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Comprehensive insights in GRK4 and hypertension: From mechanisms to potential therapeutics

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Abstract

G protein-coupled receptors (GPCRs) mediate cellular responses to diverse extracellular stimuli that play vital roles in the regulation of biology, including behavior. Abnormal G protein-coupled receptor kinase (GRK)-mediated regulation of GPCR function is involved in the pathogenesis of hypertension. Among the seven GRK subtypes, GRK4 has attracted attention because of its constitutive activity and tissue-specific expression. Increasing number of studies show that GRK4 affects blood pressure by GPCR-mediated regulation of renal and arterial function. The target receptor of GRK4 is confined not only to GPCRs, but also to other blood pressure-regulating receptors, such as the adiponectin receptor. Genetic studies in humans show that in several ethnic groups, *GRK4* gene variants (R65L, A142V, and A486V) are associated with salt-sensitive or salt-resistant essential hypertension and blood pressure responses to antihypertensive medicines. In this article, we present a comprehensive overview of GRK-mediated regulation of blood pressure,

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Declaration of Competing Interest

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focusing on the latest research progress on GRK4 and hypertension and highlighting potential and novel strategies for the prevention and treatment of hypertension.

Keywords

G protein-coupled receptor; G protein-coupled receptor kinase type 4; Dopamine receptor; Angiotensin II type 1 receptor; Hypertension

1. Introduction

Essential hypertension, also known as primary hypertension and idiopathic hypertension, or simply hypertension, is a major contributing risk factor for cardiovascular diseases and other diseases with adverse clinical outcomes, including stroke, heart failure, end-stage-renal disease, and all-cause mortality (Virani et al., 2021). In the United States National Health and Nutrition Examination Survey 2015 to 2018, the prevalence of hypertension was 28.2%, 60.1%, and 77.0% among subjects 20 to 44, 45 to 64, and 65 years old, respectively (<https://www.cdc.gov/nchs/nhanes/>). The estimated expenditure for hypertension in the US from 2016 to 2017 was \$52.4 billion (Virani et al., 2021). Of 154 health conditions in the US, hypertension ranked 10th in health care costs (Dieleman et al., 2020). Hypertension is a major public health problem worldwide (Fisher & Curfman, 2018; Kostova et al., 2020).

The pathogenesis of hypertension is determined by genetic and environmental factors and their interaction (Harrison, Coffman, & Wilcox, 2021; Wang et al., 2020), in which several organs such as the arteries, brain, endocrine organs, gastrointestinal tract, heart, kidneys, and nerves are involved. Many endocrine factors or neurotransmitters, such as dopamine, angiotensin II (Ang II), endothelin, and epinephrine, regulate sodium homeostasis and vascular reactivity, via their G protein-coupled receptors (GPCRs), to maintain a normal systemic arterial blood pressure (Eroglu, Kocyigit, & Lindholm, 2020; Prieto, Gonzalez, Visniauskas, & Navar, 2021; Rianto, Hoang, Revoori, & Sparks, 2021; Yang, Villar, Jose, & Zeng, 2021). However, in hypertension, GPCR function is aberrant, causing sodium retention and attenuated vasodilation (Prieto et al., 2021; Rianto et al., 2021; Yang et al., 2021). Although the mechanisms leading to aberrant GPCR function are complex, abnormal GPCR phosphorylation plays an important role in the pathogenesis of hypertension.

GPCR kinases (GRKs) comprise seven serine/threonine protein kinases, characterized by their ability to recognize specific GPCRs and phosphorylate their intracellular elements, leading to their uncoupling from G protein subunits, also known as desensitization, promote receptor internalization, and terminate the GPCR-mediated signaling pathway (Benovic, 2021; Gurevich & Gurevich, 2020). Due to the important roles of GPCRs in the development of human diseases, GRKs have attracted considerable attention for their role in the pathogenesis of cardiovascular diseases, including hypertension (de Lucia et al., 2021; Li et al., 2021; Pflieger, Gresham, & Koch, 2019).

Among the GRKs, GRK4 may be the most versatile subtype of GRK in the regulation of blood pressure (Yang, Villar, Jones, Jose, & Zeng, 2015). Several studies have shown that GRK4 plays a vital role in the pathogenesis of hypertension and response to antihypertensive

treatment (Jeong et al., 2020; Vandell et al., 2012; Zhang et al., 2020). In this article, we present a comprehensive overview of GRK-mediated regulation of blood pressure, especially focusing on the latest research progress on GRK4 and hypertension, and highlighting potential and novel strategies for the prevention and treatment of hypertension.

