

Do Multiple Nuclear Factor Kappa B Activation Mechanisms Explain Its Varied Effects in the Heart?

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ABSTRACT

Background: Multiple studies have demonstrated the important role of the nuclear factor kappa B (NF- κ B) in cardiac pathology. However, these studies' conclusions differ regarding whether NF- κ B is protective or detrimental for heart function. This disagreement is not surprising considering the complexity of NF- κ B signaling that involves multiple components and regulation at several steps. Furthermore, NF- κ B is a pleiotropic transcription factor that receives signals from multiple pathways, including the renin-angiotensin system (RAS) and cytokines, 2 important modulators of cardiac remodeling.

Methods: In this article, we review studies related to the role and mechanisms of NF- κ B activation in the heart, particularly with regard to the RAS, inflammation, and diabetes. The objective of this review is to consolidate multiple, often contradictory, findings to develop a clear understanding of NF- κ B signaling in the heart.

Conclusions: The studies we review demonstrate that NF- κ B effects in the heart are mechanism specific and that NF- κ B signaling is cyclical. Consequently, the timing of NF- κ B measurement is critical, and studies focused on temporal changes in the NF- κ B mechanism would help clarify its multiple roles in cardiac pathophysiology.

INTRODUCTION

Cardiovascular disease (CVD) is a major cause of morbidity and mortality in the modern world. The prevalence of CVD is on the rise because of the aging population and the increase in obesity and type 2 diabetes. Although the term CVD includes a variety of diseases with multiple causes, the common outcome of CVD is cardiac remodeling that leads to heart failure. Cardiac remodeling is associated with increased oxidative stress, inflammation, and activation of hormonal systems, all of which are involved in disease progression.^{1,2}

Clinical and experimental studies have demonstrated a role of the renin-angiotensin system (RAS) in cardiac remodeling and heart failure. Chronic inflammation also has deleterious effects on cardiovascular function. However, these 2 mechanisms are not completely independent. The nuclear factor kappa B (NF- κ B) family of transcription factors is the major mediator of inflammation. Additionally, NF- κ B controls angiotensinogen (AGT) gene expression, the precursor of a bioactive peptide of the RAS, angiotensin II (AngII). AngII activates NF- κ B, thereby generating a positive feedback loop. Tumor necrosis factor- α (TNF- α), a proinflammatory cytokine, also activates and is positively regulated by NF- κ B. The observation that angiotensin receptor blockers (ARBs) and angiotensin-converting enzyme (ACE) inhibitors reduce inflammation suggests that a part of their protective mechanism may be through the NF- κ B pathway.³

Additionally, oxidative stress increases the expression of AGT through the activation of NF- κ B in hepatocytes and vascular smooth muscle cells.⁴ AngII in turn activates NF- κ B through the production of reactive oxygen species (ROS), thereby generating a feedback loop.⁵

Clinical and experimental studies have shown important roles of both the RAS and inflammation in diabetic cardiomyopathy. NF- κ B regulates both pathways, suggesting that it may be a major hub of cellular activity in diabetes. NF- κ B signaling mechanisms are complex and produce a wide variety of sometimes opposite biological effects in diverse cell types and organs.⁵ The effects of NF- κ B in the heart

