

Cardiovascular protection by SGLT2 inhibitors — Do anti-inflammatory mechanisms play a role?



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ABSTRACT

Background: Metabolic syndrome and related metabolic disturbances represent a state of low-grade inflammation, which accelerates insulin resistance, type 2 diabetes (T2D) and cardiovascular disease (CVD) progression. Among antidiabetic medications, sodium glucose co-transporter (SGLT) 2 inhibitors are the only agents which showed remarkable reductions in heart failure (HF) hospitalizations and major cardiovascular endpoints (MACE) as well as renal endpoints regardless of diabetes status in large randomized clinical outcome trials (RCTs). Although the exact mechanisms underlying these benefits are yet to be established, growing evidence suggests that modulating inflammation by SGLT2 inhibitors may play a key role.

Scope of review: In this manuscript, we summarize the current knowledge on anti-inflammatory effects of SGLT2 inhibitors as one of the mechanisms potentially mediating their cardiovascular (CV) benefits. We introduce the different metabolic and systemic actions mediated by these agents which could mitigate inflammation, and further present the signalling pathways potentially responsible for their proposed direct anti-inflammatory effects. We also discuss controversies surrounding some of these mechanisms.

Major conclusions: SGLT2 inhibitors are promising anti-inflammatory agents by acting either indirectly via improving metabolism and reducing stress conditions or via direct modulation of inflammatory signalling pathways. These effects were achieved, to a great extent, in a glucose-independent manner which established their clinical use in HF patients with and without diabetes.

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Keywords SGLT2 inhibitors; Inflammation; Metabolism; Heart failure; Cardiovascular disease

1. INTRODUCTION

Obesity and related metabolic disorders are leading global burdens which are rapidly increasing owing to the rise in sedentary lifestyle and high caloric diet [1]. Since obesity is a strong risk factor for T2D, a parallel increase in the numbers of diabetic patients have been observed which is expected to reach 366 million in 2030 [2]. Both, T2D and obesity play a major role in the development of CVD [3], which is considered the major cause of death worldwide [4].

HF is one of the most common and serious CV complications in diabetes, which is associated with a poor prognosis [5,6]. Development of HF in diabetic patients is in fact twice as frequent as in non-diabetics [7]. Even though the incidence of HF correlates to the level of haemoglobin A1c (HbA1c) [8], lowering blood glucose by conventional anti-hyperglycaemic agents failed to show convincing evidence in reducing HF risk. In fact, all glucagon-like peptide-1 receptor agonists (GLP-1 RA) have shown at least non-inferiority and some showed significant reductions in 3-point MACE (composite of CV death, nonfatal myocardial infarction, or nonfatal stroke) in patients with T2D and high CV risk in several large RCTs [9]. Several mechanisms have been postulated to explain these CV benefits including possible direct

and indirect anti-inflammatory actions [10]. However, despite these potential mechanisms, GLP-1 RA only caused modest and inconsistent reductions in HF hospitalization rate [11]. To date, SGLT2 inhibitors are the only glucose lowering agents that reduce HF risk based on the outcomes of large RCTs, despite the fact that SGLT2 expression in the heart is quite unlikely [12].

SGLT2 inhibitors are a novel class of antidiabetic drugs which cause insulin-independent glucose lowering by reducing renal glucose reabsorption and thus enhancing glycosuria [13]. The EMPA-REG outcome trial was the first RCT to show HF and other CV outcome benefits among SGLT2 inhibitors [14]. In this trial, patients with T2D and established CVD showed reduced risk of 3-point MACE by 14% driven by a significant reduction of death from CV causes, CV death by 38%, all-cause mortality by 32%, and HF hospitalization by 35% within the first few weeks of treatment. Similar reductions in HF risk were observed with subsequent trials i.e. CANVAS for canagliflozin, DECLARE-TIMI 58 for dapagliflozin and VERTIS CV for ertugliflozin, showing ca. 30% reduction of HF hospitalization compared to placebo which confirm the role of SGLT2 inhibitors in preventing or delaying HF onset [15–17]. SOLOIST-WHF was the first trial to suggest that the potential of SGLT2 inhibitors lies beyond HF prevention [18].

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List of abbreviations

ACEI	angiotensin-converting enzyme inhibitors	IRS	insulin receptor substrate
ADP	adenosine diphosphate	JAK	Janus kinase
AGE	advanced glycation end-products	JNK	c-Jun N-terminal kinases
AngII	angiotensin II	LPS	lipopolysaccharide
AMP	adenosine monophosphate	Luseo	luseogliflozin
AMPK	5' adenosine monophosphate-activated protein kinase	LV	left ventricular
AP	activator protein	MACE	major cardiovascular endpoints
ARB	angiotensin II receptor blockers	MAPK	mitogen-activated protein kinase
ASC	apoptosis-associated speck-like protein	MCP	monocyte chemoattractant protein
ATM	adipose tissue macrophages	MMP	matrix metalloproteinases
ATP	adenosine triphosphate	MRI	magnetic resonance imaging
CAD	coronary artery disease	MT	melatonin membrane receptor
Cana	canagliflozin	mTOR	mammalian target of rapamycin
cGMP	cyclic guanosine monophosphate	NAD⁺	nicotinamide adenine dinucleotide
COX	cyclooxygenase	NADPH	nicotinamide adenine dinucleotide phosphate
CV	cardiovascular	NF-κB	nuclear factor-κB
CVD	cardiovascular disease	NLRP3	nucleotide-binding domain, leucine-rich-containing family, pyrin domain-containing-3
DAMP	damage associated molecular patterns	NO	nitric oxide
Dapa	dapagliflozin	PAI	plasminogen activator inhibitor
EAT	epicardial adipose tissue	PAMP	pathogen associated molecular patterns
EMA	European Medicines Agency	PGC	peroxisome proliferator-activated receptor gamma coactivator
Empa	empagliflozin	PI3K	phosphatidylinositol 3-kinase
Ertu	ertugliflozin	PKG	protein kinase G
eNOS	endothelial nitric oxide synthase	PKR	protein kinase R
FDA	Food and Drug Administration	PTEN	phosphorylated phosphatase and tension homologue
FFA	Free fatty acids	PVAT	perivascular adipose tissue
FOXO3a	forkhead box O3a	RAS	renin-angiotensin system
FOXp3	forkhead box P3	RCTs	randomized clinical outcome trials
GLP-1 RA	glucagon-like peptide 1 receptor agonists	ROS	reactive oxygen species
GSK3β	glycogen synthase kinase 3β	SGLT	sodium glucose co-transporter
HbA1c	haemoglobin A1c	SIRT	sirtuin
HDAC	histone deacetylase	SOD	superoxide dismutase
HF	heart failure	Sota	sotagliflozin
HFpEF	HF with preserved ejection fraction	STAT	signal transducer and activator of transcription
HFrEF	HF with reduced ejection fraction	STZ	streptozotocin
Hk	human kidney	T2D	type 2 diabetes
Hs-CRP	high sensitivity C- reactive protein	TCA	tricarboxylic acid
HUVEC	human umbilical vein endothelial cells	TGF	tumour growth factor
ICAM	intracellular adhesion molecule	TLR	toll-like receptors
IFN	Interferon	TNF	tumour necrosis factor
IGF	insulin-like growth factor	Tofo	tofogliflozin
IKK	IκB kinase	UCP	uncoupling protein
IL	interleukin	VCAM	vascular cell adhesion molecule
iNOS	inducible NOS	WAT	white adipose tissue
Ipra	ipragliflozin	XO	xanthine oxidase
IRF	interferon regulatory factor	ZDF	Zucker diabetic fatty
		βOH	β-hydroxybutyrate

Sotagliflozin reduced CV death, HF hospitalization and urgent HF visits (HR, 0.67; 95% CI, 0.52 to 0.85) in diabetic patients with recent worsening HF. Recently, this therapeutic role of SGLT2 inhibitors was further supported in DAPA-HF and EMPEROR-Reduced, where dapagliflozin and empagliflozin reduced the composite of worsening HF (i.e. hospitalisation or urgent visit for HF) and CV mortality regardless of diabetes status of HF with reduced ejection fraction (HFrEF) patients [19,20]. These benefits were driven by 30% reductions in the first or recurrent HF hospitalizations [21]. Accordingly, SGLT2 inhibitors are now established as first line agents for HFrEF management in recent guidelines [22]. In 2021, EMPEROR-Preserved

has established, for the first time, the CV benefits of SGLT2 inhibitors in HF with preserved ejection fraction (HFpEF) patients with or without diabetes with similar reductions in the primary endpoint as shown in EMPEROR-Reduced [23]. The ongoing large trial; DELIVER, is expected to provide further therapeutic implications in this patient population [24]. In a press release, however, it was announced that top-line results from the study showed reductions in the primary endpoint [25]. Apart from their diuretic and natriuretic effects [26], several novel hypotheses have been proposed to explain the early remarkable CV benefits of SGLT2 inhibitors. These include increased erythropoiesis [27], improved cardiac remodelling [28], improved ion balance and

mitochondrial energetics [12] and inhibition of sympathetic stimulation [29]. While the exact mechanisms are still under debate, the anti-inflammatory effects of SGLT2 inhibitors have come more into focus based on recent findings obtained mostly from animal and cell culture studies, as discussed in the next sections.

