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Resolution of Acute Inflammation and the Role of Resolvins in Immunity, Thrombosis and Vascular Biology

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Abstract

Acute inflammation is a host-protective response that is mounted in response to tissue injury and infection. Initiated and perpetuated by exogenous and endogenous mediators, acute inflammation must be resolved for tissue repair to proceed and for homeostasis to be restored. Resolution of inflammation is an actively regulated process governed by an array of mediators as diverse as those that initiate inflammation. Among these, resolvins have emerged as a genus of evolutionarily conserved pro-resolving mediators that act on specific cellular receptors to regulate leukocyte trafficking and blunt production of inflammatory mediators, while promoting clearance of dead cells and tissue repair. Given that chronic unresolved inflammation is emerging as a central causative factor in the development of cardiovascular diseases, an understanding of the endogenous processes that govern normal resolution of acute inflammation is critical for determining why sterile maladaptive cardiovascular inflammation perpetuates. Here, we provide an overview of the process of resolution with a focus on the enzymatic biosynthesis and receptordependent actions of resolvins and related pro-resolving mediators in immunity, thrombosis and vascular biology. We discuss how nutritional and current therapeutic approaches modulate resolution and propose that harnessing resolution concepts could potentially lead to the development of new approaches for treating chronic cardiovascular inflammation in a manner that is not host-disruptive.

Keywords

Resolution of inflammation; Lipid mediators; Omega-3 fatty acids

Subject terms

Inflammation; Vascular Biology; Physiology

Disclosures None

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Inflammation and cardiovascular disease

Persistent unresolved inflammation has long been recognized to play a fundamental role in the development of chronic diseases including arthritis, autoimmune diseases, and asthma. In recent decades, it has emerged that maladaptive inflammation is causally involved in the development of cardiovascular disease (CVD) and other metabolic disorders including diabetes and obesity. Indeed, inflammation is a central feature of virtually all pathologies afflicting both the heart and the vasculature, including chronic heart failure (HF), atherosclerosis, myocardial infarction (MI), and stroke¹, as discussed in greater depth in the accompanying reviews of this series. Moreover, mounting evidence indicates that the presence of other chronic inflammatory diseases (e.g., periodontitis, rheumatoid arthritis) can exacerbate $CVD¹$. Atherosclerosis is perhaps the CVD that has been most closely linked with chronic unresolved inflammation^{2, 3}. While the development of statins revolutionized the treatment of atherosclerosis by lowering low-density lipoprotein (LDL) cholesterol, the burden of ischemic cardiovascular conditions has continued to rise globally and CVD remains the leading cause of death among Americans⁴. Furthermore, intolerance and adverse effects of statins in a growing number of patients has emphasized the need for novel therapies^{2, 4}. Many new strategies, therefore, have looked beyond lipid lowering alone and have focused instead on halting aberrant inflammatory signaling⁵. Several clinical trials are currently underway testing the efficacy of anti-inflammatory drugs in the context of CVD. One large trial, the Cardiovascular Inflammation Reduction Trial (CIRT), using low dose methotrexate, is currently recruiting patients and is specifically designed to assess whether inhibition of inflammation will lower vascular events rates^{6, 7}. Another trial, the Canakinumab Anti-inflammatory Thrombosis Outcomes Study (CANTOS), has enrolled over 10,000 patients with stable coronary artery disease (CAD) and a history of MI. Preliminary results show treatment with the IL-1β receptor antagonist decreased measures of inflammation without affecting lipids^{4, 8}. These clinical trials will provide valuable insights into the role of inflammation in CVD and may herald a new direction in treatment strategies. While invasive procedures will remain indispensable for the most severe CVD cases, innovative therapies targeting the body's innate immune system may indeed be a promising alternative for many patients. Inherent in the discovery and development of such new, effective and less invasive therapies is a more thorough delineation of the endogenous mechanisms that govern the natural resolution of acute inflammation. A novel genus of "pro-resolving mediators" potently resolve inflammation, but also enhance certain aspects of host defense, potentially overcoming limitations to traditional therapeutic anti-inflammatory agents that can be immunosuppressive⁹. This review focuses on the role of these endogenous mediators in promoting the resolution of acute inflammation, highlighting their biosynthesis and cellular targets, and discusses how defective resolution may contribute to CVD.

Acute inflammation and its resolution

In response to injury or infection, the protective program of acute inflammation and its complete and timely resolution are critical for the restoration of tissue homeostasis¹⁰. This highly coordinated and synergistic program combines the distinct actions of multiple cell types to achieve pathogen eradication and subsequent tissue repair. The acute inflammatory response can be divided into two general phases: initiation and resolution (Fig 1). Initiation

is marked by tissue edema resulting from increased blood flow and permeability of the microvasculature; processes that are mediated in part by lipid mediators (e.g., cysteinyl leukotrienes and prostaglandins) and other vasoactive products (e.g., histamine, bradykinin). Subsequently, polymorphonuclear neutrophils (PMN) migrate to the area to defend against microbial invasion. Drawn to the site of injury by exuded chemical signals including proinflammatory lipid mediators (e.g., leukotriene B_4 ; LTB₄) and chemokines, PMN traverse the vasculature through precise interactions with endothelial adhesion receptors and subsequently engulf and degrade pathogens within phagolysosomes^{11–13}. The resolution phase is already being enacted at this early point as the influx of PMN is halted at a level appropriate for the insult and is accompanied by their timely apoptosis¹⁴. Monocytes subsequently infiltrate the tissue where they differentiate into macrophages that avidly respond to pathogen-associated molecular patterns (PAMPs) and damage-associated molecular patterns (DAMPs) present in the injured tissue. Importantly, macrophages are highly responsive to so-called find-me and eat-me signals (e.g., nucleotides, externalized phosphatidylserine) released or presented by apoptotic cells such as $PMN^{10, 15}$. Uptake of apoptotic cells by macrophages (i.e., efferocytosis) is an anti-inflammatory process associated with decreased production of inflammatory mediators, thus coupling the initiation of inflammation to its ultimate resolution (Fig. 1). The timely clearance of microbes and apoptotic cells is required to prevent bystander tissue damage and to set the stage for tissue repair and regeneration, allowing for the return to homeostasis^{9, 15}. Indeed, active clearance of apoptotic cells is a key defining feature of resolution, as failed clearance can lead to cellular necrosis and exacerbated inflammation beyond the initial insult, impeding tissue repair. Macrophages persist in injured tissues longer than short-lived PMN, during which time they are continuously reprogrammed in response to local cues to facilitate tissue repair and orchestrate the delicate balance of fibrosis^{16–18}. Like innate immune cells, adaptive immune cells also play critical roles in the host response to infection, resolution of inflammation and in tissue repair^{19, 20}. Their accumulation defines the "post-resolution" phase of the inflammatory response and assures a more rapid response to subsequent exposure to the same antigens¹⁹. Interruption of this process at any point (e.g., prolonged leukocyte recruitment and survival, impairments in apoptotic cell removal, alterations in macrophage phenotype switching) could potentially lead to chronic inflammation with resultant tissue damage, excessive fibrosis and loss of function, as is seen in many CVDs like atherosclerosis and $HF^{10, 17, 21-23}$.

By its nature, the acute inflammatory response is self-limiting in part because of inherent negative feedback regulation of inflammatory signaling pathways (e.g., transcriptional repressors, endogenous receptor antagonists) when the trigger has been eliminated. However, it has recently become evident that active resolution of inflammation involves the biosynthesis of pro-resolving mediators that, as a genus, are just as diverse as the initiators of inflammation^{24–33}. Thus, critical to determining the fate of an inflammatory response is the balance of pro-inflammatory and pro-resolving mediators that are produced in the exudate in a temporal manner. Traditionally, it has been held that an excess production of pro-inflammatory mediators underlies chronic inflammation³⁴, however, mounting evidence supports the view that disruptions in endogenous pro-resolving circuits may be an equally important mechanism^{10, 34, 35}. These pro-resolving mediators actively terminate the

production of pro-inflammatory mediators, but also directly stimulate macrophage phagocytosis of both apoptotic cells and bacteria, promote egress of phagocytes from sites of inflammation, regulate PMN apoptosis, promote chemokine scavenging, and stimulate tissue repair and regeneration^{9, 36–41}. These agonist-based actions distinguish pro-resolving mediators from intrinsic negative feedback pathways and other antagonists that terminate inflammatory signaling pathways. Systems-based approaches have played a crucial role in the identification of the principal mediators of resolution⁹. As a result, a complex and everexpanding network of interrelated mediators and the cellular targets and pathways that they engage has been assembled. The discovery of these novel bioactive pro-resolving mediators represents a paradigm shift in our understanding of the dynamic regulation of acute inflammation and has led to a new era of "resolution physiology"⁹ .

