Aldehyde Dehydrogenase 2 and Heart Failure



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1 Introduction

Heart failure (HF) is a structural or functional cardiac abnormal syndrome characterized with series of symptoms and signs such as breathlessness, fatigue, pulmonary crackles, and peripheral edema. Being a terminal phase of most myocardial lesions, HF has become a leading cause of mobility and mortality worldwide, associated with heavy clinical burden and economic costs affecting over 23 million people [14]. There is an increase to 5.5% with systolic dysfunction and an increase to 36.0% with diastolic dysfunction in people 60 years or older [85]. The costs accompanied with heart failure stand 2-3% of the total healthcare system expenditure in high-income countries and are expected to increase >2-fold in the next 2 decades [34].

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The pathological mechanisms of HF include hemodynamic abnormalities, neuroendocrine cytokine system over-activation, bioenergetics defects, signal transduction pathway dysfunction, and abnormal calcium homeostasis [54]. In addition, oxidative stress and inflammatory disorders also contribute to the pathogenic process [58, 87]. HF is caused by multiple different etiologies; however, all causes have final common pathways, at least in part, independent of the original cause. Based on previous clinical trials of neurohormonal therapies, neurohormonal activation plays a pivotal role in its pathophysiology [56, 57]. Any cardiovascular diseases can cause cardiac injury; as a consequence, myocyte cells loss and the remaining myocytes become eccentric hypertrophy. Followed by neurohormonal activation, the left ventricle changes from elliptical to spherical and is characterized by functional mitral regurgitation. Afterward, left ventricular remodeling occurs with fibrosis and ventricular dilatation in which process myocardial oxygen consumption increases and myocardial contraction efficiency reduces [10, 69]. As the course progresses, concomitant renal dysfunction and gut congestion cause reduced response to diuretics and inflammatory activation, leading to worse outcomes [58, 84]. Existing therapies including angiotensin-converting enzyme (ACE) inhibitors, angiotensin receptor blockers (ARBs), β -blockers, mineralocorticoid receptor antagonists, and advanced device therapies provide symptomatic and clinical benefits; however, they do not completely solve the molecular abnormalities. Thus, it is urgent to find other effective therapeutic targets.

Recently, energetic impairment in the pathophysiology and progression of HF has caused lots of interest. Mitochondria accounts for 30% of the volume of cardiomyocytes whose dysfunction has been recognized as the key link in the development of heart failure [69]. Mitochondrial dysfunctions cover altered utilization of metabolic substrates, increased formation of reactive oxygen species, impaired mitochondrial electron transport chain activity, abnormal mitochondrial dynamics, and altered ion homeostasis [4]. In view of the above evidences, mitochondria appear to be the main target for direct improvement of cardiac function. Acetaldehyde dehydrogenase 2 (ALDH2) is firstly regarded as an alcohol metabolism enzyme, which mainly distributed in the mitochondria. Increasing evidences have supported its important cardioprotective role [44, 53, 61]. In the following review, we will summarize its beneficial role and novel insights in heart failure due to different etiologies, as well as the potential therapeutic measures.

2 Characteristics of ALDH2

Acetaldehyde dehydrogenase (ALDH) is an enzyme superfamily responsible for the catalytic oxidation of acetaldehyde to acetic acid both physiologically and pathologically [38]. It has been identified 19 isozymes till now. ALDH2 is the most active isozyme and its molecule form is a 56 kDa tetramer. The precursor protein of ALDH2 encodes from chromosome 12, enters the mitochondria under the guidance of the signal peptide, and cleaves off the signal peptide to locate in mitochondria [28]. ALDH2 is widely distributed in human liver, kidney, heart, lung, brain, and

other tissues. It contains three domains including coenzyme-binding or NAD⁺binding domain, catalytic domain, and oligomerization domain [61]. ALDH2 also processes three kinds of enzyme activities including dehydrogenase, esterase, and reductase activities detoxifying aldehyde, 4-nitrophenyl acetate, and nitroglycerin, respectively [55].

A total of 84 single nucleotide polymorphism (SNP) loci have been found on human ALDH2 gene [46]. The foremost is rs671 G > A mutation which presents in exon 12 and results in amino acid substitutions of p.Glu504Lys [74]. The genotypes of ALDH2 comprise wild homozygous (GG also known as ALDH2 *1/*1), owning normal catalytic activity of the enzyme; mutant heterozygous (GA also known as ALDH2 *1/*2), processing a decrease in enzyme catalytic activity with only 10%–45% of the wild homozygous; and mutant homozygotes (AA type also known as ALDH2 *2/*2) having only 1%–5% enzyme catalytic activity of the wild homozygous [15]. In general, the prevalence of the genetic polymorphism (ALDH2 *2/*2) is found in nearly 8% of world populations and 40% of East Asian population [103].