2. INFLAMMATION LINKS OBESITY, DIABETES AND HEART FAILURE

Obesity and related metabolic disturbances are currently considered as conditions of chronic low-grade inflammation, which contributes to the incidence and progression of insulin resistance [30]. The expanded visceral white adipose tissue (WAT) in obesity represents not only a massive lipid storage depot, but also a major source of proinflammatory cytokines (adipokines) [31–34]. Supportive evidence of the role of inflammation in obesity-associated metabolic disorders arises from the promising results of salicylates, anakinra and infliximab in ameliorating hyperglycaemia [35–39]. Free fatty acids (FFA) released from adipose tissues into the systemic circulation bind to toll-like receptors (TLR) – 2 and 4 in metabolic cells such as adipose tissue, liver and muscles, activating downstream kinases; c-Jun N-terminal kinases (JNK), I κ B kinase (IKK) and protein kinase R (PKR) [40]. These kinases mediate serine phosphorylation of insulin receptor substrate (IRS)-1 resulting in insulin resistance. Ultimately, these kinases activate inflammatory transcription factors; nuclear factor- κ B (NF- κ B), activator protein (AP)-1 and interferon regulatory factor (IRF) contributing to inflammation. The activation of nucleotide-binding domain, leucine-rich-containing family, pyrin domain-containing (NLRP) 3 inflammasome can further contribute to the inflammatory status [41], which may be explained by the reduced 5' adenosine monophosphate-activated protein kinase (AMPK) signalling by elevated FFA levels [42]. Moreover, hypoxia resulting from the rapid expanding adipose tissue is associated with increased macrophage infiltration with a phenotypic conversion from the M2 type expressing anti-inflammatory cytokines to the proinflammatory M1 [43,44].

The increased proinflammatory cytokines from these pathways together with FFA further promote insulin resistance by establishing a positive feedback loop of inflammation [30]. Normally, pancreatic β -cells can adapt initially to the reduced insulin response by increasing their proliferation and insulin secretion, a mechanism which is compromised in later stages owing to increased β cell apoptosis leading to overt T2D [45]. The resulting hyperglycaemia plays a major role in inflammation and diabetic vascular complications which were shown to be also mediated by oxidative stress [46,47]. This state of chronic low-grade inflammation contributes to the progression of CVD by inducing endothelial dysfunction and atherosclerosis or by causing direct myocardial damage in the absence of CVD (or diabetic cardiomyopathy) [48,49].

In this regard, SGLT2 inhibitors were able to ameliorate markers of inflammation in clinical and *in vivo* studies, which may possibly contribute to their CV benefits. Studies demonstrated reductions of a large set of pro-inflammatory cytokines with empagliflozin including interleukin (IL)-6, tumour necrosis factor (TNF), monocyte chemoattractant protein (MCP)-1, Interferon (IFN)- γ , P-selectin and intercellular adhesion molecule (ICAM)-1 in the hearts of Zucker diabetic fatty (ZDF) rats [50,51]. In T2D patients, treatment with canagliflozin was associated with lower serum levels of leptin and IL-6 with higher adiponectin levels compared to glimepiride [52], while reductions in high sensitivity C-reactive protein (hs-CRP) and myeloperoxidase with concomitant increase in anti-inflammatory IL-10 were observed with empagliflozin [53].

These anti-inflammatory properties of SGLT2 inhibitors could be a result of general effects on metabolism, oxidative stress and renin-angiotensin system (RAS) signalling or from its direct action on inflammatory signalling pathways.

3. GENERAL EFFECTS OF SGLT2 INHIBITORS

3.1. Reduction of oxidative stress

Reactive oxygen species (ROS) are formed from redox reactions or electron excitation [54]. In physiological levels, they play an important role in the regulation of cell signalling, autophagy, immunity and cellular differentiation [55]. However, when the production of ROS exceeds the detoxification ability of the antioxidant defence mechanism, detrimental effects on lipids, proteins, lipoproteins and DNA can occur contributing to the loss of important cellular functions, cell damage and apoptosis, in a phenomenon defined as oxidative stress [56]. In the heart, ROS overproduction occurs primarily due to altered mitochondrial functions, enhanced nicotinamide adenine dinucleotide phosphate (NADPH) oxidase and xanthine oxidase (XO) activities or due to uncoupling of endothelial nitric oxide synthase (eNOS) [57]. On the other hand, depletion of antioxidants such as superoxide dismutase (SOD), catalase, glutathione peroxidase, nicotinamide adenine dinucleotide (NAD⁺) and glutathione have also contributed to CVD and HF. There is a direct link between ROS accumulation and stimulation of inflammatory pathways. ROS can mediate endothelial dysfunction and atherosclerotic plaque formation by activating NF- κ B mediated expression of several cytokines including endothelin-1 [58], vascular cell adhesion molecule (VCAM)-1 and ICAM-1 which promote macrophage infiltration and vascular inflammation [59]. Owing to eNOS uncoupling, oxidative stress enhances inducible NOS (iNOS) production to compensate for the decreased nitric oxide (NO) production [60], a potent vasodilator which is necessary to maintain endothelial functions, prevent leucocyte adhesion and atherosclerosis [61], while maintaining normal cardiac contractions [62]. However, excess NO binds to superoxide (O²•⁻) to form peroxynitrite (ONOO•⁻), which could be further involved in lipid peroxidation [63], foam cell formation and atherosclerosis [64]. In the myocardium of HF_{REF}, ROS stimulate mitogen-activated protein kinase (MAPK) mediated NF- κ B and AP-1 transcription factors which promotes release of inflammatory cytokines, cellular apoptosis and activation of matrix metalloproteinases (MMP), which contribute to collagen degradation and left ventricular (LV) dilatation [65]. In HF_{pEF}, on the other hand, ROS mediated eNOS inactivation reduces nitric oxide-cyclic guanosine monophosphate-protein kinase G (NO-cGMP-PKG) signalling and thus leads to titin hypophosphorylation, predisposing to myocardial stiffness and diastolic dysfunction [66].

Several studies have reported the ability of SGLT2 inhibitors in mitigating oxidative stress either by inducing metabolic changes (as will be mentioned in the next sections) or by acting as an antioxidant per se. Dapagliflozin reduced the levels of the antioxidant DJ-1 and Nrf2 that was elevated as a compensatory mechanism to neutralize the excessive lipid peroxides produced in a Parkinson's disease rat model [67], while it reduced ROS and its associated apoptosis and NF- κ B activation. In postinfarcted rats, dapagliflozin reduced myocardial superoxide and nitrotyrosine levels which resulted in activation of the signal transducer and activator of transcription (STAT) 3 signalling pathway [68]. STAT3 is a key regulator of macrophage polarization, which upon activation by dapagliflozin enhanced anti-inflammatory M2 macrophage expression and IL-10 release, whereas cardiac fibrosis was attenuated.

Furthermore, SGLT2 inhibitors act on the main sources of ROS in the heart. Empagliflozin improved mitochondrial function in rats with LV dysfunction by enhancing the expression of peroxisome proliferator-activated receptor gamma coactivator (PGC) 1- α , an important mediator of mitochondrial bioenergetics [69]. On the other hand, SGLT2 inhibitors downregulated NADPH oxidase expression *in-vitro* and *in-vivo* [69,70] with subsequent reduction of H₂O₂ and superoxide levels [70], while it restored oxidant-antioxidant balance by attenuating overexpressed SOD. In LV tissue samples of HFpEF patients, empagliflozin increased NO-cGMP-PKG signalling leading to enhanced titin phosphorylation with subsequent reduction in cardiac stiffness and LV dysfunction [71–73]. At the same time, empagliflozin enhanced endothelial relaxation and function by downregulating ICAM-1, VCAM-1, TNF, and IL-6 levels [71,72]. These effects have been attributed to the inhibition of eNOS uncoupling and its associated oxidative stress, which could be related to the direct activation of SOD [73].

