Expression of cyclooxygenase 2, microsomal prostaglandin E synthase 1, and EP receptors is increased in rat oesophageal squamous cell dysplasia and Barrett’s metaplasia induced by duodenal contents reflux

T J Jang, S K Min, J D Bae, K H Jung, J I Lee, J R Kim, W S Ahn

**Background and aim:** It is known that bile acids can induce mucosal injury, stimulate cell proliferation, and promote tumorigenesis. A large body of genetic and biochemical evidence indicates that the biosynthetic pathway of prostaglandin E (PGE) may play an important role in human and rodent tumours. Therefore, we examined the expression pattern of cyclooxygenase 1 (COX-1), COX-2, and microsomal prostaglandin E synthase 1 (mPGES-1), as well as EP receptor subtypes in rat oesophageal lesions induced by duodenal contents reflux.

**Methods:** Oesophagoduodenal anastomosis was performed in rats to induce duodenal contents reflux. We examined histological changes and expression of COX-1, COX-2, mPGES-1, and EP receptor subtypes in the oesophagus by immunohistochemistry and reverse transcription-polymerase chain reaction.

**Results:** Normal control oesophageal tissues showed COX-1 expression in subepithelial stromal cells, including endothelial cells and muscular cells, and did not reveal expression of COX-2 or mPGES-1. In the case of Barrett’s oesophagus, COX-2 and mPGES-1 were predominantly in subepithelial stromal cells. mRNA levels of COX-2, mPGES-1, EP2, EP3, and EP4 were higher in the experimental groups than in controls.

**Conclusions:** We suggest that the biosynthetic pathway of PGE may play an important role in oesophageal squamous cell dysplasia and glandular metaplasia induced by duodenal contents reflux.

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eflux of duodenal contents in addition to gastric acid in humans seems to contribute to the development of oesophagitis and Barrett’s oesophagus. Experimental studies in the rat have shown that chronic duodenal contents reflux into the oesophagus induces severe oesophagitis and also plays a role as a carcinogenic factor by increasing the number of oesophageal carcinomas when a carcinogen is given simultaneously. Moreover, chronic refluxed duodenal contents per se caused squamous cell carcinoma, adenoma, adenocarcinoma, and adenocarcinoma. Although the precise mechanism by which duodenal reflux causes oesophageal injury and predisposes to neoplasia is uncertain, there is considerable evidence that bile acids can induce mucosal injury, stimulate cell proliferation, and promote tumorigenesis. Two isoforms of cyclooxygenase 1 (COX-1) and COX-2 have been characterised in mammalian and avian species. COX-1 is constitutively expressed in most tissues to maintain stable physiological conditions whereas COX-2 is transiently induced by proinflammatory cytokines and growth factors, and involved in inflammation and mitogenesis. Recent studies have shown that the constituents of gastro-oesophageal reflux, including acid and bile, can regulate COX-2 expression. COX-2 is upregulated in reflux oesophagitis, Barrett’s oesophagus, and oesophageal carcinoma. In addition, use of COX-2 inhibitors results in a reduction in the development of oesophageal adenocarcinoma induced by duodenal reflux.

COX catalyses the conversion of arachidonic acid to prostaglandin G2 (PGG2) and PGH2. PGH2 is subsequently converted to a variety of prostaglandins that include PGE2, PGD2, PGF2, PGL2, and thromboxane A2 by each prostaglandin synthase. PGE2 has been shown to induce malignant changes in epithelial cells through immunosuppression, inhibiting apoptosis, increasing the metastatic potential of epithelial cells, and promoting angiogenesis. Two segregated biosynthetic pathways have been described for PGE2 biosynthesis. These pathways synthesise PGE2 via prostaglandin E synthase (PGES) functionally linked to either COX-1 or COX-2. At least three PGES enzymes, including cytosolic PGES, microsomal prostaglandin E synthase 1 (mPGES-1), and mPGES-2 have been identified. Induced expression of mPGES-1 has been postulated to be associated with various pathophysiological events in which COX-2 derived PGE2 has been implicated, such as rheumatoid arthritis, febrile response, reproduction, bone metabolism, and Alzheimer’s disease. In addition, mPGES-1 is

