

## RESEARCH PAPER

## Synthesis of hyaluronan in oesophageal cancer cells is uncoupled from the prostaglandin–cAMP pathway

S Twarock<sup>1</sup>, K Röck<sup>1</sup>, M Sarbia<sup>2</sup>, AA Weber<sup>1</sup>, RU Jänicke<sup>3</sup> and JW Fischer<sup>1</sup>

<sup>1</sup>Institut für Pharmakologie, Universitätsklinikum Essen, Universität Duisburg-Essen, Essen, Germany, <sup>2</sup>Pathologie der Technischen Universität München, München, Germany, and <sup>3</sup>Laboratory of Molecular Radiooncology, Clinic and Policlinic for Radiation Therapy and Radio-oncology, Universitätsklinikum Düsseldorf, Düsseldorf, Germany

**Background and purpose:** Cyclooxygenase-2 (COX2) and hyaluronic acid (HA) are common in tumours and both independently promote tumour progression. Furthermore, COX2-dependent synthesis of prostaglandins (PGs) stimulates HA synthase-1 (HAS1) and HAS2 mRNA expression, together with HA synthesis via the cAMP/protein kinase A pathway in vascular smooth muscle cells. Therefore, the aim of the present study was to elucidate whether COX2-mediated PGs induce transcription of HAS isoforms in cancer cells as well.

**Experimental approach:** Human oesophageal squamous cell (OSC) carcinoma specimens were characterized with respect to HA, COX2 and CD44 expression by immunohistochemistry. OSC cell lines (OSC1, OSC2) and HeLa cell lines (D98, H21) were exposed to exogenous PG analogues (100 nmol·L<sup>-1</sup>), etoricoxib (10 µmol·L<sup>-1</sup>) and forskolin (10 µmol·L<sup>-1</sup>). Subsequently, cAMP levels, HA secretion and HAS isoform expression were determined by ELISA and real-time RT-PCR (reverse transcriptase polymerase chain reaction) respectively.

**Key results:** COX2, HA and CD44 were detected immunohistochemically in >90% of human oesophageal tumour samples. Under basal conditions, OSC1 and OSC2 cells express HAS2 and HAS3, COX2 and Gα<sub>s</sub>-coupled EP<sub>2</sub> and EP<sub>4</sub> PG receptors. Neither stimulation with the PGI<sub>2</sub> analogue, iloprost, addition of exogenous PGE<sub>2</sub> nor forskolin induced HAS1 or HAS2 mRNA expression in OSC1 and OSC2 cells. Furthermore, in HeLa cells after induction of COX2 by tumour necrosis factor α and subsequent PGE<sub>2</sub> release, inhibition of COX2 by etoricoxib did not affect HAS expression or HA secretion.

**Conclusions and implications:** We conclude that in oesophageal and HeLa cancer cells, HAS1/2 expression was not responsive to the PG/cAMP pathway.

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**Keywords:** oesophageal cancer; COX2; etoricoxib; prostaglandins; hyaluronic acid

**Abbreviations:** COX2, cyclooxygenase-2; HA, hyaluronic acid, hyaluronan; HAS, hyaluronic acid synthase; hSMC, human vascular smooth muscle cell; OSC, oesophageal squamous cell; PG, prostaglandin; PMA, phorbol myristate acetate; TNF-α, tumour necrosis factor α

## Introduction

Oesophageal cancer is a rare but severe form of gastrointestinal cancer that is differentiated into adenocarcinomas and squamous cell carcinomas of the oesophagus. Squamous cell carcinomas are found in the upper two-thirds of the oesophagus and are mainly induced by alcohol and cigarette smoke. Adenocarcinomas are usually found in the lower third of the oesophagus and are preceded by metaplasia, referred to as Barrett's oesophagus (Enzinger and Mayer, 2003).

Cyclooxygenase (COX) is the key enzyme during synthesis of prostanoids [prostaglandins (PGs), prostacyclin, thromboxane A<sub>2</sub>]. These important mediators account for a variety of physiological and pathophysiological processes such as inflammation, constriction and dilatation of blood vessels, regulation of platelet aggregation and control of calcium regulation. COX catalyses the production of PGH<sub>2</sub> from its precursor arachidonic acid. Downstream enzymes like PGE<sub>2</sub> synthase and prostacyclin synthase account for the transformation of PGH<sub>2</sub> into the final products of the prostanoid family. The COX enzyme exists in two isoforms, COX1 that is responsible for the production of baseline levels of prostanoids and COX2 that is induced under pathological conditions and sustains inflammation. Prostanoids bind to a set of nine prostanoid receptors (Hata and Breyer, 2004; nomenclature follows Alexander *et al.*, 2008).

Correspondence: Professor Dr Jens W Fischer, Institut für Pharmakologie, Universitätsklinikum Essen, Universität Duisburg-Essen, Hufelandstrasse 55, D-45122 Essen, Germany. E-mail: j.fischer@uk-essen.de

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Over-expression of COX isoforms, especially COX2, represents a well-known correlate of malignancy in a variety of cancers including those derived from colon (Kargman *et al.*, 1995), stomach (Uefuji *et al.*, 1998), lung (Wolff *et al.*, 1998), oesophagus (Zimmermann *et al.*, 1999), pancreas (Tucker *et al.*, 1999), liver (Shiota *et al.*, 1999), as well as head and neck tumours (Chan *et al.*, 1999). Treatment with COX2 inhibitors leads to decreased proliferation of a variety of tumours such as colorectal (Coffey *et al.*, 1997), oesophageal (Zimmermann *et al.*, 1999), prostate (Liu *et al.*, 1998) and breast cancer (Hong *et al.*, 1999). These effects can be reversed by addition of exogenous PGE<sub>2</sub> or PGF<sub>2 $\alpha$</sub> .

Hyaluronan (hyaluronic acid; HA), an unbranched polysaccharide, is composed of 2000–25 000 disaccharides of glucuronic acid and N-acetylglucosamine. Besides hydration and expansion of extracellular spaces, HA exhibits a plethora of physiological and pathophysiological functions that play a role in diverse processes including angiogenesis and inflammation involving signalling through HA receptors, CD44 and the receptor of HA-mediated motility (RHAMM). HA is produced by three isoforms of transmembrane HA synthase enzymes (HAS1–3) that link the two precursor molecules in an alternating manner and extrude the growing HA strand to the outside of the cell. While HAS1 and HAS2 produce HA up to 10<sup>7</sup> Da, HAS3 generates HA with an apparent molecular weight of approximately 10<sup>5</sup> Da. In this context, it is important to differentiate between HA of different molecular weights, because depending on its length, HA serves different functions (Stern *et al.*, 2006). An overproduction of HA in tumour cells such as breast cancer (Auvinen *et al.*, 1997), lung cancer (Horai *et al.*, 1981), colon cancer (Ropponen *et al.*, 1998), mesothelioma (Azumi *et al.*, 1992) or pancreatic cancer (Ringel *et al.*, 1999) results in elevated proliferation, migration, invasion, angiogenesis and resistance to apoptosis and cytostatic treatment. Urinary HA excretion is also an emerging prognostic marker for bladder cancer (Golshani *et al.*, 2007). Interestingly, COX2 and HA are often found in the same cancer cells.