3.2. Reduction of glucotoxicity

Large prospective cohort studies have repeatedly shown the strong association between the indices of glycemia and risk of CVD and mortality in diabetic patients [74–76], which was supported by evidence indicating the role of hyperglycaemia in promoting endothelial dysfunction [77–79] and atherosclerosis *in-vivo* [80]. Oxidative stress has been proposed as the link between hyperglycaemia and its vascular damage [81,82].

Glucose is normally metabolized by tricarboxylic acid (TCA) cycle to produce adenosine triphosphate (ATP) molecules required for various cellular functions [82]. The electrons generated from the mitochondrial electron transport chain are then transferred to oxygen to be reduced to water. However, in hyperglycaemia, more glucose enters the TCA cycle causing the excess electrons to be captured by coenzyme Q and finally generating superoxide. Increasing superoxide levels by hyperglycaemia stimulates mechanisms of glucotoxicity [83], all of which activate NADPH oxidase and further enhance ROS production. Thus, contributing to induced inflammation and potential risk of CVD.

Whether the anti-inflammatory effects of SGLT2 inhibitors could be partly explained by reduction of hyperglycaemia has been the subject of a considerable debate.

In ZDF rats, empagliflozin improved endothelial function by ameliorating the activation of advanced glycation end-products (AGE) to their receptors and the associated oxidative stress and inflammation in whole blood and aortic tissues [50]. This has been attributed to the glucose lowering effects and improved glucose utilization by empagliflozin. Similar results were reported in the aortic tissues and blood of streptozotocin (STZ) treated rats [84] as well as aortic rings and perivascular adipose tissue (PVAT) of STZ treated ApoE^{-/-} mice [85]. Glucose normalization by empagliflozin in STZ-treated mice mediated atherosclerosis regression in aortic roots [86], while it significantly reduced cardiac and coronary arterial fibrosis and improved aortic endothelial function in *db/db* mice [87], which have been explained by the alleviation of hyperglycaemia and associated oxidative stress. Similarly, tofogliflozin attenuated tubulointerstitial inflammation in diabetic mice via normalization of blood glucose levels [88].

On the other hand, several studies have excluded glycaemic control as the sole mechanism behind SGLT2 inhibitors benefits. First, some of these studies have utilized non-diabetic models of CVD which were generated using lipopolysaccharide (LPS) [89–91] or angiotensin II (AngII) [92,93] as inflammatory stimulants or arterial ligation to induce ischemia [68,69,94] without altering blood glucose levels. In this view, recent RCTs such as DAPA-HF, EMPEROR-Reduced and EMPEROR-Preserved showed CV benefits in HF patients without diabetes which

support the glucose-independent mechanisms of SGLT2 inhibitors [19,20,23]. Second, *in-vitro* experiments which were carried out on myocardial cells [70] - currently known as non-expressors of SGLT2 - ensure that glucose transport into the cells is not affected by SGLT2 inhibition. Third, superior anti-inflammatory effects have been reported with SGLT2 inhibitors compared with other anti-diabetic agents with similar glucose lowering effects [95,96].

In fact, lowering glucose levels has substantially reduced the risk of microvascular complications in diabetic patients [97,98], while the benefits regarding macrovascular complications are still uncertain. In a meta-analysis, intensive glycaemic control reduced risk of non-fatal myocardial infarction and coronary heart disease by about 15% without affecting rate of stroke, HF and all-cause mortality [99]. In agreement with that, another meta-analysis comprised of four large RCTs; VADT, UKPDS, ACCORD and ADVANCE reported glucose-lowering associated reduction in MACE without affecting all-cause mortality [100]. In fact, the ACCORD trial showed an increased mortality rate with intensive therapy in diabetic patients [101], while UKPDS was the only trial among the four to show a 13% reduction in death from any cause which was manifested over 10 years of follow-up [102]. This could be related to the healthier patient population, less intensive glycaemic control and longer follow-up period utilized in this trial as compared to others [103]. On the other hand, the benefits observed by SGLT2 inhibitors in the recent RCTs diverge in the first few weeks of treatment which contradicts the long-term period required for glycaemic control to manifest improved CV outcomes as observed with other antidiabetic agents in the UKPDS trial.

3.3. Reduction of hyperuricemia

Uric acid is the end product of endogenous and dietary purine metabolism which is disposed mainly through the kidneys [104]. Elevation of serum uric acid could be attributed to the imbalance between uric acid production and excretion, which are regulated by several enzymes such as the xanthine oxidase (XO). In the context of HF and CVD, the incidence of hyperuricemia is common owing to the reduction of uric acid excretion and increasing its production in HF [105,106], the increased use of diuretics and low dose aspirin which stimulate uric acid tubular reabsorption [107], and the association of hyperuricemia to common CV risk factors e.g. diabetes, hypertension and insulin resistance [108–110].

Elevated serum uric acid was shown to be correlated to inflammatory markers in several clinical studies [111–113]. Activated inflammatory pathways could be a result of the underlying elevated oxidative stress caused by uric acid stimulation of NADPH oxidase [114–116], which is believed to be mediated by uric acid activation of RAS in some *in-vitro* studies [117,118]. In mice with unilateral ureteral obstruction, high uric acid levels enhanced fibrosis through activation of ROS/NLRP3/IL-1B signalling which was reversed by allopurinol (XO inhibitor) administration [119]. Similarly, allopurinol reversed ROS mediated uric acid activation of NLRP3 inflammasome that was responsible for endothelial injury in a chronic kidney disease rat model [120]. Moreover, elevated uric acid levels directly stimulated NF- κ B mediated pro-inflammatory cytokine release in mouse kidneys and *in-vitro*, which was blocked by inhibiting tubular uric acid transporters [121]. The same effect of uric acid on NF- κ B was associated with increased incidence of dyslipidaemia and hyperglycaemia in rats [122] and was responsible for reduced NO levels and endothelial dysfunction in human umbilical vein endothelial cells (HUVEC) [123]. Exposure to uric acid increased CRP-mediated human vascular smooth muscle cell migration and reduction of NO from HUVEC, which could be partially

explained by MAPK activation [124]. Incubation of the cells with probenecid, an inhibitor of uric acid cellular entry, reversed these effects. On this background, elevated uric acid levels have been associated with increased CV risk, mortality and incidence of HF in several large cohort studies [125–127]. Thus, it is not surprising to find that several studies have attributed the improved CV and HF outcomes to lowering uric acid levels after administration of XO inhibitors in patients with high CV risk [128,129].

SGLT2 inhibitors have been shown to reduce the levels of circulating uric acid regardless of diabetic status [130,131] and in patients with CV risk [27,132]. Since these agents reduce glucose transport through SGLT2, the resulting high glucose concentration in proximal tubules facilitates glucose exchange with intracellular uric acid through GLUT9, thus increasing uric acid elimination [133]. In this view, attenuation of hyperuricemia by SGLT2 inhibitors has been proposed as a potential mechanism of their CV benefits via reducing oxidative stress, inflammation, endothelial dysfunction and fibrosis [134].

Conversely, other studies have failed to show any improvement in CV or HF outcomes upon treatment with XO inhibitors [135–137]. In fact, patients treated with these agents might even show a trend of increasing CV mortality and hospitalization [135,138]. This evidence supports the fact that elevated uric acid is just a biomarker of oxidative stress and does not stimulate ROS or inflammatory cytokine release per se which precipitate myocardial injury [139]. In pro-oxidant conditions like in CVD, eNOS is uncoupled and becomes unable to produce NO, which beside its vasculoprotective effects plays an important role in ROS quenching [140]. Instead, XO level is upregulated to act as an alternative source of NO production together with a parallel increase in uric acid levels [141]. Thus, inhibition of XO deprives the body from an important antioxidant and NO source which could contribute to increased CV risk.

SGLT2 inhibitors, on the other hand, elicit a fasting-like state probably due to glycosuria, which in turn activates sirtuin (SIRT)1, an enzyme which plays an important role in the protection against oxidative stress and inflammation [142]. SIRT1 restores eNOS ability to produce NO [143] and consequently lowers XO and its associated uric acid levels [144]. Therefore, it can be hypothesized that the CV benefits of SGLT2 inhibitors are mainly mediated by the indirect activation of SIRT1 antioxidant and anti-inflammatory pathways, rather than by lowering serum uric acid.

