Concurrent diabetes and heart failure: interplay and novel therapeutic approaches

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Abstract:

Diabetes mellitus increases the risk of developing heart failure, and the co-existence of both diseases worsens cardiovascular outcomes, hospitalization and the progression of heart failure. Despite current advancements on therapeutic strategies to manage hyperglycemia, the likelihood of developing diabetes-induced heart failure is still significant, especially with the accelerating global prevalence of diabetes and an ageing population. This raises the likelihood of other contributing mechanisms beyond hyperglycemia in predisposing diabetic patients to cardiovascular disease risk. There has been considerable interest in understanding the alterations in cardiac structure and function in the diabetic patients, collectively termed as "diabetic cardiomyopathy". However, the factors that contribute to the development of diabetic cardiomyopathies is not fully understood. This review summarizes the main characteristics of diabetic cardiomyopathies, and the basic mechanisms that contribute to its occurrence. This includes perturbations in insulin resistance, fuel preference, reactive oxygen species generation, inflammation, cell death pathways, neurohormonal mechanisms, advanced glycated end-products accumulation, lipotoxicity, glucotoxicity, and posttranslational modifications in the heart of the diabetic. This review also discusses the impact of antihyperglycemic therapies on the development of heart failure, as well as how current heart failure therapies influence glycemic control in diabetic patients. We also highlight the current knowledge gaps in understanding how diabetes induces heart failure.

Introduction:

Cardiovascular disease (CVD) is the leading cause of death and complications in diabetic patients worldwide ¹⁻⁶, the prevalence of which is increasing despite therapeutic and pharmacological advances ^{5, 7}. Diabetes mellitus, a metabolic disorder characterized by hyperglycemia resulting from insulin deficiency (type 1) or resistance (type 2)¹¹ is a major independent risk factor in the development of heart failure^{8, 12, 13}, and is becoming a global epidemic with increasing prevalence ¹⁴⁻¹⁷. There are 2 main types of diabetes mellitus: type 1 and type 2; gestational diabetes mellitus is also part of this category but is not a major focus of this review. Type 1 diabetes mellitus, also called insulin-dependent diabetes, is an autoimmune disease where pancreatic beta cells are destroyed, and therefore, the body is unable to produce insulin ¹⁸. Type 2 diabetes mellitus, also called non-insulin-dependent diabetes, is characterized by a deficit in the function of insulin produced by pancreatic beta cells; this is also referred to as insulin resistance. Type 2 diabetes is the more common form of diabetes, and factors such as age, obesity, diet, and pre-existing hypertension affect its development and its risk of development. Therefore, due to either the elimination of insulin secretion or the reduction in insulin function, blood glucose levels are, as a result, elevated, leading to chronic hyperglycemia if left untreated. The Framingham Study was one of the first epidemiological studies to show an increased risk of heart failure in patients with diabetes mellitus ^{13, 19, 20}, with other clinical trials supporting this conclusion (see ^{21, 22} for reviews). The presence of both diabetes and heart failure in individuals leads to poor cardiovascular outcomes ^{5, 6, 23-26}. Diabetic patients have higher mortality from coronary artery disease than non-diabetics

²⁷ and show a worse prognosis ^{28, 29}, which may be associated with increased atherosclerosis ²⁷, and can lead to ischemic heart failure ³⁰. Heart failure is the major adverse cardiovascular outcome in diabetic patients ²⁰. Poor glycemic control is associated with increased risk of heart failure in individuals with type 2 diabetes ³¹, indicated by elevated hemoglobin A_{IC} levels, an index of glycemic control ³². Therefore, it is critical to mediate and treat cardiovascular conditions in diabetic patients. The prevention and treatment of cardiovascular disease and heart failure remains a considerable challenge in the treatment and management of diabetes mellitus.

Diabetic cardiomyopathy is a condition characterized by ventricular dysfunction and hypertrophy in diabetic patients independent of hypertension, ischemia, or coronary artery disease ³³. The term originated from a Rubler et al. study that identified diabetic patients with congestive heart failure without the aforementioned risk factors or other causes. Noninvasive studies show impaired diastolic and systolic function ³⁴⁻³⁷, especially with the presence of hypertension. Multiple mechanisms contribute to the development of heart failure in diabetic individuals , including increased inflammation ³⁸ and oxidative stress ^{39, 40}, changes in cardiac myocardial energy metabolism ⁴¹⁻⁴³, cardiac lipotoxicity ⁴⁴⁻⁴⁷, impaired cardiomyocyte calcium handling, and apoptosis ^{48, 49}. Diabetic cardiomyopathy may progress to either heart failure with preserved ejection fraction (HFpEF), where there is diastolic dysfunction ^{8, 52, 53}. Each phenotype can be distinguished by the various mechanisms involved in contributing to it. Hypertrophy ⁵⁴⁻⁵⁶, insulin resistance ⁵⁷, and lipotoxicity ^{58, 59} have been shown to contribute to the development of HFpEF, while

oxidative stress ⁶⁰, fibrosis ^{61, 62}, and autoimmunity caused cardiomyocyte cell death ⁶³ are involved in contributing to HFrEF. Coronary deposition of advanced glycation endproducts is involved in both phenotypes ⁶⁴. Diastolic dysfunction, which is part of the diagnosis of HFpEF ^{50, 51, 65}, can precede the heart to develop HFpEF alongside comorbidities of impaired coronary vasculature and endothelial function, and hypertrophy ^{66, 67}. Although diastolic dysfunction is mostly predominant in diabetic cardiomyopathy, systolic dysfunction may also occur in later stages of diabetic cardiomyopathy, which can contribute to the development of HFrEF ^{52, 53}. Studies have demonstrated a metabolic link between the 2 heart failure phonotypes whereby abnormal mitochondrial function and oxidative stress can lead from a HFpEF phenotype into a HFrEF phenotype by mediating cardiac hypertrophy, inflammation, fibrosis, and further endothelial cell damage, which has adverse consequences on systolic function, and ultimately more severe manifestations of diabetes and heart failure ^{54, 68-77}.

The aim of this review is to highlight the changes that occur in diabetic cardiomyopathy, alongside the mechanisms involved its development and progression. The effect of antihyperglycemic drugs on heart failure risk in diabetic individuals, and heart failure drugs on glycemic control will also be discussed, as well as novel therapeutic approaches.

Structural and functional characteristics of the failing heart in diabetics

The failing heart in the context of diabetes is characterized by multiple alterations including impairments in diastolic and subsequent systolic function ^{34-36, 50}, cardiac hypertrophy and fibrosis ⁹⁹ and impaired coronary microvascular perfusion ¹⁰⁰.

Diastolic and systolic dysfunction: Heart failure in diabetes is characterized by cardiac dysfunction, with diastolic dysfunction as a hallmark of the failing myocardium in diabetics ^{34, 101}. Echocardiography and doppler imaging assessments of diastolic dysfunction ¹⁰², have shown that left ventricular (LV) dysfunction is manifested in type 2 diabetic patients through altered LV filling ¹⁰³, abnormal LV relaxation ^{34, 104, 105}, reduced LV end diastolic volume ¹⁰⁶, and LV chamber stiffness ¹⁰⁷. Studies in type 1 diabetics demonstrate abnormalities in LV diastolic filling ^{35, 108, 109}, lower E to A ratios, prolonged isovolumetric relaxation times ³⁵, and reduction in end-systolic volumes ³⁶. Thus, there is no single parameter to indicate and quantify diastolic dysfunction . Moreover, Attali et al. showed in both type 1 and 2 diabetic patients that abnormalities in diastolic function, including increased isovolumetric relaxation time and impaired LV compliance, were not related to additional factors such as age, sex, duration of diabetes, or presence of other complications ¹¹⁰. Impaired diastolic function has also been shown in type 1 diabetic children ^{36, 111}. Additionally, the presence of hypertension can aggravate diastolic dysfunction, as demonstrated through further and severe impairment of LV relaxation and abnormal LV filling ⁵⁰. Speckle tracking echocardiography has emerged as a novel beneficial diagnostic method for early detection of LV dysfunction in diabetes, therefore being useful to detect LV abnormalities ¹¹². This method has been shown to overcome some of the limitations of transthoracic doppler imaging ¹¹³ and can be an equally, if not better, powerful approach to assessing myocardial velocities and strain ¹¹³⁻¹¹⁵. Studies have shown it's usefulness through examining LV strain in hypertensive and type 2 diabetic patients ¹¹⁶, and LV rotational mechanics in hypertensive type 2 mellitus diabetic patients ¹¹². This echocardiography method has also been utilized in animal models of diabetes, namely assessing systolic strain and contractile function in db/db mice by Li et al. ¹¹⁷ and assessing cardiac dysfunction in rat models of type 1 and type 2 diabetes mellitus by Matyas et al. ¹¹⁸.

Experimental evidence in animals complements these observations in human studies, showing a decrease in end-diastolic volume in alloxan diabetic dogs ¹¹⁹, a reduced E and A transmitral flow in *db/db* mice ¹²⁰; and an increased isovolumetric relaxation time and increased LV end-diastolic pressure in streptozotocin induced non-insulin dependent rats compared to controls ¹²¹. Additionally, Otsuka Long-Evans Tokushima fatty rats show increased deceleration time ¹²².

Systolic dysfunction is also present in diabetic cardiomyopathy, although in both human and animal studies it has been shown to take longer to develop and usually occurs after diastolic dysfunction ^{36, 120}. In human studies, this manifests mainly as a reduction in ejection fraction ^{123, 124}, along with increased LV end-systolic volume ¹²³ and reduced fractional shortening ¹²⁵. The Strong Heart Study showed systolic dysfunction to occur, as evidenced by lower LV fractional shortening and decreased stress corrected midwall shortening in diabetic patients ¹²⁶. Animal studies are consistent with this, demonstrating impaired systolic function through lower peak-developed pressures ¹²¹, +dP/dt, peak

emptying rates ¹²⁷, peak filling rates, fractional shortening ¹²⁸, and systolic blood pressure in streptozotocin induced diabetic rats.

Impaired diastolic function is either associated with normal systolic function ^{50, 104, 105, 129, 130} or can precede systolic dysfunction ³⁶. In support of this, Raev *et al.* showed diastolic damage and abnormalities to be more prevalent than systolic damage and abnormalities in type 1 diabetic patients, while systolic dysfunction occurred much later in the progression of diabetes ³⁶. However, Fang *et al.* believe the use of less sensitive techniques to measure systolic dysfunction accounts for the reason behind studies demonstrating diastolic dysfunction with normal systolic function .

Cardiac hypertrophy: Diabetic cardiomyopathy is often associated with left ventricular hypertrophy ⁹⁹. The Strong Heart Study showed an independent association between diabetes and cardiac hypertrophy ¹²⁶. Increased myocardial wall thickness can be seen in type 1 and 2 diabetes, alongside ventricular dysfunction ^{105, 126, 131}. Additionally, myocardial hypertrophy is linked to adverse cardiovascular outcomes, including being a predictor of cardiovascular death ^{132, 133}. The Framingham study demonstrated that increased LV mass is associated with increased risk of cardiovascular outcomes of mobility and mortality ¹³³. Moreover, Solomon *et al.* observed a slightly greater wall thickness in diabetic patients alongside decreased ventricle size, which they believe may be associated with higher filling pressures and diastolic dysfunction ¹³⁴. These findings have been confirmed in animal studies showing an increase in LV mass alongside impaired LV relaxation and increased chamber stiffness ¹²¹. The observed LV hypertrophy

in diabetics may precede the onset of systolic dysfunction , and can additionally be used as a diagnostic indicator in the development of heart failure in diabetics ¹³⁵.

Cardiac fibrosis: Myocardial fibrosis and collagen accumulation can manifest as a major structural alteration in the setting of diabetes ^{136, 137}, which can lead to myocardial damage and heart failure ¹³⁸⁻¹⁴¹. Multiple human studies demonstrate the presence of fibrosis in the left ventricle, alongside collagen accumulation in the interstitial and perivascular region of diabetic patients ^{99, 136, 142}. This cardiac fibrosis is associated with cardiac dysfunction ^{55, 140}, and may lead to worsened cardiac outcomes including developing congestive heart failure ^{55, 139, 143}.

Increased cardiac fibrosis in diabetes is supported by animal studies, where multiple mechanisms may be responsible for the cardiac fibrosis observed in diabetes. Otsuka Long-Evans Tokushima Fatty rats, a model of type 2 diabetes, show increased myocardial collagen content that is associated with impaired diastolic function through prolonged deceleration times and decreased early filling wave peak velocities ¹²². Streptozotocin-induced diabetic rats have increased collagen and interstitial fibrosis due to oxidative stress, alongside decreased cardiac contractility ^{144, 145}. Additionally, streptozotocin-induced diabetic mice show a time dependent increase in LV collagen content, alongside impaired diastolic and systolic function ¹³⁸. This is suggested to be due to reduced matrix metalloproteinase 2 (MMP2) levels. Spiro *et al.* also showed an increase in type IV collagen in the myocardium of diabetic rats ¹⁴⁶. The increase in cardiac collagen in diabetes may be due to increased transforming growth factor–B1 (TGF-B1)

receptor II expression ^{71, 122, 147}. Another factor that can mediate this collagen accumulation and fibrosis development in diabetes is the myocardial accumulation of advanced glycosylation end products (AGEs) ^{148, 149} which will be discussed in more detail in a subsequent section of this review.