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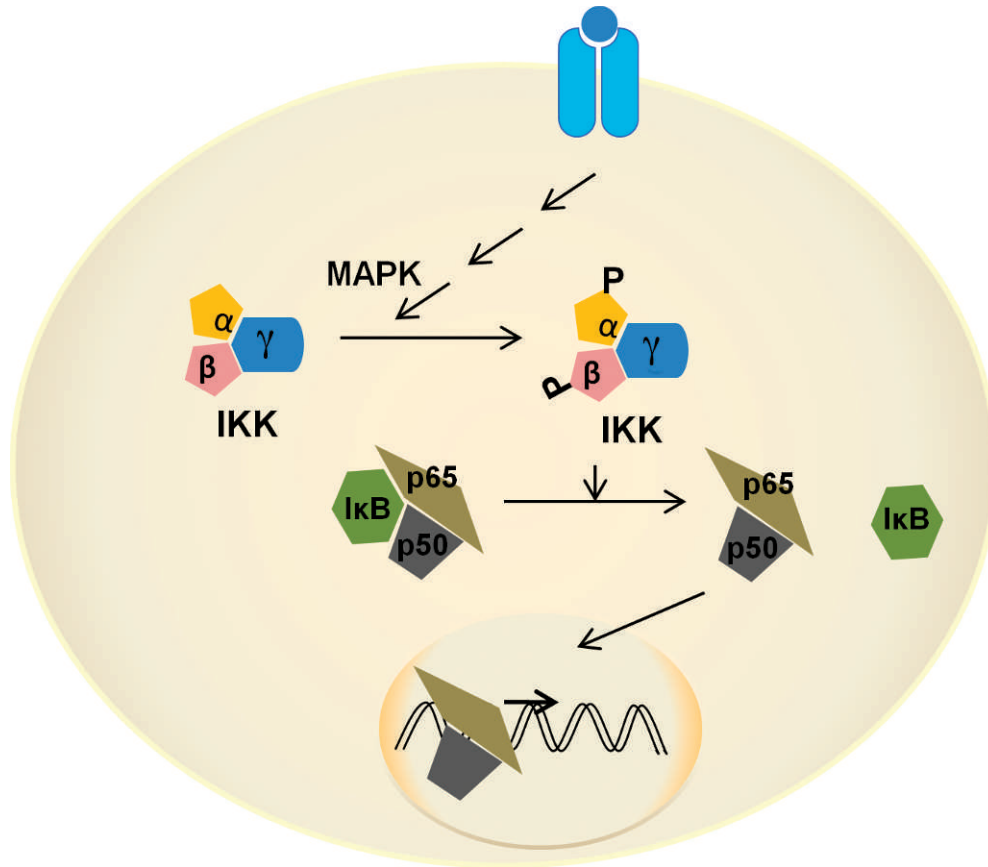


Figure 1. Canonical nuclear factor kappa B (NF- κ B) signaling. In the canonical pathway, phosphorylation of I κ B kinase α/β by mitogen-activated protein kinase (MAPK) is followed by phosphorylation of I κ B- α that occurs in an inactive complex with p50/p65. Phosphorylated I κ B- α is released and degraded in the cytoplasm. The active heterodimer of p50/p65 enters the nucleus to regulate expression of multiple genes. I κ B, inhibitory kappa B; IKK, inhibitory kappa B kinase; P, phosphorylation.

are likewise diverse, with some studies pointing to cardioprotective effects and others demonstrating cardioprotection by blocking NF- κ B activity.⁶⁻¹⁴ In this article, we review different mechanisms that activate NF- κ B in the heart and demonstrate how crosstalk between inflammation and the RAS, through NF- κ B, might influence disease progression.

NF- κ B SIGNALING PATHWAYS

The NF- κ B family consists of 5 members: p50/p105 (NF- κ B1), p52/p100 (NF- κ B2), p65 (RelA), RelB, and c-Rel. The member proteins form homo- or heterodimers, of which the p50/p65 heterodimer is the most abundant and is responsible for the majority of NF- κ B canonical transcriptional activity (Figure 1). Homodimers of p50 subunits have been associated with inhibitory transcriptional activity.¹⁵ Generally, NF- κ B dimers associate with an inhibitory- κ B (I κ B- α) protein that keeps the dimer in the cytoplasm in an

inactive state. NF- κ B activation begins with the activation of an I κ B kinase (IKK) complex that consists of catalytic subunits IKK- α and IKK- β and the scaffolding subunit IKK- γ (the NF- κ B essential modifier [NEMO]). Several mitogen-activated protein (MAP) kinases that also include NF- κ B-inducing kinase (NIK) activate IKK through the phosphorylation of IKK- α and IKK- β . IKK- β has higher activity than IKK- α for I κ B- α and is considered important in the canonical pathway.⁵ In the canonical pathway of NF- κ B activation, I κ B- α is phosphorylated at serine residue (Ser)32,36 and/or Tyr42 and separated from the p50/p65 dimer, allowing the dimer to translocate to the nucleus and bind to cognate DNA sequences.