Thus, both COX2 and HA are involved in promoting a variety of malignancies. Recently, it was discovered that PGE<sub>2</sub> and prostacyclin induce HAS1 and HAS2 mRNA expression in human vascular smooth muscle cells (hSMCs). This was proposed to be the outcome of an EP<sub>2</sub> or IP receptor-mediated activation of protein kinase A (PKA) and an increase in cAMP, that in turn binds to the cAMP-responsive element, a transcription site in the HAS2 promoter region (Monslow *et al.*, 2004; Sussmann *et al.*, 2004; van den Boom *et al.*, 2006). Based on these findings, we proposed, as a working hypothesis, that COX2-dependent PGs induce expression of HAS isoforms that subsequently give rise to HA-rich pericellular matrices, thereby promoting the malignant phenotype of cancer cells. The aim of the present study was to test this working hypothesis in oesophageal cancer cells.

## Methods

### Cell culture

The oesophageal cancer cell lines OSC1 and OSC2 (oesophageal squamous cell) and the HeLa cell lines H21 and D98 were

described previously (Defilippi *et al.*, 1987; Sarbia *et al.*, 1997). The cells were routinely maintained as monolayer cultures in RPMI-1640 supplemented with 10% foetal bovine serum, L-glutamine, penicillin and streptomycin at 37°C and 5% CO<sub>2</sub>. For experiments, cells were subjected to serum withdrawal for 24 h and subsequently treated with the test compounds.

### Immunohistochemical analysis of human tumour samples

The tumour samples were collected for diagnostic purposes and were examined by a senior pathologist (MS). Informed consent was obtained from patients or relatives. The present study was performed according to the Declaration of Helsinki. Immunohistochemical investigations were based on 69 patients (55 male; age range: 35–81 years) with squamous cell carcinoma of the oesophagus that underwent oesophageal resection without prior radio- or chemotherapy between 1978 and 1991 at the Department of Surgery, University of Düsseldorf. Subsequently, the resection specimens were fixed in buffered 4% formaldehyde. Pathological tumour stage was determined at the Department of Pathology, University of Düsseldorf according to standard procedures. For the purpose of this study, all cases were restaged according to the current TNM classification (Wittekind *et al.*, 2002). Accordingly, 26 cases were in pathological stage IIa (37.7%), 8 in stage IIb (11.6%) and 35 in stage III (50.7%).

For each of the tumours, one paraffin block including representative, non-necrotic tumour areas was selected. Three tissue cylinders with a diameter of 0.6 mm per tumour were punched from these areas and brought into a recipient paraffin block by using a tissue arraying instrument (Beecher Instruments, Silver Spring, MD, USA). Sections (4  $\mu$ m) from the tissue microarrays were mounted on Superfrost-plus slides (Langenbrinck, Teningen, Germany, Order-No 030060) for the subsequent immunohistochemical study.

Sections were immunohistochemically stained by using antibodies against COX2 (Cayman Chemical, Ann Arbor, MI, USA, 1:150) and CD44v (Sigma-Aldrich, Munich, Germany, HPA005785, 1:500). HA was detected by using biotinylated HABP (Seikagaku Corp., Tokyo, Japan; 2  $\mu$ g·mL<sup>-1</sup>) as described (Ripellino *et al.*, 1985). This staining procedure on paraffin-embedded sections detects expression of both, intracellular and extracellular HA. The sections were de-paraffinized and rehydrated in graded alcohol. Subsequently, for COX2 and CD44 detection, the sections were subjected to citrate buffer (pH 6.0) in a steamer at 100°C for 20 min. Primary antibody was applied overnight at 4°C. After application of the secondary antibody, antigen detection was performed by using diaminobenzidine as chromogen. Finally, the sections were counterstained with haematoxylin. Negative controls were treated identically to the tumour samples including antigen retrieval procedure with the exception that the primary antibody was replaced by non-immune serum. In negative controls no signals were detectable.

Using light microscopy, expression of HA, COX2 and CD44 was determined by a senior pathologist (MS). Cases were considered positive for expression when cancer cells in at least one out of three tissue cylinders per tumour case showed a distinctive extracellular/cytoplasmic (HA), cytoplasmic (COX2) or membranous expression pattern (CD44). Cases

were considered negative if cancer cells in all analysable tissue cylinders per case showed no immunoreactivity.

#### RNA isolation

Total RNA from treated cells was isolated by using Tri-Reagent (Sigma-Aldrich) according to the manufacturer's protocol. The RNA was quantitated by spectroscopic analysis at 260 nm with an Eppendorf photometer.

#### Quantification of gene expression

The expression levels of HAS1, HAS2 and HAS3 were analysed by real-time RT-PCR (reverse transcriptase polymerase chain reaction). For real-time RT-PCR quantification, total RNA (1 µg) was used for cDNA synthesis, and specific primers for human HAS isoforms were generated according to the known sequences HAS1 (NM\_001523.1), HAS2 (NM\_005328.1), HAS3 (NM\_005329.2) as follows: HAS-1: fwd 5'-TACAACCAG AAGTTCCTGGG-3', rev 5'-CTGGAGGTGTACTTGGTAGC-3'; HAS-2: fwd 5'-GTGGATTATGTACAGGTTTGTGA-3', rev 5'-TCCAACCATGGGATCTTCTT-3'; HAS-3: fwd 5'-GAGATGTC CAGATCCTCAACAA-3', rev 5'-CCCCTAATACTACTGCACAC-3'; GAPDH: fwd 5'-GTGAAGGTCGGAGTCAACG-3', rev 5'-TGAGGTCAATGAAGGGGTC-3'. The relative mRNA expression levels were determined by using GAPDH as house

keeping gene and the  $2^{-\Delta\Delta C_T}$  method. Values were then expressed as fold of respective controls.

