



Abnormal kynurenine pathway of tryptophan catabolism in cardiovascular diseases

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Received: 17 November 2016 / Revised: 26 February 2017 / Accepted: 8 March 2017 / Published online: 17 March 2017
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Abstract Kynurenine pathway (KP) is the primary path of tryptophan (Trp) catabolism in most mammalian cells. The KP generates several bioactive catabolites, such as kynurenine (Kyn), kynurenic acid (KA), 3-hydroxykynurenine (3-HK), xanthurenic acid (XA), and 3-hydroxyanthranilic acid (3-HAA). Increased catabolite concentrations in serum are associated with several cardiovascular diseases (CVD), including heart disease, atherosclerosis, and endothelial dysfunction, as well as their risk factors, including hypertension, diabetes, obesity, and aging. The first catabolic step in KP is primarily controlled by indoleamine 2,3-dioxygenase (IDO) and tryptophan 2,3-dioxygenase (TDO). Following this first step, the KP has two major branches, one branch is mediated by kynurenine 3-monooxygenase (KMO) and kynureninase (KYNU) and is responsible for the formation of 3-HK, 3-HAA, and quinolinic acid (QA); and another branch is controlled by kynurenine amino-transferase (KAT), which generates KA. Uncontrolled Trp catabolism has been demonstrated in distinct CVD, thus, understanding the underlying mechanisms by which regulates KP enzyme expression and activity is paramount. This review highlights the recent advances on the effect of KP enzyme expression and activity in different tissues on the pathological mechanisms of specific CVD, KP is an inflammatory sensor and modulator in the cardiovascular system, and KP catabolites act as the potential biomarkers for CVD initiation and progression. Moreover, the biochemical features of critical KP enzymes and principles

of enzyme inhibitor development are briefly summarized, as well as the therapeutic potential of KP enzyme inhibitors against CVD is briefly discussed.

Keywords Tryptophan · Kynurenine pathway · Aortic aneurysm · Atherosclerosis · Aging · Obesity · Diabetes

Abbreviations

1-MT	1-methyl tryptophan
3-HAA	3-hydroxyanthranilic acid
3-HK	3-hydroxykynurenine
AA	Anthranilic acid
AAA	Abdominal aortic aneurysm
AHR	Aryl hydrocarbon receptor
Ang II	Angiotensin II
DCs	Dendritic cells
HFD	High-fat diet
IDO1	Indoleamine-2,3-dioxygenase 1
IFN- γ	Interferon-gamma
IR	Insulin resistance
KA	Kynurenic acid
KAT	Kynurenine amino-transferase
KMO	Kynurenine-3-monooxygenase
KYNU	Kynureninase
KP	Kynurenine pathway
Kyn	Kynurenine
MMPs	Matrix metalloproteinases
oxLDL	Oxidized low-density lipoprotein
pDCs	Plasmacytoid dendritic cells
PLP	Pyridoxal 5'-phosphate
QA	Quinolinic acid
TDO	Tryptophan 2,3-dioxygenase
Trp	Tryptophan
VCAM-1	Vascular cell adhesion molecule-1

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VSMC Vascular smooth muscle cell
XA Xanthurenic acid

Introduction

The essential amino-acid tryptophan (Trp) is either used for protein synthesis or metabolized to a variety of bioactive molecules via the serotonin pathway or the kynurenine pathway (KP) which accounts for the metabolism of more than 95% of dietary Trp [1] that involves several vitamin B6- or B2-dependent enzymes (Fig. 1). KP is the major pathway for Trp catabolism in human [2, 3], animals [4, 5], and some fungi and bacteria [6, 7]. Generation of KP catabolites is triggered by the enzymes tryptophan 2,3-dioxygenase (TDO) [8], which is mainly expressed in liver and brain with high substrate specificity for L-tryptophan [9], or indoleamine-2,3-dioxygenase (IDO), the first and rate-limiting enzyme catabolizing both D- and L-tryptophan [10]. In general, TDO mediates basal KP, whereas IDO may control KP under disease conditions [11]. IDO has two isoforms IDO1 and IDO2 [12, 13], and IDO1 is widely expressed at low levels under normal conditions and the major isoform contributing to Trp degradation [14]. IDO2 is primarily expressed in mouse liver and kidney [12], as well as human liver, spleen, placenta, kidney, and brain, but not in the heart [15]. IDO1 is inducible, while IDO2 appears

to be basally expressed [5]. IDO catalyzes the conversion of Trp to N-formylkynurenine, which is then catabolized to the pivotal KP intermediate kynurenine (Kyn). From Kyn, one branch of the KP leads to the formation of kynurenic acid (KA) or xanthurenic acid (XA) catalyzed by kynurenine amino-transferases (KATs, KAT I–IV), one family of the key enzymes in KP (Fig. 1); and another branch results in the biosynthesis of oxidized nicotinamide adenine dinucleotide (NAD⁺) via 3-hydroxykynurenine (3-HK) and quinolinic acid (QA) [16]. Within this branch, another key enzyme kynureninase (KYNU) directly catalyzes the hydrolysis of Kyn or 3-hydroxykynurenine (3-HK) to form the anthranilic acid (AA) or 3-hydroxyanthranilic acid (3-HAA), respectively [17, 18] (Fig. 1). Human KYNU presents a 20-fold higher affinity for 3-HK than for Kyn, favoring the generation of 3-HAA over AA [19]. KP is well controlled under physiological conditions but altered as part of the activated immune response. Dysregulated KP has been linked to various diseases, including neurodegenerative disorders (such as Huntington, Alzheimer, and Parkinson disease) [8], multiple sclerosis [20], depression [21, 22], schizophrenia [23, 24], diabetes [25], cancer [26, 27], and inflammatory bowel disease [28]. Emerging evidence indicates that catabolites in KP of Trp catabolism play critical roles in cardiovascular physiology and pathology [29–31], in addition to regulating immune system functions [2, 32] and inflammation [7,

