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Akt mediated mitochondrial protection in the heart: metabolic and survival pathways to the rescue

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Abstract Cardiomyocyte death is now recognized as a critical factor in the development of heart disease. Mitochondria are not only responsible for energy production to ensure that cardiac output meets the body's energy demands, but they serve as critical integrators of cell survival signals. Numerous stressors are known to induce cell death by necrosis and/or apoptosis mediated through mitochondrial dysregulation. Anti- and pro-apoptotic Bcl-2 family proteins regulate apoptosis by controlling mitochondrial outer membrane permeability, whereas opening of the mitochondrial permeability transition pore (PT-pore) induces large amplitude permeability of the inner membrane and consequent rupture of the outer membrane. Akt is one of the best described survival kinases activated by receptor ligands and its activation preserves mitochondrial integrity and protects cardiomyocytes against necrotic and apoptotic death. The mechanisms responsible for Akt-mediated mitochondrial protection have not been fully elucidated. There is, however, accumulating evidence that multiple Akt target molecules, recruited through both transcriptional and post-transcriptional mechanisms, directly impinge upon and protect mitochondria. In this review we discuss mechanisms by which Akt activation can effect changes at the mitochondria that protect cardiomyocytes and attenuate pathophysiological responses of the heart.

Keywords Akt · Mitochondria · Heart · Hexokinase-II

Introduction

As a result of symbiotic invasion of eukaryotes by eubacteria more than 1.5 billion years ago, mitochondria have evolved to become the critical organelles for generation of cellular energy (Dyall et al. 2004) as well as gatekeepers in the control of cell survival (Kroemer et al. 2007). Heart tissue is amongst the most rich in mitochondria as there are profound energy demands on oxidative phosphorylation to support the beat-to-beat contraction/relaxation cycle. Adult cardiomyocytes do not proliferate and accordingly their loss through cell death plays a crucial role in heart diseases involving ischemic injury and the development of heart failure (Adams et al. 2000; Dorn and Brown 1999; Gustafsson and Gottlieb 2008; Hayakawa et al. 2003; Murphy and Steenbergen 2008; Nakagawa et al. 2005; Nakayama et al. 2006; Scarabelli et al. 2006; Weiss et al. 2003; Yussman et al. 2002). It has become apparent in recent years that Bcl-2 family proteins and the permeability transition pore (PT-pore) are important regulators of the mitochondrial death pathway that is activated by stress in cardiomyocytes (Gustafsson and Gottlieb 2008; Halestrap and Pasdois 2009; Murphy and Steenbergen 2008). The proapoptotic Bcl-2 family proteins induce mitochondrial outer membrane permeabilization (MOMP) (Reed 2008). Opening of the PT-pore at the inner membrane causes mitochondrial depolarization, the loss of small molecular weight substances from the matrix, and rupture of the outer mitochondrial membranes due to osmotic mitochondrial swelling (Baines et al. 2005; Crompton 1999; Crow et al. 2004; Green 2005; Green and Kroemer 2004; Gustafsson and Gottlieb 2003; Nakagawa et al. 2005; Weiss et al. 2003). The consequences of activation of these two pathways are reported to be different, and their interrelationship remains somewhat controversial. Bcl-2 family proteins are

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now believed to more specifically regulate apoptosis (Baines et al. 2005; Green 2005; Nakagawa et al. 2005). MOMP induced by apoptotic Bcl-2 family proteins leads to release of apoptotic molecules such as cytochrome c, leading to caspase-9/caspase-3 activation and DNA fragmentation. In contrast, the PT-pore is primarily associated with necrotic death (Baines et al. 2005; Nakagawa et al. 2005). Both apoptotic and necrotic cell death have been observed in the heart subjected to ischemic injury, in congestive heart failure and in cardiomyopathy, although the contribution of these two types of death is controversial and may differ depending on the types and pathological stages of stress. Regardless, preservation of mitochondrial integrity is essential not only to maintenance of metabolic homeostasis but also to survival of cardiomyocytes.

Protective signaling molecules are recruited in response to stress to prevent cell death. Akt is a serine/threonine kinase with well documented targets controlling diverse cellular functions, including protein synthesis, energy metabolism and cellular survival (Ceci et al. 2007; Fujio et al. 2000; Matsui et al. 1999; Matsui et al. 2001). Preservation of mitochondrial integrity has been shown to be involved in Akt-mediated protection (Lai et al. 2003; Miyamoto et al. 2008; Plas et al. 2001) and recent studies reveal that Akt has direct inhibitory effects not only on apoptotic Bcl-2 family proteins but also on the PT-pore (Ahmad et al. 2006; Davidson et al. 2006; Juhaszova et al. 2004; Miyamoto et al. 2008). Interestingly it is becoming increasingly clear that survival/death signaling and metabolic signaling share regulatory molecules as evidenced for example, by an unexpected physiological role of BAD, an Akt target molecule, on glucose metabolism in β -cells (Danial et al. 2008). Other molecules recently reported as regulators of Akt mediated mitochondrial protection include glycogen synthase kinase 3 (GSK-3) and hexokinase-II. The mechanisms by which Akt confers mitochondrial protection in the heart will be discussed below.