Impaired coronary microvascular perfusion: Abnormalities in coronary artery function and circulation are highly prevalent in diabetes ¹⁵⁰, which may predispose the diabetic myocardium to cardiac damage and disease, including ischemia due to impaired blood circulation and flow ^{150, 151}. Coronary flow reserve (CFR) is reduced in both type 1 and 2 diabetic patients ^{100, 152}, along with a reduction in coronary vasodilation ¹⁵³ due to reduced nitric oxide (NO) production ^{154, 155}. Marciano *et al.* showed that type 2 diabetic patients without coronary artery disease have impaired coronary microvascular function, demonstrated by a lower CPT-CF ratio (cold pressure test to coronary flow), compared to non-diabetic individuals ¹⁵⁶. Additionally, Bagi et al. showed enhanced superoxide production and decreased NO production, leading to reduced coronary dilation in coronary arterioles isolated from db/db mice ¹⁵⁷. Hyperglycemia may be a cause for this, as shown by an association between CFR and HBA1c levels ¹⁵⁸. Additionally, Durante et al. showed lower coronary flows in diabetic BB rats in response to stimulation by noradrenaline, calcium, or tachycardia ¹⁵⁹. Coronary microvascular perfusion may be further impaired by the presence of hypertension-induced vascular lesions ¹⁶⁰. A reduction in coronary capillary density in the diabetic myocardium has also been observed, due to lower angiogenesis as a result of decreased vascular endothelial growth factor (VEGF) expression ¹⁶¹⁻¹⁶³. VEGF and VEGF receptor mRNA and protein expression were shown

to be significantly decreased in diabetic and insulin-resistant non-diabetic rats ^{161, 162}, and was accompanied by decreased myocardial perfusion and LV dysfunction ¹⁶². Together, this suggests that structural abnormalities occur alongside functional abnormalities in the coronary microvasculature in diabetes. However, it is not clear which precedes the other, as some studies suggest structural changes in coronary arterial vasculature may be involved in causing further cardiac dysfunction ¹⁶⁴ and also concurrently progress as diabetic cardiomyopathy progresses ^{162, 165}. Additionally, Giordano et al. showed that VEGF is a critical determinant of cardiac function as a VEGF knockout mouse model resulted in contractile dysfunction ¹⁶⁶. Therefore, further studies need to be done to fully elucidate the interplay in sequence of events between abnormalities in coronary capillary density and cardiac function.

Additionally, impaired coronary flow reserve and vasodilation may be an early marker of atherosclerosis, which can lead to progressive deterioration of the myocardium ¹⁵¹ and an increase in the risk of cardiovascular disease ¹⁵² and ischemia ¹⁵³. Impaired vasodilation in diabetes ¹⁵⁵, which may also be due to NO production inhibition due to hyperglycemia ¹⁶⁷ through the generation of oxygen-derived free radicals ¹⁶⁸, can also lead to arterial atherosclerosis . Therefore, this abnormal coronary flow may greatly increase the risk and likelihood of myocardial ischemia ^{153, 159, 169}.

Underlying mechanisms contributing to the development of diabetic cardiomyopathy

Insulin resistance: Insulin resistance is one of the early contributing factors to the development of diabetic cardiomyopathy¹⁷⁰. The decreased efficacy of insulin to lower blood glucose levels occurs as a result of hyperinsulinemia-mediated excessive insulin receptor signaling or downregulation of insulin receptor signaling. This insulin resistance contributes to a number of adverse changes in the heart that include alterations in cardiac energy metabolism, increased inflammation and hypertrophy, lipotoxicity, glucotoxicity, alterations in mitochondrial function and ROS production, accumulation of advanced glycation products and O-GlcNAcylation, alterations in cardiac cardiomyocyte Ca²⁺– handling, systemic hyperglycemia and hyperlipidaemia¹⁷¹, and neurohormonal changes (all of which are discussed below).

It is important to note that cardiac insulin resistance precedes the development of cardiac dysfunction and heart failure. A study in mice with heart failure developed diastolic dysfunction at 2 weeks and systolic dysfunction at 3 weeks. Notably, the decline in function was preceded by significant cardiac insulin resistance which was determined via serial and direct measurements of insulin-stimulated glucose metabolism in isolated working hearts ⁴³. These findings are supported by epidemiological studies that also found that insulin resistance is a predictor, rather than a biomarker, of heart failure. In a study of 1187 elderly men that did not have congestive heart failure, the epidemiological study between 1990 and 1995 found that insulin resistance significantly increased the risk and predicted congestive heart failure ¹⁷². Another study in 431 50-year-old men with a 20-year follow-up, patients that developed heart failure at age 70 presented increased

plasma proinsulin at age 50, signifying that insulin resistance preceded cardiac dysfunction ¹⁷³.

Insulin signaling begins with insulin binding to the insulin receptor, resulting in activation of the insulin receptor substrate-1/2, PI3K/PKB (Akt) activation, GLUT4 translocation to the cell membrane, stimulation of mitochondrial glucose oxidation, and inhibition of fatty acid oxidation ¹⁷⁴. Cardiac muscle biopsies from type 2 diabetic patients have depressed PI3K/PKB signaling and decreased GLUT4 expression¹⁷⁵. In addition to decreased translocation of GLUT4, impaired PI3K engagement and stimulation of Akt also occur, due to increased phosphorylation of the serine residue on IRS-1/2¹⁷⁶.

Activation of forkhead box-containing proteins regulates insulin signaling, leading to insulin resistance. FoxO proteins are elevated in mice with high-fat diet-induced diabetes, which downregulates IRS1, consequently leading to decreased Akt signaling, insulin resistance and the development of diabetic cardiomyopathy¹⁷⁷. Also important in the regulation of insulin signaling, and is perturbed in diabetic cardiomyopathy, is the E3 ubiquitin ligase – mitsugumin 53 ¹⁷⁸. In support of this, cardiac-specific overexpression of mitsugumin 53 in mice results in severe diabetic cardiomyopathy and insulin resistance, due to degradation of the insulin receptor and IRS-1. Mitsugumin 53 overexpression is involved in transcriptionally upregulating PPARα, contributing to lipid accumulation¹⁷⁹. This accumulation of lipid intermediates (diacylglycerol and ceramides) contributes to the development of insulin resistance^{180,181}. Conversely, decreasing

myocardial levels of ceramide and diacylglycerol is accompanied by improvements in insulin sensitivity and myocardial glucose utilization¹⁸².

Altered cardiac energy metabolism: The heart has a very high energy demand despite having very low ATP stores (ATP levels effectively turnover in the heart every 5-10 seconds)¹⁸³. The heart has the ability to continually generate large amounts of ATP from various energy substrates, including fatty acids, glucose, lactate, ketone bodies and amino acids, regardless of workload, nutritional status, and hormonal status ^{183, 184}. However, this metabolic flexibility is impaired in many forms of heart disease, including diabetic cardiomyopathy¹⁸³. Insulin resistance results in an increase in myocardial fatty acid oxidation rates in diabetic cardiomyopathy and impaired glucose oxidation rates (Figure 1) ¹⁸⁵⁻¹⁸⁷. This increases myocardial oxygen consumption, decreases cardiac efficiency and strongly correlates with impaired cardiac contraction and diastolic function ^{41, 42, 84, 189}.

Multiple mechanisms contribute to the increased reliance of the heart on fatty acid use during diabetes. The first such mechanism is increased supply of fatty acids to the heart. The lack of insulin suppressive action on adipose tissue results in the release of fatty acids from adipocyte to the circulation, leading to elevation of blood plasma free fatty acid levels. Fatty acid delivery and uptake to the heart is also increased in diabetes, due to an increase in cardiac myocyte lipoprotein lipase activity and increases in sarcolemmal CD36 protein expression, respectively ^{190, 191}. The increased uptake of fatty acids across the sarcolemma is facilitated by at least three proteins, namely, CD36, FA transport

protein (FATP), and FA binding protein plasma membrane (FABPpm) ¹⁹². In STZ-induced type 1 diabetic rats and type 2 *db/db* mice an upregulation of CD36 and FABPpm protein expression occurs ^{193, 194}. While CD36 alone accounts for more than half of the total fatty acid taken up by cardiomyocytes ¹⁹⁵, both its expression and membrane localizations are increased in diabetes^{193, 195}.

Activation of transcription regulators such as peroxisome proliferator-activated receptors (PPARs) can promote expression of genes that facilities fatty acid uptake, storage and oxidation in the heart ¹⁹⁶⁻¹⁹⁸. Myocardial PPAR α expression is increased in type 2 diabetes, and mice lacking PPAR α are protected from the development of diabetic cardiomyopathy ^{41, 199}. In contrast, a recent study found no differences in the risk of cardiac dysfunction between wildtype and PPAR α deficient mice subjected to a low dose of streptozotocin (STZ) ²⁰⁰. The discrepancy in the findings could be due to the different methodology followed for inducing Type 1 diabetes (1 single injection of high STZ dose vs. 5 daily injections of lower dose) and/or the different time points of cardiac function assessment (6 weeks vs 9-12 weeks post-STZ administration). Interestingly, it has been shown that PPAR α / γ alterations may contribute to cardiac dysfunction independent of changes in fatty acid oxidation or lipid storage in non-diabetic animals ^{201, 202}.

One effect of increased PPARα in diabetes is an increase in mitochondrial carnitine palmitoyltransferase I (CPT-1) expression, a key enzyme involved in mitochondrial uptake and oxidation of fatty acids ²⁰³. In addition, perturbations in CPT-1 regulation also occur in diabetes. CPT-1 is inhibited by malonyl CoA produced from acetyl CoA by acetyl CoA

carboxylase (ACC) ²⁰⁴. Activation of AMP-activated protein kinase (AMPK) in diabetes inhibits ACC activity. Decreased ACC activity with parallel increases in malonyl-CoA decarboxylase activity ²⁰⁵ decreases malonyl CoA levels, resulting in decreased inhibition of CPT1 and accelerated fatty acid oxidation rates Figure 1) ²⁰⁶⁻²⁰⁸. Post-translational modification of fatty acid oxidative enzymes also occurs in diabetes, resulting in an increase in fatty acid oxidation ²⁰⁹. Increased acetylation of major fatty acid metabolic enzymes due to decreased SIRT3 also leads to up-regulation of fatty acid oxidation and impaired glucose metabolism in the heart ²¹⁰⁻²¹². In addition to increases in myocardial fatty acid oxidation, an increase in myocardial triacylglycerol content is seen in diabetics ²¹³. Increased myocardial uptake of fatty acids leads to the increased accumulation of lipids and their intermediate metabolites, such as long and short fatty acyl-CoAs, diacylglycerol ^{196, 214}. However, studies on the turnover of endogenous fatty acid in hearts from diabetic animals have generated variable results. We have shown an increased myocardial lipolysis rate in diabetic hearts irrespective of exogenous fatty acid concentration while endogenous synthesis rate remains unaffected ²¹³. This is further supported by a ¹³C-NMR isotopic enrichment study in diabetic rat hearts ²¹⁵. In contrast, others reported reduced or unchanged lipolysis and increased synthesis in the hearts of diabetics in the presence of high levels of exogenous free fatty acids ^{216, 217}. On the other hand, decreased levels of myocardial phospholipids is seen in diabetes together with impaired synthesis ²¹⁸. While the precise contribution of altered phospholipid metabolism is less clear in diabetes, various studies have suggested the etiologic role of phospholipid (membrane lipid) metabolic dysregulation in lipotoxic cardiomyopathy and other forms of myocardial dysfunction ^{219, 220}.

In contrast to the increased myocardial uptake and oxidation of fatty acids seen in diabetes, myocardial glucose transport, glycolysis and glucose oxidation are decreased in diabetes ²²¹⁻²²⁴. Total myocardial GLUT4 and GLUT1 expression are decreased in diabetes ²²⁵. Decreased myocardial glycogen content along with a reduced myocardial glycogen synthesis rate and impaired glycogen synthase enzyme activity is also reported in hearts of diabetics ²²⁶⁻²²⁸. Insulin deficiency or resistance compromises the glucose transport and utilization in the heart. Also, the presence of excess fatty acid derivatives, such as fatty acyl CoA, diacylglycerol, and ceramide, leads to the inhibition of insulin signaling in the heart ^{229, 230}. In addition to decreased glucose transport, inhibition of cardiac phosphofructokinase (PFK-1), the rate limiting enzyme in glycolysis, is seen in diabetes ^{231, 232}. PFK-1 is inhibited allosterically by high levels of citrate, high ATP levels, and increases in NADH, that are derived from increased fatty acid oxidation. Glucose oxidation is also decreased, due in part to increases in fatty acid oxidation, which inhibits the rate limiting enzyme of glucose oxidation - pyruvate dehydrogenase (PDH) ^{187, 221, 233}. PPAR- α activation also suppresses glucose uptake and utilization by increasing the expression of pyruvate dehydrogenase kinase 4 (PDK-4), which inhibits PDH ^{47, 234}.

Similar trends of fatty acid and glucose metabolic shifts have been also observed in type 1 and 2 diabetic patients ^{59, 235-238}. In ³¹P and ¹H magnetic resonance spectroscopy studies, a significant reduction in myocardial glucose utilization accompanied by reduced myocardial energetics (phosphocreatine to ATP ratio (PCr/ATP) and increased myocardial fatty acid metabolism and triacylglycerol content is seen in type 2 diabetic

patients ^{236, 237}. The increased rates of myocardial fatty acid oxidation persist even after insulin treatment in human type 2 diabetic patients ²³⁵. The levels of circulating free fatty acids is also negatively correlated with altered PCr/ATP ratios in patients with diabetes ²³⁹. Earlier studies recognized myocardial PCr/ATP ratios as a predictor of cardiovascular mortality in patients with dilated cardiomyopathy ²⁴⁰. Likewise, increased myocardial fatty acid utilization with a concomitant decrease in glucose utilization is seen in type 1 diabetic patients ^{59, 238}.

Since fatty acids and glucose are the two important fuels for the heart, their balanced use is critical for maintaining normal contractile function. As a result, the decreased "metabolic flexibility" and increased reliance of the heart on fatty acid as source of energy is associated with impaired myocardial function in diabetes ^{222, 241-244}. Enhanced fatty acid oxidation increases myocardial O₂ consumption and decreases cardiac efficiency ²⁴⁵. Enhanced fatty acid oxidation also alters the mitochondrial NADH to NAD⁺ ratio and acetyl CoA levels which can further modify several intracellular signaling processes²¹⁰. Although evidence suggests a detrimental effect of increased fatty acid oxidation on heart function in diabetes, there are still opposing views on the role of glucose and fatty acid alterations and pathologic significance in non-diabetic heart failure. Relatively few studies have combined diabetes. By combining high fat feeding with pressure overload hypertrophy in mice we observed a marked decrease in insulin-stimulated glucose oxidation that was associated with both diastolic and systolic dysfunction ^{246, 247}. Furthermore, nutritional

strategies that increase insulin-stimulated glucose oxidation are accompanied by a decreased severity of heart failure ^{246, 247}.