Pro-resolving mediators

The discovery of bioactive mediators with potent inflammation-resolving actions in experimental models of acute inflammation was a seminal development that provided compelling evidence that resolution is an active process rather than a passive one as traditionally thought. Self-resolving inflammatory exudates were shown to contain structurally unique families of signaling molecules that are temporally produced and when added back in experimental models of acute inflammation, potently enhance the resolution phase. Like the initial phases of inflammation, enzymatically produced lipid mediators derived from polyunsaturated fatty acids (PUFA) play a central role in resolution, due in part to their rapid production by distinct immune cell subsets. During the transition from inflammation to resolution, lipid mediator class switching takes place in which initial formation of pro-inflammatory mediators induce the production of enzymes that enable the reprogramming of leukocytes to generate specialized pro-resolving lipid mediators (SPM) from the same PUFA precursors (Fig. $1)^{42}$. For instance, 5-lipoxygenase (5-LOX) is required for the generation of potent leukocyte chemoattractant, LTB₄ from omega-6 PUFA arachidonic acid (AA). When 12-LOX is also expressed (e.g., during platelet:leukocyte interactions), the epoxide intermediate in $LTB₄$ biosynthesis (i.e., $LTA₄$) can be converted into the family of lipoxins. Lipoxins can also be produced with cooperation between 5-LOX and 15-LOX, the latter of which can be induced in PMN by prostaglandin E_2 (PGE₂), or in macrophages by Th2 cytokines (e.g., IL-4). Lipoxin A_4 (LXA₄), was the first lipid mediator shown to possess both anti-inflammatory and pro-resolving actions⁴³ and, along with subsequently discovered members of the lipoxin family (i.e., LXB4), has been the subject of extensive and rigorous study. In addition to omega-6 PUFA serving as precursors to lipoxins, omega-3 PUFAs emerged as an additional source of novel SPM. Indeed, after the initial discovery of LXA4, the enzymatic conversion of the omega-3 PUFAs eicosapentaenoic acid (EPA) and docosahexaenoic acid (DHA) to novel families of SPM – resolvins, protectins and maresins – was identified $44-47$. This mechanism could have evolved to ensure adequate production of SPM in organisms with different PUFA in their diets. Importantly, SPM are evolutionarily conserved from flatworms (e.g., *Planaria*) to humans and several studies have documented endogenous formation of these mediators in healthy individuals, with lower levels observed in patients with Alzheimer's disease, peripheral artery disease and asthma (for review, see 25). In addition to lipid mediators of resolution, other important mediators of

the resolution program have emerged and include proteins and bioactive peptides (e.g., annexin A1, galectin 1, melanocortins) and gases (nitric oxide, hydrogen sulfide, carbon monoxide). Readers are referred to recent reviews covering each of these mediators in δ detail^{24–33}. Here, we focus primarily on the biosynthesis and biological actions of resolvins in the inflammation-resolution program.

Biosynthesis of Resolvins

The use of liquid chromatography-tandem mass spectrometry (LC-MS/MS)-based approaches were critical in the identification and structural elucidation of resolvins and related SPM⁹. Applying these methods to generate a lipid mediator profile of self-resolving inflammatory exudates in mice yielded the first evidence of a novel bioactive mediator generated from EPA during resolution⁴⁴. Later named resolvin E1 (RvE1), its biosynthetic pathway proceeds via conversion of EPA to 18R-hydroperoxyeicosapentaenoic acid (18R-H(p)EPE) by acetylated cyclooxygenase-2 (COX-2) or cytochrome P_{450} enzymes, which is subsequently converted to RvE1 by 5-LOX (Fig. $2)^{44,48}$. The epoxide intermediate formed during RvE1 biosynthesis is hydrolyzed by leukotriene A_4 hydrolase (LTA₄H), the same enzyme that produces pro-inflammatory $LTB₄⁴⁹$. This mechanism provides an additional point of control in the balance between pro-inflammatory mediators and SPM. The formation of RvE1 precursor, 18-H(p)EPE, was identified in hypoxic endothelial cells and can be transformed to RvE1 through leukocyte:endothelial interactions. The complete structural elucidation of RvE1 was carried out via total organic synthesis and matching with endogenous material, which enabled assessment of the individual bioactions of RvE1. These actions include the reduction in PMN infiltration to sites of acute inflammation, blunting of inflammatory cytokine production, and the enhancement of macrophage efferocytosis $44, 48$. Additional members of the EPA-derived (E-series) resolvins have since been characterized. Similar to RvE1, RvE2 is generated from EPA via the intermediate 18R-HEPE and its subsequent 5-LOX-dependent conversion to a $5S$ -hydroperoxide^{50, 51}. However, instead of undergoing an epoxidation step, the 5S-hydroperoxide is directly converted to RvE2 (Fig. 2). More recently, a third member, RvE3, was also identified and its structure determined⁵². Unlike RvE1 and RvE2, the formation of this product does not require 5-LOX.

In addition to EPA, the other prominent omega-3 PUFA in humans and marine oils is DHA. This fatty acid serves as the precursor to the D-series resolvins, as well as the protectins and maresins^{44–48}. In the presence of 15-LOX, DHA is converted to a 17-hydroperoxide intermediate (17-H(p)DHA) which can subsequently be converted to D-series resolvins via 5-LOX. Specifically, the hydroperoxide formed via 5-LOX can undergo epoxidation to generate a 7S,8S epoxide that is then enzymatically hydrolyzed to produce resolvin D1 (RvD1) or resolvin D2 (RvD2; see Fig. $2^{40, 53}$. As with the E-series resolvins, these steps can take place within a single cell (e.g., PMN or macrophages), or can occur during cell:cell interactions (e.g., leukocyte:endothelial, PMN:macrophage). Additional structurally unique D-series resolvins, including RvD3-6, have also been identified and the complete structures and stereochemical assignments of RvD3 and RvD4 have recently been reported^{54, 55}. In murine models of acute inflammation, distinct temporal patterns of resolvin biosynthesis have been observed, highlighting additional control points in their biosynthesis (e.g., potentially novel regulation by epoxide hydrolases)⁵⁴. Interestingly, within a single cell type,

lipid mediator class switching can occur and pro-resolving lipid mediators can regulate each other. For instance, RvD1 reduces nuclear localization of 5-LOX in macrophages and diverts AA metabolism from pro-inflammatory lipid mediator, $LTB₄$, to SPM, $LXA₄$. Mechanistically, RvD1 inhibits the activation of calcium-calmodulin-dependent kinase II (CaMKII) and downstream activation of p38 and a mitogen-activated kinase (MK2), which results in reduced phosphorylation of 5 -LOX 5 ⁶. Thus, SPM biosynthesis can be amplified by positive feed-forward mechanisms, helping to tip the balance from inflammation to resolution. Indeed, in several distinct models of acute inflammation and infection, administration of resolvins blunts production of pro-inflammatory eicosanoids and increases formation of other pro-resolving mediators⁹. It is important to note that, like most autacoids, resolvins are rapidly inactivated after they have elicited their biological actions in resolution. For instance, RvE1 is a substrate for 15-hydroxyprostaglandin dehydrogenase (PGDH), generating the inactive product, 18oxo-RvE1. This suggests that accelerated inactivation could potentially underlie defective resolution in some cases and as a way to circumvent metabolic inactivation from a therapeutic standpoint, a variety of stable resolvin analogs have been formulated and shown to have enhanced biological activity in $vi\sigma^{25}$.

