

Review Article

EET Intervention on HO-1 Prevent Obesity Derived Cardiovascular Diseases

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Abstract

The increase in the prevalence of obesity represents a worldwide phenomenon and is correlated with several metabolic and cardiovascular diseases, and cardiovascular disease remains the leading cause of death worldwide. And epoxyeicosatrienoic acid (EET), correlated with heme oxygenase (HO), with emerge to be a promising strategy for pharmacological mediation. Moreover, inhibit soluble epoxide hydroxylase (sEH), the enzyme converts EET to less potent metabolite, could upregulate EET concentration, which could upregulate HO-1, working together to reduce inflammation and increase vasodilation, in order to improve endothelial and cardiac function. The EET-HO pathway has been indicated as a most potent target for reversing oxidative stress and pre-adipocyte differentiation, so as to moderate oxidative capacity in mitochondrial dysfunction. While EET agonists and sEH inhibitors are becoming one of the potential therapies, and some of them are already in clinical trials. This review serves to summarize the ability of EET and HO pathway in attenuating the clinical impairments of obesity and associated cardiovascular diseases.

Keywords: Obesity, PGC-1a; Epoxyeicosatrienoic acid; Cardiotrophin; Adiponectin; Hyperthermia

Introduction

Differential phenotype between white fat and brite fat

Adipose tissue is not only a type of connective tissue but an endocrine organ with two very important roles in the mammalian body: energy homeostasis and reproduction [1,2]. Apart from these, fat also prevent delicate organs and from mechanical stress. What should be noticed is that it is not the number of adipocytes that change in obesity (namely hyperplasia) but change in size (hypertrophy of adipocytes) [3].

Traditionally, fat can be metabolized into two difference types: white and brown fat. White fat could be found in models of obesity and metabolic syndrome, which make up the bulk of fatty tissue in the animals [4], and brown fat could be seen in the supraclavicular and spinal region of infants and rodents, and human adults also have brown fat specific depots [5]. White adipocytes are associated with very large lipid droplets with an energy storing effect [4], while brown fat is associated with smaller lipid droplets can dissipate stored chemical energy as an energy releasing phenotype [6]. Brown fat can be activated by either cold exposure or β adrenergic signalling and is linked to mitochondrial uncoupling protein (UCP1) to uncouple ATP from the electron transport chain, and lead energy consumption to combat hypothermia, obesity, and diabetes [7].

Brown fat comes from a Myf5+/Pax7+ cellular lineage, which is different from white fat, and muscle tissue also stems from this cellular lineage [8]. Although the cellular lineages are different, the core elements of the adipogenic cascade are shared by all types of fat. PPARy is the main manager of fat cell formation, and Zfp423 is a significant transcriptional factor of adipocyte lineage with bZIP, CEBP α and β to be transcriptional cofactors, and this cascade works for both white and brown fat [9,10]. Moreover, CEBPa is locked in a differentiation loop with PPARy, and one increase the other does as well [9]. Ebf2 is an important protein that recruits PPARy in brown fat, and much of the specialized brown fat function is controlled by transcriptional cofactors such as Ebf2 [11]. Although they don't bind DNA directly, they both control which targets will be bound and which will be activated [12]. The SNS plays a critical part in fat development as well, which distributes signals to both white adipose and brown adipose [13]. These different signals indicated that the effect of cold exposure is very different than food deprivation.

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Obesity has a whole body effect and can lead to diabetes, hypertension, and all known inflammatory markers are associated in obesity [14-16]. Adipocytes are a major source of TNF α and relate with recruit macrophages [17]. There are two types of macrophages: the classically activated M1 macrophages (release IL-6, are proinflammatory) and the alternatively activated M2 macrophages (release IL 10, anti-inflammatory, involved in wound healing) [18]. The ratio of M1/M2 shifted to a pro-inflammatory state in obesity body [19], and these inflammations can cut off the adipocytes oxygen, resulting in an activation of HIF1 α (oxygen sensing transcription factor) which lead to metabolic dysfunction [20]. Adipocytes are extremely beneficial and vital for homeostatic balance in normal body, but these indicated over nutrition leads to obesity, and switch the normal balance towards inflammation.

Beside from white and brown fat, there is a third category of fat has been elucidated like "beige" or "brite" fat, which is a unique type of fat distinct from the other two types of fat [21]. Brite adipocytes are not from the same cell lineage like brown fat, they are from either trans differentiation of mature white adipose or unique precursor in white cells [22].