The contribution of alterations in the use of other fuels such as ketone bodies and branched chain amino acids (BCAAs) is increasingly being recognized in heart failure pathogenesis ²⁴⁸⁻²⁵⁰. While ketone oxidation is increased in HFrEF and may be an adaptive process to maintain energy production ^{250, 251}, in diabetes myocardial ketone oxidation is impaired, and may result in a decrease in metabolic flexibility and a decrease in energy production in the heart ^{252, 253}. A decrease in BCAA oxidation in insulin resistant hearts also contributes to an impaired insulin signaling and a decrease in insulin-stimulated glucose oxidation ²⁵⁴⁻²⁵⁶.

Cardiac lipotoxicity and glucotoxicity: Under normal circumstances, the uptake and oxidation of fatty acids are finely regulated resulting in only little myocardial lipid storage. However, in diabetes a persistent elevation in circulating free fatty acids supplies the heart with excess fatty acids and promote accumulation of lipids in the cardiomyocytes (cardiac lipotoxicity) (Figure 2) $^{257, 258}$. The causative role of excess lipid accumulation in diabetic cardiomyopathy has been demonstrated using genetic or pharmacological approaches that modify uptake or oxidation of fatty acids. For instance, mice with cardiac specific overexpression of PPAR- α exhibit increased uptake and utilization fatty acid and typical features of diabetic cardiomyopathy, including ventricular hypertrophy and systolic dysfunction⁴⁷. Recently, an increased activity of lipoprotein lipase (LPL) as shown in epicardial adipose tissue from type 2 diabetic patients 259 . Interestingly, elevated activity

of LPL was associated with increased epicardial adipose tissue volume, suggesting increased fatty acids uptake. Of interest, deletion of CD36 or cardiac LpL rescues mice from a lipotoxic-induced cardiomyopathy caused by PPARα overexpression ^{260, 261}.

In contrast, overexpression of CD36, FATP1, or acyl CoA synthetase results in lipotoxocity ^{262, 263}. This lipotoxicity correlates with diastolic dysfunction and other pathophysiological findings related to diabetic cardiomyopathy ^{264, 265}. Despite their contributing role in inducing cardiac lipotoxicity in diabetes, PPARα agonists (fibrates) are still in use clinically to treat hypertriglyceridemia ²⁶⁶. In theory, lipotoxicity could arise either due to increased uptake or decreased oxidation. However, studies on pharmacological inhibition of FA oxidation or genetic manipulation of fatty acid oxidation enzymes revealed that decreased fatty acid oxidation does not actually lead to lipid accumulation ²⁶⁷. It has been hypothesized that the reduction in oxidative function may inhibit the uptake of fatty acid by feedback mechanism²⁶⁸.

In addition to fatty acid overload, cardiac lipotoxicity is also dependent on the type of fatty acids or lipids accumulated ²⁶⁸. Ceramide is one fatty acid derivative strongly associated with cardiac lipotoxicity. Inhibition of ceramide synthesis, either by deletion of serine palmitoyltransferase or pharmacologically by myriocin, results in significant metabolic and structural changes to the heart. Decreasing cardiac ceramide levels decreases heart weight, pyruvate dehydrogenase kinase 4 (PDK4) expression, fatty acid oxidation rates and left ventricular diameter, while improving glucose oxidation ^{269, 270}. Diacylglycerol (DAG) is another lipid derivative associated with cardiac lipotoxicity. Increased levels of

DAG in in the heart is associated with biochemical changes and macrovascular remodeling, indicating its possible role in the development of diabetic complications ²⁷¹. Increased levels of both ceramide and DAG can also activate and facilitate the translocation of protein kinase C (PKC) to the cell membrane ^{271, 272}. Activation of PKC by excess lipids impairs β -adrenergic signaling in the heart by phosphorylating its receptor ^{273, 274}. Phosphorylation of the β -adrenergic receptor leads to its desensitization, resulting in reduced myocardial contractility in response to catecholamines ²⁷⁵. Increased PKC activity and its translocation to the cell membrane can also attenuate insulin signaling. Previous studies have shown that PKC can phosphorylate insulin receptor substrate 1 at its serine residue and blocks insulin stimulated tyrosine phosphorylation and downstream Akt signaling ^{276, 277}.

On the other hand, although triacylglycerol is the most abundant lipid that accumulates in the heart, studies suggest that its accumulation is not associated with toxic effects in the heart ^{278, 279}. Overall, these data demonstrate that excess fatty acid storage and utilization in the heart are detrimental to heart function, although the mechanistic link between lipid accumulation and cardiomyopathy development are not clearly defined.

In contrast to lipotoxicity, less is known about glucotoxicity. Although myocardial glucose transporters are down regulated in diabetes, the heart can be still exposed to excess glucose. The increased extracellular glucose concentration results in the build-up of a glucose gradient for its transporter across the sarcolemma by mass action ²⁸⁰. As myocardial glucose oxidation is inhibited in diabetes ^{187, 221, 281}, the increased glucose flux

can lead to the accumulation of glycolysis intermediates and products. This imbalance can drive the diversion of glycolytic intermediates into pathological pathways in diabetes including protein kinase C stimulation, the hexosamine pathway, the polyol pathway, and the formation of advanced glycation end products ^{280, 282}. Increased glucose uptake in GLUT4 transgenic mice also contributes to mitochondrial dysfunction via O-GlcNAcylation of the transcription factor ST1 and many electron transport chain subunits ²⁸³.

Impaired mitochondrial function: The impact of heart failure and diabetes on mitochondrial bioenergetics has long been established. Perturbations in mitochondrial oxidative metabolism and mitochondrial ROS generation occur in both heart failure and diabetes ^{83, 284-286}. The reduction in cardiac function and efficiency along with impaired cardiac mitochondrial bioenergetics in obesity and diabetes is due to, at least in part, the excessive reliance on fatty acid oxidation and increased uncoupling protein content in these hearts, that contribute to reactive oxygen species (ROS) production ^{246, 287}. Many studies have proposed that ROS overload is a major culprit of diabetic cardiomyopathy ²⁸⁸⁻²⁹¹. Mitochondria are a major source of ROS production, and increased fatty acid oxidation can promote ROS production ^{83, 246, 287, 292-294}. Excessive cardiac ROS production induces inflammation and activates many crucial mediators of pathological signaling cascades ²⁹⁵⁻³⁰², such as protein kinase C (PKC), apoptosis signal-regulating

kinase-1 (Ask1), p38 mitogen activated protein kinase (p38-MAPK), NH2-terminal Jun kinases (JNK), and JAK-STAT. Activation of these signaling cascades can contribute to

the complications of diabetic cardiomyopathy ^{303, 304}. Furthermore, a recent study demonstrated that enhanced activity of Krüppel-like factor-5 (KLF5) is linked to an increase in oxidative stress in diabetic cardiomyopathy ²⁰⁰. This occurs through an upregulation in the expression of NOX4 via direct binding to the NOX4 promoter ²⁰⁰. The accumulation of ROS, increased ceramide production and low mitochondrial abundance contributes to impaired cardiac function in the hearts of the diabetics ²⁰⁰.

There have been many studies that have looked at the efficacy of antioxidants in managing diabetic cardiomyopathy ³⁰⁵⁻³¹². Antioxidants can mitigate ROS-mediated mitochondrial uncoupling, a characteristic of diabetic cardiomyopathy, in animal studies ^{313, 314}. Similarly, antioxidants are protective against mitochondrial ROS in the failing heart ³¹⁵⁻³¹⁷. Additionally, preclinical trials specifically targeting mitochondrial ROS respiratory complexes have been positive ³¹⁸⁻³²⁰, but further clinical trials are necessary to confirm the efficacy of mitochondrial ROS scavenger under the contexts of both general heart failure and diabetic cardiomyopathy. Additionally, the use of Nrf2 activator and NOX inhibitor have shown to be effective in animal models^{321, 322}, and calorie restriction can lower ROS production and UCP expression in Type II diabetic Otsuka Long-Evans Tokushima Fatty (OLETF) rats ^{299, 323}. Unfortunately, human clinical trials have failed to replicate these observations from animal models ³²⁴⁻³²⁷.

Inflammation and hypertrophy: Diabetes leads to increases in intramyocardial inflammation, characterized by increases in cell adhesion molecules (ICAM-1, VCAM-1) and increased macrophage infiltration resulting in the release of inflammatory cytokines

(IL-1 β , IL-6, IL-18, TGF- β 1, TNF- α) ²⁹¹. Plasma concentrations of the cytokine acutephase mediators, TNF- α and IL-6, are increased in the circulation in settings of impaired glucose tolerance, and thus, inflammation has been shown to be predictive for type 2 diabetes ³²⁸⁻³³⁰. This is due to an excess level of glucose and free fatty acids stressing both pancreatic islet cells and adipocytes, resulting in the release of pro-inflammatory cytokines and chemokines into the circulation that promote inflammation in other tissues such as the heart ³³¹. Plasma TNF- α and IL-6 levels are increased and are associated with left ventricular diastolic dysfunction in patients with diabetes³³².

Systemic and local inflammation leads to fibrosis in the myocardium as well as hypertrophy and apoptosis at the level of the cardiomyocytes ³³³. An upregulation of inflammatory signaling results in macrophage infiltration, cardiomyocyte apoptosis, hypertrophy and a profibrotic response via extracellular matrix remodeling – all of which lead to impaired cardiac contractility and diabetic cardiomyopathy ³³⁴⁻³³⁶. Macrophage and lymphocyte infiltration into the cardiac cell are followed by secretion of pro-inflammatory cytokines (TNF- α , IL-6, IL-1 β , TGF β , interferon- γ), which leads to adverse cardiac remodeling.

Due to systemic accumulation of advanced glycation end products, angiotensin II and lipotoxicity, an increase in toll-like receptor 4 (TLR4) and tumor necrosis factor receptor 1 (TNFR1) occurs in diabetes, leading to secretion of pro-inflammatory cytokines and subsequent cardiomyocyte death, hypertrophy, metabolic imbalances, contractile dysfunction, oxidative stress and mitochondrial dysfunction ^{336, 337}. More specifically, high

mobility group protein B1 (HMGB1) mediates lipopolysaccharide binding to and activation of TLR4, resulting in downstream activation of nuclear factor-KB (NF-KB) and the NLRP3 inflammasome ³³⁸. Activation of the pleiotropic transcription factor, NF-KB, results in the transcription of genes that are pro-inflammatory (MCP-1, COX-2, VCAM-1), pro-hypertrophic (ANP, myosins) and pro-fibrosis (TGFβ, collagens, FN) ^{339, 340}.

Hypertrophy follows inflammation in the heart of diabetics, since cytokines can induce cardiomyocyte hypertrophy^{336, 341-343}. The pro-inflammatory cytokine TNF- α can activate the JNK and AKT/NF- κ B pathway to promote cardiomyocyte hypertrophy³⁴⁴. Activation of the NF- κ B pathway can also result in cardiomyocyte growth ³⁴⁵. IL-1 β , through IGF-1 downstream release from cardiac fibroblasts promotes cardiomyocyte hypertrophy ³⁴⁶. Furthermore, IL-6 also contributes to cardiomyocyte hypertrophy through activation of the CaMKII and gp130 pathways which then activates the STAT3 pathway³⁴⁷. Lastly, TGF- β can activate the TAK1-MKK3/6-p38MAPK pathway and PKC-ATF2 to promote cardiomyocyte hypertrophy ^{348, 349}.

Myocardial inflammation can also lead to cardiomyocyte apoptosis which subsequently contributes to cardiac remodeling. TNF- α activates both extrinsic and intrinsic apoptotic pathways, as well as NF- κ B, to promote cardiomyocyte death ^{350, 351}. Additionally, through NO synthase activation or CHOP, IL-1 β promotes apoptosis in cardiomyocytes³⁵². The NLRP3 inflammasome also induces apoptosis via caspase-1 activation ³⁵³. The inflammasome produces active caspase 1, that when activated results in cleavage of pro-interleukin-1 β and pro-interleukin-18 and the production of active cytokines. In rats with

high-fat diet and streptozotocin-induced diabetic cardiomyopathy, silencing of the NLRP3 inflammasome decreases the levels of IL-1 β , and this observation is mirrored when silencing CMKLR1, a G-protein-coupled receptor for chemerin. Concurrent silencing of both NLRP3 and CMKLR1 potentiates the decrease in mature IL-1 β , as well as the levels of pyroptosis, underlining the important role the NLRP3 inflammasome and chemerin/CMKLR1 axis plays in mediating inflammation and pyroptosis in the setting of diabetic cardiomyopathy ³⁵⁴.

Myocardial inflammation not only results in the secretion of cytokines but also pro-fibrotic factors that activate fibroblasts and promotes cardiac fibrosis³⁴¹. TGF- β , a major cardiac pro-fibrotic cytokine, activates fibroblasts which results in the production of extracellular matrix proteins, increases collagen production, and decreases extracellular matrix degradation ³³⁴. Furthermore, IL-6 can suppress mir-29 and promote cardiac fibroblast proliferation and collagen production³⁵⁵. TNF- α also similarly promotes cardiac fibrosis through WISP1 activation ³⁵⁶.

Myocardial inflammation can also impair cardiac energy metabolism as IL-6 has been shown to impair myocardial glucose metabolism via SOCS3-dependent inhibition of IRS-1 357 . Furthermore, NF- κ B activation by TNF- α can inhibit PGC-1 α and consequently, increase glucose metabolism via downregulation of PDK4 $^{358, 359}$.

