

A novel mechanism for the anticancer activity of aspirin and salicylates

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Abstract. Epidemiological studies indicate that long-term aspirin usage reduces the incidence of colorectal cancer (CRC) and may protect against other non-CRC associated adenocarcinomas, including oesophageal cancer. A number of hypotheses have been proposed with respect to the molecular action of aspirin and other non-steroidal anti-inflammatory drugs in cancer development. The mechanism by which aspirin exhibits toxicity to CRC has been previously investigated by synthesising novel analogues and derivatives of aspirin in an effort to identify functionally significant moieties. Herein, an early effect of aspirin and aspirin-like analogues against the SW480 CRC cell line was investigated, with a particular focus on critical molecules in the epidermal growth factor (EGF) pathway. The present authors proposed that aspirin, diaspirin and analogues, and diflunisal (a salicylic acid derivative) may rapidly perturb EGF and EGF receptor (EGFR) internalisation. Upon longer incubations, the diaspirins and thioaspirins may inhibit EGFR phosphorylation at Tyr1045 and Tyr1173. It was additionally demonstrated, using a qualitative approach, that EGF internalisation in the SW480 cell line may be directed to endosomes by fumaryl diaspirin using early endosome antigen 1 as an early endosomal marker and that EGF internalisation may also be perturbed in oesophageal cell lines, suggestive of an effect not only restricted to CRC cells. Taken together and in light of our previous findings that the

aspirin-like analogues can affect cyclin D1 expression and nuclear factor- κ B localisation, it was hypothesized that aspirin and aspirin analogues significantly and swiftly perturb the EGFR axis and that the protective activity of aspirin may in part be explained by perturbed EGFR internalisation and activation. These findings may also have implications in understanding the inhibitory effect of aspirin and salicylates on wound healing, given the critical role of EGF in the response to tissue trauma.

Introduction

A total of ~600,000 patients annually succumb to colorectal cancer (CRC) worldwide (1), with indications that the incidence is on the rise, particularly in Asia (2,3) and possibly in the younger population in the USA (4). Regular aspirin intake significantly reduces the risk of developing CRC and other gastrointestinal tract malignancies (5,6), which has caused debate regarding regular aspirin use as a chemopreventive agent on a population basis (7,8). Aspirin usage following a diagnosis of colon cancer also has a positive outcome: Longer survival has been reported among patients with mutated-PIK3CA CRC, but not those with wild-type PIK3CA cancer (9).

Despite vigorous investigation, the molecular basis for the chemopreventive effect of aspirin remains controversial and a large number of molecular targets of aspirin and salicylates have been suggested (10). The molecular effects on cells and tissues include but are not limited to: Inhibition of cyclooxygenase (COX) activity (11-13) and protein phosphatase 2 enzymatic activity (14), inhibition of nuclear factor (NF)- κ B transcriptional activity (15), suppression of COX-2 gene transcription (16), inhibition of I κ B kinase- β (17), increased hMLH1 expression (18), selection for DNA microsatellite stability (19), altering mammalian target of rapamycin signalling (20) and a direct allosteric activation of AMP kinase (21). More recently, the breakdown product of aspirin, salicylate, has been demonstrated to inhibit CREB-binding protein/p300 acetyltransferase activity (22).

Therefore, it is reasonable to suggest that aspirin may act pleiotropically and there is some evidence that aspirin and other non-steroidal anti-inflammatory drug (NSAIDs)

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Abbreviations: CRC, colorectal cancer; EEA1, early endosome antigen 1; MAPK, mitogen-activated protein kinase; NSAID, non-steroidal anti-inflammatory drug

Key words: epidermal growth factor, epidermal growth factor receptor, aspirin, colorectal cancer, endocytosis

may also affect the epidermal growth factor (EGF) receptor (EGFR) axis. For example, metabolites of the NSAID sulindac can inhibit EGFR signalling through the inhibition of EGFR phosphorylation and decreased EGFR expression in HT29 colon cancer cells (23). A nitro-derivative of aspirin was demonstrated to inhibit EGFR signalling in human ovarian cancer cells (24). Furthermore, in COX-1-positive ovarian cancer cells cultured *in vitro*, aspirin inhibited EGF-associated protein kinase B (AKT) and extracellular signal-regulated kinase (ERK) phosphorylation (25).

In an attempt to better understand the toxicity of aspirin to CRC cells, a number of aspirin-like analogues were previously synthesised and diaspirins were identified as having the capacity to induce apoptosis in a CRC cell line (26), inhibit NF- κ B *in vitro* and cyclin D1 expression, and suppress tumour growth *in vivo* in a murine model of CRC without evidence of apparent toxicity to the animal (27). The aim of the present study was to investigate whether aspirin and analogues, including fumaryldiaspirin (F-DiA), diflunisal and salicylates, which are common breakdown products of these compounds, are able to perturb EGF endocytosis in SW480 CRC cells (28), as these cells are known to express relatively high levels of wild type EGFR (29) compared with normal colonic epithelial primary cells, but exhibit decreased expression of COX-1 and negligible levels of COX-2 (30,31). Given the role of EGF signalling in tissue repair (32), the findings of the present study may improve our understanding of the molecular basis of the action of aspirin as a chemopreventive agent and its inhibitory effect on wound healing.

Materials and methods

Chemicals and reagents. Foetal bovine serum (FBS) was purchased from PAA Laboratories (GE Healthcare Life Sciences, Little Chalfont, UK) or Labtech International, Ltd. (Heathfield, UK). Precision Plus Protein Colour standards and nitrocellulose were obtained from Bio-Rad Laboratories, Inc. (Hercules, CA, USA). Human recombinant EGF (PHG0313) was from Thermo Fisher Scientific, Inc. (Waltham, MA, USA). Alexa Fluor 555-EGF (E-35350) was from Molecular Probes; Thermo Fisher Scientific, Inc. EGFR (D38B1) XP[®] rabbit antibody (Alexa Fluor 488-conjugate; 1:100; cat. no. 5616) and EGFR rabbit antibody (D38B1; 1:100; cat. no. 4267) were from Cell Signaling Technology, Inc. (Danvers, MA, USA). Anti-early endosome antigen 1 (EEA1) antibody (1G11) Early Endosome Marker (ab70521; 1:1,000) was from Abcam (Cambridge, UK). Anti-GAPDH antibody (sc-25778; 1:1,000) was from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA). VectaShield[®] mounting medium was from Vector Laboratories, Ltd. (Peterborough, UK). All other reagents were obtained from Sigma-Aldrich; Merck KGaA (Darmstadt, Germany), unless stated otherwise. Aspirin analogues, detailed in Table I, were synthesised in-house using previously described methods (26,27).

Cell culture. Colon cancer adenocarcinoma SW480 cells were obtained from the European Collection of Authenticated Cell Cultures (Salisbury, UK) and cultured in Leibovitz L-15 medium (Thermo Fisher Scientific, Inc.) supplemented with 10% (v/v) FBS supplemented with L-glutamine-penicillin-streptomycin

(Sigma-Aldrich; Merck KGaA). The OE33 and Flo-1 oesophageal cancer cell lines were a gift from Professor Tim Underwood (University of Southampton, Southampton, UK) and were cultured in Dulbecco's modified Eagle's medium high-glucose with phenol red (Thermo Fisher Scientific, Inc.) containing 10% (v/v) FBS supplemented with L-glutamine-penicillin-streptomycin (Sigma-Aldrich; Merck KGaA). Cells were cultured in tissue culture flasks (Sarstedt, Inc., Newton, NC, USA) at 37°C in a humidified incubator with 5% CO₂, except for SW480 cells, which were cultured in the absence of CO₂. Cells were regularly passaged at 70–80% confluence. The OE33 (also known as JROECL33) and Flo-1 cells are commercially available oesophageal cancer cell lines (33). The OE33 cell line expresses COX-2 and has been utilised to examine COX-2 inhibitors with respect to proliferation (34). Both cell lines express EGFR (35). Additionally, the cell lines were investigated as they were derived from oesophageal carcinomas, and thus represent a counterpoint to a colorectal adenocarcinoma (SW480). Another aim of the present study was to elucidate whether the sensitivity to the salicylates observed in the SW480 cell line was specific to a cell line of CRC origin only, given the reported chemopreventative activity of aspirin against oesophageal cancer; reviewed in (36,37).

Cytotoxicity assay. Cell viability was assessed by MTT reduction assay (38) with modifications (39). Briefly, SW480 cells were seeded at a density of 10⁴ cell/well in a 96-well plate and incubated overnight at 37°C. Following 24 h, the culture medium was replaced with medium containing aspirin analogues (Table I) at 0.01, 0.03, 0.1, 0.3, 1 and 3 mM, and the cells were incubated at 37°C for a further 48 h. A no-treatment control was included in these experiments. Following 48 h of incubation, the medium was aspirated from each well, the cells were washed once with fresh medium to remove drugs, and 300 μ l MTT reagent (0.5 mg/ml) was added. The cells were incubated at 37°C for 3 h and the solution was replaced with 200 μ l dimethyl sulphoxide (DMSO). The plates were incubated at 37°C for a further 30 min and the conversion of MTT into formazan crystals was measured by recording changes in absorbance at 540 nm in a Multiskan Ascent microplate reader (Thermo Fisher Scientific, Inc.). The plates were protected from light throughout the procedure. Drug dosage ranged from 0 to 3 mM. GraphPad Prism 7 (GraphPad Software, Inc., La Jolla, CA, USA) was used to plot dose-response curves and to determine half maximal inhibitory concentration (IC₅₀) values. All assays were performed in duplicates (n=3 experiments) and viable cells were expressed as the percentage relative to the negative control: Cell viability = [absorbance 540 nm (treated cells-blank)/absorbance 540 nm (control cells-blank)]x100.