Beige cells could express abundant UCP1 and a broad gene program that is unique, and they can switch between an energy storing and energy releasing phenotype depends on environmental cues and conditions [23]. These brite cells have very low basal UCP1 expression like white adipose when unstimulated, while they will have turned to have a thermogenic profile similar to brown fat once stimulated [24]. Compared to classical brown fat, these cells have a coinciding but distinct gene pattern. Brite cell (which with low basal UCP1 expression and low uncoupled respiration like white fat) can be unstimulated by

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the stimulation with β adrenergic agonists, with the elevated levels of UCP1, while uncouple respiration to those of brown fat or above [25]. Brite cells are bifunctional; they can be fit for energy storage in absence of thermogenic stimuli, and can turn back to an energy releasing mechanism when necessary [26]. Moreover, brite fat contains a higher proportion of M2 macrophages, demonstrating a lean towards a healthier metabolic phenotype [27].

There are elements could induce browning of white adipocytes such as thyroid hormone T3, irisin, and cardiotrophin [28-30]. Irisin is a hormone released by the muscle during exercise, and the irisin response may be beige specific [29]. Brown and brite adipose tissues have a very different genetic profile, but UCP1 is shared by both types even the amount is variable, because brite tissue can change UCP1 expression it needed [31].

PRDM16 is a significant transcriptional factor for brown adipocytes [32], and it was shown to be a determining factor in the muscle/brown adipose Myf5+ lineage [33]. The cells become muscle when PRDM16 is knockout and become brown adipose when it overexpressed [31].

Moreover, all types of tissue can release adiponectin, leptin, and resistin [34]. Leptin endorses inflammation [35], adiponectin is approvingly adipose specific, and resistin is highly white adipose specific which could link obesity to insulin resistance [36]. Brown adipose may have the same adipokines as white adipose with a lower concentration, and it also has its own set of "batokines" such as IL-6 [37], which indicated by the study that ablation of brown adipose has a much higher influence on systemic metabolism than simply UCP1 deletion.

There are very noticeable differences between white adipose and brown adipose, and brite adipose as a completely separate category on its own with a different set of gene, which the ability to switch between an energy releasing and storing phenotype, may hold a potential key to the obesity problem currently facing the world.

Obesity and Cardiovascular Diseases

The increase in the prevalence of obesity represents a worldwide phenomenon and is correlated with several metabolic and cardiovascular diseases. Based on several researches, obesity was emphasized to be the central role in the development of metabolic syndrome, which is characterized with obesity, hypertension, hyperglycemia and hyperlipidemia, as well as to increase the susceptibility for cardiovascular disease and diabetes [38]. Obesity, which contain more and larger adipocyte, is showed with increases chronic low-grade inflammation, and is the key point in the proliferation of interrelated hyperglycemia and endothelial dysfunction due to elevated lowdensity-lipoproteins (LDL) and oxidative stress [39,40]. High level of cardiac oxidative stress is the early stage of heart dysfunction due to obesity, and it was always after insulin resistance with altered fatty acid and glucose metabolism [41]. The alternations of electron transport chain proteins related to mitochondrial ATP production [42], and the etiology of obesity is linked to the imbalance in energy consumption and expenditure [43], leading to a decrease of the efficiency of cardiac work. Most of obese patients without hypertension will have early segmental systolic and diastolic dysfunctions even the global function is normal [44], and abnormal left ventricular energy metabolism has been detected. In obesity patients which developed into heart failure, left ventricular (LV) was observed increasing in both chamber size and wall thickness (LV hypertrophy), which leads to diastolic dysfunction and cardiac ischemia in obesity, and pericardial fat is significantly associated with LV diastolic dysfunction [45]. Therefore, obesity is a risk factor for cardiovascular morbidity and mortality, as it is associated with alterations in cardiac structure [46], and left ventricular changes [47], resulting into left ventricular hypertrophy.

Heart and adipose tissue have a connection in regulation of energy metabolism [48]. Leptin was identified as an adipocyte-secreted hormone, which functions as a peripheral signal to communicate the organism's energy reserve [49]. While the heart is also included in energy network through the regulation of cardiac hormones natriuretic peptides identified as blood pressure control [50]. The natriuretic peptides stimulate triglyceride lipolysis in adipoctyes, promote uncoupling of mitochondrial respiration and thermogenesis in brown adipocytes via p38 MAPK [51].