Inflammation can also result in endothelial and microvascular damage, resulting in myocardial ischemia and contributing to diastolic and systolic dysfunction in diabetic cardiomyopathy ^{291, 341, 360}. Furthermore, inflammation promotes ROS generation and downregulates SERCA2 (via IL-1β and IL-6), resulting in impaired Ca²⁺ handling and ultimately, diastolic dysfunction ³⁶¹. Myocardial inflammation can also depress cardiac contractility as TNF-α, IL-6, IL-1β and IL-2 exert negative inotropic effects on the heart ³⁶². Interestingly, a study in Zucker diabetic fatty rats treated with a β2-adrenergic receptor agonist decreased pro-inflammatory and pro-fibrotic responses in the heart and kidneys ³⁶³. β-arrestin can bind to the β2-adrenergic receptor and promote internalization, subsequently promoting desensitization. While this may imply a negative role in the setting of diabetic cardiomyopathy, β-arrestins have previously been reported to inhibit NF-κB activity via IκBα. As such, inhibition of NF-κB in the setting of diabetes via βarrestin and modulation of inflammatory mediators via sympathetic nervous system regulation may offer an alternative therapeutic strategy for diabetic cardiomyopathy. Further studies that investigate the interplay between β-arrestin and NF-κB in the setting of diabetic cardiomyopathy are warranted.

Hyperinsulinemia, via increased pancreatic production of insulin, follows insulin resistance in order to compensate for impaired cellular insulin actions. An excess of insulin can contribute to cardiomyocyte hypertrophy by acutely stimulating growth via the P13K/Akt-1 pathway.

Cardiac stiffness: Impairments in insulin signaling due to insulin resistance result in decreased GLUT4 translocation to the membrane and impaired PI3K/Akt signaling which

results in decreased Ca²⁺-ATPase activity, consequently increasing intracellular Ca²⁺ levels contributing to cardiac stiffness and diastolic dysfunction¹⁷⁶.

PI3K/Akt can activate endothelial NO synthase (eNOS), which results in an increase in NO that subsequently increases coronary vasodilation¹⁷⁶. However, insulin resistance decreases activation of eNOS and consequently decreases NO levels ⁸⁶. Decreased NO results in impaired coronary microcirculation, due to impairments in coronary vascular smooth muscle cell relaxation ^{364, 365}. Therefore, insulin resistance alongside hyperinsulinemia can contribute to cardiac stiffness and diastolic dysfunction.

Advanced glycation end products: Diabetes-induced chronic hyperglycemia significantly increases the formation of advanced glycation end products (AGE) in the heart³⁶⁶. Protein glycation occurs after prolonged exposure to high concentrations of glucose, where amino groups of proteins bond non-enzymatically to glucose ³⁶⁷. A correlation exists between formation of glycosylated tissue proteins in the heart and the period of hyperglycemia ³⁶⁸. Compared to non-diabetics pathologies, hearts from diabetic patients also show a higher abundance of AGE formation in the myocardium ³⁶⁹. A high abundance of AGEs also occurs in small intramyocardial arteries of the hearts of diabetic patients ³⁷⁰. This suggests that diabetes can exaggerate AGE formation and increase the susceptibility of myocardial vasculature to glycation ³⁶⁶. Formation of collagen cross-linking is a major determinant in the development of diabetic cardiomyopathy. Of relevance, an association between AGEs formation and decreased cardiac collagen solubility, and increased collagen III gene and protein expression, is seen in diabetes ³⁷¹.

³⁷². AGE-induced increases in cross-linked collagen may lead to myocardium and arterial wall stiffness and eventually atherosclerotic plaque formation ³⁷³. Moreover, AGEs are also linked with other pathological pathways in diabetic cardiomyopathy, including oxidative stress ^{374, 375} and impaired Na⁺/K⁺-ATPase activity ³⁷⁶. Chronic hyperglycemia increases both formation of AGE and expression of AGE receptors (RAGE), which in turn induces oxidative stress by activating transcription factor NFk- β ³⁷⁷. A strong association has been observed between increased RAGE elicited by diabetes and LV contractile dysfunction typical of diabetes cardiomyopathy, which is rescued by RAGE gene knockdown or blocking ^{378, 379}.

Hexosamine biosynthesis pathway O-GlcNAcylation (O-GlcNAc): O-GlcNAc is a posttranslational modification that is responsible for regulating the activity of proteins ³⁸⁰. This process is initiated when N-acetylglucosamine (GlcNAc) is attached to a serine or a

threonine residue of a peptide via an O-linkage (O-GlcNAc). The substrate for O-GlcNAc is uridine diphosphate-N-acetylglucosamine (UDP-GlcNAc), which is synthesized via the hexosamine biosynthetic pathway (HBP). It has been estimated that 5% of intracellular glucose contributes to the HBP³⁸¹, although this has been debated as to whether it is an accurate estimation for cardiomyocytes ^{382, 383}. Nevertheless, O-GlcNAc levels are closely related to glucose availability ³⁸⁰. Glucose, after entering the cell is converted to fructose-6-phosphate (F6P) by hexokinase and isomerase. F6P is further processed to uridine diphosphate-N-acetylglucosamine (UDP-GlcNAc) by 4 enzymatic reactions. UDP-GlcNAc is the substrate for O-GlcNAc transferase (OGT), which is responsible for

catalyzing O-GlcNAc to targeted proteins. Similar to other posttranslational modifications, O-GlcNAc is a highly dynamic and reversible process, with the removal of O-GlcNAc from targeted proteins being accomplished by O-GlcNAcases ³⁸⁴.

O-GlcNAc has been proposed to occur in the nucleus, cytoplasm, and mitochondria³⁸⁰, as opposed to other types of glycosylation which can take place in the extracellular matrix³⁸⁵. Chronic activation of the HBP is often associated with diabetic cardiomyopathy³⁸⁶, as evidenced by increases in both gene expression and protein levels of GFAT in the myocardium of diabetic patients ^{387, 388}. Genetic modulation of OGA to a truncated, less effective form can exacerbate O-GlcNAc, inducing a higher chance of developing diabetes ^{389, 390}. Vascular dysfunction is a common feature of diabetes, and such dysfunction can be attributed to excess O-GlcNAc of proteins, such as transcription factor Sp1 and eNOS ^{391, 392}. Both protein levels and activity of OGT are elevated in rat aortic smooth muscle cells subjected to hyperglycemia. Additionally, excessive O-GlcNAc can lead to improper Ca²⁺ handling ³⁹³. One specific target of O-GlcNAc is phospholamban, a protein regulating the function of SERCA2. Impairment of the function of phospholamban prevents the normal Ca²⁺ pumping after excitation from SERCA2, leading to improper contraction of heart muscle. The level of O-GlcNAc on cardiac proteins is carefully regulated by changes in OGT and OGA activity ^{394, 395}. O-GlcNAc may also affect complexes I, III, IV involved in mitochondrial respiration ^{394, 396}. O-GlcNAc may also impact ketone body metabolism by downregulating β -hydroxybutyrate dehydrogenase mRNA levels as well as succinyl-CoA:3-oxoacid CoA transferase protein

levels ²⁵³. Of interest, ketone oxidation is decreased in the myocardium of diabetic mice ²⁸¹.

An increasing body of evidence suggests an increase in O-GlcNAcylation levels in GlcNAcylation, and its signaling transduction negatively correlates with its O-GlcNAcylation level in adult rat cardiomyocytes ⁴⁰⁴. While circulating levels of N-terminal proteolytic fragment of histone deacetylase 4 (HDAC4) have been shown to be elevated in patients with diabetes, O-GlcNAcylation of HDAC4 is cardioprotective in a mouse model of diabetes ⁴⁰⁵. This cardioprotection is associated with a reduction in pathological Ca²⁺/calmodulin-dependent protein kinase II (CaMKII) signaling ⁴⁰⁵. O-GlcNAcylation also plays a role in regulating autophagy by modifying the synaptosomal associated protein 29 (SNAP29) ⁴⁰⁶. Increased O-GlcNAcylation of SNAP29 inhibits autophagic flux and causes further deterioration of cardiac diastolic dysfunction in STZ-induced diabetic rats ⁴⁰⁶. Furthermore, O-GlcNAcylation can modulate ionic homeostasis by targeting the activity of a number of ion channels. For example, acute hyperglycemia can enhance K⁺ channel recovery via CaMKIIō-S²⁸⁰ O-GlcNAcylation ⁴⁰⁷. Hyperglycemia also increases O-GlcNAcylation of Nav_{1.5}, which lead to the abnormal expression and distribution of Nav_{1.5}, loss of function of the sodium channel, and prolongation of the PR/QT interval in the hearts of diabetics ⁴⁰⁸. In patients with T2D, increased O-GlcNAcylation is linked to the dynamic of glucose-induced impairment of endothelial nitric oxide synthase activation in endothelial cells that could contribute to vascular dysfunction in T2D ⁴⁰⁹.

Cardiac cell death pathways: Three main pathways are involved in cell death, apoptosis, necrosis, and autophagy⁸⁷. A controlled rate of apoptosis and autophagy is necessary for removing unwanted cells ²⁹¹. However, in diabetes, cardiac cell death occurs at an accelerated rate⁴¹⁰⁻⁴¹². This is due to both a hyperactivated cellular death pathway and an impaired cellular defense mechanism ⁴¹³. Cardiac apoptosis is elevated in diabetes ^{291, 305, 414, 415}, which is important since an apoptotic rate as low as 0.023% is sufficient to induce lethal cardiomyopathy ⁴¹⁵. There are two main pathways of apoptosis: intrinsic or extrinsic. Intrinsic pathways can be initiated by various kinds of mitochondrial insult ^{87, 416, 417}. After formation of a mitochondrial permeability transition pore (mPTP), cytochrome C leaks into the cytoplasm and assemble with Apaf-1, ATP, and procaspase-9, forming apoptosome^{418, 419}. The final product activates the effector caspase: caspase-3. which will go on to cleave target proteins^{420, 421}. Additionally, p53 is able to sense damage of DNA strands and upregulate the transcription of two essential proteins: Bax and Fas⁴²². Bax is a pro-apoptotic protein that resides on the mitochondrial membrane⁴²³, whereas Fas contributes to the extrinsic cellular death pathway. Fas will act on death receptors located on cellular membrane⁴²⁴. Soluble extracellular protein, such as tumor necrosis factor- α (TNF- α) could also bind to death receptors⁴²⁵. Ligand binding initiates the assembly of multiprotein complex termed the death-inducing signaling complex (DISC), which recruits procaspase-8⁴²⁶. Such movement results in procaspase-8 activation ^{427, 428}. Caspase-8 cleaves Bid, leading to the formation of the active form, truncated-Bid (T-Bid)⁴²⁹, that is pro-apoptotic by assisting the leak of cytochrome C into the cytoplasm. There is still debate regarding the relative contribution of apoptosis versus

necrosis to cardiomyocyte cell death in dilated cardiomyopathy ⁴³⁰. It has been reported that necrotic cardiomyocytes are more dominant compared to apoptotic cells in dilated cardiomyopathy and severe aortic stenosis ^{431, 432}. In line with this, irreversible opening of the mitochondrial permeability transition pore (mPTP) also induces cell necrosis by ATP depletion, although it potentially triggers apoptosis via outer membrane rupture and cytochrome c release into the cytosol. Whether cardiomyocyte cell death by ischemia-induced mPTP opening has a significant contribution to cardiomyocyte loss and subsequent interstitial fibrosis in diabetic hearts warrants further investigation.

The cellular death pathways described above are altered in diabetic cardiomyopathy (Figure 3) . Direct exposure of high levels of glucose in myoblast H9c2 cells induces significant apoptotic cell death. The observations of hyperglycemia increases caspase 3 activation and cytochrome C release in cardiac cells are consistent with previous findings where high levels of glucose elevated the expression of Bax and its translocation from cytosol to mitochondria-enriched heavy membrane fraction in vascular endothelial cells⁴³³. On the other hand, the use of caspase-3 specific inhibitor, Ac-DEVD-cmk, can suppress hyperglycemia induced apoptosis^{433, 434}. Additionally, up-regulation of p53 in myocytes, due to hyperglycemia, occur at very early stages in the development of diabetic cardiomyopathy⁴³⁵, whereas attenuation of p53 transcriptional activity by IGF-1 prevents myocardial apoptosis in diabetic mice⁴³⁶. Besides high glucose, exposure to high levels of palmitate can also increase mitochondrial cytochrome C release, caspase-3 activation, followed by apoptotic cell death⁴³⁷. The formation of ROS and reactive nitrogen species (RNS) in the heart is another critical mediator of diabetes-induced myocardial cell death

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⁴³⁸. Both ROS and RNS may be involved in many aspects of the cell death pathway, such as activation of caspase 3, the PKC pathway, release of cytochrome C and death receptor activation ^{434, 439, 440}. The antioxidant, metallothionein (MT), can ameliorate this hyperglycemia-induced myocardial cell death ⁴⁰.

Unlike apoptosis, the role of autophagy in diabetic heart is still controversial. With some evidence suggesting that the induction of autophagy may convey protective effects ⁴⁴¹, other studies proposed that excessive autophagy may accelerate the process to heart failure ⁴⁴². Autophagy is believed to be impaired in diabetic heart. One major regulator of autophagy is insulin and impaired insulin signaling stimulates myocardial autophagy^{442, 443}. Given that many different animal models have not shown the blunted myocardial autophagy in diabetes ⁴⁴⁴⁻⁴⁴⁶, it is surprising that there has not been any approved treatment that targets autophagy specifically. However, some available medicines, such as metformin, rapamycin and resveratrol ⁴⁴⁷⁻⁴⁴⁹, have been found to promote autophagy indirectly, in addition to their main mechanism of action. Recently, Mst1 (macrophage stimulating 1) was found to be responsible for dictating the cardiomyocyte toward either apoptosis or autophagy in diabetes ⁴⁵⁰.

Study suggested that mitochondria-dependent, calcium overload-induced necrosis might contribute to the progression of heart failure ⁴⁵³. Although necrosis has been suggested to be a passive and unregulated form of cell death, targeting the pathways of necrosis has potential for treating cardio-cerebrovascular injury ⁴⁵⁴. Regulated necrosis can be

classified into many categories, including but not limited to pyroptosis and ferroptosis. Both forms have been proposed to correlate with diabetic cardiomyopathy development. Pyroptosis is characterized by formation of plasma membrane pores and extracellular release of inflammatory cytokines . High glucose promoted cardiomyocytes pyroptosis by increasing ROS production⁴⁵⁵. Elevated level of pyroptosis also induces cell death via the miR-214-3p/caspase-1/TGF-β1 pathway in diabetic mice ⁴⁵⁶. Among the many protective actions exerted by metformin, inhibition of pyroptosis by suppressing the mTOR pathway via AMPK activation, may decrease pyroptosis-induced cell death in diabetic cardiomyopathy⁴⁵⁷. Therefore, therapies targeting pyroptosis may be an effective approach. On the other hand, ferroptosis is a newly discovered form of cell death, which can be initiated by either iron overload or oxidative stress ⁴⁵⁸. Of interest, hydrogen sulfide is an endogenous gaseous signaling molecule that is capable of inhibiting ferroptosis. One recent study proposed that treatment with the ferroptosis inhibitor ferrostatin-1 can prevent hyperglycemia-induced ferroptosis ⁴⁵⁹.