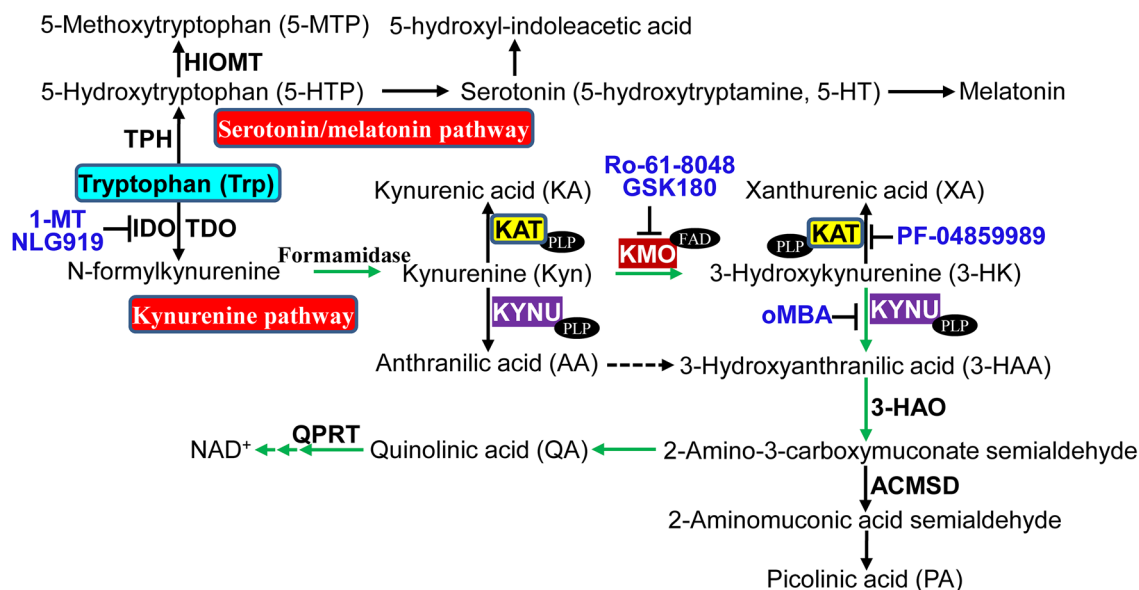


Fig. 1 Major enzymes, catabolites, and inhibitors in tryptophan metabolism. Tryptophan is metabolized via two major pathways, kynurenine pathway and serotonin/melatonin pathway. *3-HAO* 3-hydroxyanthranilate 3,4-dioxygenase, *ACMSD* 2-amino-3-carboxymuconate semialdehyde decarboxylase, *FAD* flavin adenine nucleo-

tide, *HIOMT* hydroxyindole O-methyltransferase, *oMBA* o-Methoxybenzoyalanine, *QPRT* quinolinate phosphoribosyltransferase, *TPH* tryptophan hydroxylase. Refer to the text for the expanded form of abbreviations

33]. Here, we briefly review the recent research progress on the role of dysregulated KP in cardiovascular diseases (CVD) development and the regulation of CVD risk factors on KP, highlight KP functions as the inflammatory sensor and modulator in the cardiovascular system, as well as discuss some controversies and raise questions in this field.

Role of dysregulated KP in CVD

CVD is the most common cause of death in the world. Evidence indicates that KP catabolites play important roles in cardiac pathophysiology. The expression of IDO1, the first and rate-limiting enzyme in KP, is tightly controlled and confined to certain cell types, including endothelial cells (EC) [30, 34, 35] vascular smooth muscle cells (VSMC), macrophages [36], leucocytes [37], and dendritic cells (DCs) [38], all of which are present in the artery wall. Proinflammatory modulators, such as lipopolysaccharide (LPS), tumor necrosis factor (TNF), interleukin 1 (IL-1), and IL-2, induce IDO1 expression in many cell types [39]. Notably, interferon-gamma (IFN- γ) released from activated CD4⁺ T cells robustly induces IDO1 expression contributing to Trp catabolism [39, 40], where Trp is converted to immunomodulating compounds, collectively termed kynurenines [41]. Epidemiologically, tryptophan catabolites, Kyn, AA, 3-HK, and XA are reported to associate with CVD mortality [42]. 3-HAA was identified as a precursor of cinnabarinic acid, a novel endogenous ligand of the transcription factor aryl hydrocarbon receptor (AHR) [43], which may contribute to cardiotoxicity, vascular inflammation, and endothelial dysfunction [44]. Here, we will discuss the effects of major catabolites in KP on the main CVD (Table 1).

KP and atherosclerosis

Atherosclerotic cardiovascular diseases including coronary heart disease and cerebrovascular disease are the most common forms of CVD. There are several reports about abnormal KP and atherogenesis. In the Tampere Vascular Study, elevated IDO expression was observed in the macrophage-rich cores of human advanced atherosclerotic plaques [36]. IDO activity in blood has a significant positive correlation with more advanced atherosclerosis in both sexes (Table 2) [60]. These data suggest that activated KP may play an important role in atherogenesis. However, treatment with KP catabolite 3-HAA for 8 weeks significantly decreases plasma cholesterol and triglyceride levels, and inhibits the uptake of oxidized low-density lipoproteins (oxLDL) by macrophages, and the resultant inhibition of atherogenesis in hypercholesterolaemic low-density lipoprotein receptor-deficient (LDLr^{-/-}) mice [61]. Furthermore, systemic IDO inhibition by 1-methyl tryptophan (1-MT) enhances vascular inflammation with upregulation of vascular cell adhesion molecule-1 (VCAM-1) and monocyte chemoattractant protein-1 (MCP-1/CCL2), and increasing CD68⁺ macrophage infiltration in vascular area, which leads to a significant increase in atherosclerotic lesions in the aortic arch and root of high-fat diet (HFD)-fed apolipoprotein E knockout (ApoE^{-/-}) mice [45]. Interestingly, the increased VCAM-1 expression is not limited to the plaque but also found in VSMC of the tunica media. Moreover, IDO inhibition-induced acceleration of atherosclerosis and vascular inflammation can be reversed by exogenous administration of the Trp catabolite 3-HAA [45]. All these data strongly suggest that IDO-mediated KP catabolite 3-HAA blocks atherogenesis (Fig. 2). It will be important to investigate 3-HAA levels in plasma and local atherosclerotic areas of mice and to identify if endogenous 3-HAA can be used as a biomarker for atherosclerosis initiation and progression. In

Table 1 Function and CVD association of key catabolites in kynurenine pathway

Catabolites	Function	Associated disease	References
3-HAA	Activator of aryl hydrocarbon receptor (AHR), induction of Tregs from naïve CD4 ⁺ T cells, and T-cell death	Vascular inflammation, atherogenesis	[43–46]
3-HK	ROS generator, cell apoptosis	EC dysfunction	[30, 47–49]
AA	Inhibits proinsulin synthesis	T1D	[50]
Kyn	Vasodilation, neurotoxicity and brain degeneration, endogenous ligand of AHR	Hypotension, heart failure, insulin resistance	[29, 51–53]
KA	Antagonist of NMDA receptor and α 7 nicotinic acetylcholine receptor, antioxidant, AHR agonist, ligand for GPR35	Ischemic post stroke, hypertension, preeclampsia, myocardial infarction, maternal hypertension, unstable atherosclerotic plaque, T2D, schizophrenia	[54–57]
QA	Agonist of the NMDA receptor, elevates Ca ²⁺ , ROS generator, neurotoxicity	Aging, T1D	[53]
XA	Induces apoptosis, inhibits BH ₄ synthesis	Diabetes, insulin resistance	[58, 59]

NMDA N-methyl-D-aspartate, ROS reactive oxygen species. For definitions of other abbreviations, please see the main text

Table 2 Cardiovascular diseases and their risk factors associate with alternations of KP catabolites and enzymes

