CYP450-derived oxylipins mediate inflammatory resolution

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Resolution of inflammation has emerged as an active process in immunobiology, with cells of the mononuclear phagocyte system being critical in mediating efferocytosis and wound debridement and bridging the gap between innate and adaptive immunity. Here we investigated the roles of cytochrome P450 (CYP)-derived epoxy-oxylipins in a well-characterized model of sterile resolving peritonitis in the mouse. Epoxy-oxylipins were produced in a biphasic manner during the peaks of acute (4 h) and resolution phases (24–48 h) of the response. The epoxygenase inhibitor SKF525A (epoxI) given at 24 h selectively inhibited arachidonic acid- and linoleic acid-derived CYP450-epoxy-oxlipins and resulted in a dramatic influx in monocytes. The epoxI-recruited monocytes were strongly GR1^{+}, Ly6c^{hi}, CCR2^{hi}, CCL2^{hi}, and CX3CR1^{lo}. In addition, expression of F4/80 and the recruitment of T cells, B cells, and dendritic cells were suppressed. sEH (*Ephx2*)$^{-/-}$ mice, which have elevated epoxy-oxylipins, demonstrated opposing effects to epoxI-treated mice: reduced Ly6c^{hi} monocytes and elevated F4/80^{hi} macrophages and B, T, and dendritic cells. Ly6c^{hi} and Ly6c^{lo} monocytes, resident macrophages, and recruited dendritic cells all showed a dramatic change in their resolution signature following in vivo epoxI treatment. Markers of macrophage differentiation CD11b, MerTK, and CD103 were reduced, and monocyte-derived macrophages and resident macrophages ex vivo showed greatly impaired phagocytosis of zymosan and efferocytosis of apoptotic thymocytes following epoxI treatment. These findings demonstrate that epoxy-oxylipins have a critical role in monocyte lineage recruitment and activity to promote inflammatory resolution and represent a previously unidentified internal regulatory system governing the establishment of adaptive immunity.

oxylipins | resolution | monocyte | phagocytosis | epoxygenase

Monocytes and monocyte-derived macrophages play a critical role in chronic inflammation, in part via the production and release of lipid mediators (1). One such lipid precursor, arachidonic acid, is metabolized into families of biologically active mediators by the cyclooxygenase, lipoxygenase, and cytochrome P450 (CYP) pathways (2, 3). CYPs metabolize arachidonic acid by: (*i*) an epoxygenase activity that catalyzes the conversion of arachidonic acid to epoxyeicosatrienoic acids (EETs); (*ii*) a lipoxygenase-like activity that metabolizes arachidonic acid to midchain hydroxyeicosatetraenoic acids (HETEs); and (*iii*) ω- and ω-1-hydroxylase activity, which produces ω-terminal HETEs (3). In addition to arachidonic acid, CYPs with epoxygenase activity can also metabolize alternative polyunsaturated fatty acids such as linoleic acid and docosahexaenoic acid into a series of products including epoxyoctadacamonoenoic acids (EpOMEs) and 19,20-epoxydocosapentaenoic acid (EpDPE), respectively, whose functions remain poorly understood (3–5).

The main polyunsaturated fatty acid-metabolizing CYPs belong to the CYP2 family, in particular the CYP2J and CYP2C subfamilies (3, 4, 6, 7). Moreover, these CYP-lipid–metabolizing enzymes are the primary sources of eicosanoids in small blood vessels, the kidney, liver, lung, intestines, heart, and pancreas (3, 7). In most organs, EETs and related epoxygenase products are metabolically unstable and are rapidly metabolized. The major pathway that regulates EET metabolism is that catalyzed by epoxide hydrolases (8), which convert EETs to less biologically active dihydroxyeicosatrienoic acids (DHETs) (9). EpOMEs similarly get converted into dihydroxyoctadecenoic acids (DiHOMEs), whereas 19,20-EpDPE gets converted into 19,20-dihydroxydocosapentaenoic acid (DiHDPA). Elevating the levels of endogenous CYP products by disrupting (knockout) or inhibiting soluble epoxide hydrolase (sEH) reduces neointima formation (10), atherosclerosis, abdominal aortic aneurysm, dyslipidemia (11), hypertension (12), and diabetes (13) in different mouse models, all of which to some extent one could argue have a degree of nonresolving inflammation.

Over the last 15 y there has been a vast increase in our knowledge of fatty acid mediators that regulate inflammatory processes, particularly newly identified mediators such as the resolvins that mediate the resolution of inflammation (14–16). However, unlike cyclooxygenase and lipoxygenase products, the roles of CYP450 pathways in chronic inflammation remain unclear. The arachidonic acid products of the CYP epoxygenases, the EETs, can regulate

Significance

A number of lipid mediators are known to contribute to inflammatory resolution. Fatty acid metabolites of cytochrome P450 (CYP) enzymes are found in abundance; however, their roles in inflammatory resolution are not known. Targeted lipidomics revealed that CYP450-epoxy-oxylipins were present during acute inflammation and inflammatory resolution. Using mice lacking soluble epoxide hydrolase, the major metabolizing pathway for CYP450-derived fatty acid mediators, and CYP450 epoxygenase inhibition specifically during resolution, we show that CYP450-derived lipids dramatically limit the accumulation of inflammatory monocytes during resolution. Moreover, all cells of the monocyte lineage examined showed a dramatic alteration in their proresolution phenotype following epoxygenase inhibition. These findings demonstrate that the CYP450-epoxy-oxylipins pathway has a critical role in monocyte lineage recruitment and resolution activity during inflammatory resolution.

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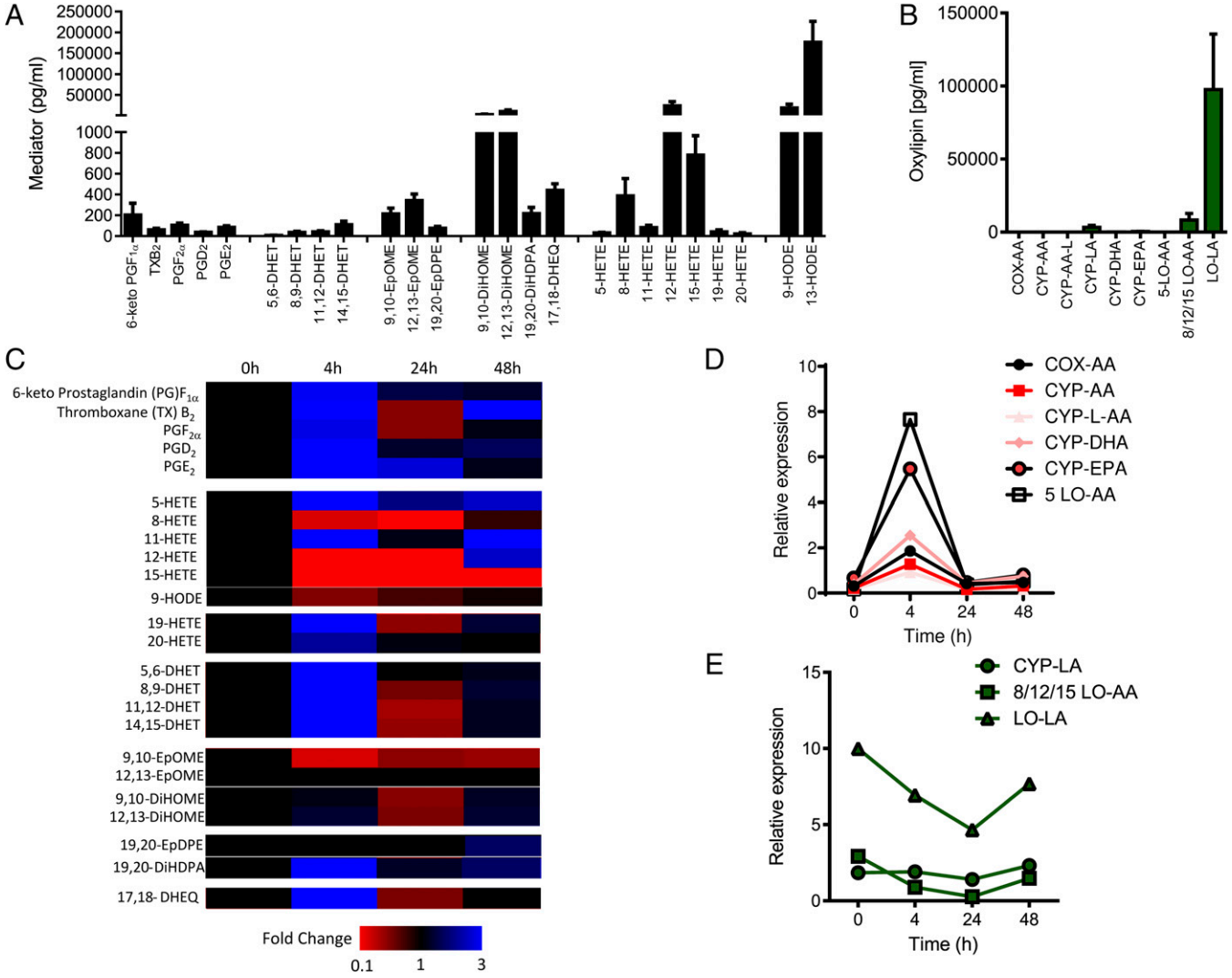