Immunoblot analysis. Cells (5x10⁴) were plated in 6-well tissue culture plates and incubated overnight at 37°C. Cells incubated with compounds (Table I) for the times stated were washed with ice-cold PBS at 4°C, gently scraped and suspended in 2X Laemmli sample buffer containing 10% β -mercaptoethanol (40) and 0.01% protease inhibitor cocktail (100X; 5871S; CST Biological Reagents Co., Ltd., Shanghai, China). The samples were then heated to 100°C for 20 min, centrifuged (18,000 x g; 5 min; room temperature),

Table I. List of compounds.

Common name, synonym	Chemical, formal name	Laboratory number	IC ₅₀ (mM)	Source
Salicylic acid, SA	2-hydroxybenzoic acid	Not applicable	2.6	Commercially available
Aspirin, <i>ortho</i> -aspirin	2-acetoxybenzoic acid, acetylsalicylic acid	PN502	1.8	Commercially available
Diaspirin, DiA	Bis-carboxyphenylsuccinate	PN508	0.8	(27)
Fumaryl diaspirin, F-DiA	Bis-carboxyphenylfumarate	PN517	0.2	(27)
Not applicable	<i>m</i> -bromobenzoysalicylic acid	PN524	0.4	(27)
Not applicable	Isopropyl <i>m</i> -bromobenzoysalicylate	PN529	ND	(27)
<i>meta</i> -aspirin	3-acetoxybenzoic acid	PN548	3.8	Synthesised in this study
<i>para</i> -aspirin	4-acetoxybenzoic acid	PN549	4.9	Synthesised in this study
Thioaspirin, <i>ortho</i> -TASP	2-acetylthiobenzoic acid	PN590	0.2	Synthesised in this study
<i>m</i> -thioaspirin, <i>meta</i> -TASP	3-acetylthiobenzoic acid	PN591	0.5	Synthesised in this study
<i>p</i> -thioaspirin, <i>para</i> -TASP	4-acetylthiobenzoic acid	PN592	0.8	Synthesised in this study
Diflunisal, DIF	5-(2,4-difluorophenyl) salicylic acid	Not applicable	0.08	Commercially available

IC₅₀, half maximal inhibitory concentration; ND, not determined.

analysed by discontinuous SDS-PAGE with a 10% resolving gel and electro-transferred using a Bio-Rad Mini-Trans Blot electrophoretic transfer cell (Bio-Rad Laboratories, Inc.) according to the manufacturer's instructions onto polyvinylidene difluoride membranes (Immobilon®; IPVH00010; Merck KGaA). The membranes were incubated with blocking buffer [5% (w/v) bovine serum albumin (BSA) and 0.1% (v/v) Tween-20 in TBS (TBST), or 5% non-fat milk (w/v) and 0.1% (v/v) TBST, for GAPDH] for 1 h at room temperature, followed by incubation with the following specific antibodies: EGFR rabbit-antibody (1:1,000), rabbit phosphorylated (p)EGFR Y1068 antibody (ab5644; 1:1,000; Abcam), rabbit pEGFR Y1045 antibody (2237S; 1:1,000; Cell Signaling Technology, Inc.), rabbit pEGFR Y1172 antibody (ab135560; 1:2,000; Abcam) and GAPDH rabbit antibody in blocking buffer and incubated overnight at 4°C. The membranes were washed extensively with TBST and incubated with horseradish peroxidase-conjugated anti-rabbit immunoglobulin (Ig)G (7074S; 1:2,000; Cell Signaling Technology, Inc.) for 3 h at room temperature. The membranes were then washed with TBST, prior to being visualised with Thermo Scientific™ Pierce™ ECL-plus reagent (80196; Thermo Fisher Scientific, Inc.) according to the manufacturer's instructions, using CL-Xposure™ X-ray film (34088; Thermo Fisher Scientific, Inc.). ImageJ software v1.52 (National Institutes of Health, Bethesda, MD, USA) was employed for densitometric analysis of western blots. Band intensities were quantified (n=3 experiments) and pEGFR or EGFR levels were calculated relative to GAPDH, the loading control. pEGFR levels were also calculated relative to total EGFR. Data points represent mean ± standard error of the mean. Statistical analysis was performed using one-way analysis of variance followed by Dunnett's post hoc test.

Microscopy. For EGF clustering, SW480 cells were seeded onto glass coverslips at ~0.3x10⁶ cells/well and cultured in Leibovitz L-15 medium containing 10% (v/v) FBS supplemented with

antibiotics to 70-80% confluence in 6-well tissue culture plates. Cells were then cultured in serum-free L-15 medium with antibiotics for 48 h. Monolayers were cooled to 4°C and incubated for 60 min in the presence of 100-500 ng/ml Alexa Fluor 555-EGF at 4°C. For experiments testing compounds, serum-starved cells were pre-incubated at 37°C for 30 min in the absence or presence of compounds (Table I), followed by 15 min at 4°C prior to the addition of an equal volume of cold (4°C) serum-free culture medium containing 200 ng/ml Alexa Fluor 555-EGF and incubation for 60 min allowing EGF binding. Prolonged incubation previously facilitated recruitment of ligand-receptor complexes into clathrin-coated pits (41,42). Cells were warmed to 37°C over 25-30 min to stimulate EGF and EGFR internalisation, washed with PBS, fixed for 5 min with cold acetone/methanol (-20°C; 1:1, v/v) and the coverslip was then placed onto a drop of VectaShield mounting medium prior to immediate epifluorescence or confocal microscopy at x400 magnification. For preliminary experiments, cells were incubated at 37°C with 1 mM aspirin, Di-A, F-DiA or with 0.5 mM PN529 (Table I) for 30 min in serum-free medium or with an equivalent final concentration of DMSO as described above, prior to the addition of fluorescently-labelled EGF. Epifluorescence microscopy was carried out using an Olympus BX61 microscope with TXRED filter set (excitation, 580-590 nm; emission, 615-630 nm; Olympus Corporation, Tokyo, Japan). For confocal microscopy a Zeiss LSM510 Meta or a Zeiss LSM880 with Airyscan (Zeiss GmbH, Jena, Germany), equipped with an argon excitation laser (458, 488 and 514 nm), using a 63 x 1.4 NA oil immersion objective. DAPI (Dako; Agilent Technologies, Inc., Santa Clara, CA, USA) was used as a nuclear counter stain (room temperature, ≥30 min) in certain experiments

For EGFR clustering, SW480 cells were cultured on glass coverslips in 6-well tissue culture plates to 70% confluence and serum-starved for 48 h. For experiments testing compounds, cells were pre-incubated at 37°C for 30 min in the absence or presence of the drug, followed by 15 min at 4°C prior to

the addition of equivalent volume of cold (4°C) serum-free culture medium containing 250 ng/ml (stock) recombinant (r)EGF (cat. no. E9644; Sigma-Aldrich; Merck KGaA, Darmstadt, Germany) to achieve a final concentration of 125 ng/ml rEGF. The cells were then incubated for 60 min at 4°C to allow EGF binding. The cells were warmed to 37°C over 25 min to stimulate EGF clustering and EGFR internalisation, washed with PBS, fixed for 5 min with acetone/methanol (-20°C; 1:1, v/v) and incubated with blocking buffer for 60 min at room temperature. The cells were then incubated with anti-EGFR (D38B1) XP® rabbit antibodies (Alexa Fluor 488-conjugate; 1:100) for 60 min in blocking buffer at 37°C, washed with PBST (0.02%) and the coverslips were placed onto a drop of Prolong Gold antifade reagent with DAPI mounting medium (cat. no. P36935; Invitrogen; Thermo Fisher Scientific, Inc.) prior to fluorescence microscopy (x400). Aspirin, DiA, F-DiA and PN529 were added at a final concentration of 0.5 or 1 mM to independent wells in 6-well plates. The controls without aspirin analogue were fixed following EGF binding for 60 min (no EGF internalisation) or following warming of the cells for 25 min at 37°C (with EGF internalisation).

For EEA1 staining, SW480 cells were seeded onto glass coverslips at $\sim 0.3 \times 10^6$ cells/well and cultured to $\sim 70\%$ confluence in 6-well tissue culture plates and serum-starved for 48 h. For experiments for preliminary testing of compounds, serum-starved cells were preincubated at 37°C for 30 min in the absence or presence of compounds, followed by 15 min at 4°C, prior to addition of the equivalent volume of cold serum-free culture medium containing 200 ng/ml Alexa Fluor 555-EGF (final concentration of 100 ng/ml) and incubation on ice for 1 h to allow EGF binding. In later experiments, the concentration was reduced to 20 ng/ml. Cells were warmed to 37°C over 30 min to stimulate EGF internalisation, washed with cold (4°C) PBS thrice and fixed for 5 min with acetone/methanol (1:1, v/v) at -20°C on ice. Cells were washed with cold PBS and blocked in blocking buffer [3% (w/v) BSA and 0.2% (v/v) Tween-20 in PBS] for 2 h at room temperature in the dark. Anti-EEA1 antibody was added to cells at 1:1,000 dilution and incubated at 4°C overnight with gentle rocking. Cells were washed with cold (4°C) blocking buffer [3% (w/v) BSA and 0.2% (v/v) Tween-20 in PBS] thrice and incubated with goat anti-mouse IgG H&L-fluorescein isothiocyanate (ab6785; 1:1,000; Abcam) secondary antibody for 2 h at room temperature. Cells were then washed with cold blocking buffer [3% (w/v) BSA and 0.2% (v/v) Tween-20 in PBS] thrice and with cold PBS prior to placing coverslips onto a drop of VectaShield mounting medium. Microscopy was performed using a Zeiss LSM 880 confocal microscope, equipped with 405 and 561 nm excitation lasers using a 40x/1.30 NA oil immersion DIC M27 objective.