3.4. Enhancing ketonemia

In diabetic failing hearts, there is an increased reliance on FFA oxidation for energy production owing to reduced glucose utilization on the basis of insulin resistance [145]. Although FFA produce more ATP than glucose, this comes with the price of increased oxygen consumption and ROS generation which can further exacerbate HF [146]. Against this background, enhancing ketogenesis by SGLT2 inhibitors may in part explain their cardioprotective effect, since ketone bodies are more energy efficient than FFA; yielding more ATP molecules per molecule of oxygen utilized [147]. Ferannini et al. have proposed that empagliflozin-associated glycosuria is responsible for the reduction of the insulin/glucagon ratio, which enhances hepatic FFA oxidation and subsequent elevation of circulating β -hydroxybutyrate (β OHB) [148] to be eventually utilized by the failing heart as a metabolic stress defence [149]. This hypothesis was experimentally proved in non-diabetic animal models of HFrEF, where enhanced myocardial utilization of ketone bodies by empagliflozin was associated with attenuation of adverse cardiac remodelling [69,150] and diastolic dysfunction [73]. In that light, β OHB also improved cardiac output, ejection fraction and

reduced systemic vascular resistance in HFrEF patients [151]. Interestingly, Lopaschuk and Verma have proposed a counterargument, stating that increased β OHB by empagliflozin is probably due to reduced myocardial ketone body utilization, a process which could be maladaptive on long term [152].

Regardless of the mechanism of increase, it was suggested that elevated cardiac β OHB increased histone acetylation and expression of oxidative stress resistant factors; melatonin membrane receptor (MT)2 and forkhead box O3a (FOXO3a) by inhibiting histone deacetylase (HDAC) [153]. Beside its antioxidant effect, HDAC inhibition by β OHB can also mitigate cardiac and extra cardiac inflammation. Rats treated with valproic acid, an HDAC inhibitor, showed reduced ventricular levels of NF- κ B, TNF, IL-1 β and ROS which attenuated cardiac hypertrophy and fibrosis [154]. Another HDAC inhibitor, suberoylanilide hydroxamic acid, reduced a diverse range of inflammatory cytokines, which diminished cardiovascular fibrosis and stiffness in hypertensive rats [155]. Acetylation of the MAPK phosphatase-1 by HDAC inhibitors attenuated MAPK signalling *in-vitro*, which was associated with reduced inflammation and mortality in LPS-treated mice [156]. Furthermore, HDAC inhibitors enhanced production of anti-inflammatory regulatory T cells through the acetylation of forkhead box P3 (FOXP3), a key transcription factor for regulatory T cells development [157]. These inhibitors also enhanced macrophage polarization to the anti-inflammatory M2 phenotype [158] through modulation of the glycogen synthase kinase 3 β /phosphorylated phosphatase and tension homologue/phosphatidylinositol 3-kinase or GSK3 β /PTEN/PI3K signalling pathway [159].

Enhanced adiponectin expression by β OHB was associated with reduced pro-inflammatory cytokines in 3T3-L1 adipocytes, mediated by the direct epigenetic modification of β OHB on histone H3K9 of the adiponectin gene [160].

Furthermore, modulation of the NLRP3 inflammasome by β OHB has been reported in several studies [161–163]. Youm et al. was the first to show the impact of β OHB on the NLRP3- inflammasome and its mediated secretion of IL-1 β in human macrophages and mice [164]. Later, Byrne et al. showed the important role of high β OHB in inhibition of NLRP3 inflammasome activation with subsequent reduction in pro-inflammatory cytokine levels and macrophage infiltration into cardiac tissues of HF mice [165]. In this study, elevated β OHB was associated with reduced cardiac remodelling and improved diastolic filling parameters.

Thus, inhibiting HDAC and NLRP3 inflammasome or enhancing adiponectin expression by elevated β OHB could contribute to the anti-inflammatory effect of SGLT2 inhibitors. However, only limited studies have shown the impact of SGLT2 inhibitor-induced ketonemia on inflammation and associated CV risk.

Recently, Kim et al. showed that empagliflozin attenuated NLRP3 inflammasome activation and the secretion of IL-1 β which could be attributed to the elevated levels of β OHB in macrophages of diabetic patients with CVD [166]. In a HFrEF mouse model, elevated β OHB by empagliflozin inhibited mitochondrial protein hyperacetylation resulting in reduced NLRP3 inflammasome assembly and subsequent cytokine release [167]. These effects were associated with reduced BNP levels, cardiac fibrosis and stiffness, while they improved exercise tolerance. In diabetic patients with history of CVD, elevated β OHB resulted in improved left ventricular diastolic functions which was reflected by reduced E/e' in echocardiography after treatment with tofogliflozin [168]. The anti-inflammatory and antioxidant effects of increased β OHB might explain, at least partially, the improvement in cardiac function in these patients and potential reduction of their HF risk.

3.5. Reduction of adipose tissue mass and associated pro-inflammatory cytokine release

Obese patients with metabolic syndrome are characterized by WAT hypertrophy in the face of increased demand of energy storage together with impaired angiogenesis, hypoxia and adipocyte apoptosis leading to macrophage infiltration and inflammation [169]. Subsequently, adipose tissue differentiation is impaired which predisposes to ectopic fat accumulation in tissues such as the heart and blood vessels. Therefore, obesity-induced inflammation can promote CVD progression through the secretion of pro-inflammatory cytokine into the circulation from distant adipose tissue depots in an endocrine manner or from the adjacent PVAT and epicardial adipose tissue (EAT) through paracrine release [170].

PVAT surrounds large arteries and veins as well as small resistant vessels, while it is completely absent from the cerebral circulation [171]. These adipose tissues regulate vascular tone in healthy conditions by releasing several relaxing factors such as adiponectin, NO and hydrogen sulphide. In obesity, however, PVAT loses its anti-contractile functions which can predispose to endothelial dysfunction and atherogenesis. In PVAT dysfunction, NO production is reduced owing to enhanced eNOS uncoupling which subsequently contributes to ultimate generation of superoxide [172,173] promoting vasoconstriction and vascular remodelling [174]. Proinflammatory adipokines such as leptin and resistin can further exacerbate oxidative stress in the vascular endothelium by activating NADPH oxidase [175]. Furthermore, PVAT release of visfatin and resistin was associated with enhanced ICAM-1 and VCAM-1 expression which can mediate leucocyte infiltration of vascular endothelial cells and atherosclerosis [176,177]. In clinical studies, PVAT inflammation predisposed to coronary artery vasospasm and increased atherosclerotic burden in patients with vasospastic angina and coronary artery disease (CAD), respectively [178,179].

EAT lies in close proximity to the heart i.e. between the myocardium and the pericardial visceral layer [180] acting as an energy source to the underlying myocytes by providing FFA, while it exerts antioxidant effect by the released adiponectin [181]. Like PVAT, the accumulated EAT in obesity releases various proinflammatory mediators such as leptin, resistin and other adipokines with reduced adiponectin production [182]. These changes were associated with increased myocardial inflammation promoting cardiac fibrosis and remodelling, which can contribute to impaired myocardial contractility [183]. In a similar manner, EAT can influence the underlying coronary vasculature promoting atherogenesis and microvascular rarefaction [184,185]. In this regard, EAT contributed to increased left ventricular mass and worse diastolic functions in HF patients [186–188].

SGLT2 inhibitors have shown consistent reductions in body weight of 1–3 kg in the first weeks of treatment [189] even in obese patients without T2D [190], which could be maintained up to 4 years [191]. Although the impact of weight loss on HF outcomes has been so far seen sceptically [192], the reduction of visceral fat content associated with dapagliflozin and ipragliflozin reduced LV mass compared to placebo and metformin, respectively, which highlights the impact of VAT inflammation on CV remodelling [28,193].

