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# Basal and inducible anti-inflammatory epoxygenase activity in endothelial cells

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# ABSTRACT

The roles of CYP lipid-metabolizing pathways in endothelial cells are poorly understood. Human endothelial cells expressed CYP2J2 and soluble epoxide hydrolase (sEH) mRNA and protein. The TLR-4 agonist LPS (1 µg/ml; 24 h) induced CYP2J2 but not sEH mRNA and protein. LC–MS/MS analysis of the stable commonly used human endothelial cell line EA.Hy926 showed active epoxygenase and epoxide hydrolase activity: with arachidonic acid (stable epoxide products 5,6-DHET, and 14,15-DHET), linoleic acid (9,10-EPOME and 12,13-EPOME and their stable epoxide hydrolase products 9,10-DHOME and 12,13-DHOME), docosahexaenoic acid (stable epoxide hydrolase product 19,20-DiHDPA) and eicosapentaenoic acid (stable epoxide hydrolase, but did not affect LPS, IL-1 $\beta$ , or phorbol-12-myristate-13-acetate (PMA)-induced TNF $\alpha$  release. In contrast, inhibition of soluble epoxide hydrolase by AUDA or TPPU inhibited basal, LPS, IL-1 $\beta$  and PMA induced TNF $\alpha$  release, and LPS-induced NF $\kappa$ B p65 nuclear translocation. In conclusion, human endothelial cells contain a TLR-4 regulated epoxygenase CYP2J2 and metabolize linoleic acid > eicosapentaenoic acid > arachidonic acid > docosahexaenoic acid to products with anti-inflammatory activity.

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# 1. Introduction

Arachidonic acid is metabolised to families of biologically active mediators by cyclooxygenase, lipoxygenase and cytochrome P450 (CYP) pathways [1,2]. Unlike COX and lipoxygenase products, the roles of CYP pathways in vascular biology remains poorly understood. CYPs metabolise arachidonic acid in several ways: (i) epoxygenases metabolise arachidonic acid to epoxyeicosatrienoic acids (EETs); (ii) lipoxygenase-like CYPs generate mid-chain hydroxyeicosatetraenoic acids (HETEs); and (iii) the  $\omega$ - and  $\omega$ -1-hydroxylase CYPs produce  $\omega$ -terminal HETEs [2]. In addition to arachidonic acid, these lipid metabolising CYPs can also metabolise linoleic acid (LA), docosahexaenoic acid (DHA) and eicosapentaenoic acid (EPA),

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to structurally related epoxy-products [3]. CYP2 family members are the main fatty acid-metabolising CYPs. Of this larger family, the CYP2J and CYP2C subfamilies represent the most important fatty acid epoxygenases [2–6].

In most organs, epoxygenase products are short-lived due to rapid metabolism. The major pathway of EET metabolism is via epoxide hydrolases [7], which convert EETs to dihydroxy derivatives called DHETs [8]. Although other enzymes possess epoxide hydrolase activity, soluble epoxide hydrolase (sEH) appears to be the key enzyme in this pathway, as genetic disruption or pharmacological inhibition of sEH elevates EETs, and reduces neointima formation [9], atherosclerosis, abdominal aortic aneurysm, and dyslipidaemia in hyperlipidaemic mice [10]. Knockout or inhibition of she also causes hypertension [11] and diabetes [12] in mouse models. The elevation of EETs by sEH inhibitors is now considered a potentially important therapeutic goal for management of cardiovascular and inflammatory disorders [13].

Primary endothelial cells use complex growth media and change phenotype with passage. EA.Hy926 are commonly used stable human cell line derived from human umbilical vein

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*Abbreviations:* AA, arachidonic acid; DHA, docosahexaenoic acid; DHET, dihydroxy eicosatrienoic acid; DHOME, dihydroxy-octadecenoic acid; DiHDPA, dihydroxy-docosapentaenoic acid; EET, epoxyeicosatrienoic acid; EPA, eicosapentaenoic acid; EPOME, epoxy-octadecenoic acid; HETE, hydroxyeicosatetraenoic acid; LA, linoleic acid; sEH, soluble epoxide hydrolase.

