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Depolarization and decreased surface expression of K⁺ channels contribute to NSAID-inhibition of intestinal restitution

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ARTICLE INFO

Article history: Received 4 January 2007 Accepted 26 March 2007

Keywords:
Non-steroidal anti-inflammatory
drugs
Intestinal epithelial cells
Membrane potential
Potassium channels

ABSTRACT

Non-steroidal anti-inflammatory drugs (NSAIDs) contribute to gastrointestinal ulcer formation by inhibiting epithelial cell migration and mucosal restitution; however, the drugaffected signaling pathways are poorly defined. We investigated whether NSAID inhibition of intestinal epithelial migration is associated with depletion of intracellular polyamines, depolarization of membrane potential (E_m) and altered surface expression of K⁺ channels. Epithelial cell migration in response to the wounding of confluent IEC-6 and IEC-Cdx2 monolayers was reduced by indomethacin (100 μM), phenylbutazone (100 μM) and NS-398 (100 μM) but not by SC-560 (1 μM). NSAID-inhibition of intestinal cell migration was not associated with depletion of intracellular polyamines. Treatment of IEC-6 and IEC-Cdx2 cells with indomethacin, phenylbutazone and NS-398 induced significant depolarization of E_{m} , whereas treatment with SC-560 had no effect on $E_{\rm m}$. The $E_{\rm m}$ of IEC-Cdx2 cells was: -38.5 ± 1.8 mV under control conditions; -35.9 ± 1.6 mV after treatment with SC-560; $-18.8 \pm 1.2 \text{ mV}$ after treatment with indomethacin; and $-23.7 \pm 1.4 \text{ mV}$ after treatment with NS-398. Whereas SC-560 had no significant effects on the total cellular expression of $K_v1.4$ channel protein, indomethacin and NS-398 decreased not only the total cellular expression of $K_v1.4$, but also the cell surface expression of both $K_v1.4$ and $K_v1.6$ channel subunits in IEG-Cdx2. Both K_v1.4 and K_v1.6 channel proteins were immunoprecipitated by K_v 1.4 antibody from IEC-Cdx2 lysates, indicating that these subunits co-assemble to form heteromeric K_v channels. These results suggest that NSAID inhibition of epithelial cell migration is independent of polyamine-depletion, and is associated with depolarization of $E_{\rm m}$ and decreased surface expression of heteromeric $K_{\rm v}1$ channels.

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1. Introduction

Non-steroidal anti-inflammatory drugs (NSAIDs) are among the most commonly used drugs worldwide despite their welldocumented gastrointestinal (GI) toxicity. Adverse GI effects of NSAIDs including GI ulceration [1–4] result in 100,000 emergency admissions, high medical costs and 17,500 deaths per year in the US alone [5,6]. Despite extensive investigation,

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the mechanisms responsible for NSAID-associated GI damage are incompletely understood. A growing body of data suggests that inhibition of mucosal cyclooxygenase (COX) and loss of protective prostaglandins (PG) are insufficient to explain NSAID enteropathies [3]. Moreover, the recently recognized untoward cardiac effects of COX-2 inhibitors suggest that the development of less ulcerogenic NSAIDs through manipulation of COX-selectivity is not a viable therapeutic strategy [7,8].

To date, NSAIDs have been hypothesized to diminish normal mucosal defense mechanisms and to promote ulcer formation not only by inhibiting cyclooxygenase (COX), but also by diminishing nitric oxide synthesis, uncoupling oxidative phosphorylation, decreasing intracellular pH, reducing surface hydrophobicity, and inducing apoptosis [3,9–13]. It has also been shown that NSAIDs contribute to ulcer formation and delay ulcer healing by inhibiting intestinal epithelial cell migration and mucosal restitution [14–16].

With respect to the latter mechanism, the epithelial cells of the GI mucosa normally form a protective barrier to pathogens and toxins that may be present in the gut lumen. These surface epithelial cells are continuously exposed to noxious agents that may cause mucosal injury. Superficial mucosal defects typically are repaired rapidly by migration of epithelial cells into the defect from the periphery of the wound. This process, called mucosal restitution, is a primary repair modality in the GI tract and is critical to preserving mucosal integrity and to suppressing ulceration [17-20]. Restitution occurs within minutes to hours following injury in vivo or in vitro, because this process requires cell migration but not cell proliferation or differentiation [17-20]. While it is established that NSAIDs promote ulceration by interfering with restitution [14–16], the signaling pathways mediating this effect have not been fully elucidated.

In contrast to the dearth of information regarding the effects of NSAIDs on intestinal restitution, extensive knowledge of the molecular basis for polyamine-modulation of this process exists [21–25]. Depletion of intracellular polyamines inhibits intestinal cell migration by decreasing the expression of delayed rectifier potassium channels, thereby depolarizing membrane potential (Em), decreasing the driving force for calcium-influx, and inhibiting downstream [Ca²⁺]_i-dependent signaling events [21-25]. NSAIDs have been shown to deplete the polyamine content of immortalized colon cancer cells by enhancing polyamine catabolism [26]. However, to our knowledge, there have been no experimental efforts to link NSAID inhibition of intestinal epithelial migration to depletion of intracellular polyamines, depolarization of membrane potential or surface expression of K+ channels. The present investigation was designed to identify potential mechanisms for the GI toxicity of NSAIDs by defining those relationships.

2. Materials and methods

2.1. Reagents

Cell culture media was obtained from American Type Culture Collection (ATCC, Manassas, VA) or JRH Biosciences. Serum was obtained from Invitrogen (Calsbad, CA) and chemicals were obtained from Sigma (St. Louis, MO) unless stated otherwise. Matrigel was purchased from BD Biosciences (Bedford, MA). DiBAC₄(3) was obtained from Molecular Probes (Invitrogen, Carlsbad CA). SC-560 (5-(4-chlorphenyl)-1-(4-methoxyphenyl)-3-(trifluoromethyl)-1 H-pyrazole) and NS-398 (N-[2-(cyclohexyloxy)-4-nitrophenyl]-methanesulfonamide) were obtained from Cayman Chemical (Ann Arbor, Ml). AVE0118, (12'(benzyloxycarbonylaminomethyl) biphenyl-2-carboxylic acid 2-(2-pyridyl)ethylamide) was obtained from Aventis Pharma (Frankfort, Germany).