*Alterations in cardiac Ca*²⁺ *handling:* One of the early perturbations in Ca²⁺ homeostasis in diabetic cardiomyopathy that precedes LV dysfunction is a slow decay of the Ca²⁺ transient ⁴⁶⁰⁻⁴⁶³. Possible mechanisms that contribute to the occurrence of these disarrangements in Ca²⁺ handling include perturbations in the activity of SERCA2a ^{465,} ⁴⁶⁶, as well as malfunctions in Ca²⁺ handling proteins due to posttranslational modifications, namely AGEs ⁴⁶⁷, O-GlcNacylation ⁴⁶⁸ and carbonylation ^{305, 469}. Impaired

Ca²⁺ handling between the sarcoplasmic reticulum and the mitochondria and alterations in Ca²⁺ influx and efflux to/from the cytosol and extracellular tissue and reduced activity of phospholamban and ryanodine receptors also contribute to the Ca²⁺ mishandling in diabetic cardiomyopathies ^{461, 470-472}. Ca²⁺ reuptake via SERCA2a is impaired in hearts from diabetic rats ⁴⁷³⁻⁴⁷⁵ and ob/ob mice ^{476, 477}. Interestingly, overexpression of SERCA2a improves Ca²⁺ handling in animal models of diabetic cardiomyopathy ⁴⁷⁸. A recent study also demonstrated that insulin resistance impairs SERCA2a activity and cardiac function via inhibiting protein kinase B/striated muscle preferentially expressed protein kinase (SPEG) signaling ⁴⁷⁹. It has also been shown that oxidative stress in the hearts of diabetics could contribution to the development of diabetic cardiomyopathy via impairing SERCA2a activity ⁴⁸⁰, and that enhancing SERCA2a activity is associated with improve cardiac function in the hearts of diabetics ⁴⁷⁹⁻⁴⁸¹.

Diabetes is also accompanied by alterations in contractile proteins that are associated with the changes in contractile function in the diabetic heart $^{136, 482, 483}$. The decrease in contractile function in the diabetic heart is also positively liked to the decrease in cardiac ATPase activity $^{472, 484, 485}$. Along with the disturbances in cardiac ATPase proteins, it has also been shown that there are disturbances in isomyosin distribution and shifts from V₁ to V₃ in the diabetic heart $^{486-488}$. There are also decreases in Ca²⁺ sensitivity along with troponin T-band shift in the diabetic heart $^{489, 490}$.

Neurohormonal mechanisms: The role of the renin-angiotensin-aldosterone system and endothelin-1 system in the pathophysiology of both heart failure and diabetes has
long been recognized ⁴⁹¹⁻⁴⁹³. Diabetes is accompanied by an upregulation of the reninangiotensin-aldosterone pathway that causes an increase in afterload, an important contributor to cardiac remodeling in diabetic cardiomyopathy . Consistent with this, a number of animal studies have shown that inhibiting activity of the renin-angiotensinaldosterone system limits the progression of diabetic cardiomyopathy ^{492, 494, 495}. As a result, angiotensin-converting enzyme (ACE) inhibitors or angiotensin receptor antagonists are recommended to treat heart failure in diabetic and non-diabetic patients ⁴⁹⁶. Moreover, diabetes is also associated with alterations in systemic autonomic function and perturbations in cardiac rhythm ^{497, 498}. Despite the detrimental effect of these irregularities in the neurohormonal system, there are no therapeutic approaches presently used to target this system in the setting of diabetic cardiomyopathy.

Changes in cardiac gene regulation: In diabetic cardiomyopathy there is differential expression of several genes involved in inflammation, fibrosis, insulin signaling, cell death and metabolism (Figure 4) ^{499, 500}. The advancements in microarray technology facilitates extensive gene expression profiling to uncover genetic mechanisms of diabetic cardiomyopathy and its therapeutic implication. Genes that are often dysregulated in diabetic cardiomyopathy are discussed in the respective section and are summarized in Figure 4.

Studies both in type 1 and type 2 diabetes have shown abnormal cytosolic Ca^{2+} homeostasis and decreased SERCA2a expression in cardiomyocytes along with diminished contractile function ⁵⁰¹. This is important in the development of diabetic

cardiomyopathy, as SERCA2a gene transfer or overexpression can reduce diabetesrelated contractile dysfunction, hypertrophy, and can differentially modulate the expression of genes involved in insulin signaling, glucose metabolism and cardiac remodeling ⁵⁰². Genes involved in inflammation and immune response are also affected by diabetic cardiomyopathy. For instance, IL6 and STAT3 genes are upregulated in patients with diabetic cardiomyopathy. On the other hand, downregulation of SOCS3 (Suppressor of cytokine signaling 3) is observed in diabetic cardiomyopathy patients compared to healthy controls ⁵⁰³.

Mitofusin 1 and 2 (Mfn1 and Mfn2) are mitochondrial dynamics proteins that controls fusion of the mitochondrial outer membrane ⁵⁰⁴. In *db/db* diabetic mice hearts, Mfn2 is down-regulated and contributes to an imbalance in mitochondrial dynamics. On the other hand, Mfn2 overexpression relieves diabetic cardiomyopathy by promoting mitochondrial fusion ⁵⁰⁵.

Activation of PPAR- α expression, a transcription regulator, also occurs in diabetic cardiomyopathy ^{47, 506}. Importantly, over expression of PPAR- α activates genes involved in cardiac fatty acid utilization, while suppressing genes in glucose metabolic pathways. This suggests that dysregulation of PPAR- α expression contributes to the metabolic derangements observed in diabetic cardiomyopathy ⁴⁷. Increased mitochondrial biogenesis has also been implicated in diabetic cardiomyopathy, and PPAR α -dependent activation of PGC-1 α may be a key driver of mitochondrial biogenic response in diabetic cardiomyopathy ⁵⁰⁶. An increase in PGC-1 α gene expression occurs in hearts of *db/db*

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mice ⁴¹. In addition, the up-regulation of PPAR α -dependent PGC-1 α is associated with increased expression of proteins of the mitochondrial electron transport chain and oxidative phosphorylation such as nuclear receptors families NRF-1 and NRF-2 and mtDNA transcription and replication (mtTFA) ⁵⁰⁶.

Recently, changes in the levels of non-coding RNAs have been recognized as important mediators of altered gene expression. The largest portion of the genome consists of non-coding RNAs. Although they are not directly transcribed to protein products, these RNAs regulate the transcription and post-transcriptional processing of many proteins. These regulatory RNAs consist of microRNAs (miRNA), long non-coding RNAs (IncRNA) and circular RNAs (circRNAs) ⁵⁰⁷. Over 4,500 IncRNA genes, and 2,000 microRNA genes has been identified in human genome alone ⁵⁰⁸. Although the function of the majority of non-coding RNAs are still unknown, mounting evidence suggests that these molecules play a significant role in a number of diseases processes and many of them are dysregulated in diabetic cardiomyopathy ^{499, 500, 509, 510}. Thus, their differential expression and role in diabetic cardiomyopathy pathogenesis is being actively investigated, partly because they may be potential biomarkers and therapeutic tools to treat diabetic cardiomyopathies.

LncRNAs are noncoding RNAs longer than 200 nucleotides in length. In addition to regulating other RNA functions, lncRNAs play an important role in epigenetic regulation by interacting with histone modifiers or chromatin remodelers or DNA ⁵¹¹. These lncRNAs also forms nucleic acid-protein complexes thereby regulating the activity or localization of these proteins or serves as a precursor for other miRNAs and circRNAs ⁵¹². Differential

expression of IncRNAs has been reported in diabetic cardiomyopathy and their abnormal expression has a role in promoting or inhibiting the development of diabetes. In diabetic cardiomyopathy there are significant alterations of a large number of IncRNAs that control apoptosis ⁵¹³⁻⁵¹⁵, fibrosis ⁵¹⁶ and inflammation ^{513, 517}. Detailed data on IncRNA changes and their function are summarized in Table 1.

Alterations in many microRNAs (miRNAs) are also linked to changes in gene expression patterns in diabetic cardiomyopathy. MiRNAs are short non-coding RNAs that regulate gene expression by binding to the 3' untranslated region of target messenger RNAs (mRNAs) ⁴²⁶. Upon binding, miRNAs repress gene expression by destabilizing or degrading the target mRNAs. To date, over 2650 mature miRNAs have been identified in humans that are implicated in various diseases ⁴²⁷. The role of various miRNAs in mediating diabetic cardiomyopathies have been studied broadly and is summarized in Table 1. Of importance, these studies have suggested the contribution of specific miRNAs to hypertrophic ⁵¹⁸⁻⁵²², fibrotic ⁵²³⁻⁵²⁵, apoptotic ⁵²⁶⁻⁵²⁹, inflammatory and oxidative stress ^{525, 530-532} changes in diabetic cardiomyopathy.

Circular RNAs (circRNAs) are produced during the processing of pre-mRNA ⁵⁰⁸. They are involved in the regulation of pre-mRNA splicing and RNA polymerase II ^{533, 534}. Analysis of the circRNA expression profiles in diabetic cardiomyopathy has shown differential regulation of several circRNAs in tissues from db/db mice hearts ⁵³⁵. Upregulation of these

circRNAs has also been shown in association with myocardial fibrosis ⁵³⁶⁻⁵³⁸ and pyroptosis ⁵³⁹.

Epigenetics mediated dysregulation of gene expression also contributes to the development of diabetic cardiomyopathies. Modification of histone proteins by lysine acetylation is a major epigenetic mechanism that regulates expression of many genes. For instance, increased acetylation of cardiac histone H3 leads to increased mRNA expression of multiple cardiomyopathy-related genes, together with cardiomyocyte hypertrophy, in diabetic mice ⁵⁴⁰. Increased acetylation of histone H3 and H4 in diabetes also leads to the recruitment of inflammatory genes promoters, including TNF- α and COX-2 ⁵⁴¹. Augmented histone acetylation at promoter regions of natriuretic peptide genes is also associated with increased expressions of ANP and BNP in the heart of diabetics ⁵⁴². Diabetes specific alterations in DNA methylation is also associated with altered in the phenotype of the heart in diabetes ⁵⁴³. This suggests that diabetes associated epigenetic modification may be an independent risk factor for diabetic cardiomyopathy.

The effect of antihyperglycemic drugs on diabetic cardiomyopathy severity

Although anti-hyperglycemic drugs significantly improve glycemic control in diabetic patients, use of these therapies does not necessarily equate to a reduced risk of developing heart failure ^{544, 545}. This highlights that lowering blood glucose alone is not sufficient to prevent diabetic cardiomyopathy development ⁵⁴⁵. However, a number of anti-

hyperglycemic drugs can alter the course of cardiovascular complications in the diabetic (Table 2). The impacts of these therapies on glucose and fatty acid oxidation are also summarized in Figure 5.

Metformin: Metformin is a first-line therapy in the majority of type 2 diabetic patients. In addition to its primary role in lowering blood glucose, beneficial effects of metformin have been shown on stimulating insulin action, decreasing inflammation ⁵⁴⁶, and improving myocardial energy metabolism ^{547, 548}. However, its effect on heart failure development remains uncertain. Some studies indicated that metformin is contraindicated in diabetic patients with heart failure due to lactic acidosis ⁵⁴⁹. A recent systematic review of 9 RCTs studies that examined metformin on heart failure related outcomes in patients with or without diabetes suggest some beneficial effects of metformin, but the overall evidence were not strong enough to make a solid conclusion about metformin decreasing heart failure severity ⁵⁵⁰. Other studies have suggested that metformin therapy does not decrease the risk of heart failure development ^{551, 552}.

Sulfonylureas: Sulfonylureas, especially older generation ones, increase the risk of adverse events in type 2 diabetic patients and are associated with a greater prevalence of hypoglycemia ^{553, 554}. A meta-analysis of 115 selected trials showed that sulfonylureas are associated with increased mortality, although major adverse cardiovascular events (MACE) did not appear to be affected ⁵⁵⁵. Another meta-analysis investigating the association of metformin and sulfonylureas on both all-cause and cardiovascular mortality in type 2 diabetic patients, showed that combination therapy resulted in an increase in

relative risk for cardiovascular hospitalization, as well as fatal and nonfatal events⁵⁵⁶. Similarly, a retrospective cohort analysis investigating the addition of insulin or a sulfonylurea in diabetic patients suggests that sulfonylureas increase the risk of non-fatal cardiovascular outcomes and all-cause mortality ⁵⁵⁷. This was also seen in a metformin and sulfonylurea combination therapy study of type 2 diabetic patients, in which patients newly treated with sulfonylureas possessed a higher risk for adverse cardiovascular events ⁵⁵⁸. Recently, a meta-regression analysis of 18 studies on the risk of cardiovascular events associated with sulfonylureas found that there was an increased risk of cardiovascular mortality and events with sulfonylurea treatment ⁵⁵⁹. In a network meta-analysis, 167,327 patients were studied to evaluate the risk of cardiovascular events with different sulfonylureas. Gliclazide and glimepiride were shown to have a lower risk of both cardiovascular-related mortality and all-cause mortality versus glibenclamide ⁵⁶⁰. Therefore, differences in the risk of mortality exist within the class of sulfonylureas. This is further reinforced by a cohort study of patients with type 2 diabetes on monotherapy with sulfonylureas, where glyburide and glimepiride did not increase the risk of adverse cardiovascular events versus glicazide, glipizide and tolbutamide⁵⁶¹.