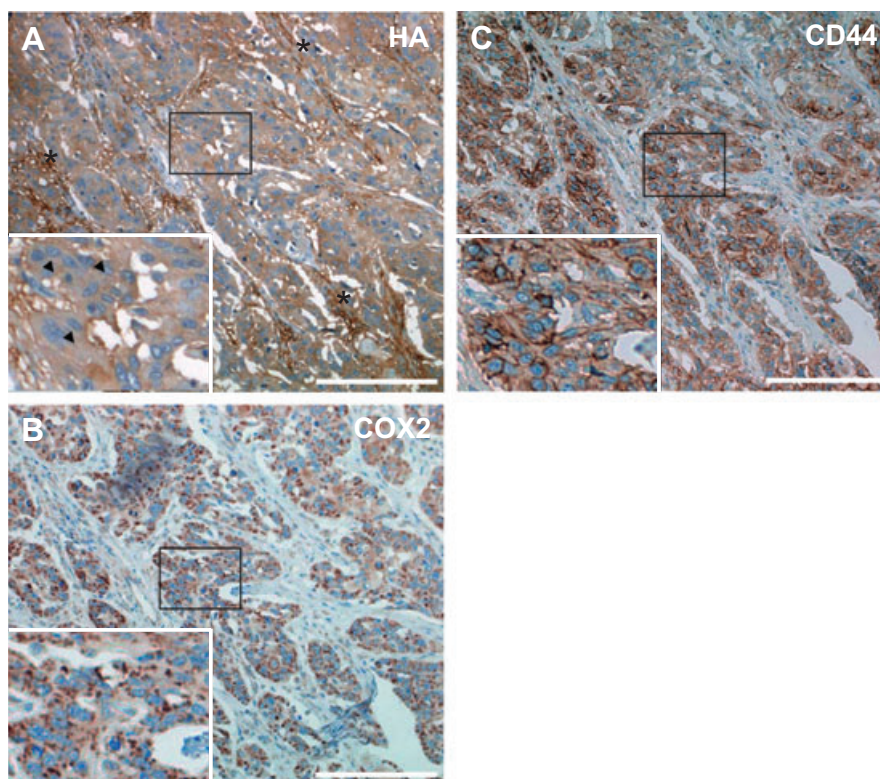
Prostaglandin receptors were detected by RT-PCR using the following sequences: EP1: fwd 5'-GCGCTGCCCCATCTTCTCC-3', rev 5'-GGTACTGCAGCTCATAGC-3'; EP2: fwd 5'-GCTGGA CTATGGGCAGTACG-3', rev 5'-AACAGGAGGCCTAAGGA TGG-3'; EP3: fwd 5'-TCGGGCTCTCCTCGTTGTC-3', rev 5'-AGTGAAGCCAGGCGAACAGC-3'; EP4: fwd 5'-ACTACG TGGACAAGCGATTG-3', rev 5'-TCACAGAAGCAATTCGGA TG-3'; IP: fwd 5'-GCGTCCTTCTTGCGCGCTGCCCTGCT GG-3', rev 5'-GGTCCCCCATCTCACTGCTGCTGGTCAGG-3'.

#### Determination of the HA concentration

Cells were plated at a density of  $10^5$  cells per well in a 6 well plate and allowed to adhere for 24 h. The cells were cultured with or without stimulus for 6–24 h. HA released into the culture medium was quantified by using a commercially available assay based on HA-binding protein according to the manufacturer's instructions (Corgenix, Broomfield, CO, USA). Secreted HA was normalized to total cellular protein.

#### Quantification of PGI<sub>2</sub> and PGE<sub>2</sub> levels

Cell culture supernatants were collected at the indicated times and the concentration of 6-oxo-PGF<sub>1α</sub>, the stable hydrolysis



**Figure 1** Immunohistochemical detection of hyaluronic acid (HA), cyclooxygenase-2 (COX2) and CD44 in human oesophageal cancer tissue. Shown are sections of human oesophageal tumours, representative for 69 cases. To assess the spatial relationships of HA, COX2 and CD44, consecutive sections were stained. HA (A) was detected in both tumour cell parenchyma and stromal cells; asterisks indicate accumulation of HA in tumour stroma; inset, magnification of the indicated area of a tumour cell cluster demonstrating HA within the cancer cells as well (arrowheads). COX2 (B) and the HA receptor CD44 (C) were detected mainly in tumour cell islands. Taken together, the majority (>90%) of the specimens displayed a strong expression of all three proteins. The micrographs show representative sections (scale bar marks 200 µm).



product of PGL<sub>2</sub>, was determined by radioimmunoassay (RIA) as described previously (Sussmann *et al.*, 2004). PGE<sub>2</sub> was quantified by use of a commercially available ELISA assay (Prostaglandin E<sub>2</sub> EIA Kit – Monoclonal, Cayman Chemical, Ann Arbor, MI, USA).

*Measurement of intracellular cAMP concentrations*

Cells were seeded in 6 well plates in serum-free medium for 72 h. Then, the cells were washed twice with 2 mL of a balanced salt solution containing 130 mmol·L<sup>-1</sup> NaCl, 5.4 mmol·L<sup>-1</sup> KCl, 1.8 mmol·L<sup>-1</sup> CaCl<sub>2</sub>, 0.8 mmol·L<sup>-1</sup> MgCl<sub>2</sub>, 5.5 mmol·L<sup>-1</sup> glucose and 20 mmol·L<sup>-1</sup> HEPES, pH 7.3. Stimuli were added for 10 min. The reaction was stopped by removing the buffer and by addition of ice-cold ethanol (96%). Following evaporation of the ethanol, intracellular cAMP levels were determined by an RIA as described previously (Schröder and Schrör, 1993). All experiments were carried out in triplicate.

*Data analysis*

Data are presented as means ± SEM of the indicated number of independent experiments. Statistical comparisons between groups were performed by using one-way ANOVA followed by the Bonferroni post test. A *P*-value <0.05 was considered significant.