Image acquisition and processing. Images were acquired and processed with Olympus compatible software (IPLab Imaging software v3.1; BD Biosciences, San Jose, CA, USA) or Zeiss software [ZEN 2.3 lite (blue edition), or Zeiss LSM Image Examiner Version 3.2.0.115; both Carl Zeiss AG, Oberkochen, Germany] and minor brightness, contrast and tonal alterations were performed with Adobe Photoshop CS6 (Adobe Systems, Inc., San Jose, CA, USA) or GIMP (<https://www.gimp.org/>).

For semi-quantitative determination of EGF internalisation, the goal of development of the proximity and density quotient (PDQ) was to establish a metric that provides a quantitative estimate of the degree of localisation of labelled EGF to the cell nucleus. A way of doing this would be to determine the mean density value of the protein molecules and another density value in the vicinity of the nuclei and calculate a quotient. A visual inspection of the data to be analysed demonstrated an effect, whereby the clustering around one nucleus would have a major effect on the overall density around a neighbouring nucleus. For this reason, it is necessary to count only those molecules, which are unambiguously within the influence of the nucleus being studied. This is achieved by defining the region of interest for each nucleus. Each molecule is allocated to a region defined by the closest nucleus. Within this region, the evenness of the distribution of proteins is measured by counting them within an area around each nucleus. The outer area is a torus extending from the full radius of the measurement area to half of that radius. The inner area is a disc within the torus. The PDQ for an individual nucleus is the ratio of the number of molecules counted in the outer area to those counted in the overall area. A larger PDQ indicates reduced clustering of molecules around the nucleus. Formally, the calculation of the PDQ is performed as follows: A set of nuclei, N , and a set of protein molecules, M were defined. A set of protein molecules associated with each nucleus, n , was identified as:

$$M_n = \{m \in M : n \in N \wedge (\forall n_- \in (N - \{n\}) : \text{dist}(m, n_-) > \text{dist}(m, n))\},$$

with $\text{dist}(m, n)$ as the Cartesian distance of the molecule m from the centroid of the nucleus n .

Then, sets of molecules proximate to each nucleus were identified as:

$$P_n = \{p \in M_n : \text{dist}(p, n) < 2k \text{ rad}(n)\}$$

$$Q_n = \{p \in M_n : \text{dist}(p, n) < k \text{ rad}(n)\},$$

with $\text{rad}(n)$ as the mean radius of the nucleus and k as an empirical constant, selected to maximise discrimination of PDQ in the control sets.

The PDQ for a molecule is calculated as the ratio of the difference of the cardinalities of P_n and Q_n and the cardinality of P_n .

$$PDQ_n = \frac{|P_n| - |Q_n|}{|P_n|}$$

The tools used to calculate the PDQ were CellProfiler (43) and the Python programming language (<https://www.python.org/>). CellProfiler was used to identify, count and locate the two objects of interest, nuclei and protein molecules. The staining of the slides provided a clear contrast between the red-rendered proteins and the blue-rendered nuclei, thus the first operation was to split the image by colour channel. The nuclei required processing in order to be reliably and unambiguously identified, which included removal of objects <50 pixels across, smoothing and thresholding. Without the latter steps, the segmentation algorithm would split single nuclei into several objects. Identification of protein molecules required suppression of objects <3 pixels. It is to be expected that different slide processing would require tuning of the image processing in order to reliably identify the two types of objects. CellProfiler provided tables of data listing each of the

objects identified, its position and size. A Python program was written, which implemented the algorithm described above. It was used to further process these data. The value of k , which provided best differentiation between the positive and negative control sets, was 4.

Compound synthesis. Full details of all the compounds referred to in the present study are provided in Table I. Aspirin analogues were, wherever possible, synthesised as previously described (26,27). *Meta*-aspirin (PN548) and *para*-aspirin (PN549) were made by a standard aspirin synthesis using 3-hydroxy and 4-hydroxybenzoic acids as starting materials, in lieu of salicylic acid. The three isomeric thioaspirin analogues (PN590, PN591 and PN592, represented as *ortho*-TASP, *meta*-TASP and *para*-TASP respectively) were prepared in good yields (~70%) from the corresponding mercaptobenzoic acids, by reaction with acetic anhydride in ice-cold sodium hydroxide solution, as previously described (44). In the present study, sodium hydroxide, rather than the potassium salt, was used without observing any changes. While the previously published method (39) only describes the preparation of the *meta*- and *para*-isomers (*meta*-TASP and *para*-TASP) it was demonstrated that the method also yields the *ortho*-isomer (*ortho*-TASP). The crude products in each case were demonstrated by thin layer chromatography and melting point comparison to be suitable for use without any further purification. Details of the additional compounds synthesised specifically for use in the present study are provided below. All percentage yields were comparable to the published values and within 15%. NMR spectra were recorded on a Jeol ECS-400 instrument.

***Meta*-aspirin (3-acetoxybenzoic acid):** White powder; melting point (mpt), 131-134°C (literature, 131-134°C); ^{13}C -NMR (CDCl_3) δ =171.19, 169.32, 150.75, 130.88, 129.68, 127.72, 127.33, 123.53, 21.14 ppm; Infra Red (IR), 3,400-2,200, 1,758, 1,675, 1,584 cm^{-1} .

***Para*-aspirin (4-acetoxybenzoic acid):** White powder; mpt, 183-185°C (literature, 187°C; 190-194°C); ^{13}C -NMR (CDCl_3) δ =171.31, 168.96, 155.06, 131.96, 126.88, 121.85, 21.25 ppm; IR 3500-2200, 1754, 1677, 1602 cm^{-1} .

Thioaspirin (*ortho*-TASP): White powder; mpt, 122-125°C (literature, 125.5-126°C) (45); ^{13}C -NMR (CDCl_3) δ =193.27, 171.48, 136.82, 132.88, 132.66, 131.92, 129.49, 129.40, 30.45 ppm; IR 3,380-1,960, 1,694, 1,676, 1,582, 1,565 cm^{-1} .

***Meta*-thioaspirin (*meta*-TASP):** White powder; mpt, 153-156°C (literature, 152-153°C) (44); ^{13}C -NMR (CDCl_3) δ =193.29, 171.09, 139.85, 136.16, 131.19, 130.43, 129.46, 128.88, 30.38 ppm; IR 3,380-1,990, 1,685 (overlapping thioester and -COOH C=O spectra), 1,583, 1,573 cm^{-1} .

***Para*-thioaspirin (*para*-TASP):** White powder; mpt, 200-202°C (literature, 202.5-203.5°C) (44); ^{13}C -NMR (CDCl_3) δ =171.31, 168.96, 155.06, 131.96, 126.88, 121.85, 21.25 ppm; IR 3,380-1,950, 1,683 (overlapping thioester and -COOH C=O spectra), 1,592, 1,566 cm^{-1} .

Results and Discussion

EGF and EGFR internalisation. EGFR is an ~170-kDa transmembrane receptor tyrosine kinase, belonging to the ErbB family of signalling receptors, whose ligands include

EGF and transforming growth factor- α , with roles in cellular proliferation, survival and differentiation (46,47). Dysfunction in the EGFR axis as a consequence of receptor overexpression, amplification and/or mutation has been reported in colorectal, breast, brain, lung, pancreatic, oesophageal and head and neck neoplasias (48-52). EGFR and NF- κ B signalling are intimately linked (53,54) and it has been reported that the aspirin analogues DiA and F-DiA can perturb NF- κ B activity in SW480 cells and that salicylates can potentially antagonise wound healing. Therefore, the aim of the present study was to investigate the effect of aspirin and aspirin-like molecules (26,27) on EGF endocytosis in the context of a rapid cell biological effect, in contrast to many experiments that examine the consequences of compound exposure of cells and tissues over protracted time frames (days). Compound toxicity was determined by the MTT assay and the IC_{50} values are presented in Table I. The aspirin analogues synthesised and tested contain modifications to either the number of rings, to the position of the acetyl group and/or have a substituted thiol group. Diflunisal is a difluorophenyl derivative of salicylic acid and is a clinically utilised NSAID with analgesic and anti-inflammatory properties.