On the same note, T2D patients treated with SGLT2 inhibitors exhibited reduced EAT mass [194–196], which was assumed to be independent of glycaemic control in some studies [197,198]. This was confirmed by the positive findings observed even in non-obese, non-diabetic HFREF patients using cardiac magnetic resonance imaging (MRI) [199]. The observed EAT reductions were much greater than the changes (if any) in body weight, VAT and subcutaneous fat. This was attributed to the high turn-over rate of FFA in EAT which could explain their higher

sensitivity to SGLT2 inhibitors. Loss of EAT by SGLT2 inhibitors showed parallel reductions in pro-inflammatory cytokines [195,196], which might explain the observed reductions of interstitial myocardial fibrosis and aortic stiffness by empagliflozin in nondiabetic patients with HFREF [200]. These findings contributed to improved diastolic functions, pulsatile load and ventricular pressure. Conversely, Gaborit et al. showed no change in EAT volume after treatment with empagliflozin in T2D patients [201], which could be attributed to the different cardiac imaging modalities used as compared to the previously mentioned studies [202]. This emphasizes the importance of utilizing more sensitive multi-slice volumetric approaches for EAT measurement using cardiac MRI instead of the single-slice methods such as computed tomography [203]. However, Gaborit et al. did not exclude the possibility of improved EAT phenotype by empagliflozin that could positively influence cardiac functions [201]. In this regard, dapagliflozin improved EAT stromal cell differentiation, reduced pro-inflammatory cytokines and enhanced EAT secretome contributing to the healing of endothelial cells in patients undergoing cardiac surgery without affecting EAT mass [204].

On the other hand, few studies showed paradoxical effects of SGLT2 inhibitors on PVAT *in vivo*. Luseogliflozin attenuated neointimal hyperplasia after wire injury in mice receiving high fat diet by reducing adipocyte size of PVAT which was associated with increased adiponectin levels and reduced macrophage infiltration [205]. Mori et al. also showed that ipragliflozin reduced inflammation, adipocyte apoptosis and macrophage accumulation in abdominal PVAT of obese diabetic mice resulting in suppression of adverse vascular remodelling [206]. Surprisingly, ipragliflozin in this study increased adipocyte size and lipid storage capacity owing to increased insulin sensitivity which was designated as “healthy adipose tissue expansion”. Another study, however, questioned this cardioprotective effect of SGLT2 inhibitors in a rat model of metabolic syndrome [207], since tofogliflozin failed to enhance PVAT-mediated vasorelaxation with no associated improvement in cardiac function and heart weight.

The mechanisms behind adipose tissue loss with SGLT2 inhibitors cannot be simply explained by caloric loss due to glycosuria [208]. Alternatively, the low insulin/glucagon ratio mediated by the reduced plasma glucose levels with SGLT2 inhibitors shifts energy metabolism towards fat utilization, fatty acid oxidation and ketogenesis in T2D patients [209], which could be AMPK dependent [210]. In obese mice, empagliflozin was associated with enhanced WAT browning by increasing the expression of uncoupling protein (UCP)-1 [208], a protein responsible for energy dissipation and thermogenesis in brown adipose tissue [211]. Together with increased adiponectin levels, enhanced M2-macrophage polarization and its associated sympathetic stimulation could be the mechanisms behind empagliflozin-mediated UCP-1 expression [208]. Sawada et al. showed that in high fat fed mice, tofogliflozin promoted glycogen depletion which subsequently mediated, at least partly, sympathetic stimulation via liver-brain-adipose-neural axis [212]. The activation of this neurological pathway was associated with enhanced lipolysis in WAT and eventually weight loss.

3.6. Attenuation of RAS signalling

RAS plays a key role in the development of hypertension and CVD which is mediated, to a great extent, by the pro-inflammatory effects of AngII signalling pathway [213]. Initially, AngII sets the stage for the inflammatory process by enhancing vascular permeability through increasing BP or by releasing prostaglandins and vascular endothelial growth factors [214]. Subsequently, the role of AngII in leucocyte recruitment to the vascular wall comes into play. AngII increases

proinflammatory cytokine and chemokine release together with the expression of adhesion molecules such as E-selectin, ICAM-1 and VCAM-1 thus enhancing leucocyte endothelial interaction. Thus, AngII can contribute to endothelial dysfunction and progression of atherosclerosis [215,216]. Furthermore, AngII is involved in the production of growth and fibrotic mediators which can augment thrombus formation such as plasminogen activator inhibitor (PAI)-1 [217], or mediate vascular remodelling and hypertension by promoting tumour growth factor (TGF)- β /Smad signalling [218]. The underlying mechanism of the inflammatory effects of AngII is primarily attributed to enhanced ROS production through AngII mediated activation of NADPH oxidase [219] and XO [220] or its associated mitochondrial dysfunction [221].

Further, AngII enhances NF- κ B activity by promoting its DNA binding or by inducing its nuclear translocation through the degradation of its inhibitor I κ B [222]. Against this background, inhibition of AngII signalling by angiotensin-converting enzyme inhibitors (ACEI) or angiotensin II receptor blockers (ARB) blunted the expression of proinflammatory cytokines in patients with CAD [223], hypertension [224] and HF [225] which could explain the decreased mortality and CV benefits of these agents in high risk patients [226,227]. Similarly, the deleterious effects of RAS activation were attenuated by the administration of SGLT2 inhibitors in diabetic and non-diabetic models. Empagliflozin blocked *in-vivo* AngII mediated inflammation by inhibiting NF- κ B and MAPK activation, which consequently

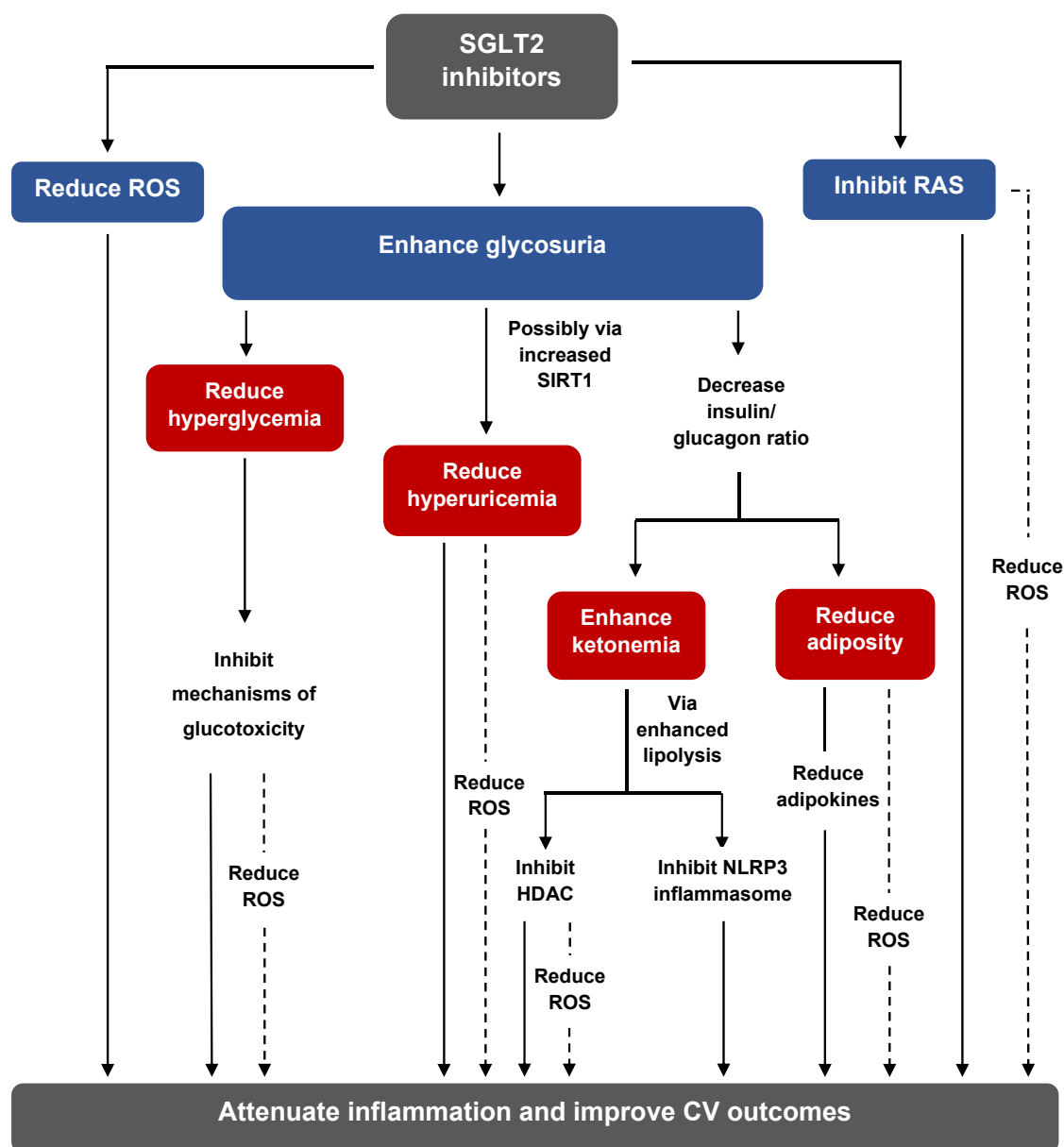


Figure 1: General effects of SGLT2 inhibitors indirectly contributing to reduced inflammation. Apart from being glucose lowering agents, SGLT2 inhibitors act by modulating other metabolic pathways through their glycosuric effects which mediate reduction of uric acid, increased ketone bodies and adipose tissue loss. Systemically, they interfere with ROS generation and RAS signalling by acting on their major sources. SGLT2 inhibitors can, therefore, indirectly attenuate inflammatory signalling via these mechanisms which might contribute to their established CV benefits. Abbreviations: HDAC, histone deacetylase; NLRP3, nucleotide-binding domain leucine-rich-containing family, pyrin domain-containing-3; RAS, renin-angiotensin system; ROS, reactive oxygen species; SGLT2, sodium glucose co-transporter 2; SIRT1, sirtuin1.