3 Beneficial Role and Mechanisms of ALDH2 in Heart Failure

Experimental and clinical studies have proved that ALDH2 plays a pivotal role in heart failure via maintaining cellular homeostasis [13, 17]. Evidences from our group and others have verified ALDH2 is closely related to various etiologies of heart failure, such as coronary artery disease (CAD) [80], hypertension [71], alcoholism [73], and other susceptibilities [102, 106]. Next we will review the pathological role and underlying mechanisms of ALDH2 involved in heart failure.

3.1 ALDH2 and Ischemic Heart Failure

Numerous studies have shown that ALDH2 has cardioprotective effects in ischemic with or without reperfusion myocardial injury. The expression of ALDH2 was found decreased in infarction border zone [90]. Cardiac function deteriorated in ALDH2 knockout posts myocardial infarction (MI) mice as evidenced by increased left ventricle (LV) cavity, LV end-diastolic pressure, and infarct size [80]. Meanwhile, there were more apoptotic cells in the non-infarcted LV region as compared with wild-type (WT) MI mice. In contrast, overexpression of ALDH2 in the heart could alleviate these injures. Ma, et al. found that ALDH2 induced autophagy during ischemia and inhibited autophagy during reperfusion which reduced hypoxic and reoxygenation cell death [53]. These data demonstrated that ALDH2 deficiency aggravated mitochondrial dysfunction and increased cardiomyocyte apoptotic cell death.

DNA methylation is a process adding methyl groups to cytosine residues into DNA sequences, which can prevent transcription factors from entering the gene regulatory region, thereby inhibiting gene transcription [65]. Increasing studies indicated that DNA methylation provided a potential molecular basis on energy metabolism between environmental and genetic factors interaction and might contribute to myocardial injury [40]. We also found DNA methylation at CpG sites (CpG1, CpG2, and CpG7) in the upstream sequence of ALDH2 promoter was upregulated post MI. These abnormal hypermethylations at the CpG sites downregulated ALDH2 enzyme activity and aggravated ischemic damages [90]. DNA methylation reflected the upstream mechanism of ALDH2 regulating cardiac function after ischemia.

We also demonstrated the downstream mechanisms of ALDH2 in protecting cardiac from ischemic injures. ALDH2 could increase the intracellular levels of 4-HNE, which could exacerbate apoptosis by inhibition of HSP70, phosphorylation of JNK, and activation of p53 [80]. 4-HNE is also a diffusible product of membrane lipid peroxidation and relates to oxidative stress-induced cell death [70]. Mitochondrialderived ROS also attacked polyunsaturated fatty acids, leading to membrane lipid peroxidation, thereby increasing reactive aldehydes [7, 8]. We demonstrated both experimentally and clinically that ALDH2 was vital in regulating microenvironment homeostasis. ALDH2 could promote angiogenesis post chronic ischemia. ALDH2 deficiency inhibited tubelike construction formation of hypoxia endothelial cell through HIF-1α/VEGF pathway, deteriorating perfusion recovery in ischemia tissue, while overexpression of ALDH2 promoted angiogenesis. Furthermore, clinical data suggested that the dysfunction of ALDH2 due to gene variant was an unfavorable factor for revascularization in patients with chronic total occlusion (CTO). Therefore, targeting ALDH2 activity may be a potential therapeutic strategy for chronic ischemic heart failure, and we have used this achievement to guide clinical decision-making [49].

3.2 ALDH2 and Stress-Induced Heart Failure

Endoplasmic reticulum (ER) stress refers to an increase in unfolded and misfolded proteins in the ER that disrupts the homeostasis in response to cellular stressors, such as heat, hypoxia, metabolic starvation, angiotensin II, and tumor necrosis factor- α [29, 42]. Substantial evidences indicated ES stress as important target for the treatment of cardiovascular disease, including ischemia/reperfusion injury, atherosclerosis, cardiac hypertrophy, and heart failure [29, 82]. Our results suggested that ALDH2 deficiency aggravated cardiac contractile dysfunction following activation of ES stress, manifested as descend of ejection fraction and fractional shortening. NADPH oxidase (p47phox subunit) increased in ALDH2 knockout mice, suggesting that ALDH2 might regulate Akt signaling pathway through p47phox NADPH oxidase-dependent manner against ER stress and ER stressinduced apoptosis [47].