2. Classification and characteristics of GRKs

It is well known that there are over 800 genes encoding GPCRs in the human genome. However, of the primary mediators of agonist-dependent phosphorylation of GPCRs, only seven GRKs (GRK1–7) have been identified (Benovic, 2021; Gurevich & Gurevich, 2020; Pflieger et al., 2019). The GRKs comprise a family of seven serine/threonine protein kinases characterized by their ability to recognize and phosphorylate, specifically, agonist-activated GPCRs (Chaudhary & Kim, 2021; Harris, 2012; Santos-Otte et al., 2019). According to their amino acid sequences and ternary structural homologies, the seven mammalian GRKs can be classified into three sub-groups: the GRK1-like subfamily, also called as visual GRK subfamily, which includes rhodopsin kinase GRK1 and visual pigment kinase GRK7; the GRK2-like subfamily, otherwise known as β -adrenergic receptor (β -AR) kinase subfamily, which includes GRK2 (β -ARK1) and GRK3 (β -ARK2); and the GRK4-like subfamily, which includes GRK4, GRK5, and GRK6 (Benovic, 2021; Gurevich & Gurevich, 2020; Pflieger et al., 2019).

The seven GRKs share certain characteristics but are distinct enzymes with specific regulatory properties. They have a similar basic structural architecture, which consists of an amino terminal (N-terminal) domain (~185 amino acids), a central highly conserved catalytic domain (~270 amino acids), and a carboxyl terminal (C-terminal) domain (~105 to 230 amino acids). Each GRK has a well-conserved N-terminus, which is considered to be vital for receptor binding and selective recognition of the activated GPCR. GRKs also contain a regulator of G protein signaling (RGS) homology (RH) domain (~120 amino acids) (Chaudhary & Kim, 2021; Hullmann, Traynham, Coleman, & Koch, 2016; Yu, Sun, Jiao, & Lee, 2018). Differences among GRKs are most notable in the C-terminal domain, which is responsible for their membrane localization. For example, GRK1 and GRK7 are prenylated at the C-terminus; however, GRK2 and GRK3 have a pleckstrin homology domain, which facilitates recruitment to the membrane via interacting with G $\beta\gamma$ subunits. Within the GRK4-subfamily, GRK4 and GRK6 are palmitoylated, whereas GRK5 contains a positively charged lipid-binding amphipathic helix and binds phospholipids via its C-terminus (Homan, Glukhova, & Tesmer, 2013; Ribas et al., 2007). By contrast, the central domain that is involved in its kinase catalytic function is highly conserved among different GRKs. It should be noted that how GRKs recognize GPCRs still remains unclear. A recent study showed direct evidence that upon receptor activation, the N-terminus of GRK1 forms a helix that anchors into the open cytoplasmic cleft of rhodopsin (Chen et al., 2021). The mechanism by which other GRK subtypes recognize GPCRs and vice-versa needs further studies.

The seven GRK subtypes have different tissue distributions. The four GRK subtypes, GRK2, GRK3, GRK5, and GRK6, are widely expressed in mammalian tissues (Benovic, 2021; Gurevich & Gurevich, 2020; Pflieger et al., 2019). By contrast, the other three subtypes,

GRK1, GRK4, and GRK7, are found in a limited number of tissues. Among them, GRK1 and GRK7 are found almost exclusively in the retina, whereas GRK4 has very limited distribution in a few organs or tissues (see below) (Benovic, 2021; Gurevich & Gurevich, 2020; Li et al., 2021; Pflieger et al., 2019; Yang et al., 2015; Zhang et al., 2020). The different tissue expressions among GRKs suggest specific properties in the regulation of GPCRs or other targets and different physiological functions in different organs or tissues.

3. Regulation of blood pressure by GRKs

Studies have shown that most GRKs, e.g., GRK2, GRK3, GRK4, GRK5, and GRK6, but not GRK1 and GRK7, are involved in the regulation of blood pressure via different mechanisms (Chaudhary & Kim, 2021; Harris, 2012; Vandell et al., 2012; Yang et al., 2015; Zhang et al., 2020).

