

**Role of Tumour Necrosis Factor Alpha (TNF α) in
Ischaemic and Pharmacological Postconditioning**

Lydia Lacerda

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Supervisor: Professor S. Lecour, PharmD, PhD

(Hatter Cardiovascular Research Institute, Faculty of Health Science, University of Cape Town)

Co-supervisor: Professor L. H. Opie, PhD; DSc; DPhil

(Hatter Cardiovascular Research Institute, Faculty of Health Science, University of Cape Town)

Declaration

I, Lydia Lacerda, hereby declare that the work on which this thesis is based is my original work (except where acknowledgement indicates otherwise), and that neither the whole work nor any part of it has been, is being, or is submitted for another degree in this or any other university.

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“It is through science that we prove, but through intuition that we discover”

Henri Poincare

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Abbreviations

α	alpha
β	beta
δ	delta
γ	gamma
ADAM	a disintegrin and metalloprotease
ADP	adenosine diphosphate
AMI	acute myocardial infarction
ANOVA	analysis of variance
ATP	adenosine triphosphate
Bad	Bcl-2 associated death agonist
Bcl-2	B-cell lymphoma-2
Bcl-x1	B-cell lymphoma-x1
CaCl ₂	calcium chloride
COX-2	cyclooxygenase-2
cGMP	cyclic guanosine monophosphate
cIAP2	cellular inhibitor of apoptosis protein 2
CK	creatine kinase
CVD	cardiovascular disease
DADLE	D-Ala ² , D-Leu ⁵ -enkephalin

DD	death domain
DED	death effector domain
EDTA	ethylene diamine tetra-acetic acid
ERK	extracellular signal-regulated kinases
ERK1/2	extracellular signal-regulated kinase 1/2
eNOS	endothelial nitric oxide synthase
FADD	fas-associated death domain
FAN	factor associated with neutral sphingomyelinase activation
FFA	free fatty acids
GAPDH	glyceraldehyde-3-phosphate dehydrogenase
GDP	guanosine diphosphate
Gi	G inhibitory
GIK	glucose, insulin, potassium
GSK-3 β	glycogen synthase kinase-3 beta
GTT	glucose tolerance test
H ₂ O ₂	hydrogen peroxide
HCD	high carbohydrate diet
HDL	high density lipoprotein
HMGB-1	high-mobility group box-1

IL-1	interleukin-1
IL-6	interleukin-6
ip	intraperitoneal
IPC	ischaemic preconditioning
IPostC	ischaemic postconditioning
I/R	ischaemia/reperfusion
JAK	janus kinase
KCl	potassium chloride
KH_2PO_4	potassium dihydrogen phosphate
LDL	low density lipoprotein
MAPK	mitogen activated protein kinase
MEK	mitogen activated protein kinase extracellular regulated kinase
MEKK	mitogen activated protein kinase extracellular regulated kinase kinase
mK_{ATP}	mitochondrial potassium-sensitive adenosine triphosphate channel
MPG	mercaptopyrionylglycine
mPTP	mitochondrial transition pore
NaCl	sodium chloride
NADPH	nicotinamide adenine dinucleotide phosphate
NaHCO_3	sodium bicarbonate

NaH ₂ PO ₄	sodium dihydrogen phosphate
ND	normal diet
NFκB	nuclear factor kappa B
NIDDM	non-insulin dependent diabetes mellitus
NO	nitric oxide
NOS	nitric oxide synthase
oxLDL	oxidised low density lipoprotein
PAR-2	proteinase activated receptor-2
PCI	percutaneous coronary intervention
PCTA	percutaneous transluminal coronary angioplasty
PI3-Kinase	phosphatidylinositol 3-kinase
PKB/Akt	protein kinase B
PKC	protein kinase C
p70s6K	p70-ribosomal s6 kinase
PKG	protein kinase G
RAGE	receptor for advanced glycation end products
RIP	receptor interacting protein
RIPC	remote ischaemic preconditioning
RISK	reperfusion injury salvage kinases

ROS	reactive oxygen species
sarcK _{ATP}	sarcolemmal potassium-sensitive adenosine triphosphate channel
SAFE	survivor activating factor enhancement
SODD	silencer of death domain
STAT	signal transducer and activator of transcription
STZ	Streptozotocin
TACE	TNF α converting enzyme
TLR	Toll-like receptors
TNF α	tumour necrosis factor alpha
TNFR1	tumour necrosis factor alpha receptor 1
TNFR2	tumour necrosis factor alpha receptor 2
TRADD	TNF receptor associated death domain
TRAF	TNF receptor associated factor
TTC	2,3,5-triphenyltetrazolium chloride
VEGF	vascular endothelial growth factor
VDAC	voltage dependent anion channel

Abstract

Aims: Ischaemic postconditioning (IPostC) is a powerful protective mechanism that activates prosurvival intrinsic signalling cascades to limit reperfusion injury but the exact signalling pathway involved in this cardioprotective effect still remains unclear.

In ischaemic preconditioning, a powerful prosurvival path, known as the Survivor Activating Factor Enhancement (SAFE) pathway, has recently been discovered to protect against reperfusion injury. The SAFE pathway involves the activation of the cytokine, tumour necrosis factor alpha (TNF α) and the signal transducer and activator of transcription-3 (STAT-3) during ischaemic preconditioning. The aim of this thesis was to investigate whether ischaemic and pharmacological postconditioning can protect via activation of the SAFE pathway. In addition, we explored the influence of obesity and diabetes in the successful effect of ischaemic postconditioning, as these two pathologies are often associated with patients with myocardial infarction.

Methods: Isolated Langendorff perfused mouse hearts from TNF knockout (TNF $^{-/-}$), TNF receptor 1 or receptor 2 knockouts (TNFR1 $^{-/-}$ or TNFR2 $^{-/-}$), and cardiomyocyte-specific STAT-3 knockouts (STAT-3 $^{-/-}$) and their respective wild types were subjected to a global ischaemia and were postconditioned with either short periods of ischaemia, TNF α (0.5 ng/ml), adenosine (100 μ M), bradykinin (100 nM) or the specific delta opioid agonist, D-Ala2', D-Leu5'-enkephalin acetate salt (DADLE) (100 nM). Infarct size was measured using triphenyltetrazolium chloride staining technique and computerized planimetry. Western blot analysis of nuclear, cytosolic or mitochondrial proteins was performed to evaluate activation of STAT-3, Akt and extracellular signal-regulated kinase 1/2 (Erk1/2), during reperfusion.

To explore the influence of obesity and diabetes on IPostC-induced cardioprotection, TNF-WT and TNF $^{-/-}$ animals were fed with a high carbohydrate diet for 11 weeks to induce obesity. Diabetes was induced by administration of a single intraperitoneal injection of streptozotocin (180mg/kg).

Results: IPostC or pharmacological postconditioning with bradykinin or DADLE reduced infarct size in TNF-WT, TNFR1 $^{-/-}$ and STAT-3-WT mice, but not in TNF $^{-/-}$, TNFR2 $^{-/-}$ and STAT-3 $^{-/-}$ mice. Adenosine reduced infarct size in TNF-WT, TNF $^{-/-}$, and STAT-3-WT but failed to protect STAT-3 $^{-/-}$ mice. TNF α -postconditioning mimicked IPostC in TNF-WT and STAT-3-

WT but not in $TNF^{-/-}$ or $STAT-3^{-/-}$ hearts. Both IPostC and TNF-PostC increased phosphorylated STAT-3 in the nucleus, while Akt and Erk1/2 were only activated after an IPostC stimulus. Similarly, bradykinin and DADLE increased phosphorylated STAT-3 in the mitochondria.

IPostC protected hearts against ischaemia-reperfusion in obese or diabetic WT mice. Surprisingly, the protective effect of IPostC was partially restored in $TNF^{-/-}$ obese mice but the infarct-sparing effect was absent in diabetic $TNF^{-/-}$ mice.

Conclusion: In this thesis we have delineated a novel prosurvival pathway activated in IPostC to limit ischaemia-reperfusion injury. This pathway, known as the SAFE pathway, requires activation of $TNF\alpha$ and activates STAT-3 via TNFR2. Pharmacological postconditioning with bradykinin and opioids mimics IPostC via the SAFE pathway. Although adenosine does not require $TNF\alpha$ in pharmacological postconditioning, STAT-3 is still required, thus confirming the existence of multiple protective pathways in the heart to protect against reperfusion injury. In our experimental conditions, obesity and diabetes do not influence the outcome of ischaemic postconditioning but $TNF\alpha$ appears to play a deleterious role in obesity.

The existence of multiple alternate pathways in IPostC and pharmacological postconditioning at the time of reperfusion may have potential therapeutic application against ischaemia-reperfusion injury. Furthermore, we show that the protective effect of ischaemic postconditioning is maintained in a modestly obese and diabetic mouse model suggesting that, relative to the severity of the disorder, obese and diabetic patients should be considered for this therapy.

A. INTRODUCTION

Transporting blood throughout the body to each organ and tissue, throughout our lives, requires an efficient and healthy heart. As we grow older, the heart becomes progressively less efficient in maintaining a healthy circulatory system. However, even as our age increases, the heart, as is the case with any muscle, benefits from a healthy lifestyle, coupled with regular age-appropriate exercise. An individual who lives such a lifestyle could expect to live to be a “ripe old age”. Unfortunately, with the modern world’s fast pace of living and working, people are exposed to a number of risk factors which promote cardiovascular diseases resulting in a shortened lifespan as well as being a burden on the social system of a country. Risk factors such as hypertension, diabetes, elevated cholesterol levels, just to mention a few, all contribute eventually to a narrowing of the arteries, compromising the circulatory system. The flow of blood and therefore, the oxygen supply and essential nutrients, to the muscle cells of the heart, is decreased resulting in a “heart attack” or myocardial infarction.

The heart is a resilient organ and possesses intrinsic survival mechanisms in an attempt to protect itself. Recently, the heart has been shown to possess a remarkable ability to adapt to conditions of stress and this has been one of the focal points of intense research over the past 15 years in the hope that the underlying mechanisms may be harnessed for therapeutic development.

In this thesis we propose to explore the activation of a major intrinsic cytokine, tumour necrosis factor alpha (TNF α), as a possible tool to protect the heart against stressful conditions observed during myocardial infarction. This study will be performed in healthy animals or in obese/diabetic animals, two pathologies often triggering myocardial infarction.

1. Ischaemic Heart Disease

Ischaemic heart disease is characterized by a reduced blood supply (ischaemia) to the heart muscle, usually due to coronary artery disease (atherosclerosis of the coronary arteries).

In the adult human body there are 96,600 kilometers of arteries, veins and capillaries pulsating at 100,000 rhythmic, muscular heart beats per day. These arteries, veins and capillaries, together with the heart, make up the cardiovascular system. Ischaemic heart disease affects individuals in their prime of life, reducing their life expectancy and their contribution towards the economy of a country substantially and is thus perceived as a growing global burden. Ischaemic events can occur as a consequence of either a thrombosis (blood clot) or narrowing and hardening of the arteries as a result of plaque accumulation. Atherosclerotic plaques within coronary arteries cause rapid occlusion in coronary artery disease resulting in the heart suffering a loss of blood flow and oxygen supply with subsequent death of the cardiomyocytes, the muscle cells of the heart. During occlusion of

the artery or arteries supplying the myocardium, the demand for oxygen may exceed the supply and myocardial ischaemia develops. Rupture of a plaque may result in a secondary thrombus formation in a coronary artery. If left untreated, this form of coronary artery disease culminates in an acute reduction of blood supply to the myocardium and is characterized by the rapid development of myocardial necrosis (Jennings, 1991).

The number one cause of death worldwide is cardiovascular disease (CVD), with more people dying annually from coronary artery disease than from any other cause (WHO, 2009). Low and middle income countries seem to be more affected than developed countries with 82% of cardiovascular related deaths occurring in the lesser developed countries and almost equally in men and women. By 2030, it is predicted that almost 23.6 million people will die from coronary artery disease – mainly from ischaemic heart disease and stroke (WHO, 2009) and the leading cause of death in the industrialized world is already ischaemic heart disease. Ischaemic heart disease is projected to become the major worldwide cause of death and disability by 2020 – only a decade away!! (WHO, 1996).

“As long as Africa remains impoverished, a major rise in coronary artery disease is unlikely” – this was the general perception about ischaemic heart disease in Africa in 1997 as expressed by Walker et al (Walker, 1998). However, in 2006, the World Bank reported that Africa was making economic progress. Unfortunately, along with economic improvements, there is a rise in chronic diseases of lifestyle such as ischaemic heart disease, a consequence of epidemiological transition. Ischaemic heart disease is rapidly becoming a leading cause of death in sub-Saharan Africa and in 2008 it was ranked as the 8th leading cause of death in men and women of this region (Menash, 2008). The progression of ischaemic heart disease in the ranking is attributed largely to lifestyle and behavioural changes associated with urbanization of the region (Menash, 2008). Ischaemic heart disease is responsible for the greatest number of cardiovascular fatalities in males and the second largest number of cardiovascular related deaths in females, in South Africa, (Figure 1). More alarmingly, a report issued by the South African Medical Research Council in 2005, states that ischaemic heart disease was the leading cause of death in the total population of the Western Cape (one of the provinces of South Africa) in a single year (Bradshaw, 2005).

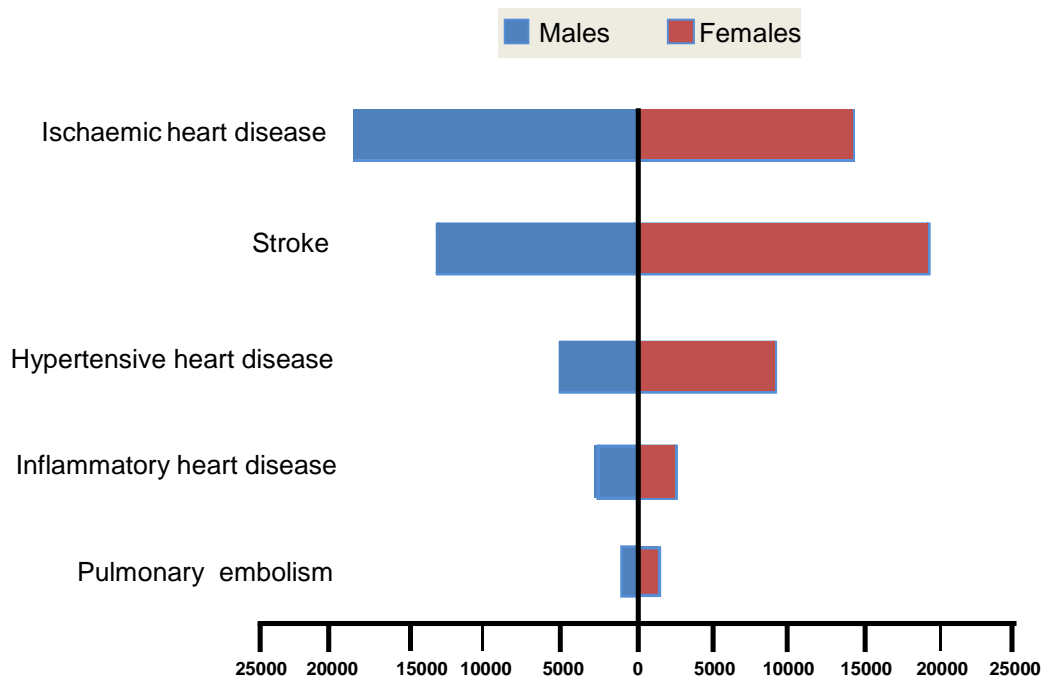


Fig 1: Ischaemic heart disease. Estimated to be the greatest cause of CVD mortality in 2000. (adapted from Norman, 2000)

1.1 Major risk factors for ischaemic heart disease

The major cardiovascular risk factors are as follows:

1.1.1 Atherosclerosis and dyslipidaemia

One of the main causes of coronary artery disease is atherosclerosis (Hansson, 2005). Atherosclerosis is an inflammatory disease in which the inflammatory system interacts with metabolic risk factors, triggering atherosclerotic plaques in the arterial tree. Myocardial infarction occurs when the atheromatous process prevents blood flow through the coronary artery and disruption of the plaque is considered dangerous as it releases harmful thrombotic material from the centre of the plaque into the circulation (Hansson, 2005). Dyslipidaemia is one of the causes of atherosclerosis, with low density lipoprotein cholesterol (LDL) contributing to the formation of plaques. Endothelial disruption and erosion also plays a role in plaque formation, contributing to ischaemic heart disease, (Figure 2).

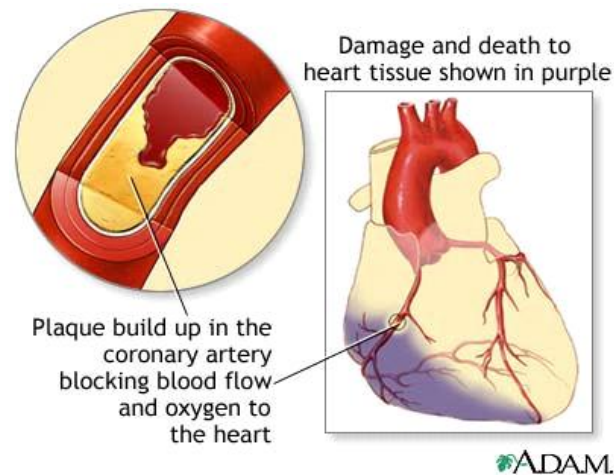


Fig 2: Plaque build up in coronary artery. Loss of blood flow and oxygen supply will eventually lead to morphological changes in the heart and ischaemic heart disease. (www.whatisheartdisease.us/images)

The cholesterol that accumulates in atheromatous lesions is derived primarily from plasma low density lipoprotein (LDL), which accounts for 60-75% of plasma cholesterol, giving LDL a powerful and direct relationship with coronary heart disease (Nestruck and Davignon, 1986). Oxidative stress and smoking are just two of the many factors which can contribute to damage of the endothelium. Consequently, excess production of oxidised LDL-cholesterol (oxLDL) occurs in the bloodstream. In an attempt to reduce the amount of oxLDL cholesterol to normal levels, macrophages engulf the oxLDL and in doing so become “foam cells”, (Figure 3). However, if there is too much oxLDL-cholesterol present, the macrophages accumulate against the vessel wall in what will become an atherosclerotic plaque. Eventually this wall of plaque may either block the vessel completely or it may rupture, resulting in the release of pro-coagulant factors and the formation of a thrombosis (Libby, 2002). Both of these events inhibit the flow of blood to the heart and result in myocardial infarction and ischaemic heart disease, (Figure 4).

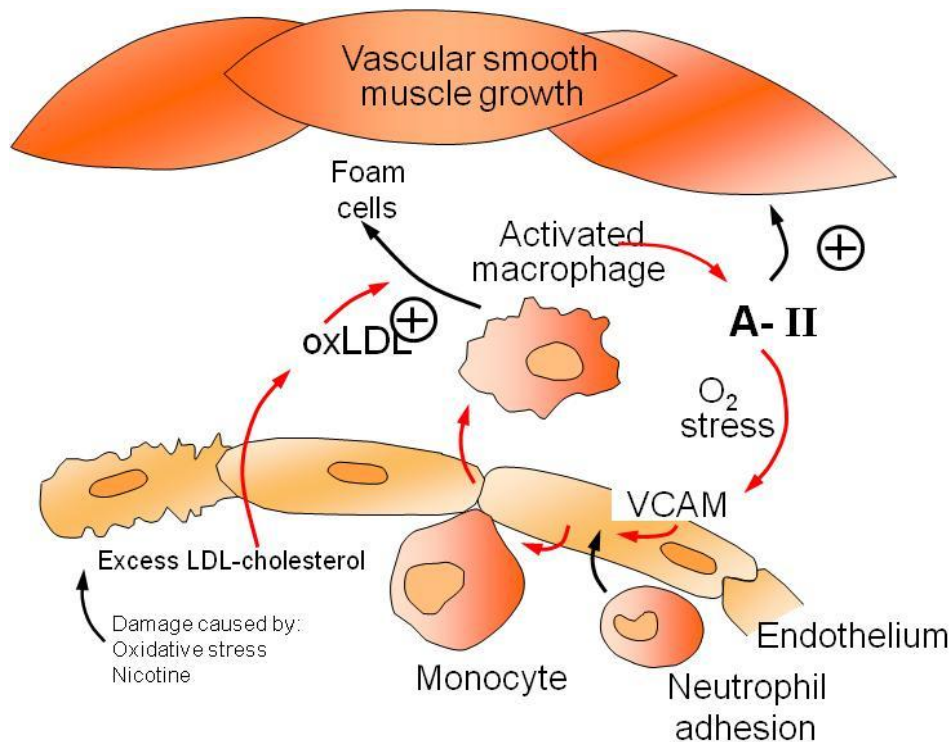


Fig 3: Formation of atherosclerotic plaque. Damaged endothelium favours the infiltration and oxidation of excess LDL-cholesterol. The oxidised LDL-cholesterol (ox-LDL) is taken up by the macrophages in an attempt to keep plasma cholesterol homeostasis. However, in the presence of excess ox-LDL the macrophages accumulate against the vessel wall and form an atherosclerotic plaque. A-II: angiotensin II; VCAM: vascular cell adhesion molecule, (adapted from Opie, 2004).

High density lipoprotein (HDL) cholesterol, also known as the ‘good’ cholesterol, helps to clear excess LDL-cholesterol from the blood back to the liver (a process known as “reverse cholesterol transport”, thereby reducing the risk of atherosclerosis. Many studies have shown that there is an increased risk of ischaemic heart disease in individuals with low levels of HDL (Assmann et al., 1996; Curb et al., 2004; Gordon et al., 1989; Gordon et al., 1977; Sharrett et al., 2001; Turner et al., 1998). However, there is also a direct relationship between increased HDL-cholesterol and decreasing cardiovascular risk with epidemiological data indicating that for every increase of 0.03 mM/L of HDL-cholesterol, the risk for cardiovascular events decreases by 2-3% (Gordon et al., 1989).

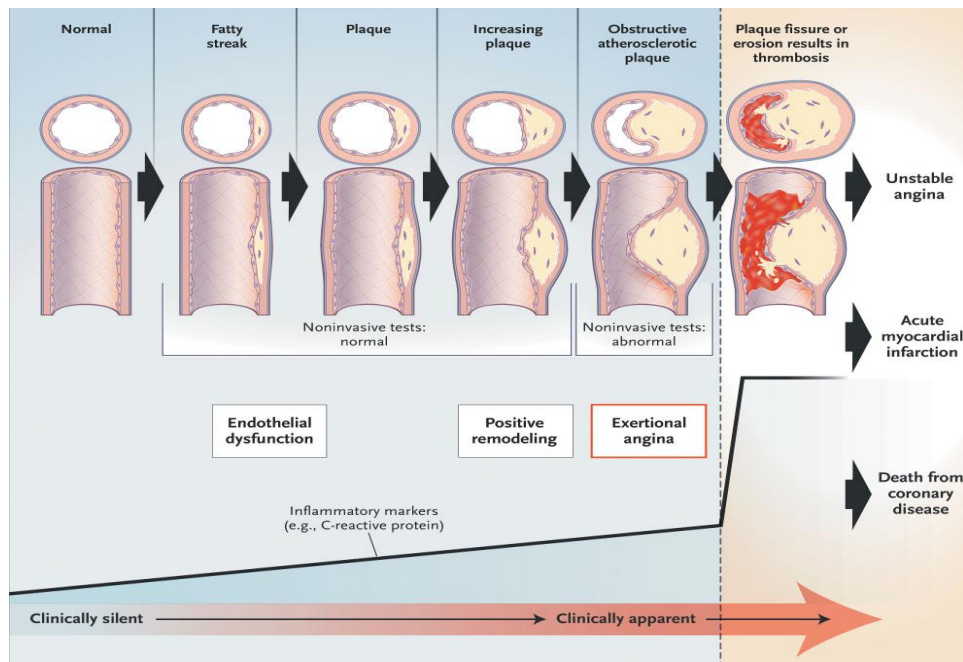


Fig 4: Typical progression of coronary atherosclerosis. At first, the atherosclerotic plaque tends to stay external to the lumen, which allows the coronary artery diameter to be maintained. As the plaque encroaches into the lumen, the diameter of the artery decreases until eventually the flow of blood is inhibited. Luminal narrowing of more than 65 to 75% may result in angina and myocardial infarction, (Abrams, 2005).

1.1.2 Obesity

Obesity has been defined by the World Health Organisation as a disease in which excess body fat has accumulated to such an extent that health may be adversely affected (WHO, 2000).

An imbalance between energy intake and expenditure results in excess energy being stored in fat cells which enlarge or increase in number, and this may eventually result in overweight and obesity. A healthy lifestyle does not necessarily mean a healthy heart if one carries excess weight, especially around the abdomen. Obesity is increasing, worldwide, at an alarming rate, and just 5 years ago, there was an estimated 1.3 billion people overweight or obese (WHO, 2005). Data from 14 years ago already revealed that this global trend towards overweight and obesity was being followed by South Africa with an disturbing number of citizens becoming overweight (South Africa Demographic Health Survey, 1996). Increasing rates of childhood and adolescent obesity are also a cause of great concern as once children have become overweight or obese, interventions to reverse these conditions are rarely successful (Pinhas-Hamiel and Zeitler, 2000). Current definitions designate overweight in adults as those having a body mass index

between 25.0 and 30.0 kg/m² and obesity as a BMI of greater than 30.0 kg/m² (Poirier et al., 2006). Distribution of body fat, or adipose tissue, appears to play a significant role in determining the presence or absence of metabolic dysfunctions that predispose to chronic diseases (Poirier, 2003). The skinfold test is another means of defining obesity by determining a person's body composition and body fat percentage. Metabolic syndrome, which consists of a cluster of diabetogenic and atherogenic abnormalities, correlates closely with CVDs and diabetes. Much has been written about the metabolic syndrome with some cardiologists favouring its existence and others questioning it. It appears that of all the parameters suggested to define metabolic syndrome, the one of most importance is that of the waist-circumference measurement (Alberti et al., 2005; Opie, 2006). A common finding in the obese state is a low-grade systemic inflammation with increased circulating levels of a number of cytokines, amongst which is tumour necrosis factor alpha (TNF α) (Bullo et al., 2003; Roytblat et al., 2000; Yudkin et al., 1999). Anti-inflammatory therapies targeting TNF α activity have indicated that blocking TNF α may interrupt the inflammatory cascade that occurs with abdominal obesity (Bernstein et al., 2006).

A large number of clinical problems are associated with being obese and these can be subdivided into two groups: those that are associated with excess adipose tissue and those that are linked to the metabolic effects of the increased adiposity (Bray, 2004). The diseases associated with the latter group include coronary heart disease, hypertension, type 2 diabetes mellitus and certain kinds of cancer (Garfinkel, 1985; Jooste et al., 1988; Lew, 1985; Manson et al., 1995; Pi-Sunyer, 1990; Pi-Sunyer, 1991; Steyn et al., 1990). Accumulation of abdominal fat increases the probability of cardiovascular disease and mortality (Donahue et al., 1987; Ducimetiere et al., 1986; Lapidus et al., 1984; Larsson et al., 1984). The metabolic effects of obesity result from the increased release of free fatty acids (FFA) and the production of adipokines which are secreted by several factors, including the cytokines TNF α and interleukin-6 (IL-6), hormones such as leptin and adiponectin and other adipokines such as resistin and angiotensinogen. These factors can impair insulin signalling and alter glucose homeostasis as well as lipid metabolism (Arner, 2003; Nieto-Vazquez et al., 2008). Obesity is therefore, now being considered as a chronic state of low-intensity inflammation, with one of the major health consequences being the increased risk of ischaemic heart disease (Larsson et al., 1981).

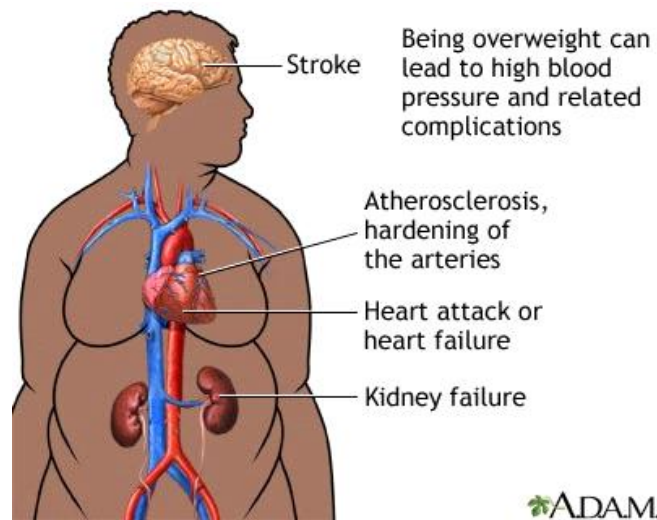


Fig 5: Cartoon illustrating possible complications arising from obesity. (www.clarion.org)

1.1.3 Diabetes

Type 1 diabetes: *the pancreatic beta cells of the body fail to produce insulin as the body's immune system has attacked and destroyed them. At present, this type of diabetes requires the afflicted person to inject insulin.*

Type 2 diabetes: *results from insulin resistance, a condition in which cells fail to use insulin properly and is sometimes combined with an absolute insulin deficiency. At first, the pancreas produces more insulin to compensate, but eventually it loses the ability to secrete insulin in response to meals. Being overweight and lack of exercise increase the risk of developing this type of diabetes.*

Diabetes and its associated complications have become a worldwide health problem. The prevalence of diabetes is rising, reaching epidemic proportions in some population groups. Type 2 diabetes is responsible for much of the increased occurrence of diabetes but there is also evidence that the incidence of type 1 diabetes is on the increase, although to a lesser extent than type 2 diabetes (Podar et al., 2001). In addition, impaired glucose tolerance is on the increase with about 197 million individuals worldwide having this disorder, predominantly as a result of obesity and the associated metabolic syndrome (Hossain, 2007). Abnormalities in both the action and secretion of insulin are responsible for the pathogenesis of diabetes. Insulin resistance is defined as the diminished ability of the cell to respond to the action of insulin with a subsequent

increase in lipolysis, release of glycerol and free fatty acids (FFA) into the circulation, ultimately leading to diabetes, (Figure 6).

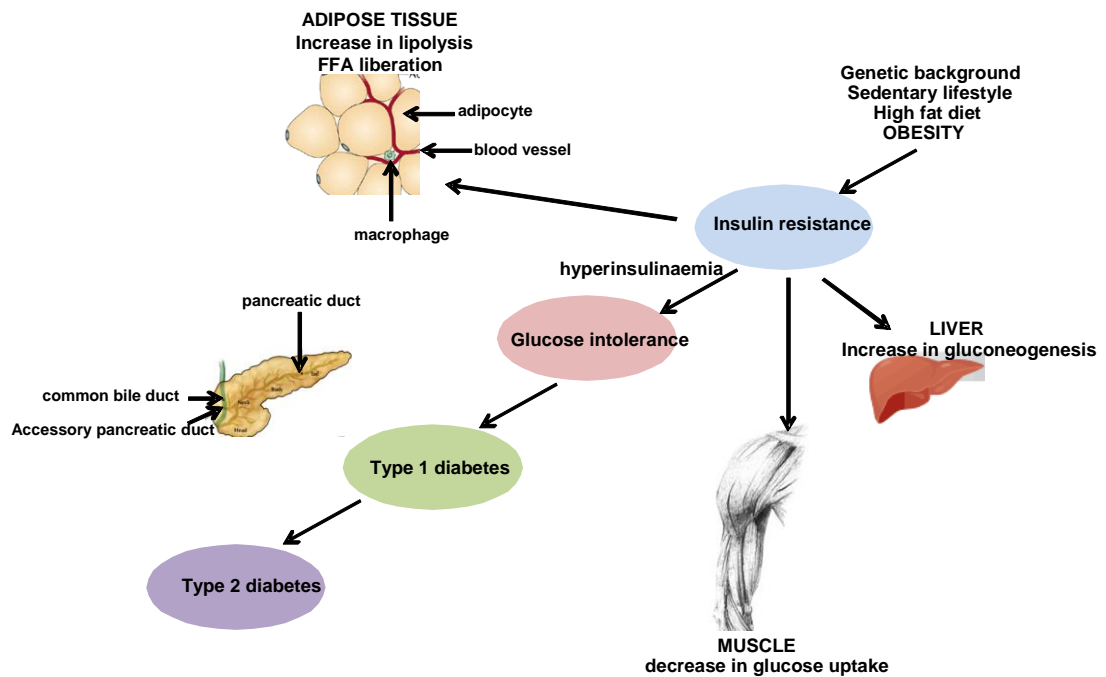


Fig 6: Schematic representing the different steps in the development of type 1 and type 2 diabetes, (adapted from Nieto-Vasquez, 2008).

The development of insulin resistance has been linked to both genetic and environmental factors, with obesity being an important contributor. Most studies thus far have focussed on type 2 diabetes. However, type 1 diabetes also has serious cardiovascular consequences which occur in a much younger age group (Orchard et al., 2006). The powerful effect of the duration of type 1 diabetes should be taken into consideration when regarding cardiovascular risk associated with this type of diabetes. In the intervening years between diagnosis and the patient's 3rd decade, 20 years or more have passed and a considerable amount of damage to the cardiovascular system may have occurred. The development of type 1 diabetes is shown in Figure 7:

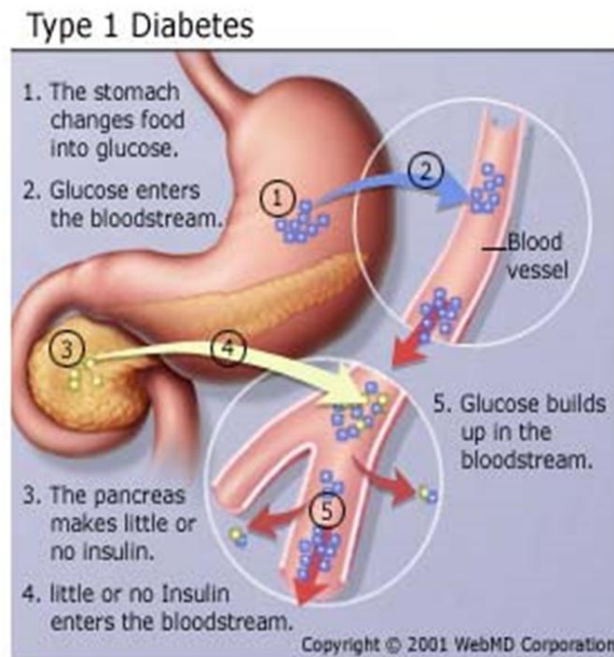


Fig 7: Development of type 1 diabetes: As production of insulin by the pancreatic cells is impaired, the level of glucose in the blood increases, leading to type 1 diabetes. (www.medicinenet.com)

Both type 1 and type 2 diabetic individuals are prone to developing ischaemic heart disease, including acute myocardial infarction and subsequent complications (Aguilar et al., 2004; Ramani et al., 1996; Stevens et al., 2004; Zairis et al., 2004). More than 50% of deaths in diabetic patients have been attributed to ischaemic heart disease (Kannel and McGee, 1979) and the mortality rate from acute myocardial infarction is almost doubled in diabetic individuals compared to non-diabetics (Abbott et al., 1988). Although it is well established that the long-term outcome of diabetic patients with ischaemic heart disease is worsened, studies have been contradictory as to whether the diabetic heart responds well or not to treatment in the event of an ischaemic episode. For instance, hearts from streptozotocin-induced diabetic rats were more resistant initially to ischaemia/reperfusion, but over a period of 8 weeks, this effect was reversed (Tosaki et al., 1996). This initial positive response to ischaemia-reperfusion was also seen in a type 2 diabetes model with an *in vivo* study (Liu et al., 1993). Indeed, in an alloxan-induced diabetic rabbit model, a decrease in infarct size was observed 8 weeks after the alloxan injection, compared to controls (Hadour et al., 1998).

The majority of clinical studies have however, demonstrated a worse outcome from acute myocardial infarction in patients who are already compromised by diabetes (Osario, 2010).

1.1.4 High blood pressure (hypertension)

Hypertension has been defined as when the arterial blood pressure rises above 140 mmHg for systolic pressure and 90 mmHg for diastolic pressure (Kaplan et al., 2010).

Elevated arterial blood pressure is the most common cause of enlargement of the left ventricle of the heart (Diamond et al., 2005; Ferdinandy, 2007; Massie and Shah, 1997; Prisant, 2005). Experimental evidence suggests that the hypertrophied myocardium is more susceptible to ischaemia-reperfusion damage than a normotensive heart (Attarian et al., 1981). However, no difference in infarct size between hypertensive and normotensive animals was demonstrated either *in vivo* in a rat heart infarct model (Ebrahim et al., 2007) or *in vitro* in a rat model of coronary occlusion (Speechly-Dick et al., 1994).

2.0 Reperfusion and reperfusion injury

Reperfusion is defined as the restoration of blood flow to an organ or tissue that has had its blood supply cut off. Reperfusion injury refers to the damage done to the organ or tissue caused when blood supply returns to the tissue after a period of ischaemia.

In the event of an acute myocardial infarction, rapid restoration of the blood flow to the ischaemic area is one of the best ways to salvage the heart. Provided that reperfusion occurs shortly after coronary artery occlusion, it is effective in reducing the infarct size thus limiting the damage inflicted by the myocardial infarct (Maroko et al., 1971). Despite the undoubted benefit of reperfusion of blood to the ischaemic tissue, reperfusion itself is not risk free. A cascade of adverse reactions can occur during reperfusion, (see review (Yellon and Downey, 2003)) which paradoxically results in an exacerbation of tissue injury known as “reperfusion injury”.

Restoration of the blood flow in the clinical setting involves either thrombolytic agents or cardiac catheterization (angioplasty), in which a stent is inserted into the occluded vessel followed by gentle inflation of a balloon to rupture the plaque or clot which is causing the blockage.

The destructive role played by reperfusion has been the subject of much debate amongst clinicians and scientists (Ferrari and Hearse, 1997; Kloner, 1993; Ovize, 1997; Przyklenk, 1997; Yellon, 1992).

There is little doubt that unremitting ischaemia causes cell death but until recently, the predominant view was that if ischaemia itself was the culprit in cell death as a result of adenosine triphosphate (ATP) depletion and its numerous consequences, then reperfusion was

essential to restore ATP synthesis, thereby rescuing the uninjured cells or those which were reversibly injured.

Others, however, held the view that the death of cells irreversibly injured by ischaemia was merely accelerated by the restoration of blood flow. Yet another point of view was that reperfusion injury was caused by the sudden influx of blood releasing reactive oxygen species and other harmful substances. Agreement has, however, been reached that 4 forms of reperfusion injury exist (Ovize, 1997):

- i) myocardial stunning,
- ii) reperfusion arrhythmias,
- iii) vascular reperfusion injury and
- iv) lethal reperfusion injury

Lethal reperfusion injury is a term used to describe myocardial cell death caused by reperfusion, and is the type of reperfusion injury addressed in this thesis, (Figure 8).

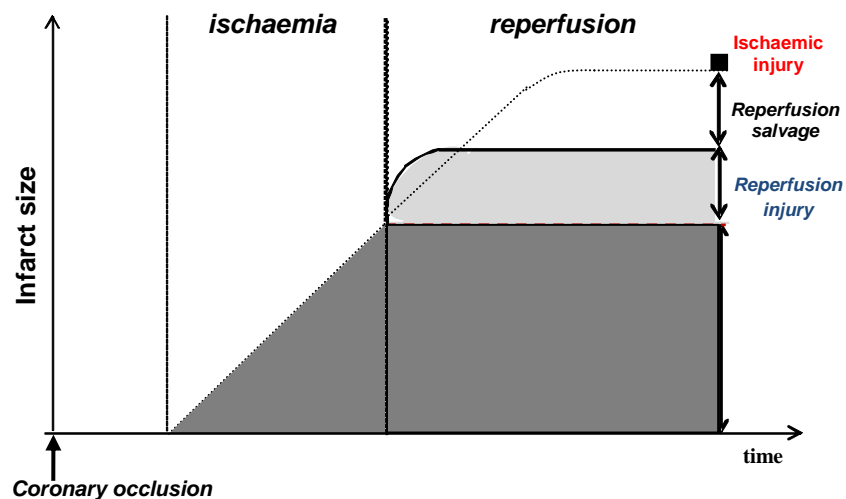


Fig 8: Lethal reperfusion injury. During ischaemia some of the cardiomyocytes are irreversibly injured (dark grey), whilst even more damage is done during the reperfusion period (light grey), (adapted from Thibault, 2007).

2.1 Mechanisms resulting in reperfusion injury

Reperfusion injury is associated with all the signs of necrotic cell death, such as release of intracellular proteins and major morphological changes which include swelling of the mitochondria. Studies over the past 10 years have revealed that mitochondria play a critical part in determining the fate of the cell after cellular stresses, (see review Kroemer et al., 2007). One of the critical components of mitochondrial response to stress is the activation of the mitochondrial permeability transition pore (mPTP) (Bernardi et al., 2006; Halestrap and Brenner, 2003). The mPTP is a non-specific, multimeric pore structure that bridges the outer and inner mitochondrial membranes. The true composition of the pore remains uncertain and is the subject of much debate. Cyclophilin-D is the only known component of the mPTP and is inhibited by cyclosporine-A (Crompton et al., 1988; Gomez et al., 2009). The inner mitochondrial membrane is impermeable to solutes while it remains closed under normal physiological conditions. During conditions of cell stress, with high matrix calcium ion accumulation, accompanied by depletion of adenine nucleotide, elevated inorganic phosphate and oxidative stress, the mPTP opens in the inner mitochondrial membrane, releasing cytochrome C and promoting cell death. These are exactly the conditions which occur during reperfusion after a period of ischaemia (Javadov et al., 2003). The mechanisms determining whether the pore opens or not during ischaemia-reperfusion have been extensively studied over the last 20 years with most investigators now being of the opinion that it is the conditions during early reperfusion which trigger the opening of the pore and that if the pore remains closed during these first early moments of reperfusion this will protect against cell death (Crompton et al., 1987; Di Lisa and Bernardi, 2006; Di Lisa et al., 2001; Duchen et al., 1993; Halestrap et al., 2004; Halestrap et al., 1998; Lemasters et al., 1997)

Griffiths et al, demonstrated that if the pore remains closed at the onset of reperfusion, a distinct reduction in infarct size is observed in experimental ischaemia-reperfusion (Griffiths and Halestrap, 1993). The past 5 years have seen the proposal of mPTP inhibition as being the ultimate cardioprotective target, with mounting evidence that it may be a crucial point of convergence for many of the cardioprotective signalling pathways. If this proves to be the case, administration of cyclosporine-A, a known mPTP antagonist, will be a simple and practical procedure, which could be delivered in the ambulance en route to the hospital and reduce the subsequent reperfusion injury which occurs with percutaneous coronary intervention (PCI) or with administration of a thrombolytic agent (Gomez et al., 2009).

In summary, this necrotic cell death, which has been termed “lethal reperfusion injury”, is a major obstacle to reperfusion therapy without which the ischaemic tissue cannot be salvaged. In addition, it has been suggested that the greater the degree of the initial ischaemic insult, the more severe the reperfusion injury will be and that if the ischaemic insult lasts long enough, no recovery is possible (Vinten-Johansen et al., 2005). Reperfusion has thus been dubbed as a “double edged sword” and a reassessment of reperfusion injury and its mechanisms is clearly needed to maximise the benefits of reperfusion and limit infarct size.

3.0 Conditioning the heart to limit reperfusion injury

3.1 Ischaemic preconditioning

Ischaemic preconditioning is defined as brief periods of ischaemia followed by reperfusion which occur prior to the prolonged ischaemic insult.

While investigating adenosine triphosphate (ATP) levels in a canine model of the failing heart, Murry and Jennings serendipitously discovered a biological mechanism which harnessed the intrinsic ability of the heart to protect itself (Murry et al., 1986). By applying four brief episodes of alternating reperfusion and occlusion prior to the infarct-inducing episode of myocardial ischaemia, Murry was able to demonstrate an astonishing reduction of 75% in the infarct size of canine hearts (Murry et al., 1986). This type of cardioprotection became known as “ischaemic preconditioning” (IPC) and a flurry of activity followed with the reproducibility of IPC being proved in multiple experimental settings using various animal models (Kuzuya et al., 1993; Marber et al., 1993; Murry et al., 1986; Przyklenk et al., 1993).

Further studies revealed the existence of two “windows” of protection in IPC. The first window, termed as “classic preconditioning” occurs immediately after initiating IPC and lasts for up to 3 hours. The second window of protection, known as SWOP or delayed preconditioning, occurs 24 to 72 hours after the initial application of IPC and is thought to protect the heart to a lesser extent than classical IPC, (see review (Yellon and Downey, 2003)).

3.1.1 Mechanisms of IPC:

Following the discovery of IPC, intensive research to delineate the mechanism involved has been undertaken but the exact signalling pathway still remains unclear. The major signalling pathways which have been implicated in IPC are illustrated in the following schematic, (Figure 9) and discussed below:

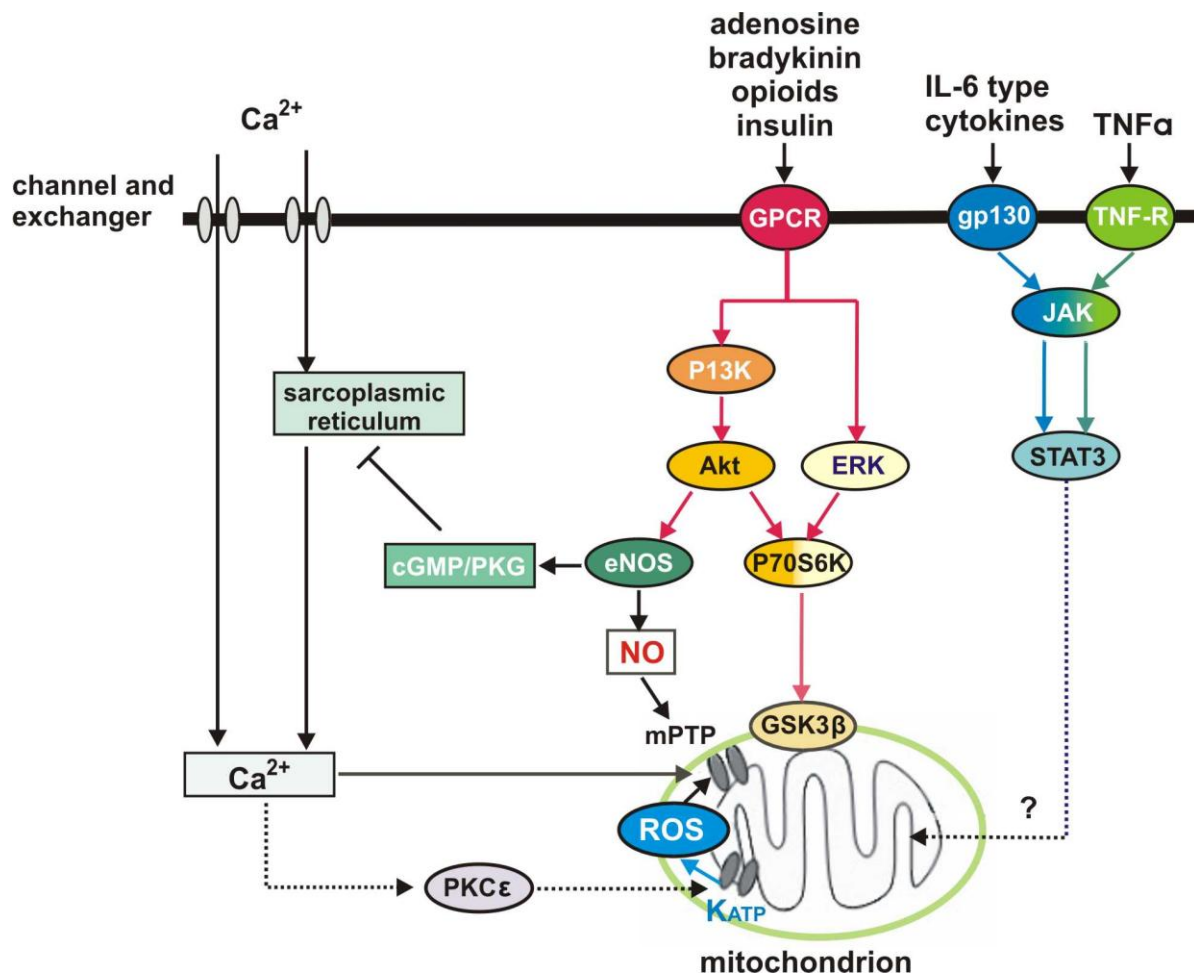


Fig 9: Diagram depicting suggested prosurvival signalling pathways involved in classic ischaemic preconditioning. IPC invokes the activation of signal transduction cascades by autocooid triggers and kinases. ROS targets the mK_{ATP} and the mPTP. Abbreviations as defined in text, (adapted from Ovize, et al, 2010).