Fig. 1. LC/MS/MS analysis of oxylipin-generating pathways in the mouse peritoneal cavity during acute inflammation and resolution initiated by zymosan A. (*A*) Absolute levels of oxylipins determined in naive peritoneal cavity lavage fluid of male C57BL/6 mice. (*B*) Sum of oxylipin-generated COX, CYP450, CYP450-lipoxygenase–like (CYP-L), or LO pathways using either arachidonic acid (AA), linoleic acid (LA), DHA, or EPA as a substrate in the naive peritoneal cavity. (*C*) Heatmap showing fold changes in oxylipin formation following zymosan A treatment (1 mg, i.p.) from 0 to 4 h (peak of acute inflammation), 24 h, and 48 h (resolution). (*D* and *E*) Fold-normalized square root of oxylipins produced by COX-AA (prostanoids), CYP-AA (DHETs), CYP-L-AA (19- and 20-HETE), CYP-DHA (19,20-EpDPE and 19,20-DiHDPA), CYP-EPA (17,18-DHEQ), and 5-LO-AA (5-HETE) pathways (*D*) and CYP-LA (EpOMEs and DiHOMEs), 8/12/15-LO (8-, 12-, and 15-HETE), and LO-LA (HODEs) pathways (*E*) in response to zymosan A over 48 h. Pathways in *D* and *E* represent two distinct responses to challenge with zymosan A. Data represent the mean ± SEM from $n = 4$–8 mice per group.

vascular tone, smooth muscle cell mitogenesis, platelet aggregation, steroidogenesis, and endothelial and vascular smooth muscle cell activation (4, 5, 7, 17–19). We recently published that in human monocytes and macrophages, epoxygenases and some of their arachidonic acid products were antiinflammatory through their ability to activate the peroxisome proliferator-activated receptor (PPAR), in particular PPARα (20, 21). Overexpression of epoxygenase enzymes CYP2J2 and CYP2C8 or genetic disruption of sEH (sEH$^{-/-}$) inhibits LPS-induced pulmonary inflammation (22, 23), and sEH$^{-/-}$ mice or treatment with sEH inhibitors is highly effective against inflammatory and neuropathic pain (24–27).

Monocytes are heterogeneous in mice and in humans (28). In mice, monocyte subsets can be divided based on the expression of Ly6c, Gr1, CC-chemokine receptor 2 (CCR2), and CX3C-chemokine receptor 1 (CX3CR1). Ly6c^{hi} monocytes are Gr1^{+}, CCR2^{+}, and CX3CR1^{lo}, whereas Ly6c^{lo} monocytes are Gr1^{-}, CCR2^{-}, and CX3CR1^{hi} (29, 30). Lipid mediators that regulate the recruitment and phenotype of monocytes are poorly understood.

Herein, using a sterile model of inflammatory resolution dependent on monocyte recruitment, we found that CYP-epoxygenase products not only accumulate in a temporal manner with monocyte recruitment but also limit proinflammatory monocyte recruitment and promote a proresolution phenotype in cells of the monocyte lineage.

Results

CYP450-Derived Lipid Metabolites Form Part of the Coordinated Eicosanoid Fatty Acid Metabolite Response in Acute Inflammation and Resolution. The naive mouse peritoneal cavity contains high levels of linoleic acid [9- and 13-hydroxy-octadecadienoic acid (HODE)] and arachidonic acid (8-, 12-, and 15-HETE)-derived lipoxygenase products as well as 9,10- and 12,13-DiHOME, which are epoxide diols of the linoleic acid epoxygenase products 9,10-EpOME and 12,13-EpOME (Fig. 1). Cyclooxygenase (COX), CYP450, 5-lipoxygenase (LO), and eicosapentaenoic acid (EPA)/docosahexaenoic acid (DHA)-derived mediators, including prostanoids, leukotrienes, EETs, and their metabolites,

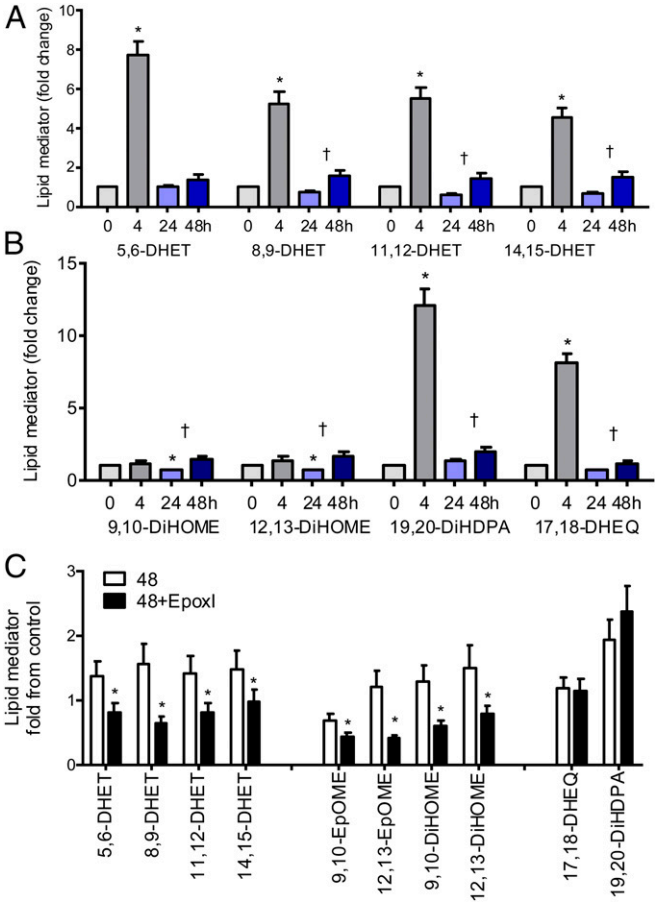

Fig. 2. Epoxygenase products form in a biphasic manner mirroring acute inflammation and inflammatory resolution: selective inhibition of CYP-AA and CYP-LA products by epoxI. (*A* and *B*) Mean fold change in (*A*) CYP-AA products: DHETs and (*B*) CYP-LA (DiHOME), CYP-DHA (DiHDPA), and CYP-EPA (DHEQ) products, 0–48 h. Each mediator was normalized to its paired levels found in the naive cavity in each experiment; the levels found in the naive cavity were given an arbitrary value of 1. $^{*}P < 0.05$ by one-sample *t* test from 0 h. $^{\dagger}P < 0.05$ by unpaired *t* test between the values observed at 24 and 48 h. (*C*) Mean fold-change in epoxygenase product formation in the presence or absence of epoxI (30 mg/kg). EpoxI was given at 24 and 36 h. Epoxy-oxylipins were measured at 48 h and are represented as fold change from the respective mean levels observed in the naive cavity at 0 h. $^{*}P < 0.05$ by unpaired *t* test. Data represents mean ± SEM.

were found at much lower levels (Fig. 1). The liquid chromatography/tandem mass spectrometry (LC/MS/MS) assay we used was primarily designed to measure CYP450 and related products. We do not currently measure some of the other newly characterized lipid species known to regulate inflammatory resolution such as the resolvins, but clearly enzymatic pathways and lipid precursors for these pathways are all also present. During the acute inflammatory response elicited by zymosan (4 h), there was the expected burst of prostanoids, increases in 5-, 11-, 19-, and 20-HETE, and also increases in the CYP450 epoxygenase-derived oxylipins 5,6-, 8,9-, 11,12-, and 14,15-DHET and 17,18-dihydroxy-EEQ (DHEQ) (Fig. 1 *C* and *D* and *SI Appendix*, Fig. S1). The DHETs are the epoxide hydrolase diols of the P450 metabolites and could readily be detected, whereas parental EETs were not seen. The 15/12-lipoxygenase products initially found at high levels in the naive cavity dropped dramatically during acute inflammation (9- and 13-HODE and 8-, 12-, and 15-HETE), with only 9-HODE and 8-HETE returning to baseline levels over the 48-h time course (Fig. 1 *C* and *E* and *SI Appendix*). After acute inflammation, which is driven primarily by polymorphonuclear cells (PMNs), resolution begins and is associated with an accumulation of proresolution monocytes peaking 24–48 h after the initial inflammatory insult. After a drop at 24 h, this influx in monocytes was associated with a second phase of epoxy-oxylipin production of arachidonic acid (DHETs), linoleic acid (DiHOMEs), EPA (17,18-DHEQ), and DHA (19,20-DiHDPA) epoxygenase products (Fig. 2 *A* and *B*). Moreover, this phase was associated with the biosynthesis of new lipid products, as the addition of a selective epoxygenase inhibitor (epoxI; SKF525A; 30 mg/kg, i.p.) to zymosan-treated mice from 24 h onward led to a selective and specific inhibition of arachidonic acid and linoleic acid epoxy-oxylipins (Fig. 2*C*). EpoxI had no significant effect on COX, LO, or CYP450-lipoxygenase–like products (*SI Appendix*, Table S1), indicating the epoxI used is highly selective for the CYP-epoxygenase pathway.