In preliminary experiments to assess compound effects on EGF internalisation, a commercially available fluorescence-labelled, pH-insensitive conjugate of EGF was used. Serum-starved SW480 cells were incubated with aspirin analogues, cooled on ice to inhibit endocytosis and allowed EGF-conjugate binding for 1 h prior to tracking the internalised EGF by confocal and DIC microscopy (Fig. 1) and immunofluorescence analysis (data not shown). In SW480 cells incubated in the absence of compounds, following ~25 min of warming to 37°C, EGF was observed to internalise with clustering often in a perinuclear location, with the nucleus clearly visible (Fig. 1). However, in the presence of aspirin, DiA and F-DiA, internalisation and perinuclear clustering was inhibited. In cells incubated with DiA, substantial clustering was noted to persist at or near the membrane, whereas with F-DiA, staining was notably diffused. These data suggested that compounds based on salicylate may perturb EGF internalisation and affect signalling, given that correctly regulated EGF internalisation and signalling via endosomes are associated (55,56). To examine this phenomenon further, serum-starved SW480 cells were incubated with compounds prior to stimulation with recombinant human EGF (125 ng/ml) and EGFR localisation was probed by immunofluorescence analysis using a commercially available anti-EGFR monoclonal antibody. Peripheral, presumably membranous staining, was clearly visible in the absence of warming (Fig. 2A), indicating that the EGFR is robustly expressed in the SW480 cell line, as previously reported (29). Internalised punctate staining of the EGFR demonstrated close proximity to the nucleus when the cells were warmed to 37°C within 25 min (Fig. 2B). In the presence of aspirin, DiA and F-DiA, during the internalisation of the receptor, staining was substantially more diffuse with all compounds (Fig. 2C-E). Furthermore, when cells were incubated with PN529 (isopropyl *m*-bromobenzoysalicylate), a toxic brominated salicylate (27), EGFR staining was markedly dissimilar compared with the control cells (Fig. 2B and F). Clustering of labelled EGFR to perinuclear locations was consistently observed following a 25-min incubation in the

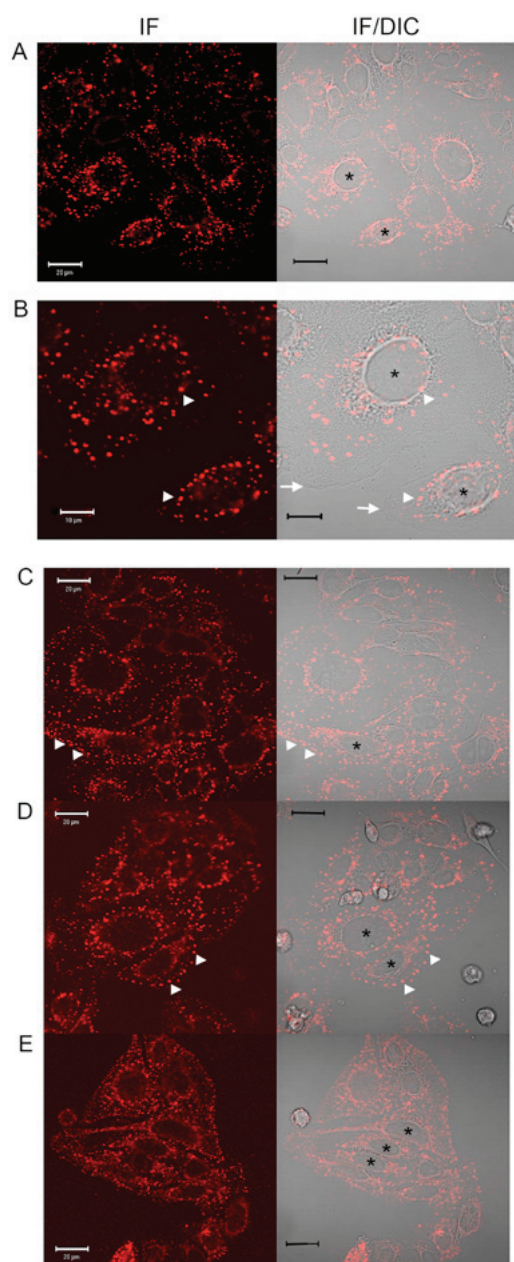


Figure 1. Effects of aspirin, DiA and F-DiA on EGF internalisation in SW480 cells. Internalisation of Alexa Fluor 555-EGF following 25 min at 37°C examined by (A and B) confocal microscopy (left panel) and highlighting the clustering, with merged DIC image (right panel). Internalisation of Alexa Fluor 555-EGF following 25 min at 37°C following preincubation for 30 min at 4°C with (C) 1 mM aspirin, (D) 1 mM DiA or (E) 1 mM F-DiA and stimulation by warming of the cells to 37°C. *Indicates nuclei for the interpretation of DIC microscopy, arrowheads indicate tagged EGF and arrows highlight the plasma membrane. DiA, diaspirin; F-DiA, fumarylidaspirin; EGF, epidermal growth factor; DIC, differential interference contrast; IF, immunofluorescence.

absence of compounds in cells warmed to 37°C. However, in the presence of aspirin and, more prominently, in the presence of DiA, F-DiA and isopropyl (m-bromobenzoysalicylate), discrete perinuclear clustering was substantially inhibited. These data suggest that aspirin and analogues rapidly perturb EGFR internalisation, and this observation may not solely be caused by a reduction in binding, as staining observed with fluorescence-tagged EGF was robust even following aspirin, F-DiA, PN524 or *ortho*-TASP treatment (data not

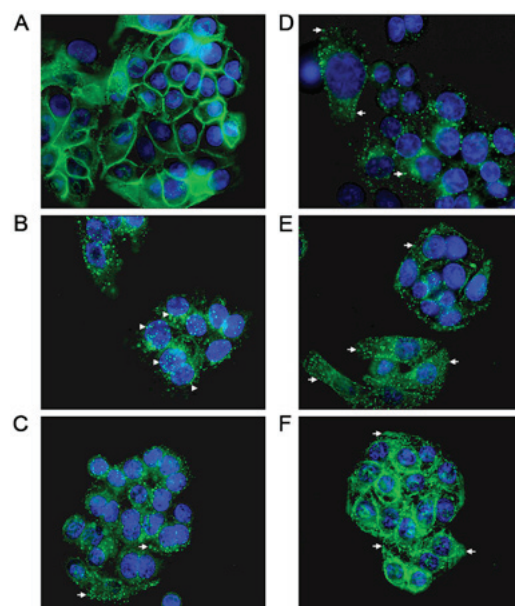


Figure 2. Immunofluorescence analysis of the effects of aspirin and aspirin-like analogues on EGFR internalisation in SW480 cells. Serum-starved SW480 cells were incubated with aspirin or diaspirin compounds and human EGF. The samples were warmed to stimulate EGF internalisation and stained using the EGFR XP[®] rabbit antibody (Alexa Fluor 488-conjugate; green). (A) Vehicle control at 4°C with cells incubated with 125 ng/ml EGF. Cells stimulated with 125 ng/ml EGF and warmed to 37°C over 25 min with (B) no preincubation, (C) 1 mM aspirin, (D) 1 mM DiA, (E) 1 mM F-DiA and (F) 1 mM isopropyl-bromobenzoysalicylate preincubation. Cells were fixed and observed by epifluorescence microscopy. Blue highlights DAPI-stained DNA. Pronounced clustering of the EGFR to a perinuclear location occurs upon warming cells (arrowheads). In the presence of compounds, the clustering is less perinuclear and/or more diffuse (arrows). Magnification, x600. EGFR, epidermal growth factor receptor; EGF, epidermal growth factor; DiA, diaspirin; F-DiA, fumarylidaspirin.

shown). Whilst the concentration of the tested compounds in the experiments may appear high in comparison with other chemotherapeutic agents, a range of 1-5 mM or above is not uncommon in *in vitro* experiments investigating the molecular action of aspirin or salicylate (21,22,57,58), with levels of 0.5-2 mM also reported as being physiologically or therapeutically relevant by a number of investigators (59-62).

EGF endocytosis in the presence of salicylate derivatives. Ligand binding leads to conformational changes in EGFR and EGFR dimerization (63-65), autophosphorylation of the cytoplasmic tyrosine kinase domain with subsequent downstream signalling through the RAS [mitogen-activated protein kinase (MAPK)], phosphoinositide 3-kinase (AKT) and janus kinase (JAK; signal transducer and activator of transcription factor 3) pathways (53,66,67). Concomitantly, the ligand receptor complex is internalised via clathrin-dependent and -independent routes, depending on EGF levels, with clathrin-mediated endocytosis prolonging signalling (68-70). Signalling from endosomes with activated EGFR may persist (71,72) and the activated EGFR may ultimately be recycled or degraded (73). Indeed, internalised endosomes may be considered as signalling platforms (56). An early destination for internalisation is the EEA1-positive endosomal compartment depending on whether or not a clathrin-mediated endocytosis occurred (74). Clustering of late endosomes is

at a perinuclear location and mislocalised endosomes can compromise signalling quality (55). In addition, it has been suggested that EGFR can translocate to the nucleus and signal therein (75-78).

EEA1 is a Rab5 effector, intimately involved in tethering, docking and fusion of early endosomes (79) and mediating crosstalk between signalling pathways (80). It was therefore examined whether spatial and temporal regulation of the internalisation of EGF was perturbed by examining the colocalisation of Alexa Fluor 555-labelled EGF (100 ng/ml) and EEA1 following a chase in the absence and presence of aspirin and aspirin analogues. This was performed while additionally controlling pH alterations with HEPES (10 mM at pH 8; Fig. 3) or phosphate buffers (data not shown) to minimise acidification caused by the addition of the aspirin derivatives. Buffering did not appear to attenuate the effects exhibited by aspirin and aspirin analogues. Peripheral EGF staining, interpreted as membrane-localised, was apparent without a chase, whereas EGF internalisation to a perinuclear location occurred in the absence of compounds upon warming to 37°C in the vehicle control group (Fig. 3). In the presence of compound at 0.5 mM, a concentration used to reduce any non-specific effects caused by acidification, and under buffered conditions, aspirin or aspirin analogues rapidly perturbed EGF internalisation (Fig. 3A). Colocalisation of EEA1 and EGF was consistently enhanced, as evidenced by dual staining, when cells were incubated with F-DiA, suggesting that the pathway of internalisation of EGF in the presence of F-DiA is substantially altered in comparison to aspirin and PN524, a brominated salicylate exhibiting an intermediate toxicity between aspirin and F-DiA (27) to SW480 cells. Furthermore, endosomal EEA1 staining was enhanced, when cells were incubated with F-DiA, based on the intensity of the fluorescence in the perinuclear location.

