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Targeting GPCR-G $\beta\gamma$ -GRK2 signaling as a novel strategy for treating cardiorenal pathologies

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Abstract

The pathologic crosstalk between the heart and kidney is known as cardiorenal syndrome (CRS). While the specific mechanisms underlying this crosstalk remain poorly understood, CRS is associated with exacerbated dysfunction of either or both organs and reduced survival. Maladaptive fibrotic remodeling is a key component of both heart and kidney failure pathogenesis and progression.

G-protein coupled receptor (GPCR) signaling is a crucial regulator of cardiovascular and renal function. Chronic/pathologic GPCR signaling elicits the interaction of the G-protein G $\beta\gamma$ subunit with GPCR kinase 2 (GRK2), targeting the receptor for internalization, scaffolding to pathologic signals, and receptor degradation. Targeting this pathologic G $\beta\gamma$ -GRK2 interaction has been suggested as a possible strategy for the treatment of HF. In the current review, we discuss recent updates in understanding the role of GPCR-G $\beta\gamma$ -GRK2 signaling as a crucial mediator of maladaptive organ remodeling detected in HF and kidney dysfunction, with specific attention to small molecule-mediated inhibition of pathologic G $\beta\gamma$ -GRK2 interactions. Further, we explore the potential of GPCR-G $\beta\gamma$ -GRK2 signaling as a possible therapeutic target for cardiorenal pathologies.

Keywords

Heart Failure; Kidney injury; Cardiorenal syndrome; Fibrosis; Signal Transduction

1. Introduction

Cardiovascular diseases (CVD) involve heart and blood vessels and include coronary artery disease (CAD), stroke, hypertension, congenital heart disease, cardiomyopathy, etc. [1]. CVD is the leading cause of death worldwide that accounts for more than 17.3 million deaths per year [2]. In 2013, CVD represented about one of every three deaths in America. Over 85 million Americans are living with some type of CVD or the after-effects of stroke.

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Projected costs of CVD including the cost of health care services, medications and loss of productivity totals more than \$600 billion in 2015, and it is expected to grow to more than \$1200 billion by 2030 [3].

Heart failure (HF) is a chronic, progressive condition in which the heart muscle is unable to pump enough blood to meet the body's metabolic demands. HF arises as the final manifestation of many CVDs such as coronary artery disease, congenital malformations and hypertension. About 5.7 million adults in the United States are affected by this debilitating disease [3]; HF treatment costs the nation an estimated \$30.7 billion each year [4]. Notwithstanding significant advances in HF treatment and management realized with β -adrenergic receptor (β -AR) blockers, angiotensin receptor blockers, angiotensin converting enzyme (ACE) inhibitors, aldosterone inhibitors, and diuretics, conventional pharmacological therapies only impede the progression and death due to HF, but do not cure it causatively [5]. Taking into consideration the steady growth of aging and diabetic populations, deeper understanding of the molecular and cellular processes that contribute to the disease pathogenesis, along with development of innovative therapeutic strategies allowing the causative cure of HF, are indispensable.

Multiple pathophysiological mechanisms contribute to HF development and progression, including neurohumoral activation [6], G-protein-coupled receptor desensitization and down-regulation [7], [8] and [9], and extracellular matrix (ECM) mediated pathologic remodeling [10]. Moreover, cardiac pathologies in HF are frequently accompanied by worsening renal function, which is known to be a strong predictor of increased mortality in HF patients [11] and [12]; this is defined as Cardiorenal syndrome (CRS) type II. In the present review, we discuss recent advances in exploring GPCR signaling as a possible therapeutic target in cardiac disease and as a potential link between failing heart and kidney, with the particular emphasis on small molecule targeting of G-protein $\beta\gamma$ subunit - GPCR kinase 2 (G $\beta\gamma$ -GRK2) components of GPCR signaling.

2. G-protein-coupled receptor signaling

G-protein-coupled receptors (GPCRs), also known as seven-transmembrane domain receptors, represent a conserved family of receptors that sense molecules outside the cell that activate intracellular signal transduction pathways and consecutive cellular responses. GPCRs are integral proteins comprised of an extracellular N-terminus, seven transmembrane (7-TM) α -helixes (TM-1 to TM-7) connected by three intracellular (IL-1 to IL-3) and three extracellular loops (EL-1 to EL-3), and an intracellular C-terminus [13]. Ligand binding to an extracellular active site of the receptor induces a conformational change in the GPCR which allows for coupling with heterotrimeric guanine-nucleotide regulatory proteins (G-proteins) [14]. G-proteins are heterotrimers of α , β and γ subunits known as G α , G β and G γ , respectively. The heterotrimeric G-proteins are rendered inactive when reversibly bound to Guanosine diphosphate (GDP) but active when bound to Guanosine triphosphate (GTP) [15]. Receptor activation facilitates the exchange of GDP for GTP on G α subunits that result in dissociation of the G α from G $\beta\gamma$ subunits to mediate downstream signaling pathways [16]. Dissociated G α subunits signal via activation of an effector molecule, such as adenyl cyclase (AC) or phospholipase C β (PLC β) to produce second messengers such as cyclic

adenosine 3['], 5['] monophosphate (cAMP), diacylglycerol (DAG), or inositol 1, 4, 5triphosphate (IP3), respectively. These second messengers modulate a variety of downstream processes, particularly regulation of contractility, hypertrophy, and apoptosis in the heart [15]. Ga proteins are classified into the families Ga_s, Ga_i, Ga_q, and Ga_{11/12} [15] with respect to downstream signaling molecules and modulated physiological processes. Dissociated G $\beta\gamma$ subunits target a wide range of signaling pathways involved in receptor desensitization and down-regulation, ion channel activation, enzyme activity modulation, cell division, transcription and cellular organelle function [17], [18], [19] and [20].