*Materials*

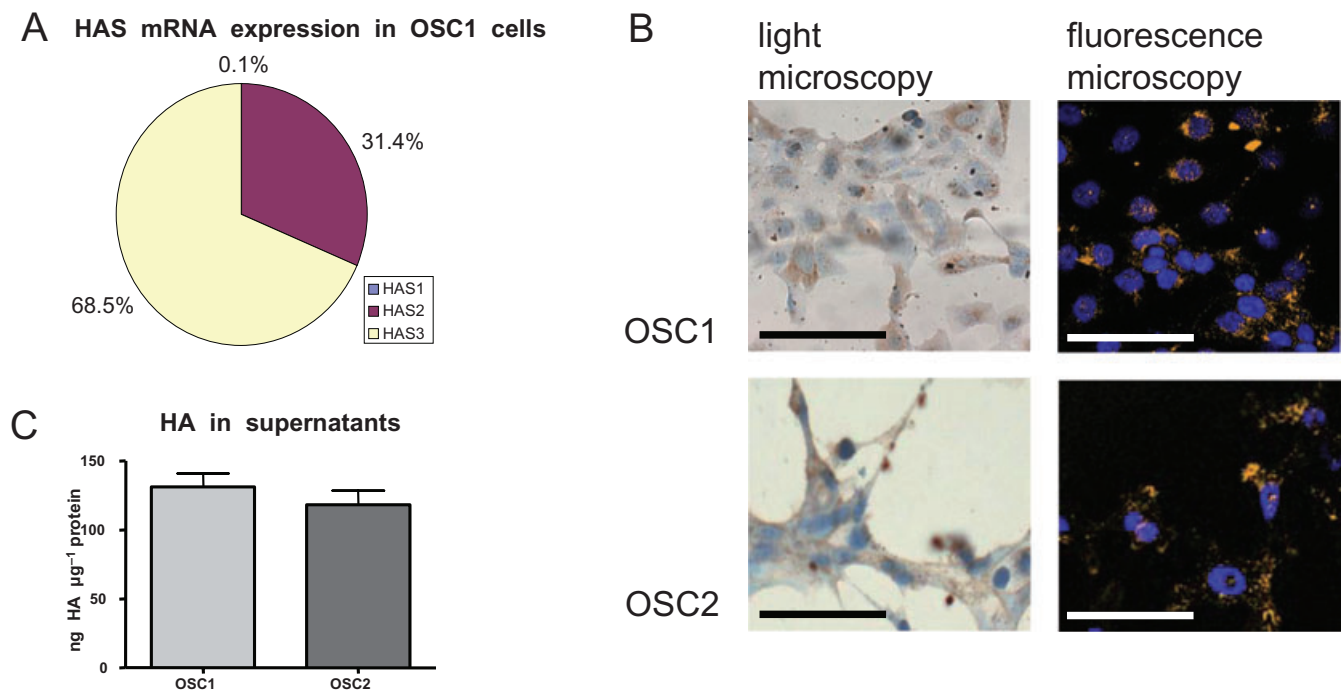
Reagents for RT-PCR and real-time RT-PCR were obtained from Qiagen (Hilden, Germany) and Invitrogen (Karlsruhe,

Germany). All cell culture reagents were obtained from Invitrogen or Sigma-Aldrich (Munich, Germany). Iloprost was kindly provided by Schering AG (Berlin, Germany). Forskolin, PGE<sub>2</sub> and human recombinant tumour necrosis factor α (TNF-α) were purchased from Sigma-Aldrich. Etoricoxib was bought from WITEGA Laboratorien Berlin-Adlershof GmbH (Berlin, Germany).

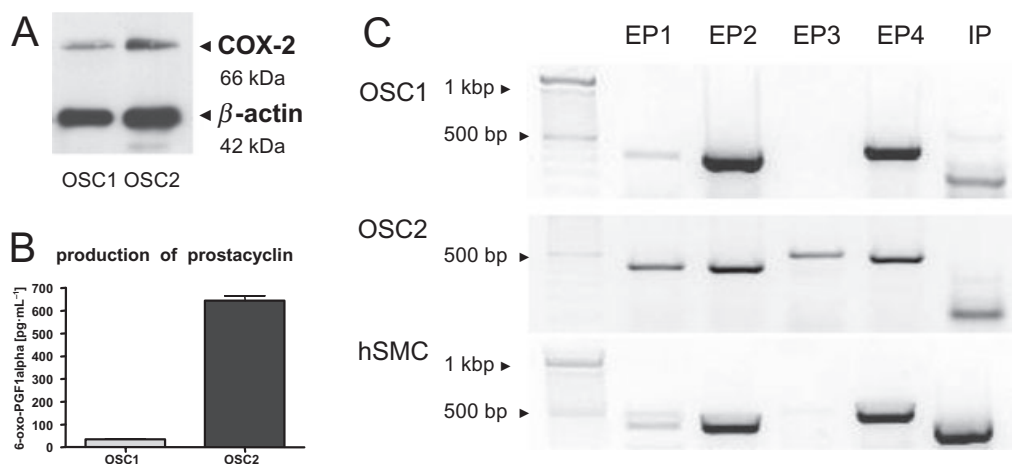
**Results**

*Expression of HA, COX2 and CD44 in human oesophageal cancer sections*

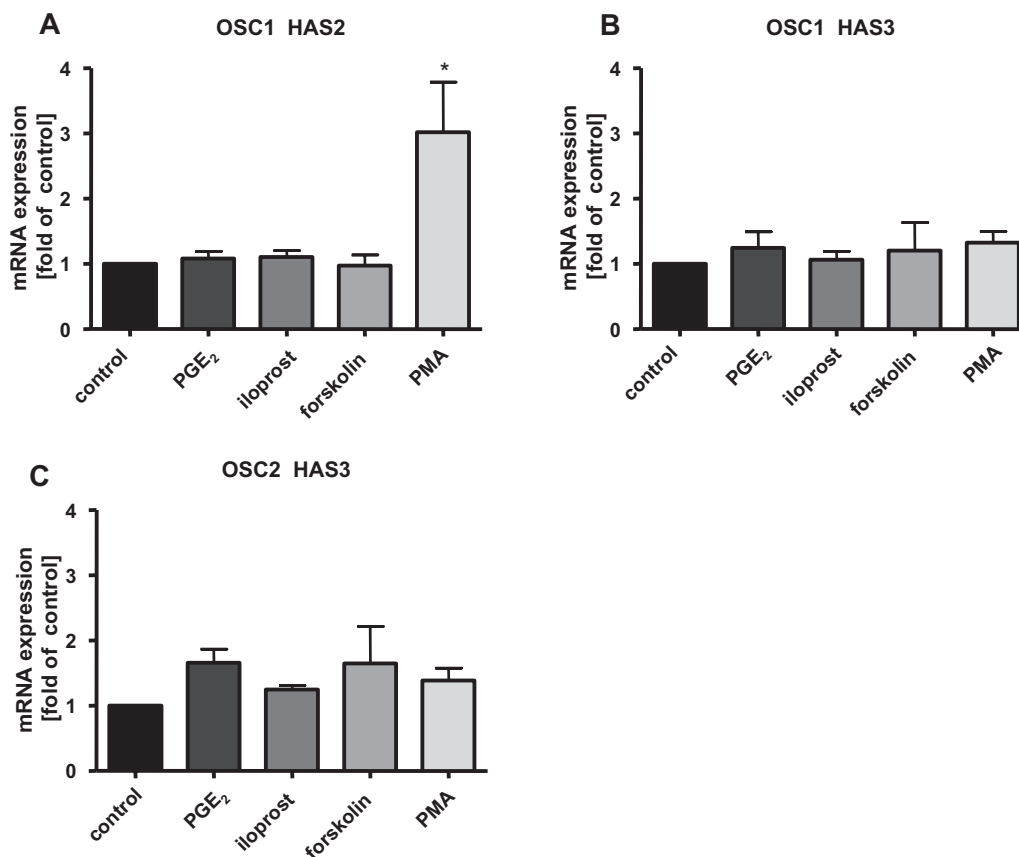
Due to technical reasons, that is, detachment of tissue sections during immunohistochemical staining, only a subset of the 69 cases could be evaluated for expression of HA (61/69), COX2 (61/69) and CD44 (58/69). Expression of COX2 and CD44 was found in 100% of informative tumour samples, whereas accumulation of HA was found in 57 out of 61 samples (Figure 1A–C). HA was detected in tumour parenchyma (Figure 1A, inset, arrowheads) and even pronounced in stromal cells (Figure 1A, asterisks). Because there is evidence that cancer cells tend to express large CD44 variants (Ponta *et al.*, 1994; 2003), an antibody detecting long CD44 variants was used. In consecutive sections CD44 and COX2 staining was observed in the cancer cells, whereas the stroma was predominantly negative (Figure 1B,C). Taken together, in the majority of cases co-localization of HA, COX2 and CD44 was observed in the cancer cells. Therefore, we