Persistent pressure overload such as refractory hypertension is a significant risk factor for heart failure and sudden death. Series of pathological cardiac remodeling were characterized by increased myocardial cells, "fetal gene program" activation, cytoskeletal reorganization, and irreversible systolic dysfunction [35, 51]. Autophagy is an important homeostatic pathway in degrading damaged proteins and intracellular organelles [24]. It has been found to be involved in pressureinduced heart failure [22]. Autophagy-related signaling pathways involved in pressure overload included the classic AMPK-mTOR-autophagy, Beclin-1dependent pathway, Akt/mTOR/FoxO3a signal pathway, and PI3K/Akt signaling [48, 89, 96, 98]. In early compensated cardiac hypertrophy after transverse aortic constriction (TAC), autophagy played an adverse role with worse cardiac function and severer mitochondria damage. In this process, ALDH2 acted through the regulation of PI3K/PTEN/Akt signaling [95]. Besides, ALDH2 deficiency further inhibited autophagy during decompensated cardiac hypertrophy accompanied with inactive Beclin-1-dependent autophagy signaling [71]. Therefore, autophagy works in both early stage and late stage of ALDH2 regulation in pressure-overload adaptive response.

3.3 ALDH2 and Alcoholic Heart Failure

Alcoholic cardiomyopathy is characterized by a dilated left ventricle and reduced myocardial contractility due to a long-term history of heavy alcohol consumption. It was estimated that about one-third of alcoholics suffered from varying degrees of alcoholic cardiomyopathy and approximately half of them resulted in death within 4 years [43, 101]. Mitochondrial defects, cell death, heart rate variability, and cardiac remodeling would eventually result in heart failure [31].

As early as the twentieth century, it was found about 50% of Asians had facial flushing after drinking alcohol, and these individuals were tested having an inactive form of mitochondrial ALDH2 [88]. rs671 in ALDH2 gene was the most influential genetic variant linked to alcohol consumption [83]. ALDH2 deficiency aggravated alcoholic myocardiopathy by weakening acetaldehyde-biogenic amine condensation products detoxification, breaking intracellular Ca2+ homeostasis, increasing apoptosis, upregulating autophagy, as well as impairing mitochondrial function [52, 66, 72, 73, 101]. Besides, our study explored the role of ALDH2 in low-to-moderate alcohol consumption. On one hand, we confirmed the cardioprotective effects of low-to-moderate alcohol consumption which manifested by elevated HDL-c levels and upregulated HO-1 expression in the myocardium. However, the benefits were disrupted when ALDH2 was deficient, possibly by activating ROS-dependent apoptosis and RIP1/RIP3/MLKL-mediated necrosis [73]. A Guangzhou biobank cohort study genotyped rs671 of ALDH2 in 4867 men. Diastolic blood pressure and HDL cholesterol which associated with ALDH2 variants were attenuated after adjusting for alcohol use. The result suggested the apparent associations between physical activity and alcohol use in ALDH2 variants population [3]. Even moderate alcohol use was found associated with subclinical adverse effects with greater left ventricle mass and more impaired diastolic functions in subjects carrying ALDH2 variants, especially among East Asians [36]. Individuals with inactive isoforms of ALDH2 should be warned to avoid drinking alcohol, even for social or occupational promotion.

3.4 ALDH2 and Diabetes Mellitus-Related Heart Failure

The increasing morbidity and mortality of heart failure are related to the increase in aging, obesity, and diabetes mellitus in a large part. The prognosis of heart failure in patients with diabetes is much worse than that in patients without diabetes. The early symptom of diabetic cardiomyopathy is diastolic dysfunction and can gradually develop into systolic dysfunction. Metabolic disorder is an important feature of diabetic cardiomyopathy, manifested as reduced glucose uptake and increased fatty acid utilization accompanied by oxidative stress, inflammation, cardiomyocyte apoptosis, and myocardial fibrosis [37]. Beyond these, mitochondrial dysfunction, impaired mitochondrial and cardiomyocyte calcium handling, endoplasmic reticulum stress, and reduced nitric oxide bioavailability were also implicated in the development and progression of diabetic cardiomyopathy [39].