3.1.2 Gi-protein coupled receptors (GPCR)

G-proteins are situated within the lipid bilayer of cells and they can move freely, thus allowing a large number of G-proteins to be activated by a single occupied receptor complex (Lubert, 1988). Gi-protein coupled receptors constitute a large and diverse family of proteins, the primary function of which is to transduce extracellular stimuli into intracellular signals.

Two types of G-proteins have been identified: the G stimulatory protein (Gs), which is cholera toxin sensitive, and the G-inhibitory protein (Gi) which is pertussis toxin sensitive. However, controversy surrounds the role of the Gi protein in cardioprotection. Thornton et al, used an *in vivo* rabbit model and Miura et al, used a canine model to investigate the involvement of the Gi-proteins in the protective effects of IPC. Both groups found that IPC-induced cardioprotection was conferred via the Gi-proteins (Miura et al., 1997; Thornton et al., 1993). In contrast, Liu and

Downey were unable to demonstrate a role of the Gi-proteins in IPC-induced protection in an open chest rat model of myocardial infarction (Liu and Downey, 1993).

G-coupled protein receptors activated by adenosine, bradykinin, and the opioid receptors will be further discussed under “Cardioprotective agents and ischaemic preconditioning” in Section E.

3.1.3 The RISK pathway

A powerful cardioprotective signalling pathway termed as the “Reperfusion Injury Salvage Kinase (RISK) pathway” consists of a group of prosurvival protein kinases, including phosphatidylinositol 3-kinase (PI3-kinase)/Akt and extracellular signal-regulated kinases, p44/p42 (Erk1/2) (Hausenloy and Yellon, 2004).

i) Phosphatidylinositol 3-kinase/Akt

The PI3-kinase/Akt signalling cascade is activated by the stimulation of a wide range of receptors, including those of growth factors and G-protein coupled receptors (Cross et al., 2000). In the setting of ischaemia-reperfusion, signalling through PI3-kinase has been demonstrated to confer protection against ischaemia-reperfusion injury by inhibiting apoptosis in cardiomyocytes subjected to hypoxia via the activation of Akt (Matsui et al., 1999).

In 2002 Mocanu et al, first demonstrated a link between ischaemic preconditioning, PI3-kinase /Akt and infarct limitation in the isolated perfused rat heart by individual infusion of the inhibitors, wortmannin and LY294002, during the preconditioning protocol (Mocanu et al., 2002).

ii) Extracellular signal-regulated kinase

The role of Erk1/2 has been extensively studied in the setting of cardioprotection. Erk 1 (p44) was not activated by global ischaemia in a perfused rat heart model but ischaemia-reperfusion increased the phosphorylation and kinase activity of both Erk1 and Erk2 (p42) in the nuclear fraction (Mizukami and Yoshida, 1997). In an *in vivo* rat model of ischaemia-reperfusion, activation of Erk1 was not triggered by ischaemia alone but was activated by 5 – 30 minutes of reperfusion (Omura et al., 1999).

Hausenloy and colleagues demonstrated that IPC resulted in the activation of both PI3-kinase/Akt and the Erk1/2 signal cascades at the time of reperfusion (Hausenloy et al., 2004b). They reported that inhibition of PI3-kinase at the time of reperfusion resulted in the activation of Erk1/2 which, in turn, activated p70-ribosomal s6 kinase (p70s6K). However, when Erk1/2 was inhibited, the PI3-kinase/Akt pathway was activated but p70s6K remained inactive, therefore

suggesting a cross talk between the kinases (Hausenloy et al., 2004b). During the preconditioning stimulus, Erk1/2 was phosphorylated and activated in the isolated rat heart and this was associated with a decrease in infarct size (Fryer et al., 2001). In contrast, Mocanu et al, found that infusion of PD98059, the Erk1/2 inhibitor, during the IPC stimulus of the isolated rat heart, did not block the protection afforded by preconditioning, therefore suggesting that Erk1/2 was not involved in the PI3-kinase–induced reduction in infarct size during IPC (Mocanu et al., 2002).

By 2007, the downstream targets of the RISK pathway included other cardioprotective kinases such as protein kinase C (PKC), protein kinase G (PKG), p70-ribosomal s6 kinase (p70s6K) and glycogen synthase kinase-3 beta (GSK-3 β) (Hausenloy and Yellon, 2007b) (Figure 10).

3.1.4 Protein Kinase C

The first kinase to be examined in detail in ischaemia-reperfusion was PKC. Of the 11 isoforms of protein kinase C, PKC alpha (α), PKC delta (δ) and PKC epsilon (ϵ) have been implicated in IPC in the rat heart (Yoshida K et al., 1997), while PKC ϵ and PKC gamma (γ) have been implicated in IPC in the rabbit heart and mouse heart (Gray et al., 2004; Ping et al., 1997). Enhanced membrane PKC activity in myocardial ischaemia was first demonstrated by Prasad et al, in 1992, using a protein kinase activity assay of the cytosol and membrane fractions of hearts which had been subjected to ischaemia or ischaemia-reperfusion (Prasad and Jones, 1992).

Wei et al, demonstrated in an *in vivo* rat ischaemia-reperfusion model, that when PKC β inhibitors were administered for a period of 2 weeks prior to experimentation, protection was conferred against cardiac microvascular ischaemia-reperfusion injury, thus implicating the β isoform of PKC in cardioprotection (Wei et al., 2010).

Activation and subsequent translocation of PKC has been reported as necessary for the protective effects of IPC (Ping et al., 1997). Administration of inhibitors of PKC prior to and/or during IPC abolished the cardioprotective effect of IPC in most animal models (Pain et al., 2000; Yellon and Downey, 2003). Furthermore, IPC could be mimicked by administration of a PKC agonist prior to an I/R insult (Speechly-Dick et al., 1994). Other studies in which agonists of PKC were used in isolated rat hearts and in cell culture (cardiomyocytes) supported these findings (Gray et al., 1997). It has been suggested that the PI3-kinase/Akt pathway, nitric oxide (NO), the mitochondrial potassium-sensitive adenosine triphosphate channel (mK_{ATP}) channel and ROS are upstream of PKC activation in the preconditioned heart (Baines et al., 1997; Ping et al., 1999a; Tong et al., 2000; Wang., 1999). Hausenloy et al, however, speculated that PKC may contribute to the cardioprotective effect of IPC by the phosphorylation of PI3-kinase/Akt at the

time of reperfusion (Hausenloy et al., 2004b). Downstream targets of PKC include p38 mitogen activated protein kinase (p38 MAPK), c-Jun-NH₂-terminal kinases (JNK), and extracellular signal-regulated kinase 1/2 (Erk1/2) (Maulik et al., 1996; Ping, et al., 1999b; Ping, et al., 1999c).

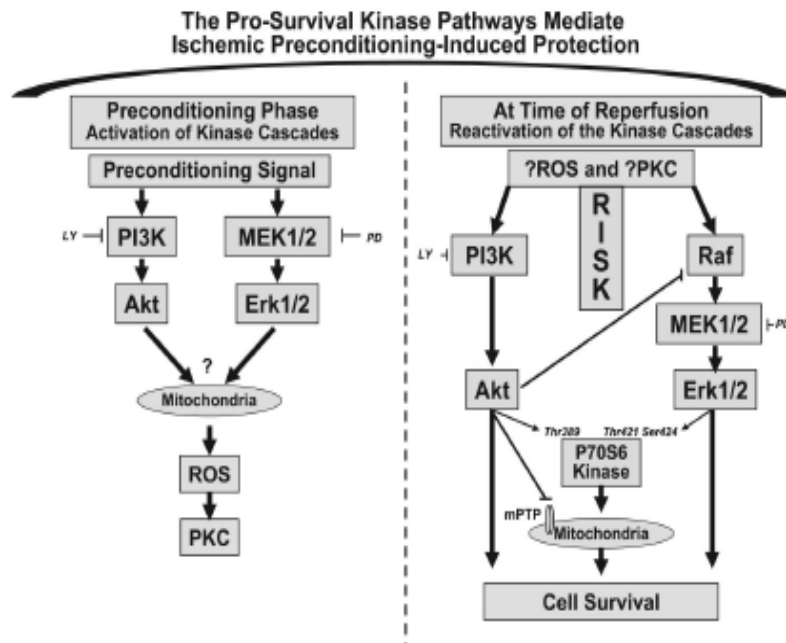


Fig 10: Hypothetical scheme outlining two phases of kinase cascade activation in response to IPC. During IPC, mitochondrial reactive oxygen species are released and PKC is activated. These events reactivate the PI3-kinase/Akt/p70S6K and Erk1/2/p70S6K cascades which comprise the reperfusion injury salvage kinase (RISK) pathway, at reperfusion. The RISK pathway mediates cellular survival through several possible mechanisms, which may include inhibition of mitochondrial permeability transition pore (mPTP) opening. MEK1/2: MAPK1/2. Further abbreviations as defined in the text. (Hausenloy, 2004)

3.1.5 Mitogen activated protein kinases and c-jun-NH₂-terminal kinases

Activation of the mitogen activated protein kinases (MAPK), p38 and c-Jun-NH₂-terminal kinases (JNK) to confer protection at the time of reperfusion gave conflicting results, with some studies reporting cardioprotective roles of these kinases at the time of myocardial reperfusion (da Silva et al., 2004) and others suggesting a detrimental role (Sun et al., 2006). A study conducted in a conscious rabbit model demonstrated that IPC stimulation increased the activity of JNK and p38 MAPK (Ping et al., 1997). This increase, together with the protective effect of IPC, was abrogated by PKC inhibition (Ping et al., 1999d). Earlier studies in rabbit and rat hearts had reported an increased activation of p38MAPK following an IPC stimulus and also during the index ischaemia in isolated rat and rabbit hearts (Maulik et al., 1996; Maulik et al., 1998; Nakano et al., 2000; Weinbrenner et al., 1997). In contrast, an *in vivo* study in pigs demonstrated that the activation of p38 MAPK was unaltered (Behrends et al., 2000) and a later

study reported that in the *in vivo* canine heart, the activation of p38 MAPK was actually decreased (Sanada et al., 2002). Therefore, the exact role of p38MAPK in IPC still needs to be clarified.

3.1.6 Nitric oxide - cyclic guanosine monophosphate - protein kinase G signalling cascade

Many studies, in different animal models, have reported that inhibition of nitric oxide (NO) is deleterious in ischaemia-reperfusion (Gonon et al., 2000; Maulik et al., 1995; Ockaili et al., 1999; Rakhit et al., 2000; Xi and Kukreja, 2000). In these studies, different nitric oxide agonists were administered before initiation of ischaemia. The protection afforded by the activation of nitric oxide was abrogated when antagonists were administered during the ischaemic period and continued for the first few minutes of reperfusion. A study done in the isolated rat heart demonstrated a reduction in infarct size when the vasodilator, adrenomedullin, was given during early reperfusion and this effect was lost if nitric oxide synthase (NOS) was inhibited (Hamid and Baxter, 2005). Similar results were observed in the mouse heart, after administration of adrenomedullin prior to reperfusion (Hamid et al., 2007).

Evidence for the generation of NO and peroxynitrite during ischaemia-reperfusion has been provided by numerous studies (Csonka et al., 1999; Wang and Zweier, 1996; Weselcouch et al., 1995; Yasmin et al., 1997), but no increase in tissue NO could be demonstrated during the preconditioning protocol itself (Csonka et al., 1999). In 2000 Lochner et al, using an isolated rat heart model, demonstrated a role for NO and cyclic guanosine monophosphate (cGMP) as a trigger in classic preconditioning (Lochner et al., 2000).

NO generation results in the activation of soluble guanyl cyclase and an accumulation of cyclic guanosine monophosphate (cGMP), with subsequent activation of protein kinase G, a cGMP-dependent protein kinase (Hausenloy and Yellon, 2006). PKG is considered to be an important mediator of cardioprotection in IPC, (see review (Burley et al., 2007)). Activation of PI3-kinase/Akt phosphorylates endothelial nitric oxide synthase (eNOS), producing NO (Cohen et al., 2000; Critz et al., 2005; Oldenburg et al., 2002) which has been implicated in the protection of endothelial cells (Dimmeler et al., 1999). NO, in turn, has been shown to protect by inhibiting the opening of the mPTP (Balakirev et al., 1997).

3.1.7 Reactive oxygen species

Free radicals are molecules or atoms with unpaired electrons in their outer shell. Free radicals that originate from oxygen are called reactive oxygen species (ROS) and are highly reactive.

ROS are produced by several enzymes and biochemical processes in the body. In the cardiovascular system the main sources of ROS are: a) Nicotinamide adenine dinucleotide phosphate (NADPH) oxidase, (b) the enzyme, xanthine oxidoreductase, (c) nitric oxide synthase (NOS) and, (d) the mitochondrial cytochromes (Donzelli et al., 2006; Grisham et al., 1999; Mason et al., 2000; Wink and Mitchell, 1998). However, in the myocardium itself, the mitochondria appear to be the main source of ROS production (Di Lisa et al., 2007).

Production of ROS during ischaemia has been shown in isolated cardiomyocytes and in isolated hearts (Becker et al., 1999; Kevin et al., 2003). A recent *in vivo* model of endothelial ischaemia-reperfusion injury in the arm demonstrated for the first time, in humans, that endothelial function after I/R injury is affected by ROS production (Loukogeorgakis et al., 2010). At the onset of reperfusion, a sudden burst of ROS occurs, which has been shown to contribute to myocardial injury and cardiomyocyte death (see review (Hausenloy, 2009)).

In an isolated rat heart model IPC reduced the excessive amount of ROS production at the onset of reperfusion, suggesting a deleterious role for the excess ROS in the early stages of reperfusion (Crestanello et al., 1996). In contrast, a study by Baines et al, suggested that ROS may also play a critical role in the signal transduction pathway underlying IPC. This group demonstrated that the antioxidant, N-2-mercaptopyrionylglycine (MPG), blocked the infarct-sparing effect of IPC in both *in vitro* and *in vivo* rabbit heart experiments, therefore suggesting that the presence of ROS was necessary for protection (Baines et al., 1997). In fact, a moderate release of ROS following the preconditioning stimulus leads to a decrease of ROS production at the onset of reperfusion (see review (Hausenloy, 2009)).

3.1.8 The JAK-STAT pathway

The Janus Kinases (JAKs) are a family of tyrosine kinases associated with membrane receptors. Ligands such as TNF α , interleukin 6 (IL-6) or growth factors activate the membrane receptors enabling the transphosphorylation and activation of two adjacent juxtaposed JAKs which can then subsequently activate the signal transducer and activator of transcription-3 (STAT-3) during ischaemia-reperfusion (see review (Lecour, 2009a)). STAT-3 is then phosphorylated at the tyrosine level, enabling it to form homodimers or heterodimers that subsequently translocate to the nucleus, resulting in gene transcription, (Figure 11).

The JAK-STAT pathway

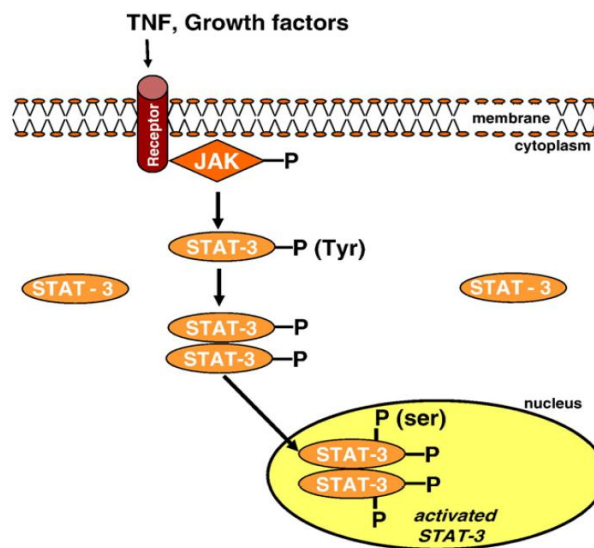


Fig 11: The JAK-STAT-3 signalling pathway. JAK2 is activated in response to a stimulus. Activated JAK 2 recruits STAT-3 from the cytoplasm and phosphorylates STAT-3 on a tyrosine residue. This phosphorylation allows STAT-3 to form homodimers and translocate to the nucleus where it upregulates transcription. STAT-3 is further phosphorylated on a serine residue to improve DNA interaction. JAK=Janus kinases, Ser=serine, Tyr=tyrosine, (Lecour, 2009).

The first study to implicate STAT-3 after acute myocardial infarction was performed in 2001 by Negoro and colleagues. In this experiment, the activation and increased phosphorylation of STAT-3 was observed for up to 24 hours after ligation of the left coronary artery in rats (Negoro et al., 2001) and the inhibitor of the JAK/STAT pathway, AG490, suppressed the phosphorylation of STAT-3 with a resulting increase in the number of apoptotic cells, implicating STAT-3 as a protective agent in ischaemia (Negoro et al., 2001).

Using an *in vivo* left coronary artery ligation model in mice, Bolli's group was the first to demonstrate the phosphorylation and translocation of STAT-3 to the nucleus in the delayed phase of ischaemic preconditioning (Xuan et al., 2001). In early ischaemic preconditioning, JAK2 and STAT-3 are activated following the IPC stimulus in isolated working rat hearts and this effect was almost completely abrogated with the inhibitor of JAK2, AG490 (Hattori et al., 2001). Furthermore, cardiac-specific STAT-3 knockout mice cannot be protected with an ischaemic preconditioning stimulus (Smith et al., 2004).

Confirmation of the role of STAT-3 in ischaemic preconditioning has also been shown with pharmacological protective agents such as cardiotrophin-1 (Brar et al., 2001), erythropoietin

(Nishihara et al., 2006; Rafiee et al., 2005) or cannabinoid agonists (Montecucco et al., 2009). Preconditioning mimetics such as TNF α (Suleman et al., 2008) and opioids (Gross et al., 2006; Gross et al., 2007) also activate STAT-3 during the trigger phase of preconditioning. Several downstream effectors of the JAK/STAT pathway have been identified, including growth factors such as vascular endothelial growth factor (VEGF), pro survival proteins, proteins involved in cell proliferation such as Bcl-2, Bcl-x1, Mcl-1, Fas, cyclin D1, E1, p21 and other transcription factors (Amit-Vazina et al., 2005; Ikezoe et al., 2004; Liu et al., 2006; Marques et al., 2007; Torbenson et al., 2002; Zhang et al., 2007). In the context of ischaemic preconditioning, STAT-3 increases the anti-apoptotic gene, Bcl-2 and reduces the pro-apoptotic gene Bax (Hattori et al., 2001). Also, STAT-3 has been implicated in the mediation of preconditioning protection via the phosphorylation and therefore, the inactivation, of the pro-apoptotic factor Bad (Lecour et al., 2005b). Other possible targets of STAT-3 may be mitochondrial connexin 43, an important component of the signal transduction cascade of IPC-induced protection (Ozog et al., 2004; Schulz et al., 2007), Akt and GSK3- β (Gross et al., 2006; Juhaszova et al., 2004).

3.1.9 The SAFE pathway

An alternative and, possibly parallel, signalling pathway to the RISK path has recently been implicated in IPC. Lecour et al, have named this pathway the Survivor Activating Factor Enhancement (SAFE) pathway, (Figure 22) which includes the cytokine, tumour necrosis factor alpha (TNF α) and the transcription factor STAT-3 as triggers and mediators (Lecour, 2009a). This pathway will be discussed in further detail in the section 4.8 and in Section D.

3.1.10 Receptors for Advanced Glycation End Products (RAGE)

Advanced glycation end products (AGEs) are formed by the nonenzymatic glycation and oxidation of proteins and lipids. The signal transduction receptor for AGEs is known as RAGE.

A range of signal transduction cascades is activated by RAGE, including the MAPKs, PI3-kinase/Akt, JAK-STAT, NF κ B, GSK-3 β and production of ROS via NADPH oxidase (Aleshin et al., 2008; Bierhaus et al., 2001; Huang et al., 2001; Kim et al., 2008; Marsche and others, 2007; Shang et al., ; Wautier et al., 2001).

Multiple cell types express RAGE, including cardiomyocytes and ischaemia-reperfusion is a potent generator of AGEs. In a study using ischaemic rat hearts, expression of RAGE was

significantly upregulated and pretreatment of the rats, 3 days prior to experimentation, with soluble RAGE (which acts as a decoy to bind the ligands of the receptor thus blocking it) reduced ischaemic injury and improved functional recovery of the myocardium (Bucciarelli et al., 2006). Moreover, the same group demonstrated that isolated and perfused hearts from RAGE null mice subjected to ischaemia-reperfusion were strikingly protected against I/R damage (Bucciarelli et al., 2006). RAGE can also modulate hypoxia/reoxygenation injury in adult murine cardiomyocytes via JNK and GSK-3 β signalling pathways (Shang et al.). High-Mobility Group Box-1 (HMGB-1), a nuclear factor released by necrotic cells, signals via RAGE. When mice were preconditioned with HMGB-1 in an ischaemia-reperfusion model, TNF α levels were decreased and lipopolysaccharide (LPS) tolerance was induced, suggesting that the RAGE receptor was involved in ischaemic preconditioning (Aneja et al., 2008).

3.1.11 Proteinase activated receptor-2

Proteinase activated receptors (PAR-2) are a recently described, novel family of seven transmembrane G-protein coupled receptors which are activated by protease cleavage (Cirino et al., 2000; Nystedt et al., 1994)

PAR-2 is expressed in vascular tissue and highly vascularized organs (Bohm et al., 1996; Nystedt et al., 1994; Nystedt et al., 1995; D'Andrea et al., 1998). The first evidence that PAR-2 was expressed in the heart and that it also contributes in the reduction of myocardial ischaemia-reperfusion injury was provided by Napoli et al, (Napoli et al., 2000). In Napoli's seminal study, an agonist of PAR-2 was added to the perfusate of the isolated heart 15 minutes into the stabilisation period and continued for 5 minutes into reperfusion. This study demonstrated that activation of PAR-2 resulted in an improved functional response associated to an improvement of metabolic parameters of the heart during experimental ischemia-reperfusion injury. Napoli and colleagues subsequently used an isolated perfused rat heart model to demonstrate that enhanced activation of PAR-2 during the IPC stimulus, with a PAR-2 activating peptide, improved the protective effect of ischaemic preconditioning (Napoli, 2002). A study conducted in rat neonatal ventricular cardiomyocytes, demonstrated that in the heart, PAR-2 stimulated p38 MAPK and ERK1/2, elevated calcium concentration and also activated JNK and Akt (Sabri et al., 2000). PAR-2 has also been implicated in vessel relaxation via the NO pathway (Cheung et al., 1998; Cicala et al., 1999; Emilsson et al., 1997) and endothelin-1 (Magazine et al., 1996) (Figure 12).

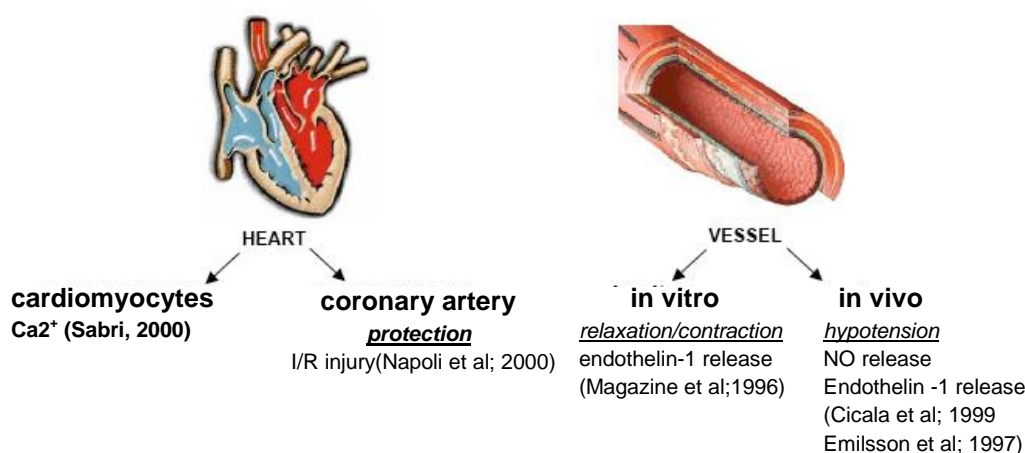


Fig 12: In vitro and in vivo effects of PAR-2 activation, (adapted from Bucci, 2005).

In addition, activation of PAR-2 significantly reduced the infarct size 24 hours after administration in a mouse model of acute myocardial ischaemia (Lim et al., 2007). The cardioprotective effect of PAR-2 agonists has been linked to PKC and Erk1/2 activation (Blackheart et al., 1996), increased TNF α levels in cardiac tissue (Bopassa et al., 2006a) and manganese superoxide dismutase activity (Camerer et al., 2000).

3.1.12 Glycogen synthase kinase-3 beta

GSK-3 β is a serine/threonine kinase that was originally identified as an enzyme that phosphorylates and downregulates glycogen synthase, the rate limiting enzyme of glycogen metabolism (Rylatt et al., 1980). Subsequently, it has been shown that the role and importance of GSK-3 β extends far beyond intermediary metabolism but the nomenclature remains in use.

Phosphorylation of GSK-3 β results in its inactivation which has been reported to be antiapoptotic. Inhibition of GSK-3 β plays an important role in the mediation of cardioprotection via phosphorylation by proteins such as the tyrosine kinases, Akt, PKC and activation of the mitochondrial ATP-sensitive potassium channel (mK_{ATP}) and the mPTP (Juhaszova et al., 2004). A study using a pressure overload hypertrophic rabbit heart model subjected to ischaemia-reperfusion, demonstrated that contractile function after ischaemia in hypertrophied hearts treated with a GSK-3 β inhibitor was better preserved and that tolerance to ischaemia-reperfusion injury in the hypertrophied myocardium was improved in the presence of the inhibitor (Barillas et al., 2007). Similarly, inhibitors of GSK-3 β added in Langendorff perfused rat hearts subjected

to a regional ischaemia-reperfusion insult at two different perfusion pressures showed an improved infarct size (Mozaffari et al, 2008).

Numerous studies have implicated inhibition of GSK-3 β in the cardioprotective effects of IPC (Gross et al., 2004; Gross et al., 2006; Juhaszova et al., 2004; Tong et al., 2002). In one of these studies, evidence implicating the involvement of GSK-3 β in the protective effect of IPC was provided with the use of two inhibitors of GSK-3 β , lithium and SB216763, administered during the IPC protocol in an isolated rat heart model of ischaemia-reperfusion (Tong et al., 2002). From this study it was reported that phosphorylation and inactivation of GSK-3 β occurred during IPC which would be consistent with a role for GSK-3 β in preconditioning. In addition, this group linked PI3-kinase to the cardioprotective effect of phosphorylation of GSK-3 β (Tong et al., 2002). In 2004, Juhaszova et al, suggested the inhibition of GSK-3 β as a “master signalling switch” which leads to the convergence of the various prosurvival pathways to the putative end effector of protection, the mPTP, (Juhaszova et al., 2009) (Figure 13).

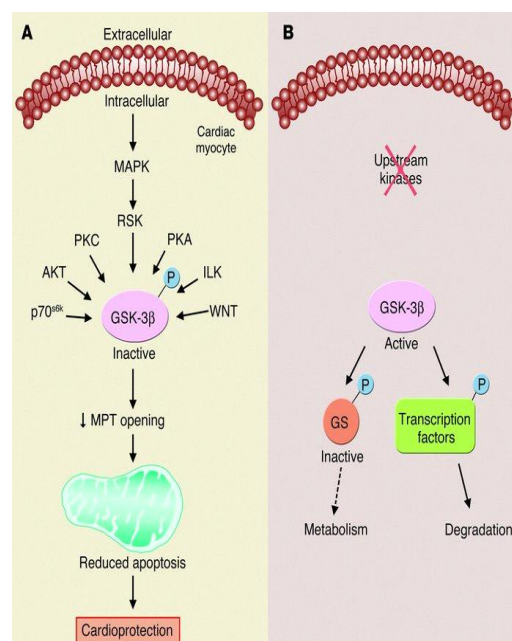


Fig 13: Scheme showing various kinases that phosphorylate and inactivate GSK-3 β . In the absence of upstream kinases, unphosphorylated GSK-3 β is active, phosphorylating and deactivating downstream targets such as glycogen synthase, or phosphorylating transcription factors and targeting them for degradation. GS = glycogen synthase; RSK = p90 ribosomal S6 kinase; ILK = integrin-linked kinase; WNT = homologues of Wingless; P = phosphorylation, (Murphy, 2004).

3.1.13 Role of the mitochondria, the mitochondrial ATP-sensitive potassium channel and the mitochondrial permeability transition pore in IPC

Mitochondria are spherical or rod shaped organelles found within the cytoplasm of eukaryotic cells and often referred to as the “powerhouse of the cell” since they act as the site for the production of adenosine triphosphate (ATP) which is vital as an energy source for several cellular processes.

Many prosurvival signalling cascades appear to converge to the mitochondria and given that the mitochondria are the “powerhouse” of the cell, this is no surprise. It is still a matter of debate as to how the prosurvival signals actually terminate at the mitochondria.

i) Mitochondrial Adenosine Triphosphate-sensitive potassium channel

Two potassium adenosine triphosphate-sensitive potassium channel subtypes coexist in the myocardium. One is located in the sarcolemmal membrane ($\text{sarcK}_{\text{ATP}}$) and the other in the inner mitochondrial membrane (mK_{ATP}). These channels allow coupling of membrane potential to cellular metabolic status and can be pharmacologically modulated by potassium channel openers and inhibitors.

Pharmacological agents which open the mK_{ATP} channel protect the ischaemic myocardium by direct protection of cardiac myocytes (Weselcouch et al., 1995). A study using an isolated canine gracilis muscle model demonstrated that when a mK_{ATP} opener was administered by direct infusion into the artery supplying the muscle, the infarct size was significantly reduced, therefore suggesting that mK_{ATP} activation is an important protective mechanism in this tissue (Weselcouch et al., 1995).

Garlid's group perfused isolated rat hearts with diazoxide, a potent opener of the putative mK_{ATP} channel, prior to ischaemia, demonstrating that opening of the channel was cardioprotective (Garlid et al., 1997). In addition, pharmacological activation of both PKC and MAPK which induce cardioprotection can be blocked with 5HD, a mK_{ATP} channel inhibitor, thus indicating that the channel may be downstream to these preconditioning mediators (Hopper et al., 2000; Yue et al., 2002). Other studies have demonstrated that PKC activates both the $\text{sarcK}_{\text{ATP}}$ and the mK_{ATP} channels (Hu et al., 1996; Sato T et al., 1998) but the actual role of these channels in IPC is controversial (Hanley P J., 2005). It should be noted that the mK_{ATP} channel has never been cloned and that much of the evidence obtained which has implicated the mK_{ATP} channel as playing a critical role in IPC was obtained using pharmacological inhibitors (Garlid et al., 1997; Liu et al., 1998b).

ii) Mitochondrial permeability transition pore

The mPTP is a large non-selective ion channel in the inner mitochondrial membrane. Mitochondrial permeability transition is an increase in the permeability of the mitochondrial membranes to molecules of less than 1500 Daltons in molecular weight. Mitochondrial permeability transition results from opening of mitochondrial permeability transition pores.

Under normal physiological conditions the mitochondrial inner membrane is impermeable to most molecules. During stressful conditions, such as ischaemia-reperfusion, the mPTP opens, allowing the transfer of molecules to the inner mitochondria (Hausenloy et al., 2004a) causing swelling and rupture of the outer mitochondrial membrane (Halestrap et al., 2004). Irreversible opening of this channel in response to the sudden return of flow of blood at reperfusion is considered to be responsible for cardiomyocyte death by uncoupling oxidative phosphorylation (Hausenloy, 2009).

Transient opening and closure of the mitochondrial permeability transition pore (mPTP) at the time of reperfusion has emerged as an important target for cardioprotection in IPC. In 2002, Hausenloy et al used an isolated rat heart model of IPC to demonstrate that an infusion of cyclosporine-A (CsA) at reperfusion, inhibited the opening of the mPTP, mimicking IPC-induced protection (Hausenloy et al., 2002). Subsequent studies have also implicated the closure of the pore at reperfusion in the protective effect of preconditioning (Hausenloy et al., 2004a; Javadov et al., 2003). Although the mechanism for this effect still needs to be elucidated, it may involve certain components of the RISK pathway such as Akt, Erk1/2 or GSK-3 β (Abdallah et al., 2006; Bopassa et al., 2006b).

3.1.14 Applications of ischaemic preconditioning

A robust and reproducible protection is exerted by the phenomenon of IPC and this has been demonstrated in the heart of every animal species tested thus far e.g. rabbits, dogs, ferrets, pigs, guinea pigs, sheep, mice, chicken and man (see review (Lawson and Downey, 1993)). However, these experimental results, done mainly in healthy, young animals may not be reproducible in humans as comorbidities, such as diabetes, obesity and left ventricular hypertrophy often accompany ischaemic heart disease in humans and these may influence the outcome of IPC (Moolman et al., 1997; Tsang et al., 2005). Moreover, the need to intervene before the index ischaemic event limits the clinical application of IPC as an acute myocardial infarction (AMI) cannot yet be predicted. The application of IPC in the clinical arena is, therefore, limited to scenarios in which the index ischaemia can be reliably predicted, such as in patients undergoing

surgery with extracorporeal circulation. Yellon et al, were the first to report that IPC induced by intermittent cross-clamping of the aorta in coronary artery bypass graft surgery preserved the patient’s ATP levels during the subsequent global ischaemic period (Yellon et al., 1993). Brief episodes of ischaemia are experienced by many patients prior to an acute myocardial infarction and it is theoretically possible that this pre-infarct angina may precondition the heart. Hence, it has been demonstrated that the presence of angina before AMI is associated with a smaller infarct size (Kloner et al., 1995). The discovery of “remote ischaemic preconditioning” in which either an upper limb or the leg can be used to apply IPC (usually by repeated, intermittent blood pressure cuff inflations and deflations) has revived the interest in IPC. This non-invasive remote form of IPC, known as RIPC, has been used successfully in children undergoing corrective heart surgery (Cheung et al., 2006), patients undergoing coronary artery bypass grafts (Hausenloy et al., 2007; Venugopal et al., 2009) and in elective percutaneous coronary interventions (Hoole et al., 2009).

3.2 Ischaemic postconditioning

Ischaemic postconditioning has been defined as a series of brief mechanical interruptions of reperfusion applied at the onset of reperfusion (Zhao et al., 2003).

Ischaemic postconditioning (IPostC) was first proposed in 2003 by Vinten-Johansen’s group as an attempt to offer an intervention which could be applied at the time of myocardial reperfusion to protect patients diagnosed with an acute myocardial infarction, (Figure 14).

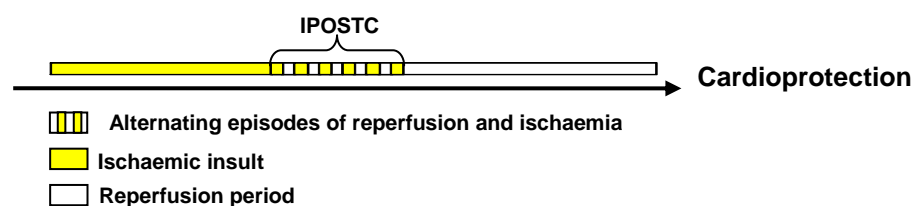


Fig 14: Schematic representation of ischaemic postconditioning. IPostC is initiated at the onset of reperfusion. The number and duration of the cycles depends upon the animal species.

As in IPC, many experiments have demonstrated that IPostC reduces infarct size in hearts of several different animal species such as: rabbit, mice, rats dogs, and pigs (Bopassa et al., 2006b; Darling et al., 2005; Heusch et al., 2006; Iliodromitis et al., 2006; Tsang et al., 2004; Yang et al., 2005; Zhao et al., 2003). The degree to which the heart is protected by ischaemic

postconditioning appears to be similar or slightly less than that obtained by ischaemic preconditioning (Fantinelli and Mosca, 2007; Halkos et al., 2004; Kin et al., 2004; Zhao et al., 2003) but the effect of IPostC is not enhanced by IPC (Halkos et al., 2004). As the clinical application of IPC was limited, the IPostC phenomenon has opened a novel era of research to limit reperfusion injury.

In their seminal paper, Zhao and colleagues, demonstrated that interruption of myocardial reperfusion with 3 x 30 second episodes of myocardial ischaemia and reperfusion in a canine model, resulted in a dramatic reduction in myocardial infarct size, comparable to that seen with IPC (Zhao et al., 2003). Thus, ischaemic postconditioning (IPostC) evolved as a method to reduce reperfusion injury. The protective effect of IPostC is clearly demonstrated in figure 15 which shows a section from a mouse heart subjected to IPostC compared to a heart which received no intervention.



Fig 15: Ischaemic postconditioning reduces infarct size. **A:** heart subjected to ischaemia-reperfusion with no intervention. **B:** heart postconditioned with alternating cycles of reperfusion and ischaemia. Red area = healthy tissue; White area = infarcted tissue.

3.2.1 Mechanisms of IPostC: A remarkable assortment of molecular mechanisms, similar to those implicated in IPC, has been demonstrated to contribute to the protective effects of IPostC. A schematic showing some of the shared signalling cascades is shown in Figure 16.

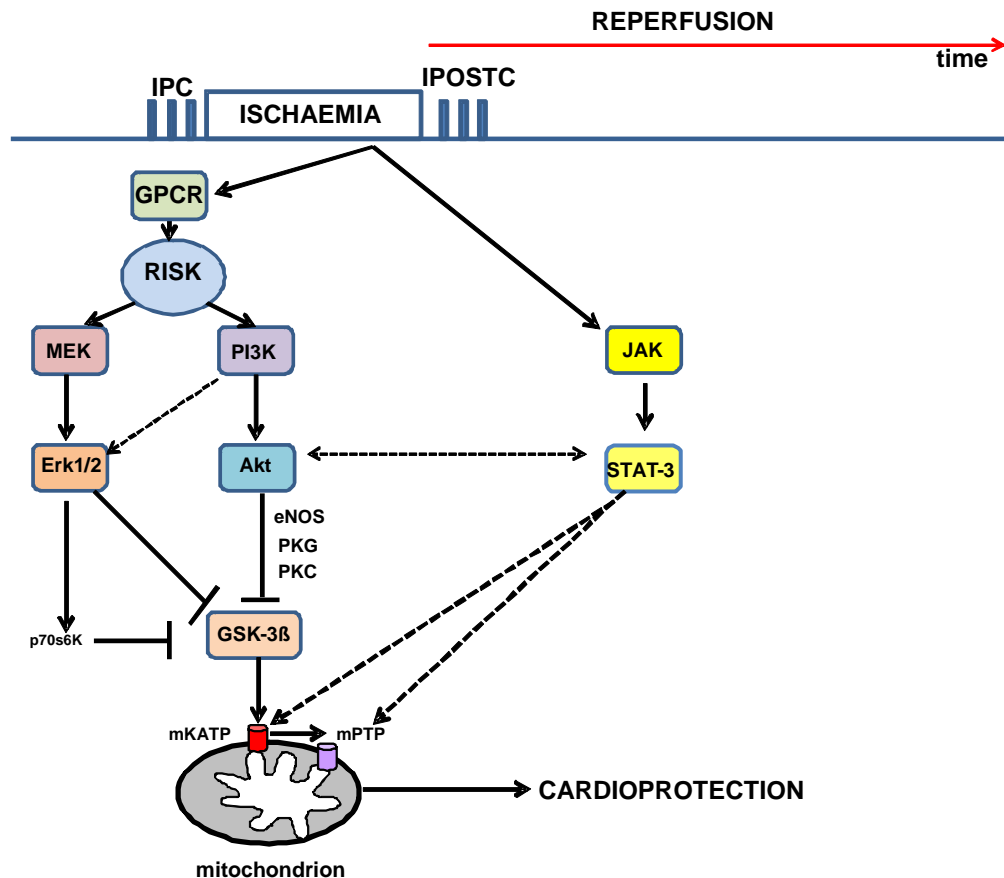


Fig 16: Hypothetical scheme outlining a few major signalling mechanisms common to IPC and IPostC. Abbreviations as defined in the text.

A number of signalling pathways which have been recruited by IPostC have been identified. These can be divided into the following:

3.2.2 Gi-protein coupled receptors

The human genome has been estimated to have an extremely large number of GPCRs and based on their sequences as well as on their known or suspected functions, there are estimated to be five or six major classes of GPCRs (Kroeze et al., 2003). Depending on the type of G protein to which a receptor is coupled, a variety of downstream signalling pathways can be activated (Marinissen and Gutkind, 2001; Neves et al., 2002).

The role of adenosine, bradykinin and opioids receptors will be discussed in section 3.3 related to pharmacological pre- and postconditioning.

3.2.3 The RISK pathway

In an isolated perfused rat heart, inhibition of PI3-kinase at reperfusion with wortmannin or LY290042 abolished the protective effect of IPostC (Tsang et al., 2004). Similarly, addition of PD98059, an inhibitor of Erk1/2 at the onset of reperfusion suggested that the protective benefits of IPostC were dependent on the activation of Erk1/2 (Yang et al., 2004b). Subsequent studies in both healthy and diseased animals as well as human atrial muscles have confirmed the role for Akt and Erk1/2 in IPostC (Feng et al., 2006; Sivaraman et al., 2007; Zhu et al., 2006). Potential distal signalling elements of the RISK pathway are suggested to be PKG, PKC ϵ and GSK-3 β (Burley and Baxter, 2009; Gomez et al., 2008) as already described in section 3.1.4.

However, a few recent experimental studies conducted in swine and rabbit models have failed to show components of the RISK pathway as mediators of IPostC-induced protection, therefore suggesting the existence of alternative pathways (Darling et al., 2005; Skyschally et al., 2009).

3.2.4 The JAK-STAT pathway

Phosphorylation of STAT-3 was first implicated in the cardioprotective effect of IPostC in 2006 by Lecour's group (Suleman et al., 2006). In isolated hearts, the protective effect of IPostC was abolished in the presence of the JAK2 inhibitor, AG490, and western blot analysis after 15 minutes of reperfusion revealed that IPostC decreased the phosphorylation of STAT-3 in the cytosol, therefore suggesting activation and translocation of STAT-3 to the nucleus after IPostC. The role of STAT-3 in IPostC was confirmed by Boengler et al in an *in vivo* STAT-3 knockout mouse model (Boengler et al., 2008). However, the evidence of interaction between the JAK/STAT pathway and the PI3-kinase pathway at the onset of reperfusion and during IPostC is inconclusive (Goodman et al., 2008) and the mechanism responsible for the activation of the JAK/STAT pathway in IPostC is unknown.

3.2.5 Nitric oxide-cGMP-PKG signalling cascade

The protective effects of IPostC have been demonstrated to be sensitive to pharmacological inhibition of the NO-cGMP-PKG pathway (Penna et al., 2006a; Yang et al., 2005). It is believed that, in response to IPostC, activated PKG opens the mK_{ATP} channel through PKC ϵ (Costa et al., 2005) which is presumably activated via the Akt component of the RISK pathway and eNOS (Tsang et al., 2004).

3.2.6 ROS signalling

IPostC reduced the amount of harmful ROS generation at the onset of myocardial reperfusion (Zhao et al., 2003). ROS signalling in the first moments of reperfusion, however, appears necessary for the protective effect of IPostC (Penna et al., 2006b) as inhibitors of ROS given at the onset of reperfusion, abolished the protective effect of IPostC. At present, the role of ROS signalling during IPostC still remains unclear.

3.2.7 Receptor for advanced glycation end products

A current literature search revealed no articles on RAGE and IPostC but the receptors for advanced glycation end products have been involved in protection during ischaemia-reperfusion and seem to be implicated in IPC (Aneja et al., 2008; Bucciarelli et al., 2006). Therefore, it is appropriate to suggest that RAGE may play a role in IPostC.

3.2.8 Proteinase Activated Receptor-2

Ischaemia-reperfusion injury and angioplasty balloon catheter injury are reported to increase the expression of PAR-2 (Damiano et al., 1999; Napoli et al., 2000) and PAR-2 activation has been shown to be protective in other tissues, including brain, airways and myocardium undergoing ischaemia or lipopolysaccharide challenge (Jin et al., 2005; Morello et al., 2005; Napoli et al., 2000).

Although never studied in IPostC, Vinten-Johansen's group demonstrated firstly, that PAR-2 is present in the left ventricular tissue of rat heart, in neonatal rat cardiomyocytes and in H₉C₂ rat cardiac-derived cells and secondly, that a PAR-2 agonist administered 5 minutes before reperfusion in the isolated rat heart, salvaged the myocardium via an Erk1/2 pathway (Jiang et al., 2007). These data suggested that IPostC may protect via activation of PAR-2.

3.2.9 Glycogen synthase kinase-3 beta (GSK-3 β)

Inactivation of GSK-3 β is a well-established downstream target of the RISK pathway. As in IPC, inhibition of GSK-3 β is believed to play an important role in IPostC, although experimental results have been somewhat contradictory. In an isolated rat heart model, Gross et al, first implicated inhibition of GSK-3 β at the onset of reperfusion (Gross et al., 2004). Gomez et al, reported that inhibition of GSK-3 β by postconditioning was required in transgenic mice to prevent the opening of the mPTP during reperfusion (Gomez et al., 2008). In contrast, Nishino et al, also using transgenic mice, found that inactivation of GSK-3 β was not required to confer

protection in either IPC or IPostC (Nishino et al., 2008). Skyschally and colleagues postconditioned pigs in the presence of inhibitors of the RISK pathway and found that protection was still induced in the pigs although blockade of the RISK pathway resulted in a reduced GSK-3 β phosphorylation and p70s6K (Skyschally et al., 2009).

However, as previously mentioned, GSK-3 β could possibly be the central point where a number of protective signalling cascades converge and its cellular location close to the mPTP may be of significance in transferring prosurvival signals (Juhaszova et al., 2009).

3.2.10 Mitochondrial involvement in IPostC

Many of the signalling pathways transmitting the cardioprotective signal of IPostC appear to converge to the mitochondria. The heart is a rich source of mitochondria with approximately 40% of total cardiomyocyte volume consisting of mitochondria (Barth et al., 1992). Thus, mitochondria are both targets and sources of injury during cardiac ischaemia and reperfusion with both the putative mK_{ATP} channel and the mPTP being implicated as possible end effectors of the protective pathway (Gross and Gross, 2006; Gross et al., 2004; Hausenloy et al., 2002).

i) Mitochondrial potassium ATP channels

Pharmacological inhibition of the mK_{ATP} at the onset of reperfusion abrogated IPostC-induced protection (Donato et al., 2007; Yang et al., 2004b) but no direct studies have shown that IPostC can actually open this channel during reperfusion. A recent study by Garlid's group suggested that IPostC generates a signalosome which conveys the signal from the G-protein coupled receptors to the opening of the mK_{ATP} leading to the inhibition of the opening of the mPTP (Quinlan et al., 2008).

ii) Mitochondrial permeability transition pore

Several studies suggest a critical role of the mPTP in IPostC. Isolated rabbit heart mitochondria subjected to a standard IPostC protocol, are more resistant to calcium-induced opening of the mPTP, therefore suggesting that IPostC inhibits mPTP opening at reperfusion (Argaud et al., 2005). Bopassa et al, subsequently reported that inhibition of PI3-kinase/Akt with wortmannin or LY294002 at the onset of reperfusion abolishes the inhibitory effect on the pore's opening of IPostC, implicating IPostC in the mediation of pore inhibition via the activation of PI3-kinase/Akt (Bopassa et al., 2006a). Also, mice lacking cyclophilin-D, the only known component of the mPTP, cannot be protected by IPostC (Lim et al., 2007).