**Figure 2** Characterization of the hyaluronan system of oesophageal cancer cell lines (OSC1 and OSC2). (A) Expression of HAS2 and HAS3 in OSC1 and OSC2 cells was determined by quantitative real-time RT-PCR. In both cell lines, HAS1 mRNA expression was only barely detectable, whereas HAS2 and HAS3 could be detected in OSC1 cells at a ratio of about 1:2. In OSC2 cells, only HAS3 mRNA was detected (data not shown). (B) Affinity cytochemical staining showed membrane-bound patches of HA. Detection was performed by light microscopy (left column) and fluorescence staining (right column). The micrographs show representative reactions (scale bar marks 50 μm). (C) The quantification of HA in the supernatants of OSC1 and OSC2 cells revealed approximately equal amounts of secreted HA normalized to protein in both cell lines. Representative data from experiments are shown. HA, hyaluronic acid, hyaluronan; HAS, hyaluronic acid synthase; OSC, oesophageal squamous cell; RT-PCR, reverse transcriptase polymerase chain reaction.



**Figure 3** Characterization of the COX2/PG system in OSC1 and OSC2 cells. (A) As shown by immunoblotting, COX2 was more abundantly expressed in OSC2 cells than in OSC1 cells. One representative immunoblot out of three is shown. (B) To determine the amount of PGs secreted by OSC1 and OSC2 cells, the prostacyclin metabolite, 6-oxo-PGF<sub>1α</sub>, indicative of PG synthesis was determined in the supernatant. OSC2 cells produced approximately 12-fold more prostacyclin than OSC1 cells ( $n = 3$ , mean  $\pm$  SEM). (C) Expression of PG receptors by OSC1 and OSC2 cells was assessed by RT-PCR. Human smooth muscle cells were examined as control. All cells expressed EP<sub>1</sub>, EP<sub>2</sub>, EP<sub>4</sub> and IP receptors. Only OSC2 cells additionally expressed marked amounts of EP<sub>3</sub> receptor. Receptor bands were found at the expected sizes, EP<sub>1/2</sub> (450 bp), EP<sub>3/4</sub> (580 bp), IP (380/520 bp). Shown are the results of one representative experiment out of three. COX2, cyclooxygenase-2; hSMC, human vascular smooth muscle cell; OSC, oesophageal squamous cell; PG, prostaglandin; RT-PCR, reverse transcriptase polymerase chain reaction.



**Figure 4** HAS isoform expression in OSC1 and OSC2 cells. Cells were stimulated for 6 h with the prostacyclin (PGI<sub>2</sub>) analogue, iloprost (100 nmol·L<sup>-1</sup>), PGE<sub>2</sub> (100 nmol·L<sup>-1</sup>), forskolin (10 μmol·L<sup>-1</sup>) or with PMA (100 nmol·L<sup>-1</sup>) as a positive control. Shown are the results of one real-time RT-PCR out of three to five independent experiments (mean  $\pm$  SEM, \* $P < 0.05$ ). The HAS2/3 mRNA expression was normalized to GAPDH. Surprisingly, neither PGE<sub>2</sub> nor iloprost or forskolin induced HAS2 mRNA in either cell line [OSC1, (A); OSC2, not shown]. Only PMA led to a strong increase of HAS2 mRNA transcription in OSC1 cells (A). HAS3 mRNA levels were not altered by any stimulus (B,C), HAS1 mRNA was not detectable (not shown). HAS, hyaluronic acid synthase; OSC, oesophageal squamous cell; PG, prostaglandin; PMA, phorbol myristate acetate; RT-PCR, reverse transcriptase polymerase chain reaction.

analysed *in vitro* whether COX2-dependent PGs would induce HA production in human oesophageal cancer cells.

#### Characterization of the HA system in the oesophageal cancer cell lines OSC1 and OSC2

Two oesophageal cancer cell lines exhibiting a different expression pattern of HAS isoforms were used in the subsequent experiments. In OSC1 cells, HAS2 and HAS3 mRNA were highly expressed at a ratio of about 1:2, while HAS1 mRNA was undetectable (Figure 2A). OSC2 cells, in contrast, only expressed HAS3 (not shown). HA was regularly found in both cell lines as pericellular HA as well as secreted HA in their supernatants. As shown by affinity cytochemistry, pericellular HA was distributed in both cell types as membrane-bound patches (Figure 2B), and both cell lines produced also approximately the same amount of HA (Figure 2C).