Elevated GRK2 expression and activity are associated with increased blood pressure (Cohn et al., 2009; Murga et al., 2019; Napolitano et al., 2012). Hypertensive humans and several animal models of hypertension have elevated expression and activity of GRK2 in lymphocytes, aortas, and vascular smooth muscle cells (VSMCs) (Cohn et al., 2009; Gros et al., 2000; Izzo et al., 2008; Napolitano et al., 2012; Zhao, Vanhoutte, & Leung, 2015). Our previous study also found that maternal diabetes mellitus-programmed hypertension in the offspring is caused by increased GRK2 activity (Luo et al., 2018). Mice with VSM-targeted GRK2 overexpression have impaired (β -AR-mediated vasodilation and increased resting blood pressure (Eckhart, Ozaki, Tevaearai, Rockman, & Koch, 2002). Pharmacological blockade of GRK2 and (β_1 -AR interaction prevents the development of hypertension in spontaneously hypertensive rats (SHRs), suggesting that inhibition of GRK2 activity could be a strategy for treating hypertension and protecting target organs (Polhemus et al., 2016; Rainbow et al., 2018; Sun et al., 2021). GRK2 downregulation/inhibition also enhances cardiac insulin sensitivity and mild heart hypertrophy with preserved systolic function, that is accompanied with repressed expression of genes related to pathological hypertrophy (Lucas et al., 2014). However, there are a few reports that could be considered as inconsistent with a major role of GRK2 in some hypertensive models (Avendaño et al., 2014; Ciccarelli et al., 2013; Cohn et al., 2008; Oliver et al., 2014; Tutunea-Fatan et al., 2018; Tutunea-Fatan, Caetano, Gros, & Ferguson, 2015). For example, nitric oxide production in adult GRK2 hemizygous mice protects against Ang II-induced hypertension (Avendaño et al., 2014). GRK2 expression is decreased in the aortas of N(G)-nitro-L-arginine methyl ester (L-NAME)-induced hypertensive rats, but unchanged in the mesenteric arteries of SHRs (Oliver et al., 2014). The reasons leading to the differences are still unknown, which need to be elucidated in the future.

Transgenic mice with cardiac-restricted expression of GRK3 have elevated systolic blood pressure and increased cardiac output, associated with cardiac myocyte α_1 -AR hyper-responsiveness (Vinge et al., 2008). Although α -ARs are important regulators of vascular resistance and GRK3 is expressed in the VSMCs, it is still unclear whether or not GRK3 can regulate blood pressure by affecting vascular resistance. An overall blood pressure lowering effect of GRK3 activation is supported from clinical evidence showing an inverse correlation

between GRK3 mRNA expression in lymphocytes and ambulatory systolic and diastolic blood pressure in normotensive and hypertensive humans (Oliver et al., 2010).

VSM-specific overexpression of GRK5 in mice causes hypertension (Keys, Zhou, Harris, Druckman, & Eckhart, 2005). In SHRs, the intramyocardial gene transfer of the amino-terminal region of GRK5 reduces GRK5-mediated exacerbation of cardiac hypertrophy in a blood pressure independent manner (Hullmann et al., 2014; Sorriento et al., 2010; Sorriento et al., 2018). GRK5 and β_1 -AR expressions are increased in the left ventricles of rats with hypertension induced by L-NAME and in peripheral blood mononuclear cells from humans with heart failure. By contrast, GRK2 expression is increased in all tissues with increased β_2 -AR expression (Montó et al., 2015). Thus, GRK5 may regulate β_1 -AR whereas GRK2 may regulate β_2 -AR. In addition, GRK5 polymorphism (41Q > L) is associated with a decreased risk of cardiovascular outcomes in hypertensive patients treated with atenolol and hydrochlorothiazide (Lobmeyer et al., 2011). This GRK5 polymorphism may be also protective in heart failure by inhibition of β -AR signaling (Liggett et al., 2008).

Renal GRK6 expression is decreased in hypertensive subjects (Xu, Watanabe, Felder, & Jose, 2001). GRK6 deficient mice have enhanced coupling of dopamine D₂-like receptors in the striatum, which suggests that GRK6 negatively regulates these receptors (Gainetdinov et al., 2003). In the intestinal cell line IEC-6, GRK6 and GRK4 are responsible for the homologous desensitization of dopamine D₁-like receptors (Fraga, Jose, & Soares-da-Silva, 2004). In hypertensive Sprague-Dawley (SD) rats, caused by coarctation of the abdominal aorta, expression of GRK6 in endothelial cells from the common carotid arteries is repressed (Wang et al., 2017). In rats with SHR heart failure, there is a subcellular redistribution of GRK6 from the intercalated discs to the cytoplasm of cardiac myocytes (Yi et al., 2005). In addition, in rats, overexpression of GRK6 in the 6-hydroxydopamine-injured striatum induces the internalization of D₁ receptor (D₁R) and normalizes the D₁R signaling by promoting its desensitization (Ahmed et al., 2010). However, the role of GRK6 in human essential hypertension remains to be determined.