Recently, a new family of mediators was identified wherein the epoxide intermediate in RvD1 and RvD2 biosynthesis can be enzymatically conjugated to glutathione to generate sulfido-conjugated resolvins³⁸. They contain glutathione at the C8 position and can be subsequently converted to cysteinylglycinyl conjugates by γ -glutamyl transferase. These novel mediators were identified in infectious inflammatory exudates, human spleens and in human leukocytes (e.g., PMN and macrophages), as well as apoptotic PMN. Like other SPM, these resolvin-conjugates enhance bacterial clearance in vivo and macrophage phagocytosis. Interestingly, they have distinct roles in promoting tissue regeneration in Planaria. Given these new roles, they were named "resolvin-conjugates in tissue regeneration; RCTRs". It is noteworthy that similar biosynthetic routes were also identified for maresin and protectin families of SPM^{38, 57}.

Pro-resolving receptors

Like other classic lipid mediators and chemokines, SPM elicit their biological actions through activation of specific G-protein coupled receptors (GPCRs) in a stereoselective manner (Fig. 2). The pro-resolving and anti-inflammatory actions of $LXA₄$ and $RvD1$ are mediated via specific signaling through ALX (also termed formyl peptide receptor 2; FPR2) and previous orphan, GPR32 (also termed D-resolvin receptor 1; DRV1)^{25, 58}. Given that LXA4 and RvD1 are structurally similar but generated from different PUFA precursors, that they activate the same receptors and promote reciprocal biosynthesis of each other provides additional evidence that pro-resolving pathways are intimately linked. Specific binding to these receptors was unequivocally demonstrated using radiolabeled material and both homoand heteroligand competition. Moreover, their pro-resolving actions were validated in mouse models of acute inflammation with targeted overexpression or genetic deletion of the ALX/ $Fpr2^{59-61}$. We note that there is no murine homolog of human GPR32 identified to date. The actions of RvD1 in counter-regulating PMN recruitment during acute peritonitis are ablated in $ALX/Fpr2$ -deficient mice⁵⁹. In human macrophages, RvD1 enhances phagocytosis of both apoptotic PMN and opsonized zymosan, actions that are potentiated by overexpression

of ALX/FPR2 and GPR32, and attenuated with their knockdown⁵⁸. Recently, it has been shown that RvD3 and RvD5 also activate human GPR32^{37, 54}. Given that these mediators are generated at distinct times during resolution of inflammation54, this mechanism may serve to preserve GPR32 ligands during different stages of resolution. Interestingly, ALX/ FPR2 is also activated by other pro-resolving mediators, such as annexin A1, as well as proinflammatory mediators (e.g., serum amyloid A; SAA). Depending on the stimulus, signaling through this receptor can elicit either pro-inflammatory or pro-resolving effects. Emerging evidence indicates that dimerization underlies in part these distinct and unconventional roles⁶². Specifically, annexin A1 and $LXA₄$ induce ALX homodimer formation and this leads to increased production of IL-10 in monocytes. In contrast, proinflammatory agonists, such as SAA, can block this homodimerization. In PMN, annexin A1, an annexin A1 peptide mimetic $(Ac_{2,26})$, and LXA₄ can induce formation of ALX heterodimers with FPR1. These heterodimers elicit distinct signaling pathways (JNK/ caspase-3) that override pro-survival signals in PMN, such as SAA, through the same receptor^{62, 63}. As noted, this is important because timely apoptosis and clearance of PMN is required for resolution of inflammation and prolonged PMN survival can cause excessive tissue damage.

A distinct GPCR denoted chemoattractant receptor 23 (ChemR23; also termed E-resolvin receptor, ERV), which shares 36% sequence identity with ALX/FPR2, is activated by RvE1 in a stereoselective manner⁴⁸. The anti-inflammatory signaling properties of ChemR23 were also demonstrated in studies interrogating a different agonist, chemerin peptide, and in mice with genetic deletion of the receptor⁶⁴. Subsequent investigations further substantiated ChemR23-mediated signaling by RvE1, including stimulation of macrophage phagocytosis via the phosphatidylinositol 3-kinase (PI3K)/Akt pathway and activation of the translational regulator, ribosomal protein S665. Interestingly, not all of RvE1's actions can be ascribed to ChemR23. Indeed, RvE1 potently blocks PMN migration in vitro yet these cells do not express ChemR23⁴⁸. Additional studies determined that RvE1 binding to human PMN is displaced by $LTB₄$ and an antagonist to the $LTB₄$ receptor (BLT1), but not by chemerin peptide, indicating that BLT1 is a target for RvE1 on PMN66. This study also demonstrated that the actions of RvE1 on PMN may be due, in part, to an attenuation of pro-inflammatory $LTB₄$ signaling, which was subsequently confirmed in *Blt1*-deficient mice⁶⁶. Like RvE1, RvE2 also binds BLT1 and blocks $LTB₄$ -stimulated actin polymerization in PMN^{50, 51}. Importantly, although many of the downstream actions of RvD1 are similar to those of RvE1, it does not bind ChemR23 or BLT158.

Using an unbiased β-arrestin based screening approach, a recent study identified that RvD2 activates a distinct GPCR denoted GPR18 (also termed D-resolvin receptor 2; DRV2), which among innate immune cells, is expressed on human PMN, monocytes and macrophages⁶⁷. Specific binding to GPR18 was confirmed using radiolabeled RvD2 and binding competition studies. Similar to the actions of RvD1, RvD2 increases phagocytosis of apoptotic PMN by macrophages, which is enhanced with GPR18 overexpression and decreased with GPR18 knockdown. Similarly, the potent actions of RvD2 in limiting PMN infiltration and enhancing clearance of E. coli and S. aureus in vivo are abolished in $Gpr18$ -deficient mice⁶⁷. Taken together, it is clear that the actions of resolvins have significant overlap yet remain specific and are mediated by distinct receptors. The multiplicity of cellular sites of action of

resolvins reinforces the concept that these mediators are vital in orchestrating the complex, coordinated response that governs the active process of inflammation-resolution. The identification of these pro-resolving receptors has, and will continue to, illuminate the mechanisms governing resolution, including the specific signaling pathways and transcriptional programs engaged by pro-resolving mediators.

Increasing pro-resolving lipid mediators with nutritional and pharmacologic approaches

Given the potent bioactions of resolvins and related SPM during acute inflammation, there has been significant attention and effort dedicated to developing strategies for increasing their production endogenously. Perhaps the most obvious approach would be to increase the substrate. As discussed, the omega-3 PUFA EPA and DHA serve as the precursors for resolvins and other SPM. In rodent models of inflammation, administration of EPA or DHA increases resolvin production²⁵. Mice with transgenic expression of a gene from Caenorhabditis elegans that enables endogenous generation of omega-3 PUFA from omega-6 PUFA (denoted *Fat-1*), are protected from inflammatory conditions such as colitis and pathological retinal angiogenesis and have elevated endogenous levels of resolvins^{68, 69}. These protective actions were recapitulated with exogenous delivery of resolvins, establishing causality in this relationship68, 69. There have been numerous studies investigating the protective effects of increased dietary intake of omega-3s on a diverse range of inflammatory diseases in humans, including asthma and arthritis, as well as CVD, based in large part on the fact that the typical Western type diet is deficient in omega-3 PUFA70, 71. One landmark clinical trial, the Gruppo Italiano per lo Studio della Sopravvivenza nell'Infarto miocardico (GISSI) Prevenzione study, enrolled more than 11,000 recent MI patients and reported that 1g of daily omega-3 supplementation for three and half years significantly reduced the risk of cardiovascular death⁷². More recently, GISSI also found greater survival in HF patients treated with omega- $3s^{73}$. Though these trials were able to demonstrate the beneficial effects of omega-3 supplementation, the mechanism of this protection remains unclear. In humans, several studies have shown that intake of omega-3 PUFA increases production of resolvins and/or their biosynthetic pathway markers (e.g., 18-HEPE and 17-HDHA) in plasma and/or serum⁷⁴⁻⁷⁹. In a very recent study, treatment with Lovaza (omega-3 ethyl esters) in patients with stable CAD increased levels of $RvD6$, 17R-PD1 and $RvE2^{80}$. Whether production of SPM is causally related to the protective actions of omega-3s in humans requires further clinical studies and it is also important to note that some chronic inflammatory diseases are associated with defects in SPM production^{25, 81}. Given that, unlike hormones, SPM are locally produced and have actions locally in inflammatory exudates (i.e., autacoids), the relevance of their measurement in plasma is still unclear. Given that some clinical trials have failed to show improvements in cardiovascular endpoints with omega-3 PUFA supplementation⁷¹, it is intriguing to speculate that altered downstream conversion to bioactive SPM could be related to variable results in such trials. Such variation could also arise from the diversity in dosing and preparations (e.g., ethyl esters vs. triglycerides). This is particularly relevant when one considers that the appearance of SPM is dependent upon multiple biosynthetic checkpoints and enzymatic degradation, rather than a passive linear increase with omega-3 intake.