CVD and risk factors	Alternation of KP catabolites	Samples	KP enzymes	Samples	References
Atherosclerosis	AA↓, 3-HAA↓, KA↑	Serum	IDO↑, IDO activity↑	Plaque, serum	[36, 63]
Myocardial infarction	KA↑, 3-HK↑, AA↑, 3-HAA↑	Plasma	IDO↑	Plasma	[64, 65]
Coronary artery disease	Kyn↑, 3-HK↑	Urine	IDO activity↑	Urine	[66]
Stroke	KA↑	Serum	Active IDO1	Serum	[67]
Diabetes	KA↑, AA↑, XA↑, 3-HAA↑, Kyn↑, PLP↓	Plasma, urine	IDO↑, KAT activity↑	Plasma	[25, 50, 68–70]
Hypertension	Kyn↑, KA↓	Rat brain	Mutation in KAT I, KYNU mRNA↑, KYNU activity↓	Brainstem	[71–73]
Obesity	Kyn↑, KA↑, QA↑	Serum	IDO1↑, KYNU↑, KMO↑, KAT III↑	Omental adipose tissue	[74–77]
Aging	Kyn↑, KA↑, 3-HK↑, AA↑, QA↑	Rat brain	IDO1↑, TDO↓	Brain striatum blood	[78–82]
Smoking (nicotine)	KA↑, PLP↓	Brain	N/A		[83]

Refers to the main text for the expanded form of abbreviations

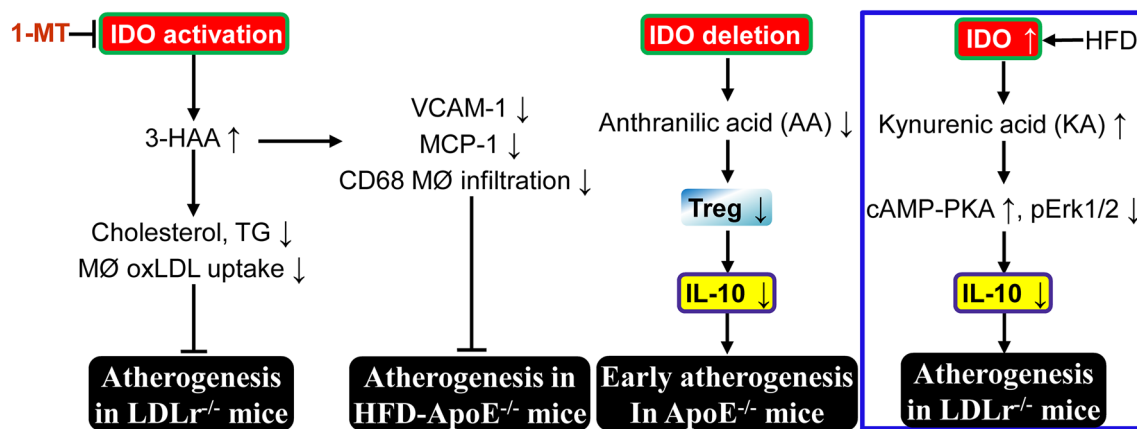


Fig. 2 Abnormal kynurenine pathway and atherogenesis. Elevation of 3-hydroxyanthranilic acid (3-HAA) or anthranilic acid (AA) by IDO activation inhibits vascular inflammation and subsequent atherosclerosis in $LDLr^{-/-}$ or $ApoE^{-/-}$ mice. However, kynurenic acid

(KA) elevation by IDO upregulation enhances vascular inflammation and exacerbates atherogenesis in $LDLr^{-/-}$ mice (blue box). $M\phi$ macrophage, PKA protein kinase A, $Treg$ regulatory T cell. For definitions of other abbreviations, please see the main text. \downarrow , inhibits

addition, 3-HK level correlates with intima media thickness of the carotid artery [62].

On one hand, IDO exerts a protective effect against atherogenesis. It was reported that IDO1 deficiency accelerates early (at 15 weeks of age) atherosclerotic lesion formation in the aortic root of $ApoE^{-/-}$ mice fed with a regular chow diet, through downregulation of the anti-inflammatory cytokine IL-10 in peripheral blood, spleen, and lymph node B cells [84]. IDO1 deletion does not significantly alter serum cholesterol levels in $ApoE^{-/-}$ mice. However, IDO1 deletion for 15 weeks increases infiltration of $CD68^{+}$ macrophages and $CD4^{+}$ T cells in atherosclerotic plaque [84]. Moreover, IL-10 production in cultured splenic B cells is stimulated by 3,4-dimethoxycinnamoyl anthranilic acid (3,4-DAA), an orally active synthetic derivative of the tryptophan catabolite AA [85], but not

Kyn. Finally, 3,4-DAA treatment inhibits inflammation and lesion formation after perivascular collar-mediated arterial injury in $ApoE^{-/-}$ mice, and decreases cytokine and chemokine production in ex vivo human atheroma cell cultures. These data imply that compensated upregulation of IDO may play protective function from atherogenesis at an early stage via AA generation (Fig. 2). Importantly, IDO1-expressing aortic plasmacytoid dendritic cells (pDCs) protect against atherosclerosis through induction of regulatory T cells [38]. In addition, IDO1 may play a critical role in maintaining plaque stability [84].

On the other hand, HFD dramatically increases IDO activity in macrophages and VSMC of aortic sinus, as well as circulating levels of KA and QA in atherosclerosis-prone $LDLr^{-/-}$ mice compared with chow diet. Furthermore, levels of Kyn, KA, and QA are higher in spleen extracts

of LDLr^{-/-} mice fed with HFD [86]. Interestingly, it was demonstrated that IDO1 deficiency inhibits atherogenesis in aortic root of LDLr^{-/-} mice fed with HFD compared with LDLr^{-/-}IDO1^{+/+} littermates via IL-10 upregulation in macrophages and FMS-related tyrosine kinase 3 (Flt3) ligand-induced bone-marrow-derived DCs [86]. Moreover, KP catabolite KA is responsible for the IL-10 reduction in macrophages via activation of the cAMP-dependent pathway and inhibition of Erk1/2 phosphorylation [86] (Fig. 2). However, deletion of both IDO1 and IDO2 does not alter IL-10 levels in macrophages in response to LPS + IFN γ compared to WT [87]. In addition, high concentrations of KA in atheromatous plaques are associated with an unstable plaque phenotype in human atherosclerotic lesions, whereas there is no detection or low levels of KA in stable fibrous plaques [86]. It is of interest to test whether endogenous KA can work as the biomarker for atherosclerosis development and plaque instability.

The reason for the discrepancy of IDO1 function in atherogenesis may be due to the different animal genetic background, different cell types involved, and specific KP downstream catabolites. Nevertheless, this contradictory evidence highlights the importance of clarifying the function and the underlying mechanisms by which IDO1 expression and activation in atherogenesis. Further investigation with tissue-specific gene knockout/knockin mouse models is needed to understand the role of these particular catabolites in different cells in atherogenesis under different disease condition.