Moreover, we know that BMI and fat mass are main independent causes of plasma nephroblastoma overexpressed (NOV) concentration, and NOV is a circulating protein that is also detected in diverse human tissues including the adrenal cortex, central nervous systems, kidney, musculoskeletal, heart and blood vessels [52], indicating the importance of NOV during cardiac development and vascular homeostasis [53]. NOV belongs to CCN family, which composed of six members, NOV is CCN3 {PMC5055498}. Studies found out there is a significant gender effect on plasma NOV concentration with women displaying a higher level of circulating NOV compared to men [52]. And it indicated that the plasma lipid profile was connected to plasma NOV after adjustment for gender. It is notable that plasma NOV is also related to physical activity, which also control plasma triglycerides [54]. Moreover, Wnt1 inducible signalling pathway protein 1 (WISP-1/CCN4) is a novel adipokine, which is upregulated in obesity, and induces a pro-inflammatory response in macrophages in vitro [55]. Wnts are signaling proteins that could control diverse biological processes, such as cell proliferation and angiogenesis [56]. While Wnt1 is secreted from primary human endothelial cells, and Wnt1/b-catenin signalling stimulates angiogenesis [57]. Angiogenesis is vital for the restoration of the blood supply to the infarct area, and it has influence on the revascularization of the perinecrotic area leading to the recovery of viable myocardium [58]. While noncanonical Wnt signalling like wnt5a, upregulated in human visceral fat compared with subcutaneous fat in obese individuals, contributes to obesity-associated metabolic dysfunction by increasing adipose tissue inflammation [59,60]. And as NOV, wnt and inflammation are linked, it is important to keep the balance so as to treat obesity and associated diseases.

Now these days, obesity becomes one of the new health problems especially when it occurs in cardiovascular diseases and browning the cardiac and peri-vascular adipose tissue is important in modulate cardiovascular risk [61] (Figure 1).

Mitochondrial Function and ROS in Obesity Induced Cardiovascular Diseases

Mitochondria in obese indexes showed decreased bioenergetics capacities and fatty acid oxidation, leading to lipid accumulation to muscular tissue [62]. And mice fed with high fat diet showed respiratory capabilities impaired by mitochondrial dysfunction [63], and the mitochondrial deficiency caused by insufficient antioxidant defences and overproduction of ROS become a potent source for energy-dependent disturbances such as inflammation [64]. ROS can be generated from many sources, such as nitric oxide (NO) synthases, cytochrome P450 enzymes and nicotinamide adenine dinucleotide phosphate (NADPH) oxidases [65]. ROS act as a signalling molecule in regulation of cellular events such as cell growth. However,





overexpression of ROS is associated with cell dysfunction and death [66], and mitochondria have been suggested to be the major intracellular site of ROS production [67]. While oxidative stress in mitochondrial dysfunction lead to downregulation of NO synthesis via upregulation of angiotensin II and 20-HETE, induces hypertension and vasoconstriction, oxidizes both low and high-density-lipoproteins, facilitates adipocyte differentiation and promotes inflammatory signalling [68-70].

Mitochondria are the primary source of intracellular energy, efficient mitochondrial function is critical [71]. The function of mitochondrial network depends on quality control, referring to fusion and fission. Mitofusin 1 and 2 (Mfn1 and 2) facilitate the mitochondrial fusion process [72], while COX-1 is related to the mitochondrial oxidative phosphorylation (OXPHOS) [73]. Brown adipose tissue is specialized to expend energy as heat by uncoupling respiration with its unique mitochondrial membrane embedded protein uncoupling protein-1 (UCP1), a process known as nonshivering thermogenesis [26], which increases heat production through an uncoupling oxidative metabolism from ATP production [74].

Proper mitochondrial function is necessary in tissues and organs in high energy demand, and multiple studies have indicated that mitochondria play a vital role in energy production, extending furthermore into thermogenesis, fatty acid oxidation, heme biosynthesis, as well as cell signalling [75,76]. The multifactorial nature of mitochondrial biology parallels the numerous pathophysiological fluctuations underlying metabolic detriments thought to develop as a result of mitochondrial dysfunction and disturbances in energy homeostasis [75,77].

Uncoupling of the electron transport chain in dysfunctional mitochondria results in overproduction of ROS, downregulation of ATP, extensive cell damage, and apoptosis of cardiomyocytes [78,79]. And in turn, the unbalance of nutrient and oxygen supply and undergo metabolic adaptation in cardiovascular diseases, which mitochondria is extremely sensitive to, leading to a progressive reduce of the mitochondrial function and loss of mitochondria structural integrity, associated with abnormalities in the respiratory chain and ATP synthesis and increase oxidative stress [80]. And the oxidative stress will make adipocyte into white fat, and deregulation of this process with failing heart and damaged mitochondria will make the situation more adverse.