GPCRs respond to extracellular signaling mediated by an extensive amount of agonists such as hormones, proteins and lipids, and participate in a comprehensive variety of physiological processes [21]. In particular, GPCRs play an important role in local and systemic regulation of cardiac function. Specifically, cardiac β -adrenergic receptors (β -ARs) are prominent regulators of cardiovascular chronotropy and inotropy [22] and [23]. Furthermore, GPCRs mediate a variety of functions in the kidney, and inappropriate activation and regulation of GPCRs may lead to kidney disease [24]. In this review, we focus on $G\beta\gamma$ -mediated signaling as a crucial component of HF pathogenesis and as a potential therapeutic target in cardiorenal pathologies.

3. β-Adrenergic receptor signaling in healthy and diseased heart

As mentioned above, cardiac β -ARs represent crucial regulators of cardiac contractile function. In response to sympathetic nervous system (SNS) activity released via mediators, catecholamines (CA) epinephrine (Epi, also named adrenaline) and norepinephrine (NEpi, also named noradrenaline), β -ARs modulate the rate and force of myocardial contractions [8]. There are three β -AR subtypes identified in mammalian hearts: β 1, β 2, and β 3-ARs [25]. Both β 1- and β 2-ARs are coupled to the downstream excitatory G α_s protein, which generally results in the activation of adenylyl cyclase (AC) and the generation of cyclic AMP (cAMP), eliciting positive chronotropic and inotropic responses. Upon chronic stimulation, β 2-ARs also couple to the inhibitory G α_i protein, which has been reported to exert a cardioprotective effect during cardiac injury [26].

In healthy human myocardium, the predominant β -ARs subtypes are the β 1- and β 2-ARs, which are present in an approximate 80:20 ratio, respectively with only a relatively minor contribution of β 3-ARs [27]. Under physiological conditions, β -ARs account for regulation of both heart rate and contractility [14, 28]. In HF pathogenesis, excess SNS activation and subsequent catecholamine overdrive is initiated as an adaptation to compensate for decreased heart rate and cardiac contractility and to maintain mean arterial pressure (MAP) [29]. Initially, the elevated SNS activity increases heart rate and contractility through β -AR stimulation. However, maladaptive effects of the elevated SNS activity including myocardial ischemia, pathologic hypertrophy, arrhythmogenicity, myocardial necrosis and apoptosis contribute substantially to disease progression [22], [30], [31] and [32]. This maladaptive response results partially from down-regulation and desensitization of cardiac β -ARs due to chronic CA stimulation [15]. In failing hearts, heightened CA β -AR stimulation induces selective down-regulation of β 1-ARs and consequent alteration of the β 1-AR to β 2-AR ratio

from an 80:20 distribution to a ratio of 60:40 [27, 33]; the remaining β 1-ARs and β 2-ARs in failing hearts prevail in a desensitized condition [30].

Cardiac β -AR signaling regulation involves activation-dependent and -independent mechanisms of desensitization [8]. Homologous, agonist-mediated, activation-dependent desensitization is accomplished by an active form of a G-protein-coupled receptor kinase (GRK) that is translocated to the adrenergic receptor after binding with the activated membrane-associated G $\beta\gamma$ subunit to phosphorylate the agonist-occupied receptor [34]. An alternative, activation-independent pathway, known as heterologous desensitization, is accomplished through the activity of a downstream signaling product of β -AR activation or other GPCR signaling events. In both cases, phosphorylated β -AR is bound by β -arrestin molecules which block the access of heterotrimeric G proteins to the receptor thereby uncoupling it and attenuating β AR signaling in the heart [35, 36].

4. Gβγ-GRK2 signaling manipulation as a strategy to treat cardiac disease

4.1. GRK2: structure, subcellular localization and function in the heart

GRK2 (aka β-adrenergic receptor kinase, βARK) belongs to a family of serine/threonine kinases that share common structural and functional features. Seven mammalian GRKs that have been characterized so far are classified into three subfamilies according to their sequence and structural similarity: (1) the rhodopsin kinase subfamily (GRK1 and GRK7); (2) the βARK subfamily (GRK2 and GRK3); and (3) the GRK4-like subfamily (GRK4, GRK5, GRK6) [37]. Within the cardiovascular system, GRKs 2, 3 and 5 are known to be expressed and play a role in GPCR phosphorylation [38], with GRK2 as a predominant GRK isoform in the heart [39].

GRKs are characterized by a tri-domain structure, with the conserved central catalytic domain and two flanking domains variable in structure in different GRK subfamilies [40]. GRK2's amino (N)-terminal domain that is responsible for receptor recognition and activity regulation contains a regulator of G protein signaling (RGS) homology (RH) domain that has been demonstrated to interact with $G\alpha_q$ proteins [41]. The carboxyl (C)-terminal domain of GRK2 determines membrane targeting and subcellular localization of the enzyme. This domain contains a pleckstrin homology (PH) domain that binds G $\beta\gamma$ subunits [42]. Under basal conditions, GRK2 is distributed primarily in the cytoplasm. Upon GPCR activation, GRK2 is translocated to the plasma membrane via binding with the activated G $\beta\gamma$ subunits. GRK2-mediated phosphorylation of the GPCR causes β -arrestin recruitment to the receptor and consequent inhibition of dissociated G-proteins from coupling to the receptor/ β -arrestin complex and further attenuation of downstream signaling [43]. Moreover, β -arrestin-bound receptors are targeted for clathrin-coated pits in the cell membrane that are internalized and either degraded in intracellular lysosomes or recycled back to the cell surface [44].

Apart from the classical mechanism of modulating GPCR signaling in the heart and extracardiac tissues, GRK2 may have other functions independent of GPCR phosphorylation. Recently emerging data suggest the concept of an extensive "GRK2 interactome" that refers to GRK2 interactions with other intracellular proteins such as actinin, clathrin, calmodulin, caveolin, tubulin, Akt, HDAC6 and ERK1/2 [39] and [45].