3.2.11 Clinical application of ischaemic postconditioning

Within 2 years following its initial description, IPostC proved to be effective in clinical studies. Several sequences of inflation/deflation of the angioplasty balloon applied after opening of the infarcted vessel, reduced infarct size, improved coronary blood flow and improved myocardial reperfusion in patients with acute myocardial infarction (Laskey, 2005; Ma et al., 2006; Staat et al., 2005; Yang et al., 2007). However, the application of IPostC in these studies was strictly restricted to patients undergoing primary percutaneous intervention (PCI) as the protocol was invasive.

Therefore, it is important to delineate the signalling pathways which are responsible for the cardioprotective effect of IPostC. Knowledge of these signalling cascades may pave the way for manipulation of pharmacological targets to achieve the same end result as IPostC. Cyclosporine-A, which prevents the opening of the mPTP at reperfusion has already been used in the clinical setting with a promising outcome. An important “proof of concept” study, provided evidence in humans that inhibition of the mPTP at reperfusion in patients with acute myocardial infarction is associated with reduction in infarct size (Piot et al., 2008).

3.3 Pharmacological pre- and postconditioning – the next step in limiting infarct size

Given the invasive nature of ischaemic pre and postconditioning and its limited use to elective PCI patients or patients undergoing cardiac surgery, pharmacological agents mimicking the protective effects of IPostC were sought and found. Included amongst the protective pharmacological agents were: adenosine, bradykinin and opioids.

3.3.1 Adenosine

Adenosine is a nucleoside composed of a molecule of adenine attached to a ribose sugar molecule moiety. It is a purine nucleoside endogenously expressed in the body and modulates many biological processes after binding to one of the adenosine receptors.

Increased release of adenosine has been reported during ischaemia and early reperfusion (Schrader et al., 1977). Adenosine, infused directly into the left descending coronary artery in a canine model reduced the deleterious effects of ischaemia by decreasing infarct size. This protective effect was attributed to the vasodilatory effects of adenosine (Engler, 1987; Olafsson et al., 1987; Takeo et al., 1988). Adenosine is one of three autacoids released by ischaemic tissue, the other two autacoids being bradykinin and opioids.

3.3.2 Adenosine receptors

There are 4 adenosine receptor subtypes, A₁, A_{2A}, A_{2B} and A₃ with all of the receptors being expressed in different cell types of the heart and blood vessels. Cardioprotection has been attributed mainly to the A₁ and A₃ receptors, (Black et al., 2002; Reichelt et al., 2005) with the A₃ receptor appearing to share a few similarities with the A₁ receptor, both having been associated with different kinase systems such as PKC, MAPK's, PI-3 kinase and Akt (Dana et al., 2000; Headrick et al., 2003; Mubagwa and Flameng, 2001). However, evidence also exists showing that the A_{2A} receptor is associated with cardioprotection (Norton et al., 1992; Xu et al., 2001). Adenosine and adenosine receptor agonists are the most widely studied cardioprotective ligands, but as yet, there is no definite agreement as to which receptor subtype contributes to cardioprotection during ischaemia or reperfusion phases.

3.3.3 Preconditioning with adenosine

The cardioprotective effect of IPC has been successfully mimicked by adenosine administration in various animal models tested, and in human atrial trabeculae (Cleveland et al., 1997; Liu et al., 1994; Liu et al., 1991). In a seminal study using *in vivo* rabbit hearts, Liu et al, infused two different non-selective adenosine antagonists directly before the IPC protocol was applied and each successfully blocked the IPC-induced protection (Liu et al., 1991). This data suggested that adenosine released during preconditioning stimulates the A₁ receptor. A few years later Liu et al, using a selective A₁ receptor antagonist and a selective A₃ receptor agonist, provided evidence demonstrating that the protection of IPC was not exclusively mediated by the A₁ receptor in heart but could also involve the A₃ receptor (Liu et al., 1994).

3.3.4 Postconditioning with adenosine

Downey's research group was the first to implicate adenosine in IPostC (Yang et al., 2005). In an isolated perfused rabbit heart, the presence of a non-selective A₁ receptor antagonist, an inhibitor of the PI3-kinase signalling cascade, wortmannin, and an inhibitor of guanyl cyclase abolished the infarct-sparing effect of IPostC and their data suggest that postconditioning-induced protection requires the adenosine A₁ receptor to activate PI3-kinase and guanyl cyclase (Yang et al., 2005). Similarly, Kin and colleagues used an isolated perfused mouse heart model to demonstrate that IPostC reduces the level of adenosine released during reperfusion (Kin et al., 2005a). This same group confirmed their data in an open chest rat model of myocardial infarction using A₁, A₂, or A₃ receptor antagonists, administered 5 minutes before reperfusion in the presence or absence of IPostC. Interestingly, IPostC in the rat protected the heart, reduced

the adenosine washout at 2 minutes of reperfusion and reduced infarct size and this effect was not abolished by the A₁ receptor antagonist. However, the A₂ and A₃ receptor antagonists did abrogate the protective effect of IPost, therefore suggesting that IPostC involved endogenous activation in the heart of the A₂ or A₃ receptor but not the A₁ receptor subtype (Kin et al., 2005a).

3.3.5 Proposed mechanisms in adenosine-induced cardioprotection

Contrary to the proposal by Cohen et al, suggesting that adenosine-induced protection was not dependent on opening of the mK_{ATP} (see review (Cohen and Downey, 2008)), intracoronary infusion with adenosine and dipyridamole (a drug which inhibits thrombus formation and also causes vasodilation) before occlusion in a canine model of IPC produced a significant reduction in infarct size which was abolished by pretreatment with glybenclamide, an ATP-dependent potassium channel antagonist. This suggested that activation of the A₁ receptors in the heart produces myocardial preconditioning via the opening of the K_{ATP} channels (Auchampach and Gross, 1993). The mechanism of protection may, therefore, be dependent on whether adenosine acts as a trigger or mediator of IPC.

PKC ϵ has also been implicated as a mediator in adenosine's preconditioning signalling pathway (Cohen and Downey, 2008). Cohen et al, hypothesized that in postconditioning, adenosine activates PKC ϵ which subsequently activates the A_{2B} receptor, Erk1/2, Akt and p70s6K, therefore inhibiting GSK-3 β and preventing the opening of the mPTP (Cohen and Downey, 2008). PI3-kinase and guanyl cyclase are involved in adenosine-induced postconditioning (Yang et al., 2005) and other investigators have also implicated the components of the RISK pathway in adenosine-induced cardioprotection (Kis et al., 2006; Park et al., 2006a; Yang et al., 2004a).

3.3.6 Clinical application of Adenosine

The ability of adenosine to limit reperfusion injury has been tested in two clinical trials involving patients with AMI. In the AMISTAD-I trial, patients undergoing fibrinolysis were randomised to receive either adenosine (70 μ g/kg/min for 3 hours) or a normal saline infusion within 6 hours of infarction onset. Infarct size was reduced by 33%, relative to the area at risk, (p<0.085). However, the percentage reduction in infarct size appeared to depend upon which area of the heart was infarcted. In addition, adenosine showed no significant effects on secondary endpoints and clinical outcomes such as death, stroke, congestive heart failure, re-infarction and cardiogenic shock (Mahaffey et al., 1999).

A second, larger AMISTAD trial (AMISTAD-II) sought to assess two doses of adenosine infusion, 50 and 70 µg/kg/min given for 3 hours, with treatment starting 15 min before the start of fibrinolysis or PCI, but still within 6 hours after infarction onset. Both doses reduced infarct size with the higher dose achieving a 59% reduction compared to 15% reduction with the lower dose. Despite the reduction in infarct size, no difference was observed in the survival rate between placebo-treated or adenosine-treated patients (Ross et al., 2005). The inconsistencies in the results of the two AMISTAD trials lead to major reservations about the reliability of infarct size quantification in clinical studies and also cast doubt about the protective effect of adenosine and selective adenosine receptor agonists at reperfusion in experimental studies (see review (Burley and Baxter, 2009)).

3.3.7 Bradykinin

Bradykinin is a nonapeptide kinin formed from a plasma protein, high-molecular-weight (HMW) kininogen, by the action of kallikrein; it is a very powerful endothelium-dependent vasodilator that helps to enlarge or open blood vessels. This lowers blood pressure, allowing the blood to flow more smoothly throughout the body. In addition, bradykinin constricts smooth muscle and stimulates pain receptors.

Similar to adenosine, bradykinin is elevated during and after an ischaemic insult (Baumgarten et al., 1993; Pan et al., 2000; Schulz et al., 1998).

3.3.8 Bradykinin receptors

Cardiomyocytes have two bradykinin receptors, a constitutive B₂ receptor and an inducible B₁ receptor (Bhoola et al., 1992; Marceau et al., 1998). Induction of the B₁ receptor in the left ventricle of rats occurs 6 hours after reperfusion, and increases up to 4 fold after 24 hours (Tschope et al., 2000; Tschope et al., 1999). Bradykinin was first associated with IPostC when two different pharmacological antagonists of the bradykinin 2 receptor abolished the protective effect of IPostC in the isolated rat heart (Penna et al., 2007).

3.3.9 Preconditioning with bradykinin

The cardioprotective effect of bradykinin has been demonstrated in various animal models such as mouse, rat, rabbit and pig hearts (Bell and Yellon, 2002; Penna et al., 2007; Penna et al., 2008; Tobe et al., 1991; Yang et al., 2004a).

Using B₂ receptor knockout mice, or a rat model deficient in high-molecular-weight (HMW) kininogen, Yang et al, demonstrated that IPC fails to protect against ischaemia-reperfusion,

therefore suggesting an important role for the B₂ receptor in IPC (Yang et al., 1997). The exact role of the B₁ receptor in cardioprotection is unknown although knockout of this receptor in female mice had no effect on remodelling after a myocardial infarction (Xu et al., 2005). Bradykinin can also mimic IPC in the rabbit heart. Exogenous bradykinin protected to a similar extent compared with IPC and the B₂ receptor blocker completely abrogated the protection (Goto et al., 1995). In addition, bradykinin can mimic the effects of IPC in humans undergoing percutaneous transluminal coronary angioplasty (Leesar et al., 1999). However, it is unknown whether bradykinin receptor inhibitors can abrogate the effect of IPC (Gross and Gross, 2006).

3.3.10 Postconditioning with bradykinin

Using an isolated rat heart model of postconditioning, Penna et al, demonstrated that intermittent infusion of bradykinin at the beginning of reperfusion stimulates a postconditioning-like cardioprotection (Penna et al., 2007). The bradykinin-induced protection was abrogated in the presence of a B₂ receptor antagonist, implicating this receptor in the protective effect. Similarly, isolated hearts from mice in which the bradykinin B₂ receptor, the bradykinin B₁ receptor or the adenosine A₁ receptor had been knocked out, showed that the protective effect of IPostC was partially attenuated in the B₁ receptor knockout but was abrogated in the absence of the B₂ receptor and the adenosine A₁ receptor (Liu et al., 2008). Using three distinctive strains of gene knockout mice, this study provided the first conclusive evidence showing IPostC-induced cardioprotection could be triggered by activation of multiple types of cell membrane receptors, which include adenosine A₁ and bradykinin B₁ and B₂ receptors.

3.3.11 Proposed mechanisms in bradykinin-induced cardioprotection

Bradykinin, given at reperfusion protects via PI3-kinase/Akt and Erk1/2 pathway and by opening the mitochondrial ATP-sensitive potassium channels with the generation of reactive oxygen species that act as second messengers to activate PKC (see review (Critz et al., 2005)). Studies in rat hearts with intermittent application of bradykinin at reperfusion, have demonstrated a sizeable reduction in infarct size with activation of NO, cGMP, PKG, mK_{ATP} opening, and ROS production being implicated in the protective pathway (Oldenburg et al., 2004).

An interesting study by Penna and colleagues in the isolated rat heart, reported the novel finding that postconditioning required endogenous activation of cyclooxygenase (COX) and release of prostacyclin GI₂ (PGI₂) and that bradykinin, given as a postconditioning mimetic, also triggered the COX activation and induced PGI₂ formation during the late phase of reperfusion (Penna et

al., 2008). This data suggested that ischaemic postconditioning and intermittent bradykinin postconditioning were not solely dependent on the activation of the RISK pathway but that it also depended on the late release of PGI₂ via activation of COX (Penna et al., 2008).

Quinlan and co-workers proposed another mechanism in which bradykinin interacted with its B₂ receptor, inducing the formation of a signalosome (a vesicular caveolar signalling platform). This would then phosphorylate an unknown receptor on the outer mitochondrial membrane. As the terminal kinase of bradykinin is protein kinase G (PKG), this would phosphorylate the unknown receptor at a serine/threonine residue. The signal would then be transmitted to activate protein kinase epsilon₁ (PKCε₁) on the inner mitochondrial membrane. PKCε₁ is situated close to the mitochondrial K_{ATP} channel and causes this to open with a subsequent increase in hydrogen peroxide (H₂O₂) production. H₂O₂ then activates a second PKCε - PKCε₂, which would then inhibit the opening of the mitochondrial permeability transition pore (mPTP) thereby leading to cardioprotection (Quinlan et al., 2008). However, the “signalosome” hypothesis remains a working hypothesis and requires further validation.

3.3.12 Clinical application of bradykinin

In a small clinical trial, thirty patients were randomised to receive an intracoronary infusion of bradykinin or normal saline for 10 minutes prior to percutaneous transluminal coronary angioplasty (PTCA). The purpose of this study was to see if bradykinin could reproduce the protective effect of IPC. Electrocardiogram (ECG), chest pain score and regional wall motion during inflation of the balloon were used to evaluate the effect of bradykinin infusion. Bradykinin-treated patients fared significantly better than the control subjects suggesting that intracoronary infusion of bradykinin before PTCA renders the myocardium relatively resistant to subsequent ischaemia. The extent of this bradykinin-induced protection was comparable to that afforded by IPC in control subjects. In addition, no adverse effects of the bradykinin infusion were reported, suggesting that bradykinin could be used in selected patients undergoing PTCA to reduce ischaemia-reperfusion injury (Leesar et al., 1999).

3.3.13 Opioids

Opioids are endogenous autacoids, released by ischaemic tissues. They are pain-attenuating peptides that occur naturally in the brain.

Endogenous opioid peptides are synthesised and released into the circulatory system after myocardial infarction. In the rat, the ventricular tissue was shown to have much higher levels of

preproenkephalin (an opioid) compared to the other organs, indicating that the heart may have a very significant endogenous opioid system (Romano et al., 2004).

3.3.14 Opioid receptors

Opioids exert their numerous pharmacological functions after fixation to their opioid receptors. Three opioid receptor subtypes have been cloned. In adult ventricular cardiomyocytes only two receptor subtypes have been reported, the kappa (κ) and the delta (δ) (Krumins et al., 1985; Ventura et al., 1989; Wittert et al., 1996; Zhang et al., 1996). However, the presence of the mu (μ) receptor is also suggested in human atrial trabeculae (Bell et al., 2000). There appears to be a converse relationship between the opioid receptors and the β -adrenergic pathway, in that when opioid receptors are inhibited, cardiac responsiveness to β -adrenergic stimulation is increased (Pepe et al., 1997). Conversely, when agonists of β -adrenergic receptors are used, the number and affinity of opioid receptors is increased (Pepe et al., 1997) which indicates a feedback mechanisms between these two systems. The protective effect of IPostC has recently been linked to the endogenous activation of the opioid receptors in a series of experiments by Zatta and colleagues (Zatta et al., 2008).

3.3.15 Preconditioning with opioids

Opioid release from ischaemic tissues appears to be involved in ischaemic preconditioning as well as in remote ischaemic preconditioning (Sadat, 2009). In a rat model, morphine, given as a preconditioning stimulus, decreased the infarct size in a similar manner to that of IPC and glibenclamide blocked the cardioprotection, implicating the involvement of the mK_{ATP} channel in this protective effect (Schultz et al., 1996). Other studies in the isolated rat heart model have demonstrated the ability of morphine to deliver the same degree of cardioprotection when administered just prior to reperfusion as it does when given prior to ischaemia (Gross et al., 2004). The protective effect of IPC in the canine heart and rat hearts can be mimicked by non-selective opioid agonists, selective κ -receptor agonists, or selective agonists for the δ -receptor, while μ specific agonists failed to induce protection (Headrick et al., 2003; Schultz et al., 1997; Schultz et al., 1996; Wang, 2001). Alternatively, opioid antagonists have been demonstrated to abrogate or partially abrogate the infarct-sparing effect of IPC (Chien and Van Winkle, 1996; Schultz et al., 1996; Wang, 2001).

3.3.16 Possible mechanisms in opioid-induced preconditioning

Preliminary evidence suggests that opioid preconditioning is mediated via endogenous opioid receptors, since the protective effects are inhibited by both naloxone, an opioid receptor antagonist, and methiodide, a peripheral acting naloxone derivative (Gross and Gross, 2006).

Isolated rat and mouse models of ischaemia-reperfusion were unable to be protected with a selective κ -opioid agonist in the presence of wortmannin, a PI3-kinase/Akt inhibitor and 5-HD, a blocker of the mK_{ATP} channel (given prior to reperfusion) thus implicating the activation of PI3-kinase/Akt and the mK_{ATP} channel in opioid preconditioning (Peart et al., 2008). Downey et al, have reported that opioid-induced protection occurs through a complex pathway which includes PI3-kinase/Akt, NOS, guanyl cyclase, PKG, opening of the mK_{ATP} and activation of PKC by redox signalling (see review (Downey et al., 2007)).

3.3.17 Postconditioning with opioids

Morphine can also protect the heart during reperfusion as shown in a study by Chen et al, in which morphine was given for the first 10 minutes of reperfusion in isolated rat hearts (Chen et al., 2008). This group demonstrated that morphine can mimic ischaemic postconditioning via activation of the κ -opioid receptor and the mK_{ATP} channel and they suggest that the δ -opioid receptor is not involved (Chen et al., 2008).

3.3.18 Possible mechanisms in opioid-induced cardioprotection

Morphine, one of the major opioids, appears to only exert a beneficial effect when administered during the ischaemic index or during early reperfusion (Gross, 2003). In contrast, the selective δ -opioid agonist, DADLE, mimicked the effects of ischaemic preconditioning in human atrial trabeculae (Tomai et al., 1999). A recent study in isolated rat hearts has provided evidence that opioids signal via the JAK/STAT pathway, with JAK2 mediating Akt activation, GSK-3 β and the inhibition of the mPTP (Gross et al., 2006). Another study in which morphine was administered just prior to reperfusion in the isolated rat heart, demonstrated a protective effect of morphine via inhibition of GSK-3 β (Gross et al., 2004). In 2009, Jinkun et al, reported that morphine prevented opening of the mPTP through a signalling pathway involving NO/cGMP/PKG/Zn²⁺/GSK-3 β in cardiomyocytes (Jinkun et al., 2009). In addition, an elegant experiment using human atrial trabeculae obtained from patients undergoing coronary artery bypass surgery revealed that the δ opioid receptor agonist, DADLE, appeared to protect human muscle from simulated ischaemia via the opening of the mK_{ATP} (Bell et al., 2000).

3.3.19 Clinical applications of the opioids

A very recent study in humans has reported a combined protective effect of morphine and remote IPostC when treatment was initiated directly following PCI in patients with acute myocardial infarction (Rentoukas et al., 2010).

4.0 Tumour necrosis factor alpha (TNF α)

The principle of an antihumoral response of the innate immune system in vivo was already recognised in 1893 by the physician, William B Coley. His inspiration came from the observation that when his cancer patients had an infection, they were more likely to recover from the cancer. Injecting his patients with a concoction containing the bacteria, *Streptococcus pyogenes*, reduced the tumours to a better extent when the patient developed a full-blown infection accompanied by a raging fever (Coley, 1893). This work was, however, not well documented and when Coley died in 1936, his treatment method died with him, having been replaced by radiation therapy. It is only in 1975 that Carswell and colleagues identified an endotoxin-induced serum factor that caused necrosis of tumours and they named this endotoxin “tumour necrosis factor alpha (TNF α)” (Carswell, 1975). Subsequently, a protein, named cachectin, was isolated from endotoxin-treated cells and was so named because it was presumed to play a role in the molecular basis of cachexia. Later on, cloning of TNF α and the protein, cachectin, confirmed that they were identical (Beutler et al., 1985). This soluble, multifaceted cytokine is produced upon activation of the innate immune system, mainly in macrophages but subsequent studies have shown that many other cells including cardiomyocytes, also express TNF α (Benigni et al., 1996; Frangogiannis et al., 1998). Two isoforms of TNF exist, TNF α and TNF β . TNF α is first expressed as a transmembrane protein and subsequent studies have provided evidence that membrane TNF α is converted to a soluble homotrimeric form by proteolytic cleavage, a process known as ectodomain shedding, allowing this cytokine to have more widespread effects (Beutler et al., 1985).

Studies in various culture systems have reported ADAM17 (a disintegrin and metalloproteinase 17) as a sheddase of TNF α and its receptors (Black et al., 1997; Moss et al., 1997; Peschon et al., 1998; Reddy et al., 2000). In 2007, Bell et al demonstrated that ADAM17 plays a primary role in TNF α , tumour necrosis factor receptor 1 (TNFR1) and tumour necrosis factor receptor 2 (TNFR2) shedding. However, deficiency of ADAM17 did not abrogate the shedding, suggesting that ADAM17 facilitates the shedding but is not a crucial element for shedding to occur (Bell et al., 2007). ADAM17 is also known as TACE (TNF alpha converting enzyme). An important determinant of shedding activity is thought to be the localisation of ADAM17 to a perinuclear compartment which raises the possibility that the intracellular environment may also be implicated in ADAM17-mediated ectodomain shedding (Peiretti, 2003).

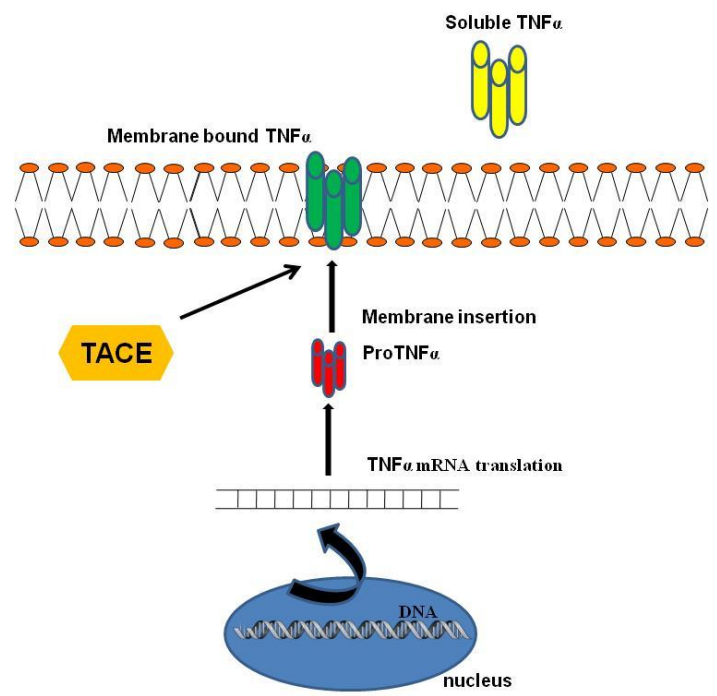


Fig 17: Production and release of TNF α : TNF α is first synthesised as a transmembrane proTNF α , allowing TNF α to exist in its membrane bound form. TACE then cleaves ProTNF α , releasing the soluble form of TNF α . TACE=TNF α converting enzyme; mRNA=messenger ribonucleic acid, (adapted from Meldrum, 1998).

4.1 TNF α receptors

Once released from the cell, TNF α docks with one of two TNF α receptors – a 55 kDa high affinity receptor, known as TNFR1 and a 75 kDa low affinity receptor, TNFR2 (Beutler, 1994; Vassilli, 1992). TNF ligand is believed to achieve all its different cellular and pathophysiological

effects by its binding to either the TNFR1 or TNFR2 receptor subtype. Like the ligand, the two receptors can undergo ectodomain shedding, generating soluble, functional receptors.

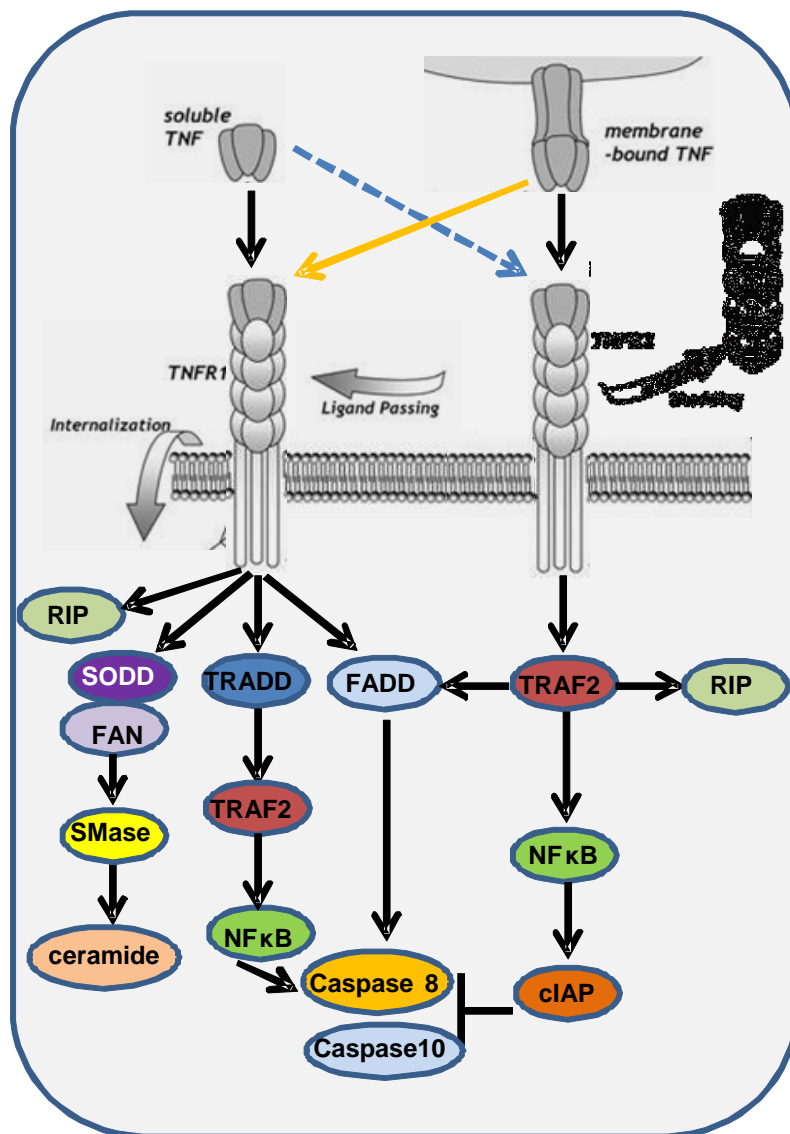


Fig 18: Simplified diagram of some of the signalling pathways modulated by TNFR1 and TNFR2 subtypes, (adapted from MacEwan, 2002).

TNFR1 contains a death domain (DD) motif which is critical in the death-inducing activity of this receptor (Tartaglia et al., 1993). A number of associating proteins and related molecules also contain a DD and signalling for cell death appears to be the function of these proteins and molecules. The silencer of death domain (SODD) is one such molecule and it binds to the DD in TNFR1 preventing the binding of other DD proteins (Jiang et al., 1999). When TNFR1 is stimulated by TNF α , the SODD dissociates from the DD and this allows access to the DD for

other activator proteins. An important element in the activation of neutral sphingomyelinase is an adaptor protein called factor associated with neutral sphingomyelinase activation (FAN) (Adam et al., 1996; Adamklages, 1998). This adaptor protein is responsible for the generation of smaller sphingolipid molecules containing ceramide which play a key role in signalling (Kolesnick and Kronke, 1998). Both TNFR1 and TNFR-associated death domain (TRADD) are found at the perinuclear-Golgi complex but TRADD only associates with TNFR1 once at the plasma membrane (Jones et al., 1999; Ledgerwood et al., 1999). It then recruits tumour necrosis factor receptor-associated factor (TRAF2) to TNFR1 (Hsu et al., 1995). TRAF2, in turn, recruits nuclear factor kappa B (NF κ B) (Vince et al., 2009). However, in addition to promoting NF κ B signalling, TRAF2 also has anti-apoptotic signalling roles, with the ring domain of TRAF2 being required for protection against TNF-induced apoptosis (Vince et al., 2009).

In addition, TRADD recruits the downstream signalling adaptor molecules, Fas-associated death domain (FADD) and RIP (receptor interacting protein). The role of RIP remains uncertain at this point. FADD contains a death effector domain (DED) that interacts with caspase 8 and a number of other molecules that regulate cell death mechanisms (Muzio et al., 1996) (Figure 18).

TNFR2 does not have a DD motif but still recruits adaptor proteins including TRAF2. The structural and functional characterization of TNFR2 has not, to date, been studied in depth. However, the membrane-bound form of TNF α is suggested to be a potent activator of TNFR2, in contrast to the soluble form, which has a greater affinity for TNFR1 (Grell et al., 1995). In the immune system, TNF α would activate its receptors by cell – cell interactions. How the binding characteristics of membrane-bound TNF α at TNFR1 and TNFR2 compare to soluble TNF α still remains to be elucidated. Minor changes in the structure of the TNF α sequence can lead to major changes in its binding characteristics and therefore determine to which receptor it will bind (Lewis et al., 1991). Lewis et al, demonstrated that TNFR1 and TNFR2 are equally activated by murine TNF α , whereas human TNF α only stimulates TNFR1 and has no effect on TNFR2 in the mouse (Lewis et al., 1991). It has been suggested that most of the *in vivo* effects of TNF α may be mediated by membrane-bound TNF α with both receptors being activated, rather than the soluble form which would activate TNFR1 more robustly than it does TNFR2 (MacEwan, 2002). If this is so, than the physiological role of TNFR2 may be more important than first thought. TRAF2 has a direct affinity for TNFR2. Under certain conditions, FADD and RIP are also thought to be able to bind indirectly to TNFR2 via TRAF2. TRAF2 can also interact with NF κ B which controls the transcription of a gene called cellular inhibitor of apoptosis protein 2 (c-IAP2). cIAP2 binds to TRAF2 thereby inhibiting the activation of caspases 8 and 10 and

subsequent apoptosis (MacEwan, 2002). This could provide a platform for the beneficial effects of TNF α (Figure 18).

4.2 TNF α and the heart

In physiological conditions, the primary role of TNF is the regulation of immune cells. As TNF α is involved in cellular differentiation, growth and apoptosis (Sack et al., 2000), it is relevant to consider whether this pluripotent cytokine is involved in normal cardiac development. The cardiac phenotype of the TNF α -deficient mice has only minor differences in cardiac mass compared with wildtype littermate controls (Smith et al., 2001) and cardiac studies in mice deficient in either of the TNF receptors showed no discernable congenital cardiac defects (Kurrelmeyer et al., 2000). These data, taken together, imply that normal cardiac morphogenesis does not require TNF α signalling, or that alternate signalling pathways compensate for lack of TNF α .

TNF α concentration is low in healthy hearts, where it is located mainly in endothelium and resident mast cells (Schulz, 2008). The discovery in the 1990s that TNF α is expressed in all nucleated cell types of the myocardium, including the cardiomyocyte itself, renewed interest in TNF α and its role in cardiac pathophysiological states such as myocardial infarction and heart failure (Levine, 1990; Herskowitz, 1995; Dorge, 2002).

Myocardial TNF α is expressed in direct response to many different forms of cardiac injury, including acute myocardial infarction (Mann, 2003; Yokoyama, 1997; Maury and Teppo, 1989). Elevated plasma levels of TNF α have also been reported in several studies conducted in heart failure patients (see review (Batista et al., 2009)). Transgenic mice, designed to overexpress cardiac TNF α have highlighted the maladaptive effects of TNF α in the heart. While wildtype counterparts remained healthy, the mice overexpressing TNF α developed myocarditis (Kubota et al., 1997).

Both human and rat cardiomyocytes possess the two TNF α cell surface receptors, TNFR1 and TNFR2. (Krown, 1995; Torre-Amione *et al.*, 1995). Recently, the existence of a mitochondrial binding protein for TNF α has been proposed, therefore suggesting that TNF α may be delivered directly from the cell surface to the mitochondria independently of TNFR1 and TNFR2 (Busquets, 2003; Ledgerwood, 1998).

4.3 TNF α and ischaemia reperfusion

Many studies have documented myocardial TNF α production during acute myocardial ischaemia with or without reperfusion (Irwin et al., 1999; Meldrum et al., 1998a; Meldrum et al., 1998b).

Preformed TNF α is released within minutes from the resident mast cells and macrophages during myocardial ischaemia, but with persistent ischaemia, TNF α also originates from the cardiomyocytes (Dorge et al., 2002) and is also one of the multi-potent cytokines known to be released by ischaemia-reperfusion (Sharma and Das, 1997). TNF α levels have also been demonstrated to be increased at the onset of reperfusion in an ischaemia-reperfusion model (Cain et al., 1998) and other studies revealed that in adenosine mediated cardioprotection this upregulation of TNF α was prevented (Cain et al., 1998; Meldrum et al., 1997).

Initially, this cytokine was regarded as only having deleterious effects, contributing to myocardial dysfunction and cardiomyocyte death in ischaemia-reperfusion injury (Giroir, 1994; Latini, 1994; Levine, 1990). In a study conducted in mice, Maekawa et al, demonstrated that once reperfusion was instituted, a deficiency in TNF α signalling resulted in smaller infarct sizes (Maekawa et al., 2000). Confirmation of this putative detrimental effect of TNF α was forthcoming in a study conducted in wildtype mice using an identical ischaemia-reperfusion protocol in the presence of anti-TNF antibodies (Belosjorow et al., 2003; Gurevitch et al., 1997; Maekawa et al., 2002).

In contrast to the detrimental effects of TNF α , it was reported that cultured cardiomyocytes are protected against hypoxic injury when exogenous TNF α is added (Nakano et al, 1998) while larger myocardial infarct sizes were observed in mice deficient in both TNF receptors compared to their wildtype littermate counterparts following an acute myocardial infarction (Kurrelmeyer et al., 2000). Both these studies implicated TNF α in cardioprotection. A plausible explanation for these opposing roles of TNF α may be the existence of an intrinsic cell survival programme in response to ischaemia which is abrogated by the inflammatory response at reperfusion.

The concept, therefore, that selective, rather than total, inhibition of TNF α may be more feasible (Abraham et al., 1995) was further explored in 2002 by Sack, who proposed that a low acute concentration of TNF α would be protective whereas a high and sustained concentration would be maladaptive (Sack, 2002).

In humans, TNFR1 has been implicated in the initiation of the negative inotropic effects of TNF α in adult cardiomyocytes (Torre-Amione et al., 1995). In addition, Monden et al, demonstrated that TNF α was toxic via TNFR1 but was protective via TNFR2 in a murine model of myocardial infarction (Monden et al., 2007). Although administration of either a soluble TNF receptor or an anti-TNF α antibody abrogates the development of heart failure in experimental animals, paradoxically, the use of anti-cytokines in attempts to neutralize TNF α has not proved successful in patients with chronic heart failure (Anker and Coats, 2002; Mann, 2002).

Higuchi et al, bred TNFR1^{-/-} and TNFR2^{-/-} mice with cardiac-specific overexpression of TNF α and found that the absence of TNFR2 exacerbates heart failure and reduces survival whereas the removal of TNFR1 blunted heart failure and improved survival, therefore suggesting that TNFR2 signalling played a protective role in the pathogenesis of heart failure (Higuchi et al., 2004). Additionally, deletion of TNFR1 and TNFR2 in mice subjected to myocardial infarction demonstrated that inhibition of TNFR1-mediated pathways attenuated ventricular dysfunction and improved survival with a downregulation of other proinflammatory cytokines, whereas inhibition of TNFR2 was deleterious, exacerbating ventricular dysfunction accompanied by upregulation of TNF α and other proinflammatory cytokines (Monden et al., 2007).

Thus, it is appropriate to suggest that the role played by TNF α in ischaemia-reperfusion may be dependent on which TNF receptor is activated and the concentration of TNF α as well as the time it is present in the system (Deuchar et al., 2007; Flaherty et al., 2008; Lecour et al., 2002; Monden et al., 2007; Schulz, 2008). The function of TNF α in ischaemia-reperfusion remains controversial, with this pleiotropic cytokine's ability to act in both maladaptive and adaptive roles (Figure 19).

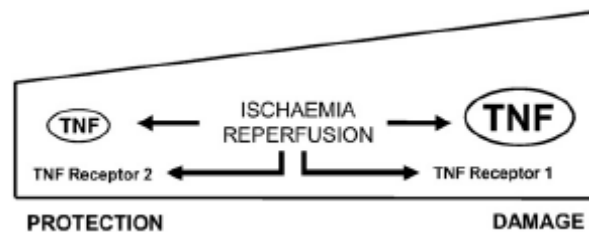


Fig 19: TNF α in myocardial ischaemia-reperfusion: protective versus damaging effects, (Lecour, 2009).

4.4 TNF α and ischaemic preconditioning

Persuasive evidence exists for a role of TNF α in ischaemic preconditioning. In 2002, Smith et al, used mice in which endogenous TNF α production was knocked out to demonstrate that the lack of TNF α precluded the protective effect of IPC. However, pharmacological preconditioning with adenosine or diazoxide was still able to reduce infarct size (Smith et al., 2002). Subsequent studies also demonstrated a lack of protection in mice in which TNF α had been knocked out (Ichikawa et al., 2004; Labruto et al., 2005). Pre-administration of TNF α , 24 hours prior to TNF α preconditioning, rescued the cardioprotective effect in TNF knockout mice, suggesting that the signalling effector system downstream of TNF α may be downregulated in TNF α

deficient mice (Smith et al., 2002). Transient low concentrations of TNF α given as a preconditioning mimetic prior to the ischaemic insult protected the heart against ischaemia-reperfusion in the isolated rat heart model (Lecour et al., 2005a; Lecour et al., 2002). In rats, mice or rabbits, classic ischaemic preconditioning was associated with increased levels of pre-ischaemic TNF α (Belosjorow et al., 1999; Smith et al., 2002) but a decreased level of endogenous TNF α was released during the ischaemia-reperfusion injury phase (Belosjorow et al., 1999; Kimura et al., 2006). Interestingly, preconditioning-induced cardioprotection with exogenous TNF α can only be achieved with a washout period prior to the sustained ischaemia, therefore suggesting that the time of TNF α production is another factor to be considered (Lecour et al., 2002).

4.5 TNF α and ischaemic postconditioning

Anaesthetized rats subjected to coronary artery ligation demonstrated that IPostC reduced levels of plasma TNF α during the reperfusion phase in a manner similar to that of IPC (Kin, 2008). However, it is presently unknown whether TNF α is involved in the protective effect of ischaemic postconditioning and the main focus of this thesis was to delineate the role of TNF α as a postconditioning mimetic.

4.6 Proposed mechanisms in TNF α -induced cardioprotection

In a manner similar to IPC, TNF α -induced preconditioning activated the sphingolipid pathway, PKC and the mK_{ATP} channel (Lecour et al., 2002). The inhibitors of these various pathways have been demonstrated to abolish the protection afforded with TNF α when given during the preconditioning stimulus (Lecour et al., 2002). In addition, the free radical scavenger, mercaptopropionyl glycine (MPG) also blocked the infarct-sparing effect of TNF α , implicating ROS in the prosurvival signalling cascade (Lacerda et al., 2006; Lecour et al., 2005a). One of the triggers for IPC is myocardial p38 MAPK which can be activated by TNF α . However, Tanno et al, demonstrated that TNF α -induced protection in the murine heart is independent to p38 MAPK activation (Tanno et al., 2003). Critical targets for downstream protection are activation of the mPTP (Gao et al., 2005), the Ca²⁺ activated potassium channel (Gao et al., 2005) and the phosphorylation of NF κ B (Somers et al., 2005).

4.7 TNF α -induced cardioprotection is independent of the RISK pathway

Phosphorylation and activation of PI3-kinase and Erk1/2 at the time of reperfusion in both ischaemic pre-and postconditioning has been shown to protect against reperfusion injury as part

of the RISK pathway (see review (Hausenloy et al., 2005)). However, strong evidence is emerging in the literature in support of the hypothesis that protection against reperfusion injury does not always require activation of the RISK pathway. The protective effect of TNF α in isolated rat hearts could not be abolished by the Erk1/2 inhibitor, PD98059, given during the preconditioning stimulus (Lecour, 2005b;). Similarly, administration of wortmannin, an inhibitor of PI3-kinase/Akt during the preconditioning stimulus did not abrogate the protective effect of TNF α in a study conducted in isolated cardiomyocytes and the isolated rat heart (Suleman, 2008).

4.8 TNF α initiates the SAFE pathway via JAK/STAT-3 signalling

Cardiac-specific STAT-3 knockout mice could not be protected by the preconditioning mimetics, adenosine, diazoxide and TNF α administered before the index ischaemia (Smith, 2004). A study by Lecour et al, provided evidence that both TNF α and early IPC phosphorylated STAT-3 in the early phase of reperfusion in the isolated rat heart model and that AG490 administered at the time of reperfusion, abrogated both IPC and TNF α -induced protection, thus linking TNF α and STAT-3 in a prosurvival signalling pathway (Lecour et al., 2005b). In addition, AG490, the JAK/STAT antagonist, administered to isolated hearts from STAT-3^{-/-} mice before the preconditioning stimulus, was shown to prevent phosphorylation of Akt (Suleman, 2008).

The activation of STAT-3 was demonstrated to co-incide with the activation of the RISK pathway during ischaemic preconditioning (Lecour et al., 2005b) but interestingly, TNF α -induced protection failed to activate the RISK pathway. This, therefore, implied that the JAK/STAT pathway may be activated independently of the RISK pathway.

The role of TNF α and its cognate receptors has not been extensively studied in the postconditioning setting although there is now a growing body of evidence implicating the signalling of the innate immune response to the events underlying myocardial adaptation to ischaemia (Valeur and Valen, 2009). However, it should be emphasized that there is very little known regarding TNF α and its effects during the first few critical minutes of reperfusion. Many studies have examined the effect of TNF α when given just before the ischaemic insult (preconditioning), but to date, no studies could be found in the literature which investigated the protective role of exogenous TNF α at reperfusion or the possible protective role of the TNF α /JAK/STAT signalling cascade in postconditioning. Only one study, to date, has reported that serum and myocardium levels of TNF α were demonstrated to be decreased at reperfusion in a rat model, resulting in protection for the heart by reducing apoptosis and preventing excess

ROS production (Kin, 2008), thus implicating a low concentration of TNF α in the protective effect of postconditioning. A schematic depicting the activation of the RISK and SAFE pathways following ischaemia-reperfusion is shown in figure 20.

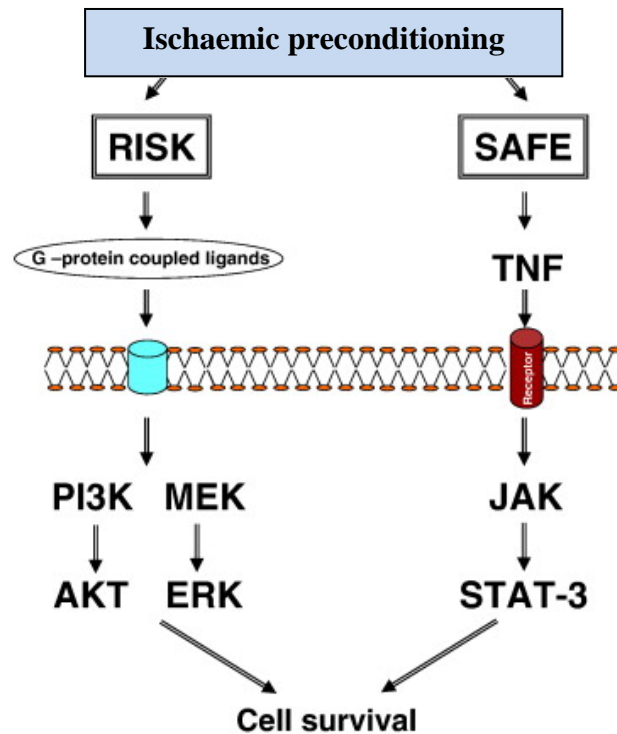


Fig 20: Activation of the RISK and SAFE pathways following ischaemia-reperfusion. Ischaemic preconditioning increases TNF α levels resulting in TNF α binding to its cell surface receptors activating the JAK/STAT pathway as part of the SAFE pathway. Similarly, activation of PI3-kinase and MEK activate Akt and Erk1/2 respectively, as part of the RISK pathway. Both pathways enhance cardioprotection against ischaemia-reperfusion. Abbreviations as defined in text, (adapted from Lecour, 2009).

4.9 TNF α , obesity and diabetes

TNF α is a multifaceted cytokine that exerts potent pro-inflammatory effects in metabolic and inflammatory disorders such as obesity and insulin resistance. An important fact is that obesity does not occur in isolation from other metabolic disorders. Chronic activation of the innate immune system is a feature of obesity which is associated with a low grade inflammation of white adipose tissue (WAT). Subsequently, this can lead to insulin resistance, impaired glucose tolerance and even diabetes (Bastard et al., 2006). The first detectable defect in type 2 diabetes is insulin resistance when insulin levels are greatly increased while glucose levels remain normal (Hotamisligil et al., 1995).

TNF α down-regulates insulin-stimulated glucose uptake by its effect on glucose transporter 4, insulin receptor autophosphorylation and insulin receptor substrate-1. TNF α is a powerful autocrine and paracrine regulator of adipose tissue. It can inhibit insulin action and insulin receptor signalling and this is thought to be mainly mediated by TNFR1 (Kanety et al., 1996; Liu et al., 1998a; Peraldi and Spiegelman, 1997). Investigators have also examined whether the TNF receptors play a role in the development of glucose homeostasis. These studies concluded that the receptors did not play a role in the induction of insulin resistance associated with obesity, but that the presence of both receptors may be required for normal glucose homeostasis (Schreyer et al., 1998). However, TNF α appears to have no significant effect on the development of dietary obesity, as the development of marked obesity was noted in mice which lack endogenous TNF α (TNF $^{-/-}$) and were fed a high fat, high caloric diet (Uysal et al., 1997). These mice remained highly insulin sensitive compared to their wild type counterparts. Furthermore, the insulin concentrations in obese TNF $^{-/-}$ mice were significantly lower than those of the obese TNF-WT and they remained at the same level as that of lean mice on a normal diet (Uysal et al., 1997). These results demonstrated that insulin resistance associated with dietary obesity is reduced in the absence of TNF α . Interestingly, genetic depletion of TNF α from macrophages in wild type mice does not protect these mice from developing obesity-related insulin resistance (De Taeye et al., 2007).

TNF α has also been linked to the development of type 1 diabetes (Yang et al., 1994). Treatment with non-toxic doses of TNF α in neonatal non-obese diabetic mice for 21 – 24 days resulted in the onset of type 1 diabetes being accelerated, whereas inhibition of TNF α for the same time period resulted in a complete absence of diabetes (Yang et al., 1994).