We screened the mouse Cyp2 family to investigate which enzymes were present in the resolution-phase inflammatory cell populations, and found Cyp2c44, Cyp2j5, Cyp2j6, Cyp2j9, and Cyp2j13 epoxygenases along with *Cyp2u1* (HETE and DHA metabolism) and Cyp2s1 (xenobiotic and retinoic acid metabolism) (Table 1 and *SI Appendix*, Fig. S2).

Epoxygenase Inhibition During Resolution Induces a Profound Accumulation of Ly6c^{hi} Monocytes. Epoxygenase inhibition was accompanied by a threefold increase in monocyte numbers during the 24- to 48-h period of inhibition and a small but significant increase in PMNs (Fig. 3*A*). These epoxI-recruited monocytes were strongly GR1^{+} (Ly6c and Ly6g antigens; Fig. 3*B*), and Ly6c^{hi} expression was confirmed by RT-PCR (Fig. 3*C*). Moreover, these recruited cells were CCR2^{hi} (Fig. 3*D*), CX3CR1^{lo} (Fig. 3*E*), and CCL2^{hi} (Fig. 3*F*) at the mRNA level. CCL2 peptide was found to be highly elevated in the peritoneal cavity of epoxI-treated mice (vehicle 8 ± 8 pg/mL; epoxI 266 ± 190 pg/mL CCL2; $n = 3$). Substantial levels of CCL2 were produced exclusively from cells elicited from epoxI-treated mice ex vivo (Fig. 3*G*), suggesting they could be a major source of this elevated CCL2. EETs are too short-lived to be given as a bolus dose in vivo, and therefore to test whether EETs were antiinflammatory in mouse monocytes, we added back authentic 14,15-EET (1 μM; 7 h) to zymosan-elicited cells ex vivo. 14,15-EET

Table 1. Expression of mouse Cyp2s in the resolving peritoneal cavity of mice

Cyp	Liver	Exudate
Cyp2j5	+	+/–
Cyp2j6	+	++
Cyp2j9	+/–	+
Cyp2j13	Not tested	+/–
Cyp2c29	+	–
Cyp2c37	ND	ND
Cyp2c38	+	–
Cyp2c39	–	–
Cyp2c40	ND	ND
Cyp2c44	+++	++
Cyp2c50	++	–
Cyp2c54	++	–
Cyp2c55	+	–
Cyp2a1	+++	–
Cyp2u1	ND	+
Cyp2s1	ND	+

Mice were treated with zymosan (1 mg, i.p.) for 24 h. The presence of mouse CYP2 mRNAs was determined by RT-PCR, using mouse liver as a positive control. The data summarize the findings from $n = 6$–8 mice. The number of + signs indicates the relative expression level; +/– indicates that not all samples tested showed positive expression. ND, not detected.

reduced Ccl2 (Fig. 3*H*), inducible nitric oxide synthase (iNOS) (Fig. 3*I*), and IL-12 (Fig. 3*J*) mRNA expression in these cells ex vivo. Addition of 14,15-EET was not universally antiinflammatory in these elicited cells, as TNFα mRNA expression remained unaltered (Fig. 3*H*). These results are consistent with our previous findings, which reported an antiinflammatory role of epoxygenases in classically activated human monocytes and macrophages in vitro (21).

sEH$^{-/-}$ Mice Have an Opposing Phenotype to Epoxygenase-Inhibited Mice During Resolution. To confirm the role of epoxy-oxylipins during resolution, we compared the findings of epoxI-treated mice with sEH$^{-/-}$ mice. sEH is considered the main pathway for epoxy-oxylipin metabolism and inactivation. Compared with epoxI-treated mice, which have reduced epoxy-oxylipins, sEH$^{-/-}$ mice have elevated oxylipins (31). In zymosan-treated mice, Ly6c^{hi} monocytes peaked at 48 h with epoxI treatment (*SI Appendix*, Fig. S3), whereas at 48 h in sEH$^{-/-}$ mice, Ly6c^{hi} monocytes were reduced compared with wild-type controls (Fig. 4*A*). There was a small increase in Ly6g^{+} cells at 48 h with epoxI, whereas in sEH$^{-/-}$ mice, Ly6g^{+} monocytes were slightly reduced compared with wild-type controls (Fig. 4*B*). With epoxI treatment, although there was no change in numbers of macrophage (F4/80)-positive cells (5.4 ± 0.5 compared with 5.9 ± 2.2 cells per milliliter, $\times 10^6$), there was a reduction in the levels of F4/80 expression on the epoxI-elicited macrophages (Fig. 4*C*). In corollary, the opposite was found in sEH$^{-/-}$ mice: CD11b^{++} F480^{++} cells were greatly elevated compared with wild-type controls (Fig. 4*C*). CD19^{+} B cells (Fig. 4*D*) and CD3^{+} T cells (Fig. 4*E*) were reduced by epoxI and elevated in sEH$^{-/-}$ mice. There was also a similar trend to reduction of MHCII^{+} CD11c^{+} dendritic cells by epoxI, with a significant elevation of these MHCII^{+} CD11c^{+} dendritic cells in sEH$^{-/-}$ mice. Therefore, in all of the major indices of inflammatory cell accumulation tested, epoxy-oxylipin inhibition with epoxI showed opposing actions to epoxy-oxylipin elevation with the use of global sEH$^{-/-}$.

Cells of the Monocyte Lineage Have a Disrupted Resolution Phenotype Following Epoxygenase Inhibition. Using a recently identified novel panel of quantitative (q)RT-PCR of resolution monocyte markers (32), we examined the total inflammatory cell population for a changes in the resolution phenotype. The total epoxI-elicited cell population was Timd4^{lo}, Tgfb2^{lo}, and Plxdc2^{lo} and IL1f9^{hi}, CD86^{hi}, and Ms4a7^{hi} compared with cells from animals treated with vehicle/zymosan alone (*SI Appendix*, Fig. S4). At indicated time points there was also a significant difference in Ccnb2 (increased with epoxI) and Aspa and Stfa2l1 (decreased with epoxI) (*SI Appendix*, Fig. S4). Because epoxy-oxylipins appear to have the most dramatic effect on monocyte lineage cell recruitment, we examined these targets in FACS-sorted Ly6c^{hi} and Ly6c^{lo} monocyte populations, resident macrophages, and recruited dendritic cells (DCs) (Fig. 5). Normal-resolution Ly6c^{hi} monocytes were Plxcd2^{hi}, Aspa^{hi}, and F5^{hi}; Ly6c^{lo} monocytes were Ccnb2^{hi}, Timd4^{hi}, and Stfa2l1^{hi}; resident macrophages were IL1F9^{hi}; and dendritic cells