2.2. Cell culture

The IEC-6 cell line was purchased from ATCC at passage 13 (Manassas, VA). These cells, developed by Quaroni et al. [27], are non-tumorigenic and retain the character of undifferentiated intestinal crypt cells. Cell culture conditions were similar to those described previously by others [28,29]. The basic medium consisted of DMEM supplemented with heatinactivated fetal bovine serum (FBS, 5%), insulin (10 μ g/ml) and gentamicin (50 μ g/ml). Cells were maintained at 37 °C in a humidified atmosphere of 5% CO₂ in air. Passages 15–20 were used for the experiments described below.

A line of IEC-6 cells stably transfected with the Cdx2 transcription factor (IEC-Cdx2) was obtained from Dr. E. Suh (University of Pennsylvania, Philadelphia, PA). This cell line was developed and characterized initially by Suh and Traber [30], and then characterized further by others [23,31,32]. The forced expression of the Cdx2 gene in IEC-6 cells results in development of a differentiated phenotype [23,31]. The LacSwitch expression vector system (Stratgene, La Jolla, CA) is used to direct conditional expression of the Cdx2 gene, with isopropyl- β -D-thiogalactopyranoside (IPTG) serving as the inducer for gene expression. IEC-Cdx2 cell stocks were maintained in the same basic culture medium used for IEC-6, and grown in media supplemented with 4 mM IPTG for 16 days to induce the conditional expression of Cdx2 before experiments.

IEC-6 or IEC-Cdx2 cells were seeded at a density of approximately $6.25 \times 10^4 \, \text{cells/cm}^2$ on 35 or 60 mm plates thinly coated with Matrigel and were grown to a confluent monolayer prior to the introduction of media containing experimental treatments for: optical analysis of cell migration; western analysis of K+ channel expression; assay of intracellular polyamines; and sharp electrode measurement of $E_{\rm m}$. Cells were seeded in 96 well plates at a density of 65,000 cells per well for assessment of $E_{\rm m}$ by DiBAC₄(3) fluorescence.

Both IEC-6 and IEC-Cdx2 cells were used in our initial investigations of drug effects on cell migration and $E_{\rm m}.$ Subsequent experiments focused on elucidating the relationship between NSAID-treatment and $K_{\rm v}$ channel expression were performed using IEC-Cdx2, exclusively.

2.3. Treatment protocols

The experiments performed utilized NSAIDs with variable effects on intestinal restitution and variable COX-selectivity, as shown in Table 1 [33–36]. Drug effects were evaluated over the following ranges of concentrations, as indicated in the text or figure legends: indomethacin $(1–100 \mu M)$; phenylbutazone

Table 1				
Inhibitor	IC ₅₀ (μM) COX-1	IC ₅₀ (μM) COX-2	COX selectivity ^a	Ulcerogenic potential
Indomethacin (human)	0.19	0.44	0.4	+++
Indomethacin (horse)	0.036	0.43	0.08	+++
SC-560 (human)	0.005	1.4	0.003	0
NS-398 (rat)	125	5.6	22	(+) slows ulcer healing
Phenvlbutazone (horse)	6.15	3.79	1.6	+++

^a Ratio of IC_{50} for COX-1 to IC_{50} for COX-2 (positively correlated with COX-2 selectivity such that a ratio < 1 indicates COX-1 selectivity and a ratio > 1 indicates COX-2 selectivity). Based on references [33–36].

 $(1-100 \, \mu \text{M})$; NS-398 $(1-100 \, \mu \text{M})$, SC-560 $(10 \, \text{nM}-1 \, \mu \text{M})$. DMSO $(0.1\% \, \text{v/v})$ was the vehicle control used for comparison to NSAIDs. Monolayers were exposed to NSAIDs for 72 h prior to wounding, measurement of $E_{\rm m}$ or performance of biochemical assays, as indicated in the figure legends. In some experiments, an inhibitor of polyamine synthesis, α-difluoromethylornithine (DFMO, 5 mM), was added to experimental media [37]. AVE0118, $(10^{-6} \, \text{M})$ was employed as an antagonist of voltage-gated K^+ (K_{ν}) channels [38].

2.4. Assessment of cell migration

Treatment-induced changes in IEC-6 and IEC-Cdx2 cell migration were assessed using a wounding assay. A razor blade was used to remove approximately one-third of the monolayer after 72 h of exposure to media containing drug. The created cell border (scratch line) was marked. Micrographs (Nikon ACT-1) were captured immediately after creation of this defect and again 4 h later. Three methods of image analysis (Nova Prime, Bioquant, Nashville TN) were used to measure cell migration. First, the linear distance (μM) from the scratch line to the point of maximum cell migration was measured. Second, the number of cells migrating into the defect was counted and expressed in terms of cells (120,000 µm²). Finally, a standardized rectangular region of interest (120,000 µm²) was created and positioned at the scratch line; the percentage of this region occupied by migrating cells was measured over time. All experiments were done in triplicate and two fields per plate were examined and measured, as described previously.

2.5. Determination of polyamine content

IEC-6 cells were washed with chilled PBS containing protease inhibitors (P-8340, Sigma), mechanically lifted using a cell scrapper in 5% deoxycholate, transferred to cryotubes and snap frozen using liquid nitrogen, then thawed and sonicated for three pulses of 10 s in ice. These lysates were centrifuged for 15 min at 10,000 rpm (5415F, Eppendorf, Brinkman Inst., Westbury, NY). The supernatant was collected and stored at $-70\,^{\circ}\text{C}$ until analysis.