**Abbreviations:** COX, cyclooxygenase; mPGES-1, microsomal prostaglandin E synthase 1; PG, prostaglandin; BrdU, bromodeoxyuridine; RT-PCR, reverse transcription-polymerase chain reaction; GAPDH, glyceraldehyde 3-phosphate dehydrogenase; VEGF, vascular endothelial growth factor; HIF-1α, hypoxia inducible factor 1α; NSAIDs, non-steroidal anti-inflammatory drugs
over expressed in colorectal cancers, non-small cell lung cancers, and endometrial cancers. A recent study showed that mPGES-1 directed cellular transformation was accompanied by changes in the expression of a variety of genes related to proliferation, morphology, adhesion, and the cell cycle. PGF₂α mediates its effects, in part, through G protein coupled PGE receptors, designated EP₁, EP₂, EP₃, and EP₄.¹² EP₁ and EP₃ knockout mice showed a reduction in the number of aberrant crypt foci that develop in mice following azoxymethane treatment.³³ Moreover, mRNA expression of EP₂ and EP₄ was increased in human cervical and endometrial cancer tissues compared with normal tissues.³⁴ ³⁵

We examined the pattern of COX-2 and mPGES-1 expression, as well as EP receptor subtypes, to elucidate the relationship between arachidonate metabolism and oesophageal tumorigenesis induced by chronic duodenal reflux.

**MATERIAL AND METHODS**

**Animals**

Thirty seven week old male Sprague-Dawley rats (Kist, Taejun, Korea) were used for the control (n = 8) and experimental (n = 29) groups. Throughout the experiment, all rats were housed in a controlled environment with a 12 hour light/dark cycle and a temperature of 22 (2) °C. After an acclimatisation period of one week, 29 experimental rats were randomly divided into five groups in a time course of four hours for each group, the control and experimental rats were killed with ether.

**Surgical technique and tissue samples**

Solid food was withdrawn for 24 hours and water for 12 hours before surgery. Anaesthesia was induced and maintained with an isoflurane-air mixture. Oesophagoduodenal anastomosis was performed according to the Clark model. In brief, a midline laparatomy was performed, and the gastro-oesophageal junction was identified and mobilised while carefully preserving the vagus nerve. The gastro-oesophageal junction was ligated, and the distal oesophagus was transected 2 mm above the ligature. A total of eight polypropylene 7-0 sutures were placed. A 5 mm transverse enterostomy was created on the antimesenteric border of the duodenum, 1 cm distal to the pylorus. An end to side oesophagoduodenostomy was performed, and the gastro-oesophageal junction was identified and mobilised while carefully preserving the vagus nerve. The abdomen incision was closed in two layers and postoperatively the rats were allowed to drink water after six hours and were fed the following day.

**Immunohistochemistry**

Serial sections of 4 μm thickness were made and spread on poly-L-lysine coated slides. Paraffin sections were immersed in xylene and hydrated using a graded series of alcohol. Antigen retrieval was performed routinely by immersing the sections in 0.01 M citrate buffer (pH 6.0) in a pressure cooker by autoclaving for 15 minutes.