Investigation of GRK2 functions beyond GPCR desensitization and down-regulation may provide new insights in understanding its role in disease pathogenesis. In the current review, we highlight recent updates relevant for GPCR- $G\beta\gamma$ signaling in HF modulation.

Understanding of the *in vivo* function of GRK2, particularly its role in cardiovascular system function and development, emerged from gene knockout studies. In 1996, Jaber *et al* demonstrated lethality of GRK2 homozygous knockout (KO) in mouse models [46]. These animals exhibited hypoplasia of the ventricular myocardium and a 70% decrease in ejection fraction and died by gestational day 15.5, presumably owing to HF. Further studies demonstrated that specific deletion of GRK2 in murine embryonic cardiomyocytes utilizing Cre recombinase expressed under the control of the Nkx2.5 promoter did not cause any apparent developmental abnormalities, suggesting that embryonic lethality of GRK2^{-/-} mice might result from extracardiac or non-cardiomyocyte effects [47].

Cardiomyocytes from adult global heterozygous GRK2 KO mice exhibited significantly enhanced cardiac contractile function compared to wild-type cells [48]. Cardiac-specific overexpression of GRK2 following myocardial ischemia/reperfusion injury (I/R) [49] caused reduced β-adrenergic signaling mediated cardiac contractility and function along with increased apoptosis [50]. These observations demonstrated that cardiac contractile function can be modulated by the level of GRK2 activity. To further evaluate the role of GRK2 in adult cardiac function, two conditional models of GRK2 ablation were generated: aMHC-Cre/GRK2^{flox/flox} for targeted KO of GRK2 specifically in cardiomyocytes in the constitutive way (at birth) and aMHCMerCreMer/GRK2flox/flox for tamoxifen-induced cardic deletion [51]. Both models resulted in positive outcomes following cardiomyocyte-restricted GRK2 ablation after myocardial infarction (MI); the aMHC-Cre/GRK2^{flox/flox} mice exhibited prevention of HF development post-MI and the aMHCMerCreMer/GRK2flox/flox mice demonstrated improved cardiac function and induced positive reverse remodeling following MI. Moreover, cardiomyocyte GRK2 KO in post-MI mice showed reduced mortality levels. Cardioprotective effects demonstrated by aMHCMerCreMer/GRK2flox/flox mice were significantly better compared to the results when the β -blocker metoprolol was used for treatment of post-MI wild-type (WT) mice over the same time period [51].

A recent study conducted by Woodall and colleagues aimed to determine the implications of GRK2 deletion in cardiac fibroblasts (CFs) prior to myocardial injury [52]. CFs significantly contribute to multiple aspects of cardiac function and disease [53], particularly via modulation of fibrotic remodeling in the heart following injury [10], [54] and [55]. To examine the consequences of GRK2 loss in CFs for cardiac function, authors used tamoxifen-inducible collagen1a2-CreER(T)/GRK2^{fl/fl} mice (GRK2 fKO mice). GRK2 ablation was achieved weeks prior to *in vivo* acute myocardial ischemia/reperfusion (I/R) injury [56]. GRK2 fKO mice subjected to myocardial I/R injury exhibited advantageous effects of pre-injury GRK2 ablation in CFs including decreased infarct size, fibrosis and apoptosis, reduced neutrophil extravasation and tumor necrosis factor a. (TNFa.) expression and secretion along with restored cardiac function. Collectively, GRK2 gene knockout studies highlighted the beneficial effects of GRK2 ablation in cardiomyocytes and CFs after myocardial injury and elicited a proposal that GRK2 should be considered as a therapeutic target for HF treatment.

4.2. GRK2 expression in cardiac disease

The first link between GRK2 and β -adrenergic receptor signaling desensitization and downregulation was established in 1993, when Ungerer et al demonstrated significant elevation of GRK2 in explanted failing human hearts at mRNA, protein and activity levels [33]. These observations initiated a series of studies performed on animal models or human tissue that aimed to delineate the role of GRK2 in cardiac disease [57]. Particularly, cardiac overexpression of GRK2 was demonstrated to be capable of direct HF induction in experimental animal models; moreover, mice overexpressing GRK2 exhibited decreased isoproterenol (Iso)-stimulated left ventricular contractility in vivo, diminished myocardial AC activity, and reduced functional coupling of β-ARs [58]. Furthermore, GRK2 expression and activity were found to be elevated in human cardiac tissue and in circulating lymphocytes, demonstrating direct correlation with the severity of HF [59] and [60]. More recent studies showed that the changes of GRK2 levels in peripheral lymphocytes mirror changes in the salutary LVAD-supported failing human heart and that these changes correlate strongly with cardiac function such that lower levels of GRK2 are associated with improved β-adrenergic signaling and myocardial function in mechanically supported failing hearts and transplanted human hearts [61] and [62]. The recent study conducted by Rengo and colleagues involved over 200 patients with HF and demonstrated that GRK2 lymphocyte level has a prognostic value for outcomes and mortality in HF patients, thus supporting the hypothesis that GRK2 levels in blood can be used as a biomarker in HF [63].

Overall, the aforementioned studies suggest consideration of GRK2 as a therapeutic target for cardiac disease and as a potential biomarker for heart function [64], [65] and [66].

4.3. Recombinant proteins as a way to inhibit Gβγ-GRK2 signaling

Excess cardiac G $\beta\gamma$ -mediated signaling leading to chronic β -AR desensitization and downregulation is a crucial component of HF pathophysiology [67]. Thus, several approaches, including genetic manipulations and pharmacological targeting, have been explored to interdict pathologic G $\beta\gamma$ -GRK2 signaling.