Another contributor to adipose dysfunction is the NAD+/ SIRT pathway [81,82]. Several studies represented the negative correspondence of SIRT expression with adiposity, insulin resistance, as well as inflammation and impaired mitochondrial quality control [83,84]. The downregulation of SIRT1 has been exhibited to subsequently decrease AMP-activated protein kinase (AMPK) [85], which acts as a vital role in the regulation of energy balance, could induce catabolic cellular states and promote fatty acid oxidation by modulating mitochondrial activity [86,87]. Whereas inhibition of AMPK could decrease adiponectin level while promote insulin resistance, increase abdominal obesity, upregulate LDL cholesterol, and reduce mitochondrial biogenesis [88-90], and the dyslipidemia subsequent from mitochondrial dysfunction may further deteriorate insulin resistance via altered fatty acid metabolism [91], which deteriorate obesity and associated metabolic diseases (Figure 1).

Epoxyeicosatrienoic Acids (EETs) Synthesis

Epoxyeicosatrienoic acids (EETs) are lipid mediators, which derived from arachidonic acid (AA) by CYP 450 pathway [92]. It could regulate blood pressure, inflammatory and glucose homeostasis [93], while it could be further hydrolyzed to lass active diols by the enzyme soluble epoxide hydrolase (sEH) [94].

Arachidonic acid (AA) is the most biologically relevant omega-6 polyunsaturated fatty acid (PUFA) and is present in phospholipids of the cell membrane, and it is able to constitute the backbone of a triglyceride, and the release of AA from phospholipids is achieved through the activity of the enzyme phospholipase A2 (PLA2) [95]. Eciosanoids are fatty acid metabolites derived from PUFAs. AA is subject to three metabolic pathyways: cyclooxygenase (COX) is responsible for the prostaglandins, and lipoxygenase (LOX) results in the leukotrienes, while cytochrome P450 (CYP) constitute a major metabolic pathway [96,97]. AA is oxidize by the CYP enzyme into two different reactions: olefin epoxidation, which generates four epoxiyeicosatrienoic acids (5, 6-EET, 8, 9-EET, 11, 12-EET and 14, 15-EET) and hydroxylation (20-HETE) [98,99]. In human, CYP2C9, CYP2C19 and CYP2J2 are the major enzymes that convert AA to EETs, while 11, 12- and 14, 15-EETs are the main products [100,101].

EETs play a vital role in regulating physiological and pathophysiological processes, which formed endogenously in various tissues and exert potent biological effect on cellular functions [102]. However, via the activity of sEH, EET is converted into dihydroxyeicosatrienoic acid (DHET), which has a decreased activity [103]. sEH is present in many mammalian tissues, including the myocardium, adipose, liver, kidney and blood vessels [93]. The C-terminal domain of soluble epoxide hydrolase (sEH) enzyme is responsible for the hydrolysis of EETs, whereas the N-terminal domain has lipid phosphatase activity [71]. Inhibition of she results in EET accumulation and retention in tissues [104], and do benefit to cardiovascular diseases [103] (Figure 2).

Heme Oxygenase and its Derived Productions

Studies have indicated that EET could upregulate heme oxygenase-1(HO-1) activity and expression [97,105], which offers protection of vascular and regulation of adipocyte formation [106]. The mechanism by which EET increases HO-1 could be related to the increase in BACH1 which is a suppressor of HMOX1, regulated by EET through glucocorticoid and AP-1 binding sites [107], which subsequently increase HO activity.

HO, an essential stress response protein, has two isoforms: HO-1(inducible) and HO-2(constitutive) [108], can be metabolized into bilirubin, carbon monoxide and iron [109], so as to decrease the injury

from heme and ROS [110,111]. HO-1 can be induced by an extremely wide diversity of drugs and chemical agents, such as aspirin, statins, eicosanoids like EET and free metals [108].

Carbon monoxide (CO) has been studied in both cell and human physiology, and has many benefit including increasing mitochondrial function in mesenchymal stem cells [112,113], improving sepsis in stem cells [114], and modulating of inflammation [115,116]. Moreover, CO can act as a gasotransmitter, a potent regulator of vascular homeostasis, decreases vasoconstriction and stimulates vasodilation, and instantaneously increases insulin secretion [117].

The HO-1 derived bilirubin can prevent oxidant-mediated cellular damage and attenuate oxidant stress [118], and have the potential benefit to protect obesity and metabolic syndrome through increasing insulin sensitivity and suppressing chronic inflammation [119]. Moreover, studies indicated that bilirubin can inhibit the oxidation of low denisity lipoproteins (LDL) [120], and is associated with a decreased risk for coronary artery disease in humans [121,122].

The degradation of heme results in the dispersion of iron, which is known to dispose the production of ROS [123]. The upregulation of HO-1 correspondences a concurrent increase in anti-oxidant ferritin [124], the toxic effects of iron-mediated oxidative damage couple a parallel rise in ferritin concentrations as a means for anti-oxidant and anti-inflammatory response [125,126] (Figure 3).