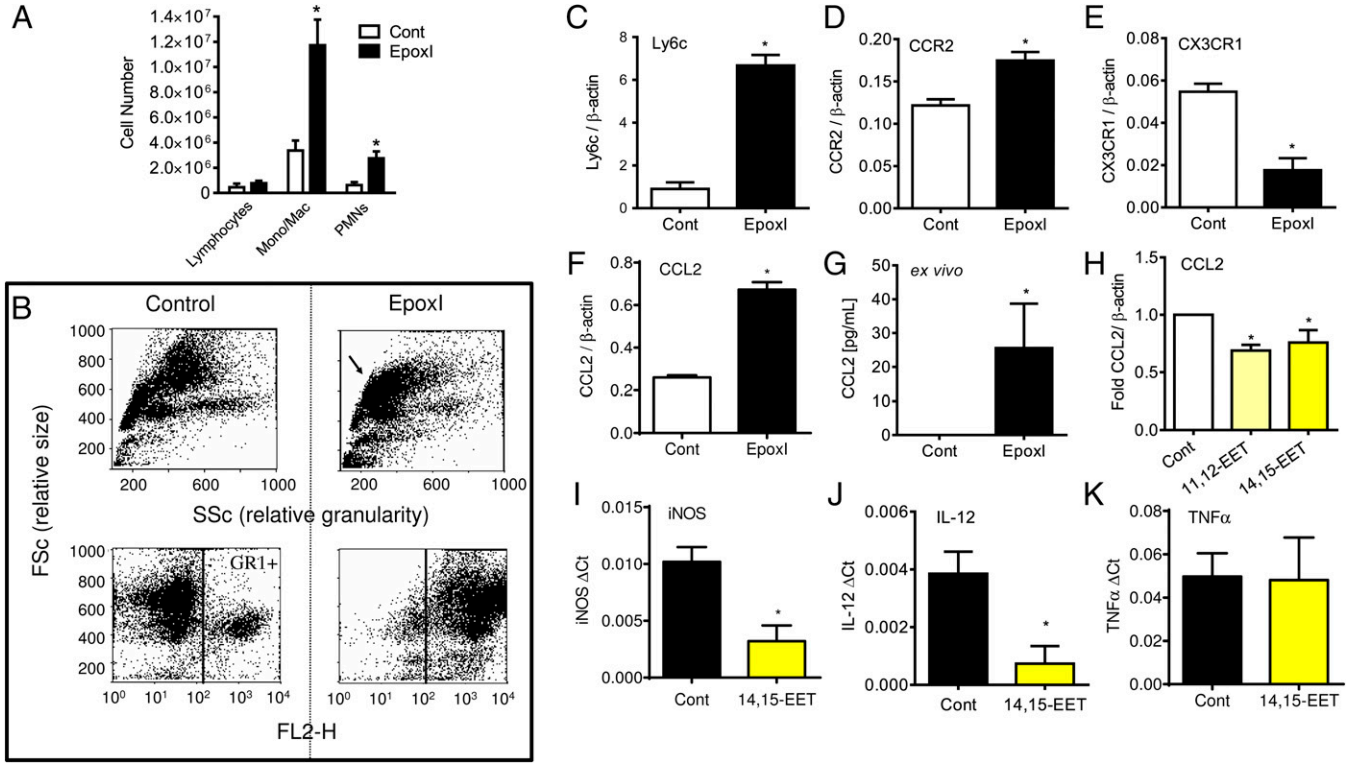

Fig. 3. Epoxygenase inhibition during resolution causes Ly6c^{hi} monocyte recruitment and Ccl2 generation. (*A*) Change in lymphocytes, monocytes/macrophages, and PMNs in the peritoneal cavity of mice 48 h postzymosan treatment. At 24 and 36 h, mice were given either sterile PBS or epoxI (30 mg/kg, i.p.). Total cell numbers were counted on a hemocytometer and the proportion of each cell type was determined by FACS. Data represent the mean ± SEM from $n = 6$ mice per group; $^*P < 0.05$ by unpaired *t* test between vehicle (Cont; PBS) and epoxI-treated mice. (*B*) Example of FACS analysis of total cell (*Upper*) and GR^{+} populations (*Lower*) of cells elicited by zymosan alone (control; *Left*) or in the presence of epoxI (*Right*) in terms of size (FSc) and granularity (SSc). (*C–F*) Relative expression of Ly6c (*C*), Ccr2 (*D*), CX3CR1 (*E*), and CCL2 (*F*) mRNA compared with β-actin in elicited cells from zymosan + control or epoxI-treated mice taken at 48 h. (*G*) CCL2 generation in cells from zymosan + control or epoxI-treated mice. Cells were elicited at 36 h and left for a further 8 h ex vivo and CCL2 was measured by ELISA. (*H–K*) CCL2 (*H*), iNOS (*I*), IL-12 (*J*), and TNFα (*K*) mRNA expression in zymosan-elicited cells at 36 h treated with 14,15-EET or 11,12-EET. Cells were elicited at 36 h and treated for a further 7 h with EETs. Data represent the mean ± SEM from $n = 3–4$ mice per group; $^*P < 0.05$ by unpaired *t* test between vehicle (Cont; PBS) and epoxI-treated mice.

were Ccr2^{hi} and IL1F9^{hi} (Fig. 5). Treatment with epoxI caused a significant down-regulation of Plxcd2 in Ly6c^{hi} monocytes and DCs; Ccna2 in Ly6c^{hi} and Ly6c^{lo} monocytes, resident macrophages, and DCs; Ccnb2 in Ly6c^{lo} monocytes and DCs; Aspa in Ly6c^{hi} and Ly6c^{lo} monocytes; F5 in Ly6c^{hi}, resident macrophages, and DCs; Tgfb2 in Ly6c^{hi} and Ly6c^{lo} monocytes, resident macrophages, and DCs; and Timd4 in Ly6c^{lo} monocytes, resident macrophages, and DCs (Fig. 5*A*). Treatment with epoxI caused an up-regulation of Ccr2 in Ly6c^{lo} monocytes, resident macrophages, and DCs; Ccl2 in Ly6c^{hi} monocytes, resident macrophages, and DCs; Ms4A7 in Ly6c^{hi} monocytes, resident macrophages, and DCs; CD86 in Ly6c^{lo} monocytes and DCs; and IL1F9 in Ly6c^{hi} and Ly6c^{lo} monocytes (Fig. 5*B*). However, Stfa2l1 was up-regulated in Ly6c^{hi} and resident macrophages but inhibited in Ly6c^{lo} monocytes. These findings collectively point toward a more inflammatory phenotype accompanying epoxygenase inhibition, with a clear induction in either or both Ccl2 and Ccr2 in all of the cell types and a decrease in resolution monocyte markers of apoptosis and repair (Timd4 and Tgfb2, respectively).

Epoxygenase Inhibition Disrupts Resolution-Phase Macrophage Differentiation and Leads to Impaired Phagocytosis and Efferocytosis. To examine functional changes in the monocyte lineage, we examined Ly6c^{+} monocytes, resident macrophages, and monocyte-derived macrophage populations. As previously shown, Ly6c^{+} monocytes were elevated, monocyte-derived macrophages were reduced, and resident macrophage levels remained unchanged (Fig. 6). Resident macrophages on a cell-per-cell level showed reduced expression of differentiation markers and phagocytosis receptors F4/80, CD11b, MerTk, Timd4, and CD103 but an increase in CD64 (Fig. 6*C*). A similar reduction in F4/80, MerTK, Timd4, and CD103 was also observed in monocyte-derived macrophages (Fig. 6*C*). In contrast, Ly6c^{+} monocytes demonstrated a small but significant increase in markers of differentiation CD11b, Timd4, and CD64, which was again accompanied by a decrease in CD103 (Fig. 6*C*).

Using the ImageStream system, we were easily able to identify phagocytosis from particles and bodies sticking to cells (Fig. 7*A*). Ly6c^{+} monocytes had a small but significant decrease in the ability to phagocytose zymosan, but not apoptotic cells (Fig. 7*B*). In contrast, resident macrophages and monocyte-derived macrophages (Fig. 7*B*) both showed greatly reduced ability to phagocytose FITC-labeled zymosan A and efferocytose apoptotic cells ex vivo.

Discussion

Our findings show that epoxy-oxylipin–generating pathways constitute an important control point limiting the accumulation of proinflammatory Ly6c^{hi} monocytes and the proinflammatory and clearance activity of cells of the monocyte lineage during inflammatory resolution (Fig. 8). Similar to cyclooxygenases and lipoxygenases, the epoxygenase pathways of arachidonic acid metabolism (*SI Appendix*, Fig. S5) are activated during both acute inflammation and resolution. There has been great recent interest in the therapeutic potential of sEH inhibitors as novel antiinflammatories. sEH inhibitors or genetic disruption of sEH in mice reduces inflammation in models of endotoxin-induced pulmonary inflammation (22), ischemia-reperfusion injury (33, 34), subarachnoid hemorrhage (35), and the murine ovalbumin model of asthma (36), and in more chronic models including atherogenic diet-induced fatty liver disease and adipose tissue (37) and atherosclerosis (38, 39). In contrast, the expression and roles of epoxy-oxylipins during inflammatory resolution have not been investigated.