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endothelial cells fused to A549 human lung epithelial cells, which retain many endothelial cell characteristics [14]. Here we have characterized the lipid metabolizing CYP, lipoxygenase and cyclo-oxygenase pathways in EA.Hy926 cells, and investigated whether they represent a simple model to test the anti-inflammatory actions of sEH inhibitors. Our results show human endothelial cells contain TLR-4 inducible CYP2J2. As a model for this induction, EA.Hy926 actively metabolizes AA, LA, DHA, and EPA to epoxygenase metabolites, and sEH inhibitors strongly inhibit inflammation-induced TNFα release.

# 2. Materials and methods

## 2.1. Materials

MS-PPOH (N-(methylsulfonyl)-2-(2-propynyloxy)-benzene hexanamide), AUDA (12-[[(tricyclo[3.3.1.13,7]dec-1-ylamino) carbonyl]amino]-dodecanoic acid) and TPPU (N-[1-(1-oxopropyl)-4-piperidinyl]-N'-[4-(trifluoromethoxy)phenyl)-urea) were from Cayman Chemical Company (Cambridge Bioscience, Cambridge, UK). SKF525A ( $\alpha$ -phenyl- $\alpha$ -propyl-2-(diethylamino)ethyl esterbenzeneacetic acid) was from Biomol (Affiniti Research Products, Exeter, UK). Human TNF $\alpha$  ELISA was from eBioscience (Hatfield, UK). Rabbit anti-p65 was from Santa Cruz (Heidelberg, Germany), while rabbit anti-CYP2J2 and anti-sEH were from Biorybt (Cambridge, UK). Taqman primers and reagents were from Invitrogen (Paisley, Renfrewshire, UK). Recombinant human IL-1 $\beta$  was from R&D Systems (Abingdon, Oxfordshire, UK). Unless stated, all other reagents were from Sigma–Aldrich (Poole, Dorset, UK).

## 2.2. Cell and tissue culture

EA.Hy926 (ATCC; LGC Standards, Middlesex, UK) were cultured in M199 supplemented with antibiotic/antimycotic mix, and 10% FBS; 37 °C; 5% CO2; 95% air. HAEC used at passage 3–4, were from PromoCell and grown according the manufacturer's recommended conditions. Human blood out-growth endothelial cells were a gift from Prof. Jane Mitchell (Imperial College London). Since FBS interferes with lipid substrate composition, and the release and detection of eicosanoids (M. Edin, unpublished observations), all experiments were performed with M199 supplemented with antibiotic/antimycotic mix, *without* FBS. Detection of eicosanoids by liquid chromatography tandem mass spectrometry was performed as previously described [27].

### 2.3. RT-PCR and real-time qRT-PCR

Total RNA was extracted using Qiagen's QiaAmp RNA Blood Mini Kit as per the manufacturer's instructions. qRT-PCR for CYP2J2, and sEH (ephx2) was measured by the Taqman qRT-PCR ddCt method and normalized to GAPDH levels, using a Corbett Rotor-Gene 6000 machine.

#### 2.4. Immunoassays, and EET measurements

TNF $\alpha$  ELISAs were performed according to manufacturer's instructions. In these experiments cell viability by MTT assays were performed as previously described [15]; and there were no significant changes between treatments groups. Immunofluorescence was performed essentially as previously described using primary antibody dilutions of 1:50–1:100 [15]. LC–MS/MS analysis of epoxygenase products in culture supernatants was as previously described [16].

# 3. Results

#### 3.1. Profile of lipid metabolizing CYPs and their products in EA.Hy926

EA.Hy926, human aortic endothelial cells, and human blood outgrowth endothelial cells contained CYP2J2 and sEH mRNA and protein (Fig. 1). The TLR-4 ligand LPS (1  $\mu$ g/ml) induced CYP2J2 mRNA (4 h) and protein (24 h) (Fig. 1). In contrast, sEH mRNA and protein levels remained relatively stable following LPS treatment (Fig. 1).