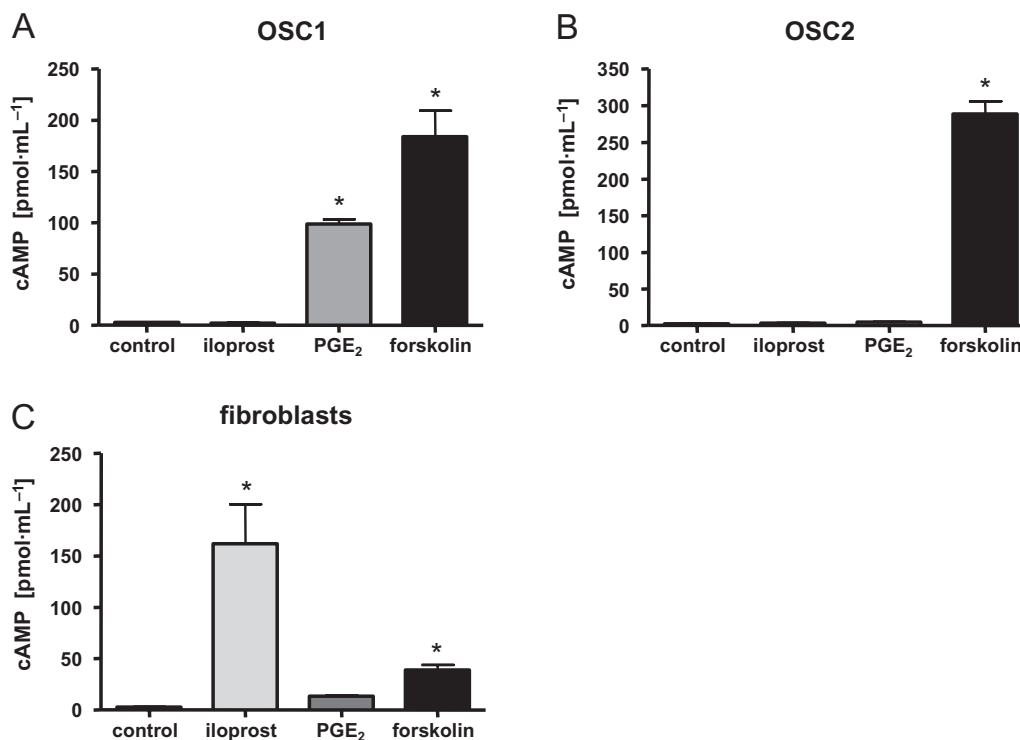
#### COX2 expression, PG production and PG receptor expression in OSC cells

Cyclooxygenase expression and activity in OSC1 and OSC2 cells were measured on protein and product level respectively. As shown by immunoblotting, COX was found to be constitutively expressed in both cell lines with OSC2 cells displaying a stronger protein expression, which is in accordance with a previous study (Zimmermann *et al.*, 1999) (Figure 3A). As an indicator for COX2 activity, the secretion of PGI<sub>2</sub> was quan-

tified. Consistent with the COX2 protein expression levels, PGI<sub>2</sub> levels were approximately 12 times higher in OSC2 cells compared with OSC1 cells (Figure 3B). Also the mRNAs of the corresponding prostacyclin (IP) and PGE receptors (EP<sub>1</sub>, EP<sub>2</sub>, EP<sub>4</sub>) were differentially expressed in both cell lines. Whereas OSC1 cells were found to abundantly express only the G $\alpha_s$ -coupled EP<sub>2</sub>, EP<sub>4</sub> and IP receptors, OSC2 cells expressed mRNA of all EP receptors and the IP receptor (Figure 3C).

#### Stimulation of OSC1 and OSC2 cells with PGs did not alter HAS isoform expression

Next we stimulated the cells with the prostacyclin (PGI<sub>2</sub>) analogue iloprost or with PGE<sub>2</sub> to assess their influence on HAS mRNA expression. Surprisingly, neither PGE<sub>2</sub> nor iloprost induced HAS1 mRNA or HAS2 mRNA in either OSC1 or OSC2 cells despite the fact that both cell lines abundantly express the IP, EP<sub>2</sub> and EP<sub>4</sub> receptors (see Figure 3C). To preclude the possibility that this deficiency is caused by a lack of cAMP induction due to inactive PG receptors, forskolin, an adenylylate cyclase activator was used to raise cAMP levels independently of PG receptor activity. However, forskolin had also no effect on the expression of the HAS2 mRNA (Figure 4A). Only phorbol myristate acetate (PMA), an activator of PKC that was used as a positive control, resulted in a strong increase in HAS2 mRNA transcription (Figure 4A). As, with the exception for PMA, none of the stimuli used was able to induce HAS2 mRNA expression, we hypothesized that HAS transcription



**Figure 5** Intracellular cAMP levels after stimulation of OSC1 and OSC2 cells and fibroblasts by iloprost, PGE<sub>2</sub> and forskolin. Intracellular cAMP levels were determined by radioimmunoassay. To examine the cAMP response, OSC1 (A) and OSC2 (B) cells were stimulated with agents known to induce cAMP levels. Human skin fibroblasts (C) were used as a positive control. Iloprost failed to induce cAMP increase in OSC1 and OSC2 cells, whereas it was able to raise cAMP levels in fibroblasts (C). PGE<sub>2</sub> stimulated cAMP production only in OSC1 cells (A), whereas forskolin led to an increase in cAMP levels in all tested cell types. Representative results from four independent experiments are shown and displayed as mean  $\pm$  SEM; asterisk indicates a *P*-value < 0.05. OSC, oesophageal squamous cell; PG, prostaglandin.

might be uncoupled from the cAMP system in oesophageal cancer cells. In addition, HAS3 mRNA levels were not altered by any stimulus in either cell line (Figure 4B,C).