Cardiac injury, mitochondria and the PT-pore

An increase in intracellular Ca^{2+} is responsible for normal cardiomyocyte contraction. In turn, mitochondrial Ca^{2+} increases in a beat-to-beat fashion in response to cytosolic Ca^{2+} transients and stimulates ATP production to meet energy demands via activation of Ca^{2+} -sensitive matrix dehydrogenases (Maack and O'Rourke 2007; Szalai et al. 2000). It has been established that cardiomyocyte injury induced by ischemia/reperfusion (I/R) results from increased cytosolic Ca^{2+} and generation of reactive oxygen species (ROS). As a result of loss of ionic homeostasis

during ischemia, supra-physiological levels of cytoplasmic Ca^{2+} lead to mitochondrial Ca^{2+} overload during reperfusion. The combination of mitochondrial Ca^{2+} overloading, oxidative stress, and the loss of adenine nucleotides potentiates opening of the mitochondrial PT-pore (Crow et al. 2004; Gustafsson and Gottlieb 2003; Murphy and Steenbergen 2008; Weiss et al. 2003; Zorov et al. 2006). Recent genetic studies demonstrate that opening of the PT-pore is also important in development of heart failure associated with dysregulated Ca^{2+} handling and β -adrenergic receptor signaling (Nakayama et al. 2007).

The molecular entities that constitute the PT-pore remain elusive and controversial. Historically, the PT-pore was believed to be composed of the adenine nucleotide translocase (ANT) in the inner membrane, the voltage-dependent anion channel (VDAC) of the outer membrane, and cyclophilin D (Cyp-D) in the matrix. This classic model has, however, been brought into question by recent genetic studies. Specifically, neither gene deletion and/or knockdown of VDAC or of ANT prevents PT-pore opening in response to mitochondrial Ca^{2+} overload (Baines et al. 2007; Kokoszka et al. 2004). A significant contribution of Cyp-D to PT-pore opening was first suggested by the inhibitory effect of cyclosporin A (Cs-A) on Ca^{2+} -induced PT-pore opening detected as mitochondrial swelling (Crompton et al. 1988) and the subsequent discovery that Cs-A binds to Cyp-D (Tanveer et al. 1996). However, as Cs-A is also a potent calcineurin inhibitor, the mechanism of the profound cardioprotection from I/R injury provided by Cs-A and the role of Cyp-D in pore formation remained ambiguous until recent studies of animals in which Cyp-D expression was genetically manipulated. It was then determined that mitochondria isolated from Cyp-D knockout mice are resistant to Ca^{2+} overload-induced PT-pore opening, and that Cyp-D null mouse hearts exposed to I/R show significantly less injury than WT mice (Baines et al. 2005; Nakagawa et al. 2005). Conversely, mitochondria isolated from transgenic mouse hearts that express elevated Cyp-D show spontaneous pore opening (Baines et al. 2005).

While these genetic studies further defined the role of Cyp-D in PT-pore regulation, there remains controversy about the role of the PT-pore in apoptotic versus necrotic death. The knockout of Cyp-D has been reported to either have no effect on (Baines et al. 2005; Nakagawa et al. 2005), or to inhibit (Chiara et al. 2008) various forms of apoptotic cell death in vitro. The intriguing possibility that Cyp-D has a role in regulating apoptotic proteins in a manner that is independent of the PT-pore has recently been suggested (Eliseev et al. 2009), and this possibility has the potential to resolve some of the current conflicts in the field.

Cardiac injury and Bcl-2 family proteins

Induction of the proapoptotic Bcl-2 family proteins is a powerful mechanism for stimulating the mitochondrial apoptotic pathway. Bax and Bak, both proapoptotic members of the family, contain three Bcl-2 homology domains (BH-domain; BH1, BH2 and BH3), but lack a BH4 domain. The BH4 domain is critical for the protective effect of the anti-apoptotic Bcl-2 proteins (Chen et al. 2002) such as Bcl-2, Bcl-X_L and Mcl-1. It has been suggested that Bax and Bak permeabilize the outer mitochondrial membrane by forming pores in response to apoptotic stimuli. MOMP induced by Bax/Bak is further activated in the presence of another class of proapoptotic Bcl-2 family proteins, the BH3-only proteins (BAD, BIM, BID, Bnip3, Nix/Bnip3L and PUMA) which facilitate Bax- or Bak-mediated pore formation and/or antagonizing anti-apoptotic Bcl-2 family proteins. The significance of Bcl-2 family proteins in cardiomyocyte survival and ischemic injury has also been established (Chen et al. 2002; Hochhauser et al. 2003; Kubli et al. 2007; Toth et al. 2006; Webster et al. 2005; Yussman et al. 2002). For instance, Bax deletion or Bcl-2 overexpression have been shown to protect the heart against ischemia/reperfusion (Brocheriou et al. 2000; Chen et al. 2001; Hochhauser et al. 2003). Expression of BNip3, a BH3-only protein, is induced by hypoxia and leads to apoptosis (Kubli et al. 2007; Webster et al. 2005) and Nix (BNip3L) is increased in hypertrophy and triggers apoptosis in the heart (Yussman et al. 2002). Ischemia/reperfusion-induced Ca²⁺ increases and subsequent activation of calpain can also lead to cleavage of Bid which induces cell death (Chen et al. 2002).

Akt and cardioprotection

Akt was originally described as an oncogenic protein and extensive studies have revealed that the activity of Akt is upregulated in many types of tumors, contributing to diminished apoptotic cell death and facilitating tumor growth. Akt regulates cellular metabolism, coordinated with its pro-survival and proliferative/growth effects. Akt is activated downstream of phosphatidylinositol 3-kinase (PI3K) in response to stimulation of receptor tyrosine kinases (Fujio et al. 2000; Matsui et al. 1999), glycoprotein 130 (Craig et al. 2001; Kuwahara et al. 2000; Sheng et al. 1997) and G-protein coupled receptors (GPCRs) (Bell and Yellon 2003; Chesley et al. 2000; Cohen and Downey 2008; Downey et al. 2007; Means et al. 2007; Naga Prasad et al. 2000; Robert et al. 2001; Zhu et al. 2001) in the heart. Akt exerts a strong cardioprotective effect in response to activation of these receptors. Early studies using adenoviral

gene transfer directly demonstrated that Akt protects heart against ischemia/reperfusion (Fujio et al. 2000; Matsui et al. 1999). Agonist induced Akt activation has also been demonstrated to be cardioprotective. For example, Akt is responsible for cardioprotection mediated by leukemia inhibitory factor (LIF) (Miyamoto et al. 2008; Negoro et al. 2001) or cardiotrophin-1 (CT-1) (Craig et al. 2001; Kuwahara et al. 2000), and Akt activation by insulin or S1P decreases I/R damage within hours in the isolated perfused heart or *in vivo* (Jin et al. 2002; Jonassen et al. 2001; Means et al. 2007). It has been also suggested that Akt is involved in cardioprotection mediated by preconditioning, a series of brief cycles of I/R prior to a sustained period of ischemia (Mocanu et al. 2002; Murphy and Steenbergen 2008; Tong et al. 2000; Uchiyama et al. 2004).

