial cells of the connecting tubule and collecting duct (Chen et al., 2017; Ransick et al., 2019), and immunoreactive cPGES has been detected in cultured mouse inner medullary collecting duct cells (Zhang et al., 2003). Therefore, its detection in mCCD_{cl1} cells is not surprising. Interestingly, mPGES2 is thought to be primarily expressed in intercalated cells (Yang et al., 2006a), whereas mPGES1 has been detected in principal cells (Chen et al., 2017; Wang et al., 2018; Ransick et al., 2019). The mCCD_{cl1} cell line used in this study is a clonal cell line derived from microdissected mouse CCD and retains features typical for CCD principal cells (Gaeggeler et al., 2005, 2011; Gonzalez-Rodriguez et al., 2007; Mansley et al., 2015, 2018, 2019). Thus, at first sight it may appear surprising that transcripts for mPGES2 but not for mPGES1 were detected in mCCD_{cl1} cells. However, mPGES1 expression is induced by cytokines and inflammatory stimuli (Murakami et al., 2002; Hao and Breyer, 2008). Therefore, basal expression of mPGES1 in CCD principal cells is probably low and may have been below the detection limit in mCCD_{cl1} cells. Low basal expression of mPGES1 is also consistent with the finding that single-cell transcriptome analysis of major renal collecting duct cell types in mouse revealed a weaker expression of mPGES1 in CCD principal cells compared with cPGES (Chen et al., 2017). Interestingly, it has been reported that mCCD_{cl1} cells show some plasticity consistent with the ability to transition between principal and intercalated cells (Assmus et al., 2018). There is little doubt that the ENaC-dependent I_{SC} observed in the present study is generated by mCCD_{cl1} cells that predominantly behave like principal cells. However, at present it is unclear whether PGE₂ synthesis and

secretion occurs in the same cells responsible for the amiloride-sensitive I_{SC} or whether this occurs in distinct interspersed cells with an intercalated phenotype that may be present in the mCCD_{cl} monolayer. It is tempting to speculate that the latter scenario may provide a mechanism by which intercalated cells can regulate principal cell ion transport function in a paracrine manner. Whether this occurs in native tissue and is physiologically relevant remains to be determined.

The biotinylation experiments performed in this study indicate that PGE₂ stimulates ENaC activity at least in part by increasing the abundance of ENaC at the apical plasma membrane. This is consistent with the finding that PGE₂ via EP₄ receptor increases cytosolic cAMP in mCCD_{cl} cells, as increased surface abundance of ENaC has previously been observed in response to an elevated cytosolic cAMP concentration (Snyder, 2000; Butterworth et al., 2005). Indeed, it is well known that vasopressin can stimulate ENaC-mediated sodium transport (Schafer and Troutman, 1990; Ecelbarger et al., 2000; Ecelbarger et al., 2001; Nicco et al., 2001) via the G_{as}-coupled V₂-receptor and subsequent activation of the AC/cAMP/PKA pathway (Schafer and Hawk, 1992; Loffing and Korbmaier, 2009; Roos et al., 2013). This involves an increased abundance of the β - and γ -ENaC subunits at the apical plasma membrane, possibly due to the insertion of ENaC-containing vesicles into the plasma membrane from a subapical pool (Snyder, 2000; Butterworth et al., 2005, 2012). Additional mechanisms probably contribute to stimulate ENaC activity following the activation of the AC/cAMP/PKA pathway (e.g., phosphorylation events; Yang et al., 2006b) and an increase in channel open probability (Bugaj et al., 2009). Thus, the stimulatory effect of PGE₂ on ENaC, like that of vasopressin, is probably more complex than simply increasing channel surface expression (Kortenoeven et al., 2015).

The conclusion that the PGE₂ effect on ENaC is mediated by an activation of the AC/cAMP/PKA pathway is further supported by the observation that the effects of PGE₂ and vasopressin were nonadditive in mCCD_{cl} cells. This indicates that the signaling pathways of PGE₂ and vasopressin converge. Importantly, the stimulatory effect of PGE₂ on ENaC activity was preserved when mCCD_{cl} cells were prestimulated with aldosterone. In contrast to the rapid effects of PGE₂ and vasopressin, the stimulatory effect of aldosterone was much slower. This slower time course of the aldosterone response is consistent with observations in isolated perfused rat CCD (Reif et al., 1986). It is not surprising, because the response to aldosterone is mediated by the mineralocorticoid receptor and involves highly complex regulatory mechanisms that are not yet fully understood but include transcriptional regulation of the channel and of regulatory proteins (Loffing and Korbmaier, 2009; Rossier, 2014).

It has been postulated that the well-documented synergistic stimulation of sodium transport by vasopressin and aldosterone (Tomita et al., 1985; Reif et al., 1986; Chen et al., 1990, 1991; Schafer and Hawk, 1992; Snyder et al., 2004; Bugaj et al., 2009) is important to achieve maximal sodium reabsorption and urine concentration due to the simultaneous stimulation of water permeability by vasopressin (Reif et al., 1986). At present, the role of PGE₂ in regulating sodium and water transport in the

distal nephron and collecting duct is less clear. It is commonly thought that prostanoids synthesized along the renal tubular system cause natriuresis by reducing medullary sodium and water reabsorption. This may be particularly relevant under conditions of elevated salt intake (Hao and Breyer, 2007) or when tubular fluid flow is high (Flores et al., 2012). Inhibition of sodium and water reabsorption by prostanoids may be due to various mechanisms such as their ability to blunt the action of vasopressin, promote medullary blood flow, and directly inhibit sodium transport in the distal nephron by reducing the activity of the Na⁺/K⁺-ATPase or ENaC (Stokes and Kokko, 1977; Hébert et al., 1990; Zeidel, 1993; Guan et al., 1998; Hao and Breyer, 2007; Flores et al., 2012). On the other hand, PGE₂ increased water flux and elevated sodium absorption in primary cultures of freshly isolated tubules (Canessa and Schafer, 1992; Wang et al., 2016). Interestingly, the expression pattern of enzymes and receptors associated with prostanoid signaling may be altered under conditions of low salt intake or volume contraction (Hao and Breyer, 2008; Li et al., 2017; Wang et al., 2018). For example, in rabbits, salt restriction markedly stimulated PGE₂ biosynthesis in the outer medulla and cortex (Davila et al., 1978; Stahl et al., 1979). This may indicate that PGE₂ is needed to minimize renal sodium excretion. Thus, there is evidence that the effect of PGE₂ on transepithelial sodium and water flux depends on the physiological or pathophysiological setting. A different responsiveness to prostanoids may also explain why indomethacin decreased blood pressure in animals on a low-sodium diet but increased blood pressure in sodium-repleted rats (Stahl et al., 1981).

In conclusion, our results demonstrate that PGE₂ can stimulate ENaC in mCCD_{cl} cells, which depends on EP₄ receptor activation and a rise in cytosolic cAMP. Furthermore, these cells can synthesize and secrete PGE₂, which acts in an autocrine/paracrine manner to stimulate ENaC-mediated sodium transport. The (patho-)physiological implications of these findings remain to be elucidated. However, the findings suggest that under certain conditions, locally generated PGE₂ may stimulate sodium absorption in the ASDN. Conversely, it is conceivable that pharmacological inhibition of PGE₂ synthesis may attenuate sodium reabsorption in the ASDN in states with increased local production of PGE₂. Additional studies are needed in native tissue and genetically modified animal models to explore a possible dual regulatory role of PGE₂ associated with its ability to inhibit or stimulate sodium absorption in collecting duct cells.

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