The noncanonical pathway involves the activation of NIK, resulting in the release of the p52/RelB heterodimer from inhibition (Figure 2).⁵ In yet another mechanism relevant to AngII-mediated NF- κ B activation, Ser536 phosphorylation of p65 results in I κ B- α -

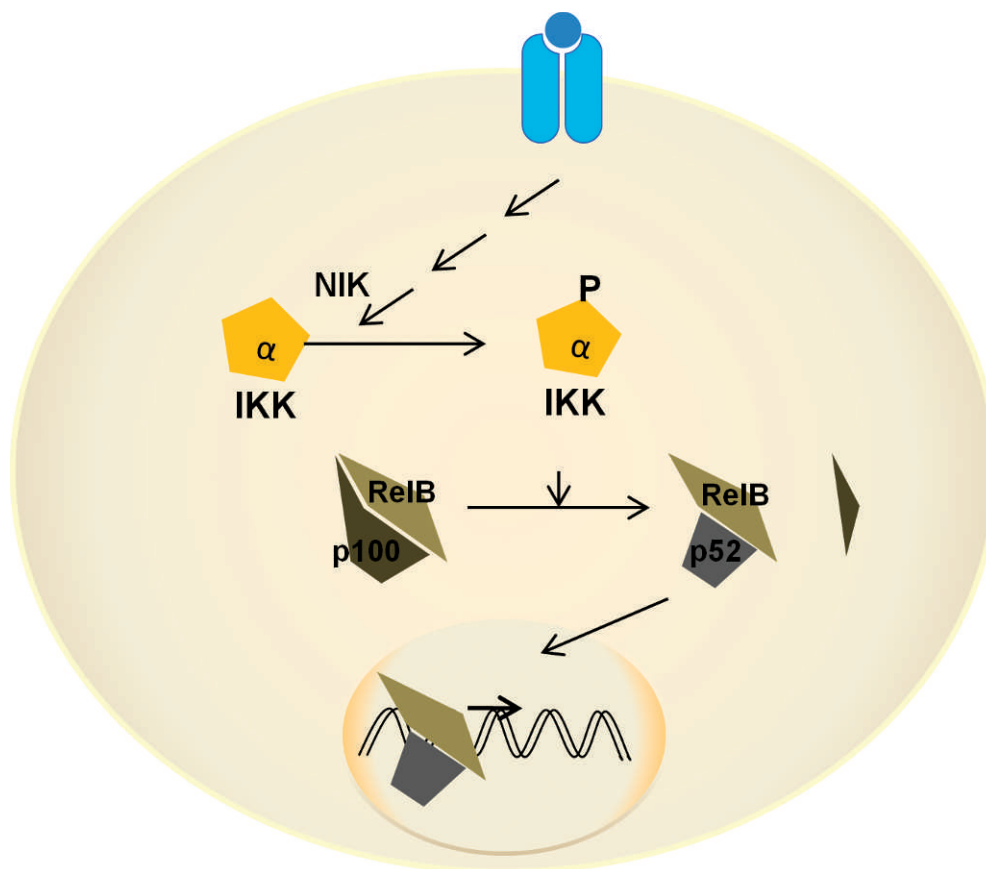


Figure 2. Noncanonical nuclear factor kappa B (NF- κ B) signaling. In the noncanonical pathway, NIK has an important role in the phosphorylation and activation of I κ B kinase α (IKK- α). IKK- α phosphorylates p100 that exists in an inactive complex with RelB. Phosphorylated p100 is partially cleaved to generate the active form p52. The p52/RelB heterodimer enters the nucleus and regulates the expression of several genes distinct from those activated by the canonical pathway. IKK, inhibitory kappa B kinase; NIK, nuclear factor kappa B-inducing kinase.