Moreover, high intake of omega-3s could also bias the utilization of EPA and DHA away from SPM and toward the formation of free radical-initiated auto-oxidation products 82 . While the evidence to date indicates that SPM and their intermediate precursors are indeed produced in humans in vivo and are demonstrable by LC-MS/MS^{74–76, 79, 83, 84}, it is noteworthy that with inadequate approaches and methodology, some studies were unable to identify endogenous SPM in plasma 85 , which have specific stereochemistry and require the use of commercially available deuterium-labeled standards for their definitive identification in human tissues. These are important considerations for the development and implementation of new therapeutic approaches and also suggest that targeted delivery of specific SPM may be a better therapeutic approach.

One important design aspect of the GISSI trials was the instruction of enrolled patients to adhere to their recommended preventive medications in addition to the omega-3 PUFA supplement⁸⁶. Included in the preventive medications for many of these patients was aspirin. Aspirin is one of the most widely used anti-inflammatory drugs, and low-dose aspirin is currently recommended by the American Heart Association (www.americanheart.org) for people at high risk of heart attack and heart attack survivors. It was traditionally held that the salubrious effects of an aspirin regimen stemmed solely from halting prostaglandin and thromboxane production87. More recently, however, additional anti-inflammatory actions of aspirin have been described. Aspirin-acetylation of COX-2 blocks the production of prostaglandin precursors, but also switches the activity of COX-2 such that it generates 15R-HETE with AA as a substrate, or 18-HEPE and 17R-HDHA with EPA and DHA as substrates, respectively. These intermediates are subsequently converted into epimeric (i.e., aspirin-triggered) lipoxins and resolvins that share the potent biological actions of the native mediators in which the chirality of the C15 or C17 hydroxyl groups in 15-HETE and 17- HDHA are predominantly in the S configuration^{44, 45, 49, 53, 88}. Thus, aspirin could be considered to be "resolution friendly" because in addition to blocking inflammation, it jump starts resolution by shifting the lipid mediator profile from pro-inflammatory eicosanoids to SPM. Interestingly, some of these mediators (i.e., RvE1) regulate platelet activation as well (*vide infra*). Of interest, a recent study demonstrated that aspirin treatment enhances reverse cholesterol transport in hypercholesterolemic mice, which reduces atherosclerosis⁸⁹. This was associated with increased production of lipoxins in the liver and exogenous delivery of a stable analog of LXB₄ recapitulated these protective effects. These results indicate that in addition to the inflammation-resolving roles of SPM, they may have additional atheroprotective actions.

In addition to aspirin, statins are a widely prescribed class of lipid-lowering therapeutics with anti-inflammatory effects in the cardiovascular system, as discussed above⁴. Results of the JUPITER trial (Justification for the Use of Statins in Prevention: An Intervention Trial Evaluating Rosuvastatin) demonstrated that administration of Rosuvastatin decreases systemic markers of inflammation and that the observed reduction in major cardiovascular events is independent of the reduction in LDL cholesterol⁹⁰. Though it is clear that statins have anti-inflammatory effects, their mechanism of action is not entirely clear. Similar to aspirin, statins have been shown to target COX-2, however rather than acetylating the enzyme, statins increase S-nitrosylation and bias the enzyme toward the formation of 15 epimeric lipoxins as well^{91, 92}. More recently, it was demonstrated that S-nitrosylation of

COX-2 via administration of Atorvastatin initiates the biosynthesis of a novel group of 13 series resolvins from omega-3 docosapentaenoic acid $(DPA)^{93}$. Human and mouse leukocytes were shown to convert DPA to these novel bioactive products analogous to other SPM including the D-series resolvins 93 . These bioactive compounds are produced in PMNendothelial co-cultures and are present in human and mouse tissues during sterile inflammation or infection⁹³. The structures of four members of this family have recently been elucidated and contain conjugated triene and diene double bonds as well as an alcohol at C13 for which they have been termed 13-series resolvins $(RvTs)^{93}$. Though their biosynthetic pathway remains to be fully detailed, it has been shown that endothelial COX-2 initiates their production during PMN-endothelial cell interactions and S-nitrosylation of the enzyme by Atorvastatin further increases their production. Conversely, selective COX-2 inhibitors diminish their formation⁹³. Along these lines, with the widespread use of these and other therapeutics that target the inflammatory response, it is critical to underscore the importance of avoiding drugs that are "resolution toxic"¹⁰. Broad anti-inflammatory treatments are insufficient in that many of the signals produced during the inflammatory phase actually play key roles in initiating resolution (e.g., prostanoids-*vide supra*)⁹⁴. Indeed, selective COX-2 inhibitors that are detrimental for CVD patients because they perturb production of protective prostanoids (i.e., prostacyclin), also impair resolution and resolvin production during acute inflammation, as do LOX inhibitors³⁹. Ideal therapies to treat inflammatory diseases would be dual in nature and blunt the inflammatory response, while promoting the resolution program 10 .

Cellular actions of resolvins in the context of CVD

Neutrophils

As the first responders to sites of microbial invasion and tissue injury, the fundamental role of PMN as drivers of inflammation has been thoroughly described⁹⁵. Their capacity as the initiators of resolution, however, is just beginning to be appreciated $29, 96$. In response to PGE_2 , PMN transition from releasing pro-inflammatory LTB_4 to pro-resolving LXA_4 in a process known as lipid mediator class switching (*vide supra*)⁴². PMN are also a prominent source of resolvins and thus, similar to macrophage subsets, PMN can be reprogrammed to promote inflammation or enact its resolution. LXA4 is critical in halting further PMN recruitment and was shown in isolated PMN to inhibit chemotaxis, as well as transendothelial and transepithelial migration (Fig. 3) 97 . As noted above, SPM (e.g., LXA₄, $RvE1$) can also override survival signals in PMN, ensuring their timely apoptosis^{41, 63}. Additionally, both lipoxins and resolvins act on macrophages to promote the efferocytosis of apoptotic PMN98, a process that reciprocally stimulates the production of both E and Dseries resolvins, maresins and protectins thereby promoting the propagation of resolution rather than pro-inflammatory signaling (Fig. $3)^{96, 99}$. Resolvins also act directly on PMN to decrease chemotaxis toward chemokine gradients (e.g., IL-8) and to reduce the surface expression of adhesion receptors (e.g., CD11b/CD18) while preventing L-selectin (CD62L) shedding40, 58. These selective actions of SPM on PMN activation, rolling, and adhesion to the activated endothelium have been demonstrated both *in vitro* and *in vivo* using human endothelial cells under flow or intravital microscopy-based imaging of the inflamed microvasculature^{40, 59, 100}. Accordingly, all SPM have been shown to blunt PMN infiltration

into sites of acute inflammation, including bacterial peritonitis, dermal inflammation and lung inflammation²⁵. In addition to blunting PMN activation and regulating apoptosis, SPM have also been shown to increase the expression of chemokine receptors (i.e., CCR5) on apoptotic PMN, thereby facilitating chemokine scavenging during resolution as these apoptotic cell/chemokine clusters are subsequently cleared by macrophages³⁶. Interestingly, in concert with blunting PMN recruitment, SPM enhance PMN-mediated bacterial containment (Fig. 3), which translates into increased survival and reduced systemic inflammation in microbial sepsis, a condition that when uncontrolled, has detrimental cardiovascular implications^{37, 40}.