Impact of EET and HO-1on Adipogenesis and Obesity

A large evidence suggested that adipocyte oxidative stress is fundamental reason in the pathogenesis of obesity, and it can cause dysregulation of inflammation-related adipocytokines, leading to cardiovascular complications [115]. Adipocytes belong to mesodermal origin and bone marrow stromal cells, and serve as a reservoir for the preadipocytes recruitment and generation [127,128]. It is well known that during adipogenesis, upregulation of ROS will differentiate adipocytes produce much more ROS than preadipocytes [129]. Morphologically, this adipocyte differentiated under oxidizing conditions are investigated by Oil Red O detection of lipid accumulation [130].

The effect of EETs on adipogenesis and the associated signaling cascades involved in adipogenesis including HO-1, adiponectin, AMPK, and pAkT have been studied in recent years [131]. And it was known that the HO system act as an cellular antioxidant defense system in obesity

and diabetes [109], whereas epoxides have anti-inflammatory [92] and antihypertensive effects [132] and EET suppression could contribute to hyperlidemic states. It is indicated that human MSC-derived adipocytes could express CYP 2J2 and produce EETs [105], which decreased adipocyte differentiation via an increase in HO-1 expression paralleled with a decrease in PPARy, C/EBPa and FAS levels, suggesting that EETs can regulate lipid metabolism in developing preadipocytes so as to inhibit and/or delay adipocyte differentiation [133]. And EET could increase adipocyte proliferation by phenotypically reprogramming adipocytes to decrease the expression of PPARy and mesodermspecific transcript (MEST), and form a module to switch to genetically reprogram the adipocyte phenotype to express less MEST and prevent hypoadiponectinemia [57,131,133]. Moreover, treat human MSCs with adipogenic media for two-week and bind with an additional 10 mM of glucose (hyperglycemic conditions) lead to an increase in adipogenic differentiation. While adipogenic differentiation decreased with the treatment of CoPP, demonstrating the predominance of preadipocytes, indicating the influence of HO-1 in the regulation of genes controlling adipocyte differentiation [134]. Therefore, MSC-derived adipocytes are not only a production of environmental stimuli and the right balance in ROS, but also the interaction of HO and EET contribute to adipogenesis and adipocyte function (Figure 4).

Impact of EET and HO-1on Obesity Induced Hypertension

Obesity is known to relate with an increase in sodium retention and volume expansion, which are the risk factors contribute to hypertension [135]. And Theken studied the relevance of EETs with hypertension in obese patients with or without CAD, and indicated that patients with obesity are associated with a low level of EET, and the CYP epoxygenase activity is suppressed and sEH metabolic activity is increased [136]. And it was studied that treat obese HO-2 KO mice with EET agonist could increase EET levels in kidney and vascular, with lowered blood pressure and decreased body weight gain and reduced subcutaneous and visceral fat, associated with an increase insulin sensitivity [106].

EETs, both autocrine and paracrine mediators, play an important role in the cardiovascular and renal systems [137]. Due to the strong vasodilatory actions within the endothelium and kidney, EETs act as an endothelium derived hyperpolarizing factor (EDHF) [138]. EETs dilate the preglomerular arterioles through Ca (2+)-activated K (+) channels in renal smooth muscle cell and hyperpolarize smooth muscle cells [139]. And EETs is also the second messengers for many paracrine and





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hormonal agents such as vasodilatory substances, namely endothelin, angiotensin, NO, and PGI2.

It is known that the pleiotropic effects of EET could upregulate HO activity, and HO system could increase EET and adiponectin [140]. The antioxidant action of HO metabolites is related to extension of small adipocyte which associated with an increased adiponectin and its downstream signals including pAMPK, peNOS and induced NO bioavailability [141,142]. Upregulation of these pathways could improve vascular function and dominate of hypertension, and the increase in biliverdin from HO-1 could prevent EET from degradation by ROS and adiponectin [121]. Moreover, HO-derived CO is a vasodilator, which could regulate basal and constrictor induced vascular tone in blood vessels [143]. And study showed that chronic CO treatment with a CO-releasing molecule (CORM) could protect vascular function [144]. High fat diet rats treated with CORM-A1-derived CO showed decrease in body fat, insulin and fasting blood glucose, while with increase in oxygen consumption and heat production [145].

These results establish the interdependence of EET and HO-1 in preventing obesity and associated hypertension, activation of these pathways could protect the vasculature against damage associated with vascular disease (Figure 4).