Endogenous peroxidase activity was blocked with 3% hydrogen peroxide for 15 minutes and then incubated with a primary antibody overnight in a humidified chamber at 4°C. Primary antibodies were polyclonal rabbit anti-COX-1 (Cayman Chemical, Ann Arbor, Michigan, USA) at a dilution of 1:2000, anti-COX-2 antibody (Cayman Chemical) at a dilution of 1:500, anti-mPGES-1 antibody (Cayman Chemical) at a dilution of 1:500, and monoclonal mouse anti-BrdU antibody (Dako, Santa Barbara, California, USA). Staining was achieved with a Dako LSAB-kit and developed with diaminobenzidine tetrahydrochloride (Dako). Sections were counterstained for five minutes with Meyer’s haematoxylin and then mounted. Human colon cancer with intense staining for both COX-2 and mPGES-1 was used as a positive control. As a negative control, rabbit and mouse IgG isotypes (Dako) were used instead of primary antibodies. BrdU labelling index was calculated by counting at least 1000 cells in a random 10 high power fields. Estimation of immunohistochemical expression of COX-1 and COX-2 was evaluated according to both intensity and area of signal: 0, absent; 1, mild; 2, moderate; and 3, severe. Expression of mPGES-1 was assessed as negative or positive.

**RNA extraction and RT-PCR**

Total RNA was extracted and purified from frozen lower oesophageal tissues using the GeneElute Mammalian Total RNA kit (Sigma according to the manufacturer’s instructions). RNA was quantified by determining absorbance at 260 nm. Total RNA (2 μg) from each sample was reverse transcribed into cDNA using Superscript RT reverse transcriptase (Life Technologies, Inc., Rockville, Maryland, USA) and random hexamer primers (Takara, Shiga, Japan). The PCR primers were as follows: glyceraldehyde 3-phosphate dehydrogenase (GAPDH) (311 bp), sense 5'-GAA CGG GAA GCT CAC TGG CAT GCC-3', antisense 5'-TGA GGT CCA CCA CCC TGG TGC TG-3'; COX-1 (443 bp), sense 5'-GAG TCT CTC CCT CCA GTT GTT CTC-3', antisense 5'-GCG AGT ATA GTA GTC CAC GTT GG-3'; COX-2 (448 bp), sense 5'-ATG CTC TTC CGA GCT GTG CT-3', antisense 5'-CAT GGG AGT TGG GCA GTC AT-3'; mPGES-1 (451 bp), sense 5'-ATG ACT TCC TGT GGG GTG ATG GAG-3', antisense 5'-TCA GAT GAT GGG GCA CTT ACC AGA-3'; EP₁ (465 bp), sense 5'-GAG GCA ACA AGC TGT GTA ACA-3', antisense 5'-AGC CAT GCC GGC CAG CAG GGC TAG-3'; EP₂ (336 bp), sense 5'-TCT GGC AGT AGC CTG AGA GCC-3', antisense 5'-GAG AGG AGG AGG AGG AGG AGG AGG-3'; EP₃ (412 bp), sense 5'-GAG AGG AGG AGG AGG AGG AGG AGG-3'; EP₄ (488 bp), sense 5'-GAG AGG AGG AGG AGG AGG AGG AGG-3'; The
PCR products were electrophoresed on 1.5% agarose gel containing ethidium bromide and then photographed under UV light.

**Statistical analysis**
The significance of differences between groups was evaluated using Fisher’s exact test and one way analysis of variance. Differences were considered to be statistically significant at \( p < 0.05 \).

**RESULTS**

**Gross findings in the oesophagus of the experimental groups**
Experimental rats showed an abnormally dilated oesophagus, and the oesophageal inner surface displayed whitish nodular patches, which were prominent in the lower oesophagus. There were superficial ulcers located mainly in the lower oesophagus. All of these macroscopic findings were present in all rats and more intense in rats exposed to reflux of duodenal contents for longer periods.

**Squamous cell lesions of the upper and lower oesophagus in the experimental groups**
The oesophagus of control rats did not reveal any pathological findings but various squamous cell lesions were seen in the upper and lower oesophagus of experimental rats. As shown in table 1, squamous cell lesions of the lower oesophagus were more severe than those of the upper oesophagus. Dysplasia in the lower oesophagus occurred at 10 weeks. To assess the biological behaviour of various squamous lesions, we performed immunohistochemical staining for BrdU because the proliferation index is often increased in dysplastic and cancer tissues. As anticipated, the BrdU labelling index of dysplasia was higher than that of normal and papillary hyperplasia (table 2).