Polyamine content was measured by liquid chromatography/mass spectrometry (LC/MS). Samples, standards and quality controls (QCs) were prepared for LC/MS by combining 300 μl of cell lysate with 30 μl of internal standard (10 $\mu g/ml$ ethambutol), and then filtered (Whatman Mini-Uniprep, PVDF filter media, 45 μm pore). A volume of 50 μl of filtered sample was injected onto the LC/MS system. The system was comprised of a P4000 narrow-bore quaternary pump with a

vacuum degasser, AS3000 auto-sampler, UV6000 photo-diode array detector and LCQ_{Duo} ion-trap mass spectrometer (ThermoFinnigan, San Jose, CA). The HPLC column (Phenomenex, Torrance, CA) was a Luna C18(2) (30 mm \times 2.0 mm, 5 μ M). Mobile phase was 10/90-acetonitrile/acetic acid (10 mM); flow rate, 0.3 ml/min. The MS used electrospray ionization to detect each ion individually, 146 (spermidine), 203 (spermine) and 205 (ethambutol) m/z.

Stock solutions of putrescine, spermidine, spermine, and ethambutol were made in water. The stock solution of ethambutol was used to make a 10 $\mu g/ml$ solution in water, which was then used as an internal standard. Standards were made fresh each day that samples were analyzed. The QCs were made in bulk prior to validation and aliquoted into 300 μl samples and stored at $-20~^{\circ}\text{C}$ until used. For the initial intraday assay validation, five QCs were randomly selected from each of the three different concentrations. For assays, two QCs at each concentration were selected at random and prepared with the standard curve and unknown samples.

Peak area ratio of analyte:internal standard was used for quantitation. The standard curve was weighted 1/x, was linear, and contained the concentrations: 0.1, 0.25, 0.5, 1.0, 1.5, 2.0, and 2.5 μ g/ml (0.3, 0.75, 1.5, 3.0, 4.5, 6, and 7.5 μ g/ml for spermine). Concentration versus peak area ratio was plotted and a linear regression equation obtained for quantitation. Prepared concentrations for QCs were 0.75, 1.25, and 2.25 μ g/ml (2.25, 3.75, and 6.75 μ g/ml for spermine).

The validation assay was considered valid if the intra-day accuracy and precision values of the QCs were within 15% of the expected values for each of the three QC concentrations. Accuracy was calculated as %accuracy = $(C_{\rm int} - C_{\rm x})/C_{\rm int} \times 100$ and precision was calculated as %precision = $C_{\rm SD}/C_{\rm x} \times 100$, where $C_{\rm int}$ was the intended concentration, $C_{\rm x}$ was the average measured concentration, and $C_{\rm SD}$ was the standard deviation of the measured concentrations. An assay run was rejected if more than two of the six QCs for a given standard curve were >±15% from their intended value or if both QCs at a single concentration were >±15% from their intended concentration. A rejected run was re-assayed with freshly prepared samples.

2.6. Determination of membrane potential

Treatment-induced changes in $E_{\rm m}$ of IEC-6 and IEC-Cdx2 cells were assessed initially by spectrophotometric determination of DiBAC4(3) fluorescence (excitation = 485 nM; emission = 527 nM; Fluoroskan Ascent FL, LabSystems Inc., Helsinki, Finland). Cells were equilibrated with 2 μ M DiBAC₄(3) for 30 min at 37 °C prior to fluorescence measurement.

In addition, the $E_{\rm m}$ of individual IEC-Cdx2 cells was measured using microelectrodes (35–55 M Ω) filled with 3 M KCl. The extracellular solution contained 132 mM NaCl, 4 mM KCl, 1 mM MgCl₂, 2 mM CaCl₂, 10 mM Dextrose, 20 mM Hepes (pH 7.4). Data acquisition was performed using an Axon Instruments 700B amplifier and multiclamp commander (Molecular Devices, Sunnyvale, CA). Recordings were made at room temperature.

2.7. Immunoblotting

Western blot analysis was used to determine the effects of NSAIDs and DFMO on K^+ channel expression. Whole cell lysates or crude membrane fractions from passages 16, 17, and 18 were resolved by SDS-PAGE, transferred to nitrocellulose membranes and probed with polyclonal primary antibodies (Alomone Laboratories, Jerusalem, Israel) directed against $K_v1.1$ (APC-009), $K_v1.4$ (APC-007), $K_v1.5$ (APC-004), $K_v1.6$ (APC-003) and $K_v10.1$ (APC-104).

Bound primary antibodies were visualized using an enhanced chemiluminescence detection system (Super-Signal, Pierce, Rockford, IL), and recorded on radiographic film. Densitometric analysis was performed using AlphaEaseFC (Alpha Innotech Co, San Leandro, CA). Equal loading of protein was confirmed by immunoblotting for actin. The detection system yielded a linear response over a ~1:20 concentration range when immunoblotting was performed using cell lysates that had been serially diluted. Actin was considered an appropriate loading control because its level of expression in IEC-6 cells is unaffected by interventions that disrupt cell migration, such as polyamine depletion [22,39]. Preliminary immunoblotting experiments (not shown) confirmed that the ratio of actin to Na+/K+ATPase in IEC-6 cell lines was not significantly affected by NSAID treatments, thereby supporting the choice of actin as the loading control in experiments designed to investigate the expression of membrane proteins.

2.8. Immunoprecipitation

To preclear non-specific binding, protein G-agarose beads (Immunopure, Pierce, Rockford, IL) were incubated with samples for 2 h at 4 °C with continuous gentle agitation, then pelleted by centrifugation (700 \times q). A monoclonal antibody directed against K_v1.4 (Sigma K1264) was added subsequently to the supernatant and mixed overnight at 4 °C. Immune complexes were then immobilized onto protein G-agarose beads, washed three times with cold RIPA buffer and eluted from the beads with SDS sample buffer containing 5% βmercaptoethanol. Immunoprecipitated proteins and supernatant fractions were analyzed by immunoblotting, as described above. The ability of the Sigma monoclonal antibody to pull down K_v1.4 was confirmed in preliminary experiments by immunoblotting the precipitated fraction with the Alomone polyclonal antibody K_v1.4; a protein band with the expected mobility of monomeric K_v1.4 (~96 kDa) was detected (data not shown). Attempts to pull-down K_v1.6 using commercially available polyclonal antibodies directed against that target were unsuccessful.