independent stimulation of p65 transcriptional activity.¹⁶ Recently, a new mechanism of NF- κ B activation through monomethylation of p65 at K37 by histone H3-lysine 4 methyltransferase, SET7/9, has been described.^{17,18} Methylation of p65 was induced by TNF- α and interleukin-1 beta (IL-1 β) and was responsible for the expression of about 25% of NF- κ B-responsive genes in TNF- α -stimulated monocytes.¹⁸ Receptor-for-advanced-glycation-endproducts ligands induced inflammatory genes in monocytes. The expression of the inflammatory genes was inhibited by small interfering RNA-mediated SET7/9 knockdown, suggesting a role of this NF- κ B activation mechanism in diabetes.¹⁸

Acetylation of p50 has been identified as yet another novel mechanism of NF- κ B activation that was demonstrated as essential for cardiac protection against ischemia/reperfusion (I/R) injury.¹⁹ In addition to the NF- κ B subunits that translocate to the nucleus

following activation, the cytoplasmic signal integrators—IKK subunits, NIK, and I κ B- α —undergo nucleocytoplasmic shuttling, suggesting that these proteins might have independent nuclear functions.²⁰

NF- κ B subunits are ubiquitously expressed and activated by multiple mechanisms and by different stimuli in diverse cell types, resulting in a wide variety of sometimes opposite biological effects. The NF- κ B canonical pathway is the most extensively studied mechanism. The canonical pathway regulates expression of about 200 genes, including genes coding for transcription factors.⁵ NF- κ B thus may have direct as well as indirect effects through these transcription factors. Direct effects further consist of early and late response genes. Some of the proteins coded by NF- κ B regulatory genes are inhibitors of NF- κ B signaling, such as I κ B- α , and produce a negative feedback control. Other genes produce proteins that are involved in generating signals that activate NF- κ B

pathways, such as nicotinamide adenine dinucleotide phosphate (NADPH) oxidase expression.²¹ Information about the gene expression profile activated by the noncanonical NF- κ B pathway is limited. However, this expression profile appears to be distinct from the canonical pathway.⁵ Similarly, the effects of the acetylation of p50 and the methylation and phosphorylation at different sites of p65 are incompletely understood.

ROLE OF NF- κ B IN CARDIAC PATHOPHYSIOLOGY

NF- κ B regulates multiple cellular processes, including cell maturation, survival, and proliferation, and systemic processes such as inflammation. NF- κ B is activated by stress and inflammatory stimuli that include cytokines, vasoactive peptides, and viral oncogenes. NF- κ B activation is generally transient because of negative feedback regulation. Although a short activation of NF- κ B might be beneficial, a prolonged activation may be detrimental by promoting chronic inflammation. Thus, the duration as well as the timing and cellular context—ie, the cell type and the stimulus—may determine the outcome of NF- κ B signaling in the heart. Several studies have demonstrated a role of NF- κ B signaling in different cardiac pathological states. Transaortic coarctation, a model of pressure overload, increased NF- κ B activity in the mouse heart that developed oxidative stress and cardiac hypertrophy.²² Adenovirus-mediated intramyocardial gene transfer of a dominant negative form of Akt or cytoplasmic superoxide dismutase attenuated NF- κ B activity and cardiac hypertrophy, suggesting a role of NF- κ B in pressure overload.²² In Dahl salt-sensitive hypertensive rats, a high-salt diet for 18 weeks increased NF- κ B activity in the heart and increased expression of TNF- α , interleukin-6 (IL-6), and IL-1 β .²³ Similarly, in spontaneously hypertensive rats (SHRs), cardiac expression of IL-6 and IL-1 β increased as did the expression and activity of p65 NF- κ B.²⁴ In the following sections, we categorize studies based on the NF- κ B subunits to determine whether different NF- κ B activation mechanisms produce different outcomes in the context of heart diseases.