**Glandular lesions of the lower oesophagus in the experimental groups**
Barrett’s oesophagus did not occur in control rats but 86% of experimental rats showed glandular metaplasia above the oesophagoduodenal junction. The results are shown in table 3. Long and atypical Barrett’s oesophagus were restricted to rats exposed to the reflux of duodenal contents for 30 and 40 weeks. Glandular metaplasia originated from the lower oesophagus because all lesions were above the oesophageal anastomosis and had an intact muscularis propria layer on histology. BrdU labelled columnar cells were located in the upper portion of Barrett’s oesophagus while in duodenal mucosa they were mainly restricted to within the isthmic portion (data not shown).

**Expression of COX-1 and COX-2 in the lower oesophagus**
Normal control oesophageal tissues showed COX-1 expression in subepithelial stromal cells, including endothelial cells and muscular cells, and did not reveal expression of COX-2 (fig 1). In the case of squamous cell lesions in the experimental groups, immunoreactivity of COX-1 was similar to that of normal controls, and there was no significant difference between histological subtypes (data not shown). COX-2 was highly expressed in dysplasia compared with normal tissue and papillary hyperplasia (table 4) (fig 1). As shown in fig. 1, COX-2 was maximally expressed around the vascular papillae of tissues showing dysplasia, and positive staining was also noticeable in the surrounding epithelial layer and basal layer. COX-1 and COX-2 were not labelled in most Barrett’s mucosa but in subepithelial stomal cells (fig 2). Of six cases of atypical Barrett’s mucosa, one (17%) showed cytoplasmic immunoreactivity for COX-2 (fig 2) and five were completely negative. We extracted and purified total RNA from frozen lower oesophageal tissues in control and experimental rats and then performed RT-PCR. In agreement with immunohistochemical results, COX-1 mRNA was expressed in both groups, and expression level of COX-2 mRNA was higher in experimental groups than in controls (fig 3).

### Table 1
Squamous cell lesions of the upper and lower oesophagus in the experimental groups

<table>
<thead>
<tr>
<th>Weeks (No)</th>
<th>Upper oesophagus</th>
<th>Lower oesophagus</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>N</td>
<td>HP</td>
</tr>
<tr>
<td>10 (2)</td>
<td>2</td>
<td></td>
</tr>
<tr>
<td>20 (12)</td>
<td>3</td>
<td>4</td>
</tr>
<tr>
<td>30 (8)</td>
<td>3</td>
<td></td>
</tr>
<tr>
<td>40 (7)</td>
<td>1</td>
<td>6</td>
</tr>
</tbody>
</table>

N; normal, HP; hyperplasia, PH; papillary hyperplasia, Dys; dysplasia.

### Table 2
Bromodeoxyuridine (BrdU) indices of squamous cell lesions of the lower oesophagus in the control and experimental groups

<table>
<thead>
<tr>
<th>No of lesions</th>
<th>BrdU indices (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Normal</td>
<td>1.0 (0.01)*</td>
</tr>
<tr>
<td>Papillary hyperplasia</td>
<td>4.2 (1.91)*</td>
</tr>
<tr>
<td>Dysplasia</td>
<td>40.7 (13.5)</td>
</tr>
</tbody>
</table>

*p < 0.05 compared with dysplasia.

### Table 3
Glandular lesions of the lower oesophagus in the experimental groups

<table>
<thead>
<tr>
<th>Weeks (No)</th>
<th>No BO</th>
<th>Short BO</th>
<th>Long BO</th>
<th>Atypical BO</th>
</tr>
</thead>
<tbody>
<tr>
<td>10 (2)</td>
<td>1</td>
<td>1</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>20 (12)</td>
<td>2</td>
<td>10</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>30 (8)</td>
<td>1</td>
<td>1</td>
<td>3</td>
<td>3</td>
</tr>
<tr>
<td>40 (7)</td>
<td>0</td>
<td>1</td>
<td>3</td>
<td>3</td>
</tr>
</tbody>
</table>