Interestingly, the cardioprotective effect of ischaemic preconditioning has been shown to be abolished in human diabetic patients (Ishihara et al., 2001). Experimental animal models have also indicated that diabetes abolishes the ability of IPC and pharmacological agents to reduce infarct size, but the mechanisms involved in this failure are unclear (Kersten et al., 2001; Kersten et al., 2000; Nawata et al., 2002; Paulson, 1997; Tosaki et al., 1996).

Cachexia, the wasting disorder of muscle and fat tissues, has previously been associated with TNF α expression, (Beutler and Cerami, 1989; Oliff et al., 1987; Torti et al., 1985) which is in contradiction to the possible link between TNF α and obesity. One explanation is the circulating levels of TNF α . These exceed 200 ng/ml in cases of cachexia whereas, in models of obesity-diabetes the circulating levels are very low, in the range of 10-200 pg/ml when detectable (Hotamisligil, 1993). Another explanation is that obesity is often associated with other hormonal abnormalities which may alter the body's response to TNF α . It has even been suggested that the

site of expression could determine the metabolic response to TNF α (Tracey, 1992). More investigative studies will be needed to elucidate how one molecule can have such diverse effects on a vast array of different pathophysiological states.

Obesity, diabetes and metabolic syndrome all alter the production of adipokines, such as leptin and TNF α . Hotamisligil and colleagues demonstrated that in human obesity, the expression of TNF α in adipose tissue was significantly elevated and that this increase in TNF α was strongly correlated with the level of hyperinsulinaemia (Hotamisligil et al., 1995).

4.10 TNF α and leptin

Leptin is an adipocyte secreted hormone involved in energy regulation and endocrine and immune function. It plays a role in innate and acquired immunity.

Both the structure of leptin and of its receptor suggests that leptin can be classified as a cytokine. Known as the “appetite suppressing hormone”, there is a strong correlation between serum leptin and TNF α levels (Margalet, 2003). Additionally, there is a strong association between circulating leptin levels and body mass index (BMI) as well as fat mass in obesity with TNF α upregulating leptin production and leptin also modulating the production of TNF α (Bastard et al., 2006). As a function of its cytokine-like structure, therefore, leptin could exert peripheral biological effects (Almanza-Perez et al., 2008).

It has been demonstrated that leptin expression is increased with administration of TNF α in rodents (Grunfeld, 1996; Sarraf, 1997). In a study conducted in humans, a positive and independent correlation between circulating levels of soluble TNFR1 and leptin was demonstrated, therefore suggesting an association between TNF α and leptin in humans similar to that seen in rodents where TNF α increases leptin expression and circulating leptin levels (Mantzoros, 1997). This possible effect of an activated TNF α system on blood leptin levels in humans could have a considerable impact on human physiology.

It has been previously reported that leptin protects the isolated mouse myocardium against ischaemia-reperfusion injury via direct action on the heart (Smith, 2006) and it is known that the JAK/STAT pathway mediates many of the metabolic actions of leptin (Frubeck, 2006). Recent evidence from isolated rat hearts demonstrated a reduction in infarct size when leptin was given at reperfusion. This cardioprotective effect was abrogated by co-administration of the JAK/STAT inhibitor, AG490, therefore implicating the JAK/STAT pathway in leptin-induced cardioprotection (Smith, 2010).

B. AIM, HYPOTHESIS AND OBJECTIVES

1.0 Aim

Ischaemic postconditioning, whereby alternating cycles of reperfusion and ischaemia are applied at the onset of reperfusion, is a powerful tool to reduce infarct size and preserve cardiac function. Delineating the exact mechanism involved in this phenomenon may lead to the development of novel therapies against the burden of ischaemic heart disease. A number of different signal transduction pathways have been reported to underlie the cardioprotective effect of IPostC. One of first signalling cascades to be linked to IPostC was the Reperfusion Injury Salvage Kinase (RISK) pathway which implicated the prosurvival kinases, phosphatidylinositol 3-kinase (PI3-kinase)-Akt and p44/p42 extracellular signal-regulated kinases (Erk1/2) in the cardioprotective effect of IPostC. The activation of these prosurvival kinases at the immediate onset of reperfusion by a diverse variety of agents such as adenosine, bradykinin and the opioids has been shown to reduce myocardial infarct size in the region of 40-50%. However, the mechanism by which these survival pathways confer cardioprotection is poorly understood.

TNF α is a cytokine implicated in inflammation, growth and apoptosis but previous work in our laboratory has demonstrated that TNF α can also protect against ischaemia-reperfusion in a dose and time dependent manner.

Therefore, the aim of the present study was to explore the role of TNF α in ischaemic and pharmacological postconditioning-induced cardioprotection.

1.1 Hypothesis and Objectives

We hypothesized that activation of TNF α is required in both IPostC and pharmacological postconditioning to confer cardioprotection and that this protective effect is mediated via the activation of TNFR2 and the transcription factor STAT-3 (Signal Transducer and Activator of Transcription). Furthermore, we investigated ischaemic postconditioning in obese and diabetic animals, two major risk factors for ischaemic heart disease that are associated with elevated levels of circulating TNF α (figure 21).

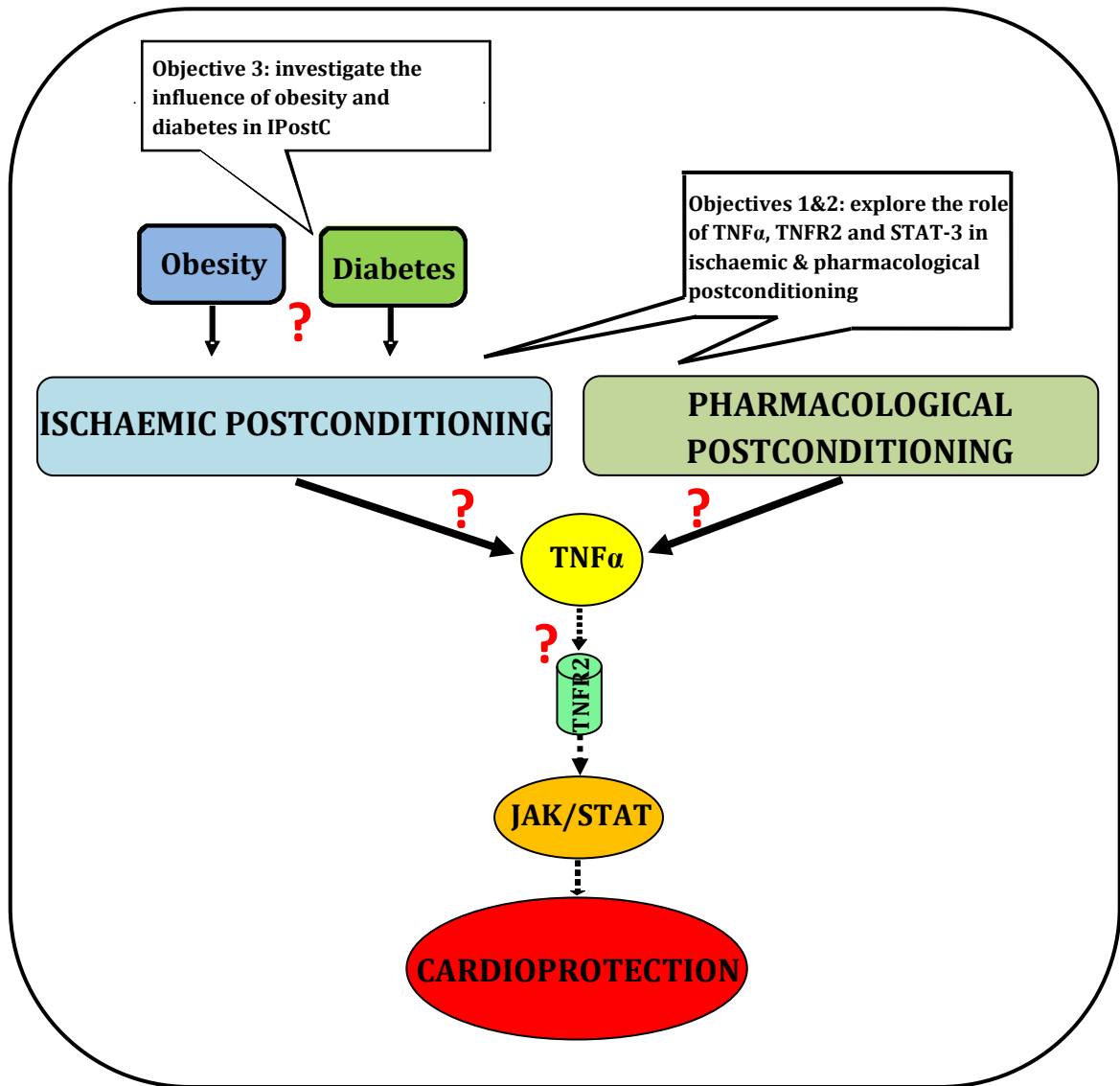


Fig 21: Schematic representation of hypothesis and objectives in ischaemic and pharmacological postconditioning. Ischaemic postconditioning activates TNF α which then binds to its TNFR2 with subsequent activation of the JAKs. This results in the dimerization of STAT-3 and its translocation to the nucleus, with the mPTP of the mitochondria being a putative end effector. Abbreviations as defined in the text.

To fulfil this aim, the following objectives will be pursued:

1) *To investigate the role of TNF α and its receptors in IPostC-induced cardioprotection.*

To achieve this objective we will use TNF deficient (TNF^{-/-}), TNFR1 deficient (TNFR1^{-/-}), TNFR2 deficient (TNFR2^{-/-}), cardiac-specific STAT-3 deficient (STAT-3^{-/-}) mice and their respective wild types. Isolated hearts will be postconditioned following an ischaemia-reperfusion insult. Should TNF α prove to be necessary in IPostC-induced cardioprotection, we would then use inhibitors of various pathways and the western blot technique to explore possible downstream targets of TNF α such as GSK-3 β , RAGE, PAR-2, Akt, Erk and STAT-3.

2) *To investigate the role of TNF α in pharmacological postconditioning.*

To achieve this objective we will use TNF deficient (TNF^{-/-}), TNFR1 deficient (TNFR1^{-/-}), TNFR2 deficient (TNFR2^{-/-}) and cardiac-specific STAT-3 deficient mice (STAT-3^{-/-}) and their respective wild types. Isolated hearts will be subjected to an ischaemia-reperfusion insult and the well-known cardioprotective agents, adenosine, bradykinin and opioids will be used as pharmacological postconditioning mimetics. Should TNF α prove to be necessary in pharmacologically-induced cardioprotection, the western blot technique will be used to explore possible downstream targets such as STAT-3.

3) *To investigate the effect of obesity and diabetes in IPostC-induced cardioprotection.*

For the obesity study, the mice will be fed a high carbohydrate diet from weaning for a total of 11 weeks. Thereafter, the isolated hearts of TNF-WT or TNF^{-/-} mice will be subjected to IPostC and infarct size will be used as the endpoint. To explore the role of diabetes in IPostC, streptozotocin will be administered to the mice to initiate onset of diabetes. At day 5 and at day 10 post streptozotocin, the isolated hearts will be subjected to IPostC and infarct size used as the endpoint. Various parameters such as body weight, heart to body weight ratio, blood glucose, plasma leptin and plasma insulin will be examined in the obese and diabetic mice.

C. MODELS & METHODS

1.0 Genetically modified animals

The best animal models to achieve our objectives were whole body TNF-deficient mice which produce no endogenous TNF α and TNF receptor1 and 2-deficient mice. A third mouse model, cardiac-specific STAT-3 deficient mice were used to determine whether the JAK/STAT pathway was involved in the protective signalling.

In this thesis all animal studies performed were approved by the Animal Research Ethics Committee of the University of Cape Town, and followed the recommendations laid down in the Guide for the Care and Use of Laboratory Animals published by the US National Institute of Health (NIH Publication No. 85 {23}, revised 1996).

1.1 Mouse Models to investigate the role of TNF α in ischaemic postconditioning

Homozygous TNF-deficient (TNF^{-/-}) mice, TNF receptor 1-deficient mice (TNFR1^{-/-}) and TNF receptor 2-deficient mice (TNFR2^{-/-}) and their respective wildtype, Black 6 x 129S (TNF-WT) were a generous gift from Dr Muazzam Jacobs and Professor Bernhard Ryffel, Department of Immunology, University of Cape Town.

Cardiomyocyte-specific STAT-3-deficient (STAT-3^{-/-}) mice from C57 Black 6 background were created in our facilities by crossing homozygous floxed STAT-3 mice with heterozygous MLC2V-driven Cre recombinase mice, as previously described (Smith et al., 2004). All mice were 12-14 weeks of age.

1.1.2 Genotyping of Mouse Models

To distinguish wildtype from TNF^{-/-}, TNFR1^{-/-}, TNFR2^{-/-} and STAT-3^{-/-} mice, tail cuts were performed at weaning (3 weeks of age) and genomic DNA extracted from the respective tails. Genotyping was performed by means of the polymerase chain reaction (PCR).

i) DNA Extraction

Approximately 1 cm of a candidate mouse tail was removed from the mouse and digested by incubating at 55°C overnight in 400 μ l tail digestion buffer (0.5M Tris {pH 8}, 0.1M EDTA, 1% SDS) with 1 mg/ml proteinase K (Roche, Mannheim) added to the buffer immediately prior to use. The following day, 400 μ l of Phenol:Chloroform:Alcohol (25:24:1)(Sigma, USA) mixture was added to each digested tail and the samples were gently inverted to mix. Cellular debris was removed by centrifugation at 13,500 rpm (Heraeus MK202, Germany) for 10 minutes

at room temperature. The bottom layer containing cellular debris was carefully removed with sterile barrier tips taking care not to disturb the interface containing contaminating proteins. 200 μ l of chloroform (Sigma, USA) was added to the remaining upper layer containing the required DNA to remove traces of phenol and the tube was again gently inverted to mix. The digests were centrifuged again at 13,500 rpm for 5 minutes. Two separate phases again occurred and the bottom phase was removed and discarded while the upper phase is once more centrifuged at 13,000 rpm for 10 minutes, forcing the remaining debris into a pellet at the bottom of the tube. The liquid phase was then transferred to a new sterile 1.5 ml eppendorff tube and 50 μ l of 7.5 M ammonium acetate (Merck) and 1 ml 100% ethanol added to each tube. The tubes were gently inverted to mix and the DNA, which appears as a white, stringy precipitate, was transferred to a clean eppendorff tube containing 500 μ l of 70% ethanol to hydrate the DNA. The samples were then centrifuged at 13,000 rpm for 10 minutes and the ethanol carefully decanted to prevent loss of the DNA pellet. DNA pellets were air dried until ethanol had completely evaporated, then 50 – 75 μ l of ultra pure RNase free water was added to dissolve the DNA. Thereafter samples were stored at -20°C until PCR analysis.

ii) Polymerase Chain Reaction (PCR)

Targeted deletion of the TNF α gene has been previously described (Marino et al., 1997). Genotyping was carried out, at random intervals of breeding, to confirm the genetic background of each species of mouse model and, in the case of TNF $^{-/-}$, TNFR1 $^{-/-}$ and TNFR2 $^{-/-}$ mice, involved testing as to whether the respective mouse's TNF α , TNFR1 or TNFR2 gene was intact or had been disrupted by the insertion of a phosphoglycerate kinase-neomycin expression cassette within the gene. Genotyping of the STAT-3 mice had to be performed at each stage of the breeding process as breeding of STAT-3 $^{-/-}$ mice gave rise to both the wild type and the STAT-3 deficient mice. Therefore, these mice had to be tested to see whether or not they were harbouring the Cre recombinase insertion as well as whether or not they were floxed.

iii) Sequences and PCR for TNF $^{-/-}$, TNFR1 $^{-/-}$ and TNFR2 $^{-/-}$ mouse models

a) TNF $^{-/-}$: Sense primer: 5'-CTA AAT GAA CTG CAG GAC GA-3'

Antisense primer: 5'-ATA CTT TCT CGG CAG GAG CA-3

Neomycin: Sense primer: 5' CTG AAT GAA CTG CAG GAC GA 3'

Antisense primer: **5' ATA CTT TCT CGG CAG GAG CA 3'**

The polymerase chain reactions were performed in 50 µl volumes made up as follows:

10xMagnesium free buffer:	10 µl
4mM MgCl ₂ :	8 µl
100µM dNTP's:	1.6 µl
Primer (TNF or Neo):	0.5 µl
β actin 3':	0.5 µl
β actin 5':	0.5 µl
H ₂ O:	26.9 µl
Taq:	0.5 µl
DNA sample:	<u>1.5 µl</u>
	50 .0 µl

PCR Conditions:

TNF:

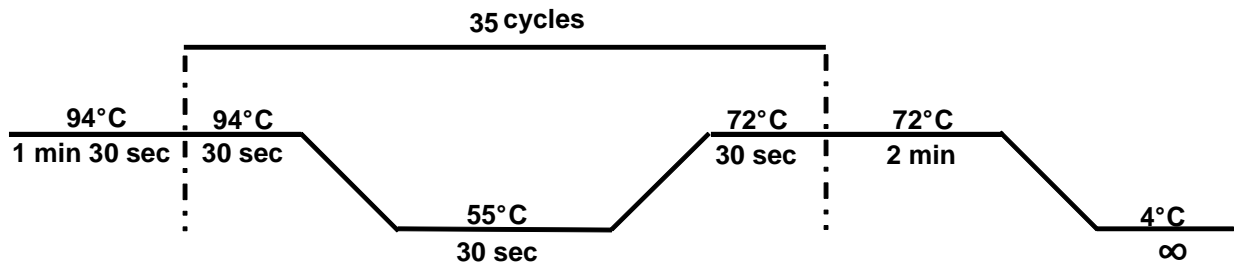


Fig 22: PCR conditions for TNF genotyping

Neo

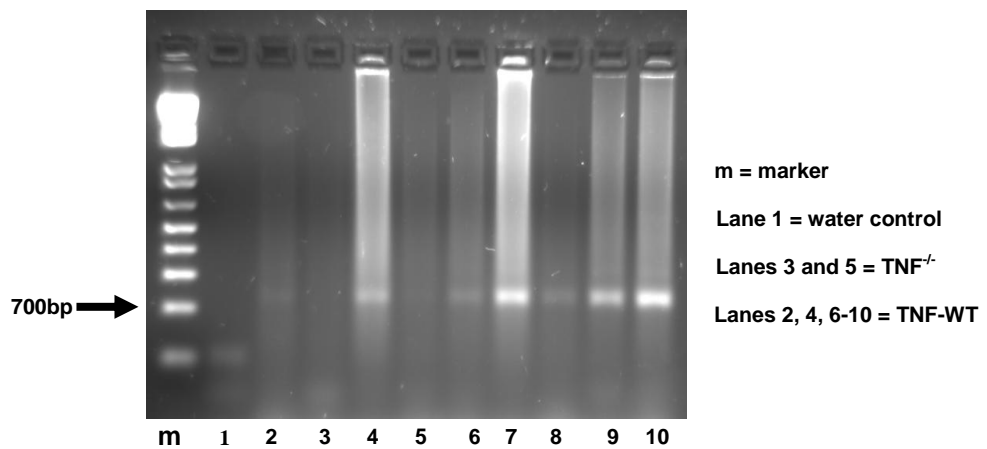
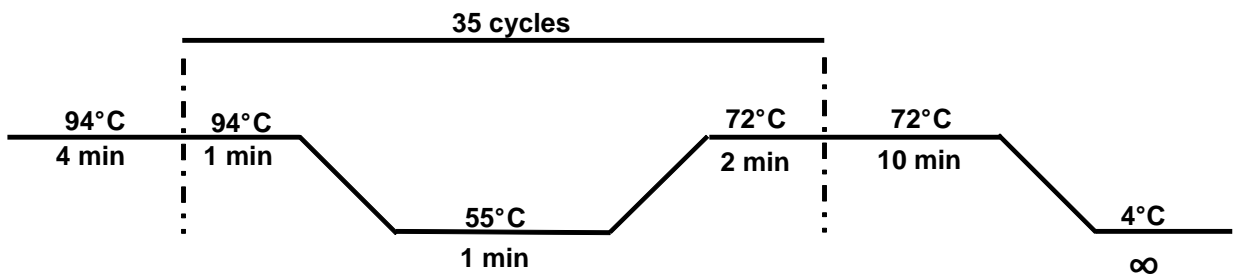


Fig 23: UV gel analysis of extracted mouse genomic DNA for the targeted disruption of the TNF α gene

b) **TNFR1^{-/-}**: TNFR1 - 4938: **5' AGA AAT CTC AG ACA ATT CTC TGC**

TNFR1 – 2883: **5' CTC TCT TGT GAT CAG CAC TG**

TNFR1 – Neo: **5' TCC CGC TTC AGT GAC AAC GTC**

The polymerase chain reactions were performed in 50 µl volumes made up as follows:

10xMagnesium free buffer:	5 µl
25 mM MgCl ₂ :	3 µl
10 mM dNTP's:	1 µl
Primers: TNFR ₁ 2883	1 µl
TNFR ₁ 4938	1 µl
TNFR ₁ Neo 34	1 µl
H ₂ O:	35.875 µl
Taq:	0.125 µl
DNA sample:	<u>2 µl</u>
	50 µl

PCR Conditions

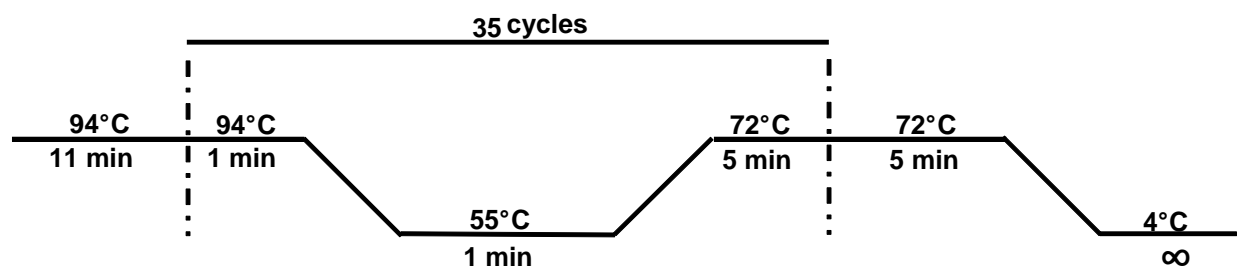


Fig 24: PCR conditions for TNFR1^{-/-}

c) **TNFR2^{-/-}: IMR0338: 5'-CCT CTC ATG CTG TCC CGG AAT**

IMR0339: 5'-AGC TCC AGG CAC AAG GGC GGG

IMR0340: 5' CGG TTC TTT TTG TCG AGA C

IMR0341: 5' ATC CTC GCC GTC GGG CAT GC

The polymerase chain reactions were performed in 50 µl volumes made up as follows:

10xMagnesium free buffer:	5 µl
25 mM MgCl ₂ :	3 µl
10 mM dNTP's:	1 µl
Primers: IMR ₃₃₈	1 µl
IMR ₃₃₉	1 µl
IMR ₃₄₀	1 µl
IMR ₃₄₁	1 µl
H ₂ O:	35.875 µl
Taq:	0.125 µl
DNA sample:	<u>2 µl</u>
	50 µl

PCR Conditions:

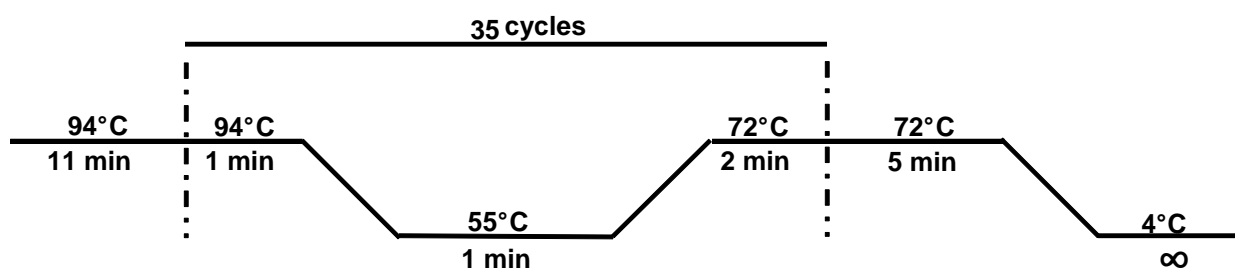


Fig 25: PCR conditions for TNFR2^{-/-}

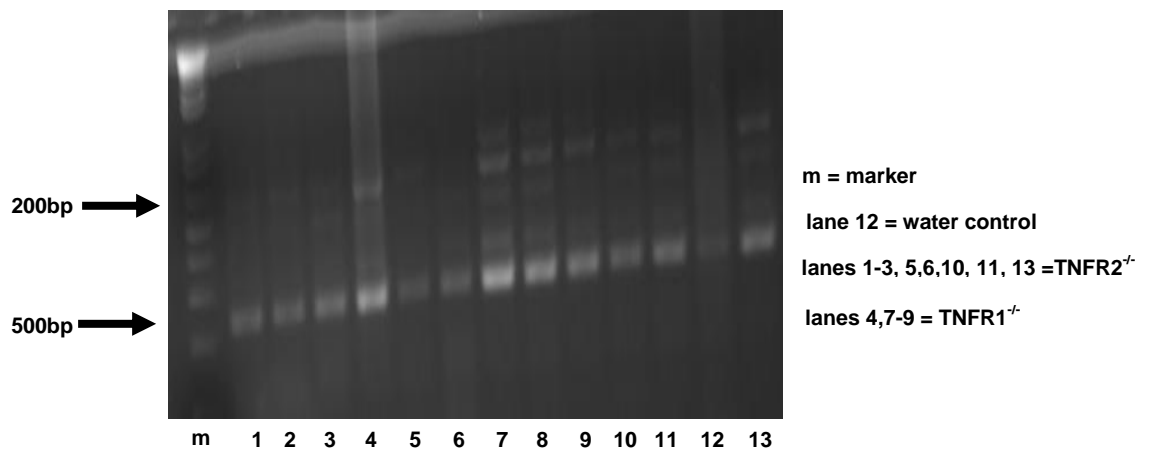


Fig 26: UV gel analysis of extracted mouse genomic DNA for the targeted disruption of the TNFR gene

iv) Sequences and PCR for STAT-3^{-/-} mouse model

To determine whether the mice were Cre positive the following sequences were used:

Cre Sense primer: 5' GTT CGC AAG AAC CTG ATG GAC A 3'

Cre Antisense primer: 5' CTA GAG CCT GTT TTG CAC GTT C 3'

The mice were also genotyped for the MLC 2V gene as a positive control for the presence of DNA. For this, the following sequences were used:

MLC 2V Sense primer: 5' GGC AAC CCT CAG ACA CCA T 3'

MLC 2V Antisense primer: 5' TGT GGA GGC TCT GGA TCA GGA C 3'

The polymerase chain reactions were performed in 50 µl volumes made up as follows:

10xMagnesium free buffer: 10 µl

4mM MgCl₂: 8 µl

100µM dNTP's: 1.6 µl

Primer: 0.5 µl

β actin 3': 0.5 µl

β actin 5':	0.5 μ l
H ₂ O:	26.9 μ l
Taq:	0.5 μ l
DNA sample:	<u>1.5 μl</u>
	50 μ l

PCR Conditions:

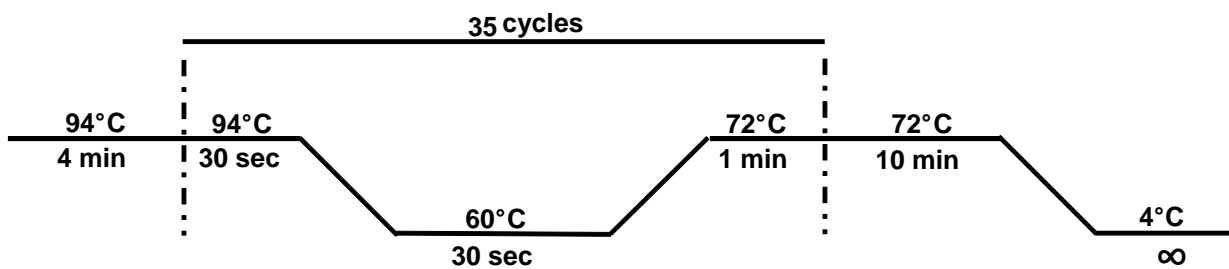


Fig 27: PCR conditions for STAT-3^{-/-} mice.

To establish whether STAT-3 was floxed, one set of primers, yielding PCR fragments of differing lengths, were used. A 250 bp fragment is seen on the gel if the STAT-3 is floxed whereas a 200 bp fragment is visualized on the gel if the STAT-3 is not floxed. If the mouse being tested is heterozygous, then a double band is seen. STAT-3 floxed or non-floxed DNA was amplified with the following primers:

STAT-3 primer (a) 5'CCT GAA CAA GTT CAT CTG TGT GAC 3'

STAT-3 primer (b) 5' CAC ACA AGC CAT ACT CTG GTC TCC 3'

The polymerase chain reactions were performed in 50 μ l volumes made up as follows:

10xMagnesium free buffer:	5 μ l
4mM MgCl ₂ :	4 μ l
100 μ M dNTP's:	0,8 μ l

Primer:	0.25 μ l
β actin 3':	0.25 μ l
β actin 5':	0.25 μ l
H ₂ O:	13.2 μ l
Taq:	0.25 μ l
DNA sample:	<u>1.0 μl</u>
	25 μ l

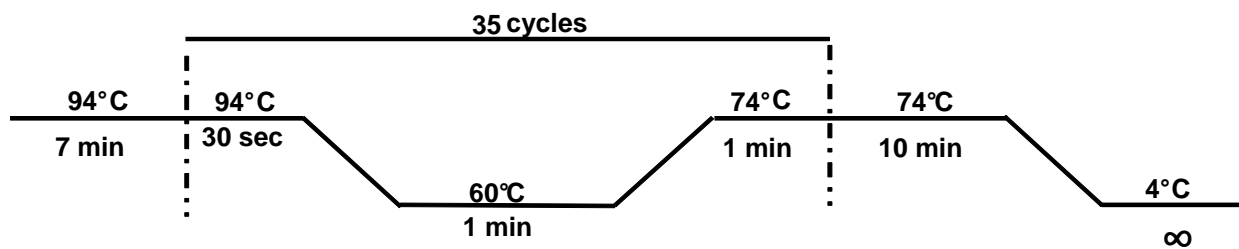


Fig 28: PCR conditions for STAT-3 floxed mice.

PCR Conditions:

1.1.3 Agarose Gel Electrophoresis

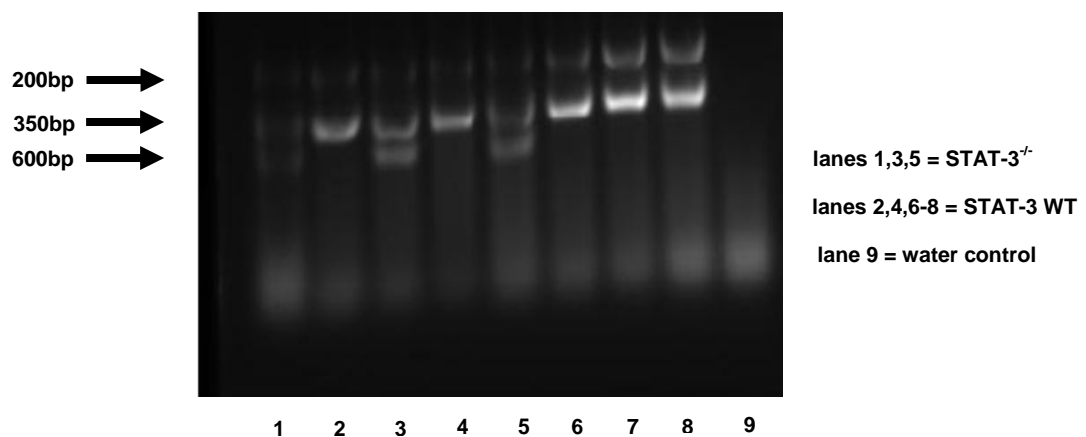


Fig 29: UV gel analysis of genomic mouse DNA showing Cre positive band (350bp) as well as the control Myosin Light Chain 2V (MLC2V) band at 600bp.

A 2% agarose gel was used to ensure optimal separation between bands. Two grams of agarose (Whitehead Scientific, RSA) was made up in 100 ml of 1 x Tris Acetate electrophoresis buffer (TAE) (0.04 M Tris-Acetate, 0.001 M EDTA) and heated in a microwave oven at 70% power until completely dissolved. Once dissolved, 1 μ l ethidium bromide was added per 100 ml gel in order to visualize the amplified DNA fragment under ultra violet light. 1 x TAE buffer was added to the gel tank once the gel had set and before removal of the combs in order to prevent the wells being damaged. Subsequently, 10 μ l of each PCR reaction was mixed with 2 μ l gel loading buffer (0.25% Bromophenol Blue-Xylene Cyanole Dye Solution, Sigma) and carefully loaded into the wells. Simultaneously, 4 μ l of a 100 bp DNA ladder (Roche, Germany) was run as a marker to verify the size of the PCR products. The gel was run at 90 V for 30 – 40 minutes and the bands were then visualized on a transilluminator (UviTec, USA). PCR results were recorded using the UviTec Geldoc apparatus with UViband software v.97 (Cambridge, UK).

2.0 Isolated mouse heart model

Adult male TNF^{-/-}, TNFR1^{-/-}, TNFR2^{-/-}, cardiomyocyte-specific STAT-3^{-/-} mice and their respective WTs were anaesthetized (pentobarbitone, 60 mg/kg ip) and heparinized (25 IU ip) to prevent the blood clotting. Once an adequate level of anaesthesia had been achieved (tested by lack of pedal reflex) the chest was opened and the sternum and attached costal cartilage excised to give adequate access to the mediastinum. Hearts were rapidly excised (Smith et al., 2002) and arrested in ice cold modified Krebs-Henseleit buffer (118 mM NaCl; 24 mM NaHCO₃; 4 mM KCl; 1 mM NaH₂PO₄; 2.5 mM CaCl₂; 1.2 mM MgCl₂; 0.5 mM di-sodium EDTA; 10 mM glucose; gassed with 95% O₂/5% CO₂ at 37°C). The aorta was cannulated and the heart then mounted on the Langendorff system. The heart was fastened by means of a 4-0 silk suture (on a 13mm curved atraumatic needle) through the apex onto a rigid lightweight lexan coupling rod to a force displacement transducer (Grass Technologies, FT02C, Mass. USA). Diastolic resting tension was adjusted to 2 g and hearts were paced at 560 beats per minute while being retrogradely perfused with modified Krebs-Henseleit buffer at a constant pressure of 110 cm H₂O (Smith et al., 2002) (see Figure 30). The developed force was recorded on a Powerlab Labchart (ADI Instruments, Australia) and coronary flow rate was measured by timed collection. These parameters were utilized as measures of cardiac function for all groups of hearts used in the protocols. Temperature was measured by placement of a fine thermocouple wire (Physitemp, NJ, USA) into the right atrium and monitored on a Digitron 2600T temperature sensor (Torquay, UK), (figure 30). Isolated hearts were subjected to a 25 minute stabilization period followed by

35 minutes global ischaemia (obtained by shutting off the flow of buffer to the heart from the aortic cannula by means of a stopcock) and a 45 minute reperfusion period (Smith et al., 2002). Coronary flow, heart rate and developed force are used to determine the viability of an isolated heart. A minimum flow of 1.5 ml/min and maximum flow of 5.0 ml/min are acceptable for coronary flow, the acceptable heart rate is between 460bpm-600bpm and the developed force is accepted when ≥ 4 .

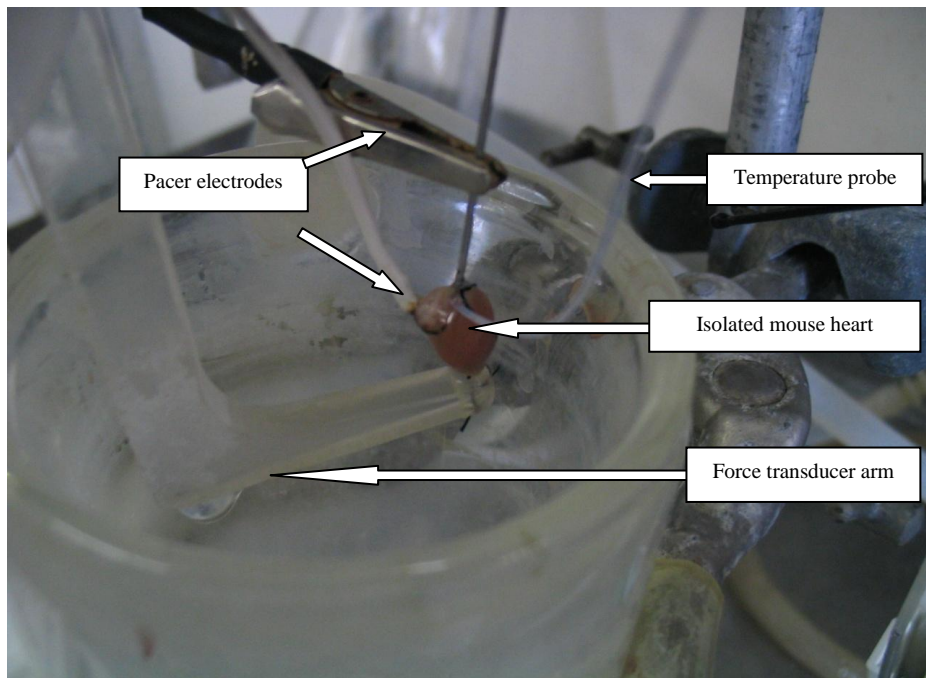


Fig 30: Mouse heart mounted on Langendorff apparatus. A cannula is inserted into aorta for retrograde perfusion. The pacer electrodes are used to pace the heart to 560 bpm and the heart pulls against a force of 2g.

2.1 Ischaemic and TNF α postconditioning protocol

The Langendorff retrograde perfusion technique was used to assess infarct size and to collect tissue for western blot analysis, where appropriate, from the TNF $^{-/-}$, TNFR1 $^{-/-}$, TNFR2 $^{-/-}$, STAT-3 $^{-/-}$ and their respective wildtypes. The isolated hearts were subjected to IPostC at the onset of reperfusion by initiating six alternating cycles of 10 seconds reperfusion, 10 seconds ischaemia. In a separate group, pharmacological postconditioning with TNF α was initiated by six alternating cycles of 10 seconds reperfusion without TNF α , 10 seconds reperfusion with 0.5 ng/ml TNF α (TNF-PostC), figure 31. To assess whether the mounting of hearts itself had an effect, a time-matched control group was subjected to the full length of the protocol on the Langendorff apparatus with no ischaemic period or IPostC.

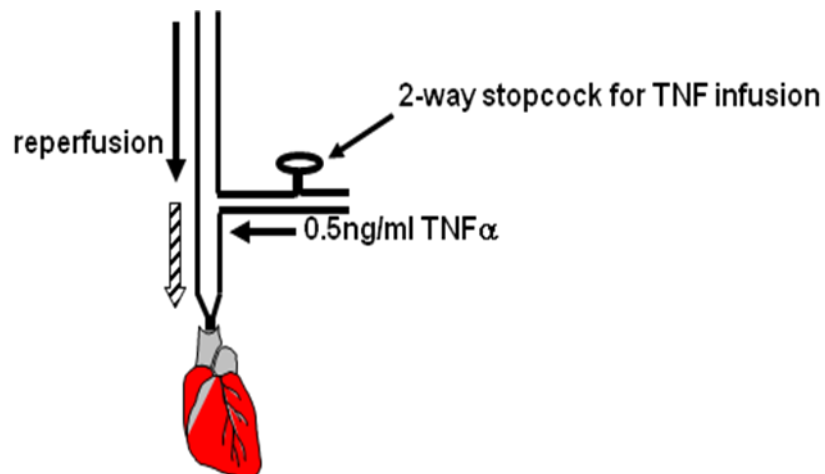


Figure 31: Protocol of pharmacological postconditioning with TNF α . Reperfusion is allowed for 10 sec, after which the stopcock is opened for 10 sec to allow the flow of 5 ng/ml TNF α to mix with the reperfusion flow, giving a final concentration of 0.5 ng/ml entering the heart. The stopcock is then closed and reperfusion with Krebs-Henseleit buffer alone takes place for 10 sec, after which the cycle is repeated a further 5 times.

2.2 Infarct size assessment

At the end of each experimental protocol, hearts were removed from the perfusion rig and infarct size assessed by triphenyltetrazolium chloride (TTC) staining as described previously (Sumeray et al., 2000). Briefly, the heart was placed in TTC stain (1% in phosphate buffer, pH7.4) for 1 minute at 37°C after which it was frozen for 24 hours to allow for easy slicing. 1mm slices, perpendicular to the apex, were then cut. Each slice includes both right and left ventricle plus septum. These slices were then compressed between thin glass plates 0.5mm apart and the sections scanned, and infarct size assessed using computerized planimetry (Planimetry+, Boreal Software, Norway), (figure 32).

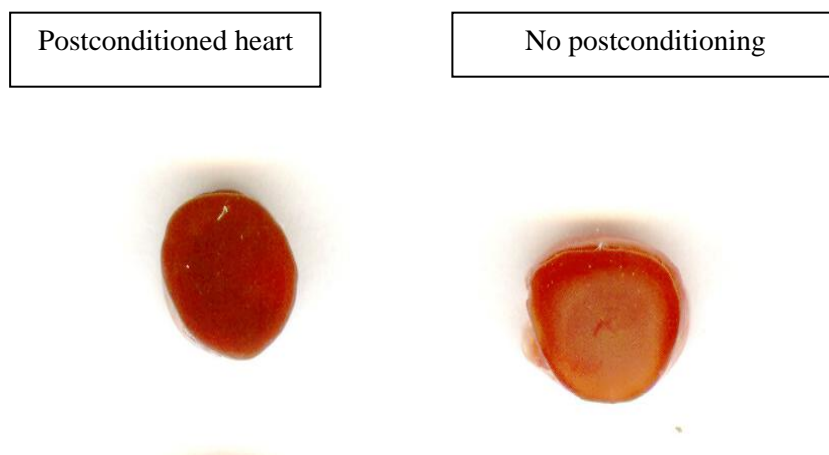


Fig 32: TTC staining for infarct size evaluation. TTC reacts with the NADPH in live tissue to give a brick red colour. There is no reaction in the infarct tissue, which remains pale and allows infarct size determination by planimetric analysis.

3.0 Nuclear and cytosolic protein extraction (adapted from Williams and Ford, 2001)

Nuclear and cytosolic proteins were extracted from the hearts by homogenization of the tissue in lysis buffer as follows:

<u>Solution A (30ml)</u>	<u>µl</u>	<u>Final Concentration</u>
1M Hepes, pH7.9	600	20 mM
2M MgCl ₂	37.5	2.5 mM
500mM EDTA	6	100 µM
100mM β-glycerophosphate	6000	20 mM
Triton X-100 (100%)	15	0.5%
100mM DTT	1500	500 µM
100mM NaVO ₄	30	100 µM
EDTA-free protease inhibitor	1200	-

100mM PMSF	300	1 mM
5M NaCl	450	75 mM
dH ₂ O	19862	-

<u>Solution B (10ml)</u>	<u>µl</u>	<u>Final Concentration</u>
1M Hepes	200	20 mM
2M MgCl ₂	12.5	2.5 mM
500mM EDTA	2	100 µM
100mM β-glycerophosphate	2000	20 mM
Triton X-100 (100%)	100	1%
100mM DTT	1500	500 µM
100mM NaVO ₄	10	100 µM
EDTA-free protease inhibitor	400	-
100mM PMSF	100	1 mM
5M NaCl	150	75 mM
dH ₂ O	6526	-

Frozen hearts were wrapped in aluminium foil, pulverized under liquid nitrogen then transferred into 900 µl of lysis buffer (Solution A). After homogenizing twice by Polytron, at setting 4, for 5 sec each time, the suspension was centrifuged at 10000 g (12000 rpm) for 5 min at 4°C. The supernatant, corresponding to the cytosolic fraction, was carefully removed and placed in a fresh tube. The remaining pellet was resuspended in 300 – 500 µl of Solution B, transferred into a

clean Eppendorf tube and centrifuged for a further 30 min at 10000 g (12000 rpm) at 4°C. The supernatant, corresponding to the nuclear fraction was removed into a clean tube.

3.1 Mitochondrial and cytosolic protein extraction (adapted from Tal M Lewin, 2008.)

Mitochondrial and cytosolic proteins were extracted from the hearts by homogenization of the tissue in lysis buffer as follows:

Lysis buffer A

100 mM Tris-HCl – pH 7.4

250 mM Sucrose

1 mM DTT

1 mM EDTA

Incubation Buffer

250 mM Sucrose

25 mM Tris

8.5 mM KH₂PO₄

Low ionic strength sample buffer (2X)

400 µl 10% SDS

400 µl glycerol

40 µl mercaptoethanol

1.16 ml 5 mM Tris (pH 6.8)

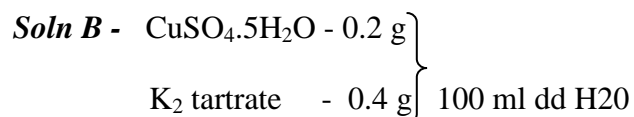
Bromophenol blue

Frozen hearts were minced finely with a pair of scissors in 1ml of lysis buffer. After homogenisation in a Dounce homogeniser, the suspension was centrifuged at 600 g for 5 min at 4°C. The supernatant was placed in a fresh tube and further centrifuged at 10 300 g (11,500 rpm) for 10 min. The resulting supernatant corresponds to the cytosolic fraction. The pellet was resuspended in 40 µl mitochondrial incubation buffer. After protein determination, an equal volume of low ionising buffer was added to the mitochondrial and cytosolic fractions.

3.2 Lowry Protein determination

Duplicate 5 μ l aliquots of nuclear, cytosolic or mitochondrial sample were added to 995 μ l of sterile distilled water (1:200 dilution) followed by addition of 1ml of the following CTC reagent.

i) **CTC reagent: Soln A** – Na_2CO_3 – 20g/100ml dd H₂O



Add **Soln A** to **Soln B** with constant stirring to prevent precipitation.

ii) **SDS**

10% SDS made by diluting a 20% solution

iii) **NaOH**

0.4 g/200ml dd H₂O

Equal volumes of all 3 reagents were mixed immediately prior to use.

The samples were immediately mixed with the CTC reagent and incubated for 10 minutes at room temperature. 500 μ l of Folin's Reagent, diluted 1:5 with double distilled water (Merck, Germany) was then added with continuous mixing, as the colour complex to be assayed forms immediately. The samples were incubated at room temperature for a further 30 minutes and optical density was read at 750 nm on a Varian 130 dual beam spectrophotometer (Peterson, 1977).

A standard curve was prepared using bovine serum albumin (Sigma, Germany) diluted to provide a concentration range from 10 – 200 μ g/ml and the range was assayed in the same manner as the samples.

3.3 Polyacrylamide Gel Electrophoresis (PAGE)

100 μ g of nuclear, mitochondrial or cytosolic protein was loaded per lane and run on a 10% (w/v) acrylamide/bisacrylamide gel at 120V for approximately 90 - 120 minutes. 7 μ l of peqGold Protein Marker IV (PEQLAB Ltd, UK) was run simultaneously to identify the size of the candidate peptides. The gel was transferred overnight (0.02 amps at 4°C) onto an Amersham PVDF membrane (Hybond, Amersham, UK). Thereafter, the membrane was fixed in methanol

for 5 minutes, air-dried and stained with Ponceau reagent (Sigma P7170, St Louis, MO) to quantitate and evaluate appropriate transfer of proteins from gel to membrane. The membrane was placed between two acetate sheets and scanned for densitometric analysis, using the UVIBand software package.