The first reported approach utilized a recombinant carboxyl (C)-terminal fragment of GRK2 comprised of 194 amino acids encoding the G $\beta\gamma$ binding domain (β ARKct) as a presumed inhibitor of G $\beta\gamma$ -GRK2 interactions. In 1994, β ARKct was expressed in COS-7 cells where it attenuated G $\beta\gamma$ mediated signaling with unaffected Ga mediated signaling, indicating its ability to discriminate between Ga and G $\beta\gamma$ pathways [68]. Subsequently, in 1995 Koch *et al* demonstrated enhanced baseline cardiac contractility *in vivo* with or without Iso stimulation in transgenic mice with cardiac-specific GRK2 overexpression [58]. Additionally, β ARKct was demonstrated to normalize β -adrenergic signaling and cardiac function in hybrid transgenic mice with cardiac-specific concomitant overexpression of both GRK2 and β ARKct [69].

Further studies demonstrated salutary effects of β ARKct on the recovery of failing myocytes function [70], [71] and [72] and prevention of cardiac dysfunction [73]. Oligonucleotide microarray left ventricular (LV) gene expression analysis performed in normal, failing and β ARKct overexpressing ("rescued") cardiac samples revealed the ability of β ARKct to

normalize gene expression changes associated with HF [74]. More recent studies performed in large animal HF models revealed preservation and amelioration of cardiac function along with normalization of CA signaling owing to stable myocardial β ARKct gene delivery [75] and [76], suggesting that inhibition of G $\beta\gamma$ -GRK2 interactions with recombinant viraldelivered β ARKct peptide is a promising therapy for HF treatment [66].

4.4. Small molecule interdiction of Gβγ-GRK2 signaling

 β ARKct inhibition of G $\beta\gamma$ -GRK2 interactions has demonstrated salutary effects on cardiac function in both acute and chronic models of HF, however, viral-based gene delivery remains a daunting therapeutic approach. In that perspective, small molecule inhibitors that could be administered systemically may represent an alternative approach to attenuate pathologic components of G $\beta\gamma$ -GRK2 signaling or interactions [64] and [66].

One of the described approaches to small molecule pharmacological inhibition of GRK2 signaling is paroxetine, the selective serotonin reuptake inhibitor, identified by Thal *et al* in 2012 [77]. This antidepressant drug binds to the active site of GRK2 and stabilizes the kinase domain, thereby inhibiting the downstream signaling. This study demonstrated increased contractility in isolated cardiomyocytes in the presence of paroxetine. Further, paroxetine was tested *in vivo* in a mouse model of MI. Schumacher *et al* recently showed that paroxetine treatment initiated two weeks post-MI results in improved cardiac function, limited ventricular remodeling and normalized SNS overdrive along with myocardial β -adrenergic system [78]. Thus, direct GRK2 pharmacological inhibition demonstrated salutary effects on HF progression.

Another potential strategy to interdict $G\beta\gamma$ -GRK2 pathologic signaling is targeting $G\beta\gamma$ subunit and inhibiting its protein-protein interactions [5] and [66]. Hence, Bonacci *et al* in 2006 performed a virtual screening of 1990 compounds from the National Cancer Institute (NCI) chemical library to identify small molecules capable of binding $G\beta\gamma$ protein interaction domain mentioned above [79]. Eighty-five identified compounds were further tested in an enzyme-linked immunosorbent assay (ELISA) for their ability to compete with a phage-displaying SIRK peptide derivative (SIGK) [80] for binding to $G\beta\gamma$ subunit. Among several tested compounds, one termed M119 (cyclohexanecarboxylic acid [2-(4,5,6-trihydroxy-3-oxo-3*H*-xanthen-9-yl)-(9CI)]) demonstrated high apparent affinity for the $G\beta\gamma$ subunit and inhibited $G\beta\gamma$ -SIGK binding *in vitro*. Moreover, pretreatment of differentiated HL-60 leukocytes with M119 resulted in interference with $G\beta\gamma$ binding to GRK2 and consequent inhibition of GRK2 translocation to the membrane, along with suppression of PLC $\beta_{2/3}$ and PI3K γ activation by $G\beta\gamma$. Thus, the small molecule M119 confirmed its ability to interfere with $G\beta\gamma$ -mediated signaling downstream of GPCRs.

Since $G\beta\gamma$ subunits are known to modulate a majority of signaling pathways, the inhibitor that selectively influences a particular subset of $G\beta\gamma$ interactions is needed for $G\beta\gamma$ -GRK2 targeting [28]. Thus, M119 was examined *in vivo* for efficacy and specificity. It has been demonstrated that inhibition of PLC β 3 that is activated by $G\beta\gamma$ subunits is associated with enhanced morphine-induced antinociception [81]. Co-administration of M119 with μ -opioid receptor agonist morphine resulted in substantial increase of morphine-dependent antinociception in wild-type mice due to M119-induced inhibition of $G\beta\gamma$ -PLC β 3

interactions, whereas M119 alone had no effects on antinociception [79]. Also, M119 demonstrated no effects on morphine-dependent antinociception in PLC β 3^{-/-} mice. Further studies showed that M119 increases analgetic potencies of morphine or μ -selective peptide, whereas it does not have any significant influence on analgesia induced by κ - or δ -opioid receptor agonists [82], corroborating the suggestion that M119 acts as a specific inhibitor of a particular subset of G $\beta\gamma$ -mediated signaling. Accordingly, we have conducted various studies to evaluate the potential of small molecule inhibition of G $\beta\gamma$ -GRK2 associations in different animal HF models.