BO, Barrett’s oesophagus.
Expression of mPGES-1 in the lower oesophagus

In the case of squamous cell lesions, mPGES-1 expression was enhanced in dysplasia compared with normal tissue and papillary hyperplasia (table 5). As shown in fig 1, mPGES-1 was expressed in stromal cells around the vascular papillae of tissues showing dysplasia. These areas corresponded to regions showing immunoreactivity for COX-2. However, all squamous cells did not display immunoreactivity for mPGES-1. In the case of Barrett’s oesophagus, immunolocalisation of mPGES-1 was subepithelial stromal cells and not mucosal cells, unlike the result observed in squamous lesions (fig 2). Expression level of mPGES-1 mRNA was higher in the experimental groups than in normal controls (fig 3).

mRNA levels of EP1, EP2, EP3, and EP4 in the lower oesophagus

We assessed mRNA levels of EP1, EP2, EP3, and EP4 in the lower oesophagus by semiquantitative RT-PCR (fig 3). mRNA levels of EP1 were negligible in both the experimental and control groups. In contrast, mRNA levels of EP2, EP3, and EP4 were higher in the experimental groups than in normal controls.

DISCUSSION

We have demonstrated a significant role for the biosynthetic pathway of PGE2 in oesophageal squamous cell dysplasia and glandular metaplasia induced by duodenal contents reflux in rats. The correlation between arachidonic acid metabolism and tumorigenesis is suggested by studies on non-steroidal anti-inflammatory drugs (NSAIDs). Long term use of NSAIDs in rheumatic patients is related to a reduced risk of various human cancers, including oesophageal cancer.36 37 A large body of genetic and biochemical evidence supports a role for COX-2 in human and rodent tumours.38 39 A recent study showed that bile acids play an important role in COX-2 expression of rat reflux oesophagitis caused by oesophago-duodenal anastomosis.40 COX-2 was expressed in 91% of

Table 4 Immunoreactivity of cyclooxygenase 2 (COX-2) in squamous cell lesions of the lower oesophagus in the control and experimental groups

<table>
<thead>
<tr>
<th>No of lesions</th>
<th>COX-2</th>
</tr>
</thead>
<tbody>
<tr>
<td>Normal controls</td>
<td>8</td>
</tr>
<tr>
<td>Papillary hyperplasia</td>
<td>27</td>
</tr>
<tr>
<td>Dysplasia</td>
<td>28</td>
</tr>
</tbody>
</table>

*P<0.05 compared with papillary hyperplasia and dysplasia.
†P<0.05 compared with dysplasia.
human oesophageal squamous cell carcinomas. In this study, COX-2 was minimally induced in papillary hyperplasia and highly expressed in squamous cell dysplasia. This lesion was also characterised by increased cell proliferation. Prostaglandins can stimulate cell proliferation but it is uncertain whether enhanced expression of COX-2 is causally linked to increased cell proliferation. However, a recent study showed that rofecoxib reduced cell proliferation in Barrett’s oesophagus by inhibiting COX-2 expression and activity. Therefore, the persistence of increased COX-2 expression in squamous cell dysplasia implies that induction of the COX-2 gene may be necessary for maintenance of the malignant phenotype characterised by increased cell proliferation. Localisation of COX-2 predominantly to stromal cells around vascular papillae and surrounding squamous epithelial cells with basal cells is consistent with published findings in human reflux oesophagitis. Recently, PGE2 production via the COX-2 catalysed pathway induced vascular endothelial growth factor (VEGF) expression by inducing hypoxia inducible factor 1 (HIF-1) stabilisation and expression. Immunostaining for VEGF and HIF-1 may be necessary to evaluate immunolocalisation of COX-2 in vascular papillae.

