Themed Section: Eicosanoids 35 years from the 1982 Nobel: where are we now?

REVIEW ARTICLE

Lipid mediators in immune regulation and resolution

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We are all too familiar with the events that follow a bee sting—heat, redness, swelling, and pain. These are Celsus' four cardinal signs of inflammation that are driven by very well-defined signals and hormones. In fact, targeting the factors that drive this onset phase is the basis upon which most current anti-inflammatory therapies were developed. We are also very well aware that within a few hours, these cardinal signs normally disappear. In other words, inflammation resolves. When it does not, inflammation persists, resulting in damaging chronic conditions. While inflammatory onset is actively driven, so also is its resolution—years of research have identified novel internal counter‐regulatory signals that work together to switch off inflammation. Among these signals, lipids are potent signalling molecules that regulate an array of immune responses including vascular hyper reactivity and pain, as well as leukocyte trafficking and clearance, so‐called resolution. Here, we collate bioactive lipid research to date and summarize the major pathways involved in their biosynthesis and their role in inflammation, as well as resolution.

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1 | INFLAMMATION AND ITS RESOLUTION

Inflammation is a protective response against infection and/or injury. However, when it becomes dysregulated as a consequence of genetic abnormalities, the ageing process or environmental factors, our immune system has the capacity to cause extensive damage. Arthritis, asthma, chronic obstructive pulmonary disease, Alzheimer's disease, atherosclerosis, and even cancer, while aetiologically disparate, are diseases unified by a dysregulated immune component. The current strategy of treating such diseases is based, largely, upon inhibiting

Abbreviations: 15-epi-LXs, 15 epimeric-LX; AA, arachidonic acid; COX, cyclooxygenase; CYP450, cytochrome P450; DHA, docosahexaenoic acid; DHETs, dihydroxy‐eicosatrienoic acids; EPA, eicosapentaenoic acid; LXs, lipoxins; MaR, maresins; NSAIDs, nonsteroidal anti-inflammatory drugs; PD1, protectin D1; PDs, protectins; RvD1, resolvin D1; Rvs, resolvins; sEH, soluble epoxide hydrolase

the factors that drive acute inflammation such as nonsteroidal anti-inflammatory drugs (NSAIDs-such as [naproxen](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=5230) or diclofenac), steroids ([prednisone](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=2866)), and "biological" drugs such as [infliximab](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=5004) (anti-TNF) and **[anakinra](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=6972)** (anti-IL-1). Although these medicines ameliorate disease symptoms, they do not bring about a "cure" and are ineffective in a significant subset of patients. Furthermore, side effects can hamper endogenous homeostatic systems, predisposing to infection. Thus, there is a need to develop more efficient and effective therapeutic agents, with one approach being to harness the body's own healing process for therapeutic gain.

Consequently, attention has turned to the other end of the inflammatory spectrum, resolution, in order to understand the endogenous processes that switch off inflammation. Our objective has been to identify novel internal counter‐regulatory systems that terminate inflammation in order to provide new targets that can be harnessed pharmacologically to push ongoing inflammation down a pro‐

resolution pathway. As a result, resolution is now been studied in great detail with clear evidence suggesting that resolution is an active process with quantifiable indices and specific requirements. Along these lines, lipid mediators have emerged as internal regulatory signals that activate many aspects of the inflammation and resolution cascade, including terminating leukocyte trafficking into tissue once the inflammatory signal has been removed, scavenging pro‐inflammatory signals as well as clearing dead cells from the resolves site. Hence, in this review, the role of lipids in the resolution cascade will be discussed.

2 | CYCLOOXYGENASE AND PROSTANOIDS

The enzyme [cyclooxygenase](http://www.guidetopharmacology.org/GRAC/FamilyDisplayForward?familyId=269) (COX) converts [arachidonic acid](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=2391) (AA) to form $PGG₂$ (Pagels et al., 1983) with the peroxidase element of the enzyme further reducing PGG_2 to PGH_2 (Hamberg & Samuelsson, 1973), which serves as a precursor for all major prostanoid mediators. There are two principal isoforms involved in the conversion of AA to prostanoids, namely, COX‐1 and COX‐2. Unlike COX‐1, which is constitutively expressed in most cells and tissues and is broadly involved in house‐keeping functions, COX‐2 is induced in response to inflammatory stimuli (Dubois et al., 1998) being expressed at sites of infection and injury with the exception of parts of the brain and kidney (Harris et al., 1994). Formation of prostanoids from $PGH₂$ occurs through the actions of downstream synthases that are expressed in a tissue and cell type‐selective fashion including PGD synthase (Shimizu, Yamamoto, & Hayaishi, 1982) PGE synthase 1, 2, and 3 (Tanaka, Ward, & Smith, 1987), PGF synthase (Hayashi, Fujii, Watanabe, Urade, & Hayaishi, 1989), prostacyclin synthase, and thromboxane A synthase (Ullrich & Haurand, 1983), which form PGD_{2,} PGE₂, PGF_{2a}, PGI₂ (also known as prostacyclin), and TXA₂ respectively. The differential expression of these downstream enzymes within cells determines the profile and levels of prostanoid production generated under resting and inflammatory conditions.

Presently, there are nine known **[prostanoid receptors](http://www.guidetopharmacology.org/GRAC/FamilyDisplayForward?familyId=58)** in mice and man. These include the PGD receptors, DP_1 and DP_2 ; the PGE_2 receptors, EP_1 , EP_2 , EP_3 , and EP_4 ; the PGF receptor, FP; the PGI receptor, IP; and the TXA receptor, TP. In addition, there are splice variants of the EP₃, FP, and TP receptors differentiated only in their C-terminal tails. All of these receptors belong to the GPCR superfamily of proteins, with the exception of $DP₂$ (also known as CRTH2), which is a member of the chemokine receptor family (Hirai et al., 2001). The IP, DP_1 , EP_2 , and EP_4 receptors signal through G_s resulting in an increased intracellular cAMP, whereas the EP_3 receptor couples to G_i to reduce cAMP, while EP₁, FP, and TP receptors signal through G_a to induce calcium mobilization.