4.5. Gβγ inhibitory treatment in acute and advanced heart failure models

Taking into account the aforementioned role of β -AR-dependent G $\beta\gamma$ -GRK2 signaling in cardiac disease pathogenesis and the proven efficiency of small molecule inhibitor M119 in selective interdiction of G $\beta\gamma$ -GRK2 interactions, we explored the effects of G $\beta\gamma$ signaling small molecule disruption in myocardial cells and in murine models of HF [83]. In this study, M119 or its highly homologous, more chemically stable analogue gallein was utilized to inhibit G $\beta\gamma$ -GRK2 interactions. M119 pretreatment followed by administration of the β -AR agonist Iso significantly enhanced AC activity and consequent cAMP generation in isolated cardiomyocytes from adult wild-type mice. Moreover, M119 increased the rate of cardiomyocyte contraction alone and in combination with Iso. Importantly, the β -AR antagonist propranolol abolished the effect of M119 and Iso on cardiomyocyte contractility, confirming the selectivity of the compound for β -AR-G $\beta\gamma$ signaling. In addition, both M119 and gallein demonstrated the ability to reduce GRK2 recruitment to the membrane of cardiomyocytes induced by Iso treatment.

To examine cardiac-specific effects of $G\beta\gamma$ inhibitory treatment initiated at the onset of HF, an acute pharmacologic murine model of HF [84] was implemented. Chronic β-AR stimulation by Iso delivered via implantable miniosmotic pumps was started simultaneously with systemic administration of M119 or vehicle and continued for 7 days. M119 treatment mitigated HF progression; particularly, M119-treated mice maintained essentially normal cardiac function and showed significantly reduced cardiac hypertrophy along with decreased level of interstitial and perivascular fibrosis, compared to vehicle-treated animals. Considering these data, $G\beta\gamma$ small molecule inhibition was administered after the onset of HF, in a transgenic mouse model of established HF generated by cardiac restricted calsequestrin (CSQ) overexpression [85]. Importantly, the CSQ transgenic mouse model recapitulates essential hallmarks of HF, including pathologic β -AR signaling [83]. One month of daily gallein administration resulted in prevention of HF progression, especially in normalized echocardiographic parameters, reduced pathologic cardiac hypertrophy and diminished expression of HF molecular markers. Moreover, M119 and gallein significantly reduced pathologically increased cardiac GRK2 protein level in Iso-pumped and CSQ animals, respectively. Overall, this study demonstrated salutary effects of $G\beta\gamma$ small molecule inhibitory treatment on both manifestation and progression of HF.

Beneficial effects of small molecular inhibitors observed in acute pharmacological and transgenic mouse models of HF elicited further interest to investigation of cardiac and systemic effects of $G\beta\gamma$ signaling inhibition. Consequently, Kamal and colleagues in 2014

examined outcomes of utilizing the small molecule $G\beta\gamma$ inhibitor gallein in a transverse aortic constriction (TAC) mouse model of pressure-overload induced cardiac hypertrophy and HF [86]. The TAC surgical model, firstly validated by Rockman et al [87], is considered a relatively clinically relevant model of HF [88] and [89]. In this study, vehicle-treated mice developed the decline in cardiac function at 8 weeks post-TAC with the concomitant worsening at 12 weeks post-TAC. Daily gallein administration for eight weeks was initiated after the establishment of HF (four weeks post-TAC). This treatment regimen, initiated after the onset of HF, alleviated cardiac dysfunction and hypertrophy along with significantly enhanced survival in the gallein-treated group compared to the vehicle-treated group. Preservation of cardiac function was accompanied by the recovery of β -AR density and reduction of GRK2 gene expression and membrane translocation. Furthermore, membrane recruitment of phosphoinositide 3-kinase γ (PI3K γ), which GPCR-induced activation was implicated in maladaptive cardiac hypertrophy and dysregulated β -AR function [28], [90], [91], [92] and [93], was reduced in gallein-treated mice compared with vehicle-treated mice. Gallein treatment also resulted in attenuated progression of cardiac hypertrophy and reduced myocardial fibrosis. Interestingly, ameliorated cardiac remodeling was accompanied by decreased phosphorylation of cardiac Akt (aka protein kinase B) and its downstream signal GSK-3 β . As mentioned above, PI3K γ -mediated GPCR dependent Akt activation and subsequent GSK-3ß Ser-9 phosphorylation lead to cardiac hypertrophy [92], [94] and [95]. Of note, authors attributed the significantly reduced expression of the fetal genes atrial natriuretic peptide (ANP) and brain natriuretic peptide (BNP) detected in gallein-treated TAC mice to elevated levels of nonphosphorylated GSK-3 β that was suggested to negatively regulate transcription and protein translation of the hypertrophic genes [96]. Other detected beneficial effects of gallein treatment on TAC HF mice were attenuated cardiac inflammatory cytokine expression and reduced myocardial apoptosis. Collectively, these data corroborate the suggestion that $G\beta\gamma$ inhibitory treatment preserves cardiac function and halts HF progression in different small animal models of HF, in part via downstream inhibitory effects on cardiac fibrosis, hypertrophic gene expression and inflammation along with promotion of cell survival.

As discussed previously, compensatory sympathetic nervous system activation and subsequent systemic release of catecholamines (CA) from the adrenal gland medullary chromaffin cells increase the rate and the intensity of cardiac contractions in response to diminished cardiac output of the failing heart [97]. Adrenal chromaffin cell α 2-ARs that belong to GPCR family are essential regulators of sympathetic outflow in HF, providing the feedback inhibition for CA release [32] and [98]. Lymperopoulos and colleagues showed that, similar to the dysregulation of β -AR signaling in failing hearts, the adrenal α 2-ARs undergo desensitization and down-regulation in response to catecholamine overdrive and concomitant adrenal GRK2 upregulation, thus contributing to HF pathogenesis [99]. Further studies revealed that delivery of adenoviral vectors containing GRK2 to adrenal glands resulted in enhancement of plasma CA levels and failure of adrenal α 2-ARs to inhibit CA secretion, whereas GRK2 inhibition using β ARKct or chromaffin cell specific GRK2 gene deletion recovers adrenal α 2-AR function, reduces CA release and attenuates HF progression [31], [99] and [100]. More recently, Jafferjee et al demonstrated that CA treatment of rat pheochromocytoma-derived or primary chromaffin cells results in GRK2

gene transcription upregulation and subsequent enhancement of a2-AR desensitization and down-regulation accompanied by elevated CA biosynthesis and release [101].