Role of the NF- κ B p50 Subunit

In mice deficient for p50 NF- κ B, p65 expression and nuclear localization were increased following the stimulation of mouse embryonic fibroblasts with lipopolysaccharide, resulting in increased IL-6 secretion compared to wild-type (WT) controls.²⁵ Further, p50 deficiency increased cardiac remodeling and systolic dysfunction in response to myocardial ischemia. These studies suggest that p50 has a protective

role in inflammation and cardiac remodeling and that p65 can function without the need for heterodimerization with p50, as in the canonical pathway.²⁵ However, in another study on p50 knockout mice, coronary artery ligation-induced ventricular dilatation over 8 weeks was less in knockout animals than in WT mice.²⁶ In an I/R model, protection of the heart by a histone deacetylase inhibitor, trichostatin A (TSA), was accompanied by increased nuclear localization of p50 NF- κ B. TSA treatment resulted in increased acetylation of p50.¹⁹ In hearts from p50-deficient mice, TSA did not protect against I/R injury, suggesting an essential role of p50 in cardioprotection.¹⁹ In a human study, the p50 NF- κ B (NF- κ B1) promoter polymorphism (ATTG₁)—associated with lower expression of p50—was correlated with heart failure in patients. The ATTG₁ genotype was associated with enhanced cardiac remodeling and impaired heart function.²⁷ Overexpression of p50 in H9c2 cells, a rat cardiomyoblast cell line, repressed expression of the c-Rel subunit of NF- κ B; c-Rel has been shown to stimulate cardiac hypertrophy and fibrosis.²⁸ Confirming these in vitro observations, hearts from p50 knockout mice were larger than WT animals' hearts and had increased c-Rel expression.²⁸ The above studies largely show a protective role of p50 in the myocardium that may be associated with an inhibitory effect on gene transcription caused by a lack of a transactivation domain.

Role of the NF- κ B p65 Subunit

In a heart failure model of coronary ligation in mice, p65 and p50 subunits of NF- κ B were translocated to the nucleus for up to 24 hours of infarction. However, only p65 was chronically activated in the myocardium as determined by DNA binding and electrophoresis mobility shift assay (EMSA).¹³ Transgenic mice overexpressing phosphorylation-resistant I κ B- α mutant showed improved survival, cardiac remodeling, and function and reduced inflammation and apoptosis.¹³ These studies suggested that persistent activation of p65 contributed to cardiac remodeling in myocardial infarction. In vitro studies using adult rat cardiomyocytes showed increased p65 nuclear translocation following cyclical stretch for 24 hours that resulted in vascular endothelial growth factor secretion.²⁹ Phenylephrine activation of NF- κ B in H9c2 cells was caused by increased phosphorylation of the p65 subunit and was associated with atrial natriuretic factor promoter activity, suggesting a role of NF- κ B in phenylephrine-induced hypertrophy.¹⁴ Similar phosphorylation of p65 and its involvement in the development of cardiac hypertrophy were demonstrated in SHRs.¹⁴ Recently, a direct physical interaction between p65 NF- κ B and the nuclear factor