The more common prostanoids, PGE_2 and PGI_2 , both enhance vasodilation (Kaley, Hintze, Panzenbeck, & Messina, 1985), oedema formation, and vascular permeability, particularly in the presence of histamine, bradykinin, and 5‐HT (Hata & Breyer, 2004). Mice that are genetically depleted for their respective receptors (IP, EP_2 , and EP_3) show reduced pleural exudation following treatment with inflammogens including carrageenan and zymosan (Yuhki et al., 2004).

Robust evidence from EP‐deficient mice has shown that the febrile response to PGE₂ arises from the actions of PGE₂ on its EP_3 receptor, which is present on sensory neurons in the periphery and brain (Dantzer, Konsman, Bluthe, & Kelley, 2000). Equally, PGE_2 is a potent pyretic agent known with elevated concentrations found in cerebrospinal fluid taken from patients with bacterial or viral infections (Saxena, Beg, Singhal, & Ahmad, 1979). While none of the prostanoids cause pain directly, $PGI₂$ and $PGE₂$ reduce the threshold of nociceptor sensory neurons to stimulation when bound to IP, EP_1 , EP3, and EP4 receptors respectively (Ahmadi, Lippross, Neuhuber, & Zeilhofer, 2002).

Prostanoids also play an important role in protecting against oxidative injury in cardiac tissue and in maintaining cardiovascular (CV) homeostasis. Indeed, their protective effect has been demonstrated in clinical studies, which found an increased risk of myocardial infarction (MI), stroke, systemic and pulmonary hypertension, thrombosis, and sudden cardiac death following the use of COX‐2‐specific inhibitors (Garcia Rodriguez, Tacconelli, & Patrignani, 2008). Furthermore, deleting specific prostanoid synthases and receptors results in an augmentation of ischaemia–reperfusion injury (Xiao et al., 2001) as well as contributing to the decline in cardiac function following MI. CV health is regulated by vasodilatory PGI₂ and pro-thrombotic TXA₂ (Bunting, Moncada, & Vane, 1983), where $PGI₂$ counterbalance the actions of TXA₂ (Grosser, Fries, & FitzGerald, 2006). Indeed, endothelial $PGI₂$ along with NO prevents TXA₂-induced platelet aggregation and thrombosis. TXA₂ is derived from platelet COX‐1 causing platelet aggregation and vascular smooth muscle contraction (Ellis et al., 1976). Clinical CV diseases including unstable angina, MI, and stroke can arise from overproduction of TXA4. Importantly, the cardio‐protective properties of aspirin can be attributed to the covalent inhibition of COX‐1 (Rocca et al., 2002).

As well as being pro‐inflammatory, many prostanoids up‐regulate intracellular cAMP triggering immuno‐suppressive effects. For example, PGE_2 and PGI_2 reduce the ability of inflammatory leukocytes to phagocytose and kill microorganisms (Aronoff, Canetti, & Peters‐ Golden, 2004), as well as inhibiting the production of downstream pro-inflammatory mediators (Aronoff et al., 2007) while, in contrast, triggering the synthesis of IL-[10](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=4975) and [IL](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=4998)-6 (Harizi, Juzan, Pitard, Moreau, & Gualde, 2002). Indeed, in a number of conditions associated with increased susceptibility to infection, including cancer (Starczewski, Voigtmann, Peskar, & Peskar, 1984), ageing (Hayek et al., 1997), and cystic fibrosis (Medjane, Raymond, Wu, & Touqui, 2005), overexpression of $PGE₂$ has been reported. Interestingly, during the very early phase of acute inflammation, $PGE₂$ indirectly exerts pro‐resolution effects by switching on the transcription of enzymes necessary for the generation of lipoxins (LXs; Levy, Clish, Schmidt, Gronert, & Serhan, 2001), resolvins (Rvs), and protectins (PDs; Hong, Gronert, Devchand, Moussignac, & Serhan, 2003). These represent other classes of lipids mediators with pro‐resolution properties.

While $PGD₂$ can elevate cAMP via its $DP₁$ receptors, $PGD₂$ may also act independently of its DP_1 and DP_2 receptors when it nonenzymically dehydrates into PGs of the J_2 series, such as PGJ₂, $\Delta^{12,14}$ $\Delta^{12,14}$ $\Delta^{12,14}$ -PGJ₂, and 15-[deoxy](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=2692)- $\Delta^{12,14}$ -PGJ₂ [15d-PGJ₂]; Clark et al., 2000). These cyclopentenone PGs form covalent attachments with reactive sulphydryl groups on intracellular regulatory proteins, which enables modulation of their function. For instance, $15d$ -PGJ₂ upon ligation to the nuclear receptor [PPAR](http://www.guidetopharmacology.org/GRAC/FamilyDisplayForward?familyId=86)-γ (Khan, 1995) decreases proinflammatory cytokine release and modifies gene expression (Jiang, Ting, & Seed, 1998) as well as directly inhibiting the actions of IκB kinase, which is responsible for the activation of NF‐κB (Cernuda‐ Morollon, Pineda-Molina, Canada, & Perez-Sala, 2001). 15d-PGJ₂, identified in rodent peritonitis resolution exudates (Rajakariar et al., 2007), independently of PPAR‐γ, can preferentially inhibit monocyte, rather than neutrophil, trafficking through differential regulation of cell-adhesion molecule and chemokine expression (Gilroy et al., 2003); regulate macrophage activation and pro‐inflammatory gene expression (Lawrence, 2002); and induce leukocyte apoptosis through a caspase‐dependent mechanism (Bishop‐Bailey & Hla, 1999). Moreover, PGD₂-derived compounds function as endogenous braking signals for lymphocytes to stimulate resolution (Trivedi et al., 2006). Table 1 summarises the prostanoids, their biological actions and concentrations at sites of inflammation.