Eighty six percent (25/29) of rats undergoing oesophago-duodenal anastomosis developed Barrett’s oesophagus but definite adenocarcinoma did not occur. This finding may be explained by the experiment of Chen and Yang. They reported that iron supplementation promoted the formation of oesophageal adenocarcinoma originated from Barrett’s oesophagus induced by the surgical techniques oesophago-duodenal or oesophagogastroduodenal anastomosis. COX-2 was over expressed in human oesophageal adenocarcinoma. Moreover, a selective COX-2 inhibitor suppressed the development of rat oesophageal adenocarcinoma induced by duodenal reflux. COX-2 expression was also increased in biopsied Barrett’s mucosa in response to pulses of acid or bile acids in an ex vivo organ culture system, which was attenuated by a selective COX-2 inhibitor. Interestingly, in metaplastic Barrett’s mucosa, COX-2 expression was observed primarily in the lamina propria, consistent with other studies; however, a shift in staining to the epithelium was observed in 17% of atypical Barrett’s mucosa, suggesting that COX-2 over expression in these cells may constitute a relatively late event.

Elevated expression of mPGES-1 has recently been demonstrated in several human cancers. Murakami et al suggested that aberrant expression of mPGES-1 in combination with COX-2 could contribute to tumorigenesis. In this study, mPGES-1 was expressed in squamous cell dysplasia, in which COX-2 was also highly expressed. It seems likely therefore that enhanced expression of mPGES-1 in addition
to COX-2 contributes to the increased amount of PGE₂. Bile acids induce COX-2 by stimulating transcription and stabilising mRNA. In contrast, mPGES-1 is not induced by bile acids. Several proinflammatory cytokines such as tumour necrosis factor and interleukin 1 induce mPGES-1. Therefore, expression of mPGES-1 restricted to stromal cells may be caused by proinflammatory cytokines produced by mucosal injury associated with the reflux of duodenal contents containing bile acids.

Eicosanoids are unstable and their activities are normally restricted to cells in the immediate vicinity that express specific receptors. Binding of PGE₂ to its receptors initiates the signalling mediated by receptor subtype specific G proteins and respective changes in second messengers. Therefore, we examined the PGE₂ receptors of the lower oesophageal sphincter by RT-PCR. Interestingly, mRNA levels of EP₂, EP₃, and EP₄ were increased in experimental oesophageal tissue showing squamous dysplasia compared with normal controls. This is in accordance with other studies.

Recently, Yang et al demonstrated an important role for the EP₂ receptor in PGE₂ induced inhibition of dendritic cell differentiation and function and diminished antitumour cellular immune responses. Another study reported the significance of PGE₂-EP₁ receptor signalling in tumour development and angiogenesis. In addition, EP₄ knockout mice showed a reduction in the number of aberrant crypt foci that develop in mice following azoxymethane treatment. In contrast, Konger et al reported that loss of EP₂ receptor in immortalised human keratinocytes resulted in increased invasiveness. They suggested that this discrepancy may be caused by differences in cell or tissue type, and the presence of negative feedback loop.

In summary, COX-2, mPGES-1, EP₂, EP₃, and EP₄ expression are increased in oesophageal lesions exposed to duodenal contents reflux, and we suggest that the biosynthetic pathway of PGE₂ may play an important role in surgically induced oesophageal squamous cell dysplasia and glandular metaplasia.

ACKNOWLEDGEMENTS

This study was supported by the Dongguk University research fund.

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REFERENCES


Table 5 Immune-reactivity of microsomal prostaglandin E synthase 1 in the squamous cell lesions of the lower oesophagus in normal controls and experimental group

<table>
<thead>
<tr>
<th></th>
<th>Positive</th>
<th>Negative</th>
</tr>
</thead>
<tbody>
<tr>
<td>Normal controls</td>
<td>0</td>
<td>8</td>
</tr>
<tr>
<td>Papillar hyperplasia</td>
<td>3</td>
<td>24</td>
</tr>
<tr>
<td>Dysplasia</td>
<td>28*</td>
<td>0</td>
</tr>
</tbody>
</table>

*p<0.05 compared with normal and papillar hyperplasia.
COX-2 and mPGES-1 in oesophageal lesions


