



# The SAFE pathway for cardioprotection: is this a promising target?

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## Abstract

The survivor activating factor enhancement (SAFE) pathway was discovered as an alternative intrinsic pro-survival signaling pathway to the reperfusion injury salvage kinase pathway for cardioprotection against ischemia–reperfusion injury. The delineation of this pathway, made of key components such as cytokines of the immune system and transcription factors, has brought major advancements in our understanding on how the heart is able to protect itself against ischemia–reperfusion injury. In this viewpoint, we describe the major steps leading to the discovery of the SAFE pathway in small animal models to date and we discuss its translation to large animals and humans.

**Keywords** Cardioprotection · Prosurvival signaling pathways · SAFE pathway · Immune system · Ischemia–reperfusion injury

## Intrinsic cardiac signaling pathways to confer cardioprotection

Following the major discovery of the ischemic preconditioning (IPC) phenomenon (whereby small episodes of ischemia–reperfusion can confer cardioprotection against a sustained ischemic insult), it became evident that the heart has the ability to activate intrinsic cardioprotective signaling pathways to protect itself against different stress situations [42]. Over the past 30 years, intense research conducted with the aim to better understand the signaling process involved in ischemic conditioning has led to the discoveries of major key prosurvival molecules and has paved the way for potential novel therapeutic targets [16]. In the 90s, the role of adenosine, protein kinase C and the putative mitochondrial potassium–adenosine triphosphate dependent (mKATP) channel were amongst the key signaling components described in the

cardioprotective signaling cascade of ischemic conditioning (see reviews [8, 9]). In 2002, Yellon's group described the reperfusion injury salvage kinase (RISK), highlighting the importance for the activation of the kinases protein kinase B (Akt) and extra regulated kinase (Erk) to limit ischemia–reperfusion injury with cardioprotective strategies [50]. At the same time, they also proposed that IPC protects the myocardium by inhibiting the mitochondrial permeability transition pore (mPTP) opening at reperfusion, thus reducing cardiomyocyte death by limiting uncoupling oxidative phosphorylation and swelling of the mitochondria [17]. In 2009, an important other step was made in the delineation of the cardioprotective signaling cascade with the discovery of the survivor activating factor enhancement pathway (SAFE) [34]. This path, triggered by the immune system, initiates the activation of a RISK-alternative pathway with a multitude of prosurvival signaling components, including the transcription factor signal transducer and activator of transcription 3 (STAT3) that targets the inhibition of the mPTP opening [4, 11, 31].

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## The discovery of the SAFE pathway, from conception to birth: a 10 year fascinating journey

Officially born in 2009, the conception of the SAFE pathway goes back to 1999 with the discovery that tumor necrosis factor alpha (TNF), a key player of the immune system, could protect against ischemia–reperfusion injury when given as a preconditioning stimulus [36, 38]. At the time, this discovery was somewhat controversial as major clinical trials with anti-TNF in ischemic heart failure were initiated (RECOVER–RENAISSANCE–RECOVER) [1]. However, most clinical trials showed an adverse effect of the anti-TNF, a result that could be partly explained by our findings that TNF, in a dose and time dependent manner, can confer cardioprotection after binding to its receptor type 2 (TNFR2) [7, 32, 49, 51]. TNF is, in fact, an important signaling component contributing to the protection of both ischemic pre- and post-conditioning [32, 53]. Deciphering the signaling targets downstream of TNF for cardioprotection, protein kinase C, reactive oxygen species and mKATP channel were identified [37, 38]. To our surprise, the protective effect of TNF was not abolished in the presence of pharmacological inhibitors for Akt and Erk, therefore, suggesting that TNF confers cardioprotection via the activation of a RISK-alternative pathway [39]. Marber’s group showed that protection with TNF is independent of p38 MAPK activation [58]. Also, the protection of TNF is abolished in the presence of an inhibitor of the Janus kinase (JAK)/STAT3 pathway or in cardiomyocytes of STAT3 knockout mice, therefore, suggesting that TNFR2 triggers the phosphorylation of JAK, which in turn, can trigger the phosphorylation of STAT3 on a tyrosine residue and its activation (after dimerization) [32, 39] (see review [45] for further details on STAT3 activation). This TNF/TNFR2/JAK/STAT3 pathway was named as the SAFE pathway and similar to the RISK path, it targets the inhibition of the mPTP opening to promote cardiomyocyte survival [4, 11, 18, 31, 34].