Thiazolidinediones (TZDs): TZDs are known to cause fluid retention and as such, can increase the risk of congestive heart failure ⁵⁶². In the Pioglitazone Clinical Trial In Macrovascular Events (PROACTIVE) study, patients with type 2 diabetes and a history of macrovascular disease were randomized to receive pioglitazone or placebo⁵⁶³. Pioglitazone increased heart failure hospitalization, although this was associated with less cardiac ischemic events ⁵⁶³. The Diabetes Reduction Assessment With Ramipril and

Rosiglitazone Medication (DREAM) study, consisting of patients with impaired fasting glucose/glucose tolerance and no known cardiovascular disease, found that while rosiglitazone reduced diabetes and the development of renal disease, it increased new-onset heart failure ⁵⁶⁴. In the Rosiglitazone Evaluated for Cardiac Outcomes and Regulation of glycemia in Diabetes (RECORD) trial, a multi-centre open-label study with type 2 diabetic patients, rosiglitazone increased the risk of heart failure or hospitalization by over 2-fold ⁵⁶⁵. As such, rosiglitazone increases the risk of heart failure and alongside other TZDs, contain serious warnings regarding the increase in fluid retention and risk of congestive heart failure ⁵⁶².

Glitazars: A dual PPAR α and γ agonist designed to concurrently treat hyperlipidemia and hyperglycemia, glitazars combine the beneficial effects of agonizing both peroxisome proliferator-activated receptors. However, glitazars present a paradox and while addressing diabetic concerns with hyperlipidemia and hyperlycemia, they have been shown to worsen congestive heart failure in diabetic patients ^{566, 567}. Specifically, muraglitazar increases major adverse cardiovascular events, congestive heart failure and death in a review of several clinical trials that included 3725 patients ⁵⁶⁸. Another glitazar, aleglitazar, while presenting effective anti-diabetic effects, also increases the risk of heart failure ⁵⁶⁹. As such, concurrent agonism of PPAR α and γ results in cardiac dysfunction, which may be due to inhibition of PGC1 α and mitochondrial biogenesis ⁵⁷⁰.

GLP-1 receptor agonists: GLP-1 agonists improve glycemic control in diabetics by mimicking GLP-1 action ⁵⁷¹⁻⁵⁷³. These include Exenatide, a partial structural analogue of

GLP-1, , with other GLP-1 analogues including liraglutide, lixisenatide ⁵⁷⁴, and semaglutide ⁵⁷⁵. Some of these GLP-1 analogues have efficacy in mediating heart failure risk in diabetics, as shown by results of multiple phase III/IV large scale double-blind randomized clinical trials. The Exenatide Study of Cardiovascular Event Lowering (EXSCEL) Trial showed that exenatide in type 2 patients with cardiovascular risk did not increase their overall risk and that the incidence of major adverse cardiovascular events was not worsened ⁵⁷⁶. The Evaluation of Cardiovascular Outcomes in Patients with Type 2 Diabetes after Acute Coronary Syndrome during Treatment with Lixisenatide (EXLXA) Trial showed similar results, where lixisenatide treatment showed no effect on major adverse cardiovascular events (MACE) in type 2 diabetic patients who had a recent acute coronary event ⁵⁷⁷. In contrast, the Liraglutide Effect and Action in Diabetes (LEADER) trial showed a lower risk of MACE, including the rate of the first occurrence of death from cardiovascular causes, nonfatal myocardial infarction, or nonfatal stroke in type 2 diabetic patients with high cardiovascular risk ⁵⁷⁸. Additionally, the Trial to Evaluate Cardiovascular and Other Long-Term Outcomes with Semaglutide in Subjects with Type 2 Diabetes (SUSTAIN-6) showed a significantly lower rate of cardiovascular death, nonfatal myocardial infarction, or nonfatal stroke in type 2 diabetic patients with high cardiovascular risk ⁵⁷⁸. However, the results of the Functional Impact of GLP-1 for Heart Failure Treatment (FIGHT) Trial showed no improved post-hospitalization clinical stability with liraglutide in recently hospitalized patients with established heart failure and reduced ejection fraction ⁵⁷⁹. Examination of the effect of liraglutide on ventricular function in stable chronic heart failure patients with and without diabetes also showed that liraglutide did not improve LVEF or systolic function, and was associated with an increase in heart rate

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and more serious cardiac adverse events ⁵⁸⁰. Combined, this calls into question the benefits of liraglutide use in preventing diabetic cardiomyopathies.

DPP4 inhibitors: Incretins-based therapy has emerged as a novel treatment approach for diabetes management, with the inhibition of dipeptidyl peptidase 4 (DPP4) being used to prevent the cleavage and inactivation of GLP-1 ⁵⁸¹. DPP-4 inhibitors increase insulin secretion from pancreatic B-cells, thereby improving insulin tolerance and glucose control ⁵⁸²⁻⁵⁸⁴. Current DPP4 inhibitors include vildagliptin, sitagliptin, and saxagliptin and are similar in their efficacy in lowering HBA1C levels ⁵⁸⁵ and improving glucose tolerance in diabetes ⁵⁸⁶⁻⁵⁸⁹. Despite the efficacy of DPP4 inhibitors in improving glycemic control, the efficacy of DPP4 inhibitors in improving heart failure outcomes in diabetics remains unclear. The Saxagliptin Assessment of Vascular Outcomes Recorded in Patients with Diabetes Mellitus (SAVOR)-Thrombosis in Myocardial Infarction (TIMI) 53 trial (SAVOR-TIMI53) showed a significant increase in the rate of hospitalization for heart failure in type 2 diabetic patients treated with saxagliptin ⁵⁹⁰. However, the Examination of Cardiovascular Outcomes with Alogliptin versus Standard of Care (EXAMINE) trial showed non-inferiority of alogliptin to placebo on major cardiovascular events in diabetic patients with recent acute coronary syndrome ^{591, 592}. Moreover, the Trial Evaluating Cardiovascular Outcomes with Sitagliptin (TECOS) showed that sitagliptin neither improved or decreased rates of cardiovascular events such as death, myocardial infarction, stroke, or hospitalization for heart failure in type 2 diabetics with pre-existing cardiovascular disease ^{593, 594}. Other studies support the results of EXAMINE and TECOS trials ⁵⁹⁵⁻⁵⁹⁸. although results from meta-analyses demonstrate conflicting evidence for the

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effect of DPP4 inhibitors on mediating cardiovascular disease ⁵⁹⁹⁻⁶⁰¹. Animals studies have also shown conflicting results on the efficacy of DPP4 inhibitors on cardiovascular disease. Sitagliptin treatment decreased LV passive stiffness and improved global LV performance in an obese type 2 diabetic mice ⁶⁰². However, long term treatment of vildagliptin showed no cardioprotective effects on cardiac function, remodeling, or infarct size in Sprague-Dawley rats subjected to myocardial infarction induced by coronary ligation ⁶⁰³. Combined, these studies suggest minimal beneficial effects of DPP4 inhibitors in reducing the risk of heart failure in diabetics, and support that certain DPP4 inhibitors may be safe in patients. However, the cardiovascular safety and efficacy of DPP4 inhibitors needs to be further elucidated.

SGLT2 inhibitors: Sodium glucose co-transporter 2 inhibitors (SGLT2i) prevent glucose reabsorption in the proximal tubules of the kidney, therefore increasing its secretion into the urine and improving glycemic control ⁶⁰⁴⁻⁶⁰⁶. Three SGLT2i approved for clinical use include empagliflozin, dapagliflozin, and canagliflozin. Recently, large-scale clinical trials have shown cardioprotective benefits independent of its antihyperglycemic effect in both type 2 diabetic and non-diabetic patients ⁶⁰⁷⁻⁶¹². The results of the Empaglifozin Cardiovascular Outcome Event Trial in Type 2 Diabetes Mellitus Patients (EMPA-REG OUTCOMES) showed a lower occurrence of death from cardiovascular causes, nonfatal myocardial infarction, or nonfatal stroke, and a reduction in overall mortality and heart failure hospitalization in empagliflozin treated type 2 diabetics patients with cardiovascular risk compared to placebo ⁶⁰⁷. The Canagliflozin Cardiovascular Assessment Study (CANVAS) and Dapagliflozin effect on Cardiovascular Events-Thrombolysis in

Myocardial Infarction 58 (DECLARE-TIMI58) trials supported the results of the EMPA-REG OUTCOMES study. The CANVAS trial showed a lower risk of cardiovascular events in type 2 diabetic patients with an elevated risk of cardiovascular disease ⁶¹², while the DECLARE-TIMI58 trial showed a reduction in cardiovascular death and heart failure hospitalization in type 2 diabetic patients with or at high risk of cardiovascular disease, although it did not reduce the rate of MACE ⁶¹¹. Interestingly, the Dapagliflozin and Prevention of Adverse-Outcome in Heart Failure (DAPA-HF) trial showed a reduction of in the risk of mortality and heart failure reduction in patients with heart failure and reduced ejection fraction with or without type 2 diabetes ⁶⁰⁹. These results are supported by the Empagliflozin Outcome Trial in Patients with Chronic Heart Failure and a Reduced Ejection Fraction (EMPEROR-Reduced) trial ⁶¹⁰. Combined, evidence from clinical trials show a safety and efficacy of SGLT2i as a therapeutic strategy to manage diabetes and associated cardiovascular disease, heart failure, and their risk.

Studies in animal models have also demonstrated cardiovascular benefits supporting the results from the major clinical trials. Empagliflozin improves cardiac contractility, by fractional area change, and improves microvascular function in *ob/ob^{-/-}* mice ⁶¹³. Additionally, empagliflozin treatment attenuates cardiac fibrosis and improves hemodynamics in hypertensive rat heart failure ⁶¹⁴. However, despite these beneficial cardiac outcomes, the exact mechanism of the cardioprotective effects of SGLT2i are unknown ⁶¹⁵. Multiple mechanisms have been proposed including diuresis/natriuresis, improved cardiac energy metabolism ²⁸¹, reduction of inflammation ⁶¹⁶, and prevention of ischemia reperfusion injury ⁶¹⁷ to name a few key mechanisms. Further studies are

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needed to fully elucidate a mechanism to explain the observed cardioprotective effects of SGLT2i in the diabetic and nondiabetic failing heart.

Insulin: While insulin is the first-line therapy to treat T1D, it is only used to manage T2D patients when oral hypoglycemic drugs and lifestyle do not establish glycemic control. It has been suggested that heart failure prevalence and cardiovascular mortality is increased in patient with T2D who receive insulin ⁶¹⁸. Evaluation of the impact of insulin therapy on cardiovascular disease in diabetic patients has been the focus of a number of recent clinical trials. For example, the ORIGIN trial (Outcome Reduction With Initial Glargine Intervention) investigated glargine's impact compared to standard care in T2D patients with high cardiovascular risk. The trial data were neutral, and the rates of incident cardiovascular outcomes were similar in the insulin-glargine and standard-care ⁶¹⁹. In addition, the DEVOTE trial (A Trial Comparing Cardiovascular Safety of Insulin Degludec Versus Insulin Glargine in Subjects With Type 2 Diabetes at High Risk of Cardiovascular Events) compared the cardiovascular safety of degludec, ultralong acting insulin, to insulin glargine in patients with T2D and high cardiovascular risk. The study showed that degludec was noninferior to glargine concerning the incidence of major cardiovascular events ⁶²⁰. While enhancing circulating insulin levels can restore cardiac insulin sensitivity in the failing heart, enhancing cardiac efficiency and reducing cardiovascular mortality, prospective studies that aim to access this possibility directly are currently lacking.

The effect of heart failure drugs on glycemic control

Renin-Aldosterone-Angiotensin inhibitors: ACE inhibitors have been shown to improve insulin resistance and glucose intolerance via increases in GLUT4 translocation (Table 3) ⁶²¹. Diabetic mice treated with the ACE inhibitor temocapril show decreases in plasma glucose and insulin levels, increases in skeletal muscle glucose uptake and increases in translocation of GLUT4 to the plasma membrane ⁶²¹. In a single blind, cross-over design study in type 2 diabetic patients that had arterial hypertension, the ACE inhibitor captopril increased insulin sensitivity and improved glycemic control ⁶²². Captopril treatment of diabetic patients also improves glucose control ⁶²³. However, while ACE inhibitor therapy improves glycemic control effects, a case-control study in diabetic patients found that ACE inhibitors are associated with an increase in hospitalization for severe hypoglycemia ⁶²⁴.

Lipid-lowering agents: Statins have a propensity to induce hyperglycemia and have been shown to cause glucose intolerance in both animals and humans. For instance, diabetic rats treated with atorvastatin or simvastatin exhibit hyperglycemia and glucose intolerance ⁶²⁵. In a meta-analysis of 9 trials of patients treated with statins, mean HbA₁c was higher by 0.12%, indicative of a modestly increased risk for diabetes with statin treatment ⁶²⁶. In another meta-analysis that investigated the effect of statin therapy on HbA₁c levels as well as fasting plasma glucose, statins increased HbA₁c ⁶²⁷. Specifically, pitavastatin improved glycemic control while atorvastatin worsened glycemic control ⁶²⁷. This diabetogenic effect was also recapitulated in a national health screening cohort of non-diabetic individuals taking statins, showing that greater adherence to the use of

statins, specifically atorvastatin, rosuvastatin, pitavastatin and simvastatin, results in increases in fasting glucose levels⁶²⁸.

B-blockers: β -adrenergic stimulation promotes insulin and glucagon release while α adrenergic stimulation inhibits insulin and glucagon secretion. Therefore β-adrenergic receptor antagonism inhibits insulin release and may worsen glycemic control especially during hypoglycemia. The selectivity of the ß-blocker yields distinct metabolic effects and certain ß-blockers can exacerbate hypoglycemic episodes by delaying glucose recover time ^{629, 630}. A retrospective study that monitored glucose in patients receiving either carvedilol or a selective second-generation ß-blocker (metoprolol or atenolol) found that ß-blockers, specifically metoprolol or atenolol, increase the odds of hypoglycemia in these hospitalized patients ⁶³¹. In hypertensive diabetic patients, treatment with propranolol or metoprolol results in mean blood sugar increases of 1.0-1.5 mM ^{629, 632}. A randomized, double-blind parallel-group trial in patients with diabetes and hypertension showed that metoprolol increases mean HbA_{1C}, but insulin sensitivity is improved with carvedilol treatment⁶³³. Third generation non-selective β-blockers (carvedilol) possess insulinsensitizing properties and improve glycemic control, while second generation β₁-selective (metoprolol) antagonism worsens glycemic control ⁶³⁴. To underline the distinct benefits between β -blockers, non-vasodilating β -blockers (metoprolol, propranolol and atenolol) have been shown to worsen glycemic control while vasodilating β -blockers (carvedilol, labetalol, nebivolol) improve glucose profiles.