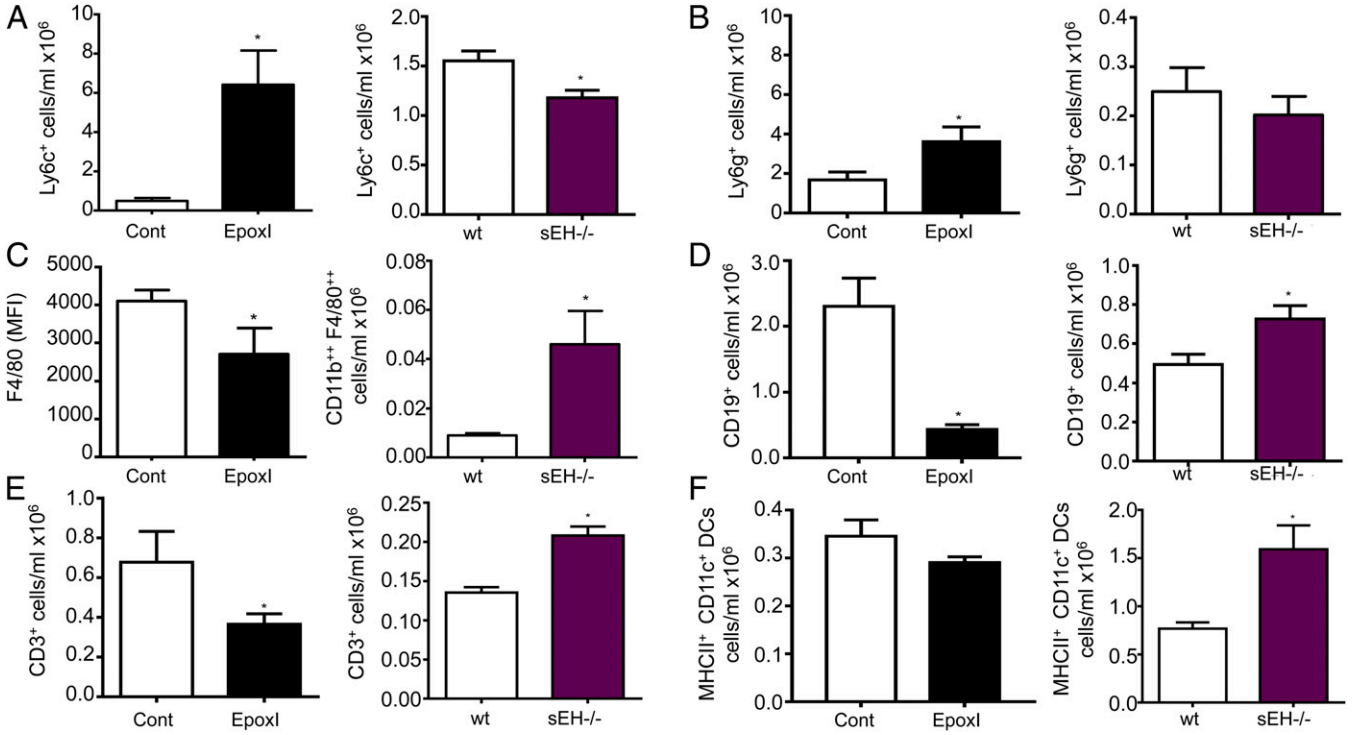

Fig. 4. Endogenous oxylipins regulate Ly6c^{hi} monocyte, F4/80 macrophage, and CD19 and CD3 cell populations during resolution. Changes in (*A*) Ly6c^{hi} cell populations, (*B*) Ly6g^{+} cell populations, (*C*) F4/80 expression (mean fluorescence intensity; MFI) (*Left*) and F4/80^{++} cell populations (*Right*), and (*D*) CD19^{+}, (*E*) CD3^{+}, and (*F*) MHCII^{+} CD11c^{+} dendritic cell populations in control and epoxI-treated mice (solid black bars; *Left*) and in wild-type (wt) and sEH$^{-/-}$ mice (*Right*). The proportion of cells in each group was determined by FACS and related back to cell numbers. Data represent the mean ± SEM from $n = 4$–5 mice per experimental group; $^{*}P < 0.05$ by two-way ANOVA or unpaired *t* test.

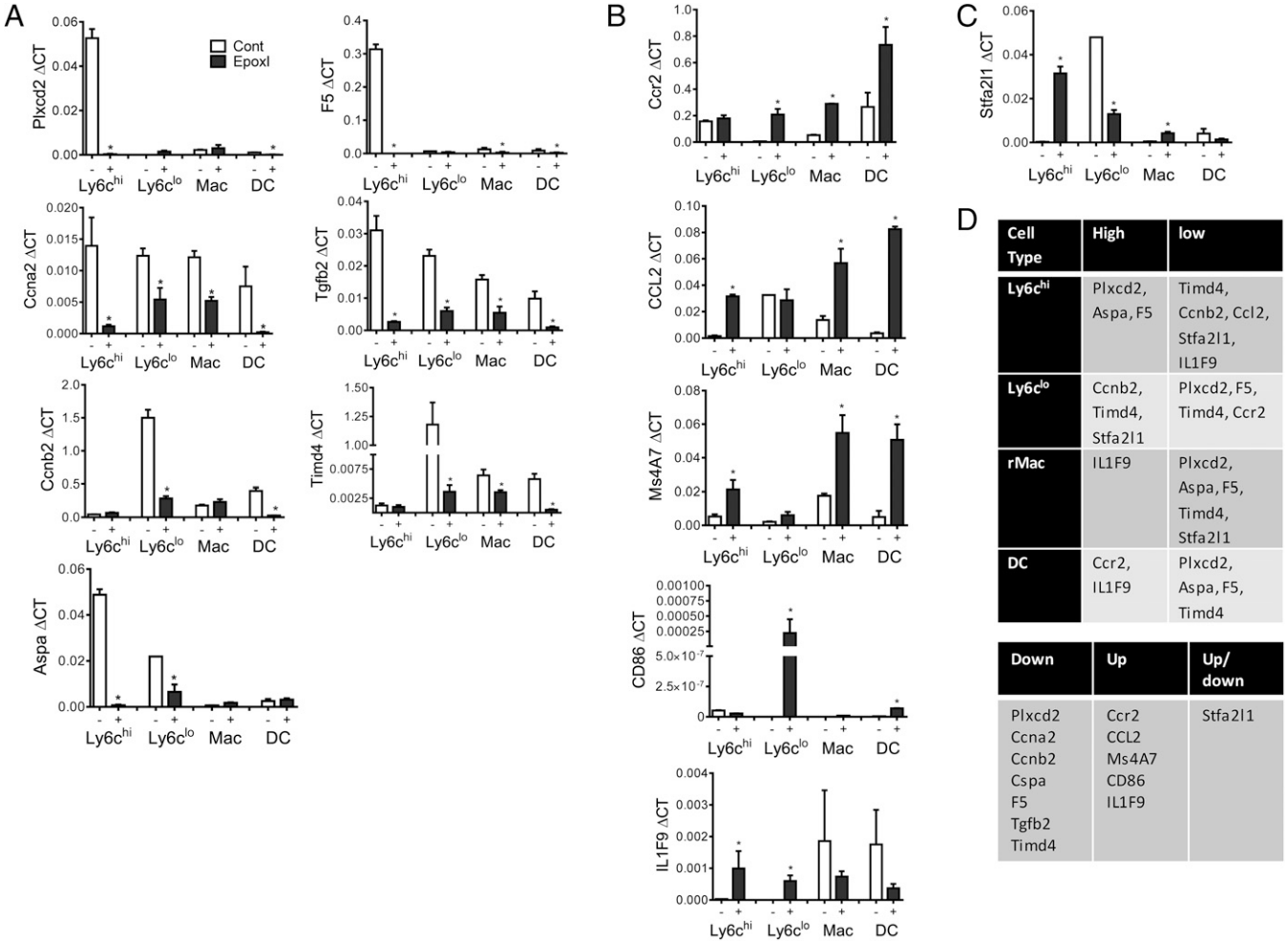

Fig. 5. Epoxygenase inhibition alters the resolution phenotype of recruited cells of the monocyte lineage. Inflammation was initiated by zymosan (1 mg, i.p.), and mice were treated with vehicle control (PBS) or epoxI (30 mg/kg, i.p.) at 24 and 36 h. Cells were collected and pooled from $n = 10$ mice and Ly6c^{hi}, Ly6c^{lo} monocytes, resident macrophages, and recruited dendritic cells were sorted on a FACSAria as detailed in *Materials and Methods*. A qRT-PCR resolution monocyte panel ($n = 3$–6) was then used to examine the phenotype of each cell type. (*A*–*C*) In the presence of epoxI, (*A*) Plxcd2, Ccna2, Ccnb2, Aspa, F5, Tgfb2, and Timd4 were found to be down-regulated in the cell types; (*B*) Ccr2, CCL2, Ms4A7, CD86, and IL1F9 were up-regulated in the cell types; and (*C*) Stfa2l1 was up- and down-regulated in a cell type-specific manner. *$P < 0.05$ by unpaired *t* test. (*D*, *Upper*) Summary of the relative basal levels of each transcript in each cell type. (*D*, *Lower*) Summary of the effect of epoxI on the different transcripts. Data represents mean ± SEM.