Taking into account the aforementioned role of adrenal a2-ARs in regulation of SNS activity in HF, we recently assessed effects of small molecule $G\beta\gamma$ signaling disruption on the adrenal gland in pathogenesis of pressure-overload induced HF [86]. Significantly decreased CA production and release, alleviated adrenal medulla hypertrophy and restored a2-AR feedback inhibition was observed 12 weeks post-TAC in the gallein-treated group compared to the vehicle-treated group. Additionally, cultured in vitro adrenal glands from gallein-treated mice exhibited significantly decreased levels of basal CA secretion. Moreover, the effects of gallein on CA generation and GRK2 expression were examined in cultured human pheochromocytoma tissue, a tumor characterized by increased CA production. Gallein treatment significantly reduced CA production in cultured pheochromocytoma slices as reflected by lowered expression of tyrosine hydroxylase, an enzyme that catalyzes the rate limiting step in CA synthesis [102], and chromogranin A, a neurokine that is synthesized and co-secreted with vesicular CA [103]. Importantly, authors observed downregulation of both GRK2 protein level and membrane translocation in cultured pheochromocytoma slices treated with gallein. Taken together, this study provides deeper insights into understanding of pathological mechanisms contributing to HF progression. In addition, the study suggested small molecule GBy inhibition as a potential systemic therapy that attenuates HF progression due to simultaneous inhibition of cardiac and adrenal G\u03b3\u03c7-GRK2 interactions [86] and [64] (see overview of approaches in Figure 1).

In view of the correlation between the elevated GRK2 activity in circulating lymphocytes and the severity of HF demonstrated in the recent studies [61], [62] and [63], further investigation will be needed to determine potential effects of $G\beta\gamma$ -GRK2 inhibitory treatment on GPCR-Gβγ-GRK2 signaling in circulating immune system cells. As Lehmann and colleagues showed, small molecule inhibition of $G\beta\gamma$ -dependent signaling with M119 and gallein blocks chemotaxis and neutrophil migration in vitro and suppresses neutrophildependent inflammation in a murine carrageenan-induced paw edema model [104]. Moreover, recent data demonstrated that GRK2 modulates T- and B-lymphocytes migration from circulatory fluids into lymphoid tissues and within the spleen via desensitization of sphingosine-1-phosphate receptor-1 (S1PR1) that is necessary for sphingosine-1-phosphate (S1P) gradient dependent movement of lymphocytes [105] and [106]. To address the role of GRK2 in immune cell migration, the authors utilized CD4-Cre/GRK2^{f/-} and Mb1-Cre/ GRK2^{f/-} mice for GRK2 ablation in T and B cell populations, respectively. GRK2-deficient T and B cells displayed resistance to S1PR1 ligand-mediated desensitization and impaired ability to enter lymph nodes compared to the control WT cells. Further studies will be required to define the influence of Gby-GRK2 signaling small molecule disruption on circulating immune cell behavior as well as signaling pathways affected by GBy-GRK2 inhibition in immune cells.

As recently reviewed, GRK2 has been found to possess non-canonical activities including regulation of molecular pathways essential for cardiac physiology and metabolism such as the insulin signaling [107] and [108]. Particularly, Lucas et al demonstrated the interconnection between GRK2 dosage and cardiac insulin sensitivity [109]. Authors

characterized insulin signaling pathway, cardiac hypertrophy and gene expression patterns in hemizygous-GRK2 (GRK2^{+/-}) mice compared to wild-type control animals. GRK2^{+/-} mice revealed improved cardiac insulin sensitivity, non-pathologic hypertrophy and significant upregulation of genes known to play a protective role in CVDs along with decreased expression of genes associated with pathological cardiac hypertrophy and devastating diseases such as diabetes and obesity. Considering the role of GRK2 in systemic insulin resistance and obesity [110], authors further investigated whether GRK2 is upregulated in cardiac tissue in adult obese or high fat diet (HFD) fed mice and found significantly increased GRK2 protein levels in both conditions. Recent research demonstrated the role of GRK2 in obesity-related cardiac remodeling and lipid accumulation [111]. Moreover, genetic ablation of GRK2 resulted in reversed glucose tolerance and global insulin sensitivity, prevention of further body weight gain, increased fatty acid metabolism and attenuated lipid accumulation and inflammation in the liver in HFD-induced mouse model of obesity and insulin resistance [112]. Importantly, recent studies have also demonstrated a key role for GRK2 in mitochondrial metabolism [113]. Overall, aforementioned studies provided a new insight into the molecular mechanisms of worsening cardiac function in CVD comorbidities and highlighted the role of GRK2 signaling in these pathologic processes. In that perspective, small molecule inhibition of GPCR-G $\beta\gamma$ -GRK2 signaling might represent a promising approach for treatment CVD co-morbidities.

5. GPCR-Gβγ-GRK2 signaling in cardiorenal pathologies

The kidney performs essential regulatory roles in the body, including waste excretion, homeostasis maintenance, fluid volume and blood pressure regulation, as well as hormone secretion. GPCRs are widely expressed in the kidney, exemplified by arginine vasopressin receptor (AVP), dopamine-1 receptor (D1-R), angiotensin II receptors and endothelin (ET)-1 receptors [24]. GPCRs are involved in regulation of numerous kidney functions including water and electrolyte transport in renal tubules, maintenance of acid-base balance and renal blood flow and filtration [24] and [65]. Dysregulation of GPCR signaling is associated with severe kidney and systemic disorders such as renal fibrosis [114], [115] and [116], acute kidney injury (AKI) [117] and [118], hypertension [119], [120] and [121], and chronic kidney disease (CKD) [122].