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Aldosterone Antagonists: Enhanced activity of the aldosterone signaling has been implicated in the development of diabetes-induced heart failure via triggering fibrosis and insulin resistance. Treatment of dilated cardiomyopathy patients with the aldosterone antagonist spironolactone resulted in a reduced collagen accumulation in the heart and improved LV function ⁶³⁵. Likewise, antagonizing aldosterone can improve diastolic function and limit fibrosis in patients with hypertensive cardiomyopathy ⁶³⁶ and metabolic syndrome ⁶³⁷. Of interest is that eplerenone is shown to limit biomarkers of inflammation and insulin resistance in patients with HIV ⁶³⁸. Aldosterone antagonists have also shown promising effects by reducing apoptosis and improving diastolic function in murine models of diabetic cardiomyopathy ⁶³⁹⁻⁶⁴¹. The impact of aldosterone antagonism on diastolic function, cardiac insulin resistance and inflammation in patient with diabetes-induced heart failure is yet to be determine.

Concluding remarks

The pathophysiology of diabetes can affect the heart through multiple mechanisms that cause structural, metabolic, and functional remodeling, leading to a well-acknowledged condition called diabetic cardiomyopathy. Diabetes-induced perturbations in insulin resistance, fuel preference, reactive oxygen species generation, inflammation, cell death pathways, neurohormonal mechanisms, advanced glycated end-products accumulation, lipotoxicity, glucotoxicity and posttranslational modifications contribute to the development of diabetic cardiomyopathies. Targeting these pathways is a potential therapeutic approach to lessening the likelihood of developing diabetic cardiomyopathies.

A number of antidiabetic therapies can also prevent diabetic cardiomyopathy and reverse cardiac dysfunction. These advancements will help achieve personalized treatment for diabetic patients by achieving glycemic control and managing comorbidities and limiting cardiovascular disease. Better clarity of the mechanisms involved in diabetic cardiomyopathy should lead to better therapeutics approaches to treat patients with diabetes and heart failure.

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Figure legends

Figure 1: Energy metabolic changes in a healthy setting versus diabetic cardiomyopathy. In the healthy heart, approximately 60% of the heart's energy comes from the oxidation of fatty acids, followed by approximately 5% by glycolysis and 25% from glucose oxidation, and 10% by ketone oxidation. However, in diabetic cardiomyopathy, due to systemic and local changes in energy substrate concentrations as well as insulin resistance, the metabolic protein machinery is perturbed and subsequently, the heart's overall energy metabolic profile is impaired. As such, diabetic cardiomyopathy results in an increase in fatty acid oxidation, decreased glucose metabolism and decreased ketone oxidation.

Figure 2: The different stages of diabetic cardiomyopathy. HFpEF, heart failure with preserved ejection fraction; HFrEF heart failure with reduced ejection fraction; HF, heart failure; AGEs, advanced glycation end products.

Figure 3: Mechanisms that contribute to diabetes-induced heart failure. While the exact pathophysiology of the diabetic-induced heart failure still not fully defined, there are a number of mechanisms that play important roles in its occurrence. This includes mitochondrial dysfunction, cardiac insulin resistance and impaired cardiac insulin signaling pathway, perturbed fuel use, low ATP levels, inflammation, advanced glycation end products, O-GlcNAcylation, cell death, neurohormonal mechanism, contractile
proteins dysfunction, oxidative stress, gene reprogramming, lipotoxicity, glucose toxicity and perturbed Ca²⁺ handling.

Figure 4: Gene expression dysregulation in diabetic cardiomyopathy. CD36: cluster of differentiation 36; PPARα: Peroxisome proliferator-activated receptor; FABP: fatty-acid-binding proteins; GLUT1: glucose transpoter 1; GLUT4: glucose transporter 4; PDK: pyruvate dehydrogenase kinase; MG53: mitsugumin 53; ANP: atrial natriuretic peptide; NF-kB: nuclear factor kappa B; TGF-β1: transforming growth factor beta 1; MMP2: matrix metalloproteinase-2; Mst1: macrophage Stimulating 1; SERCA2a: sarcoplasmic-endoplasmic reticulum Ca2+ ATPase 2a; MCP-1: monocyte chemoattractant protein-1; VCAM-1: vascular cell adhesion molecule 1; TNFα: tumor necrosis factor; SOCS3: suppressor of cytokine signaling-3; COL1A1: collagen type 1 alpha1.

Figure 5. Summary figure of various antihyperglycemic drugs and their mode of action in the context of the heart. Fatty acid, glucose and ketone body metabolism are represented in this figure with the key modes of homeostasis regulation presented. SGLT2 inhibitors inhibit SGLT2 in the proximal tubules and thus, prevent renal glucose reabsorption, promote glycosuria, decreased insulin release, increased hepatic ketogenesis and increased circulating blood ketone levels. These increased circulating ketones can subsequently modulate cardiac ketone oxidation rates. Metformin's mode of action is not well understood, although it does stimulate AMPK which inhibits ACC, decreases malonyl-CoA levels and increases fatty acid metabolism. DPP4 inhibitors prevent DPP4 from inactivating GLP-1, and thus increase GLP-1 levels to potentiate

insulin secretion from pancreatic beta cells. GLP-1 receptor agonists similarly increase GLP-1 levels to increase insulin secretion. Sulfonylureas bind to SUR1, and consequently the K-ATP channel closes, depolarizing the pancreatic islet cell and increasing intracellular calcium levels to promote secretion of insulin. Lastly, TZDs have widespread actions in the body but here we focus on its role in promoting glucose metabolism and improving insulin sensitivity via binding to PPARγ and promoting the transcription of genes involved in glucose uptake and metabolism.

Table 1: Alterations in miRNA, LncRNA, CircRNA and their role in diabetic cardiomyopathy

Non-coding	Subclasses	Status	Study setting	Altered proteins	Pathological	Reference
RNAs				/RNAs	Pathways involved	
HOTAIR	IncRNA	Downregulated	STZ induced diabetic mice,	SIRT1	Apoptosis &	513
			H9c2 cardiomyocytes		inflammation	
MALAT1)	IncRNA	Upregulated	STZ- induced diabetic rats	-	Apoptosis	642
MIAT	IncRNA	Upregulated	STZ induced diabetic rats, neonatal	miR-22-3p	Apoptosis	514
			cardiomyocytes			
Crnde	IncRNA	Downregulated	STZ induced diabetic mice, mouse	Smad3	Fibrosis	643
			neonatal cardiac fibroblasts			
LncRNA H19	IncRNA	Downregulated	STZ-induced diabetic rats,	VDAC1, miR-675,	Autophagy&	515, 517
			neonatal cardiomyocytes	EZH2 & DIRAS3	apoptosis	
Zfas1	IncRNA	Upregulated	STZ-induced diabetic mice, H9c2	miR-9 , FN mRNA,	Fibrosis	516
			cardiomyocytes	Col1&4		
miR-1	miRNA	Upregulated	STZ induced diabetic rats	Junctin	Oxidative stress	530
miR-30c, miR-	miRNA	Down regulated	Diabetic patients, diabetic rats,	p53, p21, ANP	Hypertrophy &	518
181a			and H9c2 cardiomyocytes		apoptosis	

miRNA133a	miRNA	Downregulated	STZ-induced diabetic mice, neonatal rat cardiomyocytes	ANP, BNP, MEF2A and MEF2C	Hypertrophy	519
miR-133a	miRNA	Downregulated	STZ-induced diabetic mice	TGF-β1, fibronectin (FN1) and COL4A1	Fibrosis	523
miR-150	miRNA	Downregulated	STZ induced diabetic rats, neonatal rat cardiomyocytes	р300	Hypertrophy	520
miR-373	miRNA	Down regulated	Diabetic mice , neonatal rat cardiomyocytes	MAPK signaling (ERK1/2, JNK, and p38)	Hypertrophy	521
miRNA-1, miRNA-208a	miRNA	upregulated	STZ-induced diabetic mice, diabetic patients	pro-survival Pim-1 & Caspase-3	Apoptosis	526
miR-451	miRNA	Upregulated	type 2 diabetic mice, neonatal rat cardiac myocytes	Calcium-binding protein 39 (Cab39)	Hypertrophy and contractile	522
miR-195, miR- 34a	miRNA	Upregulated	STZ induced diabetic mice	BCL-2? H9c2 cells	Apoptosis	527, 644
miR-144	miRNA	Upregulated	STZ-induced diabetic mice	Nuclear factor erythroid-2-related factor 2 (Nrf2)	Oxidative stress	531
miR-483-3p	miRNA	Upregulated	STZ-induced diabetic mice, H9c2 cardiomyocyte	Insulin growth factor 1 (IGF1)	Apoptosis	528

miR30c	miRNA	Down regulated	Type 2 db/db mice, H9c2 cardiomyocytes	BECN1	Autophagy	645
miR-30d	miRNA	Up regulated	STZ-induced diabetic rats, neonatal rat cardiomyocytes	Caspase-1, IL-1β and IL-18 & Foxo3a	Pyroptosis	646
miR-15a/b	miRNA	down-regulated	Type 2 diabetic patients, type 2 db/db mice HL-1 cardiomyocytes	(TGFβR1) and connective tissue growth factor (CTGF)	Fibrosis	524
miR-29	miRNA	dysregulated	Zucker diabetic fatty (ZDF) rats, HL-1 cardiomyocytes	Myeloid Cell Leukemia 1(MCL-1)	Apoptosis	529
miR-200b	miRNA		STZ-induced diabetic mice			
miR-200	miRNA	Upregulated	db/db mice	cyclooxygenase-2	Inflammation	532
miR-141	miRNA	Upregulated	STZ-induced diabetes,HL-1 cells	mitochondrial phosphate carrier (Slc25a3)	Mitochondrial ATP production.	647
miR-146a	miRNA	downregulated	STZ-induced diabetes	L6, TNFα, IL-1β, MCP-1, NF-κB, Col1α1, Col4α1	Fibrosis & inflammation	525

miR-301	miRNA	Upregulated	Type 2 db/db mice ,H9C2 cells	Voltage-gated potassium channel (Kv4.2)	Action potential	648
miR-193-5p	miRNA	Upregulated	Myocardial microvascular endothelial cells of type 2 diabetic Goto-Kakizaki (GK) rats	IGF2	Angiogenesis	649
circRNA_010567	crcRNA	Upregulated	db/db mice, cardiacfibroblasts	TGF-β1	Fibrosis	537
circHIPK3	circRNA	Upregulated	STZ induced diabetic mice	<i>Col1</i> α1 and <i>Col3</i> α1	Fibrosis	538
circ_0076631	circRNA	Upregulated	Type 2 diabetic patients, AC16 cardiomyocyte	Caspase 1	Pyroptosis	539

IncRNA: long non-coding RNA; miRNA: micro-RNA; circRNA: circular-RNA; STZ: Streptozotocin; VDAC1; voltage-dependent anion channel 1; EZH2: enhancer of zeste homolog 2; FN: fibronectin; Col 1&4: collagen type 1& 4; ANP: atrial natriuretic peptide; B-type natriuretic peptide; MEF2A: myocyte enhancer factor 2A; TGF-β1: transforming growth factor-beta 1; MAPK: mitogen-activated protein kinase; ERK1: extracellular signal- regulated protein kinase 1; JNK: jun N-terminal kinase; BECN1: IL-1β: interleukin-1β; IL-1β: interleukin-18; GFβR1: transforming growth factor beta receptor I; CTGF: connective tissue growth factor; IL-6: interleukin-6; TNFα: tumor necrosis factor-alpha; MCP-1: monocyte chemoattractant protein-1; Col1α1:collagen type I alpha 1; Col4α1: collagen type IV alpha 1; IGF2: insulin-like growth factor 2; Col3α1: collagen type III alpha 1