Diverse cellular Akt targets localized in various cellular compartments confer protection through transcriptional- (Brunet et al. 1999; Cook et al. 2002; Craig et al. 2001; Muraski et al. 2007) and post-transcriptional regulation of pro-survival/pro-apoptotic proteins (Datta et al. 1997; del Peso et al. 1997; Kuwahara et al. 2000; Matsui and Rosenzweig 2005; Pap and Cooper 1998; Shiojima and Walsh 2006). Stimulation of receptors that activate PI3K results in an increase in PIP₃; this drives Akt translocation to the plasma membrane where it becomes activated through phosphorylation at T308 by the upstream kinase, phosphoinositide-dependent kinase-1 (PDK1) and regulation at S473 by TORC2 (Sarbasov et al. 2005; Williams et al. 2000). In the cytosol, mTOR, GSK-3 β and FOXO are phosphorylated by Akt, regulating translation, transcription and protein degradation (Matsui and Rosenzweig 2005; Morisco et al. 2000; Shioi et al. 2002; Skurk et al. 2005). It is now evident, however, that Akt activated at the plasma membrane is not only active in the cytosol but also translocates to subcellular compartments including the nucleus and mitochondria (Bijur and Jope 2003; Camper-Kirby et al. 2001; Kunkel et al. 2005; Miyamoto et al. 2008; Sasaki et al. 2003). Translocation of activated Akt to these subcellular components can mediate diverse effects. Redistribution of activated Akt to the nucleus and to mitochondria has been demonstrated not only by fractionation and immunostaining but also by using FRET-based Akt activity reporters in non-cardiac cells (Kunkel et al. 2005; Sasaki et al. 2003). Translocation of activated Akt to mitochondria was initially reported in neuronal cells (Bijur and Jope 2003), and we demonstrated similar translocation in cardiomyocytes stimulated by LIF, a glycoprotein 130 receptor agonist (Miyamoto et al. 2008). Preservation of mitochondrial integrity is known to be involved in Akt-mediated protection (Lai et al. 2003; Miyamoto et al. 2008; Plas et al. 2001) and several target molecules through which Akt could afford mitochondrial protection have been identified.

Akt and mitochondrial protection through Bcl-2 family proteins

Phosphorylation of Bcl-2 family proteins by Akt

Apoptosis is tightly controlled by Bcl-2 family proteins through regulation of mitochondrial outer membrane permeability. BAD, a BH3-only apoptotic protein, was the first Bcl-2 family protein discovered to be phosphorylated and regulated by Akt (Datta et al. 1997; del Peso et al. 1997). Phosphorylation of BAD by Akt at Ser 136 promotes dissociation of BAD from Bcl-X_L, a protective Bcl-2 family protein. Phosphorylation of BAD at Ser136 by agonists known to activate Akt has been demonstrated in the heart and correlated with cardioprotection (Aikawa et al. 2000; Jonassen et al. 2001; Kuwahara et al. 2000; Mehrhof et al. 2001; Negoro et al. 2001; Uchiyama et al. 2004). Another Bcl-2 family protein regulated by Akt is the proapoptotic BH1-3 protein, Bax which is a key molecule in mitochondrial outer membrane permeabilization. Phosphorylation of Bax by Akt at Ser184 leads to inhibition of the conformational change required for Bax translocation to mitochondria (Arokium et al. 2007; Gardai et al. 2004; Yamaguchi and Wang 2001). Phosphorylation of Bax at Ser184 has not been reported in the heart, but deletion of the Bax gene has been shown to decrease cardiac I/R injury (Hochhauser et al. 2003). An inhibitory effect of Akt on Bcl-2 family proteins has also been suggested to be involved in the protective effect of heat shock proteins (Hsp). Hsp27 was shown in an early study to associate with activated Akt (Konishi et al. 1997). This observation was extended by recent studies demonstrating that Hsp27, like Akt, inhibits Bax activation and apoptosis and that this occurs in a PI3K-dependent manner (Havasi et al. 2008). Hsp20 association with Akt was also recently shown to prolong Akt activity, leading to sustained phosphorylation of BAD in the heart (Fan et al. 2008).

Akt and expression of Bcl-2 family proteins

In addition to modulation of activity of pro-apoptotic proteins through phosphorylation, the expression level of Bcl-2 family proteins is regulated by Akt. For example, LIF treatment has been shown to work through a PI3K/Akt pathway to prevent the doxorubicin-induced decrease in Bcl-X_L expression in cardiomyocytes (Negoro et al. 2001). Preconditioning (which activates Akt) has also been shown to prevent decreases in Bcl-2 expression level induced by I/R (Uchiyama et al. 2004). Bcl-2 is also transcriptionally upregulated by IGF-1 through CREB in an ERK- and Akt-dependent manner (Mehrhof et al. 2001). The mTOR complex-1 (mTORC1), which is downstream of Akt, controls expression of the anti-apoptotic Bcl-2 family

protein Mcl-1 at the level of translation and promotes cancer cell survival (Mills et al. 2008). FOXO has three Akt consensus sequences and phosphorylation of FOXO by Akt facilitates its nuclear export and thereby inhibits its transcriptional activity. In non-cardiac cells, FOXO transcription factors are known to enhance expression of Bim, a BH-3 only protein (Dijkers et al. 2000; Gilley et al. 2003), although the significance of Bim and its regulation by Akt/FOXO in cardiac pathophysiology has not been established. Taken together, there is abundant evidence that Akt exerts transcriptional, translational, and post-translational control of Bcl-2 family proteins which ensure that mitochondrial integrity is preserved against short-term and long-term stress.