Impact of EET and HO-1on Obesity Induced Endothelial Dysfunction and Arthrosclerosis

Endothelial cell dysfunction, verified by the diminished expression of CD31+ and/or thrombomodulin (TM) [146], located within atherosclerotic blood vessels, indicating an early feature of chronic cardiovascular diseases [147], and related to extra levels of ROS [148]. The abnormality of CD31+ gene was showed in the pathogenesis of both atherosclerosis and myocardial infarction, whereas a reduction in plasma TM was linked to an increased risk of myocardial infarction [149]. Reduced expression of CD31+ and TM in endothelial cells indicated endothelial cell death, associating with the progression of atherosclerotic heart disease [146], and the restoration of their expression could benefit atherosclerosis and myocardial infarction [149].

Inflammation, as well as endothelial cells injury, is critical contributor to atherosclerosis [150], the upregulation of IL-6 and TNF- α induced by chronic stress and angiotensin II leading to atherosclerosis [151]. Studies indicated that HO-1 suppress the proliferation of VSMCs through CO release [152] and inflammation



endothelial function, so as to prevent atherosclerosis.

alteration [153]. Monocyte chemoattractant protein-1(MCP-1) as a strong chemoattractant for macrophages turn mononuclear cells into macrophages, and nuclear factors adjust the MCP-1 transcription and expression in atherosclerotic plaque [154]. And HO-1 pathway could regulate MCP-1 secretion and ICAM-1 expression. Products of HO metabolism of heme, bilirubin and CO, upregulate EC-SOD and decrease inducible enzymes as seen in iNOS and peroxynitrite generation, and act as a countervailing influence to hyperglycemia-mediated injury in endothelial cells and in sloughing. We have also shown that decrease in iNOS and increase in EC-SOD mediated by HO-1 derived CO and biliverdin/bilirubin exerts beneficial actions in vascular protection [155].

Atherosclerosis, a progressive disease characterized by the accumulation of lipids and fibrous components in the arteries, is the major cause of mortality worldwide [156]. Widespread epidemiological evidence strongly verified the connection between the plasma cholesterol profile and the development of atherosclerosis. It indicated that high-density lipoprotein (HDL) is effectively protective against atherosclerosis, whereas low-density lipoprotein (LDL), especially oxidized (ox-) LDL, act as a trigger of atherosclerosis [157]. LDL stimulates the inflammatory response and formation of foam cells, which are fat-laden macrophages that form the fatty streaks in plaques [158]. Soluble epoxide hydrolase inhibitors could increase EET levels and reduce atherosclerotic lesion formation in mice deficient in apolipoprotein (apo) E or LDL receptor [159]. apoE-/- mice displayed advanced lesions and an obvious upregulation in the LDL/HDL ratio, and administration of sEH inhibitor extensively shrunk serum LDL concentrations and secretion of inflammatory factors [160]. And recent studies in adipose tissue have confirmed that the administration of sEH inhibitor (t-AUCB) develops the CD36-mediated recognition and ox-LDL degradation and improves cholesterol efflux by inducing ATP-binding cassette A1(ABCA1) expression [161], and ABCA1 subsequently enhanced plasma HDL concentrations and reversed cholesterol transport [162], which may also play a part factor of antiatherosclerotic effect. Meanwhile EETs accumulated in the occurrence of the sEH inhibitor and activated peroxisome proliferator-activated receptor (PPAR) y, which may be involved in sEH inhibition-induced CD36 and ABCA1 expression [162]. Moreover, EET upregulation could induce HO-1 expression and activity, altered inflammation and reverse endothelial function, so as to prevent atherosclerosis (Figure 4).

Impact of EET and HO-1 on Obesity Mitochondrial Function

Marked mitochondrial dysfunction has been observed in myocardial cells in cardiovascular diseases [163], and improve mitochondrial function is associated with an improvement in myocardial function. More than 50% of disease-related mutations in mitochondrial DNA result in cardiomyopathy in humans, and targeted mutations disturbing fatty acid transport and oxidation, high-energy phosphate transport and shutting. Protection from mitochondrial ROS, and mitochondrial DNA proofreading activity all cause profound cardiac dysfunction [164].

As the intermediary between EET and HO-1, PGC-1 α could establish homeostatic energy metabolism, normalize mitochondrial quality, upregulate mitochondrial biogenesis and respiration, adaptive thermogenesis and gluconeogenesis as well as many other metabolic processes through the activation of AMPK and SIRT [165,166]. The upregulation of this energy sensing network can stimulate the ROS formation reduction [167]. And PGC-1 α is vital to the amputation of mitochondrial ROS through regulation of the expression of plentiful enzymes that provoke ROS [168,169]. Novel statistics indicated that PGC1 α -knockdown mice have an increased exposure to neurodegeneration and hippocampus-situated oxidative stress [170]. It was reported that endogenous SIRT3-deficient showed an increased intracellular ROS, while upregulation of SIRT3 could alleviated ROS in brown adipocytes and reestablished uncoupling proteins UCP thermogenesis [87]. UCP1 and UCP2 from adipose tissue could control of energy as heat and affect energy metabolism efficiency [25]. Superior SIRT1 levels could reduce NF- $\kappa\beta$ inflammatory transcription and result in increase the level of PGC-1 α [85,171], which act as a potent suppressor of ROS through the stimulation of ERR α and the consequent generation of ROS detoxifying agents GPx1 and SOD2 [172].