Although the activation of the SAFE pathway has been confirmed with multiple cardioprotective strategies [10, 24, 26, 29, 33, 48, 62], the understanding of the exact signaling cascade of this pathway is still in its infancy and many fundamental questions remain to be answered, including the following.

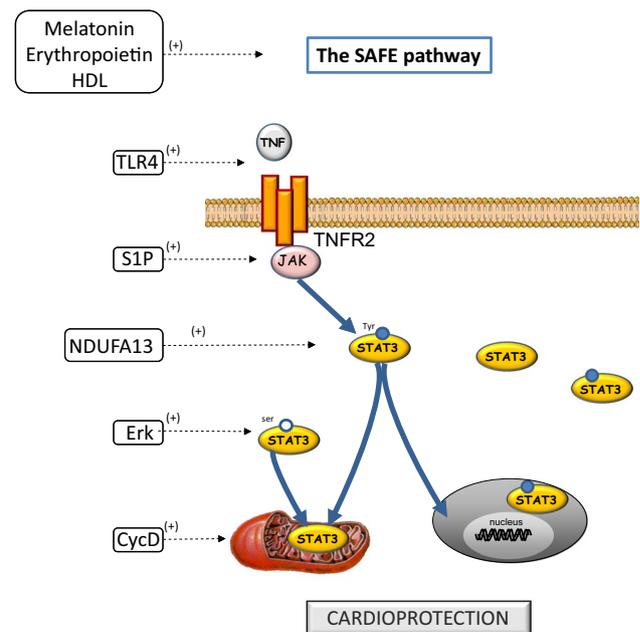
### What are the trigger signaling components upstream of TNF?

Endogenous circulating components such as sphingosine-1 phosphate, melatonin, high density lipoproteins,

and erythropoietin can activate the SAFE pathway [6, 33, 54, 59]. Although the toll like receptor 4 (TLR4) has been suggested to activate TNF as part of the SAFE path, the exact mechanism involved in TLR4 releasing TNF in the cardiomyocyte still remains to be elucidated [44] (see Fig. 1).

### What is the cellular origin of TNF?

The presence of TNF has been identified in multiple cells ranging from inflammatory cells to cardiomyocytes and fibroblasts. To identify the cellular origin of TNF that activates survival signaling cascades against ischemia–reperfusion injury, we designed cardiomyocytes specific TNF knockout mice. In these mice, ischemic postconditioning failed to protect and activate STAT3, therefore, suggesting that TNF originating from the cardiomyocytes plays a major role in the activation of the SAFE pathway for cardioprotection [30].



**Fig. 1** Major components of the survivor activating factor enhance (SAFE) pathway that promotes protection against ischemia–reperfusion injury. The activation of the SAFE pathway is triggered by the binding of TNF onto the TNF receptor 2 which, in turn, will activate the phosphorylation of JAK and STAT3. Melatonin, erythropoietin and HDL are endogenous circulatory molecules that can trigger the activation of the SAFE pathway. TLR4, S1P, NDUFA13, Erk and cyclophilin D will modulate the activation of the SAFE pathway. (+): stimulation. *CycD* cyclophilin D, *Erk* extra regulated kinase, *HDL* high density lipoprotein, *JAK* janus kinase, *NDUFA13* nicotinamide adenine dinucleotide dehydrogenase (ubiquinone) 1 alpha sub-complex 13, *Ser* serine, *S1P* sphingosine-1 phosphate, *STAT3* signal transducer and activator of transcription 3, *TLR4* toll like receptor 4, *TNF* tumor necrosis factor alpha, *TNFR2* TNF receptor 2, *Tyr* tyrosine

## How can TNF activate STAT3 after binding to TNFR2?

Whereas the binding of JAK onto the interleukin-6 receptor is well described in the literature, no binding site for JAK has been identified on the TNFR2 until now. However, there is increasing evidence that activation of STAT3 by TNF is regulated by sphingosine kinase 1 and possibly sphingosine-1 phosphate [25, 46]. It is also suggested that the dimerization of STAT3 is controlled by nicotinamide adenine dinucleotide dehydrogenase (ubiquinone) 1 alpha subcomplex 13 (NDUFA13), a subunit of mitochondria complex I through the control of hydrogen peroxide formation by complex I [23].