GSK-3 and Akt mediated mitochondrial protection

Glycogen synthase kinase 3 (GSK-3) was initially described in the context of glycogen synthesis, but it is now recognized that this kinase contributes to regulation of many other cellular functions. GSK-3 functions in the heart as both a pro-apoptotic signal and as a suppressor of hypertrophic responses. There are two isoforms of GSK-3 (GSK-3 α and GSK-3 β); GSK-3 β has been most extensively studied in the context of cardiac injury.

GSK-3 β and Akt mediated phosphorylation

GSK-3 β is an active kinase under basal conditions and Akt phosphorylation at Ser9 inhibits the kinase activity of GSK-3 β . An earlier study by Murphy and co-workers demonstrated that preconditioning induces phosphorylation and inhibition of GSK-3 β in a PI3K-dependent manner and that inhibition of GSK-3 β by LiCl or pharmacological inhibitors mimics cardiac preconditioning against I/R stress (Tong et al. 2002). Growing evidence suggests a proapoptotic effect of GSK-3 β in the heart and demonstrates that inhibition of GSK-3 β is involved in cardioprotection mediated by ischemic pre- and post-conditioning, as well as certain pharmacological interventions. Expression of a kinase-inactive form of GSK-3 β , acting as a dominant negative, protects rat ventricular myocytes from isoproterenol-induced apoptosis (Menon et al. 2007) and dominant negative GSK-3 β transgenic mice show well-compensated hypertrophy and reduced apoptosis and fibrosis during transverse aortic constriction (Hirotani et al. 2007). These findings demonstrate that inhibition of GSK-3 β has beneficial effects in the heart and suggest the possibility that inhibition of GSK-3 β by Akt-mediated phosphorylation at Ser9 could lead to cardioprotection. Indeed, a seminal study carried out using adult ventricular myocytes revealed that insulin inhibits ROS-induced PT-pore opening in a PI3K-

dependent manner and this protection could be reversed by overexpression of GSK-3 β -S9A, a mutant that cannot be phosphorylated at Ser9 (Juhaszova et al. 2004). Solloff and co-workers also reported in this study that GSK-3 β localized not only in cytosol but also at mitochondria. Phosphorylated GSK-3 β has also been reported to be increased at mitochondria by agonists such as insulin (Juhaszova et al. 2004) and erythropoietin (Ohori et al. 2008), suggesting that Akt phosphorylates and inhibits GSK-3 β actions at mitochondria. This hypothesis is supported by a recent study in which transgenic mice expressing GSK-3 β -S9A were utilized, to demonstrate that phosphorylation of Ser9 is required for Akt-mediated inhibition of PT-pore opening by post-conditioning in the heart (Gomez et al. 2008).

Recent evidence provides insights into molecular mechanisms by which inhibition of GSK-3 β leads to mitochondrial protection. Using perfused heart and immunoprecipitation/Western blotting, Nishimura et al. showed that GSK-3 β , phosphorylated in response to preconditioning or erythropoietin treatment, interacts with ANT, leading to a decrease in association of ANT with Cyp-D and resultant protection (Nishihara et al. 2007). The precise mechanism by which GSK-3 β inhibition exerts protective effects at mitochondria is controversial, however, as it has been suggested that post-conditioning protects heart mitochondria through inhibition of GSK-3 β in a Cyp-D-independent manner and that GSK-3 β actions occur upstream of the PT-pore (Gomez et al. 2008). A paper from Steenbergen's group proposed yet another model in which inhibition of GSK-3 β during ischemia adapts mitochondria to ensuing reperfusion (Das et al. 2008). Their work indicated that GSK inhibitors prevent mitochondrial ATP consumption during ischemia and thereby protect mitochondria against subsequent reperfusion. This protective effect of GSK inhibitors was suggested to occur through decreased VDAC2 phosphorylation and inhibition of ATP transport into mitochondria. The authors proposed that the decrease in ATP hydrolysis by mitochondria during ischemia leads to lower mitochondrial membrane potential, which in turn lessens Ca²⁺ sequestration into mitochondria and consequent ROS generation during reperfusion (Das et al. 2008). Thus this metabolic adjustment induced by inhibition of GSK-3 β may lead to prevention of mitochondrial Ca²⁺ overload and generation of ROS, two major inducers of the PT-pore.

Very recent studies by two different laboratories using GSK-3 α (S21A) and GSK-3 β (S9A) knock-in mice suggest that the story is more complex. Matsuda et al. (2008) demonstrated that apoptosis induced by transverse aortic constriction (TAC) was inhibited by knock-in of GSK-3 β (S9A). They further tested the role of phosphorylation of GSK-3 α using a non-phosphorylatable mutant of GSK-3 α (S21A) which enhanced apoptosis during TAC. These

results suggest that phosphorylation of GSK-3 β at Ser9 is pro-apoptotic whereas that of GSK-3 α at Ser21 is protective (Matsuda et al. 2008). This is in sharp contrast to the observations obtained in GSK-3 β (S9A) transgenic mice in which post-conditioning failed to protect the heart, leading to the conclusion that phosphorylation of GSK-3 β at Ser9 is cardioprotective (Gomez et al. 2008). Another recent study demonstrated that pre-conditioning and post-conditioning protected the heart against I/R even in GSK-3 α (S21A) and GSK-3 β (S9A) double knockin mice, suggesting that neither GSK-3 α nor GSK-3 β phosphorylation at the known Akt sites are required for pre- and post-conditioning mediated cardioprotection (Nishino et al. 2008). Interestingly this study showed that insulin still preserves the ability to prevent the PT-pore in double knock-in mice, suggesting another target of Akt for PT-pore inhibition.