Subsequently, the Ponceau stain was rinsed off with three 5 minute washes in Tris buffered saline with 0.1% Tween (TBS-T). The membrane was then blocked at room temperature in 5% fat-free milk powder made up in TBS-T. After 2 hours the membrane was vigorously hand washed in TBS-T with 3 cycles of 5 minutes. The membrane was then probed overnight at 4°C with one of the primary antibodies followed by 2 hours incubation with the appropriate secondary antibody.

3.4 Western blot analysis

Control or postconditioned mouse hearts were subjected to a 20 min stabilization period followed by 35 min of global ischemia and 15 min of reperfusion. The hearts were then snap-frozen in liquid nitrogen before being stored at -80°C until protein extraction was performed. Nuclear and cytosolic proteins were extracted from the hearts by homogenization of the tissue in a lysis buffer, as previously described (Lecour et al., 2005b; Sack et al., 1997). Phosphorylated states of Akt, Erk1/2, STAT-3 (Phospho-STAT-3, tyr705), glycogen synthase kinase-3 beta (phospho-GSK-3 β , ser 9) as well as total levels of Akt, Erk1/2, STAT-3, GSK-3 β , RAGE, PAR-2 and glyceraldehyde-3 phosphate dehydrogenase (GAPDH) were analysed by sodium dodecyl sulphate-polyacrylamide gel immunoelectrophoresis (SDS-PAGE) using antibodies from Cell Signaling Technologies. Equal loading was verified with GAPDH. Levels of phosphorylated proteins were normalized, where appropriate, to their total protein levels done in the same samples and in the same conditions but on a separate membrane. Normalization is merely the ratio of phosphorylated protein to total protein, multiplied by 100. Relative densitometry was determined with the use of computerized software package (UVI Soft, UVI Band, UVI Tech, Cambridge, UK). A minimum of 6 hearts was used per group.

3.4.1 Primary and Secondary Antibodies

i: Total STAT

Primary antibody: STAT-3 Antibody rabbit polyclonal IgG 1:4000 in 5% (w/v) fat-free milk

Secondary antibody (2 hours at room temperature): Anti-rabbit 1:3000 in 5% (weight/volume {w/v}) fat free milk

ii: Phosphorylated STAT

Primary antibody: PhosphoPlus^R STAT-3 (Tyr705) rabbit polyclonal IgG 1:1000 in TBS-T

Secondary antibody (2 hours at room temperature): Anti-rabbit 1:3000 in 5% (w/v) fat free milk

iii): Akt

Primary antibody: Akt Antibody rabbit polyclonal IgG 1:1000 in TBS-T

Secondary antibody: Anti-rabbit (2 hours at room temperature): 1:4000 in 2.5% (w/v) fat free milk

iv: Phosphorylated Akt

Primary antibody: Phospho-Akt (Ser473) Mouse mAB - 1:1000 in TBS-T.

Secondary antibody (2 hours at room temperature): Anti-mouse 1:4000 in 2.5% (w/v) fat free milk

v: Glycogen Synthase Kinase-3 Beta (GSK-3 β)

Primary antibody: GSK-3 β Rabbit mAb 1:1000 in TBS-T

Secondary antibody (2 hours at room temperature): Anti-rabbit 1:4000 in TBS-T

vi: Phosphorylated Glycogen Synthase Kinase-3 Beta (pGSK-3 β)

Primary antibody: Phospho- GSK-3 β (Ser9) 1:3000 in TBS-T

Secondary antibody (2 hours at room temperature): Anti-rabbit 1:3000 in TBS-T

vii: p44/42 MAPK (Erk1/2)

Primary antibody: p44/42 (Erk1/2) 1:1000 in TBS-T

Secondary antibody (2 hours at room temperature): Anti-rabbit 1:4000 in 2.5% (w/v) fat free milk

viii: *Phospho-p44/42 MAPK (Erk1/2)*

Primary antibody: Phospho-p44/42 MAPK (Erk1/2) (Thr202/Tyr204) 1:1000 in TBS-T

Secondary antibody (2 hours at room temperature): Anti-rabbit 1:4000 in 2.5% (w/v) fat free milk

ix: *Receptor for Advanced Glycation End Products (RAGE)*

Primary antibody: Anti-mouse RAGE Antibody 1:1000 in TBS-T

Secondary antibody (2 hours at room temperature): Anti-goat 1:4000 in 2.5% (w/v) fat free milk

x: *Protease-Activated Receptor -2 (PAR-2)*

Primary antibody: PAR-2 (SAM11): sc-13504 1:1000 in TBS-T

Secondary antibody (2 hours at room temperature): Anti-mouse 1:4000 in 2.5% (w/v) fat free milk

xi: *Glyceraldehyde-3-phosphate dehydrogenase (GAPDH)*

Primary antibody: GAPDH (FL-335) 1:1000 in TBS-T

Secondary antibody (2 hours at room temperature): Goat Anti-rabbit 1:3000 in TBS-T

For mitochondrial and cytosolic fractions, the following antibodies were used:

i: *Total STAT*

Primary antibody: STAT-3 Antibody rabbit polyclonal IgG 1:4000 in 5% (w/v) fat-free milk

Secondary antibody (2 hours at room temperature): Anti-rabbit 1:3000 in 5% (weight/volume {w/v}) fat free milk

ii: *Phosphorylated STAT*

Primary antibody: phospho STAT-3 (Ser 727) rabbit polyclonal IgG 1:1000 in TBS-T

Secondary antibody (2 hours at room temperature): Anti-rabbit 1:4000 in 2.5% (w/v) fat free milk

iii: VDAC

Primary antibody: VDAC goat polyclonal IgG 1:3000 in TBS-T (1 hour at room temperature).

Secondary antibody: Anti-goat 1:4000 in 2.5% (w/v) fat free milk (30 minutes at room temperature).

All antibodies were from Cell Signaling Technology, USA with the exception of RAGE (RD Systems, USA), PAR-2 and GAPDH (both from Santa Cruz Biotechnology, USA). Relative densitometry was determined with the use of computerized software package (UVI Soft, UVI Band, UVI Tech, Cambridge, UK). A minimum of 6 hearts was used per group.

D. TNF α , TNFR2 & ISCHAEMIC POSTCONDITIONING

*Part of the work in this section has been published in **Cardiovasc Res**, (2009) 84, 201-208
“Ischaemic postconditioning protects against reperfusion injury via the SAFE pathway”*

Part of the experiments in this section have been performed in conjunction with Sarin Somers, registered as a PhD student at the University of Cape Town

Introduction

Following acute coronary occlusion, early reperfusion is the current optimal way to rescue the heart. Although essential, reperfusion is associated with cellular damage by activation of deleterious signalling cascades, as early as in the first seconds of reperfusion (Braunwald and Kloner, 1985). Attempts in the management of reperfusion injury have recently entered the clinical arena (Piot et al., 2008; Staat et al., 2005; Yellon and Hausenloy, 2007). IPostC is still often thought to protect only via activation of the RISK pathway (Reperfusion Injury Salvage Kinases) which includes activation of the prosurvival kinases Akt and Erk1/2 at the time of reperfusion (Hausenloy et al., 2005; Kin, 2008; Tsang et al., 2004). However, recent studies demonstrated that protection with ischaemic postconditioning can occur independently of the activation of the RISK pathway, therefore confirming the existence of multiple protective pathways (Heusch et al., 2008; Montecucco et al., 2009; Schwartz, 2006; Przyklenk et al., 2008; Skyschally et al., 2009).

The activation of the SAFE pathway, which includes the cytokine TNF α , activates the transcription factor STAT-3 and is now recognized as a “RISK-free” pathway that can confer protection in ischaemic preconditioning (Lecour, 2009a; Lecour, 2009b; Lecour et al., 2002; Lecour et al., 2005b; Suleman et al., 2008). However, the role of TNF α in the SAFE pathway during ischaemic postconditioning has not yet been elucidated.

It is known that TNF α is activated during the course of reperfusion, contributing to cardiac dysfunction and apoptosis (Belosjorow et al., 1999; Maekawa et al., 2002; Mann, 2002). Although ischaemic preconditioning is associated with an increase in TNF α during the preconditioning stimulus, its protective effect is partly attributed to a reduction in myocardial TNF α levels at the time of reperfusion (Kin, 2008). However, TNF α can have adaptive effects in relation to its concentration (with low concentrations being cardioprotective and high concentrations being harmful (Kurrelmeyer et al., 2000; Lecour *et al* 2005a; Lecour *et al.*, 2005b; Lecour *et al.*, 2002) and to which of its two receptors it binds (binding to TNF receptor 1 is suggested to have an adverse effect while binding to its receptor 2 is suggested to be cardioprotective) (Monden et al., 2007; Schulz, 2008; Schulz and Heusch, 2009).

Therefore, our aim was to explore whether TNF α -signalling, through the activation of its receptor 2 (TNFR2), was involved in the protective effect of IPostC and whether this signalling mechanism activated the SAFE pathway rather than the RISK pathway, to confer protection. We

also wished to explore whether GSK-3 β , RAGE and PAR-2 were part of the signalling cascade in ischaemic postconditioning. We used an isolated heart model from wild type mice (TNF-WT) and 3 different transgenic mice from the same background, TNF knockout (TNF^{-/-}), TNF receptor 1 knockout (TNFR1^{-/-}) and TNF receptor 2 knockout (TNFR2^{-/-}) to demonstrate that binding of TNF α to its receptor 2 during ischemic postconditioning is indeed required for the protection. Pharmacological protein kinase inhibitors and mice which are deficient in the transcription factor, STAT-3 in their cardiomyocytes were used to demonstrate that, in the context of ischemic postconditioning, TNF α confers its cardioprotective effect via the activation of the transcription factor STAT-3 rather than the protein kinases Akt and Erk1/2.

Furthermore, using western blot analysis, we explored whether phosphorylation of GSK-3 β , RAGE or PAR-2 may act as downstream targets of the SAFE (TNF/JAK/STAT) pathway. A schematic of our proposed hypothesis follows in figure 33.

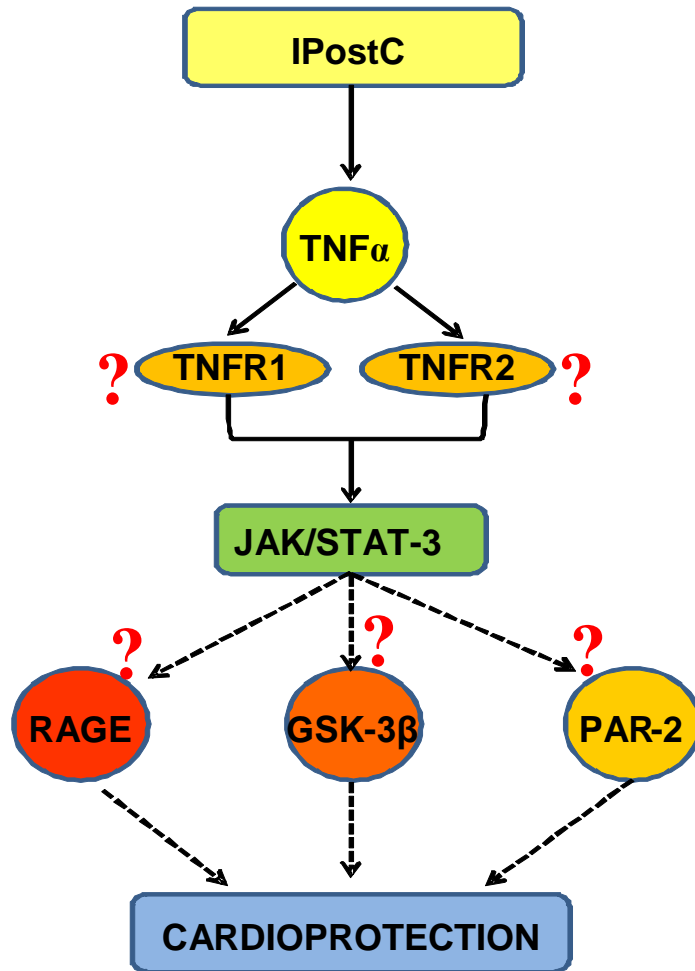


Fig 33: Schematic of proposed hypothesis. Does TNF α initiate cardioprotection during ischaemic postconditioning via the SAFE pathway? Which receptor is involved? Are RAGE, GSK-3 β and PAR-2 involved in TNF α -induced protection? Abbreviations as described in text.

2.0 Methods

All experiments were performed in accordance with the Guide for the Care and Use of Laboratory Animals published by the US National Institutes of Health {NIH Publication No. 85 (23), revised 1996}. All procedures were approved by the Faculty of Health Sciences Animal Ethics Committee, University of Cape Town.

Homozygous TNF deficient mice ($TNF^{-/-}$), TNF receptor 1 deficient mice ($TNFR1^{-/-}$) or TNF receptor 2 deficient mice ($TNFR2^{-/-}$) and their respective wildtype, Black6x129S ($TNF-WT$) were used. All these mice were a generous gift from Dr Muazzam Jacobs and Prof Bernhard Ryffel, Department of Immunology, Faculty of Health Sciences, University of Cape Town. Cardiomyocyte-specific STAT-3 deficient mice ($STAT-3^{-/-}$) from C57Black6 background were created in our facilities by crossing homozygous floxed STAT-3 mice with heterozygous MLC2V-driven Cre recombinase mice, as previously described (Smith et al., 2004). All mice were 12-14 weeks of age (weighing 25-30g).

2.1 Perfusion of isolated mouse hearts

Hearts from adult male $TNF^{-/-}$, $TNFR1^{-/-}$, $TNFR2^{-/-}$, cardiomyocyte-specific $STAT-3^{-/-}$ mice and their respective wild types were excised rapidly and perfused retrogradely using the Langendorff apparatus. Isolated hearts were subjected to a 20 minute (min) stabilization period followed by 35 min global ischemia and a 45 min reperfusion period as previously described (Smith et al., 2002)] (see Section C for protocol). At the onset of reperfusion, IPostC was initiated with 6 alternating cycles of 10 seconds reperfusion, 10 seconds ischaemia. In a separate group, pharmacological postconditioning with $TNF\alpha$ was initiated with 6 alternating cycles of 10 seconds reperfusion without $TNF\alpha$, 10 seconds reperfusion with 0.5 ng/ml $TNF\alpha$ (TNF-PostC), (see Figure 31 in Section C). A time-matched control was used to assess infarct size to determine whether the technique itself had damaged the heart. Infarct size by TTC staining demonstrated that the technique had no adverse effect.

To determine whether STAT-3, Akt or Erk1/2 were involved, AG490, an inhibitor of the JAK/STAT pathway (100 nmol/L), wortmannin (wort), an inhibitor of the phosphatidylinositol-3 kinase (PI-3 Kinase)/Akt pathway (100 nmol/L) or PD98059 (PD), an inhibitor of the Mek/Erk1/2 pathway (100 nmol/L) were given during the first 15 min of reperfusion. To confirm the role of the TNF receptors, specific antibodies to $TNFR1$ or $TNFR2$ (3 mg/L) were

given to TNF-WT mice during the ischemic postconditioning protocol, (Figure 34). The concentration of the antibodies corresponded to the ND₅₀ of the antibodies, as given by the manufacturer. A minimum of 6 hearts were used in all groups. Infarct size was evaluated by triphenyltetrazolium chloride (TTC) staining and assessed using computerized planimetry as described in Section C.

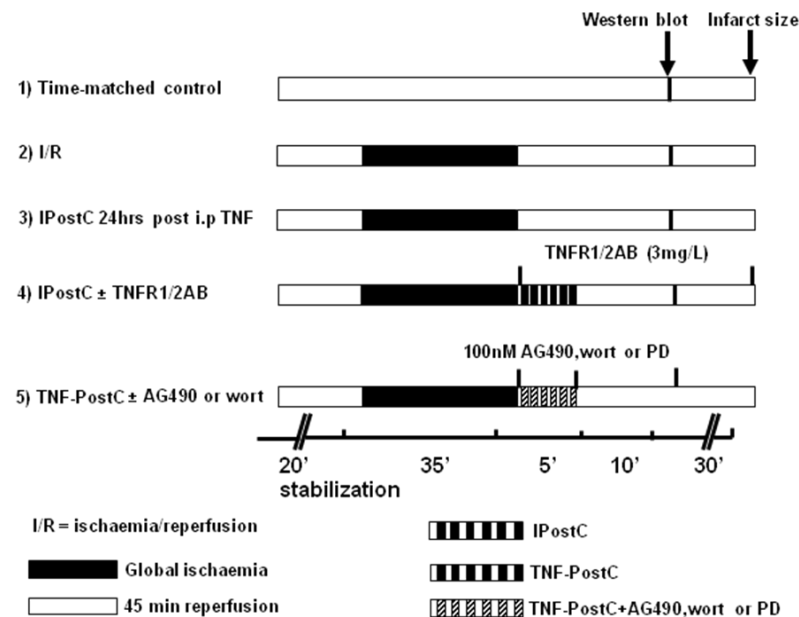


Fig 34: Schematic of experimental protocol. Hearts were removed for Western blot analysis at the timepoint indicated, in a separate experiment. TNF receptors antibodies were added throughout the reperfusion protocol. Inhibitors of PI-3 kinase, JAK/STAT and Erk1/2 were added during the postconditioning protocol as indicated. PD= PD98059; wort = wortmannin; AB = antibody; All other abbreviations as defined in text.

2.2 Ischaemic postconditioning 24 hours after administration of TNF α to TNF^{-/-} mice

In a separate set of experiments, 24 hours prior to performing the ischaemic postconditioning protocol, each TNF^{-/-} mouse was given a single ip injection of 0.5 ng TNF α . Thereafter, the isolated hearts underwent the identical ischaemic postconditioning protocol as described in the previous section.

2.3 TNF α release from heart tissue

Two minutes after the onset of postconditioning the hearts were removed from the cannula by cutting off the atria. They were then placed into an eppendorff tube and immediately immersed in liquid N₂ to snap freeze. Once frozen, the tubes were stored at -80°C until analysis. The

frozen hearts were wrapped in aluminium foil and pulverized under liquid nitrogen. The pulverized tissue was placed in 1.5 ml ice-cold phosphate buffered saline (PBS) and the suspension was then ultrasonically homogenised (Virtis, Virsonic; USA) at 50W for 10-20 seconds. Homogenates were centrifuged at 14000 rpm for 20 min at 4°C and the supernatant was collected for TNF α measurement by Elisa (kit) as per manufacturer's instructions.

2.4 Chemicals and Pharmacological Agents

Unless otherwise stated, all chemicals were obtained from Sigma-Aldrich Chemicals, Germany. Murine TNF α was obtained from PeproTech Inc., Israel; TNF Receptor monoclonal antibodies were obtained from R&D Systems, U.S.A.

2.5 Statistical Analysis

Data are presented as mean \pm SEM. Comparisons between multiple groups were performed by One-way ANOVA followed by Tukey post hoc test or Bonferroni multiple comparison test. The unpaired t test was used for TNF α release from heart tissue (Graph Pad Instat). $p < 0.05$ was considered statistically significant.

3.0 Results

3.1 Postconditioning with TNF α

➤ TNF α is required in IPostC

To determine whether TNF α is required in IPostC mediated cardioprotection, we compared infarct size (calculated as a percentage of total ventricular volume) in TNF-WT versus TNF^{-/-} mice following the IPostC protocol of 6 x 10 sec cycles of alternating reperfusion and ischaemia, (see Figure 34). IPostC in TNF-WT mice reduced infarct size from 52 \pm 3% in the ischaemic control group (I/R) to 18 \pm 2% for IPostC ($p < 0.05$; Figure 35). IPostC with 6 x 10 sec cycles in the TNF^{-/-} hearts failed to reduce the infarct size (37 \pm 3%, $p = \text{ns}$ versus TNF^{-/-} I/R group. Similar data was obtained with TNF-PostC, (Figure 35).

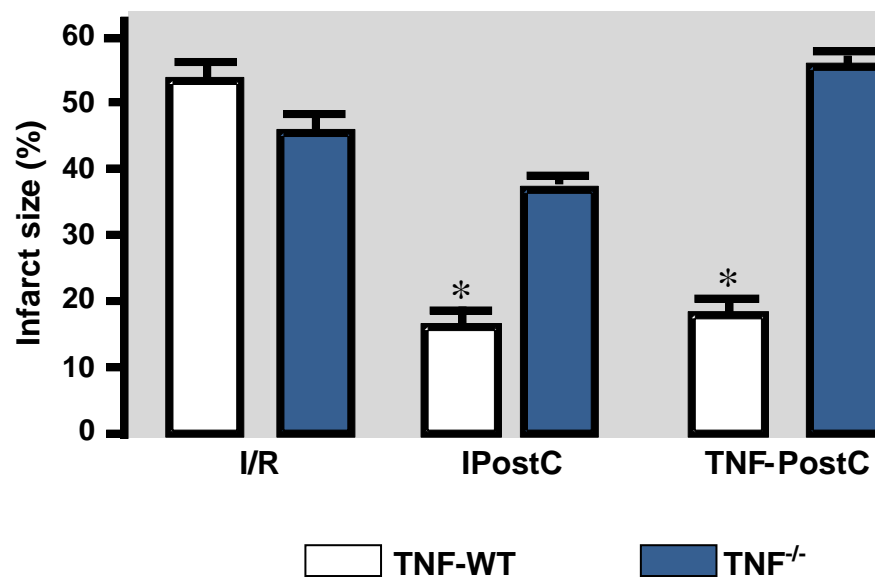


Fig 35: TNF α is required in, and can also mimic, IPostC. 6 x 10 sec cycles of IPostC reduced the infarct size in TNF-WT but not TNF^{-/-} mice, $n \geq 6$ per group, $*p < 0.05$ vs respective I/R. Similarly, the infarct size in WT was reduced by 6 x 10 sec cycles of TNF-PostC but this failed to reduce the infarct size in TNF^{-/-} mice, $n \geq 6$ per group, $p = \text{ns}$. Abbreviations are as described in the text.

➤ **TNFR1^{-/-} but not TNFR2^{-/-} mice can be postconditioned**

To investigate the role of the TNF receptors in ischaemic postconditioning, we measured infarct size in TNFR1^{-/-} and TNFR2^{-/-} mice subjected to both IPostC and TNF-PostC. Ischaemic control hearts from TNFR1^{-/-} and TNFR2^{-/-} mice had an infarct size of $42 \pm 3\%$ and $42 \pm 2\%$, respectively, which was similar to data obtained in TNF-WT mice. IPostC protocol significantly reduced the infarct size in TNFR1^{-/-} mice ($15 \pm 1\%$, n=6; $p < 0.05$ versus TNFR1^{-/-} ischaemia-reperfusion control) but failed to protect the heart in TNFR2^{-/-} mice ($38 \pm 3\%$, p=ns, versus TNFR2^{-/-} I/R control group). Similarly, TNF-PostC decreased the infarct size to $22 \pm 2\%$ in TNFR1^{-/-} hearts, ($p < 0.05$ vs its TNFR1^{-/-} I/R control group) but failed to protect the TNFR2^{-/-} hearts (p=ns, versus TNFR2^{-/-} I/R control group), (Figure 36A).

We confirmed these results by using specific neutralizing antibodies to each receptor during the ischaemic postconditioning protocol in TNF-WT mice. Ischaemic-reperfusion control hearts had an infarct size of $49 \pm 2\%$. Perfusion of the TNFR1 antibody during the IPostC protocol did not affect the protective effect of postconditioning ($19 \pm 2\%$, $p < 0.05$ versus ischaemic-reperfusion control group). In contrast, the protective effect of IPostC was lost in the presence of TNFR2 antibody ($53 \pm 1\%$, p=ns, versus ischaemic-reperfusion control group), (Figure 36B).

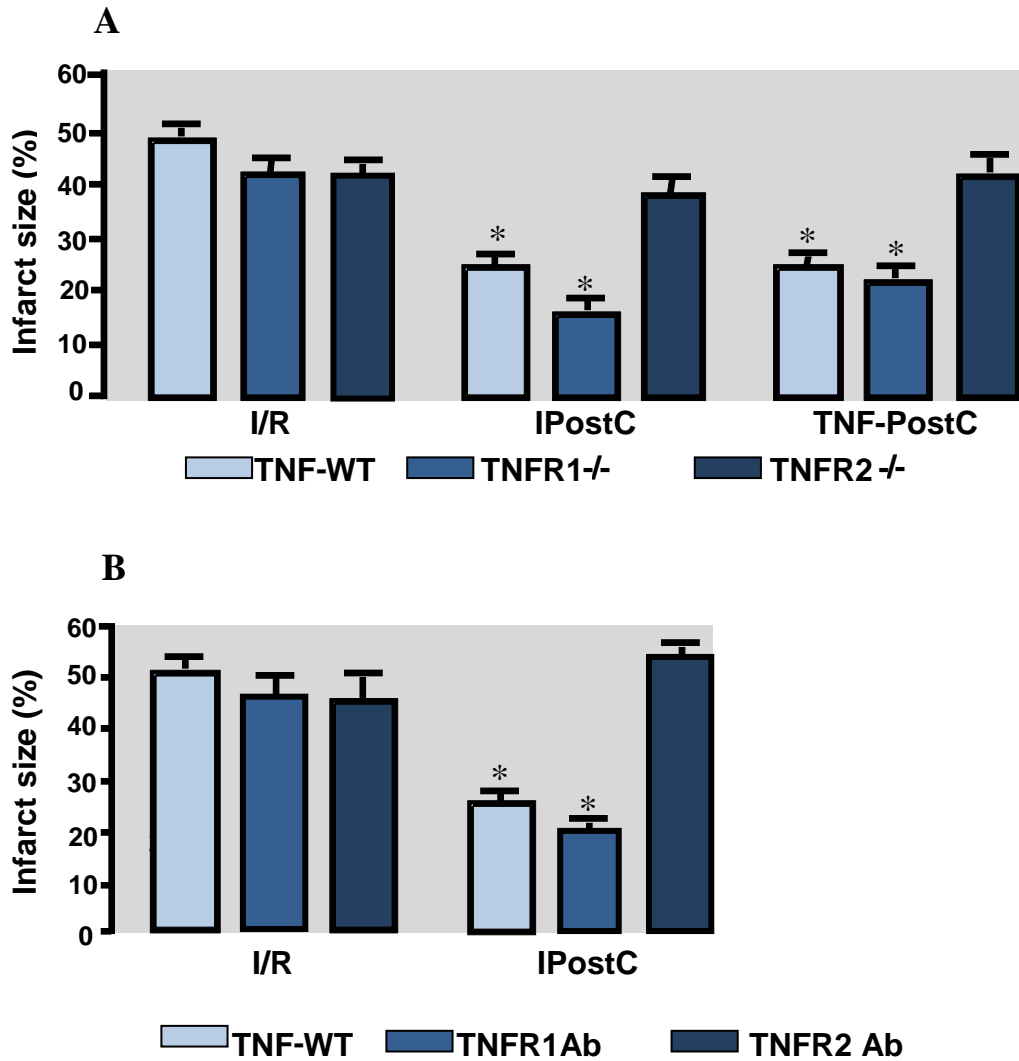


Fig 36: TNFR1^{-/-} but not TNFR2^{-/-} mice can be postconditioned. (A) 6 x 10 sec cycles of IPostC reduced infarct size in TNFR1^{-/-} mice but not in TNFR2^{-/-} mice, n = 6 per group. (B) Addition of TNFR2 antibody (Ab) but not TNFR1 Ab, abolished the infarct sparing effect of IPostC in TNF-WT hearts, *p<0.05 vs respective I/R; n = 6 per group; Abbreviations as described in the text.

Administration of TNF α 24 hours prior to ischaemia-reperfusion, restores IPostC cardioprotection in TNF $^{-/-}$ mice

A single ip injection of TNF α (0.5 ng/ml) 24 hours prior to I/R did not alter the infarct size in TNF $^{-/-}$ hearts compared to TNF-WT controls. However, the protective effect of IPostC was restored, significantly decreasing infarct size in TNF $^{-/-}$ hearts from $45 \pm 2\%$ to $16 \pm 2\%$, $p < 0.001$, (Figure 37).

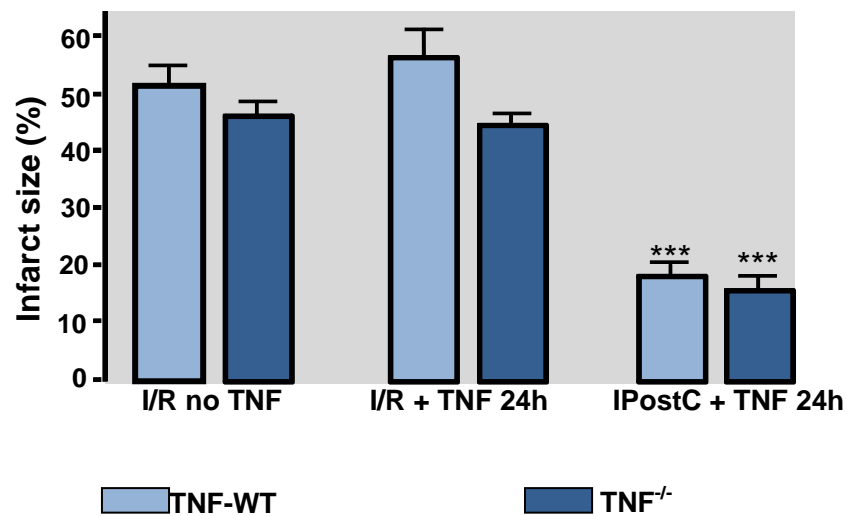


Fig 37: Restoration of protective effect in TNF $^{-/-}$ hearts. 24 hours after administration of a single ip injection of 0.5 ng TNF α , protection is restored in the isolated TNF $^{-/-}$ heart, *** $p < 0.001$ vs I/R+TNF 24h

➤ **Inhibition of STAT-3 but not Akt or Erk abrogates the protective effect of pharmacological postconditioning with TNF α**

To further investigate the mechanism responsible for the cardioprotective effect of TNF α , we measured infarct size in hearts perfused with an inhibitor of the JAK/STAT pathway (AG490), an inhibitor of the PI-3 kinase/Akt pathway (wortmannin) or an inhibitor of Mek/Erk1/2 (PD 98059), given during the first 15 min of reperfusion in IPostC or TNF-PostC hearts. The inhibitors did not affect the infarct size in the ischaemic control group. Addition of wortmannin (wort) or PD98059 did not alter the protective effect of TNF-PostC ($19 \pm 2\%$ and $20 \pm 1\%$ respectively, $p < 0.05$, versus $49 \pm 2\%$ in the ischaemia-reperfusion control group). Similarly, PD98059 did not affect the protective effect of IPostC ($29 \pm 2\%$, $p < 0.05$ versus I/R+PD;) but wortmannin completely abrogated the effect of IPostC ($43 \pm 1\%$, $p = \text{ns}$ versus $44 \pm 0.5\%$ for I/R+wort). Inhibition of the JAK/STAT pathway with AG490 during the IPostC or TNF-PostC protocols, abolished the infarct sparing effect of both postconditioning protocols ($47 \pm 2\%$ for IPostC; $p = \text{ns}$ versus $47 \pm 3\%$ for I/R and $51 \pm 2\%$ for TPostC, $p = \text{ns}$, versus $47 \pm 3\%$ for ischaemia-reperfusion control group), (Figure 38).

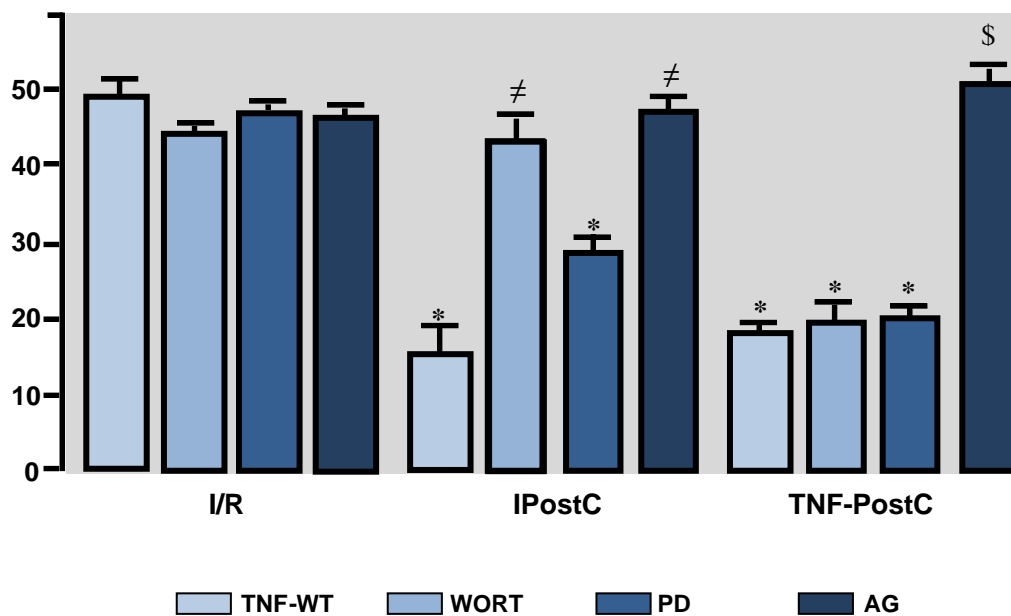


Fig 38: AG 490 abrogates TNF-PostC protection in TNF-WT mice. Inhibiting the activation of STAT-3 (by AG 490) during early reperfusion abrogates the protective effect of TNF-PostC in wild type mice. However, administration of wortmannin (a PI-3K inhibitor) or PD98059 (an inhibitor of Erk1/2) during early reperfusion did not abolish the postconditioning effect of TNF α ; $n = 6$ per group * $p < 0.05$ vs I/R alone; # $p < 0.05$ vs IPostC alone and \$ $p < 0.001$ vs TNF-PostC alone. Abbreviations as described in text.

➤ **TNF α failed to mimic ischemic postconditioning in STAT-3^{-/-} mice**

60

The infarct size in STAT-3-WT mice compared to the ischaemic control group was significantly decreased by pharmacological postconditioning with 6 x alternating cycles of 10 sec reperfusion without TNF α and 10 sec reperfusion with 0.5 ng/L TNF α initiated at the onset of reperfusion (20 \pm 1% for TNF-PostC versus 52 \pm 2% in I/R, p<0.05; Figure 39). In contrast, the STAT-3^{-/-} mice could not be postconditioned with TNF α (44 \pm 3% in I/R versus 47 \pm 1% in TNF-PostC, p = ns), (Figure 39).

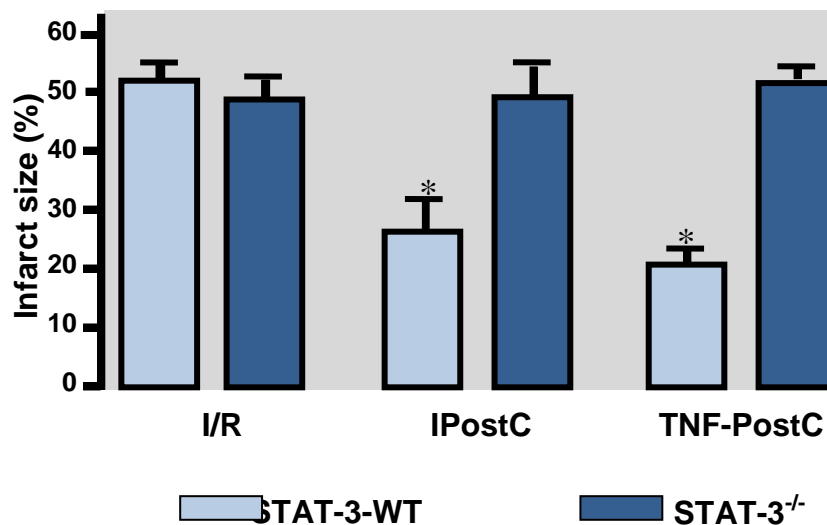


Fig 39: TNF α can mimic IPostC in STAT-3-WT but not STAT-3^{-/-} mice. 6 x 10 sec cycles of IPostC reduced the infarct size in STAT-3-WT but not STAT-3^{-/-} mice, n \geq 6 per group. Similarly, the infarct size in STAT-3-WT was reduced by 6 x 10 sec cycles of TNF-PostC but this failed to reduce the infarct size in STAT-3^{-/-} mice, n \geq 6 per group, *p<0.05 vs I/R. Abbreviations are as described in the text.

➤ **TNF α release from heart tissue after IPostC**

I/R reduced TNF α levels from 33.2 ± 4.4 pg/100mg tissue to 21.7 ± 1.9 pg/100mg tissue, $p < 0.05$
IPostC restored the TNF α levels to 36.7 ± 3.2 pg/100mg tissue, versus I/R, $p < 0.05$.

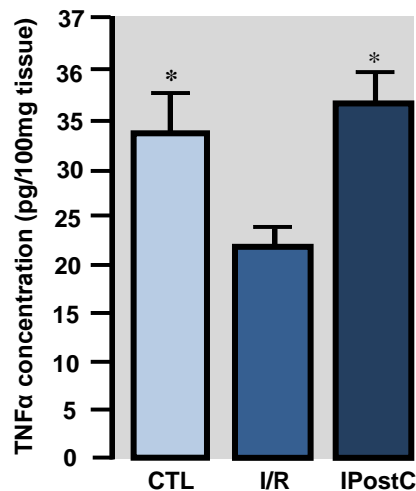


Fig 40: TNF α release by IPostC. The TNF α released from untreated hearts was 33.2 ± 4.4 pg/100mg tissue. IPostC restored TNF α release to 36.7 ± 3.2 pg/100 mg tissue whereas I/R significantly decreased the amount of TNF α released to 21.7 ± 1.9 pg/100mg tissue, $*p < 0.05$ versus I/R.

3.2 Western blot results

➤ **Levels of STAT-3, Akt, Erk1/2 and GSK-3 β in wild type and TNF $^{-/-}$ mice in nuclear and cytosolic fractions**

a) **Nuclear fraction**

We examined the levels of STAT-3, Akt, Erk1/2 and GSK-3 β during the first 15 min of reperfusion with no ischaemic episode (time-matched control) and after either IPostC or TNF-PostC. An increase in phosphorylated STAT-3 (in arbitrary units [AU] from 18 ± 2 for I/R to 54 ± 6 , $p < 0.05$), GSK-3 β (from 5 ± 0.4 for I/R to 15 ± 1 , $p < 0.05$) and Erk1/2 (from 9 ± 1 for I/R to 26 ± 2) occurred in the nuclear fraction after TNF-PostC in TNF-WT hearts. In contrast, Akt phosphorylation was decreased after TNF-PostC from 18 ± 2 for I/R to 6 ± 1 , $p < 0.05$. In TNF-

WT hearts subjected to IPostC, an increase in phosphorylation of STAT-3 (from 18 ± 2 for I/R to 46 ± 5 , $p < 0.05$) and GSK-3 β (from 5 ± 0.4 for I/R to 32 ± 2 , $p < 0.05$) was observed but the Akt level was unchanged. In TNF^{-/-} mice subjected to IPostC, a modest increase in phosphorylated STAT-3 occurred compared to the I/R group, but no increase in GSK-3 β was observed in the nuclear fraction after the postconditioning stimulus compared with the I/R group. Of note, phosphorylation of both Akt and Erk1/2 was significantly increased in the nuclear fraction by IPostC compared with I/R in TNF^{-/-} mice, (Figures 41a & 41b).

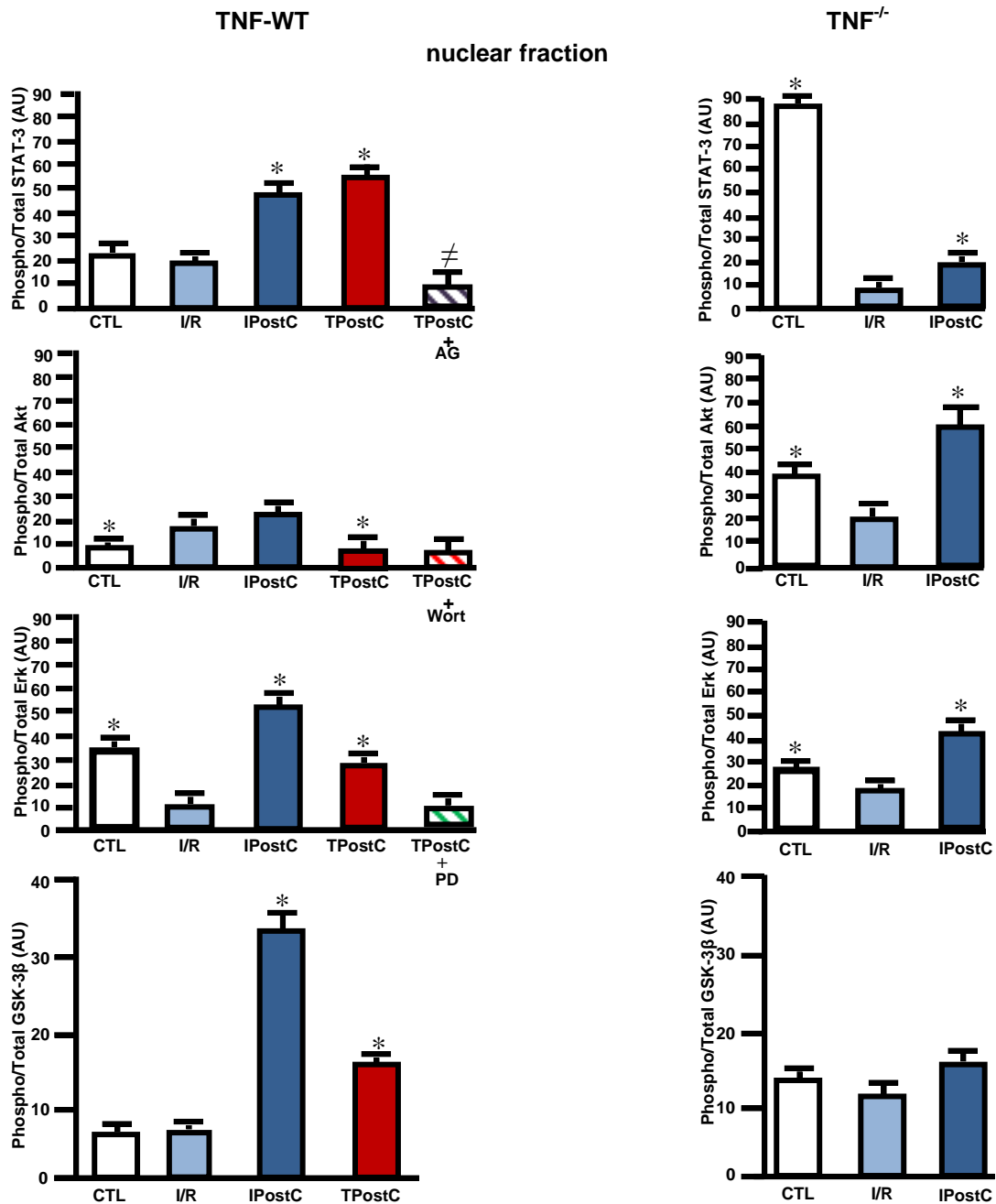
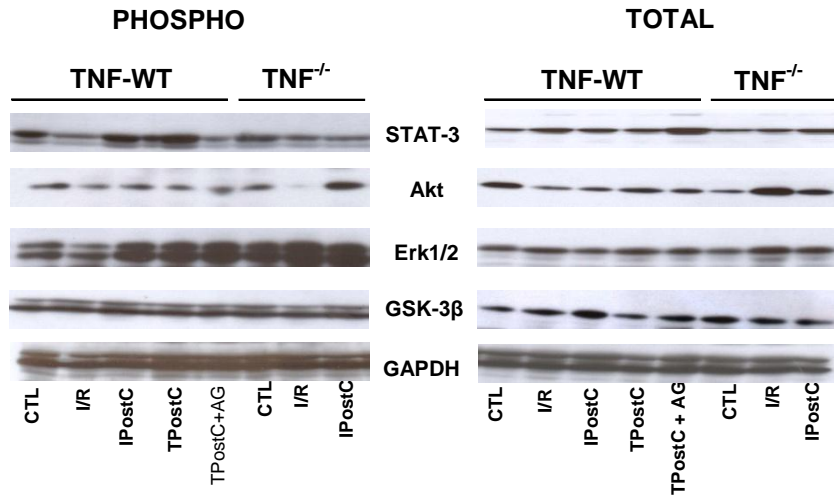


Fig 41a: Phosphorylation of STAT-3, Akt, Erk1/2 and GSK-3 β in nuclear fraction. Bar graphs of data derived from western blots of the nuclear fraction demonstrating phosphorylation of STAT-3, Akt and Erk1/2 after 15 minutes of reperfusion in isolated TNF-WT and TNF^{-/-} hearts; *p<0.05 versus I/R; #p<0.05 vs TPostC. Specific bands were quantified by densitometry and expressed in AU. Equal loading was confirmed with GAPDH Abbreviations as described in the text.



TNF-WT (nuclear fraction)

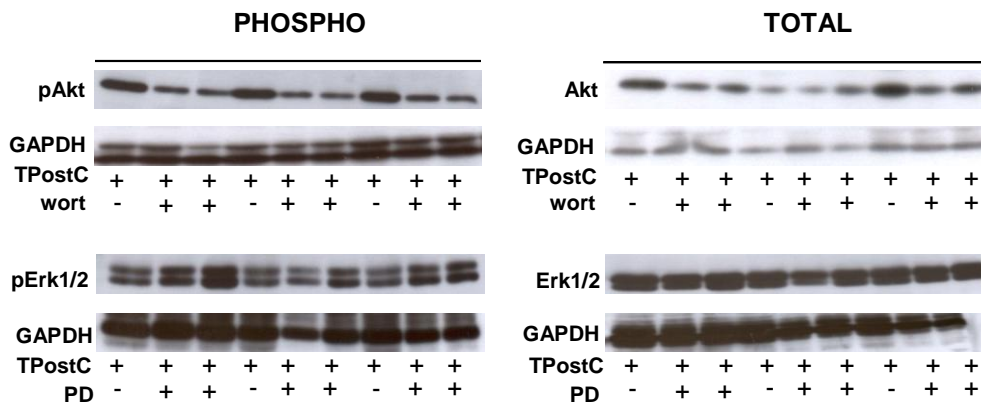


Fig 41b: Representative western blots of STAT-3, Akt, Erk1/2 and GSK-3 β in nuclear fraction. Representative western blots of the nuclear fraction demonstrating phosphorylation of STAT-3, Akt GSK-3 β and Erk1/2 after 15 minutes of reperfusion in isolated TNF-WT and TNF^{-/-} hearts; $p < 0.05$ versus I/R. Specific bands were quantified by densitometry and expressed in AU. Equal loading was confirmed with GAPDH. Abbreviations as described in the text.

b) Cytosolic fraction

In the cytosolic fraction, I/R increased phosphorylation of STAT-3 and decreased phosphorylation of Erk1/2 and Akt in wild type animals. IPostC reduced phosphorylated STAT-3 but increased the phosphorylation of both Akt and Erk1/2 in the cytosol compared to the I/R group. Pharmacological postconditioning with TNF α increased phosphorylation of Erk1/2 while phosphorylation levels of STAT-3 remained unchanged in the cytosol.

In the TNF^{-/-} mice, we were surprised to observe an increase of phosphorylation in both STAT-3 and Akt in the cytosol associated with a decrease in phosphorylated Erk1/2 following an IPostC stimulus, (Figure 42a & 42b).