2.9. Biotinylation

IEC-Cdx2 cells were incubated for 45 min at room temperature with EZ-Link sulfo-NHS-biotin (Pierce, Rockford IL) in PBS, lysed in a buffer containing 150 mM NaCl, 1 mM EDTA, 1% Nonidet P-40, 50 mM Tris–HCI, 0.5% sodium deoxycholate, 0.1% SDS, and 1% protease inhibitor cocktail, combined with 200 μl ImmunoPure Immobilized streptavidin (Pierce) and rocked overnight at 4 °C. Biotinylated proteins were pelleted by centrifugation, washed and eluted using Laemmli Sample Buffer. The starting material, supernatant, and biotinylated fractions were subsequently subjected to SDS-PAGE and immunoblot analysis as described above. Actin was used to normalize for loading. Relative cell surface expression was calculated by dividing the normalized surface protein

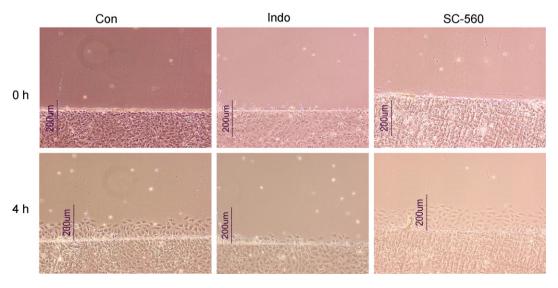


Fig. 1 – IEC-6 migration in monolayers grown in the absence (Con) or presence of NSAIDs. Drug-treated cells were exposed to either indomethacin (Indo, 100 μ M) or SC-560 (1 μ M) for 72 h before a defect was created in the monolayer (wounding). Images were captured immediately (0 h) or 4 h after wounding.

(biotinylated fraction) by the normalized total protein (biotinylated + non-biotinylated).

2.10. Statistical analysis

Data are expressed as mean \pm S.E.M. Significant differences between treatment groups were identified by ANOVA and multiple comparisons were made using LSD procedure (Statistix, Analytical Software, Tallahassee, FL). Differences were considered to be significant when $p \leq 0.05$. The numbers of replicates per treatment group and independent experiments associated with specific studies are provided in the figures or accompanying legends.

3. Results

3.1. NSAIDs and epithelial cell migration

Exposure to NSAIDs significantly reduced cell migration in response to the wounding of confluent IEC-6 and IEC-Cdx2 monolayers (Figs. 1 and 2). Maximum cell migration, measured as linear distance from the created defect 4 h after wounding, was significantly reduced in monolayers exposed to DFMO (5 mM), indomethacin (100 μ M), phenylbutazone (100 nM), and NS-398 (100 μ M) but not SC-560 (1 μ M). At these concentrations, the NSAIDs lack cytotoxic effects [40–43]. Similar results were obtained when cell migration was assessed in terms of either the number of cells in a defined (120,000 μ m²) region of interest adjacent to the created wound or the percentage of this defined region covered by cells (data not shown). As expected, results obtained using these three assessment methods were highly correlated (r^2 = 0.84).

3.2. NSAIDs and depletion of intracellular polyamines

The determination of intracellular polyamine content yielded expected results for IEC-6 treated with DFMO with and without spermadine [29,31]. As shown in Fig. 3A, exposure to DFMO significantly decreased the intracellular polyamine content of IEC-6 cells. Polyamine depletion could be counteracted by the addition of exogenous spermidine. These data support the integrity of our method for assaying cellular polyamine content.

Treatment of IEC-6 monolayers with the NSAIDs indomethacin, phenylbutazone and SC-560 failed to significantly decrease intracellular polyamine content, while NS-398 decreased spermine but not spermidine content (Fig. 3B). These data suggest NSAID-inhibition of intestinal cell migration cannot be attributed to depletion of intracellular polyamines.

3.3. NSAIDs and epithelial cell resting potential (E_m)

Treatment of IEC-6 and IEC-Cdx2 cells with DFMO to induce depletion of intracellular polyamines increased the DiBAC₄(3) fluorescence of treated cells, indicating depolarization of $E_{\rm m}$ (Fig. 4). These data are consistent with published data showing that depletion of polyamines by a 96 h exposure to DFMO induces a \sim +10 mV change in the $E_{\rm m}$ in IEC-6 cells measured by

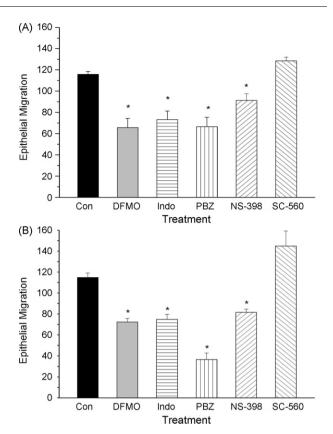


Fig. 2 – Epithelial cell migration in IEC-6 (panel A) and IEC-Cdx2 (panel B) monolayers (n = 6 per group) under control conditions (Con), after depletion of polyamines (DFMO, 5.0 mM), or after treatment with the NSAIDs indomethacin (Indo, 100 μ M), phenylbutazone (PBZ, 100 μ M), SC-560 (1 μ M) and NS-398 (100 μ M1). Monolayer cultures were exposed to drug for 72 h before wounding. Migration was measured 4 h post-wounding as the maximum linear distance from the created defect. Asterisks indicate significant difference from control (p < 0.05).

microelectrode techniques [25]. Thus, fluorescence imaging of DIBAC₄(3)-loaded cells is a useful method for detecting treatment-induced changes in the $E_{\rm m}$ of intestinal epithelial cells