A genome-wide association study contained 12,720 participants found rs671 (ALDH2) was associated with metabolic syndrome (MetS) in Han Chinese. What's more, the effects of rs671 on metabolic components were significantly correlated with drinking [108]. Previous studies showed that ALDH2 improved the contractile function of advanced diabetic cardiomyopathy by regulating Ca²⁺ homeostasis and autophagy [32, 104]. Our data found that ALDH2 deficiency impaired diastolic function in early stage of diabetic cardiomyopathy, while cardiac contractile function remained normal. In this stage, ALDH2 deficiency disrupted energy metabolism with increased AMP/ATP and ADP/ATP and decreased PCr/ATP ratio, which in turn induced activation of energy regulatory LKB1/AMPK pathway. The progressive accumulation of phosphatidylcholine and phosphatidylinositol in heart tissue induced metabolic homeostasis disequilibrium and led to deterioration of diastolic function [20, 91].

3.5 ALDH2 and Aging-Related Heart Failure

Aging is an irreversible biological process. In the cardiovascular area, age-dependent increases include left ventricular hypertrophy, diastolic dysfunction, atrial fibrillation, as well as vascular intimal thickening and vessel stiffness. Aging-related cardiac dysfunction is characterized with loss of cardiac contractile reserve, increased fibrosis and remodeling, impaired cardiomyocyte proteostasis, and loss of autophagy [21, 50]. Heart failure can be regarded as an aging-related phenotype. Aging-associated cardiac pathological changes involve oxidative stress, short telomere defect, mitochondrial damage, intracellular Ca²⁺ mishandling, etc. [2, 99].

Numerous studies suggested ALDH2 participated in the process of aging and age-related cardiovascular diseases [18, 106]; however, whether ALDH2 is beneficial or detrimental is still controversial. Wu found that ALDH2 ablation led to cardiac aging and sustained usage of Alda-1 (a specific activator of ALDH2) abrogated the aging effect [94]. ALDH2 activity was discovered to be significantly decreased in aged hearts which also demonstrated the benefits of activation of ALDH2 on retarding the aging process. In the meantime, ALDH2 was also discovered to exert age-dependent vasoprotective effects with decreased mitochondrial ROS formation and oxidative mtDNA damage [92]. On the contrary, our lab found ALDH2 overexpression such as using Alda-1 accentuated agingrelated cardiomyocyte. The dysfunction was characterized by increased contractile dvsfunction, oxidative stress, intracellular Ca2+ mishandling, and mitochondrial injury [105]. Moreover, AMPK/Sirt1 signaling cascades were found taken part in ALDH2-accenuated cardiac aging [106]. In the light of the debates, further epidemiological of different races or experimental studies will be needed to provide more evidence about the effect of ALDH2 in aging.

3.6 ALDH2 and Drug-Induced Heart Failure

Several chemical agents and drugs could impair cardiac mitochondrial function via destroying mitochondrial respiratory chain (e.g., uncoupling) or inhibiting mitochondrial enzymes. The most common agents are anticancer drugs such as anthracycline doxorubicin (DOX), cisplatin, and Trisenox; antiviral compound azidothymidine (AZT zidovudine); and several oral antidiabetic drugs such as Avandia [86].

The chemotherapy drug DOX is frequently found in inducing cardiotoxic. Left ventricular systolic pressure would significantly reduce, and left ventricular enddiastolic pressure would overtly increase after DOX treatment. ALDH2 attenuated this cardiotoxicity by inhibiting oxidative stress, decreasing the expression and activity of NADPH oxidase 2, and reducing myocardial apoptosis. In addition, DOX-induced myocardial dysfunction was severer with increased levels of 4-HNE and autophagy in ALDH2 knockout mice. Besides, these symptoms could be improved when ALDH2 activity was restored, suggesting that inhibition of 4-HNE and autophagy may be the possible mechanisms of ALDH2 against DOX-induced cardiac dysfunction [79].

Oxidative stress-induced cardiomyocyte apoptosis is also a main part in the pathogenesis of heart failure [59]. ROS production and accumulation caused intracellular redox imbalance, leading to mitochondrial dysfunction and decreased production of ATP [26]. We elucidated the relationship between ALDH2 deficiency and oxidative stress-induced apoptosis in an antimycin-induced heart failure model. Inhibiting ALDH2 activity by daidzin increased intracellular ROS levels and

apoptosis in which associated with the upregulated phosphorylation of ERK1/2, JNK, and p38-MAPK [102]. Thus, these data suggested the beneficial role of ALDH2 in drug-induced heart failure.