of activated T cells (NFAT) transcription factor was demonstrated in cardiomyocytes, resulting in synergistic activation of NFAT transcriptional activity.³⁰ Further, NFAT activity was reduced in the hearts of cardiac-specific p65 knockout mice that also showed attenuated pressure overload-induced cardiac hypertrophy.³¹ The calcineurin-NFAT pathway is known as a regulator of pathological cardiac remodeling, suggesting that activation of p65 has an adverse role in cardiac pathophysiology.³¹ Similarly, expression of G protein-coupled receptor kinase 5 (GRK5), which is increased in heart failure patients, is regulated by the binding of p65.³² Inhibition of NF- κ B by the regulator of G-protein signaling homology domain of GRK5 reduced cardiac mass in SHR and prevented phenylephrine-induced cardiac hypertrophy in Wistar Kyoto rats.¹⁴ Recently, transgenic mice with cardiac-specific overexpression of phosphorylation-deficient triple mutant I κ B- α showed attenuated nuclear translocation of p65 in response to various stimuli.³³ We demonstrated that these mice are protected from right ventricular hypertrophy resulting from monocrotaline-induced pulmonary hypertension.³⁴ Similarly, transgenic mice overexpressing myotrophin in the heart developed cardiac hypertrophy and heart failure, associated with increased NF- κ B activity as determined by p65 DNA binding.⁶ Lentivirus-mediated delivery of p65 short hairpin RNA in the hearts of these animals attenuated NF- κ B activity and regressed cardiac hypertrophy. In an inducible transgenic mouse model, with cardiomyocyte-specific expression of constitutively active IKK- β , inflammatory dilated cardiomyopathy and heart failure were observed.³⁵ Significantly, the disease could be reversed by inactivating the transgene or in vivo expression of the I κ B- α superrepressor, suggesting that IKK-activated NF- κ B in cardiomyocytes was sufficient to cause cardiomyopathy and heart failure.³⁵ Overexpression of p65 or IKK- β -enhanced p22(phox) gene promoter activity and NF- κ B decoy oligodeoxynucleotides significantly downregulated messenger RNA and protein expression of p22(phox).²¹ NF- κ B inhibitors reduced the NADPH-dependent superoxide production, suggesting that the regulation of NADPH oxidase by NF- κ B is the likely mechanism whereby proinflammatory factors induce oxidative stress in the heart.²¹ These studies show a pathological role of p65 activation in the heart.

Role of Other NF- κ B Subunits

To understand the role of the c-Rel subunit in cardiac diseases, immunohistochemical studies were performed in control and diseased human hearts from patients with end-stage ischemic or idiopathic dilated cardiomyopathy.²⁸ c-Rel was primarily localized to

cardiomyocyte nuclei in diseased tissue compared to low levels and cytoplasmic presence in controls, suggesting a pathological role of this subunit. Studies in animal models showed that the hearts of c-Rel knockout mice were significantly smaller than those in WT animals. Further, AngII-induced hypertrophy and fibrosis were significantly attenuated in c-Rel-deficient mice, suggesting that c-Rel is involved in normal and pathological heart growth.²⁸ Cardiac-specific NEMO knockout mice developed progressive eccentric cardiac hypertrophy with extensive cardiac fibrosis.¹² TNF- α -induced NF- κ B activation was largely absent in these animals; however, increased expression of TNF- α and IL-6 was observed in NEMO knockout hearts. Increased expression of TNF- α and IL-6 was accompanied by oxidative stress and reduced expression of antioxidant genes SOD1 and SOD2. This study suggested a protective role of NEMO-mediated NF- κ B signaling in the heart.¹²

DIABETIC CARDIOMYOPATHY, INFLAMMATION, THE RAS, AND NF- κ B Increased Cytokine Production in a Diabetic Heart

Given that diabetes represents a condition of chronic inflammation, the role of NF- κ B signaling in diabetic cardiomyopathy is important. In db/db mice, no increase in cardiac cytokines was observed at 11 weeks of age (after about 4 weeks of diabetes); however, plasma levels of TNF- α were significantly elevated.³⁶ A similar increase in plasma TNF- α and IL-6 were observed after 20 weeks of diabetes in db/db mice.³⁷ In these animals, cardiac NF- κ B p65 activity and p50 gene expression were significantly elevated. A high-fat diet fed to mice for 6 weeks increased their plasma and cardiac IL-6 and plasma TNF- α level in association with reduced glucose uptake and metabolism in the heart.³⁸ Acute (4 hours) IL-6 infusion to mice in the same study reproduced a similar phenotype. Furthermore, diet-induced inflammation and defective cardiac glucose metabolism were prevented in IL-6 knockout mice.³⁸ Hyperglycemia exacerbates endotoxin-induced inflammation and advanced glycation endproduct (AGE)-induced oxidative stress.^{36,39} In streptozotocin-induced diabetes models, increased cardiac levels of TNF- α , IL-6, and IL-1 β were observed after 8 weeks of diabetes in mice; increased cardiac levels of TNF- α and IL-1 β , but not interferon-gamma, were observed in rats after 6 weeks.^{40,41} Treatment with a p38 MAPK (MAP kinase) inhibitor reduced cardiac inflammation and improved systolic dysfunction but did not improve diastolic function in the diabetic mice, suggesting that factors other than cytokines may contribute to cardiac stiffness. However, an ARB attenuated both systolic