Abnormal KP and EC dysfunction

Endothelial dysfunction is featured by increased endothelium apoptosis, impaired endothelial-dependent vasomotion, and dysregulated endothelial cell activation [88, 89]. Endothelial dysfunction accounts for a significant portion of all CVD [90]. Abnormal KP is highly associated with EC dysfunction. For example, IDO-deleted pregnant mice with C57BL/6 J background display preeclampsia featured by aortic endothelial dysfunction, renal structural damage (glomerular endotheliosis), and dysfunction (proteinuria) [91]. However, the catabolites of KP responsible for preeclampsia are unknown. In contrast, our group demonstrated that KP catabolite 3-HK mediates angiotensin II (Ang II)-induced EC apoptosis and subsequent endothelial dysfunction via NAD(P)H oxidase-derived superoxide anions *in vivo*, whereas IDO1 deletion blocks Ang II action on EC [30]. KP catabolite KA elicits the adhesion of neutrophils and leukocytes to vascular endothelium via G protein-coupled receptor 35 (GPR35) in an *in vitro* vascular flow model [92]. Thus, KA may lead to abnormal EC activation and inflammation. 3-HAA inhibits the expression and activity of inducible nitric oxide synthase (iNOS)

in macrophages [93]. In addition, KP catabolite XA is a potent inhibitor of sepiapterin reductase (SPR) [58], the last enzyme in the *de novo* synthesis of tetrahydrobiopterin (BH₄), which is an enzyme cofactor essential for the activation of endothelial nitric oxide synthase (eNOS). These results suggest that elevated XA arising out of upregulated KP could attenuate BH₄ biosynthesis and consequent EC dysfunction. On the other hand, dysregulated eNOS would increase O₂⁻ production, which reacts with nitric oxide (NO) to generate peroxynitrite. Peroxynitrite then nitrates IDO at Tyr15, Tyr345, and Tyr353, and inactivates IDO. The levels of Tyr nitration and peroxynitrite inhibition on IDO activity are significantly reduced in the Tyr15-to-Phe mutant [94]. These results indicate that IDO can be nitrated and inactivated by peroxynitrite, and that nitration of IDO-Tyr15 is the most important factor for IDO inactivation. In addition, another Trp metabolite, 5-methoxytryptophan (5-MTP) (not from KP) (Fig. 1) derived from endothelium inhibits LPS-induced endothelial leakage and suppresses the overexpression of proinflammatory mediators (cyclooxygenase-2 and interleukin-6) regulated by LPS- or cecal ligation and puncture [95]. Furthermore, 5-MTP treatment protects against ligation injury-induced endothelial loss and detachment, decreases endothelial expression of intercellular adhesion molecular-1 (ICAM-1) and VCAM-1, suggesting attenuation of endothelial dysfunction [96].

KP and heart diseases

Epidemiologically, elevated levels of plasma kynurenines, including KA, 3-HK, AA, and 3-HAA, can predict the increased risk of acute myocardial infarction in patients with suspected stable angina pectoris [64]. Indeed, active IDO1 and higher concentrations of several Kyn metabolites including KA have been observed in poststroke [67], where they have been associated with increased mortality and the development of poststroke cognitive impairment [97]. Recently, urine Kyn:Trp ratio (KTR), an indicator of IDO activity, was identified as a particularly strong predictor of coronary events and mortality after elective coronary angiography [66]. The human heart itself has KAT I, its catalytic features are different from human brain KAT I or KAT II, which can synthesize KA from low concentrations of L-Kyn [98]. Kyn fluctuation markedly influences KA and its effect on cardiac activity. Blood KA levels can be employed to predict death and recurrent myocardial infarction in patients with coronary artery disease. Patients with coronary heart disease and depression have activated KP [65]. In addition, cardiac arrest patients with an initial non-shockable rhythm have higher plasma levels of KP metabolites, including Kyn, KA, and 3-HAA, than those with a shockable rhythm at intensive care unit (ICU) admission. Patients with higher plasma levels of Kyn, KA, 3-HAA,

and IDO activity have lower 24-h systolic arterial pressure. Patients died in the ICU have higher plasma KP metabolites (Kyn, 3-HAA, and KA) than those who survived. All these results indicate that the early activation of KP after cardiac arrest is an independent predictor of survival [99]. Patients with heart failure have higher plasma Kyn level [100]. High incidence of heart calcification on the endometrium proximal to the right ventricle was reported in female IDO-deficient BALB/c mice [101]. It is warranted to identify the causative function of specific KP catabolites for the particular heart diseases and to elucidate the effects of KP catabolites on the progression of different heart diseases.

KP and abdominal aortic aneurysm (AAA)

Abdominal aortic aneurysm (AAA) is a continual expansion of the abdominal aorta, a unique vascular disease, which is found in up to 9% of men aged >65 years [102], yet usually remains asymptomatic until it ruptures. Disruption of an AAA and its associated catastrophic physiological insult carries overall mortality more than 80%, and 2% of all deaths are AAA-associated [103]. Currently, there are no therapeutic strategies proven to block AAA progression and rupture, making endovascular or open surgical repair as the solely available approach [103]. Pathologically, AAA is highly associated with inflammation [104–107], smooth muscle cell apoptosis [108], and matrix degradation [109]. However, there is a paucity of knowledge on the mechanisms and factors controlling AAA growth at the early and middle stages [110]. Currently, there is no direct evidence to demonstrate that KP exerts its role in AAA development. Ang II, a principal mediator for the initiation and progression of AAA [111], strongly induces IDO expression in mouse aorta *in vivo* [30], suggesting that activated KP may play an important role in AAA development. Importantly, IFN- γ , a T-cell-derived cytokine, which is elevated in Ang II-treated mice or AAA patients [112], induces IDO1 expression in human VSMC with allogeneic T-cell inhibition *in vitro* and *in vivo* [40]. IFN- γ -induced IDO1 expression and activity in cultured human VSMC is substantially stronger than in endothelial cells (EC) or T cells [40]. Inhibition of IDO by 1-MT enhances medial vessel infiltration of allogeneic T cells and VSMC loss [40], which may associate with AAA development. Furthermore, aging, a key risk factor for AAA formation in humans [113] and animal model, increases IDO1 mRNA and activity in brain striatum of YAC128 mouse model [78]. It is well known that obesity is an independent risk factor for AAA formation in human [114] and animal model [115]. Obese adults have strikingly increased Kyn serum levels and dramatically enhanced Kyn/Trp ratio (IDO1 activity), suggesting obesity activates IDO1 [74]. All these AAA risk factors explicitly activate IDO1, suggesting that activated

KP may play a critical role in AAA initiation and development. Clinically, the level of plasma C-reactive protein (CRP), a critical biomarker for AAA incidence [116–118], is strongly correlated positively with Kyn, 3-HK, and 3-HAA, while negatively corrected with XA [42]. Finally, matrix metalloproteinases (MMPs) exert critical roles in AAA initiation and progression [119, 120]. Kyn increases MMP-1 and MMP-3 expression in dermal fibroblasts via MEK-Erk1/2 signaling pathway [121] or AHR activation [122]. Genetic and pharmacological approaches implicate the Erk1/2 pathway as a critical regulator of CCN3-dependent AAA development [123]. These results imply that catabolites from the activated KP may play a causative role in AAA formation. However, there is a missing link between the central enzymes of KP, dysregulated KP catabolites, and aortic aneurysm formation.