In response to a sterile ischemic injury in tissues such as the heart, PMN rapidly infiltrate as they would during an infection. During this process, however, excessive generation of proinflammatory mediators, as well as reactive oxygen species (ROS), are released and cause further injury to the surviving myocytes¹⁰¹. Therefore, promoting a timely removal of PMN is paramount in protecting the healthy myocardium during ischemia. Recently, it was demonstrated that treatment with RvD1 after MI in mice reduced accumulation of PMN in the ischemic myocardium, reduced fibrosis and improved fractional shortening, compared with untreated mice 102 . These actions were associated with increased SPM biosynthesis in the spleen and reduced splenic PMN numbers at day 5 post-MI. Similarly, it was previously shown that RvE1 decreases infarct size in a rat model of MI/reperfusion injury103. Proresolving mediators have also been described to be protective in ischemia/reperfusion injury outside the heart. In the kidney, in response to reperfusion after an ischemic insult, there is increased production of D-series resolvins and protectins. Treatment with resolvins prior to ischemia decreases PMN infiltration, tissue levels of myeloperoxidase (MPO), and tissue fibrosis, while administration shortly after reperfusion was also organ protective 104 . In a model of hind limb ischemia/reperfusion, resolvins protect against second-organ reperfusion injury by blocking PMN recruitment in the $\text{lung}^{67, 105}$. The endogenous protective role of SPM was demonstrated in mice lacking the ALX/Fpr2 receptor, in which ischemia/ reperfusion resulted in excessive leukocyte adhesion and emigration in the mesenteric microcirculation¹⁰⁶. As noted above, there is also emerging evidence that PMN participate in distinct phases of tissue repair as well²⁹. For instance, a very recent study demonstrated that PMN depletion impairs the development of a tissue reparative macrophage phenotype, resulting in increased apoptotic cell accumulation in the heart after MI¹⁰⁷. Along these lines, recent studies by Gronert et al. identified a distinct tissue resident and lymphoid homing PMN subset that produces SPM to blunt activation of Th1 and Th17 T cells in a murine model of immune-driven dry eye disease¹⁰⁸. Clearly, the multifaceted roles of PMN in inflammation, resolution and tissue repair deserve further study.

While mature human atherosclerotic plaques largely contain macrophages and T cells, recent studies have begun to elucidate the roles of PMN in the initiation and progression of atherosclerotic plaques, as well as in plaque rupture 109 . In humans, peripheral blood PMN counts correlate with coronary syndromes¹¹⁰. Initiation of atherosclerosis is marked by subendothelial retention of LDL and endothelial activation at sites of disturbed flow¹¹. This dysfunction is accompanied by the upregulation of adhesion molecules and production of chemokines by endothelial cells (EC) and leukocytes (e.g., interleukin 8, C-C chemokine ligand 3), which subsequently recruit PMN (in addition to other leukocytes; vide infra) to

the vessel wall. In mice, atherosclerotic lesion size is positively correlated with circulating PMN levels and conversely smaller lesions with PMN depletion 111 . Importantly, recruitment of monocytes is diminished in the absence of PMN, which is due in part to PMN-derived granule protein, cathelicidin¹¹². Diabetes exacerbates atherosclerosis in part by increasing PMN production of S100A8/S100A9, which leads to enhanced myelopoiesis and monocyte accumulation in lesions113. Moreover, a recent study found that cholesterol crystals active PMN to release neutrophil extracellular traps (NETs), which amplify immune cell activation in atherosclerosis^{109, 114}. As noted above, PMN are a rich source 5-LOX-derived LTB₄, which promotes further leukocyte recruitment. Mice lacking the $LTB₄$ receptor, BLT1 have a significant reduction in plaque formation and macrophage recruitment¹¹⁵ as did mice that received bone marrow transplants with genetically reduced levels of 5 -LOX¹¹⁶. Importantly, SPM including $RvD1$ block $LTB₄$ -stimulated expression of adhesion molecules and actin polymerization in PMN, and the E-series resolvins block activation of BLT1 by $LTB₄^{58, 66}$. PMN have also been linked with latter stages of atherosclerosis, including plaque destabilization. Plaques that are highly inflamed will continue to attract PMN to infiltrate, and the pro-inflammatory mediators and oxidants they produce may contribute to endothelial erosion and rupture of the fibrous cap. While most studies have primarily focused on other cell types (i.e., macrophages, vascular smooth muscle cells) to combat atherosclerosis, targeting PMN may offer valuable insights into halting the initiation or the eventual rupture of a plaque. All of these aforementioned features are indicative of defective resolution. In this context, it has previously been shown that RvE1 reduces PMN activation and tissue destruction in periodontitis, which is a risk factor for atherosclerosis 117 . In rabbits, periodontitis increases atherosclerosis and administration of RvE1 simultaneously decreases both periodonitits and atherosclerosis¹¹⁸. Further studies revealed that $RvE1$ reduces atherosclerosis even in absence of periodontitis, which was associated with a reduction in systemic levels of C-reactive protein (CRP). Interestingly, platelet:PMN aggregates formed during inflammation lead to SPM biosynthesis, which have been shown to reduce human platelet: PMN aggregates^{119, 120}. Given that platelet: PMN aggregates contribute to plaque inflammation¹²¹, these results highlight that SPM may have additional protective roles in regulating PMN activation during inflammation associated with atherosclerosis.

Macrophages

Both tissue macrophages and monocytes that differentiate into macrophages after being recruited to the site of injury biosynthesize and respond to SPM during resolution. Several SPM, including E-series and D-series resolvins potently enhance macrophage phagocytosis of bacteria and efferocytosis, essential functions fundamental to host-defense and resolution (Fig. $3)^{25}$. Stimulation of phagocytosis by SPM is at least partially dependent upon downstream activation of Akt, PI3K and ERK, and involves cytoskeletal remodeling associated with activation of Rho-GTPases (i.e., Cdc42, RhoA, Rac) in a receptor-dependent manner ^{65, 122–125}. Interestingly, during efferocytosis, SPM (e.g., RvD1, RvD2, RvE2) production increases, which potentiates further clearance of debris and apoptotic cells in an autocrine and paracrine manner^{39, 99}. However, in several chronic inflammatory diseases (e.g., asthma, obesity/diabetes), SPM production is disrupted and these conditions are also associated with defective resolution and efferocytosis 81 . Indeed, in diabetic wounds,

apoptotic cells accumulate and are associated with defective wound healing¹²⁶. RvD1 enhances wound healing in diabetic mice in part by enhancing efferocytosis¹²⁵. The delayed clearance of apoptotic cells in the thymus was also rescued by RvD1 and in isolated diabetic macrophages, RvD1 rescues defective phagocytosis in a receptor (ALX/Fpr2) and PI3Kdependent manner 125 . The specific mechanisms underlying impaired SPM production and efferocytosis in these diseases are incompletely understood. It is known however that the cellular location of the key 5-LOX enzyme plays an important role in dictating the predominant lipid mediators that macrophages produce. As noted above, RvD1 promotes cytoplasmic localization of 5-LOX (rather than nuclear) and shifts the production in favor of pro-resolving LXA_4 rather than the pro-inflammatory LTB_4^{56} . More recently, the mechanism of this nuclear localization was found to be governed by signaling through the myeloid-epithelial-reproductive tyrosine kinase (MerTK) receptor, mechanistically coupling efferocytosis to SPM biosynthesis¹²⁷. In mice genetically engineered to resist cleavage (and therefore inactivation) of this receptor, SPM biosynthesis is increased and resolution of acute inflammation and efferocytosis are improved 127 . In acute MI, mice deficient in efferocytosis receptors, Mertk and/or Mfge8, have decreased cardiac function and remodeling that was specifically due to abrogation of efferocytosis and not to an alteration in the circulating or tissue resident macrophages, nor the ability of monocytes to infiltrate the tissue^{128, 129}. It is, therefore, reasonable to consider that defective efferocytosis in CVD could be related to defective production of SPM, but that these alterations could be potentially rescued by exogenous administration of SPM. Whether there is a causal relationship between SPM production and efferocytosis in the heart during MI requires further study.