3 | PROSTANOIDS AND POST‐RESOLUTION BIOLOGY

Recently, we demonstrated that classical resolution may not be the end of the local immune response to infection or injury, but rather that a third phase subsequent to these exists: post‐resolution (Motwani et al., 2017). Traditionally, resolution processes were deemed successful if acute inflammation, as described by leukocyte clearance and cytokine catabolism, was terminated. However, they may have a hitherto unappreciated role in controlling adaptive immune responses and maintaining tolerance. Specifically, we found that murine innate immune‐ mediated responses to low‐dose yeast cell wall extract (zymosan, administered i.p.) or bacteria (Streptococcus pneumoniae^{ovalbumin-labelled}. i.p.) were resolved. Interestingly, these low‐dose stimuli elicited a previously overlooked second wave of leukocyte influx into tissues that persisted for weeks. These cells comprised three separate populations of Ly6c^{hi} monocyte-derived macrophages including CD11B⁺/ CD49d+ /CD115+ myeloid-derived suppressor cells, $F4-80^{10}/MHC$ -II⁺/CD11c⁺ dendritic cells, and $F4-80^{int}/CD11B^{hi}/$ CD11c[−] macrophages. In addition, tissue‐resident (embryonic‐ derived) macrophages, which disappear during the acute inflammatory response, re‐appear. These diverse populations of macrophages were observed alongside lymph node expansion and increased numbers of peripheral blood and tissue memory T and B lymphocytes. Polymorphonuclear (PMNs) were not present during this phase. One of the key events in this process is the sustained synthesis of $PGE₂$, which is derived from macrophage COX‐1/mPGES and that is triggered by IFN‐γ. It transpires that this post‐resolution phase of prostanoid biosynthesis creates a window of susceptibility to repeat infections on the one hand, while also controlling local adaptive immune processes on the other (Newson et al., 2017). The nature of these prostanoid/adaptive immune interactions is being investigated.

4 | CYTOCHROME P450

[Cytochrome P450s](http://www.guidetopharmacology.org/GRAC/FamilyDisplayForward?familyId=242&familyType=ENZYME) (CYP450s) are a family of membrane‐bound, haem-containing enzymes found in the liver, kidneys, brain, heart, CV system, and lung and are best characterized for the catalysis of NADPH‐dependent oxidation of drugs, chemicals, carcinogens, and hormones (Nelson et al., 1996). The CYP450 family contains 57 genes in humans, and although approximately one quarter of these have been shown capable of metabolizing polyunsaturated fatty acids (PUFAs), the **[CYP2J2](http://www.guidetopharmacology.org/GRAC/FamilyDisplayForward?familyId=262)** and CYP2C family members (CYP2C8, 2C9) are thought to be the major enzymes responsible for lipid mediator production (Bishop‐Bailey, Thomson, Askari, Faulkner, & Wheeler‐Jones, 2014). In addition to metabolizing AA (Figure 1), CYP450s also readily metabolize the related ω6 PUFA [linoleic acid](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=1052) (Figure 2) and ω3 PUFAs (see below also) [docosahexaenoic acid](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=1051) (DHA, Figure 3) and [eicosapentaenoic acid](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=3362) (EPA; Figure 4) into a series of related biologically active mediators (Smilowitz et al., 2013). CYP450 are capable or metabolizing PUFA substrates by epoxygenase, [lipoxygenase](http://www.guidetopharmacology.org/GRAC/FamilyDisplayForward?familyId=271) (LOX) and ω‐hydroxylase type of activities (Zeldin, 2001). The epoxygenase activity inserts a single molecular oxygen in one of the double bonds of each PUFA, for example, for AA to form one of four regioisomers of epoxyeicosatrienoic acid ([5,6](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=6304)-, 8,9-, 11,12-, or 14,15-EET; the numbers indicating the double bond in AA subject to epoxygenation; Zeldin, 2001). Each EET can be formed as either an R/S or S/R stereoisomer, with ratios of production depending on the generating CYP450. Stereoisomers of EETs may have different biological activities, but little research exists to understand the extent of these differences. CYP450s can also have LOX activity producing mid‐chain (12[R]-), and ω -hydroxylase activity producing terminal (19[S > R]-, and [20](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=4103)-) hydroxyeicosatetraenoic acids (HETEs: Roman, 2002). Once formed, epoxygenase products in particular are quickly metabolized by epoxide hydrolases (EH) or reincorporated into membranes (Zeldin, 2001). Soluble EH (sEH) and microsomal EH (encoded by the ephx2 and ephx1, respectively) combine to metabolize virtually all epoxygenase products in vivo (Edin et al., 2018). For example, EETs get converted to dihydroxy‐eicosatrienoic acids. Importantly, a number of sEH inhibitors have been developed that inhibit the breakdown of epoxygenase products to potentiate their signalling (Hwang, Wecksler, Wagner, & Hammock, 2013).

AA and related PUFA are metabolized by CYP epoxygenase and epoxide hydolases in the vascular endothelium (Roman, 2002; Zhang et al., 2001) and vascular smooth muscle. In vascular smooth muscle, AA is also catalysed by CYP hydroxylases to 20-HETE (Wang et al., 1999). Indeed, CYP4F3A in myeloid tissue catalyses the ω-hydroxylation of LTB₄ to 20-hydroxy-LTB₄, an inactivation process that is critical for the regulation of the inflammatory response (Johnson, Edson, Totah, & Rettie, 2015). However, it is unknown whether CYP4F3 is the source of 20‐HETE produced by PMNs (Bednar et al., 2000). These metabolites play a large and complex role in maintaining cardiac, renal, and pulmonary homeostasis by regulating vascular tone and reactivity, ion transport, and renal and pulmonary functions as well as growth responses (Fleming, 2007). Moreover, they have been shown to exert striking anti-inflammatory actions (Inceoglu et al., 2008), see below.