EGF internalisation (at 100 ng/ml) was analysed semi-quantitatively with PDQ measurements of the labelled EGF relative to the nucleus (Fig. 3B) in the presence of 0.5 mM aspirin, PN517 (F-DiA) or diflunisal. A larger PDQ indicated reduced clustering of molecules around the nucleus. Significant perturbation of internalisation was noted with all three compounds. It was previously demonstrated that F-DiA is significantly more effective at inducing apoptosis in CRC cells compared with aspirin, strongly reduces cyclin D1 levels and inhibits NF- κ B activity *in vitro* (27).

Given that endocytosis of the EGFR has been proposed to occur via a number of pathways, depending on the saturability of EGF ligand-induced receptor internalisation (42,81,82), the cells were additionally stimulated with relatively low EGF concentrations (20 ng/ml) under buffered conditions (HEPES). These experiments were performed to elucidate the influence of the tested compounds on EGF and EGFR internalisation and further utilised EEA1 staining (Fig. 4). Similar to experiments using 100 ng/ml EGF, aspirin and F-DiA perturbed EGF internalisation compared with the vehicle control and colocalisation of EGF with EEA1 was markedly enhanced as a consequence of a brief incubation with 0.5 mM F-DiA.

The universality of the phenomenon of perturbing EGF internalisation at 20 ng/ml was examined, with a wider range of salicylate derivatives (at 0.5 mM), including diflunisal. Dysregulation in EGF internalisation was noted with all compounds under HEPES-buffered conditions (Fig. 5)

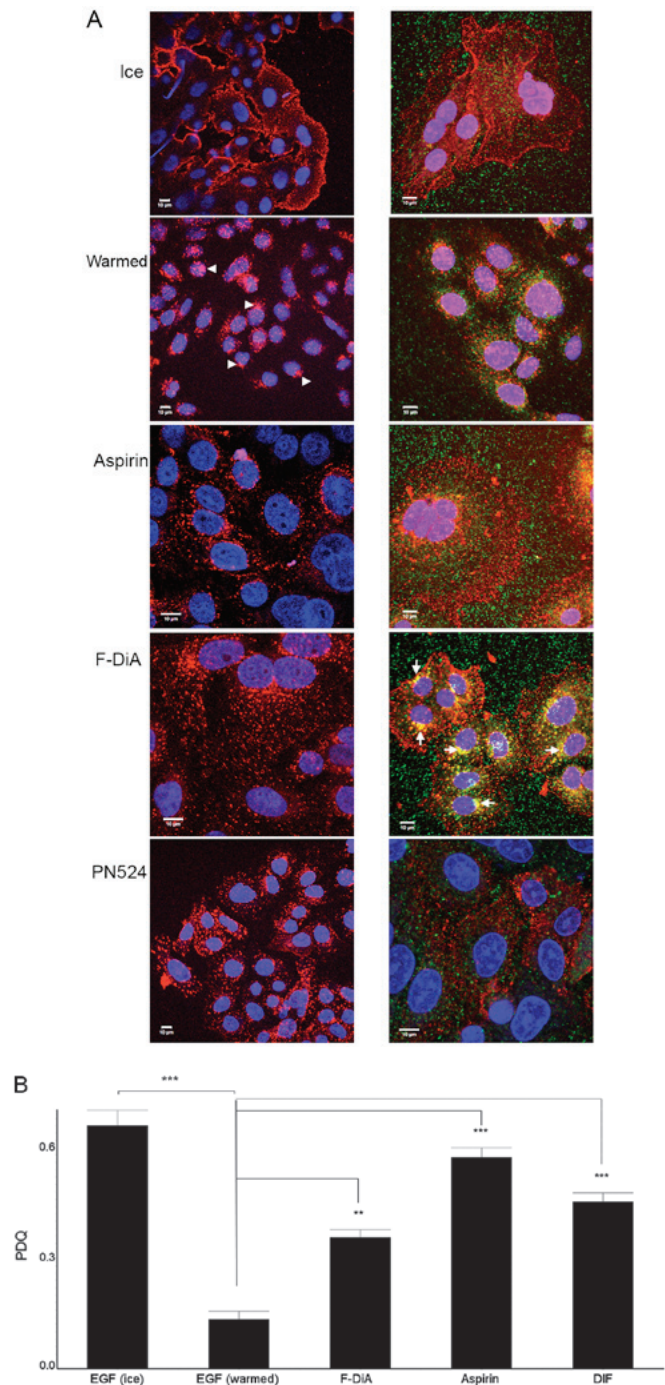


Figure 3. (A) Effect of aspirin and aspirin analogues under HEPES-buffered conditions on EGF internalisation (100 ng/ml) and EGF colocalisation with EEA1. Confocal analysis of SW480 colorectal cancer cells incubated with Alexa Fluor 555-EGF and preincubated in the absence or presence of 0.5 mM compound. Analysis was performed in the absence of compound with no warming and no EGF internalisation, and for cells stimulated with warming (37°C) with and without indicated compounds. EGF fluorescence only (left panel). Merged image for EGF and stained for EEA1 (right panel). Images were acquired at 405 nm for DAPI nuclear stain (blue), 561 nm for Alexa Fluor 555-EGF (red) and 488 nm for fluorescein isothiocyanate for EEA1 (green). Internalisation of EGF to a perinuclear location was noted upon warming to 37°C (arrowheads). In the presence of compounds, internalisation was perturbed, and with F-DiA colocalisation of EGF with EEA1 was observed (arrows). (B) Nuclear PDQ for labelled EGF was determined by image analysis of three repeated experiments; data points represent mean \pm standard error of the mean. Statistical analysis comparing the PDQ of unstimulated and warmed cells without and with compounds was performed using Kruskal-Wallis with Dunn's post-hoc test. ** $P < 0.01$ and *** $P < 0.001$. EGF, epidermal growth factor; EEA1, early endosome antigen 1; F-DiA, fumarylaldispirin; PDQ, proximity and density quotient; DIF, diflunisal.

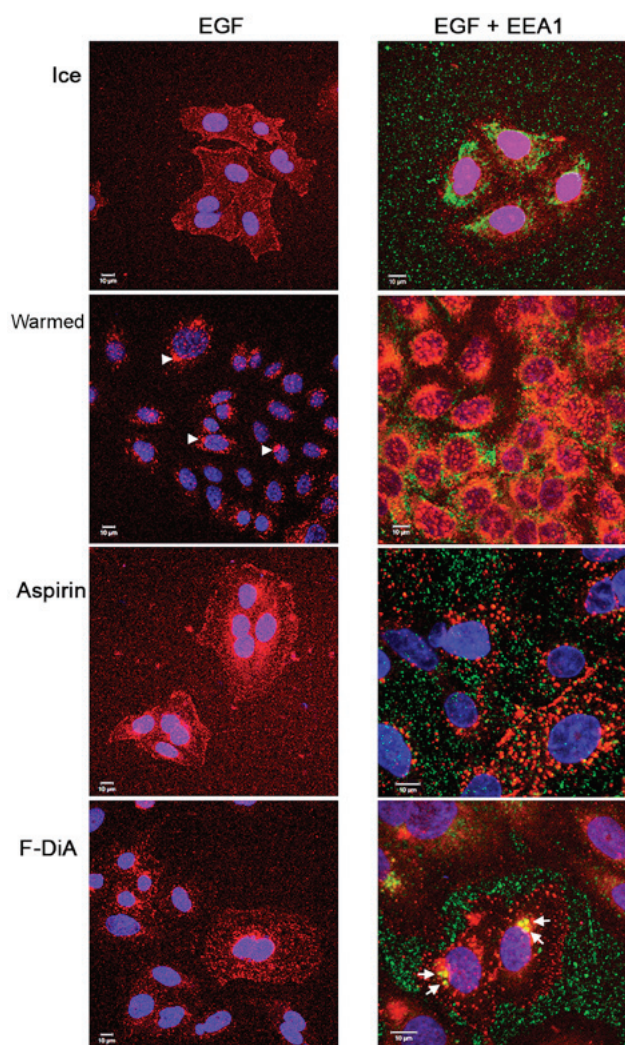


Figure 4. Effect of aspirin analogues in HEPES-buffered conditions on EGF (20 ng/ml) internalisation and EGF colocalisation with EEA1. Confocal analysis of SW480 colorectal cancer cells incubated with Alexa Fluor 555-EGF and preincubated in the absence or presence of 0.5 mM compound. Analysis was performed in the absence of compound with no warming and no EGF internalisation and for cells stimulated with warming (37°C) with and without indicated compounds. EGF fluorescence only (left panel). Merged image for EGF and stained for EEA1 (right panel). Images were acquired at 405 nm for DAPI nucleic stain (blue), 561 nm for Alexa Fluor 555-EGF (red) and 488 nm for fluorescein isothiocyanate for EEA1 (green). Representative images were taken at $\times 40$ magnification in oil/1.30 NA. oil immersion objective lens ($n=3$). Internalisation of EGF to a perinuclear location was noted upon warming (arrowheads). In the presence of compounds, internalisation was perturbed and with F-DiA colocalisation of EGF with EEA1 was also observed (arrows). EGF, epidermal growth factor; EEA1, early endosome antigen 1; F-DiA, fumarylidi aspirin.

compared with the vehicle control, including salicylic acid *per se* (Fig. 5A), aspirins with modifications to the position of the acetyl group (Fig. 5B and F) and/or compounds with an additional thiol group (Fig. 5D, G and H). When internalisation occurred in the presence of compound, it was often more diffuse than normal and the outline of the cell membrane was visible in some instances, a phenomenon particularly pronounced when the cells were incubated with diflunisal (Fig. 5E). Images were analysed and the PDQ of the labelled EGF relative to the nucleus was determined (Fig. 5I). In summary the data strongly suggested that salicylates and derivatives, at modest concentrations, rapidly altered EGF internalisation