Abnormal KP and risk factors for CVD

KP metabolites and blood pressure

Hypertension is a critical risk factor for CVD. Both Trp and Kyn have been shown to dilate pre-constricted porcine coronary arteries and mouse or rabbit aorta in a dose-dependent manner [29]. The dilating effect of Trp requires the presence of active IDO and an intact endothelium, and the effect of Kyn is endothelium-independent. Kyn-induced arterial relaxation is mediated by activation of the adenylate cyclase and soluble guanylate cyclase pathways. Intravenous administration of Kyn transiently and dose-dependently decreases the mean blood pressure in spontaneously hypertensive rats *in vivo* [29]. In addition, IDO1 is expressed in EC of resistance vessels and is a potential novel contributor to hypotension in human sepsis [51]. Thus, IDO-mediated Trp catabolites might be implicated in the regulation of the vascular tone [29]. Pharmacological inhibition of IDO by 1-MT increases blood pressure in mice but not in mice deficient in IDO1 [29]. IDO deletion deteriorates hypoxia-induced pulmonary hypertension (PH) in mice [35]. In contrast, overexpression of IDO in endothelium protects against hypoxia-induced PH in mice and monocrotaline-induced PH in rats, as well as right ventricular hypertrophy and vascular remodeling [35]. Preeclampsia, a pregnancy-specific disease of maternal hypertension, is associated with decreased IDO activity and mRNA level in placentas [124]. Moreover, elevated maternal plasma KA levels in early pregnancy (at gestational week 18) are associated with a substantially increased risk of preeclampsia in obese women [56]. However, KP is less active in preeclampsia. Clinically, median Kyn levels are higher in hypertensive than in normotensive patients (Table 2) [64]. In addition, renovascular hypertensive Wistar male

rats induced by surgical clips on the left renal artery have higher plasma levels of Kyn, KA, 3-HK, and AA, as well as higher TDO activity in liver than sham rats, suggesting that renovascular hypertension is associated with KP activation [125].

The levels of a key catabolite KA in different regions of the brain of spontaneously hypertensive rats are dramatically lower than those of normotensive rats [71]. Injection of KA, an antagonist of the $\alpha 7$ nicotinic receptor in the central nervous system, decreases blood pressure in spontaneously hypertensive rats [72]. Moreover, a missense mutation in KAT I, an enzyme that metabolizes Kyn to KA, has been identified in spontaneously hypertensive rats [73], suggesting that the mutation might inhibit KAT I activity. Additionally, KYNU catabolizing Kyn to AA or 3-HAA also controls KA level (Fig. 1). Interestingly, elevated KYNU mRNA levels were detected in the brainstem of spontaneously hypertensive rats compared with normotensive controls [126]. Furthermore, KYNU Arg188Gln mutation is linked to lower KYNU activity and essential hypertension (higher systolic and diastolic blood pressure) in the Han Chinese population [127]. All these data indicate that KA may have an anti-hypertension function.

Obesity and KP

Obesity is recognized as an independent predictor for CVD [128]. However, experimental data on potential effects of kynurenines on obesity are limited. Clinically, obese adults have dramatically elevated Kyn serum levels and significantly increased Kyn/Trp ratio (IDO activity), suggesting that obesity activates IDO [74, 129]. Furthermore, IDO activity is positively associated with body mass index (BMI), fat mass, visceral or subcutaneous adipose tissue area, and subcutaneous adipocyte size [129]. Circulating Kyn levels are positively related to BMI and a higher HOMA2-IR insulin resistance index in the D.E.S.I.R. cohort (data from an Epidemiological Study on the Insulin Resistance Syndrome). The serum levels of Kyn, KA, and QA are associated with higher BMI. The mRNA expression of several KP enzyme genes, including IDO1, KYNU, KMO, and KAT III (CCBL2) is increased in the omental adipose tissue of obese women in ABOS (Biological Atlas of Severe Obesity, ClinicalGov NCT01129297) cohort compared to lean [77] (Table 2). The mRNA expression of IDO1, KYNU, and KAT III in human primary adipocytes is induced by proinflammatory cytokines. However, KMO is not expressed in human primary adipocytes. Moreover, the mRNA level of IDO1, KYNU, KMO, and KAT III is higher in proinflammatory than in anti-inflammatory macrophages [77]. A correlation of Kyn, KA, and QA with BMI and HOMA-IR is presented in another cohort [76]. Several KP enzymes,

including IDO1, but neither IDO2 nor TDO2 is known to be more expressed in the omental and subcutaneous adipose tissues and liver of women with severe obesity compared to lean individuals [75]. This increase is due to the presence of proinflammatory T cells in the adipose tissue and also comes from the adipocyte response to inflammatory stimuli. Furthermore, the ratio of Kyn to Trp (IDO1 activity) in obese patients is higher than that in lean subjects [75]. In addition, it was observed that the serum levels of Kyn and Kyn over Trp ratio are higher in women with higher BMI and they both decrease one year after body weight loss induced by bariatric surgery [130]. Importantly, mice deficient in IDO1 are resistant to Western diet-induced obesity [131]. Mechanistically, KP catabolite Kyn, but not KA activates AHR-mediated luciferase expression in mouse hepatocyte cell, suggesting that IDO1-produced Kyn leads to obesity via AHR pathway [131].

3-HAA treatment to LDLr^{-/-} mice markedly reduces plasma cholesterol and triglyceride concentration, as well as inhibits the oxLDL uptake by macrophages. The reduction in total cholesterol may be due to the signaling modulation by peroxisome proliferator-activated receptors [61]. In another experiment, LDLr^{-/-}/IDO^{-/-} mice demonstrated a remarkable induction of serum lipids, particularly triglycerides [101]. However, the underlying mechanism by which KP catabolites regulate lipid metabolism remains elusive.

Upregulated IDO1 in both adipose tissues and liver is inversely correlated with arterial blood pressure, suggesting that IDO1 upregulation may represent a local compensatory mechanism to inhibit obesity-induced hypertension [75]. Further research is needed to investigate whether IDO1 plays a causative role in obesity-related hypertension. Interestingly, periaortic fat from rat thoracic aorta has strong staining for IDO and the brown fat marker uncoupling protein-1 (UCP1), while the white fat around rat mesenteric artery or abdominal aorta has weak staining for IDO and UCP1 [132]. Moreover, rat perithoracic aortic fat has higher IDO activity than visceral and mesenteric artery fat. Periadventitial fat inhibits Ang II-induced rat thoracic aortic and mesenteric artery maximal contraction. However, periadventitial fat does not blunt agonist (phenylephrine)-induced contraction of the abdominal aorta. The IDO inhibitor 1-MT blocks the fat-induced inhibition of Ang II-mediated contraction in the thoracic aorta but not in the mesenteric artery. The IDO product Kyn relaxes the rat thoracic aorta at high concentrations (9 mM), whereas the downstream catabolite QA relaxes the contracted rat thoracic aorta (approximately 80%) at a relatively lower concentration (1 mM). All these data imply that IDO is active primarily in the brown fat surrounding the thoracic aorta and inhibits aortic contractility [132]. It is interesting to identify catabolites from KP in the perivascular adipose

tissue which control blood vessel contraction and relaxation. Nevertheless, a causal function has yet to be demonstrated in any of these conditions.