As a stable endothelial cell line we chose to study EA.Hy926 further. Over 24 h EA.Hy926 released detectable levels of the epoxygenase products of AA [14,15-DHET]; LA [9,10-epoxy-octadecenoic acid (EPOME), 12,13-EPOME and their respective diol products 9,10-DHOME, and 12,13-DHOME]; DHA [19,20-dihydroxy-docosapentaenoic acid (DiHDPA)]; and EPA [17,18-DHET; the most abundant epoxygenase product detected under basal culture conditions] (Fig. 2A). LA but not AA lipoxygenase products 9-, and 13-HODE were also detected basally (Fig. 2B), as were the cyclo-oxygenase products PGF<sub>20</sub>, PGD<sub>2</sub> and PGE<sub>2</sub> (Fig. 2C).

When stimulated with the TLR-4 ligand LPS (1  $\mu$ g/ml; 24 h), 12,13-DHOME, 5,6-DHET and in particular 9,10-DHOME release was stimulated (Fig. 2A). LPS also stimulated release of 9- and 13-HODE, and 12- and 8-HETE (Fig. 2B), but not COX products (Fig. 2C).

# 3.2. Endogenous epoxygenase activity inhibits TNF $\alpha$ release in EAHy926 cells

EA.Hy926 cells were treated with vehicle (0.01% DMSO) or the chemically distinct epoxygenase inhibitors MS-PPOH or SKF525A (10  $\mu$ M each; 7 h). Vehicle treated EA.hy926 cells produced 19.8 pg/ml of TNF $\alpha$ . Treatment with either MS-PPOH or SKF525A induced an approximate 3-fold increase in TNF $\alpha$  production (Fig. 3A). In the presence of high concentrations of the inflammatory stimuli IL-1 $\beta$  (10 ng/ml), PMA (5 nM) or LPS (1  $\mu$ g/ml), MS-PPOH co-incubation had no additional effect on TNF $\alpha$  release (Fig. 3B).

# 3.3. Soluble epoxide hydrolase inhibitors are anti-inflammatory in EAHy926 cells

Treatment with 10  $\mu$ M AUDA significantly reduced TNF $\alpha$  secretion in the absence of inflammatory stimuli, and abolished PMA (5 nM), LPS (1  $\mu$ g/ml) or IL-1 $\beta$  (10 ng/ml) stimulated TNF $\alpha$  release (7 h; Fig. 4A). Moreover, treatment with AUDA (10  $\mu$ M) or TPPU (1  $\mu$ M) inhibited LPS (1  $\mu$ g/ml; 2 h)-induced nuclear localization of the NF $\kappa$ B p65 subunit (Fig. 4B).

# 4. Discussion

Here we show human endothelial cells contain a TLR-4 inducible CYP2J2 epoxygenase. In addition, we show EA.hy926 are a useful, simple, and reproducible human model system to study this inducible CYP epoxygenase enzyme pathway and sEH inhibitors.

Under basal conditions, the release of AA epoxygenase/sEH product 14,15,DHET, the linoleic acid products 9,10-EPOME and 12,13-EPOME (and 9,10-DHOME and 12,13-DHOME), the DHA product 19,20-DiHDPA, and the EPA product 17,18-DHET were all detectable in EA.Hy926 conditioned media. All these products are also produced by vascular tissue (e.g., rat aorta; DBB, ME, DCZ unpublished observations), and in the circulation of healthy volunteers [17]; however the actions of many of these lipid metabolites, including those produced in the largest amounts (e.g., 17,18-DHET) are still poorly understood. Exogenous



**Fig. 1.** Expression and regulation of CYP2J2 and sEH by LPS in human endothelial cells. (A) qRT-PCR (4 h) and (B) immunofluorescent analysis (24 h) of CYP2J2 and sEH expression in EA.Hy926 treated with LPS (1 µg/ml). qRT-PCR data represent relative expression compared to GADPH as mean ± SEM from n = 4 separate experiments. (C) qRT-PCR (4 h) of human blood out growth endothelial cells and (D) immunofluorescent analysis (24 h) of CYP2J2 in human primary aortic endothelial cells treated with LPS (1 µg/ml). qRT-PCR data represent relative expression compared to GADPH as mean ± SEM from n = 3 separate donors. \* indicates p < 0.05 by one-sample *t*-test between control and LPS treatment. Immunofluorescent micrographs are representative of n = 3 experiments. As a negative control, in some experiments primary antibody was omitted (2<sup>°</sup> Ab) which showed no specific staining.