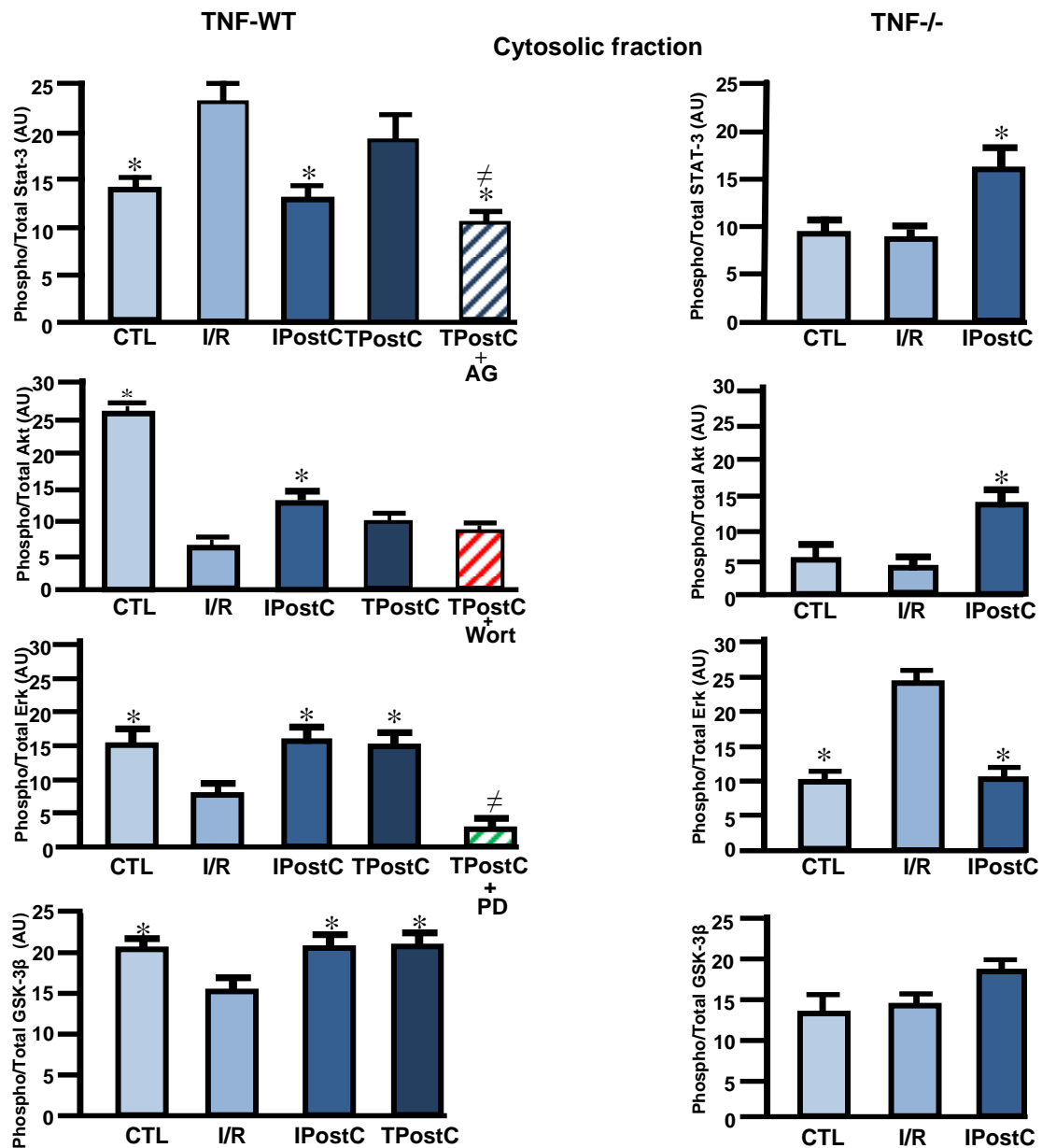


Fig 42a: Phosphorylation of STAT-3, Akt, Erk1/2 and GSK-3 β in cytosolic fraction. Bar graphs of data derived from western blots of the cytosolic fraction demonstrating phosphorylation of STAT-3, Akt and Erk1/2 after 15 minutes of reperfusion in isolated TNF-WT and TNF^{-/-} hearts; *p<0.05 versus I/R; #p<0.05 vs TPostC; n=6. Specific bands were quantified by densitometry and expressed in AU. Equal loading was confirmed with GAPDH. Abbreviations as described in the text.

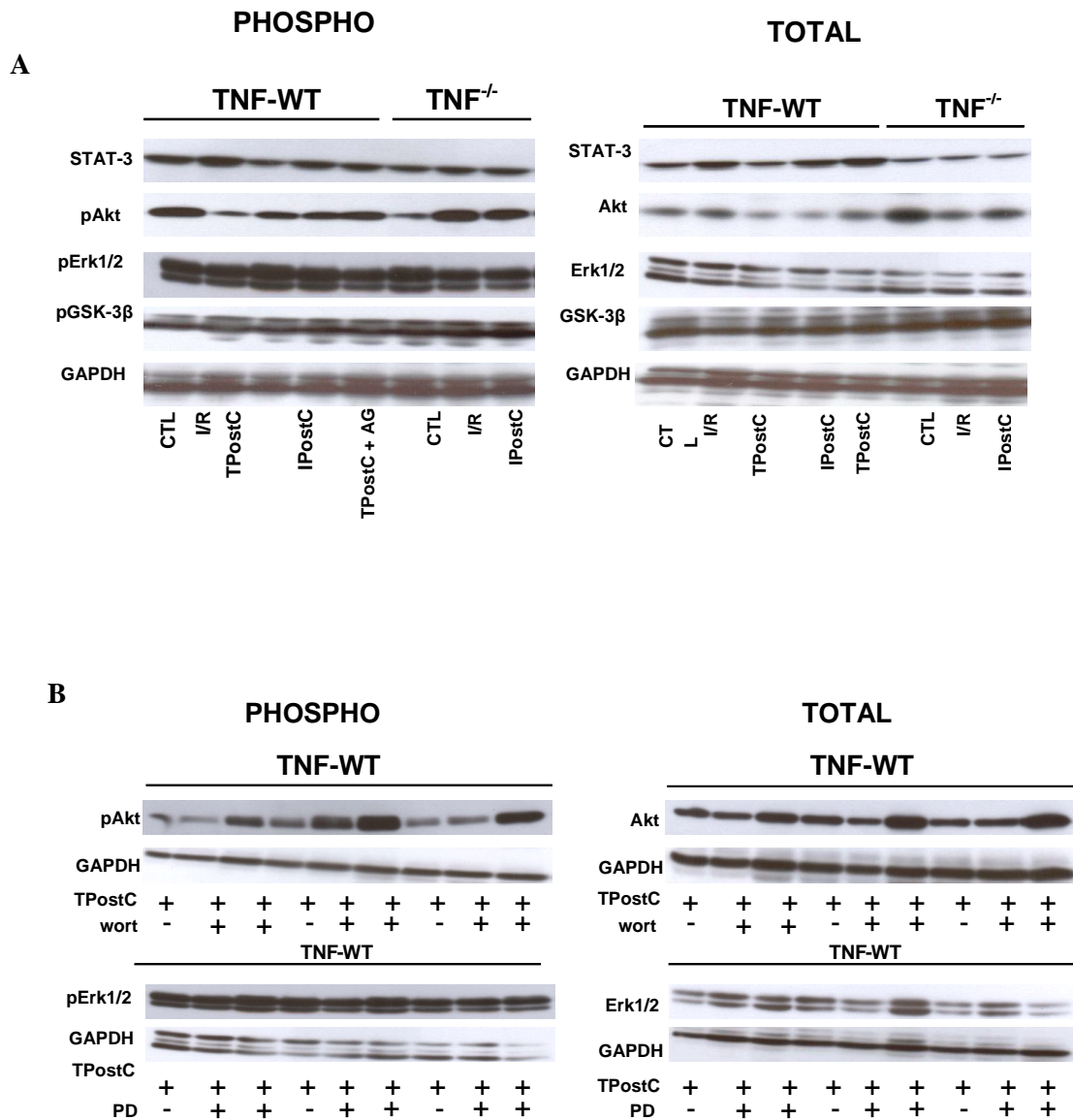


Fig 42b: Representative western blots of STAT-3, Akt, Erk1/2 and GSK-3β in cytosolic fraction. Representative western blots of the cytosolic fraction demonstrating phosphorylation of STAT-3, Akt and Erk1/2 after 15 minutes of reperfusion in isolated TNF-WT and TNF^{-/-} hearts; p<0.05 versus I/R. Specific bands were quantified by densitometry and expressed in AU. Equal loading was confirmed with GAPDH. Abbreviations as described in the text.

➤ **RAGE in TNF-WT and TNF^{-/-} mice**

a) Nuclear fraction

We examined the expression of total RAGE in nuclear fractions extracted from TNF-WT and TNF^{-/-} hearts, during the first 15 min of reperfusion after either IPostC or TNF-PostC. In TNF-WT hearts, IPostC decreased the expression of RAGE significantly in the nuclear fraction from 142.8 ± 9.6 AU in control hearts to 41.0 ± 4.5 AU in IPostC; $p < 0.05$. TPostC, however, restored the level of RAGE to a similar level seen in the controls, ($p = ns$). TNF^{-/-} hearts showed no significant difference in RAGE expression in relation to the treatment, (Figure 43).

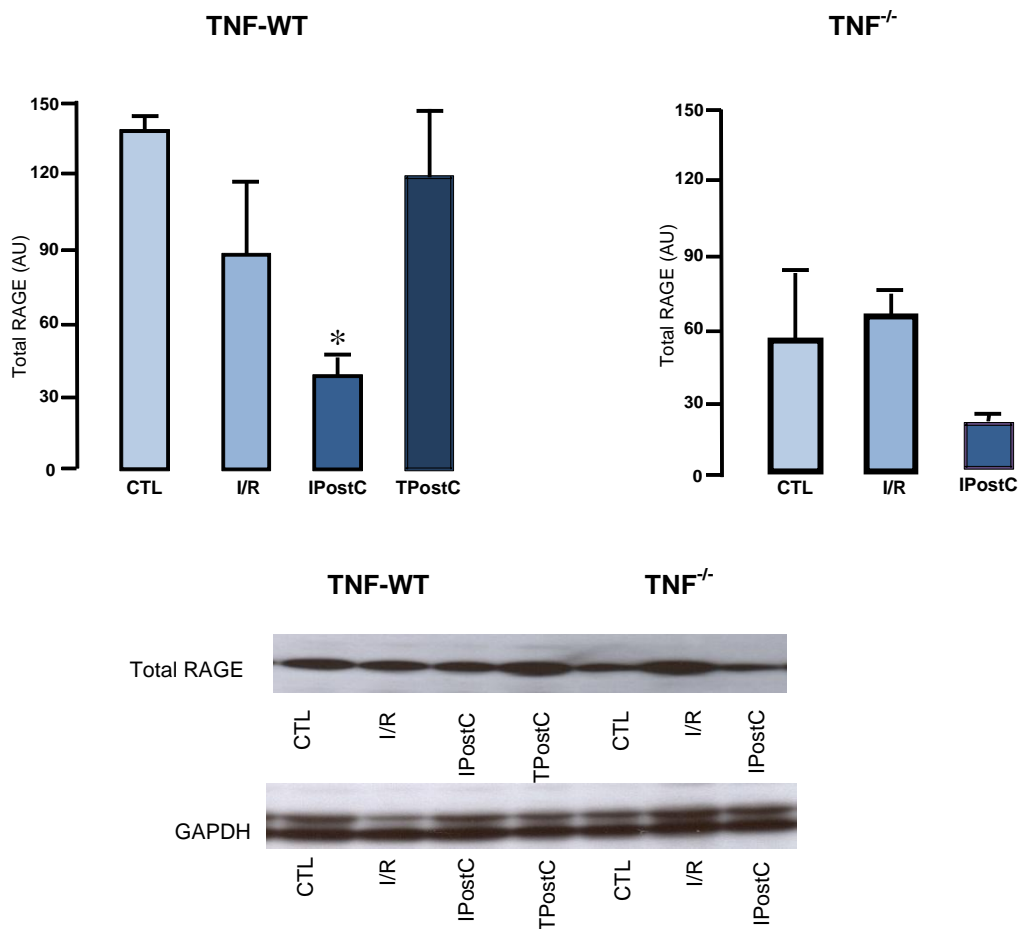


Fig 43: Expression of Total RAGE in nuclear fraction. Bar graphs of data derived from western blots and representative western blots demonstrating total RAGE expression after 15 minutes of reperfusion in isolated TNF-WT and TNF^{-/-} hearts; * $p < 0.05$ versus respective CTL; $n = 6$. Specific bands were quantified by densitometry and expressed in AU. Equal loading was confirmed with GAPDH. Abbreviations as described in text.

b) Cytosolic fraction

In the cytosolic fraction of TNF-WT hearts, RAGE expression after IPostC and TNF-PostC was significantly decreased versus the control group, from 530 ± 10 AU to 267 ± 100 AU, $p < 0.05$ and 116 ± 20 AU, $p < 0.01$, respectively. No significant differences were observed amongst the groups in TNF^{-/-} mice, $p = ns$. However, the I/R group of TNF^{-/-} hearts expressed significantly less RAGE compared with the I/R group of TNF-WT, ($p < 0.01$), (Figure 44).

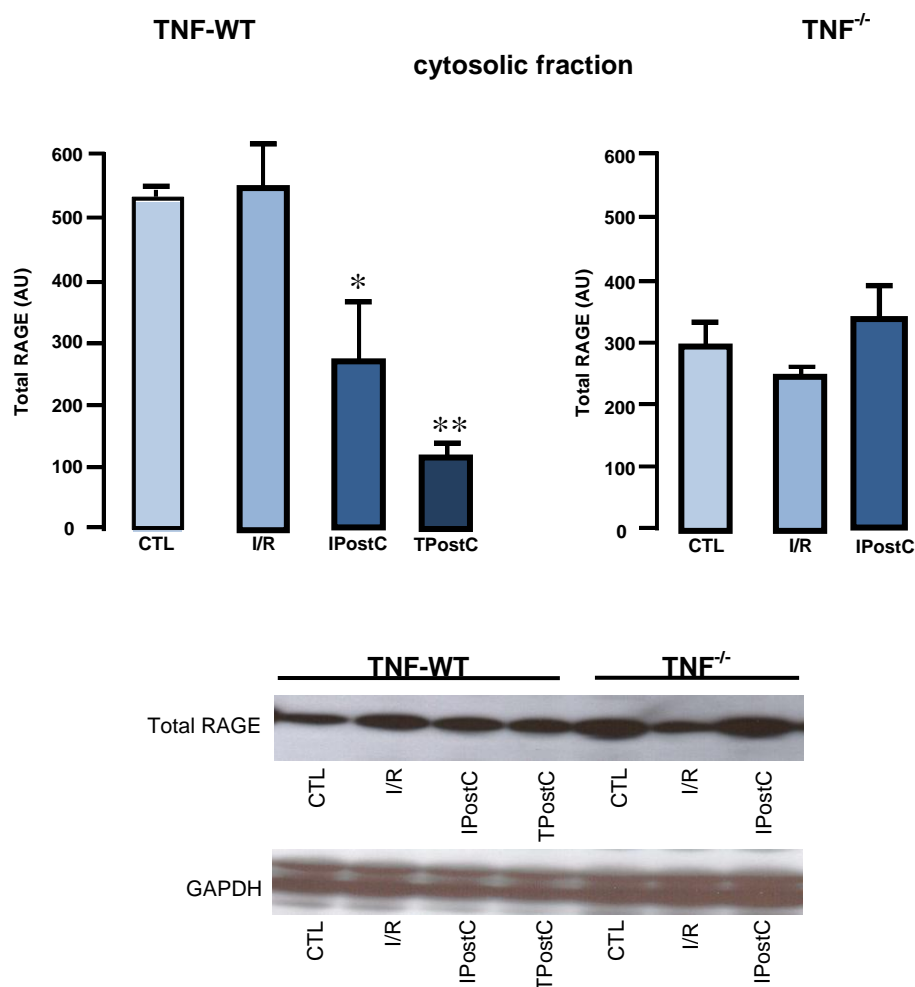


Fig 44: Expression of Total RAGE in cytosolic fraction. Bar graphs and representative western blots demonstrating total RAGE expression after 15 minutes of reperfusion in isolated TNF-WT and TNF^{-/-} hearts; * $p < 0.05$ versus TNF-WT I/R, ** $p < 0.01$ versus TNF-WT I/R. Specific bands were quantified by densitometry and expressed in AU. Abbreviations as described in text

➤ **Expression of Total PAR-2 in TNF-WT and TNF^{-/-} mice**

We examined the expression of PAR-2 in total protein extracted from TNF-WT and TNF^{-/-} hearts, during the first 15 min of reperfusion after either IPostC or TNF-PostC. In TNF-WT hearts, both I/R and IPostC decreased the expression of PAR-2 significantly to 76 ± 7 AU and 70 ± 12 AU respectively, when compared to control hearts (172 ± 21 AU) which underwent no interference, ($p < 0.05$). TPostC, however, restored the level of PAR-2 to a similar level as that of the controls, ($p = ns$). TNF^{-/-} control hearts had a significant decrease in PAR-2 expression (63 ± 5 AU) compared to hearts from TNF-WT control group (172 ± 21 AU), ($p < 0.01$). TNF^{-/-} I/R group significantly increased the PAR-2 expression to 160 ± 8 AU versus their control group, ($p < 0.05$), figures 45 & figure 46.

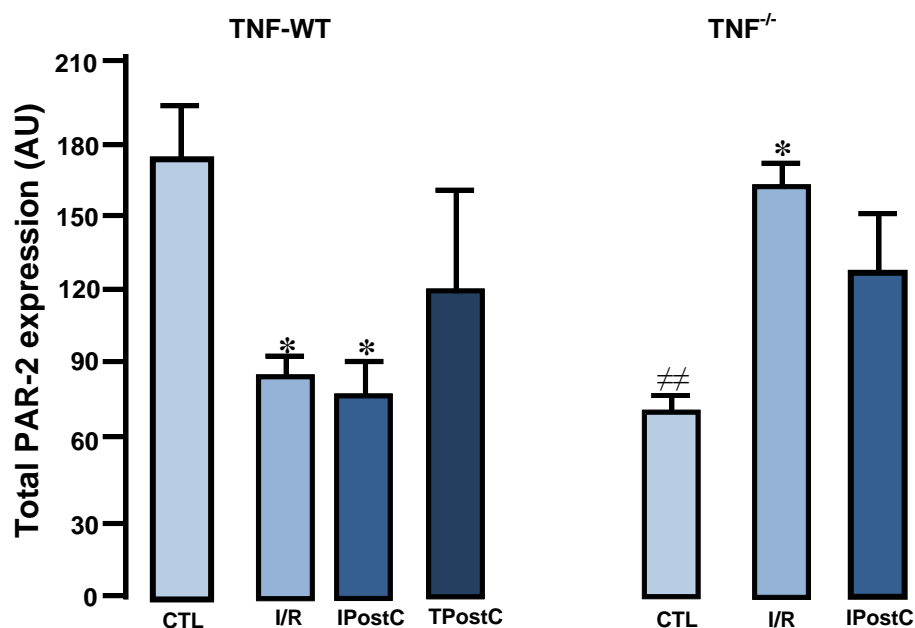


Fig 45: Expression of Total PAR-2. Bar graphs of data derived from western blots demonstrating total PAR-2 expression after 15 minutes of reperfusion in isolated TNF-WT and TNF^{-/-} hearts; * $p < 0.05$ versus respective CTL. CTL hearts from TNF^{-/-} had significantly decreased PAR-2 expression versus CTL hearts from TNF-WT, # $p < 0.01$. Specific bands were quantified by densitometry and all groups were corrected to GAPDH densities and expressed in AU. Abbreviations as described in the text.

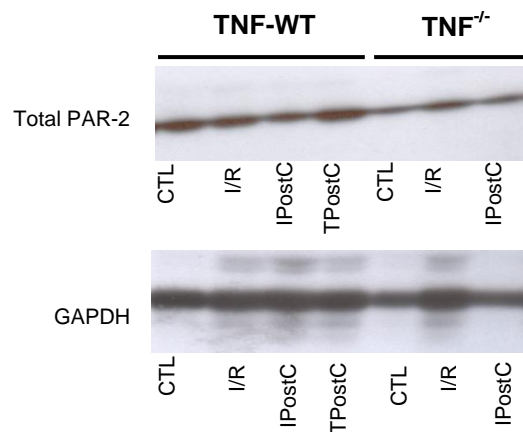


Fig 46: Representative western blots of Total PAR-2 expression. Representative western blots, demonstrating Total PAR-2 expression after 15 minutes of reperfusion in isolated TNF-WT and TNF^{-/-} hearts; $p < 0.05$ versus CTL. Specific bands were quantified by densitometry and all groups were corrected to GAPDH densities and expressed in AU. Abbreviations as described in the text.

4.0 Discussion

Using the different animal models, our data show, for the first time and possibly counter intuitively, that endogenous or exogenous TNF α , at the time of reperfusion, initiates a novel and RISK-free protective pathway in ischaemic postconditioning. In brief, six episodes of IPostC, performed by alternating cycles of reperfusion and ischaemia in TNF deficient mice, failed to protect against ischaemia-reperfusion damage. The activation of TNFR2 is implicated in the postconditioning protective effect as TNFR1^{-/-}, but not TNFR2^{-/-} mice, could be postconditioned. The protection of IPostC was reduced in the presence of wortmannin, PD98059 and AG490, while postconditioning with TNF α could only be abrogated by AG490. Furthermore, the STAT-3^{-/-} hearts could not be postconditioned. Our data demonstrate that protection with ischaemic postconditioning can be achieved via the activation of TNF α , its binding to TNFR2 and subsequent activation of STAT-3, key components of the prosurvival SAFE pathway.

4.1 Cardioprotective role of TNF α

Previously, we have consistently demonstrated, in both *in vitro* and *in vivo* experiments, that exogenous TNF α could confer a cardioprotective effect against reperfusion injury in a time and concentration dependent manner (see review (Lecour, 2009b)). Our group has shown that tissue levels of TNF α in the *in vivo* rat heart were significantly increased in the area at risk and that various doses of exogenous TNF α administered prior to the ischaemic protocol could be used as a preconditioning stimulus (Deuchar et al., 2007). Furthermore, a recent *in vivo* rat study demonstrated that the protection with IPostC is concomitant with decreased plasma levels of TNF α at reperfusion (Kin, 2008). Our present data demonstrate that TNF α , given immediately at the onset of reperfusion, confers protection in the isolated heart. This is in apparent contradiction to previous reports from isolated rat hearts, suggesting that the release of TNF α , as part of the inflammatory response during the reperfusion phase, contributes to the development of myocardial infarction, contractile dysfunction and apoptosis (Belosjorow et al., 1999; Kupatt et al., 1999). However, the recent demonstration of a biphasic kinetic pattern of TNF α release in a murine reperfusion model suggests that protection is conferred by the limited release of TNF α during the first few minutes of reperfusion (2-10min) (Reil et al., 2007). This limited amount of TNF α is then washed out and replaced with a more robust release of TNF α in the later stages of reperfusion. This subsequent increase in TNF α levels is thought to be responsible for its detrimental effects (Reil et al., 2007). In our present study we show, for the first time, in an

isolated murine heart model, that during either IPostC or TNF-PostC, the binding of TNF α to TNFR2 at the early stage of reperfusion is crucial for protection. Therefore, a possible explanation of our potentially controversial findings is that the synergetic limited early release/addition of TNF α , together with its binding to TNFR2, at the onset of reperfusion, may limit further harmful production of TNF α during the later stages of reperfusion.

Using TNF $^{-/-}$ mice in our experiments, we were surprised to observe that TNF α postconditioning in these mice failed to protect the heart. In a previous study, our research group demonstrated that the administration of TNF α to TNF knockout mice, 24 hours prior to experimentation rescued the protective effect of ischaemic preconditioning in these mice (Smith et al., 2002). These data suggest that in TNF knockout mice, the signalling pathways downstream of TNF α may be downregulated. Similar data were obtained in our postconditioning setting in the TNF $^{-/-}$ mice. In these mice the protective effect of TNF-PostC could be restored following administration of 0.5 ng TNF α , ip 24 hours prior to the experiments. Further functional studies are necessary to evaluate the TNF receptor densities and receptor-ligand interaction to fully explain this ability to rescue the protective effect in TNF $^{-/-}$ hearts.

4.2 TNF α initiates the activation of a prosurvival pathway independent of the RISK pathway

An established protective pathway, known as the RISK pathway and involving the activation of protein kinases Akt and Erk1/2 in early reperfusion, has been described in both pre- and postconditioning (Hausenloy et al., 2005; Tsang et al., 2004).

A recent study done in pig hearts reported an activation of the kinases involved in the RISK pathway following an ischemic postconditioning stimulus (Skyschally et al., 2009). However, the inhibition of these kinases did not alter the protective effect, therefore questioning the essential role of the RISK kinases in their model (Skyschally et al., 2009). Similarly, our present study, performed on a mouse model, demonstrates that pharmacological postconditioning with TNF α is independent of the RISK pathway, as neither PD98059 nor wortmannin abolished the protective effect of TNF-PostC. In contrast, the protective effect of both IPostC and pharmacological postconditioning with TNF α was abolished in the presence of AG490, the JAK/STAT pathway inhibitor. In addition, neither ischaemic nor pharmacological postconditioning with TNF α could protect in STAT-3 deficient mice.

Our new data show the existence of an alternative protective pathway during the reperfusion period of IPostC. This pathway, previously described in ischaemic preconditioning, and involving the activation of both TNF α and STAT-3, is now recognized as the SAFE pathway (Lecour, 2009a; Lecour, 2009b). Our present data strongly support the hypothesis that the recent studies describing a RISK-independent protective effect of ischaemic postconditioning may in fact activate the SAFE pathway (Przyklenk et al., 2008; Schwartz and Lagranha, 2006; Skyschally et al., 2009).

Current and past studies, done in our laboratory, suggest that inhibiting either the SAFE or RISK pathway is enough to abolish the infarct sparing effect of ischaemic pre or postconditioning (Fuglestad et al., 2008; Lecour et al., 2002; Suleman et al., 2008). Attempts to delineate a possible crosstalk between the two pathways reveal a complex interaction. Hence, in the presence of wortmannin during the ischaemic preconditioning stimulus, phosphorylation of STAT-3 was inhibited (Suleman et al., 2008). Similarly, Akt failed to be phosphorylated in STAT-3 knockout mice subjected to ischaemic preconditioning (Suleman et al., 2008). In contrast, a substantial increase of phosphorylated Akt is observed in TNF $^{-/-}$ mice subjected to ischaemic postconditioning (data from this study). Further studies are needed to characterize the crosstalk between the SAFE and RISK pathways in both ischaemic pre and postconditioning.

4.3 Possible downstream targets of SAFE pathway

Inactivation of GSK-3 β is a well established downstream target of the protective pathway (Gross et al., 2006; Hausenloy and Yellon, 2007a; Juhaszova et al., 2004; Juhaszova et al., 2009). Our present work aimed to delineate whether GSK-3 β could be a common target for the SAFE and RISK pathways. In our model, both ischaemic postconditioning and pharmacological postconditioning with TNF α increased the phosphorylation of GSK-3 β in WT mice. In contrast, we were surprised to observe that the absence of an infarct sparing effect in TNF $^{-/-}$ mice was associated with a similar level of Phospho/Total GSK-3 β as that of wild type mice subjected to the TNF-PostC protocol. Although the data from this study support the proposal suggesting that inactivation of GSK-3 β is not an essential link in the cardioprotective effect of ischaemic postconditioning, this still remains controversial (Juhaszova et al., 2009; Nishino et al., 2008). Further studies using GSK-3 β genetically modified mice will be required to confirm whether GSK-3 β is, or is not, an effective downstream target of the TNF α /JAK/STAT-3 pathway.

In contrast to previous studies (Napoli et al., 2000; Napoli, 2002) the expression of RAGE was not increased by I/R in our TNF-WT model, although it showed a slight increase in TNF^{-/-} mice. In addition, we observed that IPostC gave a significant reduction in RAGE expression whereas with TNF-PostC the expression returned to baseline levels. TNF^{-/-} mice also exhibited this significant decrease in RAGE during IPostC, suggesting that RAGE is not a downstream target of the SAFE pathway.

PAR-2 levels were decreased significantly by I/R and IPostC but were restored by TNF-PostC. In TNF^{-/-} mice I/R increased PAR-2 expression significantly. Although there was also an increase in PAR-2 expression after IPostC this was not considered significant. Our data suggest that PAR-2 is not a downstream target of the SAFE pathway.

Additional possible downstream targets of the SAFE pathway include phosphorylation of the pro-apoptotic target Bad (Deuchar et al., 2007; Lecour et al., 2005b) and the inactivation of NFκB (Kin, 2008; Nishinakamura et al., 2007). TRAF2 has also been recently implicated in the action of TNFα on NFκB by promoting the binding of TNFα to TNFR2 and discouraging the binding to TNFR1, thereby preventing TNFα-induced apoptosis (Vince et al., 2009). The inhibition of the mitochondrial permeability transition pore opening (Gao et al., 2004) is likely to be the common point at which the SAFE and RISK pathways converge.

4.4 Other alternative pathways operative in postconditioning

The presence of multiple protective pathways has already been suggested in ischaemic preconditioning (Jin et al., 2004; Lecour et al., 2005b) and it is likely that the SAFE and RISK pathways are activated in parallel with other paths in ischaemic postconditioning. Hence, sphingolipids have been proposed as members of a downstream activated path in ischaemic postconditioning (Jin et al., 2008; Vessey et al., 2008). Sphingosine-1-phosphate can activate Akt, Erk1/2 and STAT-3 in the heart (Frias et al., 2009; Radeff-Huang et al., 2004; Vessey et al., 2008). While TNFα protects via the activation of the sphingolipids, (Heusch et al., 2008) and preliminary data in our laboratory suggest that sphingosine-1-phosphate mimics ischaemic preconditioning via the activation of STAT-3 (Kelly, 2008), the existence of sphingosine-1-phosphate as an alternative path to the SAFE or RISK pathway still needs to be confirmed.

In summary, our studies have consistently shown that activation of the SAFE pathway, as represented by the activity of TNFα and STAT-3 is required in both ischaemic pre- and

postconditioning (Deuchar et al., 2007, Baxter, 2003#242 ; Lecour et al., 2002; Lecour et al., 2005b; Smith et al., 2002; Suleman et al., 2008). The specific involvement of TNFR2 in our studies suggests that this protective pathway is linked to this receptor. Overall, these results further emphasize the importance of RISK-independent pathways in cardiac protection at the time of reperfusion, which may have potential therapeutic application in the mitigation of ischaemic-reperfusion injury. A schematic of the SAFE pathway in postconditioning, based on the data from the present study, is shown in figure 47.

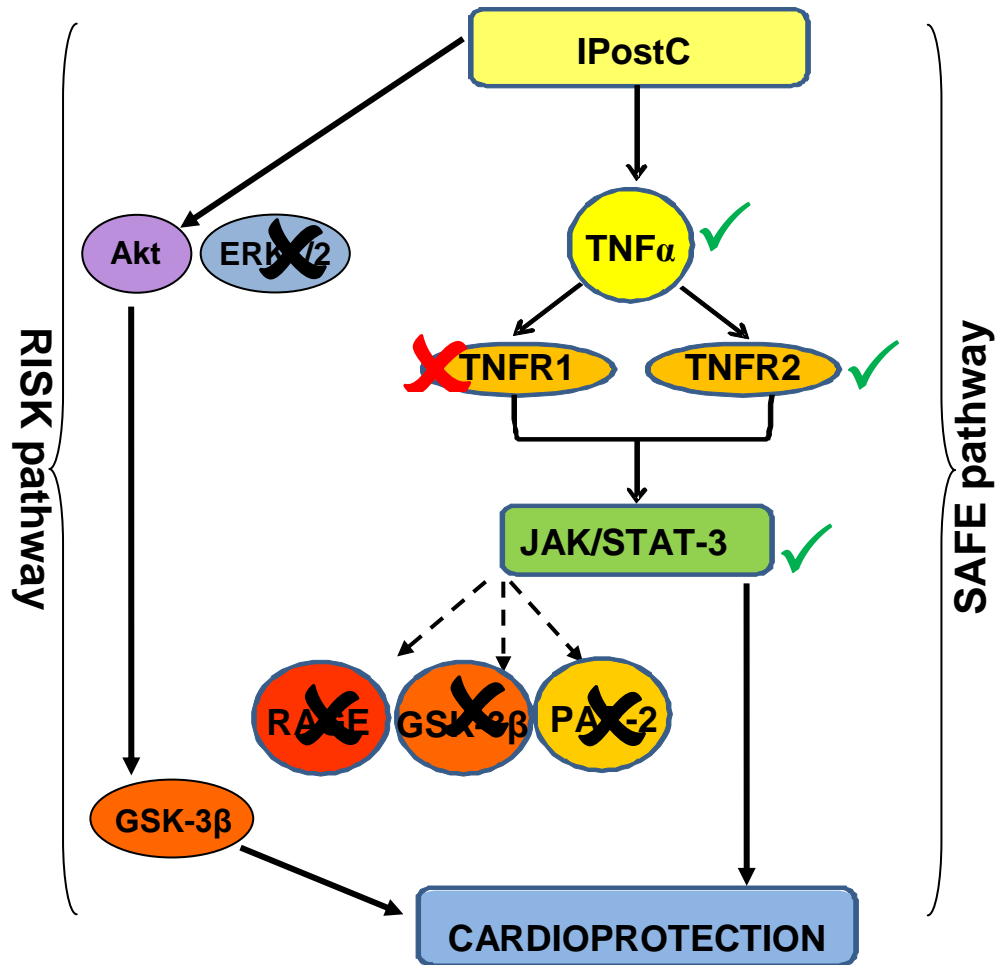


Fig 47: Schematic of SAFE pathway. TNF α activates JAK/STAT-3 via TNFR2 to confer cardioprotection. RAGE, GSK-3 β and PAR-2 are not downstream signalling targets of STAT-3 in the SAFE pathway. The RISK pathway signals through Akt and GSK-3 β .

E. CARDIOPROTECTIVE AGENTS AS POSTCONDITIONING MIMETICS

Part of the experiments in this section have been performed in conjunction with Sarin Somers, registered as a PhD student at the University of Cape Town

1.0 Introduction

Ischaemic postconditioning with intermittent episodes of ischaemia and reperfusion at the onset of reperfusion requires precise and timed applications that are impractical to apply in a clinical setting. Pharmacological agents such as bradykinin, opioids and adenosine can be substituted as preconditioning mimetics to reduce infarct size following an ischaemia-reperfusion insult but their role as postconditioning mimetics has been poorly studied (Cohen, 2001; Gross, 2006; Downey, 2007). These three mediators are known to protect via the activation of the RISK pathway (Bell, 2003; Riksen, 2004; Fryer, 2002) and several studies conducted in a preconditioning setting suggest that divergent pro-survival signalling pathways may be activated by these components. Hence, bradykinin and opioids, but not adenosine, require the activation of the mK_{ATP} channel and free radicals to confer cardioprotection in IPC (Cohen, 2001).

In the previous section, our data demonstrate that activation of the prosurvival SAFE pathway, with $TNF\alpha$ and STAT-3, acts as an alternative signalling pathway to confer cardioprotection in IPostC. The role of $TNF\alpha$ as a downstream signalling target of bradykinin, opioids and adenosine has never been investigated.

Therefore, the purpose of our study is to explore whether adenosine, bradykinin and opioids can mimic IPostC in an isolated mouse heart model. In addition, we will use $TNF^{-/-}$, $TNFR1^{-/-}$, $TNFR2^{-/-}$ and cardiomyocyte specific STAT-3 deficient mice ($STAT-3^{-/-}$) to demonstrate that this pharmacological postconditioning effect can be attributed, at least in part, to the activation of the prosurvival SAFE pathway initiated by $TNF\alpha$ and STAT-3. A recent study suggested the possibility that STAT-3 can translocate to the mitochondria after its phosphorylation in the cytosol (at the serine site) (Wegrzyn, 2009). As mitochondria are considered as the end target for the protection afforded with postconditioning (Heusch, 2010), we will use western blot techniques to investigate the possible activation and translocation of STAT-3 to the mitochondria following a pharmacological postconditioning stimulus.

Our aim, therefore, was to firstly delineate whether adenosine, bradykinin and opioids could mimic IPostC in an isolated mouse heart model and secondly, to explore whether this postconditioning effect could be attributed, at least in part, to the activation of the SAFE pathway via translocation of STAT-3 to the cardiac mitochondria, (Figure 48).

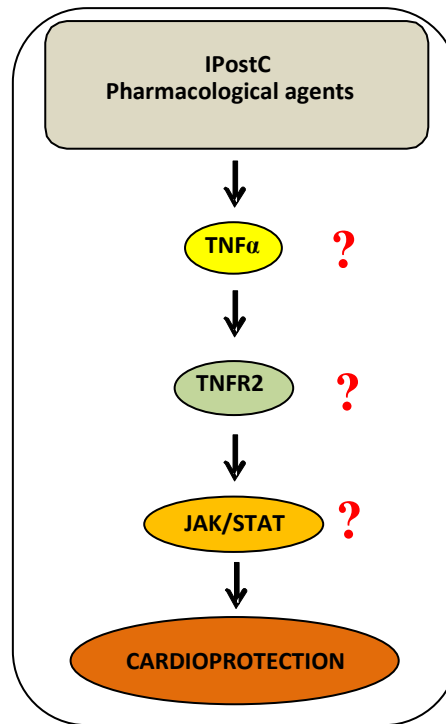


Fig 48: Schematic of proposed hypothesis. Pharmacological agents activate TNF α via TNFR2 and initiate the phosphorylation of JAK/STAT and subsequent cardioprotection.

2.0 Methods

2.1 Perfusion of mouse hearts

All animal studies performed were approved by the Animal Research Ethics Committee of the University of Cape Town, and followed the recommendations laid down in the Guide for the Care and Use of Laboratory Animals published by the US National Institute of Health (NIH Publication No. 85 {23}, revised 1996). Six mice were used per experimental group for each genotype of mouse.

Isolated hearts of adult male $TNF^{-/-}$, $TNFR1^{-/-}$, $TNFR2^{-/-}$ and $STAT-3^{-/-}$ mice and their respective WT were subjected to the identical retrograde perfusion technique as described in Section C. Postconditioning with each of the well known cardioprotective agents as postconditioning mimetics was performed in the same manner as we have described in Section D, with 6 x 10 second alternating cycles of reperfusion and reperfusion together with each of the drugs in turn as outlined in the following figure 49.

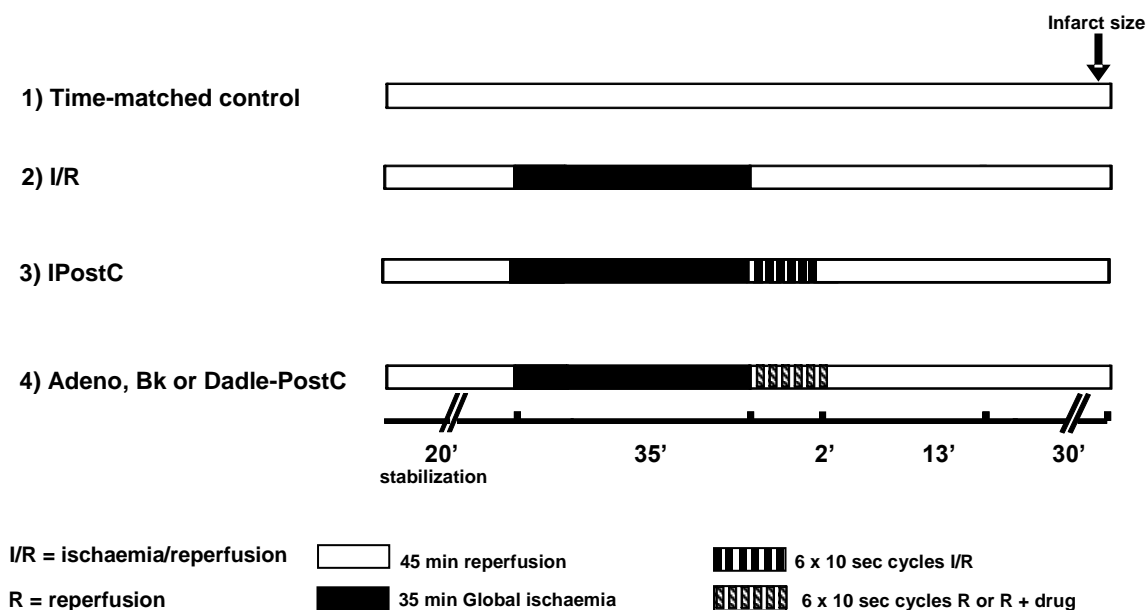


Fig 49: Schematic of Langendorff experimental model. Isolated hearts were subjected to a stabilization period of 20 min followed by 35 min global ischaemia and a 45 min reperfusion period. At onset of reperfusion, ischaemic postconditioning (IPostC) was initiated with 6 alternating cycles of 10 sec reperfusion, 10 seconds ischaemia. In a separate group, postconditioning with 100 μ M adenosine (Adeno-PostC), 100 nM Bradykinin (Bk-PostC) or 100 nM DADLE (D-PostC) was initiated with 6 alternating cycles of 10 sec reperfusion with no drug, 10 seconds reperfusion with the drug of choice.

2.1.2 Infarct size assessment

At the end of each experimental protocol, hearts were removed from the perfusion apparatus and infarct size assessed by triphenyltetrazolium chloride (TTC) staining and planimetry as described previously in Section C.

2.2 Postconditioning with classic cardioprotective agents

Adenosine (100 μ M), bradykinin (100 nM), and 100 nM of the delta opioid agonist, D-Ala 2', D-Leu 5'-enkephalin acetate salt (DADLE) were used as postconditioning mimetics in hearts from TNF^{-/-}, TNFR1^{-/-}, TNFR2^{-/-} and STAT-3^{-/-} mice and their respective wild types. Six alternating cycles of 10 seconds reperfusion, 10 seconds reperfusion with pharmacological agent was initiated at onset of reperfusion (Figure 49). A minimum of 6 hearts was used in all groups.

2.3 Western blot analysis

Control or murine hearts postconditioned with bradykinin, adenosine or DADLE were subjected to a 20 min stabilization period followed by 35 min of global ischemia and 15 min of reperfusion. The hearts were then snap-frozen in liquid nitrogen before being stored at -80°C until protein extraction was performed. To determine whether the activation of the prosurvival signalling cascade initiated the translocation of activated STAT-3 from the cytosol to the mitochondria we extracted mitochondrial and cytosolic proteins from the hearts by homogenization of the tissue in a lysis buffer, as previously described in Section C. Total and phosphorylated states of STAT-3 (Phospho-STAT-3, ser 737), were analysed by sodium dodecyl sulphate-polyacrylamide gel immunoelectrophoresis (SDS-PAGE) using antibodies from Cell Signaling Technologies and Santa Cruz Biotechnology, Inc. USA.. Equal loading for the mitochondrial fraction was verified with an antibody for the voltage dependent anion channel (VDAC). GAPDH was used as a loading control for the cytosolic fraction. The ratio of phosphorylated STAT-3 translocated from the cytosolic fraction to the mitochondrial fraction was calculated. Relative densitometry was determined with the use of computerized software package (ImageJ analysis and processing in Java). A minimum of 6 hearts was used per group.

2.4 Chemicals and Pharmacological Agents

Unless otherwise stated, all chemicals were obtained from Sigma-Aldrich Chemicals, Germany.

2.5 Statistical Analysis

Data are presented as mean \pm SEM. Comparisons between multiple groups were performed by 1-way ANOVA followed by Student-Newman-Keuls Multiple Comparison Test (Graph Pad Instat). $p < 0.05$ was considered statistically significant.

3.0 Results

3.1 Postconditioning with classic cardioprotective agents

In TNF-WT mice, ischaemic postconditioning (IPostC) reduced infarct size from $48 \pm 1\%$ in the ischaemic control group (I/R) to $16 \pm 1\%$ in IPostC, ($p < 0.001$). Bradykinin postconditioning (Bk-PostC), DADLE postconditioning (D-PostC) and Adenosine postconditioning (Adeno-PostC) reduced the infarct size in a similar manner to TNF-WT hearts, ($p < 0.001$ versus I/R). In contrast, (IPostC), Bk-PostC and D-PostC failed to protect hearts from TNF^{-/-} mice, ($p = ns$), whereas adenosine protected the TNF-deficient hearts against ischaemia-reperfusion injury, decreasing infarct size from $45 \pm 1\%$ in I/R to $19 \pm 1\%$ in Adeno-PostC, ($p < 0.001$), figure 50.

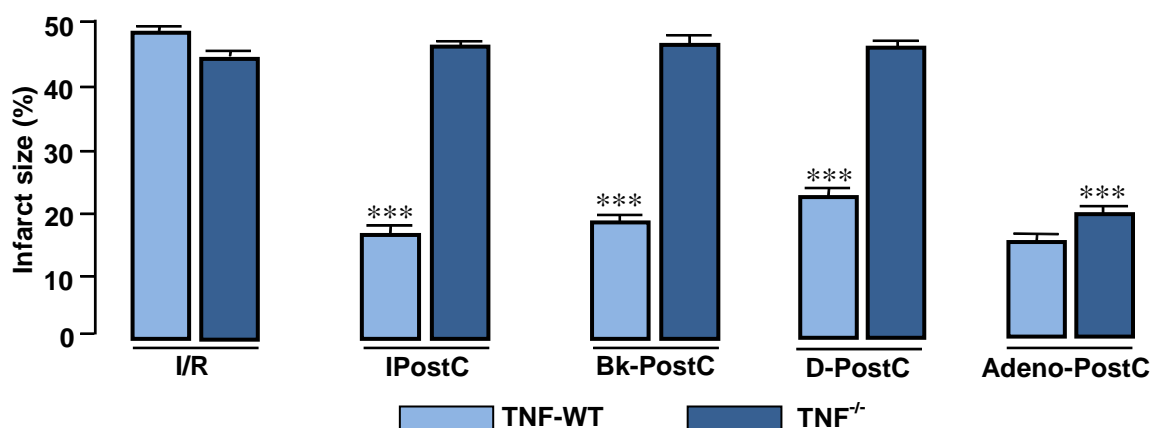


Fig 50: Adenosine protects TNF^{-/-} mice. IPostC decreased infarct size in TNF-WT from $48 \pm 1\%$ to $16 \pm 1\%$ versus I/R. Similarly, Bk-PostC, D-PostC and Adeno-PostC reduced infarct size in TNF-WT hearts to $19 \pm 1\%$, $24 \pm 1\%$ and $15 \pm 1\%$, respectively. Postconditioning with adenosine protected TNF^{-/-} hearts against ischaemia-reperfusion injury, reducing infarct size from $45 \pm 1\%$ to $19 \pm 1\%$, $***p < 0.001$ vs respective I/R control. In contrast, IPostC, Bk-PostC and D-PostC failed to protect hearts from TNF^{-/-} mice, $p = ns$. $n = 6$ per group. IPostC=ischaemic postconditioning; Bk-PostC=bradykinin postconditioning; D-PostC=DADLE postconditioning; Adeno-PostC=adenosine postconditioning.

3.2 Pharmacological postconditioning with bradykinin and DADLE protects TNFR1^{-/-} but not TNFR2^{-/-} mice

Infarct size measured after I/R in TNFR1^{-/-} and TNFR2^{-/-} were similar to WT mice. IPostC reduced infarct size in TNF-WT and TNFR1^{-/-} hearts to 19 ± 1% and 20 ± 1%, respectively versus their respective I/R controls, (***)p<0.001). In contrast, TNFR2^{-/-} hearts failed to be protected, (p=ns versus I/R). Postconditioning with 100 nM bradykinin (Bk-PostC) and 100 nM DADLE (D-PostC) significantly decreased infarct size in TNFR1^{-/-} hearts from 50 ± 1% in I/R to 21 ± 1% and 20 ± 1%, respectively, (***)p<0.001). However, TNFR2^{-/-} hearts could not be protected by either pharmacological agent, (p=ns versus I/R), figure 51.