attenuated macrophage infiltration, neovascularization and endothelial dysfunction in the suprarenal aorta [92]. In kidney tissue of diabetic mice, treatment with dapagliflozin was associated with reduced urinary AngII levels, corresponding decrease in ROS and enhanced antioxidant expression compared to voglibose, the comparator anti-diabetic agent [96]. These effects contributed to reduced inflammatory cell infiltration, tubulointerstitial fibrosis and collagen accumulation. Since both antidiabetic agents had similar glucose lowering effects, the superior antioxidant and anti-inflammatory benefits of dapagliflozin were mainly related to a glucose-independent mechanism. Similar renal protective effects have been demonstrated by canagliflozin in AngII treated mice [228]. In this study, however, RAS activation was associated with enhanced SGLT2 expression in kidney cells *in-vitro* and *in-vivo*, which was oxidative stress mediated. Thus, the protective effects of canagliflozin in this study are assumed to be a consequence of direct inhibition of SGLT2 rather than an off-target effect.

Although the expression of SGLT2 in endothelial cells is controversial, Park et al. interestingly showed that AngII enhanced NADPH oxidase activity via its action on type 1 angiotensin receptor which ultimately increased the expression of SGLT1 and SGLT2 in rat endothelial cells [229]. This signalling pathway was associated with endothelial dysfunction resulting from reduction in eNOS levels and subsequent NO release together with enhanced expression of VCAM-1, MCP-1 and tissue factor. Sotagliflozin and empagliflozin reversed these effects, highlighting the possible contribution of SGLT2 in mediating RAS outcomes. Attenuation of inflammation by SGLT2 inhibitors might be responsible for the inhibition of AngII stimulated TGF- β /Smad signalling in rats, thus ameliorating cardiac fibrosis, remodelling and diastolic dysfunction [93].

Inhibition of RAS by SGLT2 inhibitor, however, has been challenged by some studies. In patients with T2D, SGLT2 inhibitors contributed to enhanced RAS activity after 30 days, possibly as a compensatory mechanism to their natriuretic and osmotic diuretic effects [230]. Similar results were reported in diabetic patients with canagliflozin and dapagliflozin [231,232], while renin and aldosterone levels were not changed by dapagliflozin in another study of T2D [233].

Figure 1 summarizes the possible metabolic and systemic alterations which could be related to the indirect anti-inflammatory effects of SGLT2 inhibitors.

4. EFFECTS OF SGLT2 INHIBITORS ON INFLAMMATORY SIGNALLING PATHWAYS

4.1. AMPK activation

AMPK is the fuel gauge of the cell, which under stress conditions, reduces energy consumption and enhances compensatory production of ATP resulting in increased ATP/adenosine diphosphate (ADP) ratio [234]. Therefore, AMPK can regulate major metabolic pathways of glucose, lipids and proteins and is considered as a key player in autophagy. The anti-inflammatory role of AMPK has been demonstrated in diabetic patients treated with metformin which showed AMPK mediated inhibition of NLRP3 inflammasome and IL-1 β release [235]. In obese mice, AMPK deficiency enhanced macrophage polarization to the pro-inflammatory M1 type [236]. With respect to vascular cells, AMPK activation reduced TNF-stimulated NF- κ B activity [237] and the subsequent expression of adhesion molecules [238,239] in cultured endothelial cells. Again with metformin, the activation of AMPK contributed to reduced NF- κ B, iNOS and cyclooxygenase (COX)-2 expressions in vascular smooth muscle cells [240]. Furthermore, incubating fibroblasts and endothelial cells with IL-6 stimulated Janus

kinase (JAK)—STAT pathway mediated inflammation, which was inhibited upon activation of AMPK by salicylate and metformin [241]. Thus, enhancing the AMPK pathway is associated with inhibition of vascular inflammation and endothelial dysfunction, which could contribute, at least partly, to less cardiac remodelling and myocardial dysfunction [242–245].

SGLT2 inhibitors have been shown to act as AMPK activators through inhibition of complex I of the mitochondrial respiratory chain, which subsequently increases adenosine monophosphate (AMP) and ADP content [210]. Elevated AMP/ADP bind to the γ subunit of AMPK which then activates its phosphorylation at threonine 172.

In diabetic models, SGLT2 inhibitors showed anti-inflammatory effects through activation of AMPK signalling. Dapagliflozin reduced NF- κ B nuclear translocation in renal tubular human kidney (HK)-2 cells [246], while it reduced NLRP3 inflammasome activation and progression of diabetic nephropathy in mice [247], which were AMPK mediated. Through the same pathway, empagliflozin reduced the expression of IL-6, TNF and MCP-1 in the hearts of diabetic rats [51], while in cultured human cells canagliflozin reduced IL-1 β stimulated IL-6 and MCP-1 secretion probably through the inhibition of facilitative glucose uptake [248].

However, other *in-vitro* and *in-vivo* studies showed that reduced glucose levels by SGLT2 inhibitors cannot be the only mechanism explaining the AMPK-mediated anti-inflammatory effects. Inflammatory cytokine levels were attenuated [91,249] and the expression of M2 macrophages was enhanced [249] upon treatment with SGLT2 inhibitors in an AMPK-dependent manner in LPS-stimulated *in-vitro* and *in-vivo* models. Furthermore, the reduction of NLRP3 inflammasome by dapagliflozin through AMPK activation in the hearts of diabetic mice has been successfully replicated in cardiomyocytes *in-vitro* [90]. These AMPK- anti-inflammatory effects possibly contributed to inhibition of pro-fibrotic TGF- β /Smad signalling [250], reduced ventricular remodelling and improved cardiac function [90,250].

Autophagy is an important process for cellular homeostasis, by which misfolded proteins, damaged organelles and pathogens are captured by autophagosomes to be degraded by lysosomal proteases [251]. Accordingly, dysregulation of this process can predispose to inflammation [252,253] and can contribute to CVD [254–256]. Autophagy is regulated, among others, by AMPK/the mammalian target of rapamycin (mTOR) signalling [257]. Thus, modulation of this pathway restores autophagy and can contribute to the anti-inflammatory effects of AMPK activators [258,259]. Likewise, empagliflozin enhanced autophagy by stimulating AMPK and inhibiting mTOR in a mouse model of liver disease [260,261], accompanied by a concomitant reduction of IL-17/IL-23 release [261]. Moreover, the improvement of cardiac function by empagliflozin *in-vivo* was attributed to autophagosome accumulation and enhanced autophagic flux mediated by AMPK/mTOR pathway [262].

4.2. Inhibition of NLRP3 inflammasome activation

The NLRP3 inflammasome is a macromolecular protein complex which is comprised of 3 main components: NLRP3, apoptosis-associated speck-like protein (ASC) and pro-caspase-1 [263]. In response to danger signals, the inflammasome triggers pro-inflammatory cytokine secretion in two steps [264]. Priming is the first step of the inflammasome activation, where cellular debris and pathogens, also known as damage- or pathogen associated molecular patterns (DAMP or PAMP), bind to TLR and induce downstream NF- κ B mediated expression of NLRP3 and pro-IL-1 β . The second signal of DAMP or PAMP trigger several possible pathways such as enhanced K⁺ efflux

and lysosomal degradation, which finally leads to inflammasome assembly and conversion of pro-caspase-1 into its active form. The cleavage of pro-IL-1 β and pro-IL-18 by active caspase-1 to IL-1 β and IL-18 prompts the start of the inflammatory cascade.