Interestingly, treatment of IEC-6 and IEC-Cdx2 monolayers with indomethacin, phenylbutazone and NS-398, the NSAIDs previously shown to inhibit cell migration after wounding, also induced a significant increase in DiBAC4(3) fluorescence consistent with depolarization of E_m. Moreover, exposure to SC-560, an NSAID with no significant effects on epithelial cell migration, caused no significant change in DiBAC₄(3) fluorescence. Microelectrode measurements of E_m in IEC-Cdx2 confirmed that NSAID-inhibition of epithelial cell migration is correlated with drug-induced membrane depolarization. The $E_{\rm m}$ of cells treated with SC-560 (–35.9 \pm 1.6 mV, n = 8) was not significantly different from that of untreated IEC-Cdx2 cells ($-38.5 \pm 1.8 \,\text{mV}$, n = 19). In contrast, the $E_{\rm m}$ of indomethacin-treated cells (-18.8 \pm 1.2 mV, n = 6) and the $E_{\rm m}$ of NS-398-treated cells (-23.7 ± 1.4 mV, n = 15) were significantly depolarized.

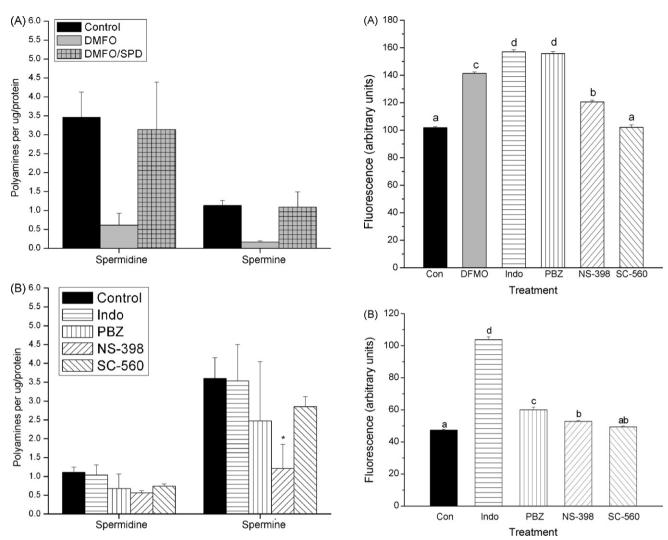


Fig. 3 – Gellular polyamine concentrations of IEG-6 exposed to NSAIDs. Panel A: polyamine content of IEG-6 cells treated with DFMO (5 mM) in the absence and presence of exogenous spermadine (SPD, 5 μ M). Panel B: IEG-6 treated with indomethacin (Indo, 100 μ M), phenylbutazone (PBZ, 100 μ M), SG-560 (1 μ M) and NS-398 (100 μ M). Cellular polyamine concentrations were measured by LG-MS at 78 h post-treatment (n = 3). Asterisks indicate significant difference from control (p < 0.05).

Fig. 4 – Relative $E_{\rm m}$ of IEC-6 (panel A) and IEC-Cdx2 (panel B) cells under control conditions (Con, n=86), after depletion of polyamines (DFMO, 5.0 mM, n=22), or after 72 h treatment with the NSAIDs indomethacin (Indo, 100 μ M, n=23), phenylbutazone (PBZ, 100 μ M, n=39), SC-560 (1 μ M, n=44) and NS-398 (100 μ M, n=22). Treatmentinduced changes in $E_{\rm m}$ were detected by monitoring the fluorescence of cells loaded with DiBAC₄(3). Increased fluorescence indicates depolarization of $E_{\rm m}$. Different superscripts indicate significant difference between groups (p<0.05).

3.4. NSAIDs and expression of K_{ν} channels

The $E_{\rm m}$ of IEC-6 and IEC-Cdx2 cells is related directly to the activity and number of K^+ channels expressed on the plasma membrane. Depolarization of $E_{\rm m}$ and inhibition of cell migration can be induced by antagonizing membrane K^+ channels [25]. Conversely, increased K^+ channel activity induces hyperpolarization and enhances the rate of epithelial cell migration [23].

It has been suggested that the E_m and migration response of IEC-6 and IEC-Cdx2 cells are determined primarily by the activity of delayed rectifier K^+ channels formed by K_v 1 (KCNA) gene products [21,23,25]. Consistent with that hypothesis, the K_v channel antagonist AVE0118 (1 μ M) impaired IEC-Cdx2 cell

migration post-wounding (Fig. 5). The inhibition of epithelial restitution by AVE0118 was comparable to that seen with the NSAIDs indomethacin, phenylbutazone and NS-398. On the basis of these findings, we hypothesized that NSAIDs depolarize $E_{\rm m}$ and inhibit intestinal epithelial cell migration by attenuating the activity of $K_{\rm v}1$ channels. Accordingly, we used immunoblotting in combination with biotinylation to determine the effects of NSAID treatment on the total expression and the cell surface expression of the $K_{\rm v}1$ channel proteins: $K_{\rm v}1.1$, $K_{\rm v}1.4$, $K_{\rm v}1.5$ and $K_{\rm v}1.6$; mRNA for these K^+ channel pore-forming subunits has been reported to be expressed by either IEC-6, IEC-Cdx2 cells or both [21,23,25].

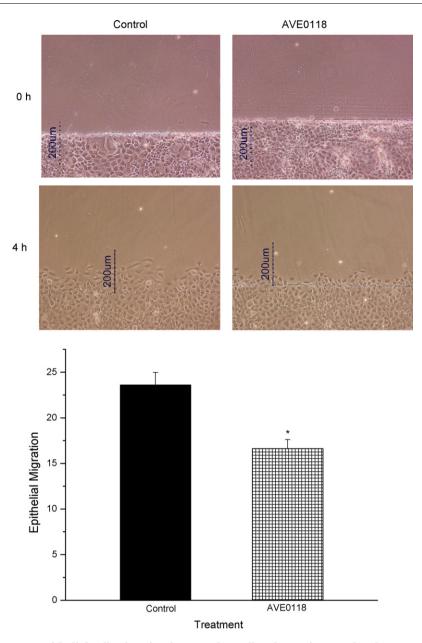


Fig. 5 – Effect of AVE0018 on epithelial cell migration in IEC-Cdx2 cells. Photomicrographs show control (left column) and drug-treated (right column) monolayers immediately (top row) and 4 h (lower row) after wounding. In the graph, epithelial migration is expressed as the percentage of a defined region adjacent to the defect that was occupied by migrating cells 4 h after monolayer wounding in the absence (Con, n = 6) and presence of the K_v channel antagonist AVE0118 (1 μ M, n = 6). Asterisk indicates significant difference (p < 0.05).