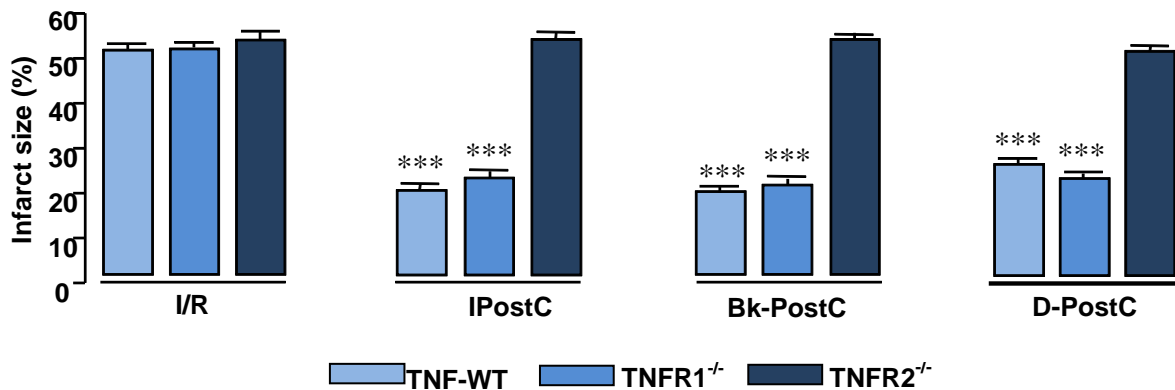


Fig 51: Bradykinin and DADLE postconditioning protect TNFR1^{-/-} but not TNFR2^{-/-}: IPostC significantly decreased infarct size in TNF-WT and TNFR1^{-/-} hearts vs their respective I/R controls. Similarly, Bk-PostC and D-PostC decreased infarct size in TNFR1^{-/-}, (***)p<0.001 versus respective I/R controls). In contrast, hearts from TNFR2^{-/-} mice failed to be protected, (p=ns); n=6 per group. IPostC=ischemic postconditioning; Bk-PostC=bradykinin postconditioning; D-PostC=DADLE postconditioning.

3.3 Bradykinin, DADLE and Adenosine fail to protect STAT-3^{-/-} hearts

IPostC in STAT-3-WT mice reduced infarct size from $48 \pm 2\%$ in the ischaemic control group (I/R) to $23 \pm 5\%$ for IPostC, ($p < 0.001$). Bk-PostC, D-PostC and Adeno-PostC reduced the infarct size in a similar manner to STAT-3-WT hearts, from $48 \pm 2\%$ in I/R to $18 \pm 3\%$, $18 \pm 2\%$ and $17 \pm 2\%$, respectively, ($p < 0.001$ versus I/R). In contrast, STAT-3^{-/-} mice failed to be protected by the pharmacological agents, ($p = ns$).

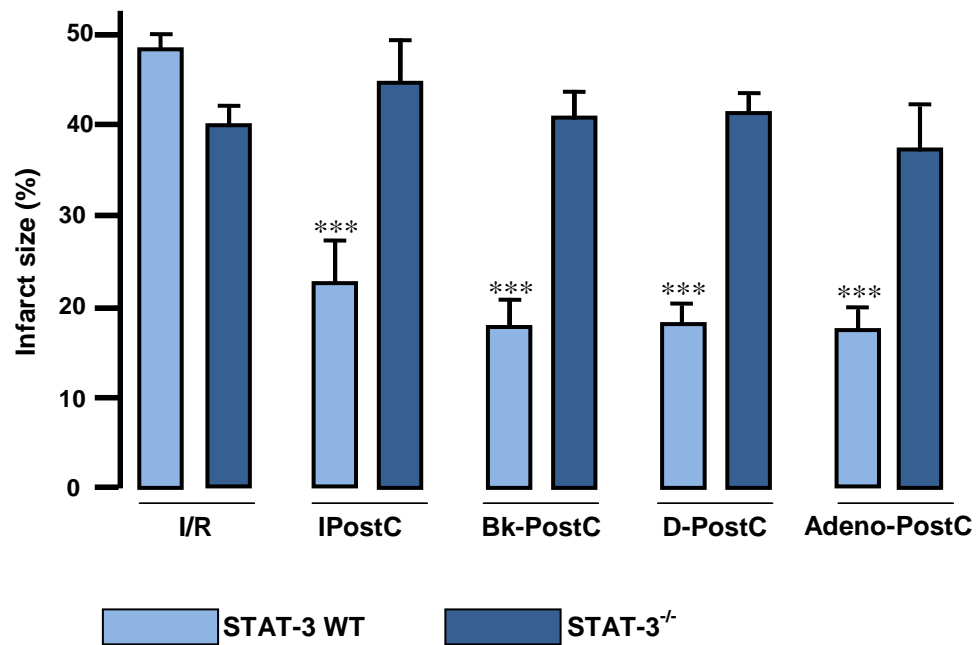


Fig 52: Pharmacological postconditioning in STAT-3-WT and STAT-3^{-/-} mice. IPostC protected STAT-3-WT from I/R, *** $p < 0.001$. Similarly, Bk-PostC, D-PostC and Adeno-PostC reduced infarct size in STAT-3-WT hearts (*** $p < 0.001$ versus I/R). In contrast, none of the postconditioning protocols protected the STAT-3^{-/-} hearts against ischaemia-reperfusion injury, $p = ns$; $n = 6$ per group; IPostC=ischaemic postconditioning; Bk-PostC=bradykinin postconditioning; D-PostC=DADLE postconditioning; Adeno-PostC=adenosine postconditioning.

3.4 Western blot analysis

➤ Levels of pSTAT-3 in mitochondrial and cytosolic fractions of TNF-WT and TNF^{-/-}

We examined the translocation of phosphorylated STAT-3 from the cytosolic fraction to the mitochondrial fraction after pharmacological postconditioning. An increase in the ratio of mitochondrial pSTAT-3 to cytosolic pSTAT-3 (expressed in arbitrary units) was observed with Bk-PostC and D-PostC from 156 ± 4 in I/R to 220 ± 2 and 280 ± 4 , respectively, ($p < 0.05$). There was no significant translocation of pSTAT-3 observed when adenosine was used as a postconditioning mimetic, ($p = ns$), figure 53. A similar trend was observed in TNF^{-/-} mice. VDAC was used as evidence for the mitochondrial fractions.

a) Mitochondrial and cytosolic fractions

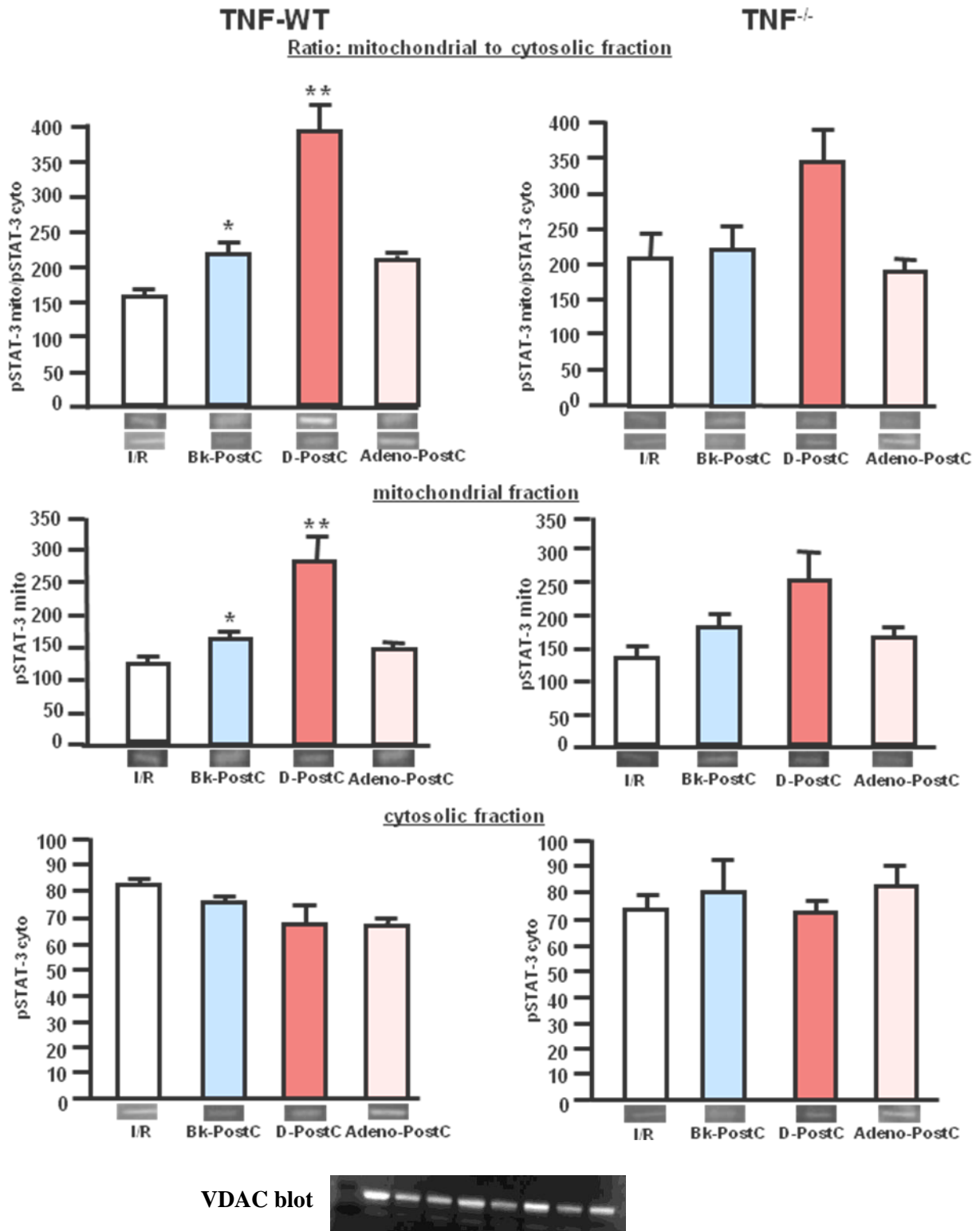


Fig 53: Phosphorylation and translocation to mitochondria of STAT-3. Bar graphs and representative western blots of mitochondrial and cytosolic fractions demonstrating the translocation of pSTAT-3, * $p < 0.05$ vs I/R; ** $p < 0.01$ vs I/R. The VDAC blot is evidence of the mitochondrial fraction. $n = 4$. IPostC=ischemic postconditioning; Bk-PostC=bradykinin postconditioning; D-PostC=DADLE postconditioning; Adeno-PostC=adenosine postconditioning.

4.0 Discussion

The novel data in this section show that bradykinin and opioids, given as true postconditioning mimetics, can mimic the cardioprotective effects of IPostC via the activation TNF α binding to TNFR2, and leading to subsequent activation of JAK/STAT-3, major components of the recently discovered SAFE pathway. Adenosine, however does not require TNF α to mimic IPostC, but activates JAK/STAT-3 to confer protection. In brief, six alternating episodes of reperfusion and reperfusion with bradykinin or DADLE in the TNF $^{-/-}$ mice failed to reduce the infarct size, whereas the protective effect was still evident when adenosine was used as a postconditioning mimetic. Postconditioning with bradykinin or DADLE protected against reperfusion injury in the TNFR1 deficient, but not in TNFR2 deficient mice. In addition, STAT-3 $^{-/-}$ mice failed to be protected against ischaemia-reperfusion by bradykinin, DADLE or adenosine. Surprisingly, TNF $^{-/-}$ hearts were still protected with Adeno-PostC, therefore suggesting adenosine-induced cardioprotection may occur through partial but not total activation of the SAFE pathway.

4.1 Cardioprotective role of adenosine

Endogenous adenosine is released during ischaemia and reperfusion with blood and interstitial concentrations being elevated after ischaemic insults. In emergency situations adenosine plays an important role in the cardiovascular system. For several decades, adenosine has been widely used for treatment of supraventricular arrhythmias. The potential of adenosine therapy for myocardial infarction has been shown in clinical trials and it has also exhibited cardioprotective effects in preclinical models of myocardial stunning and infarction (Claeys et al., 2004; Kloner et al., 2006; Marzilli et al., 2000; Quintana et al., 2003; Ross et al., 2005). However, adenosine has a short half life in the plasma and has the unwanted side effect of lowering the blood pressure. Great progress is being made in the development of agonists and antagonists which can mimic the beneficial effect of the drug without the deleterious side effects. One such agonist is GP531, a potent second generation adenosine-regulating agent, which has a better half life than adenosine with none of the harmful side effects, such as the lowering of blood pressure (Hale and Kloner, 2010).

In contrast with our data, Penna et al failed to protect the isolated rat heart when adenosine (10 μ M or 30 μ M) was given as a postconditioning stimulus (5 cycles of 10seconds) at the onset of reperfusion (Penna et al, 2009). These data suggest that adenosine may protect as a postconditioning mimetic in a time and concentration dependent manner.

Delineating the mechanisms involved in adenosine-induced cardioprotection would certainly contribute to the development of agents which would encompass the beneficial effects of adenosine without presenting its deleterious side-effects. In the present study, we found that TNF^{-/-} mice but not cardiomyocyte specific STAT-3^{-/-} mice could still be protected with an adenosine postconditioning stimulus. Our data suggest that STAT-3 was not activated as a downstream target of TNF α . This is in accordance with previous work conducted in our laboratory whereby adenosine, given as a preconditioning stimulus, was able to protect against ischaemia-reperfusion in TNF^{-/-} mice but failed to protect STAT-3^{-/-} mice (Smith, 2002; Smith 2005). In fact, STAT-3 activation with adenosine may occur via Erk1/2 activation (Frias, 2009) or PKC activation. Hence, in a late preconditioning setting, Bolli's group demonstrated that activation of PKC contributed, at least in part, to the phosphorylation of STAT-3 (Xuan, 2005; Xuan, 2007) (figure 54).

Adenosine is one of the most widely studied endogenous signalling molecules in cardioprotection, but results found in the literature have not always proved to be consistent. Further investigation into the exact signalling mechanisms involved in ischaemia-reperfusion injury is required.

4.2 Bradykinin and opioids mimic IPostC via the SAFE pathway

Bradykinin is released in significant amounts from the ischaemic myocardium to activate protective effects and plays a role in assuaging reperfusion injury.

There is almost complete agreement that B₂ receptor activation is required for bradykinin-induced protection since icatibant, a selective and specific B₂ receptor antagonist, abolishes bradykinin-induced protection in most models (see review, (Baxter and Ebrahim, 2002)). IPostC-induced cardioprotection in isolated rat hearts was abrogated in the presence of selective inhibitors of the bradykinin B₂ receptor during intermittent bradykinin administration, therefore confirming targeting of the B₂ receptors during early reperfusion is responsible for the protective effects of bradykinin (Penna et al., 2007). The signalling mechanisms downstream of bradykinin receptors involved in the acute protective actions of bradykinin are, however, poorly understood. Similarly, an ischaemic insult to the heart results in the synthesis and release of endogenous opioid peptides which have also been documented in postconditioning-induced cardioprotection (Kin et al., 2005b). IPostC-mediated protection in rat hearts has been reported to be abrogated by a selective antagonist of the opioid δ -receptor as well as a selective antagonist of the κ -receptor (Wang et al., 2007) suggesting that the protective effect of IPostC may be mediated by the δ and κ opioid receptors.

In our present study, intermittent bradykinin or opioid (DADLE) postconditioning at reperfusion required TNF α activation via TNFR2 to confer cardioprotection. Our novel data confirm the protective effect of bradykinin and DADLE as ischaemic postconditioning mimetics and we show for the first time that bradykinin and opioids activate TNF α via TNFR2. To our knowledge, opioids have not been given intermittently as postconditioning mimetics, but only towards the end of ischaemia and/or during reperfusion. Both bradykinin and opioids are known to activate the RISK pathway when given at reperfusion (Gross and Gross, 2006; Penna et al., 2007). Our present data show that both agents can activate the SAFE pathway as a RISK-free signalling mechanism, therefore confirming the existence of multiple prosurvival pathways.

4.3 STAT-3 in mitochondria

STAT-3 is a transcription factor but it is unlikely to act as one as the effects seen in our experimental conditions occur too rapidly. A new role for serine STAT-3 by its translocation to the mitochondria was suggested by Wegrzyn et al, (Wegrzyn *et al.*, 2009). This group provided evidence that STAT-3 is required for optimal function of the mitochondrial electron transport chain, which may permit STAT-3 to facilitate responses to cellular homeostasis. Our western blot analysis demonstrates that postconditioning with bradykinin or DADLE increases the ratio of phosphorylated STAT-3 in the mitochondrial fraction to that of the STAT-3 in the cytosolic fraction, therefore suggesting the involvement of mitochondrial STAT-3 in the signalling pathway of these cardioprotective agents. However, although adenosine protection required STAT-3 signalling, it did not stimulate its translocation to the mitochondria, therefore suggesting an additional mechanism of action for STAT-3 which may be linked to its activation in the cytosol (phosphorylation at the serine site) or translocation to the nucleus.

In conclusion, our data provide new evidence that bradykinin and opioids as postconditioning mimetics activate STAT-3 via TNF α and TNFR2 to confer cardioprotection (SAFE pathway). Considering the low levels of STAT-3 found in the mitochondria (Wegrzyn *et al.*, 2009), it is appropriate to suggest that the role of STAT-3 in the signalling cascade may be attributed to a “kinase-like” activity. Overall, the existence of multiple alternate pathways in ischaemic and pharmacological postconditioning at the time of reperfusion may have potential therapeutic application against ischaemia-reperfusion injury. However, further studies will be required to investigate the exact prosurvival role of STAT-3 within the mitochondria

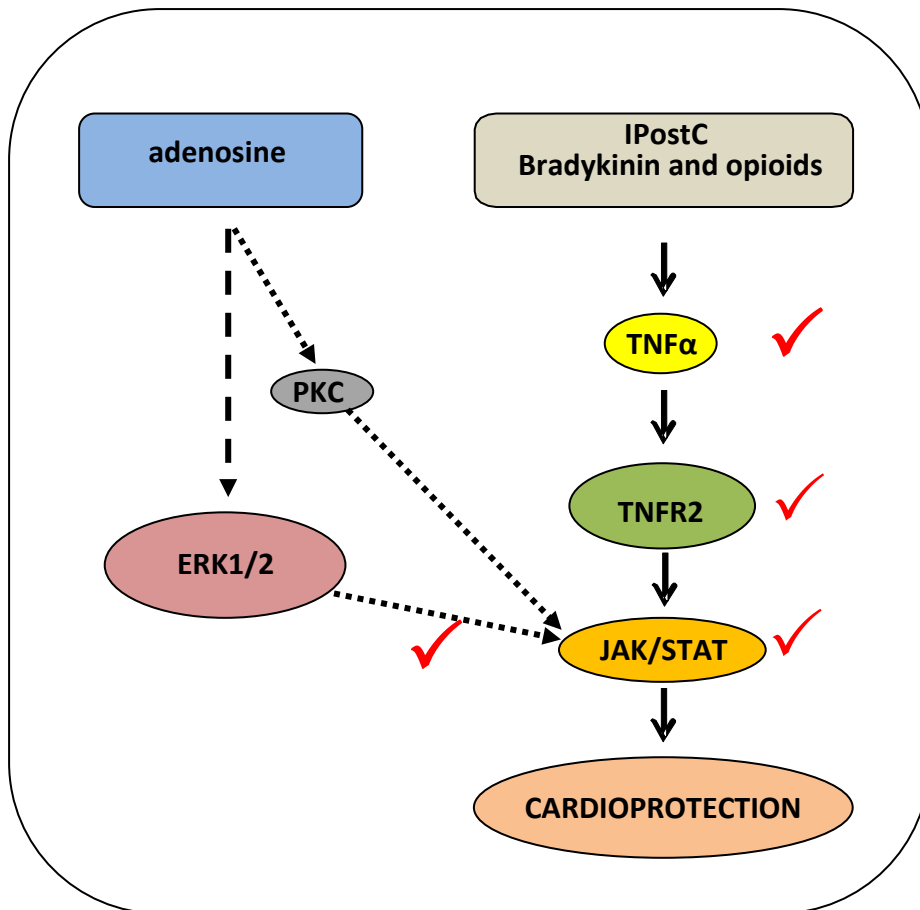


Fig 54: Schematic of our recent data. Activation of TNF α via TNFR2 by bradykinin and DADLE initiates the phosphorylation of JAK/STAT and subsequent cardioprotection. Adenosine makes partial use of the SAFE pathway via STAT-3 signalling but does not require TNF α .

F. ISCHAEMIC POSTCONDITIONING, OBESITY & DIABETES

1.0 Introduction

There is an increasing awareness that both obesity and diabetes are not only growing epidemics but also pose a serious risk factor for cardiovascular disease, in particular myocardial infarction. Prolonged exposure to environmental factors such as high fat or high carbohydrate diets and physical inactivity leads to the development of overweight and resultant obesity and diabetes (MacDonald, 1998).

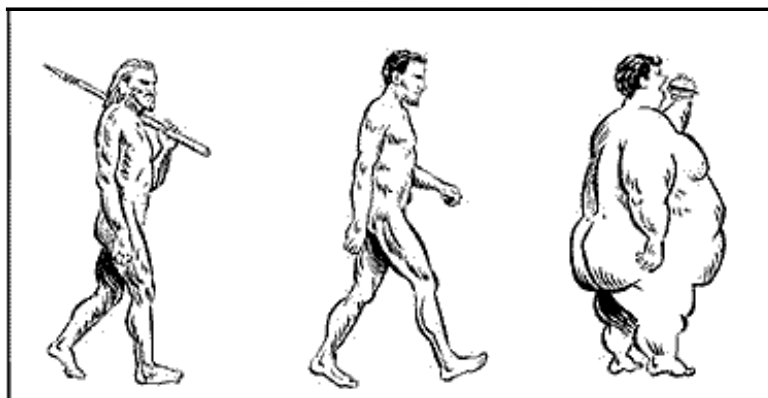


Fig 55: The evolution of man to obesity and diabetes. Fast food and lack of exercise can lead to obesity and diabetes, (adapted from www.evolutionminute.com).

Interestingly, obesity can also be considered as an inflammatory disorder, with TNF α overexpressed in adipose and muscle tissue of obese humans (Hotamisligil et al., 1995; Kern et al., 1995; Krogh-Madsen et al., 2006; Saghizadeh et al., 1996). Nearly all animal models of obesity and insulin resistance seem to produce higher levels of TNF α mRNA and protein compared to their lean counterparts (Hamann et al., 1995; Hofmann et al., 1994; Hotamisligil et al., 1994a; Hotamisligil et al., 1996). A model of obesity in TNF α -deficient mice suggested that the absence of TNF α limits the development of dietary obesity and these mice had significantly improved insulin sensitivity in both diet-induced obesity and in the ob/ob mouse model compared to their wildtype counterparts (Uysal et al., 1997). These data suggested that TNF α facilitates the development of insulin resistance associated with obesity and diabetes and it is further suggested that this effect is mediated via TNFR1 (Uysal et al., 1998).

Animal studies suggest that the ischaemic tolerance is attenuated with obesity, cholesterol and elevated glucose levels (Aasum et al., 2008; Ferdinandy., 2007). In Zucker diabetic fatty and lean Goto-Kakizaki type 2 diabetic rats, ischaemic preconditioning failed to protect the isolated hearts from ischaemia-reperfusion injury (Kristiansen et al., 2004). Similarly, Zucker obese rats subjected to ischaemia-reperfusion failed to be protected with pharmacological preconditioning (Katakam et al., 2007). The outcome of postconditioning in obese and diabetic patients is

uncertain as the recent application of ischaemic postconditioning in clinical studies has excluded patients who were diabetic. Therefore, the aim of our study was to evaluate:

- 1) whether obesity or diabetes would alter the protective outcome of ischaemic postconditioning and
- 2) whether TNF α may alter the cardiovascular risk of obese or diabetic mice subjected to ischaemia-reperfusion and postconditioning, (Figure 56).

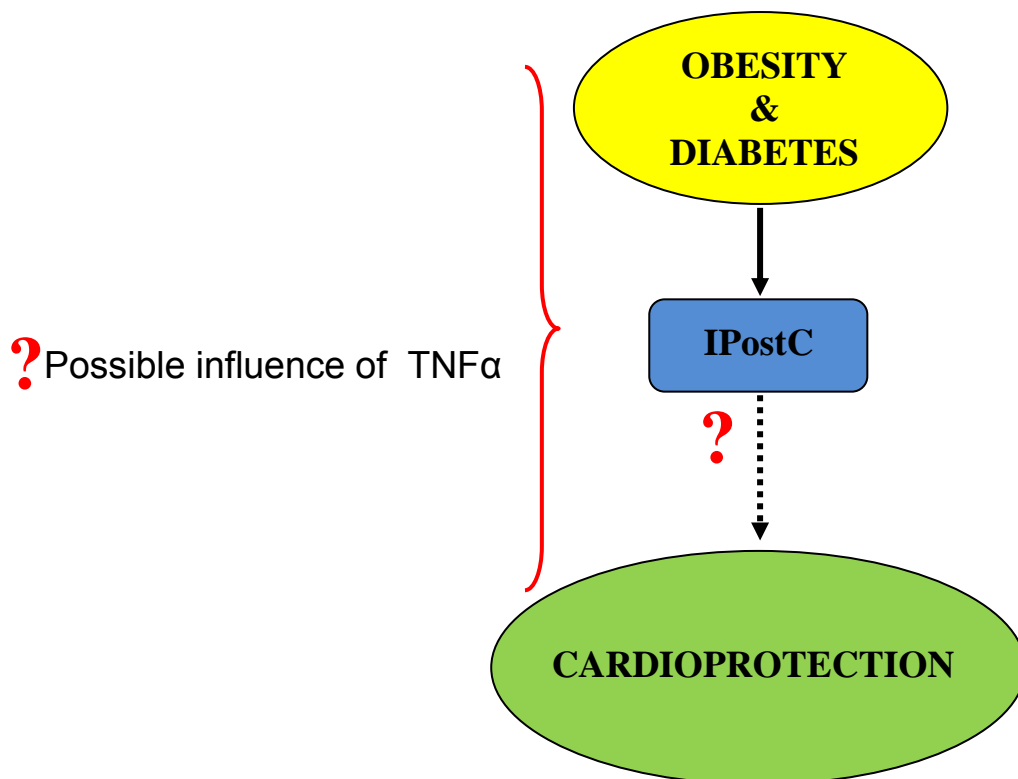


Fig 56: Influence of obesity and diabetes on the outcome of ischaemic postconditioning: does TNF α contribute to this effect?

To investigate our aims we will use TNF-WT and TNF^{-/-} mice.

- 1) A high carbohydrate diet will be fed to the mice for a total of 11 weeks, commencing upon weaning (at 3 weeks of age) to create an obese murine model.
- 2) Diabetes will be induced with a single streptozotocin injection administered intraperitoneally.
- 3) Isolated mouse hearts from the obese and diabetic animals will be subjected to ischaemia-reperfusion and an ischaemic postconditioning protocol as outlined in Section C.

- 4) Physiological parameters, such as glucose tolerance test, blood glucose, plasma insulin and plasma leptin will be analysed.

2.0 Methods

2.1 All animal studies performed were approved by the Animal Research Ethics Committee of the University of Cape Town, and followed the recommendations laid down in the Guide for the Care and Use of Laboratory Animals published by the US National Institute of Health (NIH Publication No. 85 {23}, revised 1996). The total number of animals used in this study was 24.

2.2 High carbohydrate diet (HCD) mouse model

For an obese murine model, male mice were removed from the animal unit at weaning (3 weeks of age) and placed in our facility. TNF-WT and TNF^{-/-} were separated into 2 groups each. One group of mice received a normal diet (ND) of mouse chow and the other group received a diet containing elevated carbohydrates and fats that resembles a Western-type diet (HCD), for 11 weeks, (figure 57). The high carbohydrate diet compared to normal chow, provided energy in the form of carbohydrate, protein and fats, as illustrated in the table below (Aasum et al., 2008; du Toit et al., 2008).

	Carbohydrates (%)	Protein (%)	Fats (%)
Normal diet (ND)	60	30	10
High carbohydrate diet (HCD)	69	17	14

Table 1: Energy provided by high carbohydrate diet (HCD) versus normal chow (ND).

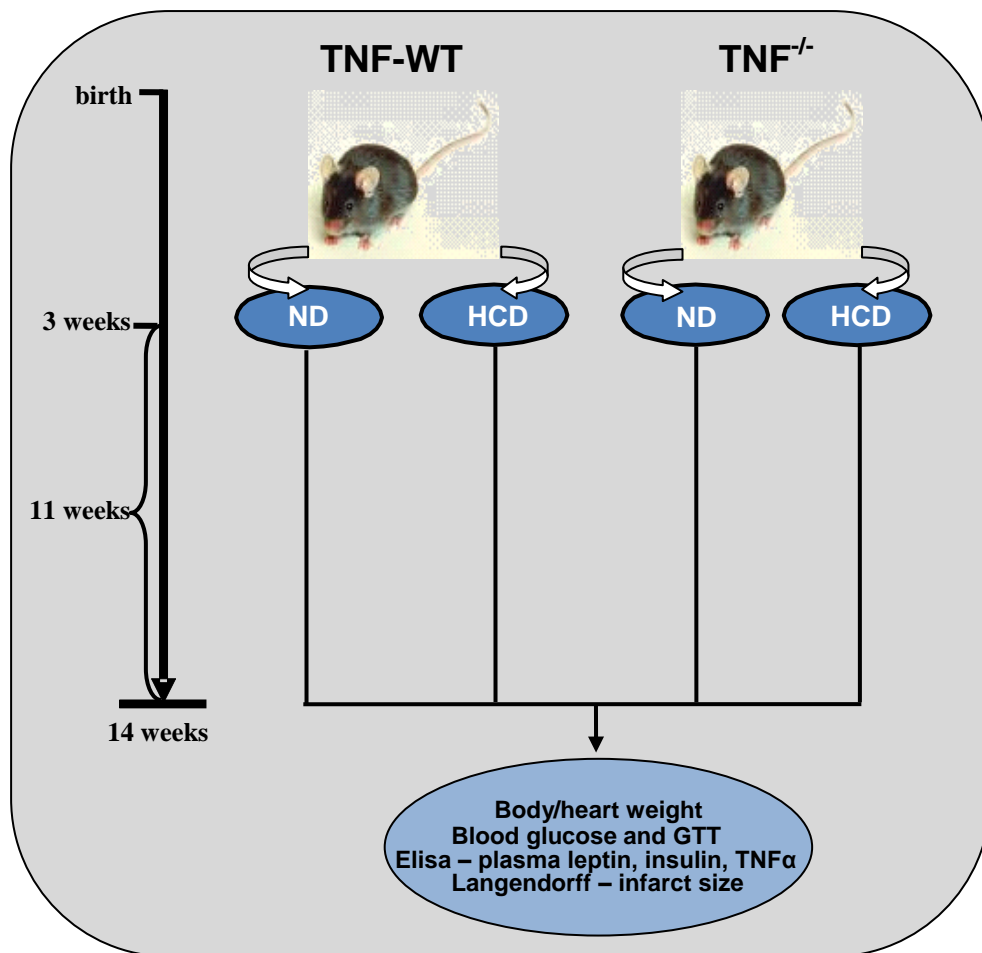


Fig 57: Creation of obese mouse model in TNF-WT and TNF^{-/-} mice.
 ND = normal mouse chow; HCD = high carbohydrate diet.

2.3 Streptozotocin diabetic mouse model (STZ)

Experimental diabetes type 1 was induced in a total of 10 mice per group by a single ip injection of 180 mg streptozotocin (STZ)/kg body weight, dissolved in 0.1 M citrate buffer (Daniel et al., 2007). Non-diabetic control animals were treated with solvent (citrate buffer) alone. Standard mouse chow and tap water were provided ad libitum for all groups. At the end of each time period, only STZ-treated mice with a blood glucose level greater than 16 mmol/l were considered as diabetic (normal blood glucose levels in mice range from 3.4 to 9.7 mmol/l). Of the 10 STZ-treated TNF-WT mice, 7 met the criteria, 2 did not achieve sufficiently high glucose levels, and 1 died shortly after receiving the STZ. No death occurred in the 10 non-diabetic control animals in each group.

2.4 Blood glucose and glucose tolerance test

Blood glucose and the glucose tolerance test (GTT) were done at 14 weeks. Mice were fed normal chow or HCD as described in Table 1. Prior to performing the blood glucose tolerance test, the mice were fasted for 7 hours, but had free access to drinking water. A 20% solution (w/v) of glucose was made up in sterile distilled water. Each mouse was then sedated with a mixture of ketamine (75mg/kg) and medetomidine (0,5mg/kg) given intraperitoneally. Sufficient depth of anaesthesia was determined by pedal reflex. Body weight was recorded for each mouse, a tail cut was done and blood glucose determined in mmol/L by means of a blood glucose monitor (Accu-Chek Active; Roche Diagnostics, Mannheim, Germany) as per the manufacturer's instructions. The Accu-Chek Active allows very small volumes of blood (1-2 µl) to be accurately assessed. A bolus of glucose was given ip (15 mg/kg) and the time of the injection noted. Thereafter, a blood glucose measurement was performed every 30 min post injection, until a decrease in the glucose level was observed (120 to 150 min).

2.5 Perfusion of mouse hearts

The HCD fed mice (14 weeks of age), the STZ-treated mice (5 and 10 days post STZ treatment) and their respective controls were anaesthetized (pentobarbitone, 60 mg/kg ip) and heparinized (25 IU ip) to prevent the blood clotting. Once an adequate level of anaesthesia had been achieved (tested by lack of pedal reflex), body weight was recorded and the hearts were isolated and perfused retrogradely as described in Section C. At the same time, blood was taken from the thoracic cavity of each mouse and placed in a chilled heparinized tube for plasma collection.

2.6 Ischaemic postconditioning

HCD fed mice and STZ-treated mice were subjected to the ischaemic postconditioning (IPostC) protocol which consisted of six alternating cycles of 10 seconds reperfusion, 10 seconds ischaemia, commencing at the onset of reperfusion as described in detail in Section C.

2.7 Ratio of heart weight to body weight

At the end of the perfusion protocol, each heart was carefully dried and weighed after staining with TTC. The heart weight to body weight ratio for each mouse was then calculated.

2.8 Insulin levels

The blood samples were centrifuged at 5000 rpm for 5 min at 4°C and the plasma was removed and frozen until required. Quantitative determination of baseline insulin levels were evaluated using the Ultra Sensitive Mouse Insulin Elisa Kit (Crystal Chem Inc; USA) as per the manufacturer's instructions.

2.9 Leptin levels

For analysis of leptin levels, blood samples were centrifuged at 5000 rpm for 5 min at 4°C and the plasma was removed and frozen until required. Quantitative determination of baseline leptin levels were evaluated using the Ultra Sensitive Mouse Leptin Elisa Kit (Crystal Chem Inc; USA) as per the manufacturer's instructions.

2.10 Chemicals and Pharmacological Agents

Unless otherwise stated, all chemicals were obtained from Sigma-Aldrich Chemicals, Germany.

2.11 Statistical Analysis

Data are presented as mean \pm SEM. Comparisons between multiple groups were performed by 1-way ANOVA followed by Tukey post hoc test or Bonferroni multiple comparison test (GraphPad InStat). Two-way ANOVA followed by Bonferroni multiple comparison test (GraphPad Prism) was performed where species or diet differed. $p < 0.05$ was considered to be statistically significant.

3.0 Results

3.1 Effect of high carbohydrate diet on physiological parameters in TNF-WT mice

HCD increased the body weight significantly in the TNF-WT mice from 29.0 ± 1.0 grams to 32 ± 0.7 grams, ($p < 0.05$ vs ND; figure 58A), but the heart weight to body weight ratio remained unchanged between ND and HCD fed mice, ($p = ns$, figure 58B). Although HCD did not affect baseline blood glucose levels, ($p = ns$, figure 58C), it improved glucose tolerance in the WT animals, ($p < 0.05$ vs ND, figure 58D), and increased plasma insulin and leptin levels to 0.64 ± 0.06 ng/ml from 0.46 ± 0.03 ng/ml in ND, ($p < 0.05$), and to 10.6 ± 0.9 ng/ml from 4.5 ± 0.4 ng/ml in ND, ($p < 0.001$, respectively, figure 58E and figure 58F).

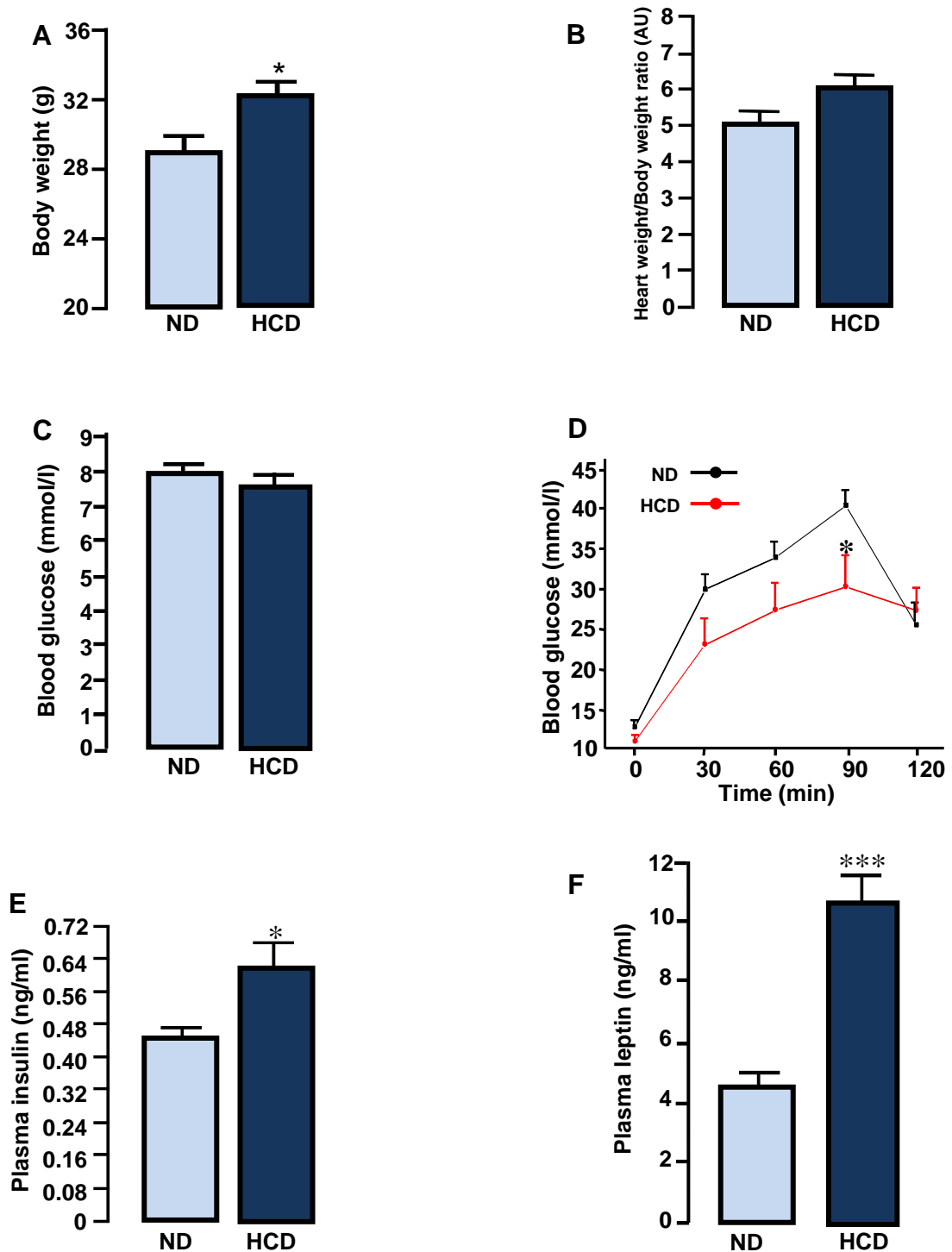


Fig 58: Effects of HCD on physiological parameters in TNF-WT mice. Body weight, plasma insulin and plasma leptin levels were significantly increased by HCD, but no significant difference was observed in heart to body weight ratio or blood glucose levels, $p=ns$; $*p<0.05$; $***p<0.001$ vs ND; $n\geq 6$ for all groups.

3.2 Effect of high carbohydrate diet on IPostC-induced cardioprotection in TNF-WT hearts

To determine whether obesity can affect the outcome of IPostC, the isolated hearts of mice fed with either a normal diet or a HCD were subjected to the IPostC protocol with 6 x 10 second cycles of alternating reperfusion and ischaemia. HCD mice subjected to I/R presented a similar infarct size compared to mice fed with ND ($p=ns$). Furthermore, IPostC conferred protection to a similar extent in HCD mice compared with ND mice versus their respective ischaemia reperfusion control groups ($p<0.001$), (Figure 59).

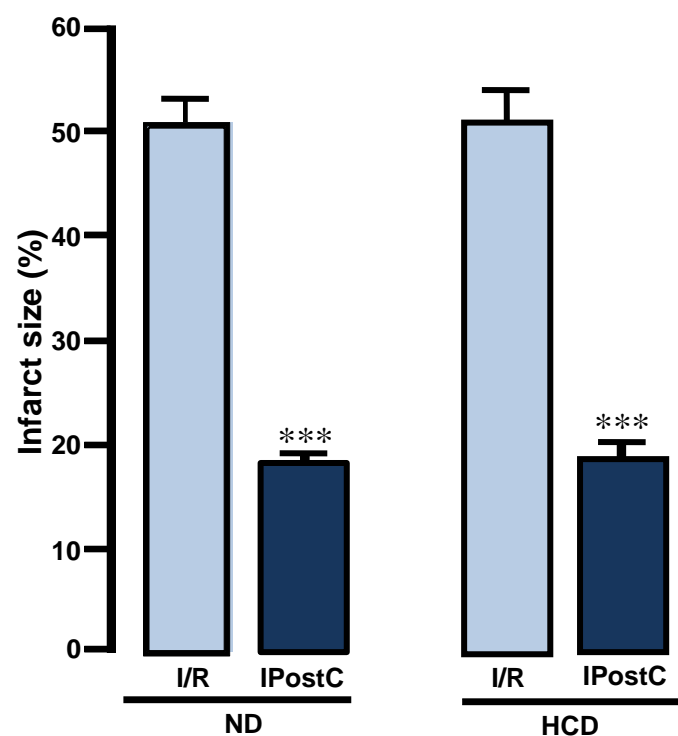


Fig 59: Effect of HCD on IPostC-induced cardioprotection in isolated TNF-WT hearts. IPostC in TNF-WT (ND) reduced infarct size from $50 \pm 2\%$ in the ischaemic control group (I/R) to $18 \pm 0.5\%$ for IPostC. The infarct size in TNF-WT (HCD) was decreased to $18.5 \pm 1.6\%$ versus $50 \pm 2\%$ for the I/R group, $***p<0.001$ vs I/R; $n \geq 6$.

3.3 Effect of high carbohydrate diet on physiological parameters in TNF^{-/-} mice

Mice in which endogenous TNF α is not produced were used to investigate the role of TNF α in obesity. TNF^{-/-} mice were fed either the normal diet or the HCD and physiological parameters were assessed 11 weeks later. Similar to TNF-WT mice, body weight was increased with HCD from 26 ± 0.4 grams to 31.5 ± 0.5 grams, ($p < 0.05$ versus ND, figure 60A). However, TNF^{-/-} mice on the HCD had a significantly lower heart weight to body weight ratio than their ND counterparts, ($p < 0.05$ versus ND, figure 60B). Baseline blood glucose and tolerance of HCD-fed TNF-deficient mice to glucose remained unchanged by HCD, ($p = \text{ns}$; figure 60C and 60D). Plasma insulin levels were increased significantly by HCD from 0.41 ± 0.02 ng/ml to 0.57 ± 0.01 ng/ml, ($p < 0.05$ versus ND, figure 60E). Similarly, the plasma leptin levels were significantly increased by the HCD from 1.3 ± 0.08 ng/ml to 11.2 ± 0.8 ng/ml, ($p < 0.05$ versus ND, figure 60F).

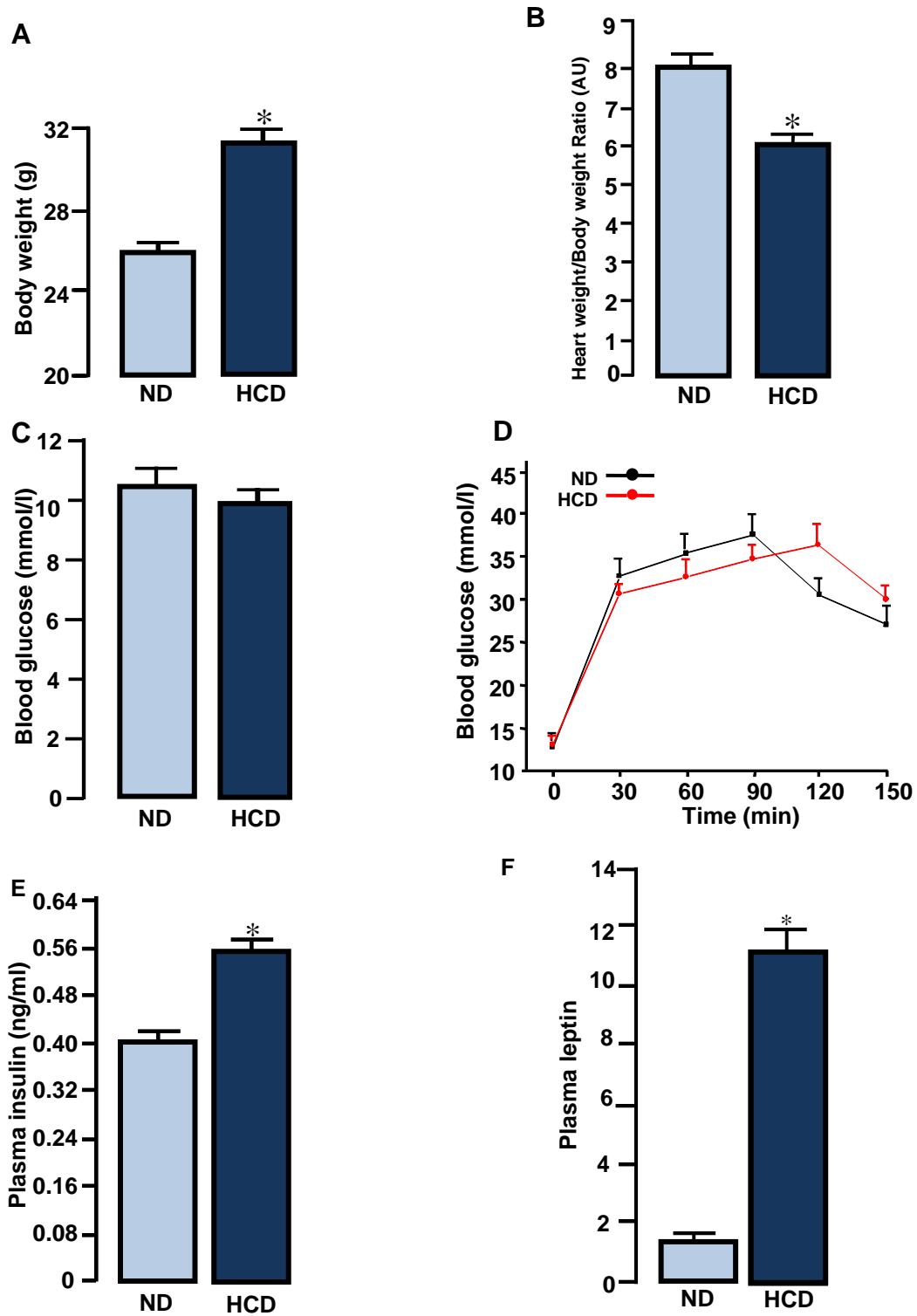


Fig 60: Effects of HCD on physiological parameters of $TNF^{-/-}$ mice. Body weight, plasma insulin and plasma leptin levels were significantly increased by HCD, but blood glucose levels remained unchanged, $p=ns$. However, heart weight to body weight ratio was partially decreased by the HCD, $*p<0.05$ vs ND; $n=6$ for all groups.