Defective efferocytosis has also been implicated as a crucial underlying contributor to the pathophysiology of atherosclerosis²². Macrophages play essential roles in all aspects of the life of an atherosclerotic plaque (Fig. 3). They are important both in the pathology of the lesion from initiation of formation to expansion, necrosis and eventual rupture, as well as in the resolution and regression of lesions^{3, 130}. Monocytes infiltrate the subendothelium of the arterial wall in response to inflammatory cytokines. Once in the plaque environment, monocytes differentiate into macrophages and attempt to clear excess oxidized lipoproteins and cholesterol from the tissue 22 . At this early stage of lesion development efferocytosis seems to function normally and the majority of plaques do not progress to a vulnerable state. In the plaques that do progress, however, both key events of resolution (cessation of additional immune cell entry and efferocytosis) are impaired 22 . This precipitates the generation of lipid-laden foam cells that fail to clear from the plaque, continue to secrete pro-inflammatory mediators and eventually undergo post-apoptotic secondary necrosis 22 . Atherosclerosis could, therefore, be viewed as a state of failed resolution of inflammation and defective clearance of plaque macrophages may underlie the progression of advanced atherosclerotic lesions, characterized by macrophage necrosis (Fig. $3)^{22}$. In a mouse model of diabetic atherosclerosis, isolated peritoneal macrophages were shown to have impaired phagocytosis and in atherosclerotic lesions, there was defective macrophage efferocytosis that was reversed by a fish-oil diet rich in SPM precursors, EPA and $DHA²¹$. Interestingly, it was also recently shown that RvD1, acting through ALX/FPR2, protects macrophages from oxidative stress-induced apoptosis during efferocytosis, in part by regulating NADPH oxidase activation and expression of apoptotic proteins, Bcl-xL and Bcl-2131. In the context

of atherosclerotic lesions, the ability of macrophages to survive and continue to phagocytose debris despite the highly oxidative environment is critically important. Furthermore, atherosclerotic apoE-null mice lacking SPM biosynthetic enzyme 12/15-LOX display exacerbated lesion formation compared with *apoE*-null mice and macrophage-specific overexpression of $12/15$ -LOX protects against lesion development¹³². Macrophages isolated from these mice produce higher amounts of $SPM¹³²$. Similarly, in rabbit models of atherosclerosis, it was shown that macrophage-specific overexpression of 15-LOX is atheroprotective, resulting in decreased lesion area¹³³. It should be noted that these enzymes produce both pro-inflammatory and pro-resolving mediators in a temporal manner and both atheroprotective and atherogenic roles of this enzyme have been reported 134 . However, as noted above, RvE1 used as an oral/topical agent decreases aortic atherogenesis, the intima/ media ratio, and inflammatory cell infiltration into plaques¹¹⁸. Additionally, recent studies documented that the ALX/FPR2 receptor has an endogenous anti-atherosclerotic role and that selective targeting of this receptor improves lesion stability and reduces lesion necrosis in mice with pre-existing atherosclerosis^{135, 136}. These results collectively demonstrate that chronic macrophage-mediated inflammation in atherosclerosis may be due in part to a failure of endogenous resolution programs.

In addition to macrophage efferocytosis, the phenotype of both monocytes and macrophages is an important determinant of resolution (Fig. 3). Recruited by specific chemokines (e.g., C-C chemokine ligand 2; CCL2) in the injured myocardium or atherosclerotic lesions, inflammatory Ly6Chigh monocytes infiltrate the tissue shortly after the initial PMN wave, as occurs during any normal acute inflammatory response¹⁰¹. After this first phase of recruitment, monocytes can differentiate into $Ly 6C^{low}$ monocytes/macrophages to facilitate tissue repair in a process dependent in part upon the transcription factor, Nr4a1137. These monocytes/macrophages facilitate repair because they are a rich source of growth factors and cytokines that promote tissue vascularization and matrix remodeling, while inflammatory monocytes/macrophages secrete inflammatory cytokines (e.g., IL-1β, IL-6, TNF-α) and are highly microbicidal¹³⁸. Too many Ly6Chigh cells or a delayed switch into reparative macrophages can lead to excessive proteolysis of the healthy myocardium surrounding the infarct, while increased production of IL-1β enhances proliferation of hematopoietic stem and progenitor cells to promote leukocytosis $139, 140$. In addition, a population of embryonicderived resident cardiac macrophages may contribute to homeostasis, as well as tissue repair in the heart during injury, and can be replaced by circulating monocytes and through local proliferation^{17, 141}. Similar to the heart, the phenotype of lesional macrophages in atherosclerosis has become an important indicator of the overall plaque environment. Macrophages have traditionally been lumped into general classically-activated (M1) or alternatively-activated (M2) designations based largely on in vitro model systems in which macrophages are polarized with LPS/IFN γ or IL-4, respectively, although this nomenclature has been revised because macrophages largely exist in a phenotype spectrum dependent upon the local milieu¹⁶. This is true for disease states like atherosclerosis, where distinct M1 vs. M2 populations are difficult to discern¹³⁰. With this inherent plasticity, agents that induce pro-resolving characteristics in macrophages may be beneficial in breaking the feed forward pro-inflammatory cycle occurring in the lesion. Notably, SPM biosynthetic enzyme, ALOX15, is highly induced in macrophages stimulated with IL-4 and these IL-4 stimulated

macrophages make more SPM overall than LPS/IFN γ -polarized macrophages⁹⁹. SPMs have also been implicated in promoting a switch from an inflammatory macrophage phenotype to a tissue reparative phenotype. For instance, it was recently shown that RvD1 blunts macrophage chemotaxis toward CCL2, while enhancing phagocytic function. In human macrophages, RvD1 potently blocks production of pro-inflammatory cytokines (e.g., IL-1β, IL-8 and CCL2) in a GPR32-dependent manner, which could help to prevent further recruitment of inflammatory monocytes¹⁴². In inflamed obese adipose tissue, $RvD1$ decreases the proportion of inflammatory macrophages and enhances macrophage phagocytosis and expression of arginase 1143, 144. In isolated macrophages, RvD1 increases expression of Arg1, Ym1, IL-10 and CD206, typical of an anti-inflammatory macrophage phenotype¹⁴³. RvD2 similarly increases surface expression of CD206, as well as CD163, in human macrophages in a GPR18-dependent manner⁶⁷. In murine peritonitis, both RvE1 and RvD1 have been shown to decrease CD11b expression on macrophages. CD11b^{low} macrophages have been reported to express lower levels of iNOS, COX-2 and MMP-9 and engulf more apoptotic cells compared with CD11 b^{high} macrophages¹⁴⁵. There have also been reports of a previously undescribed resolution-phase macrophage (rM) subset with similar features of the CD11b^{low} population and possessing a hybrid of the M1/M2 phenotypes (M2: mannose receptor expression, IL-10, TGF-β and arginase 1; M1: high iNOS and COX-2 expression)¹⁴⁶. These rM also express $A\ell\alpha I_5$, indicating that they may contribute to SPM production during resolution. As critical regulators of the resolution program, targeting the actions of macrophages may be an effective strategy to address chronic maladaptive inflammation in CVD.

