A
rachidonic acid is a polyunsaturated omega-6 fatty acid that is released in response to tissue injury. Arachidonic acid represents one of the pivotal signaling molecules involved in the initiation and propagation of diverse signaling cascades regulating inflammation, pain, and homeostatic function. Drugs developed to target these signaling pathways represent >25% of annual pharmaceutical sales worldwide. Arachidonic acid is metabolized through 3 enzymatic pathways. The cyclooxygenase (COX) pathway produces prostanooids. The lipoxygenase (LOX) pathway yields monohydroxy compounds and leukotrienes, while the cytochrome P450 (CYP450) epoxygenase pathway generates hydroxy and epoxygenoids. This group of lipid mediators, which are derived from the 20-carbon atom arachidonic acid or similar fatty acids, is collectively referred to as eicosanoids (“eicosa” means 20 in Greek). A schematic metabolic pathway of arachidonic acid is shown in the Figure. There is mounting evidence that some of these metabolic products play critical roles in cardiovascular (CV) disease.

CV disease remains one of the leading causes of death in Western societies.
Cardiac failure is the final consequence of a variety of etiologies including coronary heart disease, myocardial infarction (MI), hypertension, arrhythmia, viral myocarditis, and genetic cardiomyopathies. Once heart failure (HF) develops, the condition is for the most part irreversible. Although considerable progress has been made in the pharmacologic and device management of HF in recent decades, the mortality in HF patients remains significant. Moreover, the incidence and prevalence of cardiac failure are increasing as the population ages.

Recently, our laboratories have taken advantage of a new technique of metabolomic profiling using liquid chromatography-tandem mass spectrometry (LC-MS/MS) to elucidate the contribution of arachidonic acid metabolism in CV diseases. Metabolomics is a promising approach that has been used widely as a powerful tool in disease diagnosis, biomarker discovery, toxicity evaluation, gene function, and pharmacologic research. In this review, we will provide examples of the use of metabolomic profiling in our 2 recent studies. Liu and colleagues used a broad metabolomics approach to quantify the representative oxylipin mediators derived from arachidonic and linoleic acids mediated by COXs, LOXs, and CYP450s. Oxylipins are oxygenated lipids, and one of the most biologically important groups of oxylipins is the eicosanoid family. Specifically, Liu and colleagues applied metabolomic profiling in a murine model and identified a link between the administration of rofecoxib (Vioxx) and adverse CV events. They found a significant increase in 20-hydroxyeicosatetraenoic acid (20-HETE), a potent vasoconstrictor and the culprit for increasing risk for MI and stroke. This mechanism may be shared among other nonaspirin nonsteroidal anti-inflammatory drugs (NSAIDs). In the second instance, Li and colleagues, using a similar approach, demonstrated the beneficial effects of increasing epoxyeicosatrienoic acid (EETs) and the EETs/dihydroxyeicosatrienoic acids (DHETs) ratio by application of soluble epoxide hydrolase (sEH) inhibitors in a murine MI model.

20-Hydroxyeicosatetraenoic Acid
Rofecoxib is a potent, orally active, and selective COX-2 inhibitor that was previously approved by the US Food and Drug Administration to treat a wide
Figure. Diagram illustrating the metabolic pathways for arachidonic acid and linoleic acid. Arachidonic acid is metabolized through 3 enzymatic pathways. The cyclooxygenase (COX) pathway produces prostanoids. The lipoxgenase (LOX) pathway yields monohydroxy compounds and leukotrienes, while the cytochrome P450 (CYP) epoxygenase pathway generates hydroxy and epoxyeicosanoids. See text for acronym expansions.

The current theory on the possible mechanisms responsible for the observed adverse effects of rofecoxib is that rofecoxib reduces the production of prostacyclin $\text{(PGI}_2\text{)}$, an inhibitor of platelet aggregation (PGI2) (Figure). This results in an increase in platelet aggregation and may predispose patients to adverse CV events including MI or stroke. On the other hand, it has long been known that NSAIDs inhibit the production of the potent platelet activator thromboxane (TX) A$_2\text{}$, so these agents may have thrombolytic activities. Hence, one might expect that conventional NSAIDs are “neutral” or even beneficial to the CV system. However, a significantly increased risk for CV diseases such as MI, hypertension, and HF has been observed to be associated with the administration of some of the nonaspirin NSAIDs, including but not limited to diclofenac, ibuprofen, naproxen, and indomethacin. Thus, the current hypothesis provides an incomplete explanation for the observed adverse CV events associated with the use of NSAIDs. We reason that this may result from the fact that the dominant theory is based on monitoring only a few arachidonic acid metabolites. To evaluate the risks and benefits of selective COX-2 inhibitors and to develop safe coxibs or adjuvants to improve the safety of existing coxibs, it is critical to understand the possible interactions among different arachidonic acid metabolic pathways.