KP and diabetes

Diabetes mellitus is a major risk factor for CVD which is the principal cause of death in more than 70% of type 2 diabetic patients [133]. Despite advances in our understanding of some early markers of CVD, the mechanisms underlying the increased risk of CVD in diabetes remain incompletely delineated. Recently, there are several reports on upregulated KP in type 1 (T1D) or type 2 diabetes (T2D), as well as the diabetic cardiovascular complication. Clinically, T1D, but not T2D has higher concentrations of plasma AA (2.3-fold induction) compared with non-diabetic subjects [50]. However, the cause and consequence of AA elevation in T1D are unknown. The underlying mechanism may be the individual kynurenines that inhibit proinsulin synthesis from pancreatic islets in rat models [85] or generate complexes with insulin decreasing its biological activity [134]. Elevated plasma KA and XA are observed in both T1D and T2D compared with non-diabetic persons [25]. TDO inactive mutation attenuates high sucrose diet-induced insulin resistance (IR) in *Drosophila*, suggesting that upregulated KP mediates high sucrose diet-induced IR [135]. Levels of kynurenine-modified serum proteins (KMSP) are significantly higher in diabetic patients and particularly in type 2 diabetic patients with CVD. Moreover, KMSP level is only associated with the onset of diabetes and might participate in CVD development [136]. However, the role of KMSP in diabetes and diabetic cardiovascular complication remains elusive.

Some catabolites of the KP have been proposed to be risk factors for the IR development. Surplus dietary Trp induces IR in pigs [137]. Diabetic retinopathy patients have elevated serum Kyn [68]. mRNA expression of KAT I and KAT II is modestly reduced in skeletal muscle from type 2 diabetic patients compared with that in normal glucose tolerant humans [69]. Exercise enhances KAT activity [69] may through upregulation of skeletal muscle KATs mediated by PGC-1 α 1 [22]. As expected, bariatric surgery is associated with the improvement and even the remission of T2D. It was shown that higher levels of KA and QA 1 year after the bariatric surgery are associated respectively with T2D mitigation and better glucose homeostasis and that lower levels of XA are associated with better glucose homeostasis [130]. Recently, it was demonstrated that serum Kyn concentration in chronic hepatitis C virus (HCV) patients is highly associated with HOMA-IR and HOMA-beta scores [136], suggesting that Kyn may contribute to higher incidence of IR in HCV patients [53].

Aging and abnormal KP

Aging is a critical risk factor for CVD. Importantly, aging increases IDO1 mRNA and activity in brain striatum of YAC128 mouse model [78]. Upregulation of KP in aging may be due to the activation of IDO by age-related chronic inflammation [138]. Depletion of TDO-2 (homology to human TDO) in *Caenorhabditis elegans* suppresses the toxicity of age-related aggregation-prone proteins, including α -synuclein, β -amyloid, and polyglutamine proteins, as well as extends life span by increasing Trp levels [139]. Kyn increases with age apparently [79]. Kyn levels and IDO1 activity are highly associated with aging [80]. Moreover, KA, 3-HK, and AA are associated positively with age [64]. XA treatment induces mitochondria damage, cytochrome C release in VSMC, consequent activation of caspase-3 and -9, and resultant cellular apoptosis [59]. Further investigations are warranted to clarify the causal or resultant roles of KP catabolites on age-related CVD.

Smoking and KP

16% higher XA serum concentrations are presented in heavy drinkers than never or occasional drinkers [140]. There is no substantial change in Kyn metabolites, which was observed among smokers [140], although smoking is associated with considerably lower levels of PLP, biologically active form of vitamin B6 [141], a cofactor for KATs and KYNU. KA, an endogenous antagonist for NMDA and α 7 nicotinic acetylcholine receptor, is dramatically increased in rat hippocampus, striatum, and frontal cortex, but not in the serum by prolonged (for 10 days) subcutaneous injections or osmotic minipumps of nicotine. Nicotine has no effect on Kyn in brain or blood [83]. There is no report about the effect of smoking on KP and its function in CVD.

Therapeutic implications for abnormal KP-associated CVD

Dysregulation of KP of tryptophan catabolism has been correlated with several cardiovascular diseases. Investigation quantifying KP catabolites in local tissue, plasma, and urine samples from patients diagnosed with CVD or its risk factors has presented associations between kynurenines concentrations and pathology (Table 2). Inflammatory condition usually upregulates IDO1, KMO, and KYNU. Notably, 3,4-DAA, a synthetic derivative of the tryptophan catabolite AA, as well as 3-HAA play an anti-inflammatory function in the vascular system and protect from atherosclerosis [45, 84]. However, catabolite KA exerts a proinflammatory role [92] and enhances

atherogenesis [86]. Taken together, KP activation might be implicated in inflammation-related CVD, such as atherosclerosis, AAA, and endothelial dysfunction. Indeed, the inflammation-responsive and stress-reactive feature of KP enzymes has promoted biomarkers development and brought into focusing several promising therapeutic targets of relevance across a broad range of CVD and cancer [9]. Animal studies have demonstrated that KP enzyme inhibition, especially involving IDO1, KMO, KYNU, and KAT II, can serve as the viable drug targets for treating cardiovascular diseases. Pharmacological manipulation of the KP enzymes employing structure-based drug design has, therefore, become an attractive area of drug development. Here, we briefly summarize the biochemical features of key KP enzymes and principles of inhibitors development.

IDO regulation

IDO is expressed unremarkably in most tissue but highly active in placenta required for the maternal immune suppression of T cells to prevent rejection of fetus [142]. Usually, IDO is silent under physiological condition, but strongly induced by inflammatory mediators under disease state. IDO1 induction in macrophages and VSMC by HFD contributes to the exacerbated atherosclerosis [86]. However, deletion of IDO1-positive pDCs enhances atherosclerosis in LDLr^{-/-} mice, while aortic IDO1-positive pDCs prevent atherosclerosis [38]. In addition, IDO1 deletion accelerates early atherogenesis in aortic root and promotes plaque destabilization of ApoE^{-/-} mice [84].