Combined heart and kidney disorders, characterized by pathological interactions ("crosstalk") between affected organs, are defined as cardiorenal syndrome (CRS) [123]. Different approaches have been applied to characterization and classification of CRS [124]; according to a classification proposed by Ronco *et al* in 2008, CRS is discriminated into five types with respect to the acute or chronic pathogenesis and the initiating event [123]. While the specific mechanisms behind this pathologic crosstalk between heart and kidney remain poorly understood, CRS is associated with exacerbated dysfunction of either or both organs and reduced survival [125] and [126]. Essentially, kidney maladaptive remodeling and impaired function serve as a strong predictor of mortality in HF patients [11] and [12]. Thus, investigation of mechanisms of pathologic crosstalk between failing heart and kidney may contribute to development of novel therapeutic strategies for HF, kidney dysfunction and CRS.

Considering the role of GPCR signaling in normal physiology and pathology of both heart and kidney, we recently scrutinized the role of GPCR-G $\beta\gamma$ -GRK2 in CRS type 2 (CRS2), which is characterized as a chronic heart failure (CHF) accompanied by the development of CKD [123] and [127]. The study proposed that elevated activity of SNS and endothelin (ET) system causes desensitization and down-regulation of renal GPCRs owing to pathologic $G\beta\gamma$ -GRK2 interactions, resembling the dysregulation of β -ARs observed in HF. To recapitulate the clinical features of CRS2 progression, a non-ischemic TAC mouse model of pressure-overload induced HF was utilized. To determine the role of $G\beta\gamma$ -GRK2 signaling in kidney dysfunction besides the crosstalk with the heart, a direct bilateral ischemia reperfusion (I/R) acute kidney injury (AKI) model was also implemented [128]. Development of CKD secondary to TAC was reflected by elevated serum creatinine levels, emerged morphological and molecular signs of tubular damage and increased focal tubulointerstitial and perivascular fibrosis in the kidneys at 12 weeks post-TAC. Development of CKD in the chronic phase of HF is consistent with clinically observed consequence of CRS2 progression. Importantly, observed maladaptive changes were accompanied by elevated levels of ET-1 along with increased protein expression and membrane localization of ET receptors (ET_A and ET_B), that corroborates the proposed role of ET system in CRS. Furthermore, authors detected the elevation of $G\beta\gamma$ -GRK2 signaling in kidneys at 12 weeks post-TAC; essentially, upregulation of both ET and $G\beta\gamma$ -GRK2 signaling was attenuated by small molecule $G\beta\gamma$ inhibitor gallein treatment. Gallein pretreatment of mice subjected to AKI revealed protective effects of small molecule $G\beta\gamma$ -GRK2 inhibition on kidney function. Importantly, GRK2, ET-1 and ET_A gene expression was elevated in kidneys of both CHF and AKI I/R mice.

Overall, these data suggest the role of $G\beta\gamma$ -GRK2 interactions in both acute and chronic kidney injury and the potential mechanism underlying pathologic crosstalk between heart and kidney in CRS2. Moreover, the study provides mechanistic insight into fibrotic tissue remodeling, demonstrating the role of $G\beta\gamma$ signaling in mouse embryonic fibroblasts (MEFs) endothelin-1 induced activation and migration [127] and [129].

A study performed by White *et al* aimed to explore the role of $G\beta\gamma$ subunits in kidney remodeling after AKI I/R injury [130]. In this study, rats were treated with supraphysiologic doses of gallein (30 mg/kg or 100 mg/kg) daily for three days post-I/R I/R with minimal benefit [131]. We recently reported a gallein dose-response study in which maximal efficacy and minimal toxicity were observed at 10 mg/kg/d [86], with high-dose toxicity possibly attributable to effects on cell cycle progression [45] and [132] and cell division [19] and [20]. Further, the different treatment schemes between our studies may produce divergent outcomes. As described above, our study with a much lower (i.e. physiologic) dose of gallein pretreatment was indeed renoprotective [128].

Taking into consideration the bidirectional nature of the crosstalk between heart and kidney, Polhemus and colleagues recently examined whether catheter-based renal denervation (RDN) that is thought to reduce blood pressure owing to the disruption of sympathetic signaling [133] and [134] possesses cardioprotective effects [135]. To model the heart injury, authors subjected spontaneously hypertensive rats (SHR) representing a model of established hypertension to myocardial I/R at 4 weeks after either bilateral radiofrequency-

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RDN (RF-RDN) or Sham-RDN. Rats treated with RF-RDN exhibited a significant reduction in myocardial infarct size and substantially improved left ventricular function 24 hours following the I/R. Moreover, RF-RDN treated rats demonstrated the attenuation of myocardial oxidative stress and elevation of cytoprotective nitric oxide (NO) signaling [136] compared to the Sham-RDN treated group. Importantly, RF-RDN treatment caused the reduction in myocardial GRK2 gene expression level, particularly the decrease in GRK2 Ser670 phosphorylated protein level.

Phosphorylation at residue Ser670 of GRK2 results in the activation of downstream mitochondrial cell death pathways [137]. Interestingly, neither significant cardioprotective effects nor alterations in myocardial GRK2 signaling were detected in normotensive rats following RF-RDN treatment. These findings highlighted the importance of CA signaling for the communication between heart and kidney, suggesting involvement of GRK2 signaling in these interactions (see overview of cardiorenal crosstalk in Figure 2). Future studies will provide more mechanistic insights into GPCR-G $\beta\gamma$ -GRK2 signaling in the development and progression of heart disorders concomitant to the kidney injury. A summary of current animal models for CVD and CRS utilized in studies outlining the role of GPCR-G $\beta\gamma$ -GRK2 signaling pathway in these pathologies is provided in Table 1.