 $K_{\rm v}1.4,~K_{\rm v}1.5$ and $K_{\rm v}1.6$ channel proteins were detected routinely by immunoblotting of IEC-Cdx2 cell lysates, whereas $K_{\rm v}1.1$ was not found consistently. $K_{\rm v}1.4$ and $K_{\rm v}1.6$ but not $K_{\rm v}1.5$ were detected in the biotinylated fraction after surface biotinylation of IEC-Cdx2 cells (Fig. 6). These data indicate that even though IEC-Cdx2 cells express $K_{\rm v}1.4, K_{\rm v}1.5,$ and $K_{\rm v}1.6$ channel proteins, only $K_{\rm v}1.4$ and $K_{\rm v}1.6$ are present on the cell surface membrane to form functional channels.

The detection of $K_v1.6$ in the fraction of proteins precipitated by $K_v1.4$ antibody (Fig. 7, top panel) suggests that these two subunits co-assemble to form heteromeric K^+ channels. The specificity of the interaction between $K_v1.4$

and $K_v1.6$ was confirmed in two ways. First, the protein precipitated by $K_v1.4$ was immunoblotted with $K_v1.6$ antibody incubated with an excess of the fusion protein used to generate the antibody (Fig. 7, middle panel). In addition, the protein pulled down by an unrelated K^+ channel antibody was probed for $K_v1.6$ (Fig. 7, lower panel). $K_v1.6$ protein was not detected under either experimental condition.

The total cellular expression of $K_v1.4$ was decreased significantly by indomethacin, phenylbutazone and NS-398, the NSAIDs that impair epithelial cell migration, but not by SC-560, an NSAID with no inhibitory effects on cell migration (Fig. 8A). Cell surface expression of $K_v1.4$ was similarly affected

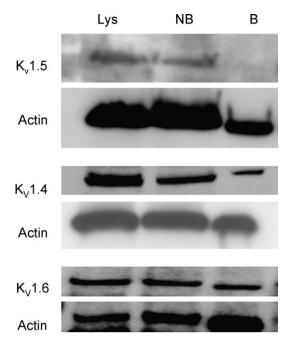


Fig. 6 – K_v1 channel expression by IEC-Cdx2 cells. Immunblots of $K_v1.5$, $K_v1.4$ and $K_v1.6$ channel protein in total cell lysates (Lys), non-biotinylated (NB) and biotinylated (B) fractions obtained from IEC-Cdx2 cells subjected to cell surface biotinylation. After detection of K_v channel protein (upper panel of each set), each blot was stripped and reprobed with antibody against actin to assess loading (lower panel of each set). The apparent molecular masses of $K_v1.5$, $K_v1.4$ and $K_v1.6$ were \sim 75, \sim 96 and \sim 56 kDa, respectively. The immunoblots shown are representative of $n \geq 3$.

(Fig. 8B). Interestingly, the NSAIDs that impair epithelial cell migration, indomethacin and NS-398, had no effect on the total expression of $K_v1.6$ (Fig. 9A), although the cell surface expression of this channel subunit was decreased significantly (Fig. 9B).

4. Discussion

Early mucosal restitution, as manifest by the rapid resealing of superficial wounds secondary to cell migration, is an important primary repair modality, critical to preserving GI mucosal integrity [17–20]. NSAIDs contribute to ulcer formation and delay ulcer healing by inhibiting intestinal epithelial cell migration and mucosal restitution [14–16]. The present study provides a potential mechanism for these ulcerogenic effects of NSAIDs.

Our data are consistent with published work [3] that suggests inhibition of mucosal cyclooxygenase (COX) and loss of protective prostaglandins (PG) are insufficient to explain NSAID effects on epithelial restitution. The COX-2 antagonist NS-398 and the non-specific COX anatagonist indomethacin impeded intestinal epithelial cell migration, whereas the COX-1 antagonist SC-560 did not. In our investigations, NSAID-inhibition of restitution was associated with depolarization of

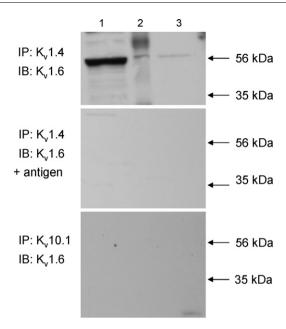


Fig. 7 – Go-association of $K_v1.6$ and $K_v1.4$ in IEC-Cdx2. Top: immunoprecipitation (IP) was performed with antibody against $K_v1.4$, then the starting material (lane 1), IP (lane 2) and supernatant (lane 3) were immunoblotted (IB) directly with antibody against $K_v1.6$. Middle: IP was performed with antibody against $K_v1.4$, then the starting material (lane 1), IP (lane 2) and supernatant (lane 3) were IB with antibody against $K_v1.6$ pre-incubated with an excess of the fusion protein encoding the epitope used to generate the $K_v1.6$ antibody. Bottom: IP was performed with antibody against $K_v10.1$, then the starting material (lane 1), IP (lane 2) and supernatant (lane 3) were IB directly with antibody against $K_v1.6$. The immunoblots shown are representative of $n \geq 3$.

epithelial cell $E_{\rm m}$ and altered surface expression of heteromeric K^+ channels, but not with depletion of intracellular polyamines. To our knowledge, this is the first demonstration of an association between the GI toxicity of NSAIDs and their effects on $E_{\rm m}$ and $K_{\rm v}$ channel expression.