and diastolic dysfunction by decreasing inflammation and normalizing matrix metalloproteinase activity in the heart.⁴² Pralnacasan, an interleukin-converting enzyme inhibitor that prevents IL-1 β activation, and a low dose of atorvastatin that reduced inflammation without affecting cholesterol levels attenuated cardiac dysfunction in the rat model.^{41,43} Similar beneficial effects of statins in diabetic nephropathy have been reported.⁴⁴ Another study using an anti-TNF- α antibody showed protection from cardiac dysfunction in diabetic rats.⁴⁵ From these studies, an inverse correlation between cytokine levels and cardiac dysfunction in the diabetic heart is clear. Because cytokines activate NF- κ B and NF- κ B increases expression of cytokines, in the following section we review studies that link hyperglycemia, cytokines, the RAS, and NF- κ B.

Activation of NF- κ B by Hyperglycemia in the Heart

Increased p65 NF- κ B activity was observed by EMSA in rat myocardium after 12 weeks of diabetes induced by streptozotocin.⁴⁶ The accumulation of AGEs has been implicated in diabetic cardiomyopathy. In a study using glycated bovine serum albumin, increased oxidative stress was observed in neonatal rat cardiomyocytes in a time- and concentration-dependent manner and was associated with nuclear translocation of p65 NF- κ B in a protein kinase C (PKC)-dependent manner.³⁹ The exposure of neonatal rat cardiomyocytes to 25 mM glucose for 48 hours resulted in increased levels of total and phospho-NF- κ B, both in the cell lysate and in the nucleus.⁴⁷ This result was accompanied by increased expression of TNF- α . The effects of glucose on NF- κ B and TNF- α could be inhibited by the PKC inhibitor Ro 31-8220, suggesting the involvement of the PKC/NF- κ B pathway. Similarly, the exposure of H9c2 cells to 33 mM glucose caused increased IKK, I κ B- α , and p65 phosphorylation and NF- κ B-responsive luciferase activity.⁴⁸

Activation of NF- κ B by Proinflammatory Cytokines in the Heart

TNF- α was not cytotoxic and did not provoke apoptosis in normal myocytes.⁴⁹ However, TNF- α caused a 2.2-fold increase in apoptosis in myocytes defective for NF- κ B activation.⁴⁹ In vitro studies in the human cardiac cell line AC16 and in vivo studies in cardiac-specific TNF- α transgenic mice showed that TNF- α activated NF- κ B signaling, as determined by decreased levels of I κ B- α , by increased nuclear levels of p65, and by EMSA.⁵⁰ NF- κ B activation resulted in decreased expression of peroxisome proliferator-activated receptor gamma coactivator 1-alpha (PGC-

1 α) and an increase in glucose oxidation rate, the increased glucose oxidation rate representing a likely mechanism in inflammatory cardiomyopathy.⁵⁰ In this regard, PGC-1 α was reported to be decreased significantly at 2, 8, and 20 weeks of diabetes in the hearts of a streptozotocin-induced diabetic mouse model.⁵¹ TNF- α expression was increased significantly in the hearts of diabetic mice compared to controls at 2 and 20 weeks but not at 8 weeks. To compare acute versus chronic stimulation with TNF- α of NF- κ B activity in the heart, mice acutely injected with TNF- α or cardiac-restricted TNF- α transgenic mice, respectively, were studied.⁵² Whereas both acute and chronic stimulation activated p50/p65 heterodimers, chronic TNF- α exposure additionally resulted in nuclear translocation of transcriptionally inactive p50 homodimers, suggesting that transcriptionally inactive p50 homodimers constituted an adaptive response to minimize the inflammatory consequences of chronic TNF- α stimulation.⁵² In vitro studies in H9c2 cells showed enhanced TNF- α expression in response to free fatty acids.⁵¹