IDO inhibition by 1-MT promotes atherosclerosis in ApoE^{-/-} mice [45]. Thus, IDO demonstrates the greatest potential as the druggable target for CVD treatment. Three major kinds of small-molecule IDO inhibitors have been used for cancer treatment in clinical trials (Table 3). For example, 1-MT (referred to as Indoximod) is the first and widely used competitive inhibitor of IDO1 [143], and 1-MT is being used as cancer immunotherapy combined with Docetaxel or Paclitaxel chemotherapy in phase II clinical trials for metastatic breast cancer [144]. There is another phase I/II trial using Indoximod (overcoming tumor-induced immune suppression) in combination with Tergenumatucel-L (activating human immune system) immunotherapy and Docetaxel to treat subjects with advanced non-small cell lung cancer (NSCLC) [145]. 1-MT was also used for atherosclerotic research in mouse [45]. Another IDO1 inhibitor, 4-amino-1,2,5-oxadiazole-3-carboximide (INCB024360, also named Epcadostat) was in phase II clinical trial for patients with myelodysplastic syndromes [146]. NLG919 (an imidazoleisoindole derivative), a potent direct IDO1 inhibitor ($K_i = 7$ nM) [147], is currently in phase I clinical trials for adult patients with recurrent advanced solid tumors [148]. In addition, peptide-based IDO1-targeting vaccines were employed in phase I trial for stage III-IV non-small-cell lung cancer patients [149]. However, there is no clinical trial with IDO inhibitors for CVD treatment.

The novel and potent IDO1 (EC 1.13.11.52) inhibitors are being developed intensively [163]. IDO1 is a monomeric, redox-sensitive, heme-containing dioxygenase

Table 3 Inhibitors of targeted KP enzymes

Targeted enzymes	Inhibitors	K_i	IC ₅₀	Testing systems	Outcomes	References
IDO1	1-Methyl D-tryptophan	34 μ M	120 μ M (HeLa)	Rabbit small intestinal IDO	Anti-tumor	[143, 150]
	INCB024360	10 nM	72 nM (IDO1), 7.1 nM (HeLa)	Tumor cells	Anti-tumor	[150, 151]
	NLG919	7 nM	75 nM	Effector T cells	Immune activation and tumor regression	[152, 153]
KMO	GSK180	N/A	6 nM	Mouse model of acute-pancreatitis	Anti-inflammation	[154]
	Ro-61-8048	4.88 nM	37 nM	Rat neuropathic pain model	Decrease neuropathic pain	[155]
	UPF-648	56.7 nM	20 nM	Human KMO	Neuroprotection	[156]
KYNU	o-Methoxybenzoyalanine	4 μ M	N/A	Rat	Sedation	[157]
	2-Amino-4-[3'-hydroxyphenyl]-4-hydroxybutanoic acid	100 nM	N/A	Recombinant Human KYNU	N/A	[158]
	S-phenyl-L-cysteine sulfoxide	140 μ M	N/A	Recombinant KYNU of <i>P. aeruginosa</i>	Potential antivirulence	[159]
KAT II	PF-04859989	N/A	23 nM	Rat brain	Anti-schizophrenia	[160]
	BFF 122	N/A	1 μ M	Rat brain	Cognitive enhancement	[161, 162]

K_i inhibition constant, N/A not available. For definitions of the abbreviations, please see the main text

enzyme, making that possible to develop competitive inhibitor binding to the heme active site [164]. As crystal structures of IDO1 [165], IDO1 bound with an inhibitor [147, 166], and IDO1 bound with substrate [167] were revealed, it is indicated that occupation of both hydrophobic pockets A and B, as well as heme iron-atom in IDO1, is critical for IDO1 inhibition [147]. Furthermore, the JK loop (a 20-amino-acid protein segment) in human IDO1 connecting the J and K helices undergoes structural rearrangement upon substrate binding. JK loop N-terminal (JK loop^N) may function as a target for post-translational modifications such as phosphorylation or a mediator for protein–protein interactions. The residue Thr379 in highly conserved “GTGG” motif of JK loop C-terminal (JK loop^C) is crucial for hydrogen bonding interactions between substrate Trp and IDO enzyme, highlighting the importance of JK loop conformation in regulating the enzyme activity. Interestingly, Tyr353 nearby helix J can be nitrated upon exposure to peroxyxynitrite, resulting in IDO inhibition [168]. The structure–activity relationships (SAR) analysis by molecular docking technologies and pharmacophore models indicates that 1*H*-indazole scaffold is a novel key pharmacophore and necessary for potent IDO1 inhibition [169]. Recently, rational drug design demonstrates that O-alkylhydroxylamines function as IDO1 inhibitors by binding heme iron with nanomolar-level cell-based potency and limited toxicity [170]. Finally, IDO2 is clearly different from IDO1 regarding substrate specificity, affinity, kinetic parameters, and inhibition profiles [171, 172]. IDO2, but not IDO1, generates a distinct signal for activation of transcription factor liver-enriched inhibitory protein (LIP) that is Trp-restoration independent [15]. All the results suggest that it is necessary to develop specific inhibitors of these enzymes. For example, D-1MT but not L-1MT strongly inhibits IDO2 but not IDO1 in overexpression T-REx cells system [15]. However, we are far from understanding the profile of IDO expression and activation during CVD initiation and development. A better understanding of IDO biology is needed to comprehend the effect of IDO inhibitors and to provide a rationale for their therapeutic application in CVD. As a note, no IDO1 inhibitor is currently being used in clinical trial for CVD treatment.

Kynurenine monooxygenase (KMO) inhibition

KMO (EC 1.14.13.9, also is referred to as kynurenine 3-hydroxylase), a pivotal enzyme in KP [4], is a flavin adenine dinucleotide (FAD)-dependent monooxygenase locating in the outer mitochondrial membrane where it catalyzes Kyn to 3-HK [156]. KMO is expressed in kidney, liver, and brain, as well as macrophages and monocytes. A crystal structure of truncated yeast KMO has been reported [156]. KMO inhibitors are being researched as

potential treatments for neurodegenerative diseases, such as Alzheimer’s, Huntington’s, and Parkinson’s diseases [162, 173, 174]. There have been no reports or detailed studies on KMO inhibition for CVD in animal or clinical trial. Reported KMO inhibitors contain an aryl ring appropriately spaced from an acidic functionally or bioisostere [175]. The most well-known and potent KMO inhibitor is Ro-61-8048, which decreases neuropathic pain intensity induced by chronic constriction injury and potentiates the analgesic properties of morphine [176]. In addition, KMO is central to the pathogenesis of acute-pancreatitis-multiple organ dysfunction syndrome (MODS). KMO inhibition by genetic deletion or a potent and specific inhibitor, oxazolidinone GSK180 (3-(5,6-dichloro-2-oxobenzo[d]oxazol-3(2 H)-yl)propanoic acid), prevents extrapancreatic tissue injury to the lung, kidney, and liver in mouse model of acute-pancreatitis (Table 3), may through upregulation of Kyn, KA, and AA, or downregulation of 3-HK and 3-HAA [154]. Therefore, KMO inhibition may prevent inflammation-related CVD including atherosclerosis and AAA. It is warranted to test the currently available KMO inhibitors for CVD treatment in animal and clinical trials.