PGC1a could stimulate biogenesis and restore mitochondrial quality control through the transcriptional co-activation of nuclear respiratory factor (NRF)-1/2 [173]. NRF-1 and NRF-2 enhance mitochondrial transcription factor A (TFAM) localization to the mitochondria, resulting in augmented mitochondrial biogenesis and respiration, so as to a simultaneous rise in cytochrome C and ATPase [174]. TFAM elevation is important in the PGC1a-facilitation of mtDNA replication [164]. Based on the role of mitochondrial dysfunction played in ROS production underlying metabolic syndrome, PGC1a and concomitant SIRT have major implications in the therapeutic mitochondrial targeting of obesity, DM, and cardiovascular dysfunction indexes [175,176].

Impact of EET and HO-1 on Obesity Induced Cardiovascular Diseases

As Theken proved that obesity individuals have low level of EETs, and the further study presented that both obese and non-obese CAD patients had significantly higher plasma EETs with a higher epoxide/ diol ratios, demonstrating that CYP epoxygenase and sEH metabolic function are altered in patients with established atherosclerotic cardiovascular disease [136]. Treatment with EET agonist 11-(nonloxy) undec-8(z)-enoic acid (NUDSA) increases adipose tissue levels of EET and HO-1, as well as serum adiponectin in high-fat diet rats, associated with a decrease in blood pressure, subcutaneous and visceral fat content and inflammation factors (TNF- α and IL-6) [177]. EET was known to upregulate the expression of wnt1 canonical signaling cascade, attenuate cardiac dysfunction and improve angiogenesis [58]. And we know that EET could decrease NOV expression in cardiac and adipose tissue, which simultaneously increase PGC-1a mediated downstream signaling, enhancing mitochondrial function and energy metabolism, and preventing the development of cardiac remodeling in cardiomyopathy [178].

EET could increase osteoblast differentiation whereas decrease adipogenesis differentiation [140,179], which was further, supported by the observation that CoPP affected adipocyte differentiation in adults rats and developed in bodyweight loss without different food consumption [180]. There are many pharmacological agents which could increase HO-1 levels and decrease adiposity like the beneficial effect of CoPP, named as apolipoprotein A1 mimietic peptides L-4F, EET, and peroxisome proliferator-activated receptor (PPAR), which also lead to a decrease in visceral subcuntaneous fat and an increase in insulin sensitivity [181-184], associated with the reduction of large adipocytes number and a lift of smaller healthy adipocytes [133,185,178].

Enhanced EET has been indicated to reduce myocardial fibrosis and inflammation, so as to reduce hypertrophy and improve diastolic

function of metabolic syndrome rats [186]. Aortic endothelial function, peNOS expression and adipose tissue markers of energy homeostasis such as pAMPK, pAkt, Sirt1, and fatty acid synthase (FAS), are restored in animals with NUDSA treatment [177,187,188]. As EET can upregulate HO-1, and we found out that CoPP induction of HO-1 decreased circulating free fatty acids and C-reactive protein, increased adiponectin, through the activation of AMPK-P13K-eNOS pathway [185,189], and adiponectin concentration in human plasma is lower in patients with clinical manifestations of CAD than in BMI- and age-adjusted control subjects independent of other risk factors {1508} [190], while high adiponectin adiponectin is related to a lower risk of CAD in male diabetic patients{1567} [191] It should be highlight that HO-1 could not increase adiponectin directly, but through HO-1 mediated antioxidant mechanism with a decreased in heme associated with increase in superoxide dismutase, which reduce ROS levels and increase adiponectin[176,181,192,193].