Endothelial cells

As the interface between the blood and surrounding tissues, the vascular endothelium plays an important early role in the inflammatory response by regulating the passage of macromolecules and immune cells across the vessel wall. As such, the temporal regulation of endothelial permeability and activation is a critical determinant of resolution of the inflammatory response. Indeed, CVDs such as atherosclerosis are associated with EC dysfunction and chronic vascular inflammation¹¹. Analogous to the dynamic regulation of leukocytes during resolution, EC activation is a self-limited process that is uniquely positioned to control the magnitude and duration of the acute inflammatory response. Indeed, continued EC activation, characterized by expression of specific leukocyte adhesion receptors such as vascular cell adhesion molecule 1 (VCAM-1), is associated with chronic inflammatory diseases such as atherosclerosis and diabetes¹¹. Accordingly, EC participate in the local biosynthesis of SPM during acute inflammation and are also direct cellular targets of SPM (Fig. 3). As noted above, the biosynthesis of most SPM occurs during cell:cell interactions in which the biosynthetic intermediates, such as 18-HEPE, can be generated in EC and transformed by adhering leukocytes expressing 5-LOX to generate SPM such as RvE144, 45. Furthermore, it was in the co-culture of human PMN and EC that the novel 13 series resolvins ($RvTs$) were recently identified⁹³. A recent study documented that $RvD1$ and RvE1 are also biosynthesized during co-culture of choroid-retinal (CR) EC with leukocytes (both PMN and monocytes), but not by CREC alone¹⁴⁷. This mechanism likely serves as a local circuit to ensure timely termination of leukocyte adhesion to EC. Indeed, EC express the known receptors for SPM, including ALX/FPR2, GPR32 and

GPR1858, 148, 149. Acting through these receptors, SPM regulate both recruitment and adhesion of leukocytes. AA-derived LXA4 downregulates VCAM-1 and additional adhesion receptors, E-selectin¹⁵⁰ and ICAM-1^{151, 152}, but also increases nitric oxide (NO) and prostacyclin (PGI₂) that potently counter-regulate leukocyte: EC interactions and platelet activation^{153, 154}. Lipoxins have further EC-specific effects including the downregulation of NADPH oxidase, which results in the decreased generation of ROS¹⁵⁵. Similarly, resolvins have also been shown to regulate leukocyte:EC interactions. For instance, RvD2 limits leukocyte:EC adhesion in a NO-dependent manner and the regulation of PMN recruitment during microbial peritonitis by RvD2 is attenuated in mice deficient in endothelial nitric oxide synthase $(eNOS)^{40}$. Both RvD1 and RvD2 limit PMN interaction with human EC prestimulated with inflammatory cytokines (e.g., TNFα, IL-1β) under shear conditions at a physiological flow rate in a receptor-specific manner^{40, 59}. In this context, resolvins block PMN capture, rolling and firm adhesion to EC and they attenuate EC production of chemokines (e.g., IL-8 and CCL2) that recruit additional leukocytes^{132, 147}. A recent study of the mechanisms whereby resolvins regulate inflammatory signaling in human EC uncovered that D-series resolvins stimulate the activation of PI3K and Akt in a receptordependent manner and thereby regulate the production of developmental endothelial locus 1 (Del-1), an anti-inflammatory protein secreted by EC that prevents PMN transmigration¹⁵⁶.

In addition to regulating EC-dependent adhesion of leukocytes during inflammation, resolvins also regulate permeability of the microvasculature. In vitro, RvD1 protects against endotoxin-induced impairment of barrier function in human EC^{157} and aspirin-triggered-RvD1 enhances the restitution of endothelial barrier function following acute lung injury iⁿ $vivo¹⁵⁸$. These results build on prior studies showing that lipoxins counter-regulate leukocyte-mediated microvascular permeability in vivo during acute inflammation stimulated by pro-inflammatory lipid mediators¹⁵⁹. Permeability of the microvasculature is increased by several inflammatory mediators that also regulate angiogenesis. Along these lines, LXA4 decreases vascular endothelial growth factor receptor 2 (VEGFR-2) phosphorylation and its downstream signaling in human EC, thereby blocking VEGFinduced angiogenesis 152 . Similarly, in a model of murine retinal neovascularization, administration of exogenous SPM (LXA4, RvD1, RvE1 and PD1) protects against pathologic angiogenesis^{69, 160, 161}. This is in contrast to some pro-inflammatory lipid mediators (i.e., 12-HETE and PGE_2) that promote angiogenesis^{162, 163}. Collectively, EC have emerged as important cellular targets of SPM during resolution, suggesting that engagement of SPM receptors might offer a new approach to reversing EC dysfunction that is associated with CVD. Future studies will be required to understand fully the role of SPM in EC biology and whether dysfunctional EC in chronic diseases is associated with deficits in resolution circuits at the level of EC:leukocyte interactions.

Vascular smooth muscle cells

In addition to their physiological role in regulating blood vessel tone, vascular smooth muscle cells (VSMC) also contribute to the pathophysiology of atherosclerosis and restenosis. In response to vascular injury, VSMC undergo a phenotype switch from a contractile to a synthetic phenotype; synthetic VSMC are more proliferative and chemotactic and are characterized by a decreased expression of contractile proteins and increased

production of pro-inflammatory mediators¹⁶⁴. During chronic vascular inflammation (e.g., atherosclerosis) this pro-inflammatory environment can lead to successive contractile to synthetic phenotype switching thereby propagating the pro-inflammatory environment. Moreover, VSMC can assume a phenotype similar to macrophages and express macrophage markers, such as CD68 and $F4/80$ in lesions¹⁶⁴. Interruption of this pro-inflammatory signaling cycle may, therefore, be a useful strategy to combat inflammatory diseases of the vessel wall. Receptors for SPM, including ALX/FPR2 and ChemR23, are expressed in human saphenous vein SMC and administration of RvE1 and 15-epi-LXA₄ counter-regulate platelet-derived growth factor (PDGF)-stimulated VSMC migration in a dose-dependent manner (Fig. 3)¹⁶⁵. These actions are due in part to direct regulation of PDGF receptor phosphorylation. Building on these in vitro studies, it was recently found that 15-epi-LXA₄ inhibits VSMC migration and intimal hyperplasia following murine carotid artery ligation and this protection afforded by 15-epi-LXA₄ is abolished in mice lacking the $ALX/Fpr2$ receptor¹⁶⁶. Interestingly, genetic deficiency of $ALX/Fpr2$ decreases stability of atherosclerotic lesions, effects that were attributed to decreased collagen production by VSMC¹⁶⁷. Given that VSMC can also contribute to stability of atherosclerotic lesions¹⁶⁴ and that it is their phenotype that dictates their contribution to atherosclerosis, these results suggest that SPM may regulate the balance of VSMC phenotypes important for regulating lesion stability. More recently, it was demonstrated that RvD1 and RvD2 inhibit proliferation, migration, monocyte adhesion, superoxide production and pro-inflammatory gene expression in human¹⁶⁸, rat¹⁶⁹ and mouse¹⁷⁰ VSMC. Similar results were observed with another SPM, maresin 1 $(Mar1)^{171}$. In a rabbit model of balloon angioplasty, local delivery of RvD2 reduces cell proliferation, leukocyte recruitment and neointimal hyperplasia¹⁶⁸. Interestingly, RvD2 and Mar1 increase M2 macrophage polarization during vascular injury, which may help to promote vessel remodeling and repair¹⁷⁰. Additionally, RvD1, delivered perivascularly via a novel biodegradable wrap reduces carotid artery neointimal formation without promoting thrombosis after carotid angioplasty in rats¹⁶⁹. In a model of human pulmonary artery hypertension, RvD1 and RvE1 are able to normalize arterial hyper-reactivity induced by pro-inflammatory mediators $172, 173$. These results add further support to the concept that most CVDs are driven by abhorrent inflammation and that stimulating resolution of inflammation could potentially restore physiological function of the vasculature.