TABLE 1 Biological actions of lipid mediators and their relative concentrations at sites of inflammation TABLE 1 Biological actions of lipid mediators and their relative concentrations at sites of inflammation

Note. E. coli: Escherichia coli. Note. E. coli: Escherichia coli.

FIGURE 1 Cytochrome p450 metabolism of arachidonic acid to epoxyeicosatrienoic acids (EETs) and their subsequent conversion by soluble epoxide hydrolase to dihydroxy‐ eicosatrienoic acids

FIGURE 2 Cytochrome p450 metabolism of linoleic acid to epoxy‐octadecenoic acids (EPOMEs) and dihydroxy‐octadecenoic acids (DHOMEs)

FIGURE 4 Cytochrome p450 metabolism of docosahexaenoic acid to epoxyeicosatrienoic acids

5 | CYP450 AND INFLAMMATION

EETs catalysed by CYPs 2C8, 2C9, and 2J2 inhibit the activation of the transcription factor NF‐κB via the IκB kinase (Node et al., 1999). Consequently, EETs may therefore have the propensity to down-regulate various cytokine-induced, pro-inflammatory, signalling pathways downstream of NF‐κB activation. This may explain how EETs prevent the adhesion of PMNs to the vascular wall by suppressing the expression of cell adhesion molecules, including [intracellular adhesion](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=6757) [molecule](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=6757)‐1 (ICAM‐1), [vascular cell adhesion molecule](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=6758)‐1 (VCAM‐1) and E‐selectin on the surface of endothelial cells in response to cytokines (TNF‐α and IL‐1α), and LPS (Fleming, 2007; Figure 5). We recently reported that epoxygenases are anti-inflammatory in human primary monocytes and macrophages (Bystrom et al., 2011), regulate M1 and M2 phenotype (Bystrom et al., 2011), and promote bacterial and lipid phagocytosis (Bystrom et al., 2013). In a mouse model of inflammatory resolution, we took this further using a CYP450 epoxygenase inhibitor SKF525A and sEH knockout mice (Gilroy et al., 2016). We reported how CYP450 epoxygenase‐derived mediators play a crucial role in controlling the infiltration of monocytes into sites of inflammation and are essential for the pro‐resolution phenotype of cells of the monocyte lineage (Gilroy et al., 2016) driving macrophage efferocytosis. Additionally, it was recently reported that EETs display analgesic bioactions during experimental inflammatory pain (Inceoglu et al., 2008). In general, CYP450‐derived epoxygenase products are anti‐atherosclerotic, vasodilatory, and anti‐inflammatory (Chaudhary et al., 2013), with the notable exception of linoleic acid‐ derived/EH product DiHOMEs. DiHOMEs have recently been shown to mediate thermal hyperalgesia (Zimmer et al., 2018) and, at high levels, are toxic to PMNs and were originally termed "leukotoxins" (Moghaddam et al., 1997).

The use of sEH inhibitors and sEH knockout mice has been invaluable to understanding the in vivo roles or epoxygenase products. Inhibiting sEH has revealed protective roles for epoxygenase products in injury‐induced vascular neointima formation (Revermann et al., 2010), atherosclerosis and aneurysm formation (Zhang et al., 2009), and inflammatory cell recruitment (Gilroy et al., 2016). sEH inhibition or overexpression of producing enzymes such as CYP2J2 are also protective in various acute inflammatory lung injury models (Revermann

et al., 2009). EETs released from platelets exert anti-thrombotic properties by inhibiting platelet aggregation induced by AA and vascular injury (Briggs, Xiao, Parkin, Shen, & Goldman, 2000). EETs can also increase the expression of tissue plasminogen activator in a cAMP‐ dependent mechanism, thus suggesting potentially important roles in controlling the fibrinolytic balance at sites of inflammation (Node et al., 2001).

The identification of epoxygenase‐product receptors has almost exclusively focused on AA‐derived EETs and HETEs, with very little research so far on other n3 and n6‐PUFA products. Putative recep-tor targets include the [TRP](http://www.guidetopharmacology.org/GRAC/FamilyDisplayForward?familyId=78) channels, the PPARs, and [GRP40](http://www.guidetopharmacology.org/GRAC/FamilyDisplayForward?familyId=24) (Bishop‐Bailey et al., 2014). EETs can directly activate PPAR‐γ in endothelial cells (Liu et al., 2005) and PPAR‐α in monocytes with EET‐mediated anti‐inflammatory effects blocked by PPAR‐γ (Liu et al., 2005) or PPAR‐α antagonists respectively. PPAR activation does not however account for all the anti-inflammatory effects of EETs. It has been suggested that the anti-inflammatory properties of EETs occurred through its ligation to a cell surface receptor. It was reported that EETs bind with high affinity to an "EET-receptor" on the surface of a monocytic cell line, belonging to a specific class of GPCRs (Behm, Ogbonna, Wu, Burns‐Kurtis, & Douglas, 2009). GRP40 can be activated by 14,15‐EET (Ma et al., 2015). However, it must be noted that GRP40 activation only occurred above 10 μM (Ma et al., 2015), whereas most biological effects occur in the nM range. These receptors are not present in all cells, and their known actions do not always correlate with the vascular and antiinflammatory activities of epoxygenase products. The identity of this receptor and its role, if any, in initiating the immuno-modulatory actions of EETs is yet to be determined. By contrast, intracellular signalling pathways are more established. Depending on the model system used, epoxygenases or its products can reduce cellular activation by inhibiting NFκB, inhibiting ERK activation, elevate cAMP, and/or induce cellular hyperpolarization (Thomson, Askari, & Bishop‐Bailey, 2012). Recently, it has also been proposed that inhibiting inflammatory endoplasmic reticulum stress may be critical for the beneficial effects of epoxygenase products, particularly in neuropathic pain.