Human monocytes in vitro express CYP2J and CYP2C enzymes, and in particular CYP2J2 can be induced by TLR4/LPS activation (21). Rodents have expanded CYP2J and CYP2C subfamilies compared with humans, and we found the epoxygenase CYP2C44 (40) along with CYP2J6 and CYP2J9 to be present in all inflammatory exudates tested, although these findings do not rule out contributions from other local stromal or vascular cells. To test the role of epoxygenases, we used the epoxI SKF525A, which we routinely use in our in vitro assays (18–21, 41) and found to have an IC_{50} for human CYP2J2 of 1 μM (20). SKF525A is routinely used in vivo within the range of 5–50 mg/kg (42–45) and, although reported as a selective epoxygenase inhibitor, SKF525A had not been examined in the in vivo setting followed by lipidomic analysis to confirm this. We chose SKF525A as our epoxI because it is soluble in aqueous solutions (in our case PBS), and so could be tolerated far better over longer periods with limited vehicle effect. In our experiments only epoxygenase products and not prostanoids or lipoxygenase products were inhibited (by ∼50%) with 30 mg/kg of epoxI (*SI Appendix*, Table S1). epoxI was given at 24 h to look at resolution specifically, so the true levels of inhibition may be higher, as the kinetics of preformed metabolite removal in the cavity over the 24- to 48-h time period is not known.

The combination of these experiments using epoxI and sEH$^{-/-}$ mice leads to the conclusion that epoxy-oxylipins are produced during resolution by resident and incoming inflammatory cells and act in a paracrine/autocrine manner to limit Ly6c^{hi} monocyte accumulation and monocyte macrophage proinflammatory phenotype. Following epoxI treatment at the onset of resolution, the elicited cells were predominantly GR1^{+}, Ly6c^{hi}, and CX3CR1^{lo} and produced large amounts of the major Ly6c^{hi} chemokine CCL2. CCL2 was undetectable from cells elicited by zymosan alone. Interestingly, although limiting Ly6c^{hi} monocyte recruitment, epoxy-oxylipins promoted the differentiation of monocytes into F4/80^{++} macrophages. In vitro, we previously found that epoxygenases and 11,12-EET promote macrophage bacterial phagocytosis, which was associated with a reduction in CD11b (41). Similarly, here resident and monocyte-derived macrophages had reduced CD11b, along with markers of the mature macrophage phenotype and phagocytosis MerTK and Timd4, the latter being particularly important for efferocytosis (46). These markers correlated with both resident and monocyte-derived macrophages from epoxI-treated mice to have a dramatic reduction in the ability to phagocytose fresh zymosan and efferocytose apoptotic cells. LyC6^{+} monocytes had small but significantly

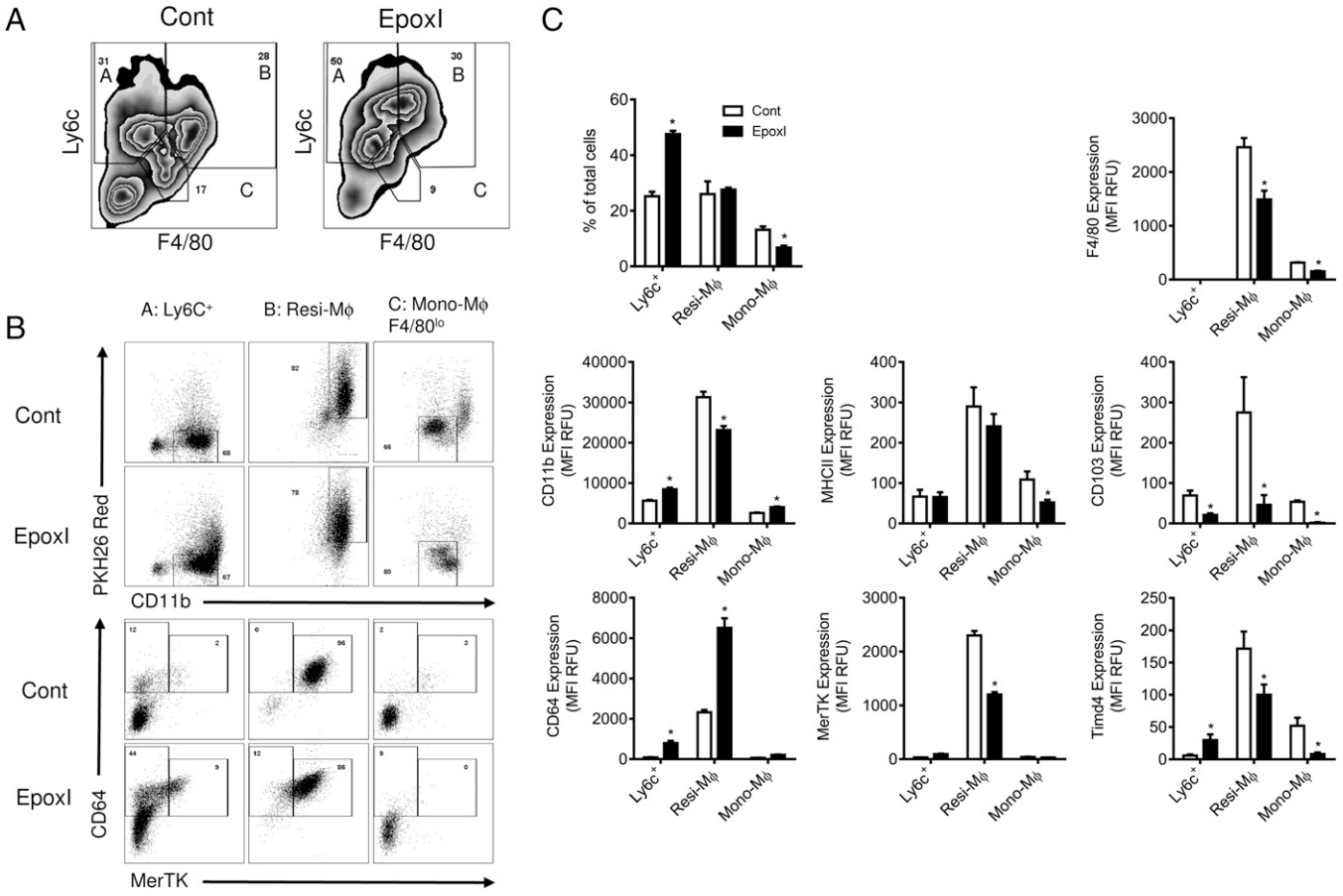

Fig. 6. Epoxygenase inhibition regulates monocyte and macrophage differentiation. Inflammation was initiated by zymosan (1 mg, i.p.), and mice were treated with vehicle control (PBS) or epoxI (30 mg/kg, i.p.) at 24 and 36 h. Cells were collected at 48 h and pooled from $n = 9$–18 mice. Ly6c$^+$ monocytes, resident macrophages, and monocyte-derived macrophages were sorted on a LSR Fortessa as detailed in *Materials and Methods*. (*A*) Representative zebra plots of the cell populations from control and epoxI-treated mice expressing Ly6c and F4/80, with labeled populations representing A: Ly6c$^+$ monocytes; B: resident macrophages (Resi-Mϕ); and C: monocyte-derived macrophages (Mono-Mϕ). (*B*) Representative dot plots of the Ly6c$^+$ monocytes, resident macrophages, and monocyte-derived macrophages from control and epoxI-treated mice expressing PKH26 and CD11b (*Upper*) and CD64 and MerTK (*Lower*). (*C*) Changes in cell numbers, and expression (MFI in relative fluorescence units; RFUs) for F4/80, CD11b, MHCII, CD103, CD64, MerTK, and Timd4 in the Ly6c$^+$, Resi-Mϕ, and Mono-Mϕ cell populations. Data are mean ± SEM from $n = 5$ mice per group; $^*P < 0.05$ by unpaired t test.

elevated CD11b and CD64 expression but reduced CD103. There was, however, little change in the ability of monocytes to phagocytose material apart from a small reduction in the ability to phagocytose zymosan ex vivo. These findings suggest epoxy-oxylipins not only limit the recruitment of proinflammatory monocytes but also limit resident and monocyte-derived macrophages' ability to clear inflammatory stimuli and apoptotic cells, further indicating a prominent role for epoxygenase pathways in promoting the resolution phenotype. Interestingly, CD19$^+$ B cells and CD3$^+$ T cells were also increased in sEH$^{-/-}$ mice (and reduced with epoxI treatment). Lymphocytes are dispensable for resolution in this model (47) but form an important part of a defense mechanism against reinfection (47). Although we have yet to investigate fully which lymphocyte subtypes are present, these findings nonetheless also provide the first evidence, to our knowledge, for a role for epoxy-oxylipins in acquired immune responses along with the roles in resolution reported here.