#### Induction of cAMP levels in response to prostaglandins

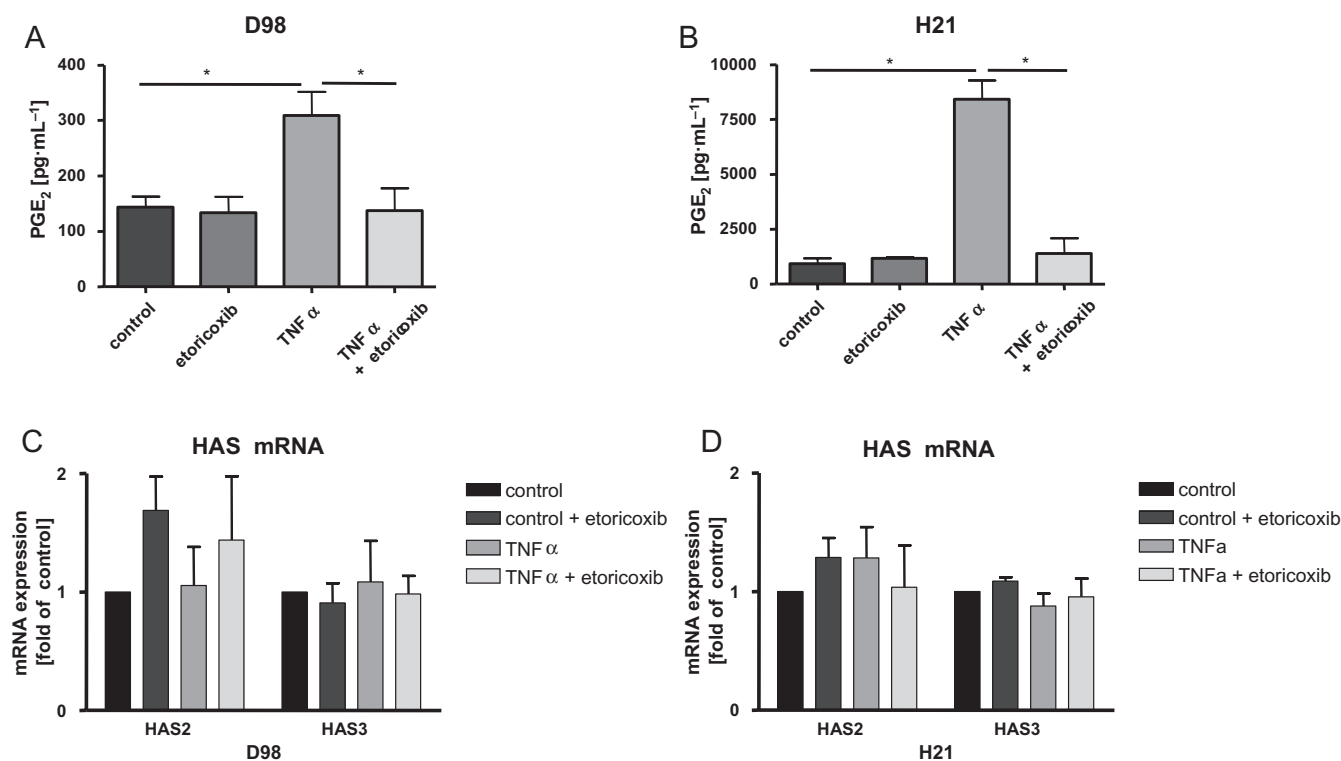
To verify that PG receptors were active and coupled to the  $G_{\alpha_s}$  subunits in OSC cells, we performed cAMP assays to measure the activation of adenylyl cyclase in response to PGs. In contrast to the positive control in hSMCs (data not shown) and fibroblasts (Figure 5C), activation of the IP receptor by iloprost did not lead to an increase of cAMP in either OSC1 or OSC2 cells, suggesting that IP receptors are inactive (Figure 5A,B). On the other hand, PGE<sub>2</sub> stimulation resulted in a strong cAMP response, however, only in OSC1 cells (Figure 5A). The lack of a cAMP response in OSC2 cells might be due to receptor down-regulation because of the high endogenous PG levels in these cells (Figure 3B). Forskolin significantly increased cAMP levels in all examined cell lines proving the presence of functional adenylyl cyclase (Figure 5A–C). Thus, in OSC2 cells, both the IP and the EP receptors (EP<sub>2</sub> and EP<sub>4</sub>) did not induce cAMP formation, which may explain why HAS2 expression was not affected by PGs in these cells. However, in OSC1 cells, PGE<sub>2</sub> strongly induced cAMP production, suggesting that cAMP-dependent activation of HAS2 gene transcription was blocked downstream of cAMP in these cells.

#### Expression of the HAS isoenzymes in HeLa cells

To verify these results also in another cellular model, cancer cells known to produce large amounts of PGE<sub>2</sub> upon cytokine-induced COX2 expression were investigated. Therefore, two HeLa cells lines (D98, H21) exhibiting a TNF- $\alpha$ -inducible, COX2-dependent PG synthesis (Figure 6A,B) (Jänicke *et al.*, 1994; Totzke *et al.*, 2003) were used to assess the effects of endogenous PGE<sub>2</sub> on HAS mRNA expression. TNF- $\alpha$  had no effect on mRNA levels of HAS2 and HAS3 isoforms in both cell lines (Figure 6C,D) despite the dramatic induction of PGE<sub>2</sub> release (Figure 6A,B). Compatible with the conclusion that autocrine release of PGE<sub>2</sub> in response to TNF- $\alpha$  did not induce HAS2 or HAS3, no effect of the COX2 inhibitor etoricoxib on HAS expression was detected (Figure 6C,D). Interestingly, etoricoxib itself caused a slight induction of HAS2 in HeLa D98 cells that was, however, not significant. Taken together, these data further substantiate our conclusion that HAS mRNA expression is not regulated by PGs in cancer cells.

#### Discussion and conclusions

Previously it has been shown in hSMC that HAS1 and HAS2 are strongly induced by stimulation of the  $G_{\alpha_s}$ -coupled IP and EP<sub>2</sub> receptors (Sussmann *et al.*, 2004; van den Boom *et al.*, 2006). Subsequently, it was reported that closure of the ductus arteriosus is also dependent on COX2-mediated PGE<sub>2</sub> synthe-



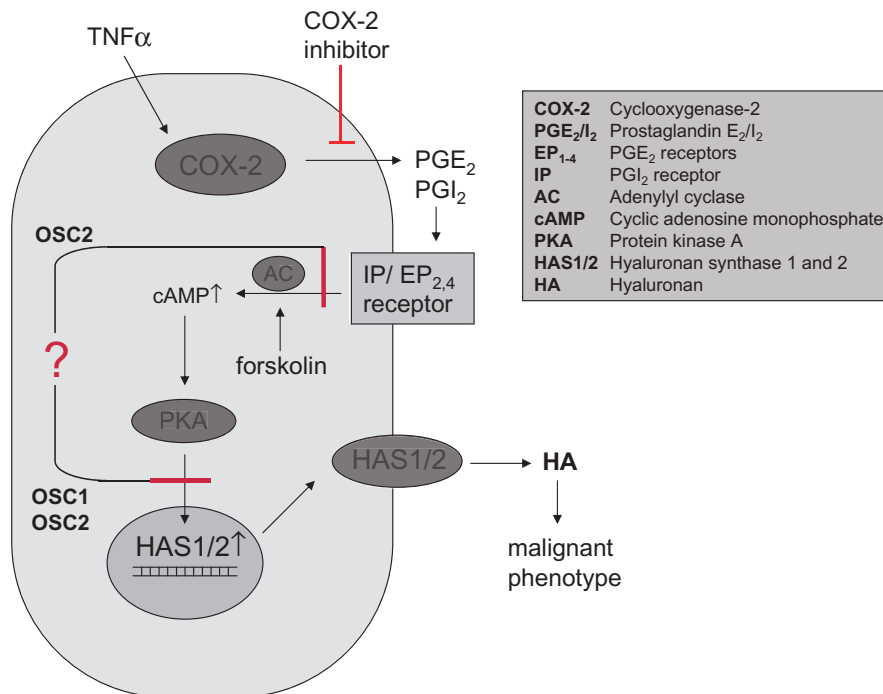
**Figure 6** HAS isoenzyme expression in HeLa cells. The HeLa cell lines D98 and H21 were stimulated with TNF- $\alpha$  (30 ng·mL<sup>-1</sup>) for 6 h in the absence or presence of the COX2 inhibitor etoricoxib (10  $\mu$ mol·L<sup>-1</sup>). The determination of PGE<sub>2</sub> levels revealed high endogenous PGE<sub>2</sub> synthesis in response to TNF- $\alpha$  that was responsive to etoricoxib (A,B). Real-time RT-PCR revealed that HAS2 and HAS3 mRNA levels were not regulated in response to COX2 and PG induction (C,D). HAS1 was not detectable (data not shown). The diagrams show representative results from three experiments (mean  $\pm$  SEM, \* $P$  < 0.05). COX2, cyclooxygenase-2; HAS, hyaluronic acid synthase; PG, prostaglandin; RT-PCR, reverse transcriptase polymerase chain reaction; TNF- $\alpha$ , tumour necrosis factor  $\alpha$ .