Activation of NF- κ B by AngII in the Heart

Experimental and clinical studies have demonstrated a significant role of the RAS in diabetic cardiomyopathy. The following studies provided evidence that AngII activates NF- κ B in the heart, thereby suggesting NF- κ B as a central pathway in diabetic cardiomyopathy. In rat neonatal ventricular myocytes, AngII activated NF- κ B signaling as determined by nuclear localization and DNA binding assay of the p65 subunit.⁵³ A specific inhibitor of PKC prevented p65 translocation to the nucleus, suggesting involvement of this pathway. Similar activation of a PKC-dependent canonical NF- κ B pathway by AngII in a time- and concentration-dependent manner was reported in adult feline cardiomyocytes and isolated hearts.⁵⁴ Increased expression of TNF- α accompanied NF- κ B activation. In adult rat cardiomyocytes, both AngII and TNF- α induced NF- κ B DNA binding activity at 30 minutes of stimulation, although TNF- α was significantly more potent than AngII.¹⁰ Using supershift assay, the study determined that AngII-induced NF- κ B DNA binding activity largely consisted of p50. Interestingly, AngII did not activate NF- κ B in cardiac fibroblasts.¹⁰ In another study with isolated adult rat cardiomyocytes, AngII increased NF- κ B p50 expression in a dose-dependent manner when applied not only to intact cells but also to isolated nuclei.⁵⁵ AngII-induced p50 expression could be partially blocked by AT₁ and AT₂ antagonists and more completely by the combination of both. Several studies have described intracellular or intracrine actions of AngII.⁵⁶⁻⁵⁹ Cardiomyocyte-restricted ex-

pression of the NF- κ B superrepressor I κ B- α delta N (Δ N^{MHC}) in mice attenuated AngII-induced cardiac hypertrophy, suggesting a requirement of NF- κ B in AngII response.¹⁰ In the rat embryonic cardiomyocyte cell line H9c2, AngII induced NADPH oxidase-activated p50 binding to the cardiac SCN5A sodium channel promoter, resulting in decreased transcriptional activity.⁶⁰ Increased NF- κ B DNA binding was observed in the hearts of rats double transgenic for human renin and AGT; the binding was prevented by pyrrolidine dithiocarbamate treatment.⁶¹ Increased NF- κ B activity in SHR was prevented by captopril treatment, likely through inhibition of the RAS.²⁴ These studies showed that AngII activates NF- κ B signaling in cardiomyocytes.

CONCLUSIONS

In the heart, NF- κ B signaling has been extensively studied, yet we are far from a complete understanding of the role of this pathway in different cardiac pathological states. Studies using genetic models have provided significant information regarding the role of the individual components of the NF- κ B signaling machinery. However, because of the multiplicity of interactions of these components, a composite picture of the effects of the addition or deletion of individual components on the overall NF- κ B mechanism is lacking. Further, most studies investigating the activation of NF- κ B in the heart have examined only 1 or 2 mechanisms, providing an incomplete understanding of the pathways involved. The presentation of 1 or 2 mechanisms as representing all NF- κ B activity has likely led to confusion regarding protective or detrimental effects of this pathway in the heart. The studies we reviewed demonstrate that the effects of NF- κ B in the heart are mechanism specific. Additionally, NF- κ B signaling is cyclical in nature and autoregulated by continuous feedback mechanisms. Therefore, the timing of the NF- κ B measurement is critical, and studies determining temporal changes in NF- κ B mechanisms would provide useful information. In this review, we were careful to identify the specific pathways studied in the heart and hope that doing so can help clarify the multiple roles of NF- κ B in cardiac pathophysiology.

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