4.1. NSAIDs and depletion of intracellular polyamines

Previous studies have shown that depletion of intracellular polyamines by DFMO inhibits epithelial migration [22,25,29,31,37] and that NSAIDs accelerate polyamine catabolism in colonic epithelial cancer cells [26]. Based on these observations, we hypothesized that NSAIDs inhibit intestinal cell migration by depleting intracellular polyamines. In the course of testing this idea, we first confirmed that NS-398 and indomethacin inhibit epithelial cell migration in wounded IEC-6 and IEC-Cdx2 monolayers, and then validated the findings of previous investigators that depletion of intracellular polyamines inhibits migration (Fig. 2). However, as shown in Fig. 3B, the NSAIDs that inhibited cell migration either had no significant effect on IEC-6 polyamine content (indomethacin and phenylbutazone) or decreased spermine but not spermidine content (NS-398). Together, these results

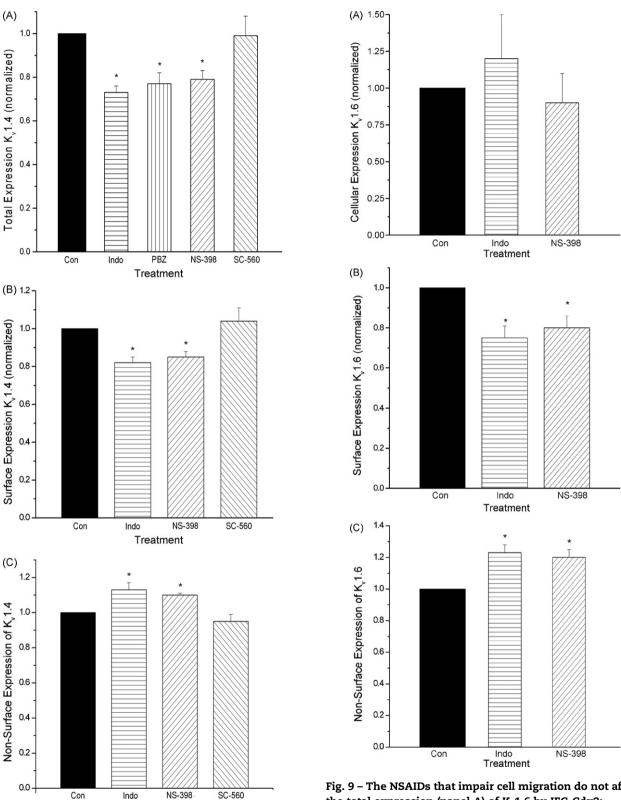


Fig. 8 – NSAIDs influence the total (panel A), cell surface (panel B) and non-surface (panel C) expression of $K_v1.4$ by IEC-Cdx2. Monolayer cultures were exposed to drug for 72 h before surface biotinylation and lysis. Drug concentrations were identical to those used in Figs. 1–4. Asterisks indicate significant differences (p < 0.05).

Treatment

Fig. 9 – The NSAIDs that impair cell migration do not affect the total expression (panel A) of $K_v 1.6$ by IEC-Cdx2; however, these drugs do diminish the cell surface expression (panel B) and increase the non-surface expression (panel C) of $K_v 1.6$. Monolayer cultures were exposed to drug for 72 h before surface biotinylation and lysis. Drug concentrations were identical to those used in Figs. 1–4. Asterisks indicate significant differences (p < 0.05).

indicate that polyamine-depletion is not a major mechanism underlying NSAID-mediated inhibition of intestinal cell migration. In light of these findings, we next considered the possibility that NSAIDs with ulcerogenic potential inhibit restitution by altering $E_{\rm m}$ independent of poyamine depletion, as discussed below.

4.2. NSAIDs and epithelial cell resting potential (Em)

The migration of IEC-6 and IEC-Cdx2 cells is influenced significantly by $E_{\rm m}$ because of the Nernstian relationship between $E_{\rm m}$ and the driving force for calcium-influx, where the latter is an important determinant of downstream signaling events that are critical for epithelial restitution [21–23,25]. Consequently, depolarization of $E_{\rm m}$ reduces Ca^{2+} influx and thereby inhibits cell migration [23,25]. In the present study, NSAIDs with inhibitory effects on cell migration induced a significant increase in DiBAC₄(3) fluorescence consistent with depolarization of $E_{\rm m}$ whereas an NSAID without negative effects on intestinal cell migration, caused no significant change in DiBAC₄(3) fluorescence (Fig. 4).

The global qualitative results provided by the fluorescence measurements subsequently were confirmed by direct measurements of $E_{\rm m}$ in individual cells using microelectrode techniques. Those studies indicated that the resting $E_{\rm m}$ of IECCdx2 cells measured in this study (-38.5 ± 1.8 mV) was similar to that reported previously (-43 ± 2.5 mV) by Rao et al. [23] and that the NSAIDs with inhibitory effects on epithelial cell migration depolarized the $E_{\rm m}$ of IEC-Cdx2 by $\sim\!15$ to 20 mV. This magnitude of depolarization was associated with differential rates of intestinal cell migration in previous investigations of epithelial restitution [23,25], supporting the contention that depolarization of $E_{\rm m}$ is sufficient to account for the inhibitory effects of NSAIDs on the migration of IEC-Cdx2 cells.

4.3. NSAIDs and expression of K_v channels

The $E_{\rm m}$ of non-excitable cells reflects the K⁺ gradient established by the Na⁺/K⁺-ATPase, a selective permeability to K⁺ and an inward "leak" driving the measured $E_{\rm m}$ from the Nernst potential for K⁺. The membrane permeability to K⁺ is the most important determinant of $E_{\rm m}$, and it can be mediated through non-gated permeation pathways and/or voltage-gated potassium channels.