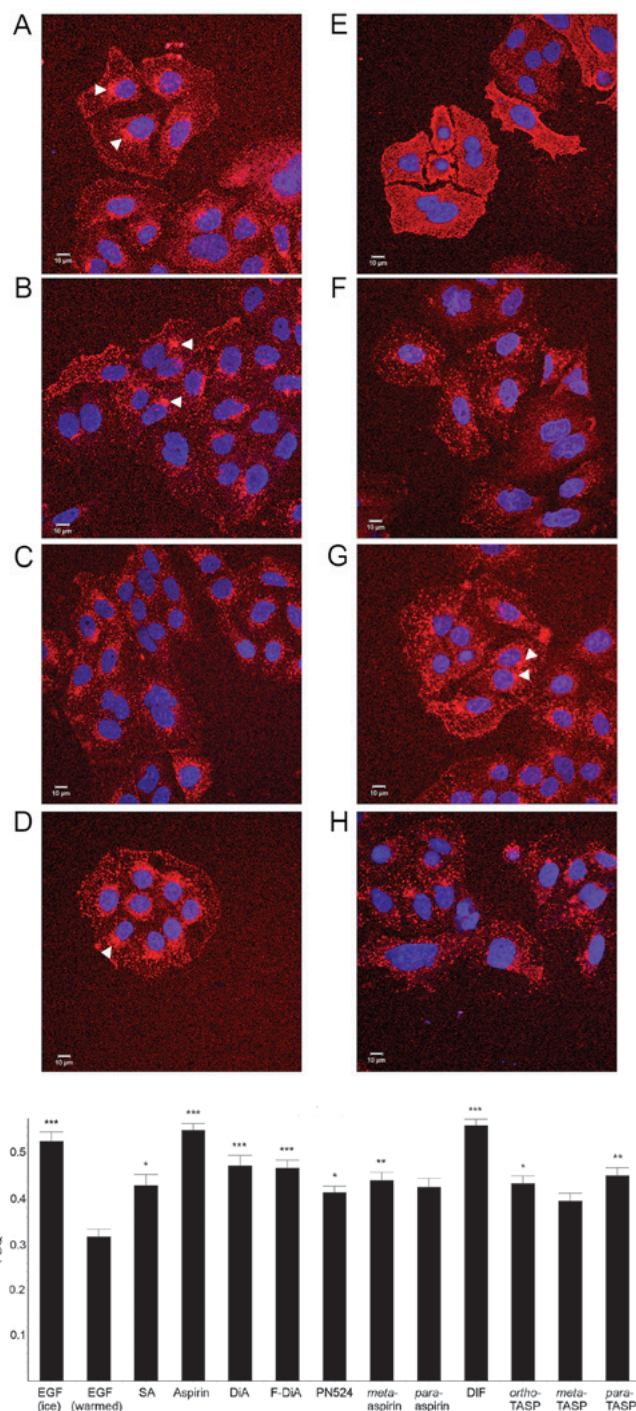


Figure 5. Effect of salicylates and aspirin analogues on EGF internalisation (20 ng/ml). Confocal analysis of SW480 colorectal cancer cells incubated with Alexa Fluor 555-EGF and preincubated in the absence/presence of 0.5 mM of the following compound under HEPES-buffered conditions: (A) SA, (B) *meta*-aspirin, (C) DiA, (D) *meta*-TASP, (E) DIF, (F) *para*-aspirin, (G) *ortho*-TASP and (H) *para*-TASP. Images were acquired at 405 nm for DAPI nucleic stain (blue) and 561 nm for Alexa Fluor-EGF (red). Representative images were taken at $\times 40$ magnification, with an oil/1.30 NA oil immersion objective ($n=3$). Limited clustering of EGF to a perinuclear location occurred upon warming cells (arrowheads), but was absent with diflunisal. (I) Nuclear PDQ of labelled EGF was determined by image analysis of three repeated experiments; data points represent mean \pm standard error of the mean. Statistical analysis comparing the PDQ of unstimulated and warmed cells with and without stated compounds was performed using Kruskal-Wallis with Dunn's post-hoc test. * $P<0.05$, ** $P<0.01$ and *** $P<0.001$ vs. EGF (warmed). EGF, epidermal growth factor; PDQ, proximity and density quotient. SA, salicylic acid; DiA, diaspirin; DIF, diflunisal; *ortho*-TASP, *ortho*-thioaspirin; *meta*-TASP, *meta*-thioaspirin; *para*-TASP, *para*-thioaspirin.

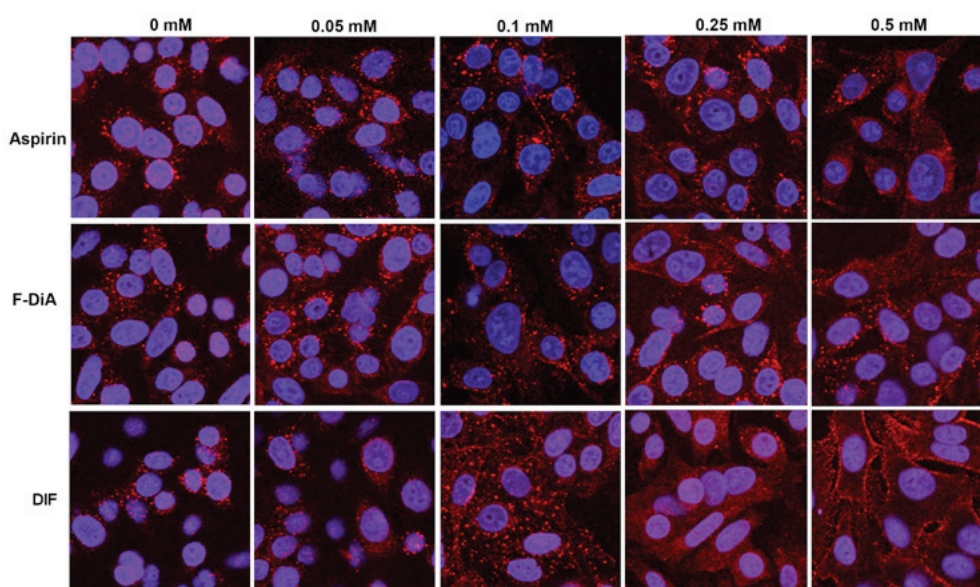


Figure 6. Effect of selected aspirin analogues on EGF (100 ng/ml) internalisation in the SW480 cell line at concentrations <0.5 mM. Confocal analysis of SW480 colorectal cancer cells incubated with Alexa Fluor 555-EGF and preincubated in the absence/presence of stated compound and vehicle control (0 mM) under HEPES-buffered conditions. Images were acquired at 405 nm for DAPI nucleic stain (blue) and 561 nm for Alexa Fluor-EGF (red). Representative images were captured at x400 magnification (n=3). EGF, epidermal growth factor; F-DiA, fumaryldiaspirin; DIF, diflunisal.

and endocytosis. The titratability of the inhibition of EGF internalisation was tested for dose-dependency in the SW480 cell line qualitatively with aspirin, F-DiA (PN517) and diflunisal (selected because of clinical relevance and cytotoxicity) at concentrations below 0.5 mM (Fig. 6), and although observable with aspirin and F-DiA, was most notably observed with diflunisal, thus arguing against a non-specific phenomenon associated with the experimental design.

Aspirin and aspirin analogue effects on EGFR and EGFR phosphorylation. Non-specific inhibition of kinase activity has been proposed as a mechanism of action for aspirin and sodium salicylate (59). Hence, compound effects on EGFR phosphorylation, focusing on cytoplasmic phosphotyrosine residues in the ligand-induced autophosphorylation domain were examined, including Tyr1045, Tyr1068 and Tyr1173. Phosphorylated Tyr1045 acts a site for c-Cbl interaction that results in receptor ubiquitylation and degradation of EGFR, and Cbl regulates receptor downregulation and signalling (83). Phosphorylation of Tyr1068 enhances binding of the growth factor receptor bound protein 2 (Grb2), which is necessary for entry into coated pits (41) and Ras/MAPK activation (84,85). Phospho-Tyr1173 is involved in binding the tyrosine phosphatase SHP1 and signalling via recruitment of the adaptor proteins SHC and Grb2, resulting in control of the ERK activity (86-88). In the present study, SW480 cells were incubated overnight with the compounds of interest and briefly stimulated with 200 ng/ml rEGF (5 min) in serum-free medium. EGFR phosphorylation status was analysed by immunoblotting (Fig. 7A). The data suggested that inhibition of EGFR phosphorylation upon stimulation, by aspirin (*ortho* and *meta*) was weak. Tyr1045 and Tyr1173 phosphorylation was more sensitive to the presence of diaspirins, including F-DiA (PN517) and PN524, and thioaspirins, including *ortho*-TASP, *meta*-TASP and *para*-TASP, compared with Tyr1068.