In summary it is currently unclear whether phosphorylation and inhibition of GSK-3 β by Akt confers cardioprotection through inhibition of the PT-pore. The conflicting findings in the literature might reflect the use of different experimental models (e.g. acute inhibition by pharmacological inhibitors vs. chronic inhibition by overexpression of mutants of GSK-3; transgenic vs. knock-in mice), interventions to induce stress (TAC, ex vivo I/R and in vivo I/R), and different methods for detection of the PT-pore opening (e.g. mitochondrial membrane potential, mitochondrial swelling, calcium retention capacity).

GSK-3 β and Bcl-2 family proteins

GSK-3 β has been reported to directly regulate Bcl-2 family proteins. Mcl-1, an anti-apoptotic Bcl-2 family protein, was originally reported by Green and co-workers to be phosphorylated by GSK-3 and degraded, an effect that was prevented by Akt expression (Maurer et al. 2006). Upregulation of Mcl-1 by GSK inhibition has been recently reported to be protective against TNF- α in cardiomyocytes (Hirovani et al. 2007). In addition to the potential for control of Mcl-1 protein stability through Akt/GSK-3 β , Mcl-1 is transcriptionally up-regulated in response to Akt activation through mTOR signaling and promotes tumor cell survival (Mills et al. 2008). GSK-3 β also phosphorylates Bax and induces an activating conformational change, a response that has been shown to be inhibited by Akt in neuronal cells (Linseman et al. 2004) and in yeast (Arokium et al. 2007). Similarly, in H9c2 cells, H₂O₂-induced BAX translocation to mitochondria was attenuated when GSK-3 β was knocked down by siRNA or when cells were stimulated with erythropoietin, an Akt agonist (Ohori et al. 2008). It has also been reported that GSK-3 inhibitor treatment increases the association of Bcl-2 with mitochondria in the heart (Das et al. 2008). Taken together, there are a multitude of possible mechanisms through which GSK-3 β could

mediate the protective effects of Akt on cell survival in the heart.

Hexokinase-II and Akt mediated mitochondrial protection

Hexokinase (HK) catalyzes the first step of glycolysis, phosphorylating glucose to glucose-6-phosphate, and is therefore of fundamental importance in cellular metabolism and physiology. There are four isoforms of HK (I, II, III and IV). HK-I and HK-II have mitochondrial binding motifs at their N-terminus and are reported to bind to the outer mitochondrial membrane through VDAC (Anflous-Pharayra et al. 2007; Azoulay-Zohar et al. 2004; Nakashima et al. 1986; Pastorino et al. 2005). HK-I is ubiquitously expressed and HK-II, while also widely expressed, is the predominant isoform in insulin sensitive tissues such as adipose, skeletal and cardiac muscle and in most cancer cells (Pastorino and Hoek 2003; Wilson 2003). Pedersen's laboratory has suggested that mitochondrial HKs play a role in facilitating ATP production through the preferential access of this enzyme to ATP generated by mitochondria, resulting in more efficient glycolysis (Mathupala et al. 2006; Pedersen 2007). Elevated glycolysis even in the presence of oxygen is one of the hallmark changes in metabolism associated with cancer cells, and was classically described as the Warburg Effect (Pastorino and Hoek 2003; Pedersen 2007).

There is increasing evidence that energy metabolism and cell survival utilize common pathways in many organ systems, and key amongst these is the recognition that Akt regulates metabolism as well as cell survival (Matsui and Rosenzweig 2005; Pastorino and Hoek 2003). Akt-mediated protection of cardiomyocytes against I/R is accompanied by enhanced sarcolemmal Glut-4 expression and glucose uptake (Matsui et al. 2001). Glyceraldehyde-3-phosphate dehydrogenase (GAPDH), a glycolytic enzyme, mediates not only changes in glycolysis but also protects cells from caspase-independent cell death (CICD) by enhancing autophagy (Colell et al. 2007). Of further interest, the apoptotic and metabolic machinery share certain signaling molecules (Galluzzi et al. 2008). For instance, a physiological/metabolic role of the proapoptotic Bcl-2 family protein BAD has been revealed in pancreatic beta cells in which phosphorylation of BAD at Ser155 inhibits its apoptotic effect but concomitantly increases glucose-fueled mitochondrial respiration (Danial et al. 2008). This increase in mitochondrial metabolism enhances glucose-stimulated insulin secretion from the beta cells by a mechanism that involves binding and stimulation of glucokinase (the HK IV isoform mainly expressed in pancreas and liver) by phosphorylated BAD (Danial et al.

2008). Whether a similar mechanism for maintenance of optimal glucose metabolism in heart could be mediated through an interaction of phosphorylated BAD with hexokinase II is not yet known.