Table 2: The effects of different classes of anti-diabetes drugs on the

cardiovascular system and the development of diabetic cardiomyopathy

Class of the therapy	Biological	Study	Main findings	References
	target of each drug class	characteristics		
		Systematic review randomized clinical trials (RCTs)	 Improved cardiovascular outcomes in early stage of diabetes No beneficial effects after hospitalization or in those with overt CVD 	550
		Post hoc analysis of SAVOR-TIMI 53 Trial (Saxagliptin and Cardiovascular Outcomes in Patients With Type 2 Diabetes Mellitus	No effect on CVD end points in patients with prior heart failure and kidney disease	650
Metformin	Unknown	Prospective cohort study	Increased risk of CVD and death in patients with T2DM at the time of first AMI May be beneficial in Post-AMI	651
		Experimental DCM (both <i>in vivo</i> & <i>in</i> <i>vitro</i>)	improved cardiac function and alleviation of apoptosis	652
		Systematic review and meta- regression analysis	Reduced mortality in patients with HFpEF	653
		Observational study	Has no effect on the risk of HF-related exacerbation	551
Sulfonylureas				
		Retrospective analysis of 20,450 type 2 diabetic patients	Increased risk of adverse cardiovascular events with sulfonylureas	553
Sulfonylureas	Stimulate insulin release from the pancreas	Meta-analysis of 115 randomized clinical trials	Increased mortality and risk of stroke with sulfonylureas in type 2 diabetics	555
		Meta-analysis of 9 observational studies	Combination therapy with sulfonylureas and metformin increased the relative risk of	556

			cardiovascular mortality and hospitalization	
		Retrospective cohort analysis of 1787,341 diabetic patients	Sulfonylureas increased the risk of non-fatal cardiovascular outcomes and all-cause mortality	557
		Retrospective cohort analysis of 5,730 patients with type 2 diabetes	Increased risk of adverse cardiovascular outcomes with sulfonylurea treatment or combination therapy of sulfonylurea + metformin	558
		Meta-regression analysis of 18 studies	Increased risk of cardiovascular mortality and events with sulfonylurea treatment	559
Glicazide and glimepiride vs. glibenclamide		Network meta- analysis of 167, 327 patients	Gliclazide and glimepiride lowered the risk of cardiovascular- related mortality and all- cause mortality than glibenclamide.	560
Non-specific, long- acting sulfonylureas (glyburide, glimepiride)		Retrospective cohort analysis of 17,604 sulfonylurea initiators	Does not increase myocardial infarction, ischemic stroke, cardiovascular deaths or all-cause mortality	561
Short-acting sulfonylureas (glicazide, glipizide, tolbutamide)			Increases myocardial infarction, ischemic stroke, cardiovascular deaths or all-cause mortality	
Thiazolidinediones				
Pioglitazone	Bind avidly to peroxisome	5238 type 2 diabetic patients treated with either pioglitazone or placebo for 4 years	Pioglitazone increases the percent of heart failure hospitalizations but associated with decreased cardiac ischemic events	563
Rosialitazone	proliferator- activated receptor gamma in adipocytes to promote	5,269 subjects with impaired glycemic control treated with ramipril, rosiglitazone or placebo	Rosiglitazone increased new-onset heart failure despite decrease diabetes	564
	adipogenesis and fatty acid uptake	4447 type 2 diabetic patients on metformin or sulfonylurea monotherapy received either	Rosiglitazone increased the risk of heart failure or hospitalization	565

		rosiglitazone or a combination of metformin and sulfonylurea		
Dipeptidyl-peptidase 4 (DPP4) inhibitors				
Saxagliptin		Type 2 diabetic patients who had a history of, or were at risk for cardiovascular events received saxagliptin or placebo and followed individuals for 2 years	Saxagliptin increased the rate of hospitalization for heart failure in type 2 diabetic patients, with no effect on any other outcomes including rate of ischemic events, myocardial infarction, or death	590
Alogliptin	Block the action of Dipeptidyl- peptidase 4 (DPP4) on incretins	Type 2 diabetic patients with a recent acute coronary syndrome (either an acute myocardial infarction or unstable angina requiring hospitalization within the previous 15 to 90 days), received either alogliptin or placebo and were followed individuals for 40 months	Alogliptin was non- inferior to placebo on major adverse cardiovascular events (MACE - rates of death from cardiovascular causes, nonfatal myocardial infarction, and nonfatal stroke)	591
Sitagliptin		Type 2 diabetic patients with pre- existing cardiovascular disease received either Sitagliptin or placebo and were followed for 3 years	Sitagliptin neither improved or decreased rates of cardiovascular events such as death, myocardial infarction, stroke, or hospitalization for heart failure	593
Sitagliptin		Type 2 diabetes received either sitagliptin or conventional therapy and were followed for 2 years	Sitagliptin did not have any additional effects on the progression of carotid IMT in participants with type 2 diabetes; no significant	595

		differences between	
		groups	
DDP-4 Inhibitors	Identified patients from Taiwan's National Health Insurance Research who were diagnosed with type 2 diabetes mellitus (T2DM) and had previous history of HF between 2009 and 2013 who did and did not use DPP-4 inhibitors.	The risks of all-cause mortality, and mortality from the combination of MI and stroke, and ischemic stroke were lower for patients receiving DPP-4 inhibitors than for those who did not receive this treatment. The risk of hospitalization for heart failure had no significant differences between groups.	596
DPP-4 inhibitors	Analysis of heart failure patients with diabetes mellitus who were hospitalized due to their heart failure and discharged from Fukushima Medical University between 2009 and 2013, and either did or did not take DPP4 inhibitors	DPP-4 inhibitor use was associated with low cardiac and all-cause mortality in heart failure patients with diabetes mellitus hospitalized for heart failure	597
DPP-4 inhibitors	A meta-analysis of randomized clinical trials to determine the effect of DPP4 inhibitors on the incidence of MACE, cancer, and pancreatitis	DPP-4 inhibitor use may be associated with a possible protection from cardiovascular events	599
DPP-4 inhibitors	A systematic review of articles with the search terms <u>DPP-</u> <u>IV</u> <u>inhibitors</u> and heart failure in full papers and abstracts published since October 2013	DPP-4 inhibitor use may increase the risk of hospitalization for heart failure, should be used with caution especially in patients with a history of CVD and heart failure	600
DPP-4 inhibitors	A meta-analysis of clinical trials assessing the effects of DPP-4 inhibitors in	DPP-4 inhibitors have a no significant effects on both all-cause and cardiovascular mortality; there is a statistically	601

		participants with type 2 diabetes mellitus	significant increase in heart failure hospitalizations with DPP-4 inhibitors in comparison with placebo or other hypoglycemic agents	
Sitagliptin		Male CD-1 mice received sitagliptin prior to 30 min myocardial ischemia and 4 h reperfusion	Sitagliptin limited myocardial infarct size post myocardial ischemia and reperfusion	654
Sitagliptin		db/db mice received either sitagliptin or vehicle for 8 weeks	Sitagliptin decreased LV passive stiffness and improved global LV performance	602
Vildagliptin		Sprague-Dawley rats pretreated with either vildagliptin or vehicle then subjected to coronary ligation to induce MI	Vildagliptin showed no cardioprotective effects on cardiac function, remodeling, or infarct size	603
Glucagon-like peptide 1 (GLP-1) receptor agonists				
Exenatide	Mimic the action of incretin on insulin release from the pancreas	Patients with type 2 diabetes, with or without previous cardiovascular disease received either exenatide or placebo and were followed for 3 years	Exenatide showed no significant differences in the incidence of major adverse cardiovascular events	576
Lixisenatide		Type 2 diabetes who had had a myocardial infarction or who had been hospitalized for unstable angina within the previous 180 days received either lixisenatide or placebo and were followed for 2 years	Lixisenatide treatment showed no significant effect on major adverse cardiovascular events (MACE) between type 2 diabetic patients who had a recent acute coronary event	577

Liraglutide	Patients with type 2 diabetes and high cardiovascular risk; individuals received either liraglutide or placebo and were followed up for 3 years	Liraglutide treatment demonstrated a lower risk of MACE, including the rate of the first occurrence of death from cardiovascular causes, nonfatal myocardial infarction, or nonfatal stroke	578
Semaglutide	Type 2 diabetic patients received either semaglutide or placebo for 104 weeks	Semaglutide lowered the rate of cardiovascular death, nonfatal myocardial infarction, or nonfatal stroke	575
Liraglutide	Patients with established heart failure and reduced LVEF who were recently hospitalized received either liraglutide or placebo and were followed for 180 days	Liraglutide showed no improved post- hospitalization clinical stability with Liraglutide in recently hospitalized patients with established heart failure and reduced ejection fraction	579
Liraglutide	Diabetic and nondiabetic patients with reduced left ventricular ejection fraction received either liraglutide or placebo for 24 weeks	Liraglutide did not improve LVEF or systolic function, and was associated with an increase in heart rate and more serious cardiac adverse events	580
	Male Sprague- Dawley rats received either exenatide analog AC3174 or vehicle two weeks post-MI for 11 weeks	Exenatide analog, AC3174, improved cardiac function, morphology and survival	655
Exendin-4	Male Sprague- Dawley rats heart received either exendin-4 or GLP- 1(9-36) amide, the primary endogenous metabolite of GLP- 1, following 45 minutes ischemia	Exendin treatment administered at reperfusion reduced infarct size and improved mechanical performance in isolated Sprague Dawley rat hearts during ischemia-reperfusion	656

		prior to 120 minutes reperfusion		
Liraglutide		Male C57BL/6 mice received liraglutide for 1 week prior to a permanent surgical ligation of the left anterior descending artery	Liraglutide treatment decreased infarct size, and improved cardiac output 4 weeks post ischemia in mice	657
sodium-glucose transport protein 2 (SGLT2) inhibitors				
Empagliflozin		Patients with type 2 diabetes at high risk for cardiovascular events; individuals received either empagliflozin or placebo once daily and were followed up for around 3 years	Empagliflozin treatment showed a lower occurrence of death from cardiovascular causes, nonfatal myocardial infarction, or nonfatal stroke, and a reduction in overall mortality and heart failure hospitalization in type 2 diabetics patients with cardiovascular risk compared to placebo	607
Canagliflozin	Inhibit sodium- glucose transport protein 2 in the kidney and prevent the kidneys from reabsorbing glucose back	Patients with type 2 diabetes and an elevated risk of cardiovascular disease; individuals received either canagliflozin or vehicle and were followed for 188 weeks	Canagliflozin showed a lower risk of cardiovascular events	612
Dapagliflozin	into the blood	Patients with type 2 diabetes who had or were at risk for atherosclerotic cardiovascular disease; individuals received ither dapagliflozin or placebo and were followed for 4 years	Dapagliflozin showed a reduction in cardiovascular death and heart failure hospitalization	611
Dapagliflozin		Patients with heart failure and reduced ejection fraction	Dapagliflozin reduced the risk of mortality and heart failure reduction in	609

		with or without type 2 diabetes; individuals received either dapagliflozin or placebo and were followed for around 18 months	patients with heart failure and reduced ejection fraction with or without type 2 diabetes	
Empagliflozin		Patients with class II, III, or IV heart failure and an ejection fraction of 40% or less; individuals received empagliflozin or placebo and were followed for 16 months	Individuals receiving empagliflozin had a lower risk of cardiovascular death or hospitalization for heart failure than those receiving placebo, regardless of the presence or absence of diabetes	610
Empagliflozin		ob/ob ^{-/-} mice received empagliflozin for 10 weeks	Empagliflozin improved cardiac contractility, by fractional area change, and improved microvascular function after 10 weeks of treatment compared to controls	613
Empagliflozin		Spontaneous hypertensive rats (SHR) fed with high fat diet received empagliflozin for 12 weeks	Empagliflozin attenuated cardiac fibrosis and improved hemodynamics in in a hypertensive rat heart failure model	614
Insulin				
Insulin	Enhances glucose uptake and oxidation	12,537 people with cardiovascular risk factors plus impaired fasting glucose, impaired glucose tolerance, or type 2 diabetes to receive insulin or standard care	Rates of incident cardiovascular outcomes were similar in the insulin-glargine and standard-care	619
		7637 patients with type 2 diabetes to receive either insulin degludec or insulin glargine	Degludec was noninferior to glargine concerning the incidence of major cardiovascular events	620

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Table 3: The effects of different classes of heart failure therapies on

glycemic control and diabetic cardiomyopathy

Class of the therapy	Biological target of each drug	Study characteristics	Main findings	References
Renin-Aldosterone- Angiotensin System inhibitors	class			
Renin-Aldosterone- Angiotensin System inhibitors	Reduce sodium and water retention	Case-control study of 6,649 diabetic patients using ACE inhibitors	Increased the hospitalization for severe hypoglycemia	624
Temocapril		Diabetic mice treated with temocapril for 14 days	Temocapril decreased improved glycemic control	621
Captopril		Single blind, cross- over study with 16 type 2 diabetic patients treated with captopril or placebo for two 3-month treatment periods	Captopril improved glycemic control and insulin sensitivity in postprandial conditions	622
		130 diabetic patients treated with captopril for 4 months	Captopril improved glucose control	623
Lipid-lowering agents				
Lipid-lowering		Meta-analysis of 9 trials	Statins increase HbA _{1C} levels	626
agents	Inhibitor of HMG-CoA	Meta-analysis of 23 trials	Statins increase HbA _{1C} levels	627
Pitavastatin	reductase, the enzyme that catalyses the first step of cholesterol synthesis	Meta-analysis of 23 trials	Improved glycemic control	627
Atorvastatin		Meta-analysis of 23 trials	Worsened glycemic control	627
Atorvastatin, rosuvastatin, pitavastatin and simvastatin		National health screening cohort of 379,865 non- diabetic individuals	Statins increase fasting glucose levels	658

		taking statins for 11 years		
ß-receptor blockers				
Atenolol, metoprolol		Retrospective study of patients receiving either carvedilol or a selective second- generation beta- blocker (metoprolol or atenolol)	Metoprolol or atenolol increased hypoglycemia	631
Propranolol, metoprolol	Reduce	20 hypertensive diabetic patients receiving propranolol or metoprolol	Propranolol or metoprolol increased mean blood sugar levels	632
Second generation ß ₁ -selective- blockers (Metoprolol)	sympathetic output in the heart	Randomized, double-blind parallel-group trial with 1235 diabetic patients treated with carvedilol or metoprolol	Metoprolol increased HbA _{1C}	633
Third generation non-selective ß- blockers (Carvedilol)		Randomized, double-blind parallel-group trial with 1235 diabetic patients treated with carvedilol or metoprolol	Carvedilol improved insulin sensitivity	633
Aldosterone antagonists				
Spironolactone	Reduce sodium and water retention	25 dilated cardiomyopathy patients with a New York Heart Association functional class of I or II were examined before and after treatment with spironolactone for 12 months	Spironolactone reduced collagen accumulation in the heart and improved LV function	635
		30 medically treated ambulatory hypertensive patients with	Spironolactone improved diastolic function and limited fibrosis	636

	diastolic dysfunction		
	and without		
	ischemia were		
	randomized to		
	spironolactone or		
	placebo for 6		
	months		
	Eighty patients with	Spironolactone	637
	metabolic syndrome	improved diastolic	
	treated with	function and limited	
	angiotensin II	fibrosis	
	inhibition, were		
	randomized to		
	spironolactone or		
	placebo for 6		
	months		

- Mitochondrial dysfunction
 DyslipIdemia
- Insulin resistance
- Impaired insulin signaling
- Hyperglycemia
- ROS generation
- Inflammation
- Fibrosis
- Hypertrophy

- · Perturbed Ca2+ handling
- Contractile protein dysfunction Ischemia
- Autophagy
- Lipotoxicity
- Glucotoxicity
- · Cell Death
- · O-GlcNAcylation

- AGEs
- Neurohormonal mechanisms
- Hypertension
- Renal Failure
- Microangiopathy
- Coronary artery disease
- Macroangiopathy













Diabetic cardiomyopathy





Glucose oxidation