4 ALDH2-Related Therapy

The following mediators have been reported mediating the cardiac dysfunction in ALDH2 deficiency: (1) ROS and toxic aldehydes [53]; (2) apoptosis pathways [80], involving caspase 3, Bcl- 2; (3) oxidative stress signaling cascade [1], involving MAP kinase cascades ERK1/2, SAPK/JNK, and p38 MAP kinase; and (4) autophagy [1], involving Beclin-1 and AMPK-mTOR. The indispensable role of ALDH2 in the pathogenesis of heart failure sheds light on the development of potential therapeutic target of it (Fig. 1). Here we summarized activators of ALDH2 and several other aspects which have already been proved to have therapeutic effects or just have a therapeutic potential on heart failure.

4.1 ALDH2 Activator

Alda-1, a small molecule activator of ALDH2, is a potential new therapeutic candidate. Alda-1 was reported to exert its cardioprotective effect through reducing oxidative stress, restoring calcium and CaMKII homeostasis, and detoxifying O_2^- induced reactive aldehydes to less reactive acids [16, 93]. It was also been

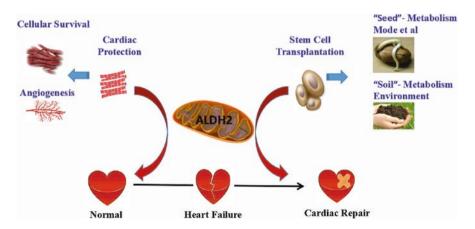


Fig. 1 A scheme depicting how ALDH2 plays the protective role in heart failure. ALDH2 protects myocardium by maintaining myocardial cell survival and angiogenesis. At the same time, ALDH2 can determine the efficacy of stem cell transplantation by increasing the quality of seed stem cells and metabolic microenvironment

reported to ameliorate pulmonary vascular remodeling in pulmonary arterial hypertension and inhibit atherosclerosis in apolipoprotein E-knockout mice [76, 97]. In addition, it played a protective role in cardiac dysfunction induced by abnormal glucose and lipid metabolism as well as DOX or 4-HNE-induced autophagic reduction and cell death [30, 79]. However, the efficiency of the activation by Alda-1 is variant, for example, Alda-1 increases acetaldehyde oxidation in wild-type ALDH2 (ALDH2*1) and East Asian variant of ALDH2 (ALDH2*2) approximately 1.5- and 6-fold, respectively [6]. Thus, the patients undergoing pathological processes such as cardiac ischemia with wild type or mutant ALDH2 might gain more benefits with the treatment of pharmacologic enhancement of ALDH2 activity [16]. Belmont conducted kinetic experiments to characterize Alda-1 on the properties of ALDH2 and found it was a complex behavior, where Alda-1 acted as inhibitor in low concentrations of aldehyde and as an activator in high concentrations [5]. Moreover, Alda-1 has exogenous and toxic characters. All factors should be taken into consideration when considering Alda-1 as an exogenous stimulator of ALDH2, including the working concentration, half-life period, continuity of stimulation, toxic, and side effects.

Protein kinase C ε (PKC ε) has been found as another ALDH2 activator via direct phosphorylation. In vitro, phosphorylation of wild-type ALDH2 recombinant protein was reported increased its enzymatic activity [16]. The enzymatic activity of the phosphorylated ALDH2*2 is 270% of the non-phosphorylated ALDH2*2 [60]. Following phosphorylation by recombinant ε PKC, there was an increase of 70% of the ALDH2 activity. Similar to Alda-1, the effect of ε PKC phosphorylation was more pronounced on ALDH2*2 mutant enzyme. T185E, S279E, and T412E were three common phosphomimetic mutations sites by ε PKC in the protection of ALDH2 against reactive aldehydes. Treatment with PKC activator upregulated ALDH2 activity, while applying PKC inhibitor had the opposite effect. Some study demonstrated that PKC ε -ALDH2 interaction had disincentive effects in 4-HNEinduced aberrant PPAR γ regulation, which suggest that PKC ε -ALDH2 regulatory axis may be a therapeutic target for treating metabolic syndrome [100].