Previous investigations of IEC-6 and IEC-Cdx2 cells have shown not only that these cells express multiple $K_{\rm v}$ channel subunits at the mRNA and protein levels, but also that delayed rectifier currents conducted by 4-aminopyridine-sensitive $K_{\rm v}$ channels play a major role in controlling $E_{\rm m}$ and epithelial cell migration [21,23,25]. Our work extends those findings. We document for the first time the cell surface expression (Fig. 6) and the co-association (Fig. 7) of $K_{\rm v}1.4$ and $K_{\rm v}1.6$ channel proteins in IEC-Cdx-2 cells. Previous investigations had shown expression of mRNA encoding these two K^+ channel subunits in intestinal epithelial cells using RT-PCR [21,23]. We also demonstrate that the experimental atrial anti-arrhythmic drug AVE0118 inhibits IEC-Cdx2 migration (Fig. 5). AVE0118 and 4-aminopyridine are structurally distinct K^+ channel antagonists; however, both drugs inhibit the delayed rectifier

currents conducted by channels composed of $K_v1.x$ subunits [38,44].

In one previous investigation, the pharmacological inhibition of K⁺ channels in intestinal epithelial cells was shown to accelerate rather than inhibit intestinal epithelial cell migration [45]. Although this result may appear to be at odds with our data, it is actually consistent with the hypothesis that K_v channels influence epithelial restitution by modulating E_m and intracellular calcium. Lotz et al. [45] studied the effects of K+ channel antagonism on epithelial restitution using monolayers of the human colon carcinoma cell line T84. Colon cancer is associated with increased expression of L-type calcium channels, and surface expression of these channels has been documented in cultured T84 cells [46]. In contrast, it is well established that IEC-6 and IEC-Cdx2 cells lack voltagegated calcium channels [21,23,25]. On this basis, antagonism of K_v channels would be expected to have opposite effects on intracellular calcium and calcium-dependent cellular events in T84 cells compared to IEC-6 and IEC-Cdx-2 cells. As described above, depolarization of $E_{\rm m}$ in the IEC-6 cell lines that lack voltage-gated calcium channels decreases the driving force for calcium entry thereby inhibiting calciumdependent cell migration processes [21-23,25]. Conversely, in T84 cells the depolarization of $E_{\rm m}$ associated with $K_{\rm v}$ channel inhibition would be expected to increase the influx of calcium through voltage-gated L-type calcium channels and thus enhance epithelial cell migration.

The co-association of K_v1.4 and K_v1.6 channel subunits in IEC-Cdx2 cells is a significant finding. The inactivation kinetics of heteromeric K+ channels containing K_v1.4 and K_v1.6 subunits differ significantly from those of homomeric K_v1.4 channels, such that channels formed by co-assembly of K_v1.4 and K_v1.6 conduct delayed rectifier K⁺ currents which exhibit slow, partial inactivation, while channels comprised of K_v1.4 conduct rapidly inactivating transient outward currents [47]. The expression in non-excitable cells of K_v1.4 channels does not confer sufficient K+ permeability to influence the membrane potential because of the rapid, complete inactivation of the currents [48]. In contrast, the expression in non-excitable cells of K_v channels that conduct delayed rectifier currents is sufficient to generate a resting $E_{\rm m}$ that corresponds to the base of the curve relating voltage to current activation [48]. Therefore, the heteromeric channels containing K_v1.4 and K_v 1.6 would be expected to contribute to the $E_{\rm m}$ of IEC-Cdx2.

The NSAIDs that inhibit cell migration and depolarize the E_{m} of IEC-6 and IEC-Cdx2 cells decrease significantly the total expression of K_v1.4 (Fig. 8A) but not K_v1.6 (Fig. 9A) protein in whole cell lysates. In contrast, these drugs decrease the cell surface expression of both K_v1.4 (Fig. 8B) and K_v1.6 (Fig. 9B) in IEC-Cdx2. These data suggest that the availability of Kv1.4 can influence the surface expression of the heteromeric channel complexes containing K_v1.4 and K_v1.6 in intestinal epithelial cells. In fact, an extensive body of work indicates not only that K_v1 channel pore-forming subunits possess distinct trafficking and surface expression properties, but also that the subunit composition of heteromeric channels determines their surface expression [49-52]. The ER export and cell-surface expression of K_v1.6 is inherently less efficient than that of K_v1.4 [50]. However, K_v1.4 enhances intracellular trafficking and surface expression of co-associated K_v1 subunits in

epithelial cells [49–52]. Thus, it is highly plausible that NSAIDs inhibit intestinal cell migration and promote ulcer formation by reducing the expression of functional delayed rectifier channels on the surface membrane of intestinal epithelial cells by decreasing the total expression of $K_{\nu}1.4$.

Additional experiments will be needed to understand the mechanistic basis for the newly documented effects of NSAIDs on the abundance of voltage gated K^+ channels on the intestinal epithelial cell surface, and to translate those findings into new strategies for mitigation of NSAID toxicity. Key questions remain to be answered: Do NSAIDs impact the cell surface abundance of K_v channels via drug effects on forward trafficking, internalization and recycling, or some combination thereof? What are the downstream effectors that link K^+ channel expression and E_m to epithelial cell? Do gastroprotective prostaglandins modulate NSAID effects on E_m and K_v channel expression? Are the effects of NSAIDs on K_v channel expression completely independent of the drugs' therapeutic effects?

4.4. Concluding remarks

In summary, NS-398 and indomethacin inhibit intestinal epithelial cell migration through the depolarization of $E_{\rm m}$ and the biogenic modulation of $K_{\rm v}$ channels. The data from this study suggest that these NSAIDs decrease the trafficking and cell surface expression of heteromeric $K_{\rm v}$ channel complexes by decreasing the total cellular expression of the $K_{\rm v}1.4$ channel subunit. The mechanisms responsible for drug-induced changes in the expression of $K_{\rm v}1.4$ remain to be elucidated. However, it is clear that depletion of intracellular polyamines is not a contributing factor. This work has identified a novel mechanism for NSAID-induced GI toxicity.

Acknowledgements

This work was supported by NIH grants P20-RR017686, T32RR017497 and T35RR007064 and the College of Veterinary Medicine at Kansas State University.

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