We used a murine model that was administered with rofecoxib for a period of 3 months. In this model, there was a dramatic decrease in bleeding time, which reflected an increase in platelet aggregability. Increased platelet aggregability has been associated with the pathogenesis of MI and stroke. The quantitative levels of 27 oxylipin mediators of the plasma from treated animals were determined using metabolomic profiling. There was a $>120$-fold increase in the plasma concentration of 20-HETE in the mice treated with rofecoxib. Moreover, a direct infusion of 20-HETE in mice also resulted in shortened bleeding time. Taken together, our data may provide a link between the use of rofecoxib and related compounds and the reported adverse CV events. This hypothesis suggests 20-HETE as a biomarker for CV risk from coxibs as well as possible strategies for attenuation of their adverse effects. For example, we predict that inhibition or down-regulation of CYP4A and or CYP4F may ablate the CV events of coxibs. In addition, this study exemplifies the use of metabolomic profiling as a promising tool to gain a more comprehensive understanding of biologic processes.

**Epoxyeicosatrienoic Acids**

The CYP450 epoxygenase products, the epoxyeicosanoids, also known as EETs, are major anti-inflammatory arachidonic acid metabolites with a variety of biologic effects. There is growing evidence supporting the notion that EETs and other epoxy and diol fatty acids play a significant protective role in the CV system. EETs have been identified as potential endothelium-derived hyperpolarizing factors (EDHFs). Major roles of EETs include modulation of both blood pressure and inflammatory signaling cascades. EETs are also associated with a number of other physiologic functions, including modulation of ion channel activity, angiogenesis, cell proliferation, vascular smooth muscle cell migration, leukocyte adhesion, platelet aggregation and thrombolysis, and neurohormone release. It has been proposed that diminished production or concentration of EETs contributes to CV disorders. A polymorphism of the human CYP2J2 gene, which is highly expressed in heart and active in the biosynthesis of EETs, encodes variants with reduced catalytic activity and is independently associated with an increased risk of coronary artery disease. Transgenic mice with cardiomyocyte-specific over-expression of human CYP2J2 demonstrate enhanced post-ischemic...
functional recovery and significant protection against doxorubicin-induced cardiotoxicity. As the protective role of EETs in CV biology has been increasingly recognized, considerable interest has arisen in developing methods to enhance the bioavailability of these compounds.

There are a variety of pathways involved in the degradation of EETs, but the major pathway is catalyzed by the enzyme soluble epoxide hydrolase (sEH). sEH converts EETs to their corresponding diols, dihydroxyeicosatrienoic acids (DHETs), thus modifying the function of these oxylipins. Over the past few years, sEH has gained considerable attention as a therapeutic target for CV diseases.

Specifically, we tested the effects of sEHIs on prevention and reversal of cardiac hypertrophy and post-ischemia remodeling, which are among the most common causes that lead to heart failure. We demonstrated that sEHIs can prevent the development of pressure-induced cardiac hypertrophy using a murine model of thoracic aortic constriction (TAC). In addition, sEHIs reversed the pre-established cardiac hypertrophy caused by chronic pressure overload, in which a high level of expression of sEH in mouse atrial and ventricular myocytes was documented.

Recently, our laboratory has also demonstrated the beneficial effects of sEHIs on the progression of cardiac remodeling using a clinically relevant murine model of MI. Using LC-MS/MS-based techniques, we documented a significant decrease in the EETs/DHETs ratio in an MI model indicating increased sEH activity, which may play a role in the progression of post-ischemia remodeling. Treatment with sEHIs resulted in the normalization of the EETs/DHETs ratio and a reduction in post-ischemia LV remodeling. Moreover, we have documented that the significant decrease in the EETs/DHETs ratio in the MI model showed a striking parallel with the changes in inflammatory cytokines at 3 weeks post-MI, which indicated a heightened inflammatory state. Additionally, the normalization of the EET/DHET ratios by sEHIs results in a reversal of the elevated cytokine levels in the MI model. Persistent inflammation involving increased levels of inflammatory cytokines plays a potential pathogenic role in the progression of LV dysfunction and remodeling in HF.