As stated above, metabolites of the CYP hydroxylases also possess anti-inflammatory properties. For instance, 16-HETE can block

FIGURE 5 Biological properties of cytochrome p450 metabolites

the adhesion of leukocytes to the microvascular endothelium (Bednar et al., 2000) while also suppressing the synthesis of LTs as well as inhibiting rises in cerebrospinal fluid pressure, which represents index of tissue damage and swelling, in thrombo-embolic model of stroke in rabbits (Bednar et al., 2000). Moreover, PMN‐ derived 20‐HETE and 16‐HETE also counteract TX‐induced platelet aggregation (Hill, Fitzpatrick, & Murphy, 1992). Therefore, it can be surmised that not only do metabolites of CYPs maintain CV and renal systems but also regulate other diverse signalling pathways pertinent to fibrinolysis, platelet aggregation, inflammation, and cellular injury.

6 | LEUKOTRIENE AND LXs— BIOSYNTHESIS

LOX enzymes include 5-, 12-, or 15-LOX and are expressed in leukocytes, platelets, and endothelial cells respectively. 5‐LOX, for instance, metabolizes AA to the slow-reacting substances of anaphylaxis (LTC₄, LTD₄, and LTE₄: potent mediators of the allergic response; Lewis et al., 1980) as well as $LTB₄$, a powerful PMN and eosinophil chemoattractant (Borgeat & Samuelsson, 1979).

To date, four subtypes of [LT receptors](http://www.guidetopharmacology.org/GRAC/FamilyDisplayForward?familyId=35) have been described, namely, the BLT_1 and BLT_2 and two for the cysteinyl leukotrienes CysLT₁ and CysLT₂. Once bound, LTs signal via a G-protein in the cytoplasm to increase intracellular calcium and block formation of cAMP, which then modulates diverse cellular activities ranging from motility to transcriptional activation. While $CysLT_1$ receptors mediate mucus secretion, oedema accumulation, and broncho‐constriction in airways (Lynch et al., 1999), the Cys-LT₂ receptors drive inflammatory responses, tissue fibrosis in the lung, and vascular permeability (Beller et al., 2004). Not surprisingly, C ysLT₁ receptors are overexpressed in patients with chronic rhinosinusitis or asthma who have aspirin sensitivity (Sousa, Parikh, Scadding, Corrigan, & Lee, 2002). By comparison, the BLT1 receptor is a high-affinity receptor for $LTB₄$ and mediates its chemo-attractant and proinflammatory properties (Tager & Luster, 2003). Although BLT2 receptors act in a similar fashion to BLT1 receptors, $LTB₄$ affinity towards BLT1 receptors is much higher.

In contrast, [LXs](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=1034) are a series of trihydroxytetraene-containing bioactive eicosanoids that were first isolated from human leukocytes in the mid‐1980s (Serhan, Hamberg, & Samuelsson, 1984). However, in contrast to LTs, which are manufactured by intracellular biosynthesis, LXs are generated through cell–cell interactions by a process known as transcellular biosynthesis. In different human cell types, during the first biosynthetic step of LX biosynthesis, LOX inserts molecular oxygen into AA. This can be achieved by two major routes—the first pathway occurs in eosinophils, monocytes, or epithelial cells and involves the oxygenation of AA at C‐15 by 15‐ LOX yielding 15S-HPETE. Secreted 15S-HPETE is then taken up by monocytes or PMNs and converted to 5,6‐epoxytetraene by 5‐ LOX, which is then hydrolysed within these cells by either $LXA₄$ or $LXB₄$ hydrolase to $LXA₄$ or $LXB₄$. Activation of this pathway

concomitantly reduces LT synthesis, which requires 5‐LOX to convert AA into LTA₄ (Claria & Serhan, 1995). The second major route of LX biosynthesis occurs in an LTA₄-dependent manner and involves platelet–leukocyte interactions. 5‐LOX within leukocytes converts AA into LTA4, which when secreted is taken up by platelets adhering on the surface of the leukocyte and is subsequently transformed to $LXA₄$ and $LXB₄$. This occurs via the LX synthase activity of human 12‐LOX (Romano & Serhan, 1992). A third pathway of LX generation was discovered following aspirin ingestion, which irreversibly acetylates COX‐2 in endothelial cells and other activated cell types; this is a property specific to aspirin and not shared with other NSAIDs. Consequently, instead of COX‐2 converting AA into PGG₂, aspirin acetylation reprograms the enzyme resulting in the transformation of AA into 15R‐HETE (C‐15 alcohol carried in the R‐configuration). This is then metabolized in a transcellular manner by adherent leukocyte, vascular endothelial, or epithelial 5‐LOX to form 15 epimeric-LX (15-epi-LX) or aspirin-trigged LXs that carry their C‐15 alcohol in the R configuration rather than 15S native LX. Although initially thought to be only aspirin triggered, a pathway of endogenous 15‐epi‐LX generation has recently been described, where neuronal sphingosine kinase 1 mediates this COX‐2 acetylation (Lee et al., 2018). Aspirin‐triggered LXs share many of the immune regulatory characteristics of native LXs.