Thioaspirin (*ortho*-TASP) was the sole compound able to inhibit phosphorylation at all sites tested herein. These findings suggested that aspirin and analogues at the concentration tested (0.5 mM), even when cells are stimulated with high rEGF concentrations (200 ng/ml), did not universally inhibit all autophosphorylation sites, which argues against aspirin or its analogues acting solely as a non-specific kinase inhibitor, as suggested by Frantz *et al* (59). Due to the sensitivity of Tyr1045 to the compounds and its role in receptor degradation, the compound effects on EGF phosphorylation at this site with a 2-h incubation period were further examined and modest changes were observed by immunoblot analysis (Fig. 7B). As aspirin was reported to normalise EGFR expression (89), the effect of aspirin and its analogues on total EGFR levels in SW480 cells was further investigated following a 24-h incubation (Fig. 7C). SW480 EGFR levels were substantively affected by diflunisal, but weaker changes were observed as a result of incubation with F-DiA (PN517), PN524 and PN529, with little evidence of a significant reduction in EGFR by aspirin or salicylate following a 24-h incubation. pEGFR levels relative to total EGFR at pY1068 and pY1173 sites was totally inhibited by *ortho*-TASP and significantly reduced at pY1045 and pY1173 phosphorylation sites by *meta*-TASP, *para*-TASP, F-DiA, DiA and PN524. The pEGFR level at pY1068, pY1045 and pY1173 was increased by aspirin, *meta*-aspirin and *para*-aspirin (Fig. 7D).

Taken together, these data suggested that the EGFR axis was sensitive to aspirin and its derivatives. These compounds may have a rapid and profound impact on EGF endocytosis and autophosphorylation (particularly *ortho*-TASP with overnight incubation) and, furthermore, to a limited extent, EGFR expression, with the exception of diflunisal. Diflunisal markedly reduced EGFR expression and perturbed EGF internalisation, an observation that, to the best of our knowledge, has not been previously reported for a prescription

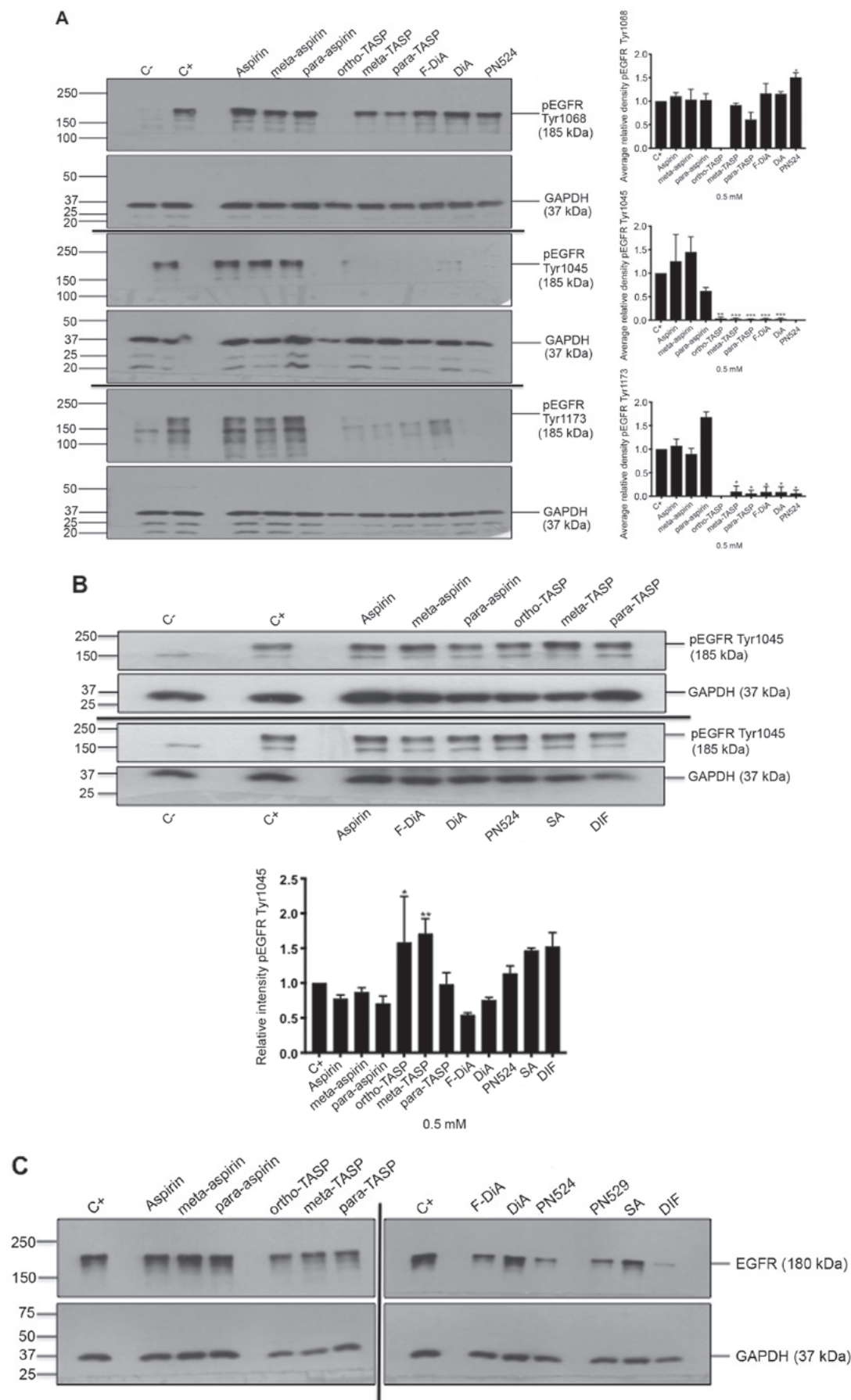


Figure 7. Effect of aspirin and analogues on pEGFR tyrosine kinase phosphorylation sites and EGFR. SW480 colorectal cancer cells were incubated with human rEGF (200 ng/ml) for 5 min following being treated with 0.5 mM compound (HEPES buffered) for (A) 24 h and probed with anti-human EGFR pY1068, pY1045 and pY1173 antibodies accompanied by corresponding histograms, (B) 2 h and probed with anti-human EGFR pY1045 and (C) 24 h and probed with an anti-human EGFR (D38B1) XP rabbit antibody.

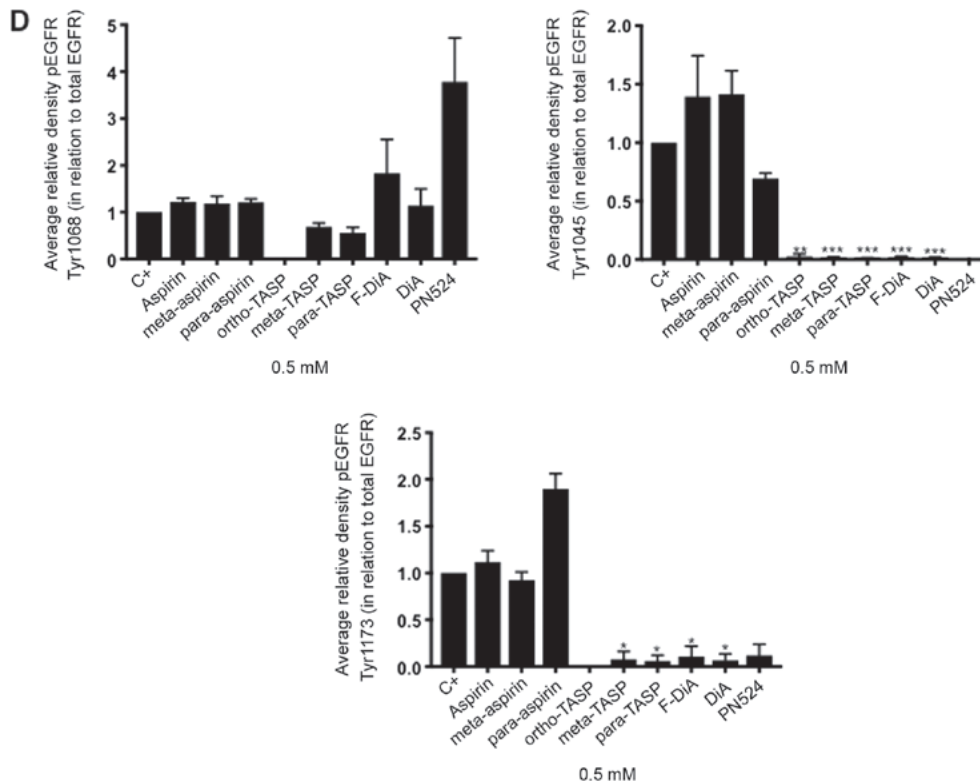


Figure 7. Continued. (D) Quantification of the effect of these compounds on pEGFR tyrosine kinase phosphorylation sites 1,068, 1,045 and 1,173 in relation to total EGFR expression. Cell lysates were analysed by SDS-PAGE and immunoblotting. pEGFR and EGFR bands migrated at ~185 and ~180 kDa, respectively, molecular weight was estimated by interpolation using Precision Plus colour standards. Band intensities were quantified relative to GAPDH. C- represents unstimulated and untreated cells; C+ represents untreated cells stimulated with rEGF for 5 min. Data presented are the mean of three individual experiments \pm standard error of the mean. * $P < 0.05$, ** $P < 0.01$ and *** $P < 0.001$ vs. C+ (one-way analysis of variance followed by Dunnett's test). EGF, epidermal growth factor; EGFR, EGF receptor; pEGFR, phosphorylated EGFR; rEGF, recombinant EGF; Y, tyrosine; SA, salicylic acid; DiA, diflunisal.

drug used in the treatment of osteoarthritis and rheumatoid arthritis. Given that endocytosis of EGF is rapid with migration of the ligand-receptor as a unit to a perinuclear location and then recycled being a canonical pathway for EGF (90) and the role of the proteasome in the processing and degradation of EGFR (91), it would be of interest to determine whether there is any association between diflunisal use and proteasomal degradation of EGFR.