## What are the main downstream targets of STAT3?

Following stimulation by TNFR2, activated STAT3 interacts with Nuclear Factor kappa B signaling to upregulate mitochondrial fusion by activation of optic atrophy 1 (OPA1) expression [43]. It is now well established that phosphorylated STAT3 is present in both the nucleus and the mitochondria during ischemia–reperfusion [56, 57, 60]. The mitochondrial pool of STAT3 is dependent on phosphorylation of a serine residue and is stabilized by cyclophilin D, thus enabling STAT3 to regulate reactive oxygen species levels, the electron transport chain activity and the mPTP activity [4, 5, 40]. In pigs, phosphorylation of the tyrosine residue of STAT3 in the mitochondria mediates cardioprotection by postconditioning [19].

## Does long-term stimulation of the SAFE pathway confer cardioprotection?

Although the transient activation of TNF and interleukin 6 at the onset of reperfusion is cardioprotective [38, 41], a chronic stimulation of the SAFE pathway is unlikely to benefit the heart [35]. In experimental ischemic heart failure, excess stimulation of TNF will contribute to inflammation, apoptosis and remodeling processes (via the TNF receptor 1) [15, 51]. Similarly, [19] inflammation and adverse cardiac remodeling is observed in mice with continuous activation of STAT3 [22]. In contrast, female cardiomyocyte STAT3 deficient mice develop postpartum cardiomyopathy which suggests that STAT3 may protect from postpartum oxidative stress [21].

## SAFE pathway and cross talk with other cardioprotective components

Although many different intrinsic signaling molecules and mechanisms have been described in the heart, it is not quite clear yet whether these molecules interact with each other

to confer cardioprotection. It is questionable as to whether the activation of both RISK and SAFE pathways is required to maximize the protection.

In fact, a cross-talk between the RISK and SAFE pathway has already been described. In STAT3 cardiomyocyte deficient mice or in the presence of a STAT3 inhibitor in wild-type mice, cardioprotective preconditioning strategies fail to protect and to activate components of the RISK pathway such as Akt and Erk [54, 55, 62]. Similarly, inhibition of the RISK pathway with Akt or Erk inhibitors results in the lack of protection of conditioning strategies and failure to activate STAT3 [54, 55]. In fact, Erk may also be a key player in the regulation of serine phosphorylation of mitochondrial STAT3 [14, 62].

Some cardioprotective strategies may require the activation of both RISK and SAFE pathways while other strategies may require only one of the two pathways for protection [24, 47, 48, 63]. However, caution needs to be taken when interpreting the published results as many studies have based their conclusion on western blot results only (absence of activation of Akt, STAT3 or Erk) performed at a very specific time point and without validating their results with the inhibition of the protein of interest. Considering that these proteins are activated at different time points during the reperfusion period, western blot data are clearly insufficient to draw any conclusion on the activation of the SAFE versus RISK pathway, unless kinetic studies with multiple time points are performed [27].

## SAFE pathway and cardioprotection: from small to large animals to humans

To consider the SAFE pathway as a potential therapeutic target for novel cardioprotective strategies, it is critical to explore whether the activation of this pathway can be found in large animals and in humans with known cardioprotective strategies. Heusch's group has been able to demonstrate the translation from small animal models to a pig model whereby the activation of STAT3 with cardioprotective strategies such as ischemic pre- and post-conditioning as well as remote ischemic conditioning is required for the protection [13, 19, 52]. In humans, demonstrating the causal role of the SAFE pathway is challenging: activation of proteins can be identified but whether these proteins actually contribute to the protection remains to be validated. Nevertheless, Heusch's group elegantly measured the activation of various proteins in cardiac biopsies of patients subjected to remote ischemic preconditioning. To their surprise, activation of STAT5 but not STAT3 was increased in the presence of the conditioning stimulus, therefore, suggesting that STAT5 may play a critical role for cardioprotection in a human setting [20]. In children undergoing tetralogy of Fallot repair

surgery, remote ischemic preconditioning was associated with the activation of both STAT3 and STAT5 [61]. The increase in cytokine levels after remote ischemic conditioning could be observed for interleukin 1 alpha but not TNF [12].

Although the activation of the SAFE pathway offers a promising target for the development of novel therapeutic approaches to limit ischemia–reperfusion injury, the exact delineation of its signaling components in humans is critical as these might differ from small animals or even pigs. In addition, very little information is known about the effect of confounders and comorbidities on the activation of the SAFE pathway. Aging, obesity and propofol alter the activation of STATs [2, 3, 28] and a better understanding on the effect of such confounders on the SAFE path are required before cardioprotective strategies targeting the SAFE pathway can be considered in the clinical setting.

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## Compliance with ethical standards

**Conflict of interest** The authors declare that they have no conflict of interest.

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