And PGC-1a has been addressed to control many aspects of oxidative metabolism, including mitochondrial biogenesis and respiration through coactivation of many nuclear receptors [194]. In cardiac cells PGC-1a could induce abundantly of genes, programming for critical enzymes in major metabolic programs required for high-efficiency ATP production [195,196], and it could upregulate over 70% of the subnets in the mitochondrial electron transport chain and the ATPase complexes, and markedly increase fatty acid oxidation [197,198]. PGC-1a expression is stimulated by ischemic conditions in cell culture, and a PGC-1a transcriptional coactivators have now arose as a dominant regulator of mitochondrial biology in the heart, and the expression of PGC-1a is suppressed in numerous heart failure diseases [199,200]. Pathologically hypertrophied heart induced by the coarctation of the aorta, have a decreased expression of both PGC-1a and its target genes of fatty acid oxidation and oxidative phosphorylation [201,202]. And after PGC-1a knockout, the heart isolated from the mice showed decreased fatty acid oxidation and heart function reduction, and the myocardial fibers presented decreased ATP synthesis rates and ATP production efficiency [203,204]. Structural analysis demonstrated an abnormal mitochondrial cristae density and cytoplasmic accumulation of lipids, indicating a reduction in fatty acid consumption paralleled with lipotoxicity from increase fatty acid uptake [203].

Moreover, PGC-1 α also appears to mediate EET-induced HO-1 activity upregulation [178,205], this mechanism has been observed categorically in adipose tissue suggesting the activity of EET in the vasculature and in the myocardium in cardiovascular diseases (Figure 5).

Soluble Epoxide Hydrolase Inhibitor and Clinical Therapy

Soluble epoxide hydrolase (sEH) could degrade epoxides into corresponding diols, which are substantially less active than the original compound. Inhibition of sEH leads to EETs accumulation and retention in various tissues [206,207], which have anti-inflammation function, vasodilatory activity and antihypertensive action, while it can promote fibrinolysis and inhibit platelet aggregation [98, 106,208-210]. While induction of CYP-epoxygenases, administration of EET analogs, or inhibition of sEH all result in upregulation of EET concentration, and could benefit cardiovascular diseases [103]. sEH is upregulated in obesity and associated diseases, with a downregulation of EET, and decreased level of EET may lead to more sEH with lass P450epoxygenases [211], so sEH inhibitors have been extensively studied for the potential value on cardiovascular therapy.

In vitro studies, administration of sEH inhibitor in the MSC culture

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indicated a decreased adipogenesis and adiposity associated with an increased level of EET, and it was known that EET can decrease MSC-derived adipocytes [133]. And MSCs, act as pleotropic cells, can differentiate to adipocytes or osteoblasts as a result of crosstalk by specific signaling pathways including HO-1/2 expression [179]. Adipocyte stem cells treated with AUDA showed decrease in adiposity with an increased effectiveness of EET [133].

In vivo studies have indicated that sEH inhibition can reverse inflammation [212], reduce the development of atherosclerosis in apoplipoprotein E knowout mice [213,214], and it turned out to lower blood pressure in angiotension-dependent hypertension [215,216]. And it was studied that Glu287Arg substitution in sEH genes, in humans with familial hypercholesterolemia, is related with an increased plasma cholesterol level [217], and sEH deficiency or inhibition in high fat diet mice has been linked to the reduction of liver steatosis and attenuation of endoplasmic reticulum stress in adipose tissue [97,218]. Weighty testing of sEH inhibitors indicated they are highly selective for sEH, lack significant toxicity, and have potential for use in humans [219,220]. Moreover, the connotation between sEH and vascular phenotypes has been investigated in 106 patients with stable CAD, and there was a significant inverse relationship between 20-HETE levels and brachial artery flow-mediated dilation, paralleling with an inverse relationship between sEH function and a combination of MCP-1 and cellular adhesion molecule score, indicating that sEH therapy might be effective in human vascular dysfunction [221].

GSK2256294A is a reversible binding inhibitor of isolated recombinant human sEH, exhibiting a good preclinical pharmacokinetic profile with high oral bioavailability in mice [222]. Concurrently, it has been detected to reduce the inflammatory response induced by repetitive exposures to cigarette smoke, indicating its dose-dependent effect with sustaining for up to 24 hours [223]. And it has the promise to become one of the vital treatment for obesity and associated diseases.

Conclusion

The obesity associated cardiovascular diseases relies heavily on EET system, implementation of the EETs in pharmaceutical synthetics may comprise a novel avenue for the treatment of metabolic insults through the restoration of mitochondrial integrity and associated adiposity. The capacity of EET with induced HO-1 could decrease inflammatory

cytokines, angiotensin II and oxidative stress, finally lead to heart function improvement, rendering it a noticeable target for clinical application. The pleiotropic effect of EET synchronously contrast the multifactorial etiology of obesity and associated cardiovascular diseases, and it acts like a master regulator of multiple components of myocardial energetics through the crosstalk with both PGC-1 α and HO-1. sEH inhibitor therapy has been test *in vitro* and *in vivo* in animals, as well as in human preclinical trials to value the adverse effects of EET. As such, the application of the EET-HO-1 module in a clinical setting may potentially serve as a potent approach to combatting the unabated global epidemic of obesity and associated cardiovascular diseases.

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