4. Regulation of blood pressure by GRK4

4.1. Features of GRK4

Of all the seven GRKs, GRK4 is the unique subtype with four splice variants (GRK4 α , β , γ , and δ) in humans (Fig. 1) (Premont et al., 1996). GRK4 α , consisting of 578 amino acids, is the longest and full-length GRK isoform. GRK4 β , consisting of 546 amino acids, has no N-terminal exon 2 (32-codon deletion), which contains the phosphatidylinositol bisphosphate (PIP₂) binding region. GRK4 γ , consisting of 532 amino acids, has no C-terminal exon 15 (46-codon deletion). GRK4 δ , consisting of 500 amino acids, the shortest isoform, has no exons 2 and 15 (Jose, Soares-da-Silva, Eisner, & Felder, 2010; Premont et al., 1996). It should be noted that alternative splicing in GRK4 β and GRK4 δ would lead to the loss of α 0 and α 1 helices of the RH domain, and GRK4 γ and GRK4 δ would lose the C-terminal end of α 10 and all of the α 11 helix, which brings the folding and stability of these three GRK4 isoforms into question (Allen et al., 2015). Unlike humans with four GRK splice variants, rats have five GRK4 splice variants (GRK4A, B, C, D, and E), while mice have only one GRK4. Rat GRK4A has 76% identity with GRK4 α , the longest of the human GRK4 splice

variants. However, the mouse and rat GRK4 sequences are 90% identical (Premont et al., 1996; Virlon et al., 1998).

Three missense single nucleotide polymorphisms (SNPs) in the coding region of GRK4 γ have attracted the most attention in the regulation of blood pressure. The SNPs in nucleotide 448, CGT to CTT (amino acid 65R > L, rs2960306); nucleotide 679, GCC to GTC (amino acid 142A > V, rs1024323); and nucleotide 1711, GCG to GTG (amino acid 486A > V, rs1801058) are associated with hypertension (Jose et al., 2010; Yang et al., 2015), which will be discussed in detail. It should be noted that the GRK4 SNPs (A142V, A486V and R65L) increase its activity, although the mechanisms are not clear at this time.

Unlike the ubiquitously expressed GRK2, GRK3, GRK5, and GRK6, GRK4 is expressed in a limited number of tissues, i.e., arteries, bones, cerebellum, heart, kidneys, small intestines, thyroid, and testes (Jose et al., 2010; Voigt, Holzapfel, Meyer, & Paschke, 2004; Yang et al., 2015). The GRK4/GRK5/GRK6 group has constitutive activity while GRK1, GRK2, GRK3, and GRK7 are activated by binding to ligand-activated GPCRs (Baameur et al., 2010; Fraga et al., 2004; Jose et al., 2010; Li et al., 2015a, b; Yang et al., 2015). The constitutive activity of GRK4 may be due, at least in part, to its ability to bind to inactive G α and G β subunits (Keever, Jones, & Andresen, 2008; Neve, 2006).

4.2. Localization of GRK4

GRK4 has a restricted expression pattern, with highest levels in the testes and myometrium and, to a lesser extent, in a few other organs, including the arteries, brain, kidneys, small intestines, and thyroid, but minimal expression in the normal heart (Jose et al., 2010; Voigt et al., 2004; Yang et al., 2015).

Due to the vital role of the kidney in blood pressure regulation, the distribution of GRK4 in the kidney has been extensively studied. In human, rat, and mouse kidneys, GRK4 is highly expressed in the subapical membranes of renal proximal tubules (S1 and S3 segments), thick ascending limbs of the loop of Henle, distal convoluted tubules, and renal resistance vessels, and much less, in glomeruli (Felder et al., 2002; Sanada et al., 2006a, b; Villar et al., 2009; Yang et al., 2020). GRK4, distributed at the plasma membrane and cytoplasm under basal conditions, becomes internalized at the perinuclear area after activation of GPCRs (Felder et al., 2002; Villar et al., 2009; Yang et al., 2020). Specifically, in the renal plasma membrane microdomains, GRK4 is distributed in non-lipid raft fractions and in lipid raft fractions (Villar et al., 2009). We have also reported that GRK4 is localized in the nuclei of human renal proximal tubule cells and in HEK-293 cells, heterologously expressing hGRK4 γ wild type or variants (Wang et al., 2016).