These findings all point to a proresolution role for epoxy-oxylipins in the proinflammatory monocyte lineage. CCL2 is known to be a central mediator of monocyte recruitment. EpoxI-treated animals produced large amounts of CCL2, as did the inflammatory cell population taken ex vivo. When we examined individual cell types, CCL2 was induced by epoxI in Ly6c^{hi} monocytes, resident macrophages, and DCs, whereas its receptor CCR2 (already expressed in Ly6c^{hi} monocytes) was also induced in Ly6c^{lo} monocytes, resident macrophages, and DCs. Using gene array analysis, we previously identified a panel of novel resolution monocyte markers (32) including the T-cell costimulator CD86, IL-1 family gene 9 (IL-1f9; IL-36), and CD20-like family member Ms4a7; these were all elevated after epoxI treatment, whereas the phosphatidylserine apoptotic cell receptors Timd4, Tgfb2, and Plxdc2 were all decreased. The roles of a number of these novel markers are still not known in resolution, but clearly epoxI-mediated changes in expression along with changes to functional clear antigens and apoptotic cells represent cells of a phenotype of highly dysregulated resolution compared with the typical resolution monocyte cell population.

Recently, the major rat monocyte epoxygenase Cyp2j4 was knocked out using a zinc finger nuclease-mediated gene-targeting approach (48). The resultant bone marrow-derived macrophages showed a highly profibrotic enriched transcriptome and activity and an induction of PPARγ (48). Although not an identical phenotype to those observed in our study, this adds to the literature showing that lipid-metabolizing CYPs have a critical role to play in the inflammatory phenotype of cells in the monocyte lineage.

EETs are too unstable to give as a bolus dose in vivo. Therefore, we tested the direct antiinflammatory effects of monocyte

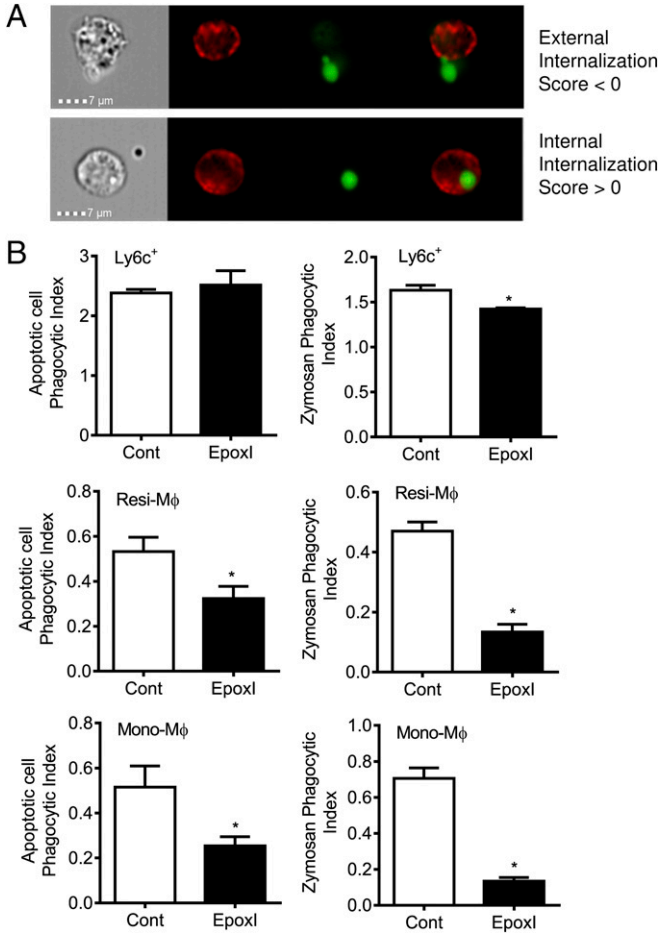

Fig. 7. Epoxygenase inhibition reduces the phagocytic activity of monocyte-derived and resident macrophages. Inflammation was initiated by zymosan (1 mg, i.p.), and mice were treated with vehicle control (PBS) or epoxI (30 mg/kg, i.p.) at 24 and 36 h. Cells were collected at 48 h and pooled from $n = 9$–18 mice. Ly6c$^+$ monocytes, resident macrophages, and monocyte-derived macrophages were sorted on a FACSAria as detailed in *Materials and Methods*. Sorted cells were then tested for their ability to phagocytose CFSE-labeled apoptotic cells (thymocytes) or FITC-labeled zymosan BioParticles over 30 min. Cells were gated using F4/80 or Ly6c on an ImageStream^X Mark II. (*A*) ImageStream analysis differentiates cells that phagocytose apoptotic cells or zymosan (internalization score >0) from those where particles or bodies just stick to the cell (internalization score 0). (*B*) Ex vivo phagocytosis of apoptotic cells (*Left*) and zymosan (*Right*) by Ly6c$^+$, Resi-Mϕ, and Mono-Mϕ cell populations from control and epoxI-treated mice. Data are mean ± SEM from $n = 3$–6 pooled samples; *$P < 0.05$ by unpaired t test.

epoxygenases ex vivo. 14,15-EET, the most abundant epoxygenase-derived EET produced in the peritoneal cavity, inhibited inflammatory cell activation of iNOS and IL-12 ex vivo from zymosan-treated mice. Interestingly, the linoleic acid epoxygenase products 9,10-EpOME and 12,13-EpOME are highly expressed in the naive cavity. The levels of 9,10-EpOME drop rapidly during acute inflammation, whereas 12,13-EPOME changes very little. Our results therefore indicate a change in substrate availability or utilization by epoxygenases from linoleic acid in homeostasis to arachidonic acid during inflammation. Linoleic acid epoxygenase products are generally considered proinflammatory and cytotoxic (49–51), whereas EETs are antiinflammatory. How this concept fits with the high levels of EpOMEs and DiHOMEs in the naive peritoneal cavity is intriguing, and points to novel as yet undiscovered homeostatic roles for these mediators in the peritoneal cavity. In contrast to EETs, which inhibited Ccl2 in monocytes ex vivo, 9,10-EpOME induced CCL2 (*SI Appendix*, Fig. S6), suggesting that substrate utilization may be an additional layer of control over the effects of the epoxygenase pathway. High EpOME levels have been found in acute respiratory distress syndrome and in patients with extensive burns (52); however, the roles of EpOMEs in inflammation and homeostasis remain relatively poorly understood, although it is becoming clearer that many of the cytotoxic effects attributed to them are in fact due to their sEH metabolite DiHOMEs (51, 53–55). Clearly more work is needed to elucidate the function of these linoleic acid epoxygenase products in homeostasis as well as pathophysiology. At this stage, we cannot rule out potential effects of DHA and EPA CYP metabolites but, because we saw no significant change in their production with epoxI, we have here focused on arachidonic acid and linoleic acid epoxy-oxylipins.

The levels of 12/15-lipoxygenase, 9,10- and 12,13-HODE, and 12- and 15-HETE were high in the peritoneal cavity, consistent with the report that high levels of resident macrophages contain 12/15-lipoxygenase in mice, where it regulates immune function. Our findings are also consistent with this report in that 12/15-lipoxygenase products drop during acute inflammation (in our experiments only 9-HODE returned to basal levels by 48 h), and support the findings that 12/15-lipoxygenase has an important homeostatic immunoregulatory role in this model (56).

In conclusion, we have characterized the epoxygenase pathways during a model of inflammatory resolution. Inhibition of epoxygenases during this resolution phase established a proinflammatory environment that allows the recruitment of proinflammatory Ly6c monocytes and macrophages with a highly dysregulated resolution/clearance phenotype, whereas mice where epoxygenase products are elevated have an enhanced resolution phenotype. An efficient and active epoxygenase pathway therefore joins similar proresolution lipid pathways such as the resolvins in the requirement for effective inflammatory resolution. Drugs that elevate epoxygenase products or epoxygenase product mimetics therefore represent novel resolution targets for chronic inflammatory disorders.

Materials and Methods

Materials. EETs were purchased from Cayman Chemical (Cambridge Bioscience). SKF525A was purchased from Biomol (Affiniti Research Products). Cytokines and the MCP-1 ELISA were obtained from R&D Systems. The TNFα ELISA was from eBioscience. Unless otherwise stated, all other reagents were purchased from Sigma-Aldrich.