**Fig. 2.** Oxylipin product profile of control and LPS stimulated EA.Hy926 endothelial cells. LC/MS/MS analysis of the (A) epoxygenase, (B) lipoxygenase and (C) cyclo-oxygenase products release (pg/ml) from EA.Hy926 untreated or treated with LPS (1  $\mu$ g/ml) for 24 h. Data represents the mean ± SEM from *n* = 3. \* indicates *p* < 0.05 for each analyte by unpaired *t*-test between control and LPS treatment.

17,18-EET acts as a vasodilator [18] via actions on the BK channel alpha subunit [19], inhibits ristocetin-induced thrombus formation [20], and exerts negative chronotropic effects and protects neonatal rat cardiomyocytes against  $Ca^{2+}$ -overload [21]. Any effects on endothelial cells themselves have yet to be described, so warrant further investigation.

We recently showed CYP2J2 is induced by LPS in human peripheral blood monocytes and monocyte cell lines [22]. CYP2J2 is expressed in endothelial cells under basal conditions and is similarly induced by the TLR-4 ligand LPS in endothelial cells. This induction of CYP2J2 was accompanied by increases in 9,10-DHOME, 12,13-DHOME and 5,6-DHET formation. These findings indicate that CYP2J2 is a more general inflammation-induced enzyme than perhaps previously thought. EA.Hy926 cells also express CYP2C8 but not CYP2C9 mRNA (Fig. 1), which showed no sign of induction with LPS. Although EPA and DHA may be rate limiting in this system for their own metabolite formation, our results indicate that the LPS inducible eicosanoids are from an inducible CYP such as CYP2J2, while the non-inducible products (e.g., 17,18-EET) may derive more from non-TLR-4 regulated enzymes such as CYP2C8.

Epoxygenase pathway metabolites regulate endothelial cell adhesion molecule expression [3]. Our results now show



**Fig. 3.** Epoxygenase inhibition induces TNF $\alpha$  release from EAHy926 endothelial cells. (A) Fold TNF $\alpha$  release (7 h) from EA.Hy926 treated with vehicle control (0.01% DMSO), or epoxygenase inhibitor, MS-PPOH (10  $\mu$ M) or SKF525A (10  $\mu$ M). Data represents mean ± SEM from *n* = 7 replicates from 4 separate experiments. \* indicates *p* < 0.05 by one-sample *t*-test between control and epoxygenase inhibitors. (B) Fold TNF $\alpha$  release (7 h) from EA.Hy926 treated with vehicle control (0.01% DMSO), 5 nM PMA, 1  $\mu$ g/ml LPS or 10 ng/ml IL-1 $\beta$  in the presence of absence of 10  $\mu$ M MS-PPOH. Data represents mean ± SEM from *n* = 5 replicates from 4 separate experiments. \* indicates *p* < 0.05 by one-sample *t*-test between control and inflammatory stimuli inhibitors. † indicates *p* < 0.05 by one-sample *t*-test between treatment in presence or absence of MS-PPOH.