4.2 MicroRNAs

MicroRNAs are a class of endogenous interfering RNAs whose primary function are regulating the expression of genes. Evidences indicated that microRNAs could participate in diverse pathophysiological processes of cardiovascular disease, including hypertrophy, apoptosis, cardiac conduction, fibrosis, and angiogenesis [25, 75]. MiR-34a has been elucidated in many cell lines and was found related to apoptosis, energy metabolism, lipid metabolism, aging, and stem cell division [12, 33]. Bioinformatics analysis produced a protein-protein interaction network in HepG2 cells and revealed that ALDH2 was a potential target of miR-34a [19]. Our study found that increased circulating miR-34a could decrease ALDH2 activity and increase cardiomyocyte apoptosis post-MI injury [23]. Although miR-34a is not a

cardiac-specific miRNA, the expression abundance of miR-34a in myocardium ranks third in all tissues [9], and it is reasonable to regard miR-34a as a diagnostic marker for MI. MiR-28 could also promote ischemia via inhibition of ALDH2 expression in myocardium [45]. In addition, Shen predicted ALDH2 as target genes of miR-224 [72]. The target sites existed in 3'UTR of ALDH2 suggesting that miR-224 downregulated the expressions of ALDH2 and finally regulated target genes in lipid metabolism. MicroRNAs might be new diagnostic indicators and therapeutic targets for heart failure patients. But it requires further studies in large cohort to assess the specificity and sensitivity of them.

4.3 Mesenchymal Stem Cell (MSC) Therapy

Mesenchymal stem cell (MSC) therapy is a promising approach in alleviating ischemic injury and promoting tissue regeneration [64]. A present meta-analysis including 64 studies strongly supported the potential of MSCs therapy for ischemic stroke [68]. Up to now, various strategies have been used to increase transplant effects in ischemic diseases, including tissue engineering scaffolds, genetic modification, and hypoxia-based pretreatment [27]; however, implanted cell dysplasia is still a problem [78]. Many factors could affect the efficiency including regenerative cell source, injectable delivery vehicles, and microenvironmental signals [67, 77]. Our results showed that host ALDH2 affected the survival of transplanted MSCs. Protein array analysis also revealed that ALDH2^{-/-} tissues expressed low levels of angiogenic factors, including cysteine-rich angiogenic inducer 61, endoglin, epidermal growth factor, fibroblast growth factor-1, angiopoietin-1, matrix metallopeptidase-3/-9, and insulin-like growth factor binding protein, all of which could enhance the tolerance of engrafted MSCs during vasculogenesis in hypoxia injury [107]. Thus, ALDH2 may be regarded as a homeostatic mediator of microenvironment by increasing local capillary density and energy supply and decreasing oxidative stress after ischemia.

4.4 ALDH Bright Cells

Autologous bone marrow-derived aldehyde dehydrogenase bright (ALDHbr) cells isolated by flow sorting express high activity of ALDH [41]. They have been applied in clinical practice to repair tissue damage and have been proven safe and efficient in patients with chronic myocardial ischemia [62]. Our observations supported the effective therapeutic effect of ALDHbr cells on ischemic myocardium; in addition, we demonstrated ALDH2 as a key mediator in the process [81]. Weakened glycolysis, mitochondrial respiratory abnormalities, and increased mitochondrial ROS gave rise to the diminished therapeutic efficacy of ALDHbr cells in ALDH2 deficiency mice rather than oxidative phosphorylation impairment. The results gave

us a hint that ALDH2 activity was a pivotal precondition in the efficacy of ALDHbr cell therapy; therefore individuals with loss of ALDH2 function are unsuitable for ALDHbr cell therapy. However, recently a clinical trial didn't find significant positive outcomes of ALDHbr cells in patients with peripheral artery disease [63]. Future investigational trial tests about cell therapy should be carried out to find new anatomic and perfusion insights.

5 Conclusion

Just as Braunwald said, heart failure is the last battle of cardiovascular diseases [11]. It is urgent to find a novel prospective to illustrate the mechanism of heart failure and to improve the prognosis. This review summarizes the roles of ALDH2 gene polymorphisms and ALDH2 enzyme activity in heart failure induced by multiple causes, such as ischemic injury, hypertension, alcohol, diabetes, and aging. Except the controversial role of ALDH2 in aging-related cardiac dysfunction, studies have suggested a cardioprotective role of ALDH2 to counteract cardiac dysfunction due to different etiologies. Emerging evidences provided new insight in understanding the epigenetic and transcriptional regulation of ALDH2 as well as the effect of ALDH2 in stem cell transplantation. Strategies aim to enhancing ALDH2 activity or expression, as well as improving mitochondrial function, will bring novel prospects for the treatment of heart failure.

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