Mitochondrial HK-II and the pro-survival effect

There is accumulating evidence that HKs play a role in protective as well as glycolytic signaling. Treatment of cells with heparin-binding epidermal growth factor (HB-EGF) increases cellular HK activity in renal epithelial cells and expression of HK-I by adenovirus protects against acute oxidant-induced epithelial cell death (Bryson et al. 2002). Similarly, retroviral overexpression of HK-II in lung epithelial-like A549 cells is protective against oxidative injury and this overexpressed HK-II is localized to mitochondria (Ahmad et al. 2002). There is also evidence for HK protection in experiments with isolated mitochondria. Addition of purified mammalian HK-I to mitochondria from rat liver, which do not express mitochondria-associated HKs, was shown to prevent Ca^{2+} -induced mitochondrial swelling and cytochrome c release. In contrast addition of yeast HK lacking the ability to bind to mitochondria was not protective (Azoulay-Zohar et al. 2004). These data indicate that mitochondrial HKs have an inhibitory effect on Ca^{2+} -induced PT-pore opening. Support for a role of endogenous HKs in protection comes from the observation that dissociation of HK-II from mitochondria sensitizes fibroblast to apoptotic death (Majewski et al. 2004a) and induces apoptosis in Hela cells through a PT-pore opening (Chiara et al. 2008). Taken together, these results provide support for the idea that mitochondrial HK-I and HK-II have the ability to protect mitochondria against ROS or Ca^{2+} overload, two well-known PT-pore inducers. HK-II also protects against apoptosis induced by Bcl-2 family proteins such as Bax by competitive inhibition of Bax binding to VDAC (Majewski et al. 2004a; Pastorino and Hoek 2003; Pastorino et al. 2002; Robey and Hay 2006).

The role of HK-II in cardioprotection has not been fully considered although HK-II is abundantly expressed in the heart. Several recent studies have demonstrated a protective role of mitochondrial HKs in cardiomyocytes (Chiara et al. 2008; Miyamoto et al. 2008; Sun et al. 2008). Overexpression of full length HK-I or HK-II was shown to protect against H_2O_2 -induced mitochondrial depolarization in neonatal rat ventricular myocytes (NRVMs) (Sun et al. 2008). Decreased protection was observed using either mutants lacking the mitochondrial binding motif or kinase-dead HK-I or HK-II (Sun et al. 2008), implicating mitochondrial binding and the catalytic activity of hexokinase in protection of NRVMs. Chiara et al. (2008) demonstrated that dissociation of mitochondrial HK-II

increases susceptibility of adult ventricular myocytes to ROS induced PT-pore opening (Chiara et al. 2008). Our work was the first to report that HK-II is phosphorylated by Akt and plays a significant role in Akt-mediated protection against Ca^{2+} overload in isolated heart mitochondria and against H_2O_2 -induced injury of NRVMs (Miyamoto et al. 2008), as described below.

Mitochondrial HKs and Akt mediated protection

The protective effect of mitochondrial HK-II has been linked to the PI3K/Akt signaling axis (Gottlob et al. 2001; Majewski et al. 2004a; Pastorino and Hoek 2003; Pastorino et al. 2002; Rathmell et al. 2003; Robey and Hay 2006). Hay and co-workers demonstrated that expression of membrane targeted constitutively activated Akt (myristoylated-Akt; myr-Akt) increased mitochondrial HK activity in Rat1a fibroblasts and suggested that HK was involved in Akt-mediated cell survival (Gottlob et al. 2001). Cells were protected against growth factor withdrawal and UV exposure, and adenoviral HK-I expression mimicked the protective ability of myr-Akt. It has also been shown that Akt inhibits tBid-induced apoptosis through increased mitochondrial HK activity, antagonizing tBid-mediated BAX and BAK activation at mitochondria (Majewski et al. 2004b). Interestingly, disruption of mitochondrial HK-II association impaired myr-Akt-mediated protection even in the absence of Bax and Bak, critical mediators of MOMP. This strongly suggests that Akt/mitochondrial HK-II has the ability to prevent cell death induced by mechanisms beyond those that regulate Bcl-2 family proteins and MOMP.

Recent work from our laboratory suggested that mitochondrial HK-II plays a crucial role in Akt-mediated inhibition of the PT-pore in cardiomyocytes (Miyamoto et al. 2008). In our study, endogenous Akt was activated by LIF, shown to accumulate at mitochondria and implicated in phosphorylation of mitochondrial HK-II. HK-II has an Akt consensus sequence (RxRxxS/T) that is conserved in rat, mouse and human. The appearance of phosphorylated Akt at mitochondria was associated with increased mitochondrial HK-II. Dissociation of HK-II from mitochondria diminished the protective effects of LIF against H_2O_2 -induced mitochondrial depolarization in NRVMs. Addition of recombinant active Akt to mitochondria isolated from adult mouse heart inhibited Ca^{2+} -induced cytochrome c release and this effect was reversed by dissociation of HK-II from the mitochondria. Thus phosphorylation of mitochondrial HK-II appears to underly Akt-mediated PT-pore inhibition in cardiomyocytes. A similar observation was recently reported in Hela cells where insulin, a strong Akt activating agonist that normally prevents cell death, was unable to antagonize cell death induced by HK-II dissociation from mitochondria (Chiara et al. 2008). These

results provide strong support for the idea that mitochondrial HK-II is a critical effector of Akt-mediated mitochondrial protection.

It is of interest that mitochondrial HK activity in the heart is increased by interventions known to activate Akt, including insulin, preconditioning and morphine (Russell et al. 1992; Zuurbier et al. 2005), although conflicting data has been shown (Doenst et al. 1998). An immunogold labeling study further demonstrated that insulin increases mitochondrial association of HK-I and HK-II and the finding that the increase is greater for HK-II than HK-I suggested differential regulation of distribution of these two mitochondrial HKs (Southworth et al. 2007). This is of particular interest as we found that there is an Akt phosphorylation consensus sequence in mouse, rat and human HK-II but not in HK-I (or HK-III). Thus Akt-mediated phosphorylation and regulation of hexokinase distribution might be specific to HK-II. It will be of interest to investigate whether the Akt/HK-II pathway plays a significant role in preservation of heart integrity against stress such as ischemia/reperfusion *in vivo*.