The sEHIs appear to change the pattern of inflammatory mediators from a state that promotes the propagation of inflammation toward one promoting resolution.

sEH has been shown to be expressed in cardiomyocytes. The expression of sEH is upregulated by angiotensin II in cardiac myocytes in vitro and in vivo, suggesting a potential regulatory role of sEH in angiotensin II-induced maladaptive hypertrophy. Finally, recent human epidemiologic studies have identified associations between variations in EET metabolic pathway genes and increased CV risk. A polymorphism leading to reduced gene activity of CYP2J2 is associated with an increased risk of coronary artery disease, and EPHX2 has also been identified as a susceptibility factor for HF. Taken together, these findings suggest that increased sEH activity and reduced bioavailability of EETs may play a significant role in the pathogenesis of HF.

Interestingly, sEHIs have been shown to indirectly downregulate the expression of COX-2 and synergize with NSAIDS toward the reduction of inflammation. This suggests that these drug combinations (NSAIDS and sEHIs) may produce a beneficial anti-inflammatory effect while reducing the required dose of COX-2 inhibitors, thus avoiding the adverse CV side effects attributed to COX-2 inhibitors.

Future Directions
Both studies on 20-HETE and EETs demonstrate the use of metabolomic profiling as a promising tool to gain a comprehensive understanding of the biologic processes. Increased sEH activity has been demonstrated in an animal model of MI, supporting the notion that sEH may play an important role in the progression of post-ischemia remodeling. However, increased expression level of this enzyme has not been directly detected in the heart. Further studies to explore the mechanism by which sEH activity is dysregulated in MI and possible involvement of other organs such as liver and kidney may help to shed new light on the molecular defects in the pathogenesis of myocardial failure. Moreover, in order to definitively determine the best therapeutic utility for sEHIs, future studies to evaluate the potential interactions of sEHIs with other pharmaceuticals are warranted. It has been shown that regulation of sEH is intimately tied to the renin-angiotensin-aldosterone system in animal models of hypertension and cardiac hypertrophy. sEHIs also synergize with COX-2 inhibitors and other modulators of the arachidonic acid cascade to exert anti-inflammatory effects. Thus, the combination of sEHIs and angiotensin-converting enzyme inhibitors or COX inhibitors may provide powerful combination drug therapies with favorable side effect profiles. Since HF is a complex clinical syndrome with diverse etiology and a wide array of pathophysiology, in order to translate the observed beneficial effects of sEHIs into clinical intervention in patient care, additional information is needed to identify whether the observed beneficial effects can be generalized to other causes of HF, such as idiopathic dilated cardiomyopathy and drug-induced HF.

HF and its comorbidities represent a major market and one of the paths to the clinic for sEH that is best supported by mechanism, animal studies, and human data. However, the length, high cost, and high risk of clinical trials for HF make this path unattractive to many pharmaceutical companies. Investigational new drug status for an sEHI could permit investigator-initiated clinical trials to address this problem. Finally, other oxylipins apart from the 0-6 arachidonic
acid metabolites may be relevant in CV disease. For example, the ω-3 fatty acids eicosapentaenoic acid (EPA) and docosahexaenoic acid (DHA) accumulate in the heart, and the epoxides of EPA and DHA are analogs of the EETs. In fact, DHA and EPA epoxides share some of the vasoactive and anti-inflammatory effects of the EETs in vitro and in some cases have been shown to be more potent. DHA and EPA epoxides are, in general, better substrates for sEH than the EETs, so it is possible that some of the cardioprotective effects of sEH inhibition are due to reduction in DHA and EPA epoxide metabolism in the heart. Intervention studies with ω-3 lipids could test the hypothesis that these natural products have a protective effect on cardiac hypertrophy.

In summary, metabolomic profiling has been shown to not only identify a potential marker of risk or effect of rofecoxib and possibly other drugs, but also to demonstrate paths to mitigate the risk of these valuable pharmaceuticals. In addition, metabolomic profiling expanded our knowledge of the role of eicosanoids in the progression of cardiac hypertrophy. This knowledge from profiling pointed to inhibitors of the sEH as possible therapeutic agents for the prevention or even treatment of this and other CV diseases. During the past few years, the use of sEHs in animal models have demonstrated that sEH have therapeutic potential in a broad range of cardiac diseases, many of which are comorbidities with hypertrophy. Although possible side effects associated with the inhibition or genetic deletion of sEH have been reported, the data obtained from several laboratories employing animal models of cardiac hypertrophy and ischemia/reperfusion support the notion that sEHs and possibly EET mimics represent promising therapeutic targets for combating detrimental cardiovascular remodeling, HF, and related diseases.

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