7 | LXs—RECEPTORS AND BIO‐ACTIONS

The biological actions of LXA₄ and 15-epi-LXs are mediated through [ALX](http://www.guidetopharmacology.org/GRAC/ObjectDisplayForward?objectId=223&familyId=23&familyType=GPCR) receptors, which are specific GPCRs, isolated and cloned in mouse, human, and rat tissues (Chiang, Takano, Arita, Watanabe, & Serhan, 2003). The ALX receptor is also known as the FPR2 receptor. Human ALX was identified and cloned in various leukocyte populations including T cells (Ariel, Chiang, Arita, Petasis, & Serhan, 2003), monocytes (Maddox et al., 1997), and tissue‐resident macrophages, synovial fibroblasts (Sodin‐Semrl, Taddeo, Tseng, Varga, & Fiore, 2000), and intestinal epithelial cells (Gronert, Gewirtz, Madara, & Serhan, 1998). LXA₄ and 15-epi-LX A_4 (but not LXB₄, LTB₄, LTD₄, or PGE₂) show high affinity towards ALX receptors (K_d = 1.7 nM). ALX receptors also have the ability to interact with other small peptides/proteins such as Ac2‐26 and glucocorticoid‐derived annexin-1, which carry out similar anti-inflammatory effects as the LXs and 15‐epi‐LXs. Studies in transgenic mice over‐expressing human ALX receptors showed that the protective and immune modulatory effects of LXs and 15‐epi‐LXs were ligand‐ and receptor‐dependent (Devchand et al., 2003). In a peritonitis model of zymosan‐induced acute inflammation, infiltration of neutrophils was substantially diminished in ALX transgenic mice compared to their wild-type equivalents (Devchand et al., 2003) with the site of LX action identified as the leukocyte/endothelial interface mediated by the generation of the anti-adhesive actions of NO (Paul-Clark, Van Cao, Moradi-Bidhendi, Cooper, & Gilroy, 2004).

15‐Epi‐LX analogues also regulate an ALX‐dependent p38/MAPK cascade, known to promote chemotaxis by inhibiting leukocyte‐

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specific AP-1 phosphorylation and activation (Ohira et al., 2004). In addition to ALX receptors, LXs also function as partial agonists to a subclass of rhodopsin receptors (Cys-LT₁) more commonly activated by LTs, mediating bioactions in several tissues and cell types other than leukocytes (Badr, DeBoer, Schwartzberg, & Serhan, 1989). At nanomolar concentrations, $LXA₄$ competes for binding with $LTD₄$ on mesangial cells (Badr et al., 1989) and HUVECs (Fiore, Romano, Reardon, & Serhan, 1993) as well as opposing the pro‐inflammatory effects of LTD₄. There is also evidence that another intracellular receptor, the Ah receptor, mediates the biological actions of LXs; Ah receptor is a ligand‐activated transcription factor that can trigger such anti-inflammatory events as the expression of suppressor of cytokine signalling 2 (Aliberti, Serhan, & Sher, 2002).

LXs are anti-inflammatory at nanomolar concentrations controlling both granulocyte and myeloid cell entry into sites of inflammation. Indeed, the ability of LXs to diminish neutrophil trafficking was corroborated when an analogue of 15‐epi‐LX was intravenously administered to BLT1 receptor knockout mice, that have dramatically elevated neutrophils in the lungs after high limb ischaemia– reperfusion (Chiang et al., 1999). Furthermore, research in our laboratory has found, in humans, that 15-epi-LXs regulate PMN influx in forearm blisters, accounting for low-dose aspirin's anti-inflammatory properties (Morris et al., 2009). Our additional work on resolving inflammation has revealed that humans fall into two categories, those who resolved their acute inflammatory responses in an immediate manner and those that show a more delayed or prolonged healing process, with the severity and duration controlled by expression of endogenous epi-LXs or ALX receptors (Morris et al., 2010). Paradoxically, while they inhibit neutrophil and eosinophil transmigration (Maddox et al., 1998), LXs promote monocyte infiltration into sites of inflammation, which when differentiated into macrophages bring about some of the key aspects of resolution and wound healing (Maddox & Serhan, 1996) without inducing neutrophil degranulation or release of other ROS (Jozsef, Zouki, Petasis, Serhan, & Filep, 2002).

Once at the site of inflammation and resolution, monocyte‐ derived macrophages are stimulated by LXs to ingest and clear apoptotic neutrophils (Godson et al., 2000), which may be facilitated by changes in the actin cytoskeleton (Maderna et al., 2002). Moreover, LXs increase levels of the anti‐inflammatory cytokine TGF‐β1, which, in turn, dampens a range of pro-inflammatory pathways (Bannenberg et al., 2005). LXs are also anti‐fibrotic, thereby improving tissue remodelling by reducing the proliferation of fibroblasts and mesanglial cells induced by a numbers of factors, including connective‐tissue growth factor, platelet‐derived growth factor, TNF‐α, LTD4, and TGF‐β (Leonard et al., 2002). 15‐Epi‐LXs exert the same biological effects as endogenously produced LXs, but with additional properties including causing increased vasorelaxation (Serhan, 1994) and endothelial cell production of anti‐inflammatory NO synthesis (Paul‐Clark et al., 2004). In addition, 15‐epi‐LX A4 inhibits TNF‐α‐induced IL‐1β in periodontitis in vivo (Hachicha, Pouliot, Petasis, & Serhan, 1999), down‐regulates suppressor of cytokine signalling 2 signalling (Machado et al., 2006), and dampens TNF‐α‐induced IL‐8 biosynthesis

(Gronert et al., 1998). As expected, LXs and 15‐epi‐LXs exert beneficial effects in a range of experimental models of inflammation and human diseases including cystic fibrosis (Karp, Flick, Yang, Uddin, & Petasis, 2005), glomerulonephritis (Munger et al., 1999), periodontitis (Pouliot, Clish, Petasis, Van Dyke, & Serhan, 2000), ischaemia– reperfusion injury (Chiang et al., 1999), various cutaneous inflammation models (Schottelius et al., 2002), pleuritis (Paul‐Clark et al., 2004), asthma (Levy et al., 2005), wound healing processes in the eye (Gronert et al., 2005), colitis, inflammation‐induced hyperalgesia in rats, and microbial infection in mice (Aliberti, Hieny, Reis e Sousa, Serhan, & Sher, 2002). See Table 1 for specialized pro‐resolving lipid mediators, their biological actions and concentrations at sites of inflammation.