There is evidence that Akt also regulates the expression of HK-II at the transcriptional level. Insulin treatment was shown to increase HK-II mRNA in adipose and skeletal muscle cell lines (Printz et al. 1993). A subsequent study by the same group showed that insulin induced upregulation of HK-II but not HK-I and this induction of HK-II was inhibited by PI3K blockers or rapamycin, suggesting a contribution of PI3K/Akt and mTOR to induction of HK-II mRNA by insulin (Osawa et al. 1996). Similar observations were reported by other groups examining the effect of insulin in 3T3-L1 adipocytes (Culbert and Tavare 2002) and colony-stimulating factor-1 in myeloid cells (Lee and States 2006). We also observed that HK-II protein expression is increased in NRVMs in response to 48 h LIF treatment or overexpression of wild-type Akt (unpublished observation), consistent with a role for Akt in HK-II upregulation in the heart. Thus, in addition to acute regulation of mitochondrial HK-II through phosphorylation, the upregulation of HK-II induced by Akt activation may contribute to protective effects of Akt under chronic stress.

How does mitochondrial HK-II prevent PT-pore opening?

Although protective roles of mitochondrial HKs have been clearly demonstrated, the mechanism by which mitochondrial HK-I and HK-II protect mitochondria are not clear. Mitochondrial HK has preferential access to ATP generated by mitochondria and the ADP produced by this reaction can be transported back into mitochondria by the ANT and used as a substrate by F_1F_0 ATP synthase to produce ATP (ADP recycling). By providing mitochondria with a constant supply of ADP (presuming an adequate supply of inorganic

phosphate), mitochondria can slightly depolarize the membrane and can thus minimize ROS production by the electron transport chain (da-Silva et al. 2004). This could contribute to protection, although this needs to be experimentally proven especially under stressed conditions. In fact, apoptosis induced by HK-II detachment from mitochondria was prevented by deletion or inhibition of Cyp-D or inhibition of the ANT, but not by an F_1F_0 ATP synthase inhibitor (Chiara et al. 2008). While VDAC is an established binding partner of HK-II at the OMM, it has been shown using VDAC KO MEF cells that VDAC is not involved in apoptosis induced by HK-II dissociation (Chiara et al. 2008).

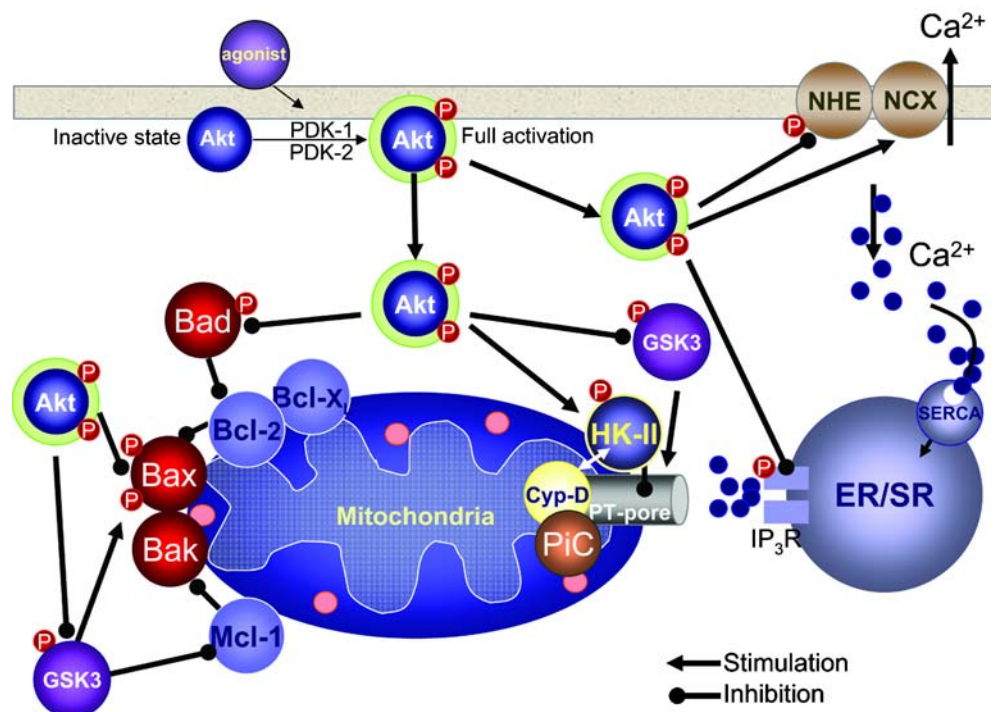
Interestingly, pharmacological inhibition or knockdown of Cyp-D resulted in release of HK-II from mitochondria (Chiara et al. 2008; Machida et al. 2006), suggesting that Cyp-D stabilizes mitochondrial HK-II binding. Cyp-D and HK-II have both been postulated to be regulatory components of the PT-pore. Gene deletion of Cyp-D blocks PT-pore opening and conversely overexpression of Cyp-D induces mitochondrial swelling and cell death in the heart (Baines et al. 2005). In contrast to the stimulatory role of Cyp-D in PT-pore induction in the heart, mitochondrial HKs play an inhibitory role in PT-pore opening. These positive and negative regulators appear to interact with each other to regulate pore opening. However, it remains unclear which molecule(s) mediate the interaction of these two PT-pore regulators. Cyp-D is a mitochondrial matrix protein

and HK-II binds to the OMM thus it is unlikely that they physically interact. Cyp-D and HK-II bind to ANT and VDAC respectively, but the recent exclusion of these proteins as necessary for PT-pore opening clearly suggests that other proteins must be involved in HK-II and Cyp-D-mediated protection. Recent data suggests that the phosphate carrier binds to CypD, and this inner membrane transporter has been implicated in formation of the PT pore (Alcala et al. 2008; Leung et al. 2008). As well, phosphate has recently been shown to be required for inhibition of PT-pore formation by CsA or by CypD ablation (Basso et al. 2008), which may imply regulation of phosphate carrier conformation by the presence of its substrate. Recently, Cyp-D was suggested to directly bind to Bcl-2 (Eliseev et al. 2009), which would require Bcl-2 to be localized to and span the inner membrane. New avenues of exploration will need to be pursued to identify the full complement of proteins involved in HK-II-mediated protection of mitochondria.