Kynureninase (KYNU) inhibition

Eukaryotic kynureninase (KYNU, EC 3.7.1.3) preferentially catalyzes the conversion of 3-hydroxy-L-kynurenine rather than L-kynurenine to 3-hydroxyanthranilic acid than anthranilic acid, as well as L-alanine [177]. KYNU is a cytoplasmic enzyme with 465 amino acids and 52 kDa, one member of the amino-transferase superfamily, contains PLP as the cofactor [178]. A conserved lysine residue Lys276 was identified to be the critical residue for cofactor covalent-binding within the pyridoxal-p-binding site consensus sequence. The reaction specificity of KYNU is determined in part by active site residues, Trp64, Gly281, and Thr282 in *P. fluorescens*, and the homologous His102, Ser332, and Asn333 in human KYNU. Asn333 can form a hydrogen bond to the 3-OH of 3-HK in the human enzyme [19, 178]. These results may help to develop the selective inhibitor for KYNU. A structural analogue of the reduced form of substrate 3-HK, 2-amino-4-[3'-hydroxyphenyl]-4-hydroxybutanoic acid, is the most potent and reversible inhibitor of KYNU from both rat and human, with a K_i value of 100 nM [158]. Interestingly, removal of the aryl amino group coupled with a reduction of the carbonyl group at position 7 of the alanine side chain greatly enhances the potency of the inhibitor [158]. Importantly, rats treated with KYNU inhibitor, o-methoxybenzoylalanine (oMBA) (50–400 mg/kg i.p.), have increased Kyn and KA levels in the brain, blood, liver, and kidney, associating with a decreased locomotor activity [157]. Recently, another structural analogue of Kyn, S-phenyl-L-cysteine

sulfoxide, was shown to competitively inhibit *P. aeruginosa* KYNU activity in vitro, suggesting a potential antivirulence strategy [159]. Since KYNU directly controls 3-HAA formation, which regulates cardiovascular cells inflammation and apoptosis, it would be interesting to exam the effect of KYNU inhibition on CVD initiation and development.

Kynurenine amino-transferases (KATs) inhibition

KATs are homodimeric PLP-dependent enzymes that mediate the irreversible and permanent transamination of Kyn to KA, a known neuroactive agent, as well as 3-HK to XA [179] (Fig. 1). Evidence suggests that abnormal levels of KA are implicated in many neurodegenerative diseases such as Parkinson's disease, Huntington's disease, Alzheimer's disease, and schizophrenia [179, 180]. Four isoforms (KAT I to IV) of this enzyme have been identified [9]. Human KAT I and KAT III have the highest sequence identity (51.7%) [179]. Human KAT I is located in the heart, it more efficiently catalyzes the transamination of hydrophobic amino acids, such as Trp. It is not inhibited by glutamine [98]. KAT II seems to be the major isoform contributing to KA formation (75%) in human brain [179]. However, KAT II is undetectable in murine heart and skeletal muscle [22]. Thus, KAT II inhibitors have been proposed as new therapeutic approaches for psychiatric and cognitive disorders [181]. Human KAT III shares 30.1% similarity of amino acids to that of KAT II and has higher mRNA expression levels in heart, liver, and kidney [182]. KAT IV is identical with glutamic-oxaloacetic transaminase 2 or mitochondrial aspartate amino-transferase [183]. It plays a major role in KA formation in mouse brain [184]. KAT (I, II, and III) protein levels in mouse gastrocnemius muscle can be induced by exercise, but not chronic mild stress [22]. Importantly, exercise training increase KATs (I, II, III, and IV) mRNA and protein levels in human vastus lateralis skeletal muscle [22, 185]. KATs are insensitive to minerals changes, but sensitive to dietary vitamin B6, which is sequestered by an intricate hydrogen bond networks and covalently binds to the conserved catalytic Lys263 to form a reactive Schiff base [186]. Thus, Lys263 in the active site of KAT II is very important for KAT II activity. Recently, the three-dimensional (3D) crystal structure of the PLP form of hKAT II at 1.83 Å resolution was reported [180, 187]. Several KAT II inhibitors have been designed to interact with Kyn-PLP imine adduct and function as irreversible inhibitors [9]. For example, Pfizer developed cyclic hydroxamic acid PF-04859989 (KAT II $IC_{50}=23$ nM) which forms an irreversible covalent bond with the enzyme cofactor PLP in the active site, leading to a highly potent complex [188]. The hydroxyl and keto moieties in the quinolone core of PF-04859989 make two critical hydrogen bonds with Asn202 and Arg399 in hKAT II.

PF-04859989 is a brain penetrant and irreversible inhibitor of human and rat KAT II, and it is highly selective for KAT II versus KAT I, III, and IV isoforms [160]. KATs inhibition may be a novel therapeutic strategy for treating preeclampsia and other CVD. However, the expression profile and pathophysiological function of KATs in cardiovascular system remain unknown. There are not many in vivo studies presented for hKAT II inhibitor. There is no KAT III or KAT IV inhibitor which was reported to date.

Conclusions and perspectives

It is still unclear whether endogenous kynurenines directly contribute to the initiation or promotion of CVD development, although emerging evidence suggests that Trp metabolism has diverse and complex effects on the cardiovascular outcomes in human. Gaining a better understanding of cell type specificity and the mechanisms of regulating KP enzyme expression/activities is extremely important because they generate and determine the balance of different metabolites. It is critical to employ pharmacological or genetic strategy with a gain of function and loss of function to clarify the causal or consequent roles of different KP catabolites in identical cells-associated typical CVD. As mentioned above, the manipulation of KP metabolism through the administration of enzyme inhibitors can serve as a novel therapeutic strategy for several CVD and their risk factors and the development of specific KP enzyme inhibitors is desirable. However, this manipulation should be controlled precisely by monitoring kynurenines levels to identify new therapeutic targets and biomarkers and assess the normalization of KP imbalances. Numerous medicinal chemistry studies are currently aimed at the design of novel, potent, and selective inhibitors for each of these aforementioned enzymes. Taken together, to target the pathway effectively, it is essential to better understand: 1, the effects of KP metabolites on initiation and progression of CVD in vivo; 2, controlling of particular KP enzymes on the levels of the bioactive metabolites in vivo; 3, unique features and regulation of KP enzymes under physiological and CVD pathological conditions; and 4, structure–activity relationship to develop the specific and potent inhibitors or activators for KP enzymes, as well as metabolites analogue to mimic or antagonize KP intermediates' function in vitro and in vivo. All deep investigation on KP would accelerate our understanding of KP in the prevention, early detection (biomarkers developing), treatment, and cure of CVD, and benefit the personalized and precision cardiovascular medicine.

Acknowledgements This study was supported by funding from the following agencies: National Institutes of Health RO1 (HL132500,

HL128014, HL110488, HL080499, HL089920, AG047776, and CA213022). This work is in part supported by the Georgia Research Alliance. Dr. Zou is a Georgia Research Alliance Eminent Scholar in Molecular Medicine.

Compliance with ethical standards

Conflict of interest The authors declare that there are no conflicts of interest.

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