8 | SPECIALIZED PRO‐RESOLVING LIPID MEDIATORS—BIOSYNTHESIS

Omega‐3 polyunsaturated fatty acids, including EPA and DHA, are known not only to maintain organ function and health but also to reduce the severity of inflammatory reactions and incidences of infection (Arita et al., 2005). Although also now known to be metabolized by COX, LOX, and CYP450 pathways into distinct lipid mediators, a novel series of ω‐3 PUFA products were identified in the resolving exudate of a mouse dorsal air pouch or peritonitis model using lipidomic and bio-informatic analysis (Lu, Hong, Tjonahen, & Serhan, 2005). These endogenous mediators are called [Rvs](http://www.guidetopharmacology.org/GRAC/LigandDisplayForward?ligandId=3934), PDs, and maresins (MaR).

EPA or DHA generate the Rvs and are categorized as members of the E‐series (from EPA) or D‐series (from DHA). Both series of Rvs were initially isolated from murine dorsal air pouches treated with EPA or DHA as well as aspirin. Transcellular formation of E‐ series Rvs occurs with the conversion of EPA to 18Rhydroxyeicosapentanoic acid by COX‐2 expressed within endothelial cells treated with aspirin. Similar to 15R‐HETE in 15‐epi‐LX formation, 18R-hydroxyeicosapentanoic acid is released from endothelial cells to neighbouring leukocytes for its conversion by 5‐LOX to either RvE1 or RvE2, via a 5 (Ariel et al., 2006) epoxide‐containing intermediate (Arita, Clish, & Serhan, 2005). This interaction is blocked by selective COX‐2 inhibition but not by indomethacin or paracetamol (Serhan et al., 2000). Although this transcellular route was proposed as the synthetic pathway for Rvs, intracellular production of Rvs and MaR have been observed in macrophages without the need for transcellular interactions. RvE1 is spontaneously produced in healthy subjects with levels increasing after treatment with either aspirin or EPA (Arita et al., 2005). D-series Rvs, aspirin-triggered resolvin D1 (RvD1; AT-RvD1), and RvD1, are synthesized via a pathway involving sequential oxygenations, initiated by 15‐LOX or aspirin‐acetylated COX‐2 in the microvasculature , respectively, followed by 5‐LOX in human neutrophils with an epoxide containing intermediate. For AT‐RvD1s, DHA is initially converted to epimeric 17R‐hydroxydocosahexaenoic acid. In the absence of aspirin, however, DHA is enzymically converted to 17S‐ hydroxydocosahexaenoic acid (Hong et al., 2003). Interestingly,

generation of E‐series Rvs can also be mediated by microbial and mammalian CYP450 enzymes, which convert EPA into 18‐HEPE. 18‐HEPE can then be transformed by human neutrophils into either RvE1 or RvE2 (Serhan et al., 2000). Hence, it is possible that microbes at sites of infection may contribute to the production of Rvs in a similar pathway.

DHA is also a precursor for the generation of PDs being enzymically converted by 15‐LOX to a 17S‐hydroperoxide‐containing intermediate. This intermediate is then converted by leukocytes into a 16(17)-epoxide that is subsequently converted in these cells to a 10,17‐dihydroxy‐containing compound (Hong et al., 2003). PDs are distinguished by the presence of a conjugated triene double bond and by their potent bioactivity. One specific DHA‐derived lipid mediator, 10,17S‐docosatriene, was termed protectin D1 (PD1), which when generated in neural tissue is called neuroprotectin D1. Moreover, PD1 exhibits tissue‐specific bioactivity as, in humans, this lipid is synthesized by peripheral blood mononuclear cells and Th2 CD4+ T-cells, while, in mice, it has been isolated from exudates and brain cells, human microglial cells (Serhan et al., 2002), and in peripheral blood (Hong et al., 2003).

9 | SPECIALIZED PRO-RESOLVING LIPID MEDIATORS IN INFLAMMATION AND RESOLUTION

One of the broader immunomodulatory properties of RvE1 is its ability to inhibit the accumulation of neutrophils and dendritic cells at sites of inflammation. This occurs by blocking trans‐endothelial migration of these cells across the microvascular endothelium as well as enhancing their clearance from inflammatory sites (Arita, Bianchini, et al., 2005). Other actions of RvE1 includes inhibition of ROS production from neutrophils in response to bacterial peptide, fMLP and TNFα (Gronert et al., 2004); inhibition of $LTB₄$ -BLT₁ receptor signalling via NF- κ B and hence the biosynthesis of pro-inflammatory chemokine and cytokines (Arita et al., 2007); enhancement of macrophage efferocytosis of apoptotic bodies (Schwab, Chiang, Arita, & Serhan, 2007); and up‐ regulation of the chemokine receptor CCR5 on late apoptotic neutrophils (Ariel et al., 2006), which, in turn, blocks chemokine signalling. RvE1 also regulates leukocyte pro-inflammatory cell surface markers including L‐selectin, while selectively disrupting TX‐mediated platelet aggregation (Dona et al., 2008), adding further insight into its anti-inflammatory/pro-resolution properties. In disease states, RvE1 suppresses Porphyromonas gingivalis‐induced oral inflammation and bone loss during periodontitis (Hasturk et al., 2006), is protective in trinitrobenzene‐sulphonic acid‐induced colitis in rodents (Arita, Yoshida, et al., 2005), and mediates re‐epithelisation of mouse cornea after thermal‐injury (Gronert et al., 2005). Taken together, RvE1 triggers various aspects of the pro‐resolution cascade ranging from the timely inhibition of granulocyte accumulation at sites of inflammation to the efferocytosis or clearance of inflammatory debris (see Serhan, 2008).