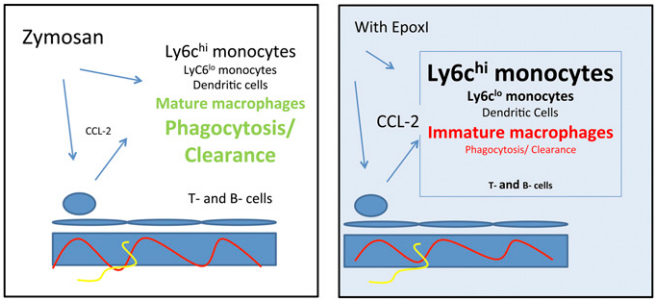

Fig. 8. Role of epoxy-oxylipins in the resolution phenotype. The resolution of zymosan-initiated inflammation involves monocytes, dendritic cells, and T- and B-cell recruitment and the differentiation of monocytes into resolution-type macrophages. In this process (as revealed by epoxI treatment and sEH$^{-/-}$ mice), epoxy-oxylipins, most likely EETs, limit Ccl2 and Ccr2 expression, Ly6c^{hi} monocyte accumulation, and T- and B-cell recruitment and encourage the formation of mature phagocytic resolution macrophages. Ly6c^{hi} monocytes, Ly6c^{lo} monocytes, dendritic cells, and monocyte-derived and resident macrophages are all activated in the presence of epoxI.

Animal Models. All experiments were completed with adult male mice on a pure C57BL/6 background. Control C57BL6/J mice were bred under standard conditions and maintained in a 12-h light/dark cycle at 22 °C and given food and tap water ad libitum in accordance with UK Home Office regulations. Mice with targeted disruption of sEH were rederived and backcrossed onto a C57BL/6 genetic background for more than 10 generations, as previously described (31, 52, 57). $sEH^{-/-}$ mice have significantly higher circulating epoxide:diol ratios compared with wild-type littermates, consistent with functional sEH disruption (57, 58). Peritonitis was induced by intraperitoneal injection of 1 mg type A zymosan in 0.5 mL PBS after 30 s of sonication on ice. Inflammatory cells were retrieved at the time points described in *Results* by injecting 2 mL sterile cell dissociation buffer (Life Technologies). Cells were counted by hemocytometer, and exudates were stored at −80 °C until further analysis. In some experiments, the epoxI SKF525A (30 mg/kg) or sterile PBS (0.5 mL vehicle) was given intraperitoneally 24 h after zymosan injection and at 12-h time points thereafter up until 96 h. These studies received institutional review board approval for the use of mice from the UK Home Office.

Lipid Analysis. Eicosanoids and other fatty acid metabolites were extracted from inflammatory exudates by solid-phase extraction and eluted in ethyl acetate, essentially as described (22); see *SI Appendix, Supplemental Methods* for extended methods. Lipids were separated by reverse-phase HPLC on 2-μm, 150-mm, 5-μm Luna C18 columns (Phenomenex) and quantified using an MDS Sciex API 3000 triple quadrupole mass spectrometer (Applied Biosystems) with negative-mode electrospray ionization and multiple-reaction monitoring, as described (12). The relative response ratios of each analyte were used to calculate concentrations while correcting for surrogate losses via quantification relative to internal standards. The sensitivity of analytes ranged from 0.25 to 25 pg per sample.

FACS Analysis and Cell Sorting. Flow cytometry and cell sorting were done on LSRII/LSRFortessa and FACSAria (BD Biosciences), respectively. Cells were incubated with Fc-blocker (AbD Serotec) and fluorescently labeled antibodies. Data were analyzed with FlowJo 7.0.1 software (Tree Star) using fluorescence minus one controls as the reference for setting gates. Antibodies were obtained from BD Biosciences unless stated [F4/80 (eBioscience), CD11b, CD11c, Ly6c, Ly6g, Gr1, CD3 (BioLegend), CD19, CD4, CD8, CD62l, CD44, MerTK, CD64, CD103, Timd4, and MHCII (eBioscience)] (59). To identify resident macrophages (59), PKH26-PCL^{red} (0.35 mL, 500 nM; Sigma) was injected intraperitoneally 2 h before injection of zymosan. In cell-sorting experiments, monocytes and macrophages were sorted from a population of $CD19^-$ and $CD3^-$ cells as either $Ly6c^+F4/80^+$ or $Ly6c^-F4/80^+$; see *SI Appendix*, Fig. S7 for the gating strategy. For identification of $Ly6g^+$ neutrophils and $Ly6c^+$ monocytes, a combination of Gr1 and anti-Ly6c or anti-Ly6g was also used. Resident macrophages were characterized as PKH^{red++}, DCs were characterized as PKH^{red-} $MHCII^+$ and $CD11c^+$, $Ly6c^{hi}$ monocytes were characterized as $LyC6^{hi}$ PKH^{red+} $MHCII^-$, and $Ly6c^{lo}$ monocytes were characterized as $Ly6c^{lo}$ PKH^{red-}, as previously described (59).

Cell Culture ex Vivo. The peritoneal lavage was treated with ACK lysis buffer (Thermo Fisher Scientific) to remove erythrocytes. After being washed, peritoneal cells were suspended in DMEM supplemented with 10% (vol/vol) FBS and 50 μg/mL penicillin/streptomycin. The cells (2×10^6) were seeded on a 12-well plate and left to adhere for 45 min in a humidified CO_2 incubator. Nonadherent cells were removed by three washes with DMEM. Remaining adherent cells ($\sim 1 \times 10^6$) were incubated in 0.5 mL DMEM in the presence or absence of 1 μM 14,15-EET or 11,12-EET or vehicle (0.3% EtOH). After 6 h, cell-free supernatants were removed and cells were lysed using TRIzol (Invitrogen) for subsequent RNA extraction.

RT-PCR. Cells analyzed by qRT-PCR were lysed and RNA was isolated using TRIzol. The panel of resolution markers Timd4 (T-cell Ig and mucin domain-containing 4), Tgfb2, Plxdc2 (plexin domain-containing protein 2), IL1f9 (interleukin 1 family member 9), CD86, Ms4a7 (membrane-spanning four domains, subfamily A, member 7), Ccna2 (cyclin A2), Ccnb2 (cyclin B2), F5 (coagulation factor V), Aspa (aspartoacylase), and Stfa2l1 (stefin A2-like 1) were measured by qRT-PCR as previously described (32). Resolution monocytes were previously found to be enriched for cell-cycle/proliferation genes as well as for Timd4 and Tgfb2, key systems in the termination of leukocyte trafficking and clearance of inflammatory cells. Ly6c, CX3CR1, Ccl2, Ccr2, Cyp2j5, Cyp2j6, Cyp2j9, Cyp2j13, Cyp2c29, Cyp2c38, Cyp2c39, Cyp2c44, Cyp2c50, Cyp2c54, Cyp2c55, Cyp2a1, Cyp2u1, Cyp2s1, and β-actin were measured by RT-PCR. Primers are detailed in *SI Appendix, Supplemental Methods*.

Efferocytosis and Phagocytosis Assays. Apoptotic cells (thymocytes) were produced from the thymuses of three naive control C57/BL mice killed 24 h presort. Harvested thymuses were passed through a 70-μm mesh and then lysed with ACK buffer for 3 min. Cells were washed twice with RPMI 1640 with 100 U/mL penicillin/streptomycin. To induce apoptosis, thymocytes were resuspended in media at 2×10^6 cells per mL and exposed to UV radiation (~300 nm) for 20 min followed by incubation for 16–24 h at 37 °C with 5% CO_2 using a Syngene UV transilluminator. Cells were then washed with PBS and labeled with 2 μM carboxyfluorescein succinimidyl ester (CFSE) according to the manufacturer's protocol (Life Technologies; CellTrace CFSE).

Sorted cell populations, 2×10^5 cells per well, were plated in 24-well plates and resuspended in X-VIVO 15 media (Lonza) containing 10% FBS, 2 mM L-glutamine, and 1× pen-step. Cells were then challenged with apoptotic thymocytes (5:1 monocyte/macrophage) or FITC-labeled zymosan A BioParticles (Thermo Fisher Scientific) (10 particles:1 monocyte/macrophage) for 30 min at 37 °C and 5% CO_2. Phagocytosis was stopped by placing the plates on ice. Cells were detached using 10 mM EDTA containing 4 mg/mL lignocaine for 20 min. Cells were stained for either F4/80 or Ly6c (allophycocyanin) for 20 min (see *FACS Analysis and Cell Sorting*, above) before being washed twice with PBS containing 2 mM EDTA. Cells were then fixed using 4% paraformaldehyde in PBS and analyzed on an ImageStream^X Mark II (Merck Millipore) (*SI Appendix*, Fig. S8).

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