3.4 Effect of high carbohydrate diet on IPostC-induced cardioprotection of TNF^{-/-} hearts

To determine whether absence of TNF α in obesity can affect the outcome of IPostC, the isolated hearts of TNF^{-/-} mice fed with either a normal diet or a HCD were subjected to the IPostC protocol as described in Section C. HCD mice subjected to I/R presented a similar infarct size compared to mice fed with ND (p=ns). Surprisingly, the hearts from TNF^{-/-} mice fed with HCD demonstrated a partial reduction in infarct size versus the I/R control, (p<0.05; figure 61).

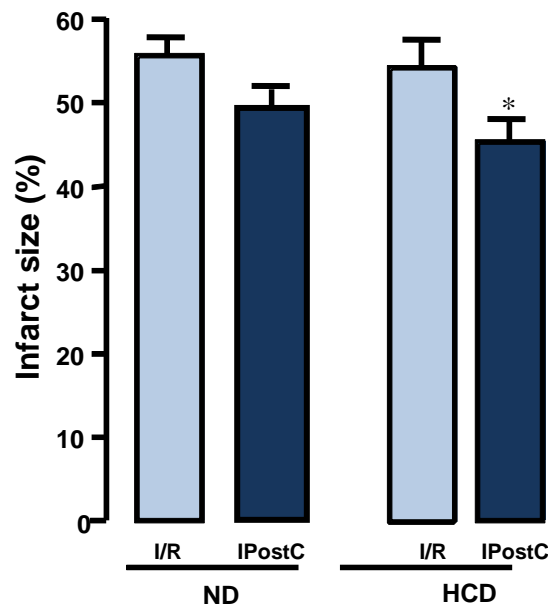


Fig 61: HCD restores cardioprotection in TNF^{-/-} mice. IPostC reduced infarct size in the HCD fed mice from 56 ± 1.8% to 45 ± 4.2% whereas mice fed the normal diet could not be protected, *p<0.05 vs I/R; n≥6.

3.5 Effects of diabetes on physiological parameters in TNF-WT mice

To create a type 1 diabetic model, TNF-WT mice were given a single intraperitoneal injection of streptozotocin (180 mg/kg body weight). Physiological parameters and experiments were performed either 5 days or 10 days post STZ administration. TNF-WT had a significant decrease in body weight 5 days post treatment, from 29.7 ± 1.3 grams to 21.0 ± 2.6 grams ($p < 0.001$ versus no STZ). However, the weight was restored by day 10, ($p = \text{ns}$ versus no STZ; figure 62A). There was no change in the heart weight to body weight ratio ($p = \text{ns}$; figure 62B). Streptozotocin injection increased baseline blood glucose at day 5 and 10, ($p < 0.01$ vs no STZ; figure 62C), but decreased plasma insulin from 1.02 ± 0.2 ng/ml to 0.53 ± 0.14 ng/ml (5 days post STZ) and to 0.36 ± 0.02 ng/ml (10 days post STZ), ($p < 0.05$ vs no STZ-treatment; figure 62D). Similar results for glucose and insulin after streptozotocin treatment have been reported (Nakamura et al., 2006). Leptin levels were also reduced by STZ at day 5 and 10 after injection from 4.5 ± 0.7 ng/ml to 0.6 ± 0.2 ng/ml and 0.23 ± 0.04 ng/ml, respectively, ($p < 0.001$ vs no STZ treatment; figure 62E)

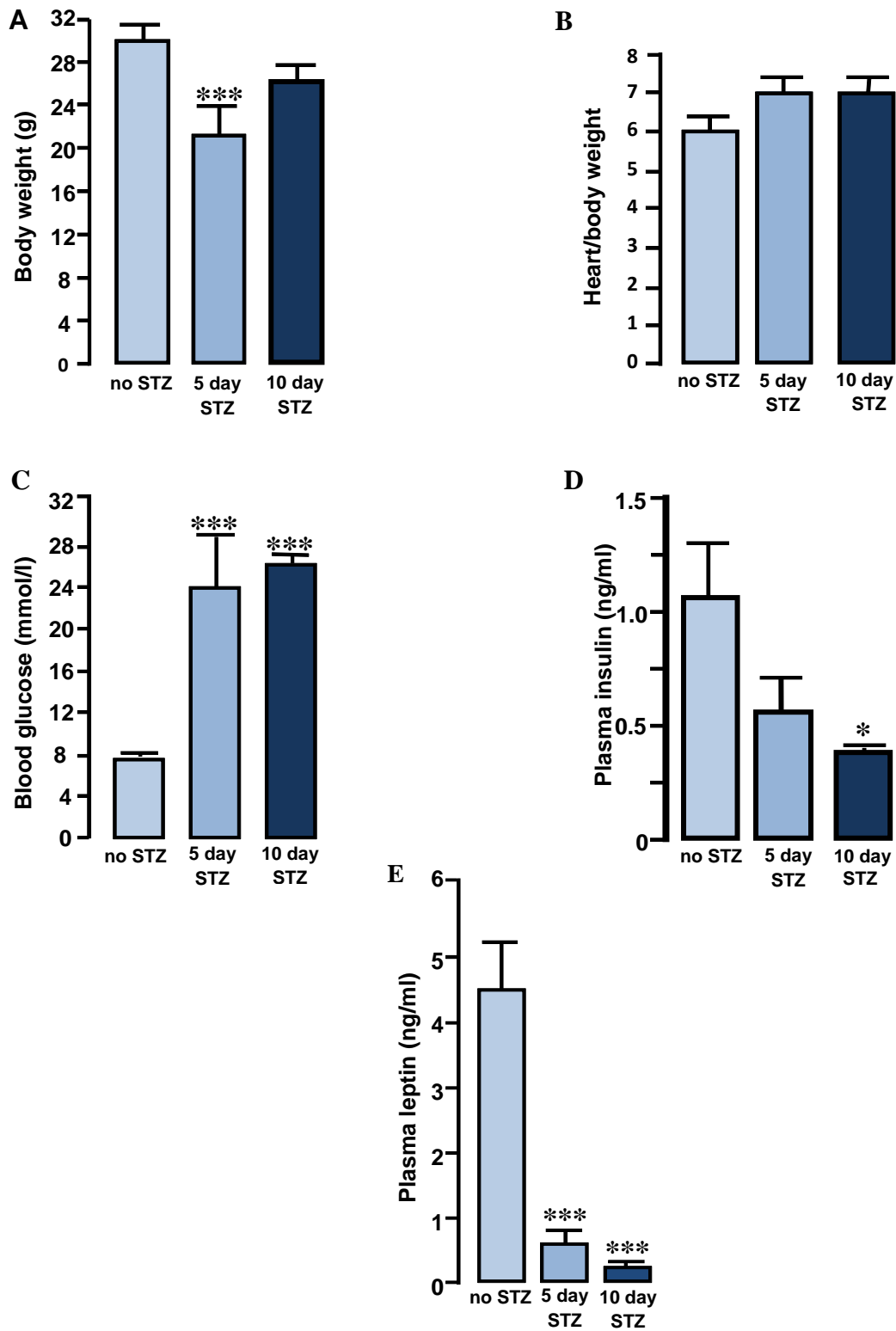


Fig 62: Effects of diabetes on physiological parameters in STZ-treated TNF-WT mice. Body weight was decreased 5 days post STZ treatment, *** $p < 0.001$ vs untreated controls and returned to normal at day 10, $p = ns$. Heart weight to body weight ratio remained unchanged at both time points, $p = ns$. Insulin and leptin levels were both decreased, respectively. STZ increased baseline blood glucose at days 5 and 10, * $p < 0.05$ and *** $p < 0.01$, vs no STZ,

3.6 Effect of diabetes on IPostC-induced cardioprotection in TNF-WT hearts

Isolated hearts from TNF-WT were subjected to the IPostC protocol. STZ-treated TNF-WT mice subjected to I/R showed a similar infarct size to the untreated I/R control, at both time points, ($p < 0.001$ vs I/R). STZ-treatment did not affect the cardioprotective effect of IPostC after 5 or 10 days versus the untreated animals, ($p < 0.001$)

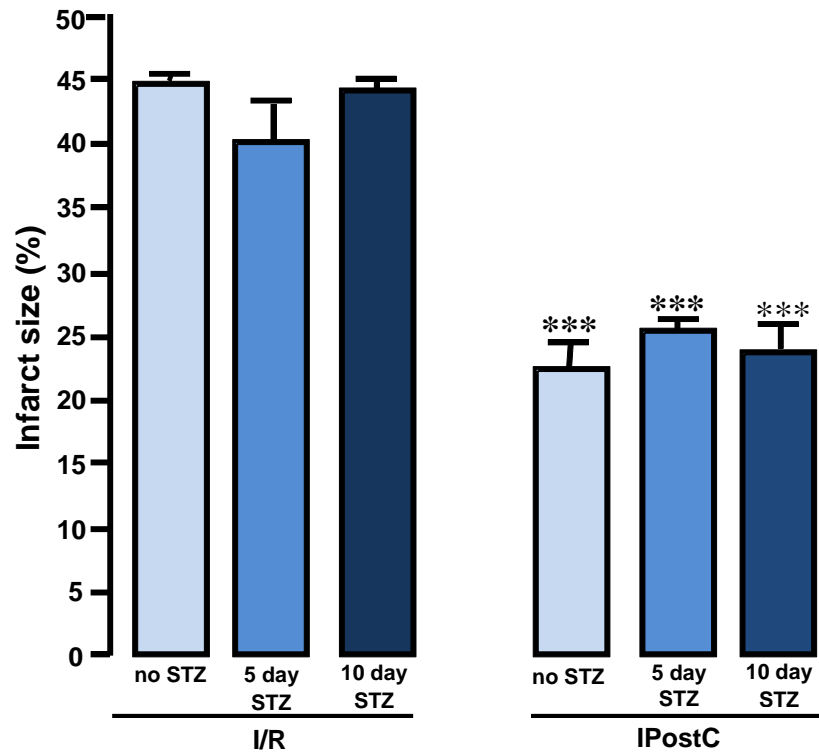


Fig 63: IPostC protects diabetic TNF-WT hearts. Infarct size was significantly decreased by IPostC in untreated animals, $***p < 0.001$ vs I/R. Similarly, IPostC reduced infarct size from $40 \pm 3\%$ to $25.3 \pm 0.5\%$ at 5 days and from $44 \pm 1\%$ to $23.6 \pm 1.8\%$ at 10 days in the STZ-treated mice, $***p < 0.001$ vs respective I/R controls; $n=6$.

3.7 Effect of diabetes on physiological parameters in TNF^{-/-} mice

To investigate whether TNF α plays a role in type 1 diabetes, TNF α -deficient mice were injected intraperitoneally with a single dose of streptozotocin (180 mg/kg body weight). Similar to TNF-WT mice, STZ administration had no effect on body weight, 5 or 10 days after STZ treatment (p=ns, figure 64A). However, the heart weight to body weight ratio was significantly increased by the STZ after 10days, (p<0.01 vs untreated, figure 64B). As expected, STZ increased baseline blood glucose at both time points (p<0.001 vs untreated, figure 64C). Although no significant difference was seen in plasma insulin levels at 5 days post STZ treatment, there was a significant increase 10 days after treatment, from 0.32 ± 0.1 ng/ml to 1.06 ± 0.3 ng/ml, (p<0.01 versus untreated control, figure 64D). The diabetic TNF^{-/-} animals also demonstrated significantly elevated levels of plasma leptin at 5 days post STZ administration, from 2.3 ± 0.1 ng/ml to 3.0 ± 0.4 (p<0.001 versus untreated control), but the levels were drastically reduced in the 10 day post treatment group to 0.3 ± 0.08 ng/ml, (p<0.05 versus untreated control group, figure 64E)

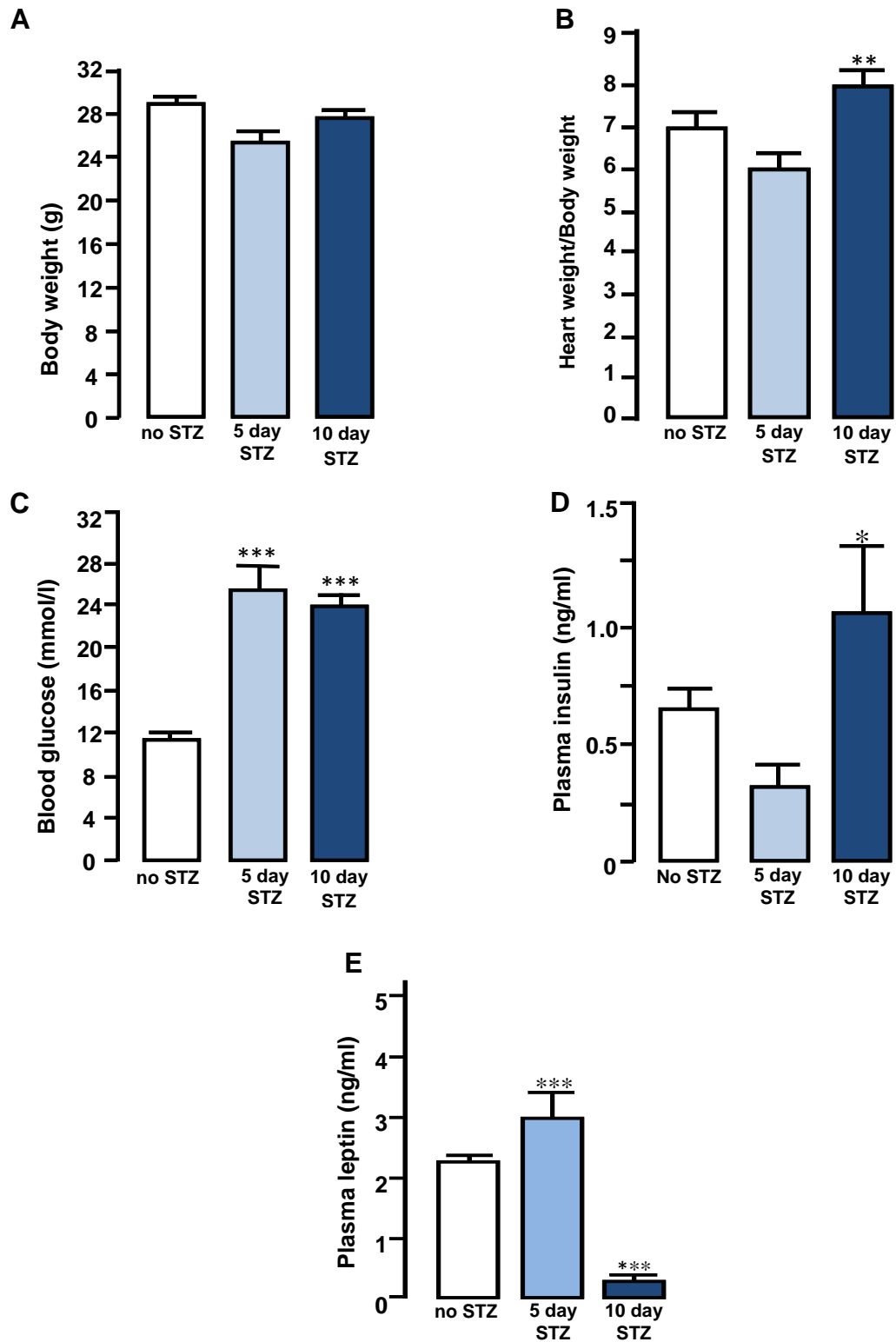


Fig 64: Effects of diabetes on physiological parameters in STZ- treated $TNF^{-/-}$ mice. Body weight remained unchanged by STZ-treatment. STZ significantly increased the heart to body weight ratio and plasma insulin levels after 10 days. Blood glucose was increased at both timepoints. Plasma leptin was significantly increased at day 5 but decreased below baseline at day 10, *p<0.05, **p<0.01, ***p<0.001vs no STZ; n=6.

3.8 Effect of diabetes on IPostC-induced cardioprotection in TNF^{-/-} hearts

Hearts were isolated from the two groups of STZ-treated mice at the appropriate time point. The hearts were subjected to the IPostC protocol of 6 x 10 sec cycles of alternating reperfusion and ischaemia. IPostC could not protect the hearts of STZ-treated animals, (p=ns vs I/R).

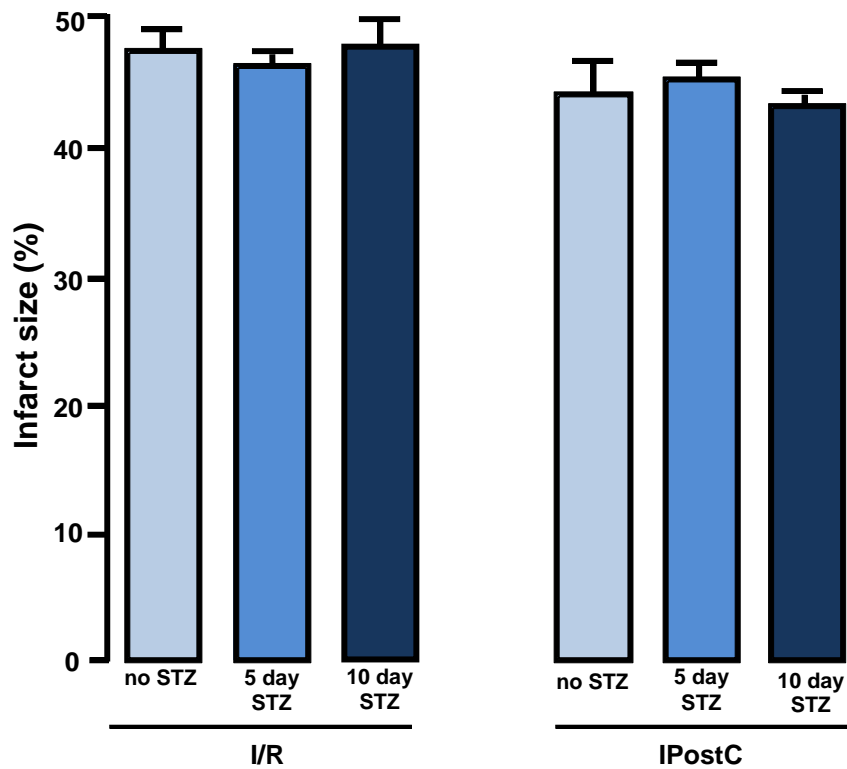


Fig 65: STZ-induced diabetes in TNF^{-/-} mice abrogates the cardioprotective effect of IPostC. Ischaemic postconditioning failed to protect hearts from TNF^{-/-} mice after both 5 and 10 day STZ-treatment, p=ns, n=6.

4.0 Discussion

As most of the patients hospitalized with myocardial infarction are obese or diabetic, it is critical to investigate whether IPostC can be considered as a successful therapy for these patients. Ischaemic heart disease accounts for more than 50% of deaths in diabetic patients (Kannel and McGee, 1979), therefore it is important to explore the cardioprotective signalling mechanisms involved in diabetes associated with ischaemia-reperfusion injury.

Our data revealed that 11 weeks of a high carbohydrate diet or the administration of a single intraperitoneal injection of streptozotocin resulted in a modest model of obesity or diabetes as demonstrated by changes in body weight, blood glucose levels, plasma insulin and plasma leptin levels. IPostC-induced cardioprotection was evident in the TNF-WT with either obesity or streptozotocin-induced diabetes, whereas in the absence of TNF α no protection occurred with the IPostC stimulus in the diabetic mice. Surprisingly, partial restoration of the cardioprotective effect was seen in the TNF $^{-/-}$ mice fed with the HCD.

4.1 Obesity/diabetes and susceptibility to ischaemia-reperfusion

The HCD used in our study is of a similar composition to the conventional Western-type diet of humans and was chosen to represent a moderately obese phenotype (Aasum et al., 2008), unlike the more severe obese models of either ob/ob mice or the db/db mice which are either leptin deficient or have no leptin receptors or the rat models with severe obesity or diabetes. Despite the possible benefits of a low carbohydrate diet, the possibility exists that there are potential adverse effects in those with ischaemic heart disease. A diet low in carbohydrates as commonly used to reduce weight, has been demonstrated to affect myocardial energy substrates, insulin signalling and increased myocardial injury following ischaemia-reperfusion injury in the isolated rat heart (Wang et al., 2008).

Obesity is associated with hyperinsulinaemia which markedly modulates the extent to which myocardial injury occurs during ischaemia-reperfusion (Jonassen, 2000; Jonassen, 2001; Zhang, 2004). Therefore, it is plausible to suggest that in obesity, the impact of high levels of circulating insulin on myocardial injury during ischaemia and reperfusion could over-shadow myocardial susceptibility to ischaemia-reperfusion injury.

Diabetes is associated with an increased mortality and reduced myocardial tolerance to ischaemia (Aronson, 1997; Aasum, 2003) and myocardial ischaemic injury has been shown *in vivo* in animal and human studies to be aggravated by diabetes and insulin resistance (Huang, 2009). However, a wide range of results, ranging from no change, increased or decreased

sensitivity to ischaemia with or without reperfusion injury has been demonstrated in several experimental studies conducted in diabetic animal models (Forrat, 1993; Liu, 1993; Vogel, 1988; Ferdinandy, 2007; Boudina, 2007).

In our obese and diabetic models, the detrimental effect of ischaemia-reperfusion on infarct size was unchanged in the TNF-WT and the TNF^{-/-} mice.

4.2 Obesity/diabetes susceptibility to IPC and IPostC

A study demonstrating the absence of protection in the heart after an IPC stimulus was conducted using rat models which have severe forms of diabetes and obesity (Zucker diabetic fatty rats and the Goto-Kakizaki type 2 diabetic rat) (Kristiansen et al., 2004). Other scientists have reported attenuation of IPC-induced protection in the diabetic hearts of rats and dogs (Tosaki, 1996; Kersten, 2000; Ravingerova, 2000). In contrast, several experimental studies have reported that the streptozotocin-induced diabetic heart is protected by IPC (Liu, 1993; Tatsumi et al., 1998; Thirunavukkarasu, 2007).

The majority of experimental investigations on postconditioning-induced cardioprotection have been done using healthy, young animals and there is a scarcity of studies exploring the role of IPostC in the presence of associated risk factors such as obesity and diabetes. Our data demonstrate that obesity, induced by a high carbohydrate diet or streptozotocin-induced diabetes did not affect the cardioprotective effect of IPostC in the wildtype animals.

Failure of IPostC to limit infarct size was reported from a study conducted in ob/ob mice, which are leptin deficient (Bouhidel et al., 2008). However, a limitation of this study was the lack of leptin in this mouse strain. In contrast to Bouhidel's study, we found that in our model of modest obesity infarct size was decreased and this was accompanied by an increase in plasma insulin and leptin levels, suggesting a cardioprotective role for these two hormones in obesity.

Postconditioning in the diabetic animal model has also not been thoroughly studied but data from experiments conducted in Goto-Kakizaki rats suggested that to protect the diabetic myocardium, the IPC stimulus needed to be increased in order to achieve the necessary threshold for cardioprotection (Hausenloy, 2006b; Tsang, 2005). A very recent study in a murine model of streptozotocin-induced diabetes (using a similar dose to ours) reported a loss of efficacy in IPostC-induced cardioprotection (Przyklenk, 2010). Possible explanations for the contradiction between this study's findings and the findings of our study are: 1) the insulin levels of the mice in Przyklenk's study were significantly lower (0.18 ± 0.08 ng/ml) than the insulin levels found in our diabetic mice (0.36 ± 0.02 ng/ml); 2) the difference in mouse species; and 3) the number of

I/R cycles performed to postcondition the heart. The disparity, therefore, in IPC and IPostC experimental studies conducted in obese and diabetic animals may possibly be due to a number of factors such as: the type of animal model, the choice of experimental protocol, the type of diabetes and the model to induce obesity or diabetes.

4.3 Role of TNF α in obese and diabetic animals

TNF α production is markedly increased in muscle and adipose tissue in obese humans and rodent models of obesity-diabetes, compared with tissues of lean individuals. Several studies have demonstrated that TNF α plays a role in mediating insulin resistance as a result of obesity (Hotamisligil, 1993; Hotamisligil, 1994a; Hotamisligil, 1994b; Kern, 1995; Hotamisligil, 1995; Saghizadeh, 1996). Weight loss or treatment with pioglitazone, an insulin-sensitizing agent, reduces TNF α levels (Hotamisligil, 1993; Hotamisligil, 1995; Kern, 1995a; Kern, 1995b; Hofman, 1994). Three factors which contribute to the control of body weight have been linked to TNF α : 1) the intake of food, 2) expenditure of energy and 3) storage of energy. Administration of TNF α in a rat model resulted in reduced food intake (Rothwell, 1993) and also inhibited gastric emptying, leading to a feeling of satiation, most likely due to activation of leptin (Arbós et al., 1992; Fargeas et al., 1993). Neutralization of TNF α by intravenous administration of a soluble TNF receptor-immunoglobulin G chimeric protein provided a significant improvement in insulin sensitivity in fatty rats (Hotamisligil, 1993; Hotamisligil, 1994a), but treatment of humans with non-insulin-dependent diabetes mellitus with a specific TNF α antibody had no effect on insulin sensitivity (Ofei, 1996). Although TNF α has been proposed as a link between obesity and insulin resistance (Hotamisligil, 2003), in our study the baseline blood glucose was unchanged by HCD in the TNF-WT and TNF^{-/-} mice but plasma insulin levels were increased, therefore suggesting the development of insulin resistance, even in the absence of TNF α .

Surprisingly, in our study the HCD partially restored the protective effect of IPostC in the TNF-deficient mice, therefore suggesting that absence of TNF α in obesity may be of benefit to the heart. In our modestly obese TNF^{-/-} mice, which could be partially protected by IPostC, the plasma leptin level was significantly elevated compared to the animals kept on the normal diet where IPostC-induced protection was abrogated, therefore suggesting that this adipokine may be implicated in a compensatory mechanism. The increased plasma leptin levels observed in our TNF^{-/-} mice correlate with an increase in body weight and the same correlation was found in mice fed a high fat diet (Ahren, 1999).

In contrast to our obese TNF^{-/-} mice, our STZ-induced diabetic TNF-WT model had significantly decreased leptin levels and the protective effect of IPostC was abolished.

It has recently been reported that the tissue preserving actions of leptin are influenced by obesity (Dixon et al., 2009). Dixon's group showed that leptin decreased the infarct size in Wistar and Zucker lean rats, which have functional leptin receptors, but the cardioprotection was lost in the Zucker obese rats in which the leptin receptors are non-functional. This data provided evidence suggesting that the tissue preserving actions of leptin are influenced by the severe obesity seen in Zucker obese rats. Thus, the degree of obesity as well as the presence or absence of TNF α may be of importance in determining the protective effects of leptin.

In experimental conditions, using a TNF α infusion to induce insulin resistance requires concentrations of TNF α which far exceed the physiological range usually observed in insulin resistant rodent models (Altomonte et al., 2003). Rodent data and human data therefore, do not always coincide and the role of TNF α in insulin sensitivity is far from being straightforward.

The presence of TNF α in obesity has been reported to contribute towards the development of cardiac hypertrophy in cardiomyocytes (Yokoyama, 1997). In support of this hypothesis, our data demonstrate a decrease in the heart to body weight ratio in TNF-deficient mice fed with a diet high in carbohydrates.

Interestingly, when TNF receptor deficient mice were fed a high fat diet the data suggested that the two TNF receptors work in concert to protect against diabetes (Schreyer et al., 1998). In this study TNFR1 knock out mice were protected against diet-induced obesity implicating TNF α and TNFR2 in protection against obesity. Our data support this study as in our TNF^{-/-} mice fed with a HCD, the body weight increased by 21% whereas in comparison, the body weight of TNF-WT increased by only 11%, therefore suggesting that absence of TNF α leads to an increase in weight and would thus, influence the development of obesity. It would be of interest to repeat our experiments in our TNF^{-/-}, TNFR1^{-/-}, and TNFR2^{-/-} animals to further delineate the role of TNF α receptors in our model.

In conclusion, our data demonstrate that the cardioprotective effect of IPostC was unaltered in a high carbohydrate diet model of obesity and streptozotocin-induced obesity. Whilst TNF α is necessary for IPostC-induced cardioprotection, maintenance of glucose homeostasis and in the control of appetite to prevent obesity it can also lead to cardiac hypertrophy. In our obesity model, absence of TNF α partially restores cardioprotection, while in our diabetic model, the absence of TNF α abrogated the protective effect of IPostC, illustrating the bidirectional effect of TNF α and the fact that the role of TNF α in obesity and diabetes related ischaemic heart disease remains a complex one. Further studies measuring the levels of TNF α or using TNF antibodies

will be required to better understand the protective/deleterious role of TNF α in these pathologies.

A schematic illustrating our present data on the influence of obesity and diabetes on the outcome of IPostC is shown in figure 66.

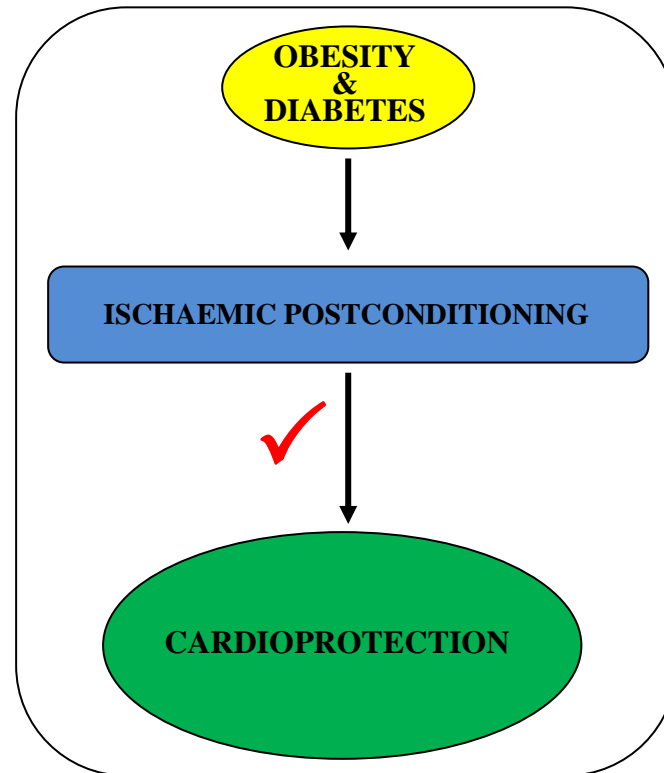


Fig 66: Schematic illustrating the effect of obesity and diabetes on the outcome of IPostC. Diabetic and modestly obese mice can be protected against ischaemia-reperfusion by ischaemic postconditioning

G. FINAL DISCUSSION - CONCLUSION

1.0 Concluding remarks

The discovery of IPostC in the last decade has opened a novel era for therapeutic targets against reperfusion injury. In IPostC, the alternating cycles of reperfusion and ischaemia are applied at the onset of reperfusion, making this a more applicable treatment for patients with myocardial infarction, either in the ambulance or on admission to hospital. However, delivering these alternating cycles requires precise and timed application which is difficult and could substantially differ with the pathophysiology of the individual. Therefore, pharmacological postconditioning, where a drug, agonist or antagonist could be administered at the onset of reperfusion would seem to be a better alternative in the quest to attenuate the injury caused by reperfusion therapy

To further understand the signalling mechanisms of cardiac postconditioning, we aimed to explore the role of TNF α in ischaemic postconditioning as well as pharmacological postconditioning. As obesity and diabetes are now considered to be burden diseases, we also wished to investigate the influence of obesity and diabetes on the protective effect of ischaemic postconditioning and explore the role of TNF α in obesity/diabetes related ischaemic heart disease.

The main and novel findings presented in this thesis are as follows:

1. IPostC confers cardioprotection in the murine model of I/R via the SAFE pathway which involves the activation of TNF α , TNFR2 and the subsequent triggering of the JAK/STAT-3 protective signalling cascade.
2. TNF α mimics IPostC independently of the RISK pathway.
3. Bradykinin and opioids mimic IPostC via the SAFE pathway and therefore require the activation of TNF α and STAT-3 for protection. Bradykinin and opioids possibly protect via translocation of STAT-3 to the mitochondria. In contrast, adenosine protects via STAT-3 but independently of TNF α .
4. Our data confirm the existence of multiple parallel prosurvival signalling pathways in IPostC, (Figure 67).
5. Cardioprotection can still be conferred by ischaemic postconditioning in modestly obese mice or in streptozotocin-induced diabetic mice.
6. Obese TNF knockout mice could be partially protected with IPostC, therefore suggesting a compensatory protective mechanism in TNF^{-/-} mice.

Our data suggest that obese and type 1 diabetic individuals may still benefit from IPostC, relative to the severity of the disease. The existence of multiple alternate pathways in IPostC and pharmacological postconditioning at the time of reperfusion may provide novel therapeutic applications against ischaemia-reperfusion injury in the clinical setting.

In summary, our data provide evidence for the alternative protective prosurvival SAFE pathway in both ischaemic and pharmacological postconditioning, involving activation of TNF α , TNFR2 and STAT-3.

We have discovered that intrinsic TNF initiates a powerful prosurvival pathway in the heart involving the activation of its receptor 2 and the transcription factor, STAT-3

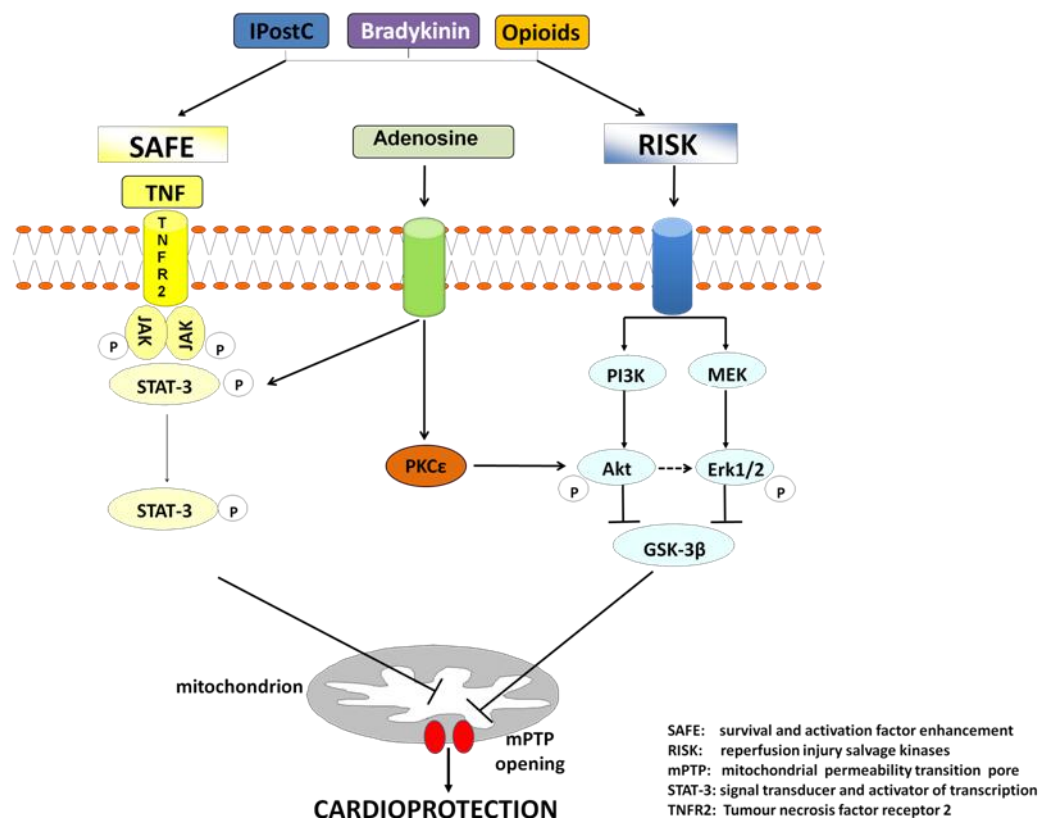


Fig 67: Proposed signalling scheme of the novel SAFE pathway and the known RISK pathway in ischaemic and pharmacological postconditioning-induced cardioprotection.

2.0 Study Limitations

The use of an isolated heart system in this thesis does not take into account the intracoronary activation of leucocytes which may result in a detrimental effect of TNF α in the setting of myocardial ischaemia. Our data will need to be validated *in vivo*. Kinetic analysis of blood and cardiac TNF α levels following ischaemic postconditioning would facilitate the understanding of the exact role of TNF α as a cardioprotective agent against reperfusion injury. However, previous data exploring the protective effect of TNF α have demonstrated a good correlation between *in vitro* and *in vivo* preconditioning experiments, therefore suggesting that the circulating leucocytes do not play a major role (Deuchar et al., 2007; Lecour et al., 2005a). Alternatively, the cellular origin of the cardioprotective TNF α may reside in cardiomyocytes, and other cells such as resident cardiac mast cells and macrophages (Dorge et al., 2002; Frangogiannis et al., 1998)

Another limitation to the study lies in the use of non-specific inhibitors, such as wortmannin and AG490. Wortmannin inhibits the PI-3 kinase signaling cascade and not Akt directly. AG490 blocks JAK2, which is upstream of STAT-3. Therefore, the use of such inhibitors can only indicate that a particular signaling cascade is blocked, but cannot pinpoint the exact downstream protein kinase involved. Further studies, either with siRNA techniques or genetically modified mice, such as Akt null mice and GSK-3 β knockout mice, may give greater specificity to the findings of the present study.

The high carbohydrate diet was given to our mice from 3 weeks of age (at weaning) for a period of 11 weeks only as we wished to keep the age of the mice consistent with the 12-14 weeks we had used in our previous experiments. Therefore, the increase in weight in our mice was modest and all the manifestations of obesity may not have yet developed. Future work should include a group in which the HCD is given for a total of 16 weeks to explore the role of severe obesity on the cardioprotective effect of IPostC, bearing in mind that an increase in age generally affects the outcome of postconditioning adversely.

Our diabetic model was created with streptozotocin which induces type 1 diabetes. Future experiments should be done with a type 2 diabetic mouse model to confirm the data observed in our present model.

3.0 Future directions

The cellular origin of TNF α is mainly in the macrophages and resident mast cells (Frangogiannis et al., 1998), but cardiomyocytes also produce TNF α under prolonged ischaemic conditions (Dorge et al., 2002). Delineation of the cellular source of TNF α production in the context of myocardial infarction may be of clinical value. For this purpose we are in the process of generating a cardiomyocyte-specific TNF $^{-/-}$ mouse, using the Cre-Lox P technology. The Cre-Lox P system of genetic targeting requires two separate mouse lines, a homozygous TNF floxed mouse and an organ specific heterozygous Cre recombinase-expressing mouse. These two mice are crossed to produce the cellular specific TNF $^{-/-}$ mouse. Using the same Cre-Lox P technology, we are also in the process of breeding macrophage-neutrophil TNF α -deficient mice. Together with the cardiomyocyte specific TNF $^{-/-}$ this should provide us with a better insight into cellular origin of the cardioprotective TNF α .

The mechanisms triggering the activation of TNF α in the SAFE pathway remain uncertain. Toll-like receptors (TLR's) play a role in the innate immune system to activate TNF α . Recent data exploring the impact of the toll-like receptors 2 or 4 in ischaemia-reperfusion, have not been conclusive as to whether TLR2 or TLR4 are protective or deleterious (Abarbanell et al., 2010; Arslan et al., 2010; Dong et al., 2010). Using our TNF-WT and TNF $^{-/-}$ mice we aim to investigate the possibility that TLR's contribute to the activation of TNF α in the initiation of the prosurvival SAFE pathway.

H. PUBLICATIONS ARISING FROM THIS WORK

Ischaemic postconditioning protects against reperfusion injury via the SAFE pathway. **Lydia Lacerda**, Sarin Somers, Lionel H Opie and Sandrine Lecour. *Cardiovasc Res* (2009) 84, 201-208.

I. PREVIOUS PUBLICATIONS

1: Ischemic and Pharmacological Preconditioning in Girardi Cells and C2C12 Myotubes Induce Mitochondrial Uncoupling. Jan Minners, **Lydia Lacerda**, Joy McCarthy, James J. Meiring, Derek M. Yellon and Michael N. Sack *Circ Res*: 2001(89): 787-792

2: TNF α -induced cytoprotection requires production of free radicals within mitochondria in C₂C₁₂ myotubes. **Lydia Lacerda**, Robert M. Smith, Lionel Opie, Sandrine Lecour. *Life Sciences*; 2006 : 2194-2201

3: Genetic depletion of cardiac myocyte STAT-3 abolishes classical preconditioning. Robert M. Smith, Naushaad Suleman, **Lydia Lacerda**, Lionel H. Opie, Shizuo Akira, Kenneth R. Chien, Michael N. Sack. *Cardiovasc Res*: 2004 (63); 611-616

4: Pharmacological Preconditioning With Tumor Necrosis Factor α Activates Signal Transducer and Activator of Transcription-3 at Reperfusion Without Involving Classic Prosurvival Kinases (Akt and Extracellular Signal-Regulated Kinase). Sandrine Lecour, PharmD, PhD; Naushaad Suleman, MS; Graeme A. Deuchar, PhD; Sarin Somers, BSc; **Lydia Lacerda**, BSc; Barbara Huisamen, PhD; Lionel H. Opie, MD, PhD. *Circulation*: 2005; 3911-3918

6: Diazoxide-induced respiratory inhibition-a putative mitochondrial K_{ATP} channel independent mechanisms of pharmacological preconditioning. Jan Minners, **Lydia Lacerda**, Derek M. Yellon, Lionel H. Opie, Christopher J. McLeod, and Michael N. Sack. *Mol Cell Biochem*: 2007; 11-18

7: Signal Transducer and Activator of transcription factor 3 is involved in the cardioprotective signalling pathway activated by insulin therapy at reperfusion. Britt N. Fuglestad, Naushaad Suleman, Crina Tiron, Tambuzai Kanhema, **Lydia Lacerda**, Thomas V. Andreasen, Michael N. Sack, Anne K. Jonassen, Ole D. Mjøs, Lionel H. Opie, Sandrine Lecour. *Basic Res Cardiol*: 2008; 444-453

8. TNF α protects cardiac mitochondria independently of its cell surface receptors. **Lydia Lacerda**, Joy McCarthy, Shazia F.K. Mungly, Edward G Lynn, Michael N. Sack, Lionel Opie, Sandrine Lecour. *Basic Res Cardiol* (article in press) DOI 10.1007/s00395-010-0113-4

J. PEER REVIEWED, PUBLISHED CONFERENCE PROCEEDINGS:

- 2007: “Ischaemic postconditioning requires TNF activation to confer protection”. **Lydia Lacerda**, Lionel Opie and Sandrine Lecour. International Society for Heart Research; Italy; *published in J Mol Cell Cardiol, 2007; Vol 42, S184.*
- 2007: “Critical role of age, strain and number of cycles in ischemic postconditioning” International Society for Cardiovascular Research; Italy; Sarin Somers, **Lydia Lacerda**, Naushaad Suleman, Lionel Opie, Sandrine Lecour. International Society for Heart Research; Italy; *published in J Mol Cell Cardiol, 2007; Vol 42, S186.*
- 2008: “TNF alpha can mimic ischaemic postconditioning”. **Lydia Lacerda**, Lionel Opie, Sandrine Lecour. International Society for Heart Research; Greece; *published in J Mol Cell Cardiol; 2008; Vol 44(4); 718 (abstract No.17).*
- 2008: “TNF α signalling – a critical role in ischemic postconditioning?” **Lydia Lacerda**. 9th Annual conference of the SA Heart association, South Africa; *published in SA Heart, Spring, 2008; Vol 5, No.4; 217.*
- 2009: “Pharmacological postconditioning with TNF alpha protects via the SAFE pathway rather than the RISK”. **L. Lacerda**, S. Somers, L.H. Opie, S. Lecour. European Society of Cardiology- Heart Failure Conference-ISHR section; France; *published in European J Heart Failure, 2010; Vol 12(1).*
- 2009: “Innate immunity and protection against reperfusion injury via the SAFE pathway”. 10th Annual Congress of SA Heart Association, South Africa; **Lydia Lacerda**, Sarin Somers, Lionel Opie and Sandrine Lecour; *published in SA Heart, Spring, 2009; Vol 6, No.4; p 272.*
- 2010: Bradykinin, insulin and opioids mimic ischaemic postconditioning via the SAFE pathway. **Lydia Lacerda**, Sarin Somers, Lionel Opie and Sandrine Lecour. Frontiers in Cardiovascular Biology, Berlin, Germany, 2010 – *published in Cardiovasc Res, Vol 87; Supplement 1, 15th July 2010.*
- 2010: Does obesity and/or diabetes influence the outcome of ischaemic postconditioning? **Lydia Lacerda**, Lionel Opie and Sandrine Lecour. 11th Annual Congress of SA Heart Association, South Africa; *published in SA Heart, Winter 2010; Vol 7, No.3; p 208.*
- 2010: HDL protects against lethal reperfusion injury via the SAFE pathway. Miguel Frias, Sarin Somers, **Lydia Lacerda**, Richard James and Sandrine Lecour. 11th Annual Congress of SA Heart Association, South Africa; *published in SA Heart, Winter 2010; Vol 7, No.3; p 202.*

K. CONFERENCE OUTPUTS

Ischaemic postconditioning requires TNF activation to confer protection. **Lydia Lacerda**, Lionel Opie and Sandrine Lecour. Physiology Society of Southern Africa, Muldersdrift, Gauteng, South Africa, 2007.

TNF α can mimic ischaemic postconditioning. **L. Lacerda**, L.H. Opie and S. Lecour. International Society for Heart Research, Athens, Greece 2008

Activation of TNF α signalling at onset of reperfusion protects the heart. **L. Lacerda**, L.H. Opie and S. Lecour. South African Medical Research Council Research Day, Parow, South Africa 2008

TNF α signalling – a critical role in ischaemic postconditioning? **L. Lacerda**, L.H. Opie and S. Lecour. South African Heart Conference, Wild Coast Sun, South Africa 2008

Pharmacological postconditioning with TNF α protects via the SAFE pathway rather than the RISK pathway. **Lydia Lacerda**, Sarin Somers, Lionel Opie and Sandrine Lecour. International Society for Heart Research – ESC meeting, Nice, France 2009

Pharmacological postconditioning with TNF alpha protects via the SAFE pathway rather than the RISK pathway. **Lydia Lacerda**, Sarin Somers, Lionel Opie and Sandrine Lecour. Physiological Society of Southern Africa, Stellenbosch, South Africa, 2009

Innate immunity and protection against reperfusion injury via the SAFE pathway. **Lydia Lacerda**, Sarin Somers, Lionel Opie and Sandrine Lecour. South African Heart Conference, Sun City, Johannesburg, South Africa. 2009.

Adenosine, bradykinin, insulin and the opioids can mimic ischaemic postconditioning: Role of the cytokine, tumour necrosis factor alpha (TNF α). **Lydia Lacerda**, Lionel Opie and Sandrine Lecour. Astrazeneca Conference, Tygerberg, South Africa, 2010

Bradykinin, insulin and opioids mimic ischaemic postconditioning via the SAFE pathway. **Lydia Lacerda**, Lionel Opie and Sandrine Lecour. Frontiers in Cardiovascular Biology, Berlin, Germany, 2010

Does obesity and/or diabetes influence the outcome of ischaemic postconditioning? **Lydia Lacerda**, Lionel Opie and Sandrine Lecour. 11th Annual Congress of SA Heart Association, South Africa, 2010

HDL protects against lethal reperfusion injury via the SAFE pathway. Miguel Frias, Sarin Somers, **Lydia Lacerda**, Richard James and Sandrine Lecour. 11th Annual Congress of SA Heart Association, South Africa, 2010

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