Accumulated lipids in blood vessels can lead to NLRP3 activation which predisposes to atherosclerosis by impairing endothelial dysfunction and promoting coagulation [265]. The cell debris released from ischemic cardiac injury afterwards act as DAMP which trigger NLRP3/IL-1 β signalling and acute inflammation leading to leucocytes infiltration to the myocardium and triggering further myocardial damage [266]. Tissue healing is initiated when the inflammasome stimulates IL-1 β release in myofibroblasts, promoting collagen accumulation, cardiac fibrosis and remodelling. Moreover, IL-1 β has been considered as cardio-depressant; impairing contractility and inducing LV systolic dysfunction which further worsens HF [267]. It is worth mentioning that the role of NLRP3 inflammasome activation in cardiotoxicity and non-ischemic injuries has also been discussed [268,269]. Thus, it is no surprise that mice deficient of inflammasome components were more resistant to developing atherosclerosis [270], while inflammasome inhibitors contributed to reducing infarct size and improving cardiac function in different *in-vivo* models [271]. Furthermore, deficiency of NLRP3 in mice resulted in lower LV dilation and fibrosis, while it showed better survival compared to wild type [272]. In HF patients, administration of anakinra (IL-1 β inhibitor) improved peak oxygen consumption and exercise capacity [267,273].

Direct effects of SGLT2 inhibitors on NLRP3 inflammasome activation have been established in non-diabetic mice via modulation of intracellular Ca²⁺ levels. Although not commonly mentioned, Ca²⁺ mobilization was considered an important activator of NLRP3 inflammasome starting from the assembly step until IL-1 β release [274]. Recently, it has been shown that extracellular Ca²⁺ can activate the inflammasome and IL-1 β release through the induction of calcium sensing receptor signalling [275]. On this account, empagliflozin improved systolic and diastolic functions in HFpEF mice, while it reduced cardiac fibrosis, mass and remodelling as well as diastolic dysfunction in HFpEF rats [94]. This was attributed to the inhibitory effect of empagliflozin on the priming and activation of the inflammasome and the resulting expression of inflammatory cytokines *in-vivo*, *ex-vivo* and *in-vitro*, mediated by lowering intracellular Ca²⁺ levels. Since the intracellular levels of both Na⁺ and Ca²⁺ are inter-related [276], empagliflozin was able to reduce Ca²⁺ levels in cardiomyocytes of HF mice by the attenuation of late Na⁺ current and its intracellular accumulation [277]. This caused inhibition of NLRP3 inflammasome activation and improved functional recovery. Similar reduction in NLRP3 mediated inflammation was observed with empagliflozin in cardiomyocytes treated with doxorubicin (a dose-dependent cardiotoxic agent), which might be explained, although not explicitly stated, by lowering Ca²⁺ content [278].

4.3. Promoting M2 macrophage polarization

Following tissue injury, innate immune response is initiated where macrophages release inflammatory cytokines that stimulate differentiation and activation of fibroblasts and other cells necessary for tissue repair and wound healing [279]. Macrophages then take another form which suppresses inflammation and ensures normal tissue function and structure. Thus, macrophages have two distinct phenotypes, M1 and M2 [280], where the abundant subtype can be decided by the nature of the surrounding microenvironment [281]. The M1 proinflammatory subtype releases TNF, IL-6 and IL-1 β , while M2 releases anti-

inflammatory cytokines and growth factors such as IL-10, TGF- β and insulin-like growth factor (IGF)-1 [282]. In the heart, aberrant inflammatory response can hinder post myocardial injury repair mechanism and can predispose to cardiac remodelling [283]. Therefore, targeting the direction of macrophage polarization towards the anti-inflammatory type or reducing the levels of pro-inflammatory M1 can be considered as therapeutic strategies to improve healing process and reduce fibrosis, remodelling and risk of HF [284,285].

SGLT2 inhibitors have been shown to enhance macrophage polarization from M1 to M2 in several studies. However, further research is needed to examine this mechanism in the context of CVD and myocardial injury.

LPS- stimulated human and murine macrophages showed increased numbers of M1 subtype and M1/M2 ratio, the effect which was completely reversed upon dapagliflozin treatment in a glucose-independent manner [89,286]. The same results were replicated by canagliflozin in LPS induced mice and macrophages [287]. The reduction of M1 phenotype was associated with attenuation of TNF, IL-6 and IL-1 β release which alleviated lung injury. Against this background, the enhanced M1 to M2 polarization reported in T2D patients might be explained by a direct effect of empagliflozin rather than by its glucose lowering alone [288].

Adipose tissue macrophages (ATM) play a critical role in aggravating inflammation and insulin resistance [289], that could be extended to ectopic fat tissues around blood vessels and heart promoting CVD, as previously discussed. Strategies to promote the anti-inflammatory ATM phenotype have been utilized to improve insulin sensitivity and reduce inflammation in obesity [290–292]. In high fat-fed mice empagliflozin enhanced M2 ATM levels in WAT, while it reduced M1-mediated inflammatory cytokines release together with reduced phosphorylation of JNK and extracellular signal-regulated kinase (ERK)1/2 [208], p38 MAPK and NF- κ B [293]. Additionally, empagliflozin inhibited infiltration of CD3⁺, CD4⁺ and CD8⁺ T cells in WAT, which is considered a critical step in M1 ATM recruitment. Interestingly, modulation of ATM polarization by ipragliflozin contributed to healthy adipose tissue expansion which was attributed to enhanced insulin signalling [294]. Potential inflammatory pathways that could be altered by SGLT2 inhibitors are presented in Figure 2.

Table 1. summarizes the different SGLT2 inhibitors available in the market and the respective evidence, if any, which supports the anti-inflammatory effects of each member.

5. SUMMARY

Growing evidence has shown the importance of SGLT2 inhibitors in attenuating inflammation in several *in-vitro* and *in-vivo* studies, which could possibly explain in part the reduced risk of plaque formation, endothelial dysfunction, cardiac fibrosis and ventricular remodelling. However, further clinical trials are needed to determine whether these anti-inflammatory effects are associated with the reduction of HF hospitalization and CV mortality observed with SGLT2 inhibitors in recent RCTs.

In fact, most of the evidence available relates these anti-inflammatory effects to the systemic and metabolic improvements of SGLT2 inhibitors, which were, to a great extent, dependent on their glucosuric effects. Although SGLT2 inhibitors are mainly investigated in the setting of T2D, it is becoming increasingly evident that the CV benefits observed are not exclusively related to the associated glycaemic control and reduced glucotoxicity. Furthermore, lowering of uric acid by