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RvE1 binds to **[ChemR23](http://www.guidetopharmacology.org/GRAC/FamilyDisplayForward?familyId=338)** (now Chemerin₁ receptors, with high affinity (K_d = 48.3 nm) resulting in the down-regulation of NF- κ B activity and consequently the synthesis of pro-inflammatory cytokines such as TNF‐α as well as modulating pathways involved in MAPK signalling (Arita, Bianchini, et al., 2005). Although it has been found in the kidney, gastrointestinal system, brain, and CV tissue and cells of the myeloid lineage, the levels of ChemR23 receptor expression are highly variable. For example, these receptors are significantly increased on human monocytes but comparatively less so on neutrophils in response to anti‐inflammatory mediators such as TGF‐β. As with ALX receptors, ChemR23 is also a receptor for peptide ligands including chemerin, which also exerts anti-inflammatory actions (Cash et al., 2008). RvE1 also interacts with the LTB₄ receptor, BLT1, and is a partial antagonist preventing neutrophil activation (Arita et al., 2007). Therefore, RvE1 couples to two distinct receptors to suppress pro‐ inflammatory mechanisms while enhancing pro‐resolution pathways.

While structurally distinct from RvE1, RvE2 is a second member of the EPA‐derived family of E‐series Rvs. In PMNs from human, it is generated at higher concentrations than RvE1, but is equipotent when given intravenously and additive when administered alongside RvE1 (Tjonahen et al., 2006). RvE2 also suppresses PMN migration into the peritoneum after zymosan (Tjonahen et al., 2006). It is still unclear what receptor RvE2 couples to, as it mediates resolution by activating Chemerin₁ receptors and antagonizing the LTB₄ receptor BLT1.

There are four members of the D-series Rvs, namely, RvD1, RvD2, RvD3, and RvD4 (Hong et al., 2003). As with RvE1, RvD1/D2 exerts both anti-inflammatory and pro-resolution properties by blocking neutrophil infiltration, while also enhancing macrophage efferocytosis of apoptotic bodies (Krishnamoorthy et al., 2010). The latter occurs via the binding of RvD1 to either ALX receptors or the orphan GPR32, which are present on the surface of monocytes and PMNs, the expression of which is up‐regulated by inflammatory stimuli including granulocyte-macrophage-colonystimulating factor and zymosan (Krishnamoorthy et al., 2010). Importantly, in a model of cecal ligation and puncture RvD2, whose receptor is GPR18 (Chiang, Dalli, Colas, & Serhan, 2015), markedly reduced PMN accumulation, bacteria numbers and proinflammatory cytokines leading to increased animal survival.

As already mentioned, in addition to D-series Rvs, DHA also acts as a precursor for the synthesis of PDs. PD1, for instance, is synthesized in the human brain and microglial (Serhan et al., 2002) and peripheral blood mononuclear cells (Hong et al., 2003). As with Rvs, PD1 may also inhibit PMN migration as well as toll-like receptor-mediated activation (Duffield et al., 2006) while suppressing Th2 inflammatory cytokines and pro‐inflammatory lipid mediator synthesis (Levy et al., 2007). PD1 also blocks T‐cell migration in vivo and promotes T-cell apoptosis (Ariel et al., 2005). PD1 is protective in experimental models of oxidative stress (Mukherjee et al., 2007), ischaemic stroke (Marcheselli et al., 2003), ischaemia–reperfusion renal injury (Duffield et al., 2006), asthma (Levy et al., 2007), and Alzheimer's (Lukiw et al., 2005). Indeed, peripheral blood mononuclear cells from Alzheimer's patients given a DHA‐rich dietary supplement show dampened

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biosynthesis of IL‐1β, IL‐6, and granulocyte‐colony‐stimulating factor (Vedin et al., 2008). As with RvE2, a receptor is yet to be identified. However, it is possible that it couples to a different receptor from that for RvE1, as its anti‐inflammatory effects are additive with those of RvE1 in vivo.

MaR1 and MaR2 are produced in tissues by macrophages via the actions of 12‐LOX, through a 13,14‐epoxide intermediate (Serhan et al., 2009). MaR1 can also be generated at sites of vascular inflammation during human platelet–neutrophil interactions via platelet 12-LOX conversion of DHA to 13S,14S-epoxy-MaR, followed by neutrophil conversion to MaR1 (Abdulnour et al., 2014). The receptors for MaR have yet to be identified. Although MaR have only been recently discovered, it has been reported similar to Rvs and PD1, MaR1 blocks the infiltration of PMNs while stimulating macrophage phagocytosis of apoptotic PMNs or zymosan.

10 | SUMMARY

Inflammation is a good thing; it kills bacteria and helps to heal wounds while imparting long-term memory against inciting antigens. Lipids play a key role in these events and come in many forms, including those that drive the cardinal signs of inflammation and those that help to restrain it and bring the response to a timely end. In fact, studying lipids and their inhibitors, NSAIDs, has given us a great deal of insight into homeostasis, immune responses to infection/injury and the wound healing process. Indeed, inflammatory onset has been a historical point of interest for the development of anti-inflammatory drug therapies. Research on the other end of the inflammatory spectrum, resolution, has provided the opportunity to harness endogenous mediators and their receptors to help drive ongoing inflammation down a pro-resolution pathway. Moreover, this is achievable without compromising host defence. Such complex manipulation of the immune system provides new opportunities to develop further pro‐ resolution therapies based upon what we have learned from studying lipids in this setting.

10.1 | Nomenclature of targets and ligands

Key protein targets and ligands in this article are hyperlinked to corresponding entries in [http://www.guidetopharmacology.org,](http://www.guidetopharmacology.org) the common portal for data from the IUPHAR/BPS Guide to PHARMA-COLOGY (Harding et al., 2018), and are permanently archived in the Concise Guide to PHARMACOLOGY 2017/18 (Alexander, Christopoulos et al., 2017; Alexander, Cidlowski et al., 2017; Alexander, Striessnig et al., 2017).

CONFLICT OF INTEREST

The authors declare no conflicts of interest.

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