sis and subsequent stimulation of HAS2-mediated HA synthesis via stimulation of the EP<sub>4</sub> receptor (Yokoyama *et al.*, 2006). Furthermore, it has been known for a long time that forskolin stimulates HA synthesis in oocytes thereby stimulating the rapid expansion of the pericellular matrix of oocytes (Fischer and Schrör, 2007). These findings suggest that the PG-mediated activation of HAS expression is a regulatory pathway of general importance. The fact that the majority of human oesophageal tumour samples were characterized by the presence of HA, COX2 and CD44 was compatible with the working hypothesis that COX2-mediated PG synthesis stimulates HA production in human tumours. However, *in vitro* in OSC1 and OSC2 cells, prostacyclin and PGE<sub>2</sub> did not induce HAS1 or HAS2 mRNA expression. In contrast, PMA, known to induce HAS2 transcription via activation of PKC (Feusi *et al.*, 1999; Pienimäki *et al.*, 2001; Anggiansah *et al.*, 2003) did up-regulate HAS transcription in both cell lines.

To investigate the reason for the absence of PG-mediated activation of HAS1/2 transcription, we analysed the prostanoïd signalling pathway in more detail. RT-PCR analyses revealed that all the necessary receptors including the EP<sub>1-4</sub> and IP receptors are expressed in both OSC cell lines. Next, we pursued the question, if a loss of PG receptor-dependent induction of cAMP levels could be the reason for the lack of HAS2 induction in response to exogenous PGs. Forskolin, a direct cAMP inducer was used as positive control. Interestingly, in contrast to hSMC and fibroblasts, iloprost failed to stimulate cAMP production in either cell line. This might indicate that the IP receptor is either not expressed at the

protein level or desensitized. PGE<sub>2</sub> was able to raise cAMP levels in OSC1, but not in OSC2 cells, which are characterized by an abundant native overproduction of PGI<sub>2</sub> and PGE<sub>2</sub> (Zimmermann *et al.*, 1999) (Figure 3B), that in turn could cause a down-regulation of active EP receptors. Forskolin led to a strong increase of cAMP levels in both cell lines showing the presence of activatable adenylyl cyclase. However, forskolin failed to induce HAS2 mRNA as revealed by quantitative real-time RT-PCR (Figure 4). In conclusion, in OSC1 and OSC2 cells, the cAMP-mediated HAS2 mRNA induction might be inhibited downstream of cAMP generation. A possibility that might be considered in future studies is that epigenetic changes such as methylation of HAS2 promoter may occur. Another possibility for aberrant HAS2 mRNA expression was pointed out by Chao and Spicer who described a naturally occurring HAS2 antisense RNA (HASNT) that leads to inactivation of HAS2 gene expression (Chao and Spicer, 2005).

To further investigate the possibility that a highly elevated endogenous PG production could induce HAS2 mRNA in an autocrine manner and in turn permanently activate the HAS system, we used the two HeLa cell lines D98 and H21 that display strong COX2 induction in response to TNF- $\alpha$  (Jänicke *et al.*, 1994; Totzke *et al.*, 2003) and abundant production of PGE<sub>2</sub>. However, also in these cells no correlation between HAS mRNA expression and PGE<sub>2</sub> production was observed as indicated by the use of the COX2 inhibitor etoricoxib. These experiments show that the observed lack of response towards PG-mediated HAS expression is not restricted to oesophageal



**Figure 7** Schematic illustration of the proposed stimulation of HAS isoform regulation by prostaglandins in cancer cells. According to our working hypothesis COX2-dependent prostaglandins (PGI<sub>2</sub> and PGE<sub>2</sub>) activate the G $\alpha_s$ -coupled IP and EP<sub>2/4</sub> receptors, raise intracellular cAMP levels and in turn activate HAS1 and HAS2 expression via PKA-mediated signals. The increased HA secretion stimulates a malignant tumour cell phenotype through signalling via HA receptors CD44 or receptor of HA-mediated motility (RHAMM). The present findings indicate that in oesophageal squamous cancer cells, OSC1 and OSC2, stimulation of HAS1 and HAS2 expression by prostaglandins is inhibited either at the level of prostaglandin receptor-stimulated cAMP induction (OSC2) and/or downstream of cAMP (OSC1, OSC2).



cancer cells and might therefore be relevant for cancer cells of varying origins.

Tofuku *et al.* showed that metastasis-associated properties of cancer cells such as invasion and proliferation are enhanced by lower molecular weight forms of HA that are characteristic of HAS3 activity (Itano and Kimata, 2002). In contrast, HAS2 is believed to deliver higher molecular weight HA that supports pericellular coat formation and thus inhibits a malignant tumour cell phenotype (Tofuku *et al.*, 2006) unless it is further processed by for example hyaluronidases (Stern, 2008). This hypothesis was corroborated by Bullard *et al.* (2003) who compared the primary colon carcinoma cell line SW480 and the corresponding lymph node metastasis cell line SW620. The expression levels of HAS3 mRNA were much higher in the metastasizing cells (Bullard *et al.*, 2003). Therefore, it is tempting to speculate that the observed absence of PG- and cAMP-dependent HAS2 mRNA induction is leading to a higher relative contribution of HAS3 to HA synthesis, which might support the malignant phenotype of OSC1 and OSC2 cells.

Taken together, the present data did not support our working hypothesis that COX2-dependent PGs induce HA synthesis in oesophageal cancer (Figure 7). In OSC2 cells, this is likely to have been caused by desensitized PG receptors, which did not lead to raised cAMP levels after stimulation. In OSC1 cells, the EP receptors induce cAMP levels in response to PGE<sub>2</sub>; however, HAS1 and HAS2 were not induced. Instead, we showed that in OSC1 and OSC2 cells, PG- and cAMP-dependent induction of HAS1/2 mRNA expression was not active. In future studies it should therefore be investigated whether epigenetic changes or natural occurring antisense RNA against HAS2 suppress the expression of HAS1 and HAS2 in these cancer cells.

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## Conflict of interest

The authors state no conflict of interest.

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