Platelets

In response to blood vessel damage, platelets are rapidly activated and are a critical component of the coagulation process that serves to re-establish a physical barrier and prevent blood loss. This process is tightly coupled with the inflammatory program and like the regulation of leukocyte trafficking, platelet activation must be tightly controlled to prevent further damage to the host. Platelet aggregation and ensuing thrombosis are significant vascular events that are linked to inflammation and atherogenesis and causative to the clinical manifestations of atherosclerosis, namely stroke and MI^{174} . In a randomized clinical trial involving healthy subjects receiving aspirin, pro-thrombotic thromboxane was reduced and there was a concomitant increase in 15-epi LXA₄^{175, 176}. Plasma levels of 15epi-LXA₄ were increased most significantly with low dose aspirin (i.e., $81mg$), the currently recommended dose for patients with CVD. Interestingly, human platelets express ChemR23

and RvE1 blocks ADP and thromboxane-stimulated platelet aggregation (Fig. 3)^{100, 177}. Further mechanistic studies revealed that RvE1 blocks ADP-mediated signals downstream of its receptor, $P2Y_{12}$, and that the effects of RvE1 are ChemR23-dependent¹⁷⁷. In a mouse burn wound model, exogenous systemic administration of RvD2 prevents thrombosis of the deep dermal vascular network and subsequent dermal necrosis. Treatment with RvD2 allowed enhanced neutrophil access to the dermis while preventing neutrophil-mediated damage due to an abrogation of pro-inflammatory mediator secretion¹⁷⁸. Lastly, it was very recently shown that SPM increase macrophage uptake of blood clots *in vitro*, indicating that SPM may play a role in clot remodeling during resolution⁸⁰. As noted above, SPM are biosynthesized during interactions of platelets and leukocytes during inflammation. Thus, these results collectively demonstrate that SPM may play key roles in regulating the resolution of thrombosis during inflammation, which could have important therapeutic implications for CVD.

Summary and future directions

A significant effort has been dedicated to the structural elucidation of pro-resolving mediators and the identification of their receptors⁹. It is becoming apparent that alterations in specific resolution pathways are associated with a wide range of chronic diseases. Chronic unresolved inflammation is now widely documented to play a causal role in the development of CVD. As such, a large effort is underway to determine whether targeting specific proinflammatory pathways could be beneficial for clinical management of CVD beyond the current standard of care (e.g., aspirin, statins, anti-platelet therapies)⁴. In contrast to traditional anti-inflammatory strategies that blunt the production of inflammation "initiators" and could potentially lead to immunosuppression, pro-resolving mediators described in this review activate distinct cellular programs to resolve inflammation without compromising host-defense⁹. In fact, pro-resolving mediators enhance host defense to both viral and bacterial infections and lower the threshold for antibiotic therapy^{37, 179}. Thus, new strategies to promote resolution of inflammation may offer unique opportunities to combat chronic inflammation associated with CVD. Pro-resolving mediators blunt excessive inflammation that can be detrimental to healthy tissues, particularly in sterile inflammation, and they also stimulate distinct processes necessary for tissue repair and regeneration. They have multiple cellular targets in the inflammatory response, including immune cells, platelets and vascular cells. Collectively, the structural elucidation of these mediators and identification of the signaling pathways that they engage could inform the development of therapeutic strategies encompassing a novel "resolution pharmacology" approach. Indeed, clinical trials with proresolving mediators in humans with other inflammatory pathologies are currently underway¹⁸⁰.

In regards to therapeutically stimulating resolution of inflammation in CVD, there are several things to consider. First, can SPM be effectively targeted to sites of local inflammation? In a rabbit model of atherosclerosis, delivery of RvE1 to the oral cavity decreased lesion formation in the aorta, suggesting that resolvins can get absorbed and have systemic effects¹¹⁸. Our group has also shown that microparticles released from activated immune cells carry SPM and as a form of biomimicry, we formulated humanized nanoparticles that showed enhanced bioactivity; these could potentially be vehicles for

targeted delivery¹⁸¹. Moreover, recent studies have begun to highlight other methodologies for local sustained SPM delivery (e.g., vascular wraps) that could be used in vascular surgery for example¹⁶⁹. Second, what are the mechanisms underlying failed resolution of inflammation? As noted in this review, several chronic diseases are associated with lower levels of SPM. Does this occur because of altered biosynthesis or enhanced metabolic inactivation? How are SPM receptors modulated in chronic diseases like CVD? Lastly, are nutritional based approaches (e.g., omega-3 PUFA) sufficient to promote resolution, or are there contexts in which resolution is perturbed regardless of substrate availability? These are all important questions for future work in the area and given that the field of "resolution physiology" is relatively new, an understanding of the mechanisms underlying failed resolution of inflammation in CVD is likely to be a fruitful area of future investigation.

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Non-standard Abbreviations and Acronyms

EC endothelial cells

VSMC vascular smooth muscle cells

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Time

Figure 1. The coordinated temporal events of self-limited acute inflammation

The ideal outcome of an acute inflammatory response is complete resolution. The inflammatory response can be divided into two general phases: initiation and resolution. Critical to progressing from initiation to resolution is the temporal switch in lipid mediators that are biosynthesized by leukocytes in the tissue; a process known as lipid mediator class switching. The earliest stage of the inflammatory response is marked by tissue edema due to increased blood flow and microvascular permeability and is mediated by the release of proinflammatory lipid mediators including the cysteinyl leukotrienes and prostaglandins. Polymorphonuclear neutrophils (PMN) infiltrate in response to lipid mediators including leukotriene B_4 and engulf and degrade pathogens. Subsequently, PMN undergo apoptosis and also switch from releasing pro-inflammatory mediators to pro-resolving mediators (e.g., resolvins) that signal the clearance of apoptotic cells by macrophages in an antiinflammatory process termed efferocytosis. In addition to promoting efferocytosis, proresolving lipid mediators halt further PMN recruitment and stimulate a pro-resolving macrophage phenotype that is important for tissue repair.

Figure 2. Biosynthesis and receptors for resolvins

The family of resolvins is divided primarily into two groups based on the omega-3 polyunsaturated fatty acid (PUFA) from which they are formed. For E-series resolvin biosynthesis, eicosapentaenoic acid (EPA) is utilized as a substrate for acetylated COX-2 or cytochrome P_{450} enzymes, giving rise to a 18-hydroperoxide intermediate that can be converted to either RvE1 or RvE2 by 5-LOX and additional enzymatic transformations, as indicated. Similarly, DHA serves as a substrate for 15-LOX, giving rise to a 17 hydroperoxide intermediate. This intermediate is subsequently converted to RvD1 or RvD2 via 5-LOX, the formation of an epoxide intermediate, and subsequent enzymatic hydrolysis. Monohydroxy fatty acids, 18-HEPE and 17-HDHA, serve as pathway markers for E-series resolvins and D-series resolvins, respectively. Once formed, resolvins elicit their potent biological actions by activating specific G-protein coupled receptors (GPCRs) in a stereoselective manner. RvE1 is an agonist for ERV/ChemR23, while both RvE1 and RvE2 block the actions of pro-inflammatory leukotriene B_4 by binding to its receptor, BLT1. Dseries resolvins are agonists for distinct GPCRs, with RvD1 binding to DRV1/GPR32 and ALX/FPR2, and RvD2 binding to GPR18.

Figure 3. Biological actions of resolvins and other SPM related to cardiovascular inflammation During acute inflammation or infection, SPM are generated during leukocyte:endothelial interactions and directly block PMN chemotaxis. By directly targeting the endothelial cells of the vasculature (e.g., post-capillary venules), SPM also prevent the capture, rolling and adhesion of PMN to the endothelial monolayer and block their extravasation across the endothelium. Additionally, they stimulate production of nitric oxide (NO) and prostacyclin (PGI2) from endothelial cells. In the tissue, SPM promote the protective actions of leukocytes by enhancing the phagocytosis of bacteria and cellular debris and macrophage efferocytosis of apoptotic PMN. In the context of atherosclerosis, SPM decrease endothelial cell production of leukocyte chemoattractants and adhesion molecules, potentially halting further recruitment of leukocytes to the vessel wall. They also potently prevent platelet aggregation and thrombosis. In the subendothelium, enhanced macrophage efferocytosis and promotion of a resolving macrophage phenotype contributes to a more stable plaque with decreased necrosis. During pathologic inward remodeling (i.e. restenosis), SPM combat neointimal hyperplasia by inhibiting vascular smooth muscle cell (VSMC) proliferation and migration while simultaneously preventing monocyte adhesion and inflammatory signaling.