Compound effects on EGF internalisation in Flo-1 and OE33 oesophageal cancer cells. It was further explored whether the phenomenon reported herein was restricted to SW480 cells. To that end and due to evidence that aspirin is chemopreventive in oesophageal cancer (92,93), Alexa Fluor 555-EGF internalisation was examined in the presence of the stated compounds with HEPES buffering in Flo-1 and OE33 oesophageal cancer cells. It was observed that aspirin and F-DiA perturbed EGF internalisation (at 20 ng/ml) in the oesophageal cancer cell lines (Fig. 8). EGF internalisation was markedly affected by F-DiA (PN517) in the Flo-1 cells, whereas a modest effect was observed with aspirin.

The present study details a rapid and hitherto unreported effect of aspirin and analogues, which are likely to yield salicylates upon metabolism within cells and tissues, on EGF endocytosis. The study has relevance to wound healing, cellular proliferation and chemoprevention. It addressed the analyses at physiologically achievable levels (≤ 1 mM) of aspirin (60,61) and analogues, and examined EGF internalisation and

EGFR phosphorylation with what is considered to be an experimentally legitimate concentration of EGF (20 ng/ml) under buffered conditions, as acidification of the cytosol may inhibit endocytosis.

Given the number of compounds tested and for the purposes of clarity the effect of compounds on EGF at unitary values of either 0.5 or 1 mM was investigated. It is recognised that these values are in certain cases $>IC_{50}$ values for SW480 cells reported herein but with the exception of diflunisal the fold-differences are relatively modest; however, plasma concentrations of diflunisal range from 150-350 μ M (22). Most significantly, the concentrations employed for salicylate and aspirin tested were markedly $<IC_{50}$ values and within a range (0.5-2 mM) that has been reported to be physiologically relevant.

Although a substantial body of published work pertains to the effects of aspirin and NSAIDs on gene expression and apoptosis in cells of cancerous origin (26,27), experiments often focus on outcomes over hours/days. It was observed that ligand-induced activation of receptor tyrosine kinases is rapid (within minutes); therefore, EGF endocytosis was investigated in the presence of aspirin/salicylates within time frames relevant to EGF signalling and endocytosis.

With respect to the findings presented herein, p38 MAPK was proposed as a potential rapid target, based on previously published work. Cyclin D1 is overexpressed in cells with EGFR mutations (94). It has previously been reported that, in SW480 cells treated with F-DiA, cyclin D1

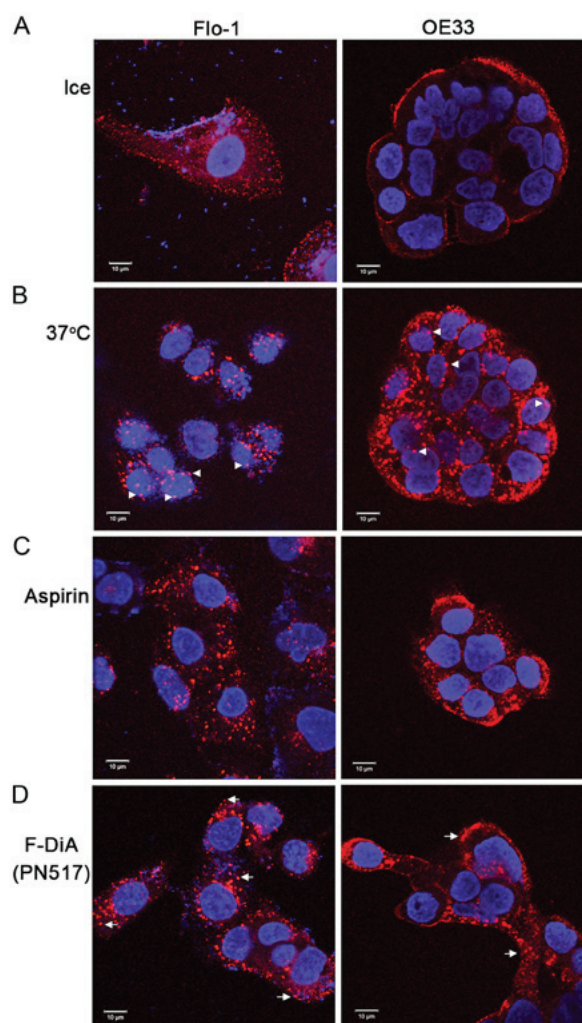


Figure 8. Effect of aspirin analogues under HEPES-buffered conditions on EGF (20 ng/ml) internalisation in Flo-1 and OE33 oesophageal cell lines. Confocal analysis of oesophageal cancer cells incubated with (0.5 mM) or without compound and fluorescence-conjugated EGF. (A) Negative control, without EGF internalisation in the absence of compound. Cells warmed to 37°C in (B) the absence of compound and the presence of (C) aspirin and (D) F-DiA. Images were acquired at 405 nm for DAPI nucleic stain (blue) and at 561 nm for Alexa Fluor 555-EGF (red). Representative images were recorded at x40 magnification using an oil/1.30 NA oil immersion objective lens. The close proximity of EGF towards the nucleus upon internalisation with warming is observed in Flo-1 cells (arrowheads). In cells incubated with compound the EGF is less centrally located (arrows), particularly with F-DiA in Flo-1 cells. EGF, epidermal growth factor; F-DiA, fumarylidi aspirin.

levels are significantly reduced and NF- κ B signalling is attenuated (27). It has previously been reported that cyclin D1 is reduced in CRC cells treated with aspirin, a phenomenon which has been reported to occur in response to p38 MAPK activation within minutes (95). Furthermore, salicylate exposure to COS-1 cells (58) can activate p38 MAPK also within minutes. A number of studies have implicated p38 MAPK as regulator of endocytic trafficking (96,97) and EGFR internalisation (98). p38 MAPK can regulate mu opioid receptor endocytosis, with the early endosomal antigen EEA1 identified as a substrate for p38 MAPK, which impacts on Rab5 activity regulating endocytosis (99). It is suggested that the rapid dysregulation of endocytosis is caused by salicylates and aspirin rapidly activating p38 MAPK and disrupting receptor trafficking, perturbing EGF

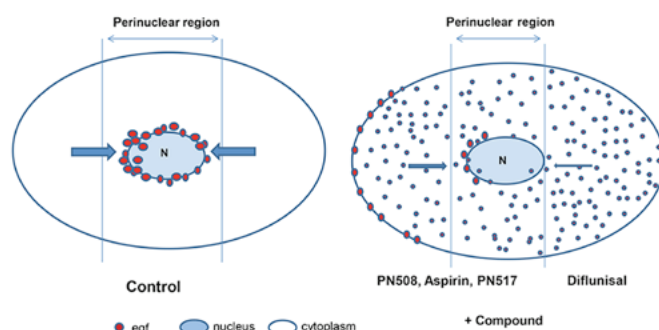


Figure 9. Schematic illustration of the effects of compounds on EGF internalisation. EGF, epidermal growth factor.

signalling and ultimately controlling cyclin D1 levels. This may be facilitated by salicylate persisting/accumulating as a result of its long half-life determined by pharmacokinetics analysis (100), and inhibition of NF- κ B and further known downstream targets. The relatively long half-life of salicylate may be relevant to the chemopreventive nature of low-dose aspirin: Accumulation in tissue reduces inflammation and may further attenuate EGF signalling. These findings suggest that aspirin and salicylates may be useful in cancer treatment, where EGFR amplification, overexpression and constitutive activation in a cancer is notable, e.g., in glioblastoma (101).

With respect to the question of non-salicylate NSAIDs also impacting on the EGFR signalling pathway, notwithstanding concerns regarding conflicting evidence regarding the efficacy of other NSAIDs (in comparison with aspirin) in reducing human cancer risk, e.g., in breast cancer (102), the evidence in the literature to date, as far as we are aware, other than work focused on sulindac affecting EGFR activation/inhibition, directly is limited. There are, however, reports of interrelationships between EGFR and PGE₂: for example, activation of EGFR by EGF has been reported to stimulate PGE₂ production (103).

Given the fundamental role of EGF signalling in cancer biology, the findings reported in the present study have relevance to the understanding of the chemopreventive activity of aspirin, particularly with respect to COX-independent pharmacology. A schematic illustration of the effect of selected compounds on EGF internalisation is presented in Fig. 9, based upon the representation of the endosomal system by Taub *et al* (55). It is suggested that further research is required to evaluate the rapid effect of salicylates on the EGFR axis, with a focus on MAPK activation and EGF/EGFR trafficking and phosphorylation. Furthermore, experimentation on EGF signalling in normal tissues exposed to aspirin or diflunisal is required to establish the physiological relevance of the observations reported herein. An analysis of the response of other growth factors and their cognate ligands to physiological concentrations of salicylates, with a particular emphasis on growth factor trafficking and recycling, appears timely. From a biochemical perspective, salicylate is a lipophilic monohydroxybenzoic acid that can interact with lipid membranes (104,105). It is also conceivable that interference of salicylates with membrane structure may affect endocytosis (106).

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Availability of data and materials

The analysed data sets generated during the study are available from the corresponding author on reasonable request.

Authors' contributions

All the authors have read and approved the final version of this manuscript. AIJB, CSK, STS, CJP and IDN designed and performed the experiments, analysed the data and wrote the manuscript. RMN wrote the Python program for proximity and density quotient analysis, and SJ contributed to data analysis and manuscript preparation.

Ethics approval and consent to participate

Not applicable.

Patient consent for publication

Not applicable.

Competing interests

CJP and IDN are named inventors for a patent for fumaryl-diaspirin as an anti-colorectal cancer agent (US patent no. 9351980).

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