Upstream of mitochondrial PT-pore formation: Inhibition of cytosolic Ca^{2+} overload by Akt

Cytosolic Ca^{2+} concentration is tightly regulated such that the intracellular Ca^{2+} concentration is kept at 100 nM~1 μ M, one to ten thousands fold lower than that in the extracellular space. Dysregulation of Ca^{2+} handling leads to cytosolic Ca^{2+} overload, followed by excessive mitochon-

Fig. 1 Akt preserves mitochondrial integrity against stress through multiple target molecules in the heart



drial Ca^{2+} sequestration and PT-pore opening. Akt appears able to prevent Ca^{2+} overloading under pathophysiological settings. For instance, we have found that LIF pre-treatment dramatically dampens the increase in cytoplasmic Ca^{2+} in NRVMs treated with H_2O_2 (unpublished observation), and we previously reported that myr-Akt expression in NRVMs reversed the decrease in expression of $\text{Na}^+/\text{Ca}^{2+}$ exchanger1 (NCX1) induced by constitutively active Gq α (Q209L) expression. NCX1 is responsible for Ca^{2+} extrusion from the cytoplasm in cardiomyocytes. Akt-mediated preservation of NCX1 expression prevents cytoplasmic Ca^{2+} overload, opening of the PT-pore and apoptosis induced by Q209L (Miyamoto et al. 2005). Akt-mediated transcriptional up-regulation of NCX1 and NCX3 has recently also been reported in a study on PC12 cells which further supports the idea that enhanced Ca^{2+} extrusion via NCX contributes to Akt-mediated protection (Formisano et al. 2008). Interestingly, the cardiac Na^+/H^+ exchanger (NHE) has been discovered to be phosphorylated by Akt leading to inhibition of its activity (Snabaitis et al. 2008). NHE imports Na^+ ions and exports H^+ ions in response to acidosis during ischemia. The resultant intracellular accumulation of Na^+ ions, in turn, reverses NCX ion transport to the Ca^{2+} import mode, leading to cytosolic Ca^{2+} overloading. NHE inhibition during I/R has been established to be protective in many experimental models (Avkiran et al. 2008) and NHE phosphorylation and inhibition mediated by Akt would provide a molecular mechanism for prevention of cytoplasmic Ca^{2+} overloading in the heart.

Another interesting new finding is that Akt phosphorylates type-I IP_3 receptors (IP_3R) which inhibits Ca^{2+} release from the ER and prevents Ca^{2+} induced apoptosis in non-cardiac cells (Khan et al. 2006; Szado et al. 2008). A consensus sequence for Akt phosphorylation (RxxRxxS/T) is present in all three types of IP_3Rs . Although the contribution of IP_3Rs to cardiac Ca^{2+} regulation has been thought to be minimal, there is compelling evidence that Ca^{2+} release from IP_3Rs can regulate cardiac cellular functions including positive inotropy, gene expression, induction of arrhythmia and regulation of pacemaking (Domeier et al. 2008; Kockskamper et al. 2008; Roderick and Bootman 2007; Woodcock et al. 2008; Wu et al. 2006). Therefore, it is probable that IP_3 -induced Ca^{2+} release plays a role in regulation of mitochondrial Ca^{2+} , as observed in NRVMs stimulated with ATP (Jaconi et al. 2000) and in the heart (Garcia-Perez et al. 2008). ER/SR Ca^{2+} overloading has been receiving increasing attention in the context of regulation of cell death, since ER/SR Ca^{2+} pools communicate with mitochondria due to the proximity of release site and mitochondria (Diwan et al. 2008; Garcia-Perez et al. 2008; Jaconi et al. 2000; Szalai et al. 1999). Notably, IP_3R expression is increased in heart disease (Go et al. 1995), thus IP_3R signaling and its regulation by Akt may

have increased importance in regulation of mitochondrial Ca^{2+} and cardiomyocyte survival under pathophysiological conditions. Finally, it is becoming clear that Bcl-2 family proteins not only function at mitochondria but also localize to and control the Ca^{2+} content of intracellular Ca^{2+} pools (Pinton and Rizzuto 2006). Apoptotic Bcl-2 family members such as Bax, Bak and Nix localize at ER/SR and increase their Ca^{2+} content, resulting in large increases in Ca^{2+} release and subsequent mitochondrial Ca^{2+} overloading (Demaurex and Distelhorst 2003; Diwan et al. 2008; Nutt et al. 2002; Scorrano et al. 2003). In contrast, anti-apoptotic Bcl-2 family proteins decrease ER Ca^{2+} content and Ca^{2+} release (Demaurex and Distelhorst 2003; Palmer et al. 2004). It will be of considerable interest to determine whether Bcl-2 family proteins play a role in SR-mitochondrial Ca^{2+} crosstalk in the heart, as demonstrated in Nix mediated cardiac cell death (Diwan et al. 2008), and whether Akt-mediated phosphorylation of Bcl-2 family proteins can lead to change in their ER/SR localization.

Conclusion

Akt provides a strong survival signal in various cell types including cardiomyocytes and exerts its protective effects through phosphorylation of diverse target molecules such as Bcl-2 family proteins, GSK-3 β and, as recently demonstrated HK-II (Fig. 1). Preservation of mitochondrial integrity is critical for regulation of metabolism and cell survival, and recent compelling evidence indicates that Akt can serve as an integrator and gatekeeper of both of these mitochondrial functions.

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