6. Conclusions

GPCR signaling modulates a wide array of physiologic processes throughout the organism, consequently its dysregulation causes maladaptive responses in different organ systems, including the cardiovascular and renal systems. Owing to the increasing clinical significance of disorders involving multiple organ systems, investigation of molecular pathways that mediate pathological crosstalk between affected organs may hold therapeutic promise. Further exploration of inhibiting the GPCR-G $\beta\gamma$ -GRK2 signaling pathway might lead to the development of novel approaches for HF, kidney injury and CRS treatment.

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Highlights

• GPCR signaling is a crucial regulator of cardiac and renal function;

- Dysregulation of GPCR-Gβγ-GRK2 signaling is associated with severe cardiac and renal disorders;
- $G\beta\gamma$ -GRK2 small molecule inhibitory treatment elicited beneficial effects in acute and advanced heart failure models;
- Gβγ-GRK2 signaling interdiction resulted in advantageous outcomes in acute kidney injury and in kidney injury concomitant with cardiac injury;
- GPCR-G $\beta\gamma$ -GRK2 signaling represents a potential mechanism underlying the crosstalk between heart and kidney in cardiorenal pathologies.



Figure 1. Novel strategies to target $G\beta\gamma\text{-}GRK2$ interactions

A: GPCR ligand binding, G-protein subunits dissociation, downstream signaling activation; B: SNS overactivation, GRK2 recruitment, GPCR phosphorylation, conformational change in GPCR; C: β -arrestin binding, G-proteins uncoupling, GPCR desensitization and downregulation, GPCR signaling attenuation. The uncoupled GPCR undergoes endocytosis and either degradation in proteasomes or recycling. Paroxetine stabilizes the kinase domain of GRK2 and thus inhibits its ability to phosphorylate GPCRs. β ARKct is a recombinant peptide that represents the G $\beta\gamma$ binding domain of GRK2 and blocks G $\beta\gamma$ -GRK2 interactions. The small molecule gallein competitively binds with G $\beta\gamma$ subunits and inhibits its protein-protein interactions, thus attenuating downstream G $\beta\gamma$ -GRK2 signaling.



Figure 2. The proposed role of GPCR-G $\beta\gamma$ -GRK2 signaling in pathologic crosstalk between organs

Decreased myocardial contractile function and reduced cardiac output in HF are initially compensated by increased activity of SNS released by CA produced in the adrenal gland. However, eventually CA overdrive elicits maladaptive effects including myocardial β -AR dysregulation via elevated G $\beta\gamma$ -GRK2 signaling. Moreover, G $\beta\gamma$ -GRK2 desensitizes adrenal α 2-ARs, thus creating and maintaining the vicious cycle of SNS overactivation that leads to further progression of HF. Our recent study demonstrated elevated activity of GPCR-GRK2 signaling in acute kidney injury detected in a pressure-overload induced murine model of HF 12 weeks after TAC. Future experiments will explore and validate the role of GPCR-G $\beta\gamma$ -GRK2 signaling in pathologic cardiorenal crosstalk.

Table 1

Current approaches for GPCR-G $\beta\gamma$ -GRK2 signaling manipulation and animal models for studying its role in the development and progression of CVD and CRS.

GPCR-G $\beta\gamma$ -GRK2 signaling manipulation	Summary of effects	Model name and reference
GRK2 knockout: • Constitutive/inducible ablation in cardiomyocytes • Inducible ablation in CFs	 Prevention of HF development, improved cardiac function, induced positive reverse remodeling Restored cardiac function, decreased fibrosis and apoptosis 	 Murine LAD ligation LV MI model [51] Murine LCA ligation LV MI model [52] and [56]
GRK2 overexpression: Constitutive overexpression in cardiomyocytes	Increased infarct size, promoted apoptosis, impaired cardioprotection	Murine LAD ligation LV MI model [49] and [50]
Recombinant protein inhibition of Gβγ-GRK2 signaling: • Recombinant carboxyl (C)-terminal fragment of GRK2 (βARKct)	 Improved cardiac contractility, mitigated left ventricular remodeling, normalized SNS activity and cardiac β-adrenergic signaling Preserved regional and global systolic function Ameliorated LV function, reversed LV remodeling and fetal gene expression, normalized neurohumoral levels 	 Rat cryoinfarction MI model [71] and [72] Ovine acute transmural MI model [75] Porcine left circumflex coronary artery (LCX) MI model [76]
Small molecule interdiction of Gβγ-GRK2 signaling: • Selective serotonin reuptake inhibitor paroxetine • Small molecule inhibitors (M119, Gallein) • Small molecule inhibitor Gallein • Small molecule inhibitor Gallein	 Improved cardiac function, limited ventricular remodeling normalized SNS overdrive and myocardial β- adrenergic system Mitigated HF progression, preserved cardiac function, decreased hypertrophy and fibrosis Attenuated HF progression, reduced cardiac hypertrophy, alleviated cardiac function Improved survival, alleviated cardiac function, attenuated cardiac hypertrophy, inflammation and apoptosis, restored myocardial β- adrenergic system, reduced CA production and adrenal remodeling May prolong renal dysfunction (likely due to supraphysiologic dose- related toxicity). Attenuated renal dysfunction and tubular damage, decreased inflammatory genes expression Attenuated renal dysfunction, tubulo- interstitial damage 	 Murine LCA ligation LV MI model [56] and [78] Murine isoproterenol acute HF model [83] and [84] Murine calsequestrin transgenic established HF model [83] and [85] Murine TAC model of pressure-overload induced cardiac hypertrophy and HF [86], [87], [88] and [89] Rat bilateral I/R acute kidney injury model [130] and [131] Murine bilateral I/R acute kidney injury model [127] and [128] Murine TAC model of pressure-overload induced cardiac hypertrophy and HF [87], [88], [89] and [127]

GPCR-G $\beta\gamma$ -GRK2 signaling manipulation	Summary of effects	Model name and reference
	concomitant with post-TAC HF	