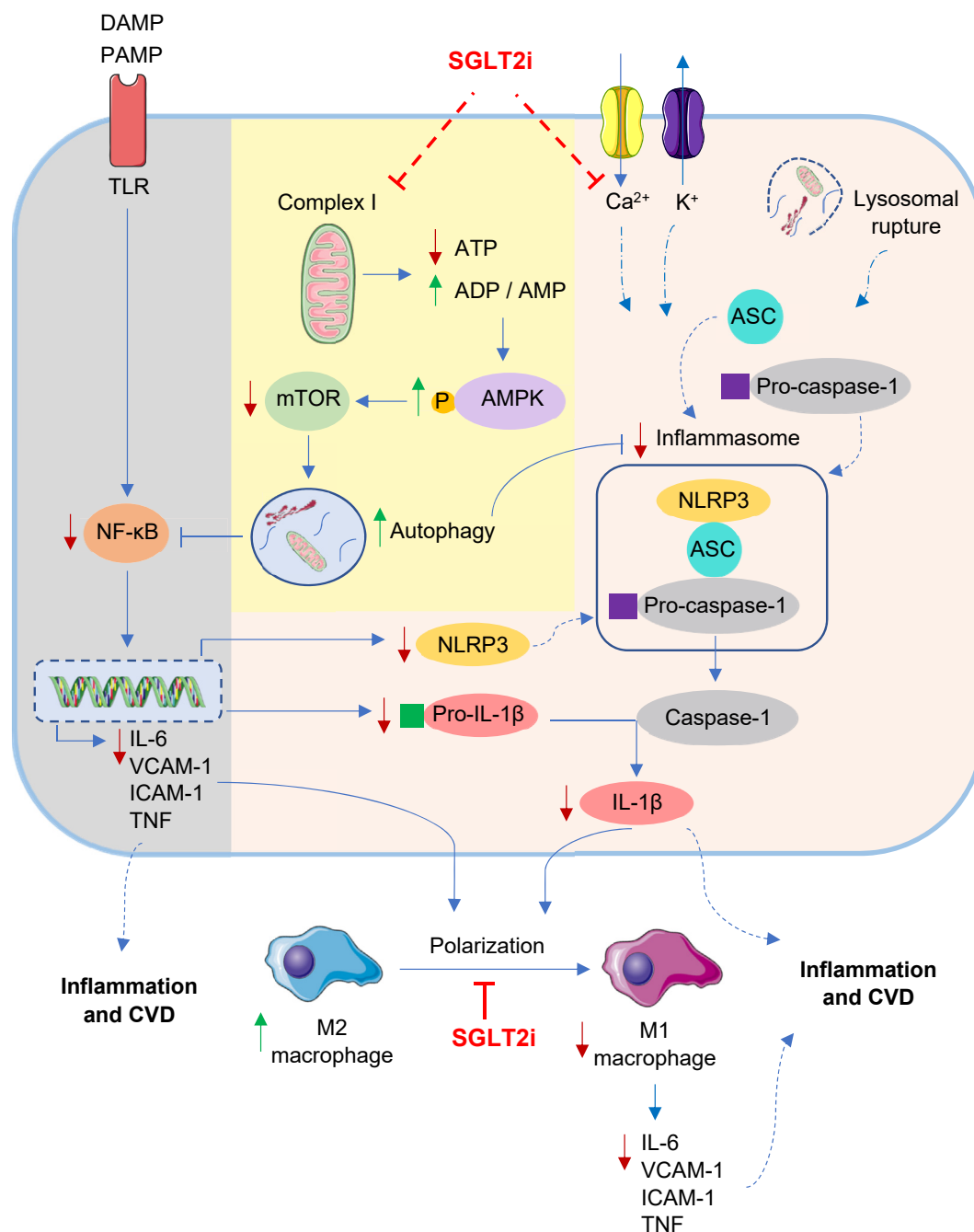


Figure 2: Inflammatory signalling pathways as potential targets of SGLT2 inhibitors. SGLT2 inhibitors suppress the mitochondrial complex I eventually predisposing to the induction of AMPK/mTOR/autophagy signalling. Autophagy directly inhibits inflammasome activation, while it also attenuates NF- κ B mediated release of several inflammatory cytokines together with NLRP3, an essential component of the inflammasome. Increased intracellular Ca^{2+} , enhanced K^{+} efflux and lysosomal degradation can signal inflammasome assembly and activation. On that note, SGLT2 inhibitors reduce intracellular Ca^{2+} levels which can contribute to the inhibition of inflammasome activation and subsequent attenuation of IL-1 β release. The overall reduction of inflammatory cytokines enhances M2 macrophage production and reduce their polarization to the pro-inflammatory M1 subtype, a process which was also directly modulated by SGLT2 inhibitors. These direct actions on inflammation can explain the CV benefits of SGLT2 inhibitors. Blue arrows, represent the normal signalling pathway; red arrows, represent decreased expression upon SGLT2 inhibitor treatment; green arrows, represent enhanced expression. ADP, adenosine diphosphate; AMP, adenosine monophosphate; AMPK, AMP-activated protein kinase; ASC, apoptosis-associated speck-like protein; ATP, adenosine triphosphate; CVD, cardiovascular disease; DAMP, damage-associated molecular patterns; ICAM-1, intercellular adhesion molecule-1; IL-1 β , interleukin-1 β ; IL-6, interleukin-6; mTOR, mammalian target of rapamycin; NF- κ B, nuclear factor- κ B; NLRP3, nucleotide-binding domain, leucine-rich-containing family, pyrin domain-containing-3; PAMP, pathogen associated molecular patterns; SGLT2i, sodium glucose co-transporter 2 inhibitors; TNF, tumour necrosis factor; VCAM-1, vascular cell adhesion molecule-1.

Table 1 — A summary of the currently available SGLT2 inhibitors and the respective evidence, if any, of their anti-inflammatory mechanism.

Member	Empa [295]	Dapa [296]	Cana [297]	Ertu [298]	Sota [299]	Ipra [300]	Luseo [301]	Tofo [302]
	C ₂₃ H ₂₇ ClO ₇	C ₂₁ H ₂₅ ClO ₆	C ₂₄ H ₂₅ FO ₅ S	C ₂₂ H ₂₅ ClO ₇	C ₂₁ H ₂₅ ClO ₅ S	C ₂₁ H ₂₁ FO ₅ S	C ₂₃ H ₃₀ O ₆ S	C ₂₂ H ₂₆ O ₆
Approval	FDA [303] EMA [304]	FDA [305] EMA [306]	FDA [307] EMA [308]	FDA [309] EMA [310]	EMA for type 1 diabetes [311]	Only in Japan, South Korea and Russia [312]	Only in Japan [313]	Only in Japan [314]
Dosing regimen	10/25 mg once daily	5/10 mg once daily	100/300 mg once daily	5/15 mg once daily	200 mg once or twice daily	50/100 mg dose once daily	2.5/5 mg once daily	20 mg once daily
Anti-inflammatory mechanism								
Reduce ROS and hyperglycaemia mediated inflammation	[50,71,84–87]	[67,68]						[88]
Reduce adipose tissue volume and associated adipokines	[199]	[195,204]				[206]	[196,205]	
Reduce AngII mediated inflammatory mediators/macrophage infiltration/fibrosis	[92,229]	[93,96]	[228]		[229]			
Enhance ketone mediated NLRP3 inflammasome inactivation	[166,167]							
Enhance M2 macrophage polarization	[208,288,293]	[89,286]	[287]			[294]		
Enhance AMPK mediated reduction of inflammatory mediators	[51,249,261]	[90,246,247]	[91,248]					
Inactivation of NLRP3 inflammasome	[94,277,278]							

References of the anti-inflammatory mechanisms are colour coded in blue for animal models, red for in-vitro and ex-vivo studies, green for human data and purple for studies with combined models.

AngII, angiotensin II; AMPK, 5' adenosine monophosphate activated protein kinase; Cana, canagliflozin; Dapa, dapagliflozin; Ertu, ertugliflozin; EMA, European Medicines Agency; Empa, empagliflozin; FDA, Food and Drug Administration; Ipra, ipragliflozin; Luseo, luseogliflozin; NLRP3, nucleotide-binding domain, leucine-rich-containing family, pyrin domain-containing-3; ROS, reactive oxygen species; SGLT2, sodium glucose co-transporter 2; Sota, sotagliflozin; Tofo, tofogliflozin.

SGLT2 inhibitors is probably not cardioprotective per se, but it is rather a reflection of oxidative stress mitigation via promoting the antioxidant SIRT1 expression or through direct inhibition of the main ROS generating mechanisms in the heart. Oxidative stress and inflammation could also be mediated by RAS activation, which was reversed upon SGLT2 inhibitor administration. Reduction of ectopic fat deposition by SGLT2 inhibitors is associated with reduced proinflammatory cytokine and adipokine release, which directly influence cardiac and vascular remodelling. Rather being a super fuel, a hypothesis which is still under debate, βOHB has been established as a specific HDAC and NLRP3 inflammasome inhibitor, which may explain the antioxidant and anti-inflammatory effects of ketonemia associated with this drug class. Although based on limited number of studies to date, the reported CV benefits could partially be a consequence of direct influence on proinflammatory signalling pathways. SGLT2 inhibitors contributed to enhanced AMPK phosphorylation which was associated with downstream inhibition of inflammatory mediators. Activation of autophagy mediated by AMPK/mTOR signalling could explain these effects. Apart from AMPK activation, SGLT2 inhibitors contributed to inhibition of NLRP3 inflammasome priming and assembly, possibly via modulation of Ca²⁺ signalling. Enhancing macrophage polarization to the M2 subtype could attribute to SGLT2 inhibitor anti-inflammatory actions, although further studies are needed to establish these findings in cardiac tissues.

In summary, among other mechanisms cardioprotective effects and major improvement of HF outcomes observed with SGLT2 inhibitors are possibly related to both direct and indirect mitigation of inflammatory signalling pathways independent of glycaemic control.

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CONFLICT OF INTEREST

None declared

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