**Fig. 4.** sEH inhibition limits basal and inflammation induced TNF $\alpha$  release from EAHy926 endothelial cells. (A) TNF $\alpha$  release (fold from control; 7 h) from EAHy926 treated with vehicle control (0.01% DMSO), 5 nM PMA, 1 µg/ml LPS or 10 ng/ml IL-1 $\beta$  in the presence of absence of the sEH inhibitor AUDA (10 µM). Data represents mean ± SEM from *n* = 8 replicates from 4 separate experiments. \* indicates *p* < 0.05 by paired *t*-test between treatment in presence or absence of AUDA. (B) EA.Hy926 were treated with LPS (1 µg/ml; 2 h) to induce p65 nuclear localization. Scale bar 50 µ. LPS induced p65 nuclear localization is absent or strongly diminished in cells treated with sEH inhibitors TPPU (1 µM; middle panels) or AUDA (10 µM; lower panels). 2nd Ab control shows immunofluorescence levels when primary antibody was omitted. Each panel was enhanced by 20% brightness and 20% contrast in PowerPoint to aid visualization. Data is representative of *n* = 3 experiments.

suppression of TNFa can be added to the repertoire of anti-inflammatory actions of endothelial epoxygenases. TNF $\alpha$  is a potent inflammatory activator of endothelial cells [23] and is critical to the pathogenesis of many inflammatory diseases, most notably rheumatoid arthritis [24]. sEH inhibitors with their anti-inflammatory and analgesic profile [25,26] may therefore represent a viable treatment for highly inflammatory disorders such as rheumatoid arthritis. Although we may assume the actions of sEH inhibition in endothelial cells is due to the direct elevation of epoxygenase products acutely, it is also important to note that following sEH inhibition, endothelial epoxygenase products also get incorporated into cell membranes (potentially for future release) as well as get metabolized via  $\beta$ -oxidation to form novel lipid species such as 10,11-epoxy-hexadecadienoic acid (16:2) and 8,9-epoxy-myristoleic acid (14:1) whose functions are not known [27], and are rarely measured.

Primary endothelial cells are responsive to IL-1 $\beta$ , LPS and PMA [28]. The fact that EA.Hy926 respond similarly further supports their use as an in vitro cell line to model physiological effects of EETs. Chemically distinct epoxygenase inhibitors SKF525A and MS-PPOH increased TNF $\alpha$  production. This suggests, similar to our recent report on monocytes [29], that EET production helps

to maintain inflammatory homeostasis. The stronger TNF $\alpha$  release induced by the classical endothelial cell activators IL-1 $\beta$ , LPS or PMA, was not further elevated by epoxygenase inhibition, suggesting a common pathway e.g., NF $\kappa$ B, critical for TNF $\alpha$  release, is being used by both pathways.

Although EA.Hy926 cells appear to be a good model to study epoxygenase products and activity, they are not without their limitations when compared to primary endothelial cells. Importantly, in our hands the cyclooxygenase system was relatively inactive, and no prostacyclin could be detected under any condition, including following LPS stimulation (Fig. 2C).

Endothelial cells and EA.Hy926 cells actively metabolize different fatty acid substrates to a variety of biologically active fatty acyl/oxy-lipins. There has been intensive interest in the mechanistic basis for the effect of diets rich in different sources of fatty acids (e.g., the Mediterranean diet, diets rich in  $\omega$ 6-polyun-saturated fatty acids, or fish oil supplementation), and their effects on cardiovascular biology [30]. Modifying substrates and diet may therefore change the balance of oxylipins and may be responsible for the purported benefits of these lifestyle modifications. Moreover, changing the substrate itself not only alters mediator production but also can change the balance of mediators produced by regulating

phospholipase activation. Interestingly, when we gave AA (10 mM) acutely to EA.Hy926 cells, we did induce AA products 5,6-DHET and HETEs, however we also strikingly induced LA product EPOMEs and DHOMEs (Fig. 2). In contrast, 14,15-DHET, 17,18-DHET, and 19,20-DiHDPA were all reduced. Therefore in this simple system, just substrate delivery to the epoxygenase enzymes is far more complex than would be originally predicted.

In summary, endothelial cells contain an anti-inflammatory TLR-4 inducible epoxygenase. In addition, EA.Hy926 cells represent a useful tool to study the anti-inflammatory actions of the epoxygenase pathway metabolites and soluble epoxide hydrolase inhibitors in vascular endothelial cells.

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### Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at http://dx.doi.org/10.1016/j.bbrc.2014.03.020.

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