The blood vessels are also critical in the regulation of blood pressure. GRK4 is expressed in the tunica media and adventitia of conductance and resistance arteries from rats and mice (Chen et al., 2014; Zhao et al., 2015). The GRK4 mRNA is well expressed in the aortas of Wistar Kyoto (WKY) rats and SHR s (Zhao et al., 2015). In fact, GRK4 is generally distributed in both large and small arterial vessels, including the thoracic aorta, superior mesenteric arteries, carotid arteries, and renal arteries; there is no difference in GRK4

expression among these vessels. Similar to the GRK4 location in renal proximal tubule cells, GRK4 is also distributed at the membrane and cytoplasm in VSMCs (Chen et al., 2014).

The heart and resistance vessels are the organs involved in the generation of arterial pressure. GRK4 is modestly expressed in the normal heart, but substantially increased after myocardial infarction (Li et al., 2021; Sanada et al., 2006a, b). Our recent study showed that in the heart, GRK4 protein is expressed to a higher extent in cardiomyocytes than in non-cardiomyocytes, such as smooth muscle cells, endothelial cells, and fibroblasts (Li et al., 2021).

4.3. Effect of GRK4 on blood pressure

GRK4 expression and activity in the kidneys and arteries are increased in the hypertensive state, relative to the normotensive state (Table 1). The increase in renal GRK4 levels in hypertension may be organ specific because there is no difference in cardiac GRK4 expression between WKY rats and SHRs (Sanada et al., 2006a, b). Moreover, unlike other GRK subtypes, such as GRK2 and GRK5, which have increased expression secondary to hypertension, GRK4 expression and activity are increased before the onset of hypertension. Therefore, GRK4 may be of importance in the development of hypertension.

A role for GRK4 in regulating blood pressure is supported by several studies using GRK4 knockout, selective renal reduction of GRK4 expression via the chronic renal cortical interstitial-selective infusion of GRK4 antisense oligodeoxynucleotides or GRK4 small interfering RNA (siRNA) delivered by ultrasound-targeted microbubble destruction, and overexpression of GRK4 variants (Table 2). In contrast to the GRK4 knockdown-mediated amelioration of blood pressure in hypertensive rodents, the expression of human GRK4 gene variants (hGRK4 γ 65 L, hGRK4 γ 142, or hGRK4 γ 486 V) in mice increases blood pressure (Table 2).

4.4. Mechanisms involved in the regulation of blood pressure by GRK4

Humans with essential hypertension have aberrant pathophysiological changes, such as increased renal sodium reabsorption and vasoconstriction and impaired arterial vasodilatation, that are not properly regulated by hormones via their receptors (Harrison et al., 2021; Kemp, Howell, Gildea, Keller, & Carey, 2020; Liu et al., 2021; Nizar, Shepard, Vo, & Bhalla, 2018; Olivares-Hernández et al., 2021; Rianto et al., 2021). Accumulating evidence show a role of GRK4 on the dysfunction of GPCRs or non-GPCRs in hypertension.

4.4.1. GRK4 regulates D₁R function by affecting its phosphorylation state—

The dopaminergic system exerts an autocrine/paracrine regulatory role on renal sodium transport and blood pressure via its five receptor subtypes (Albrecht et al., 1996; Gildea et al., 2010; Jose, Eisner, Drago, Carey, & Felder, 1996; Olivares-Hernández et al., 2021; Yu et al., 2006). Dopamine receptors, belonging to GPCRs, are classified into two families: D₁-like receptors (D₁R and D₅R), which stimulate adenylyl cyclase activity and D₂-like receptors (D₂R, D₃R, and D₄R), which inhibit adenylyl cyclase activity. D₁-like receptors are the major determinants of dopamine-mediated regulation of sodium transport (Albrecht et al., 1996; Chugh, Lokhandwala, & Asghar, 2011; Harris, 2012; Jose et al., 2010; Villar

et al., 2013; Wang et al., 2014; Yang et al., 2021; Ye et al., 2018; Yu et al., 2006). Indeed, during conditions of moderate sodium excess, more than 50% of renal sodium excretion is regulated by D₁-like receptors (Yang et al., 2021). However, in hypertensive patients and hypertensive animal models, such as the Dahl salt-sensitive rat and SHR, D₁R-mediated natriuresis and diuresis are decreased (Albrecht et al., 1996; Jose et al., 1996).

The impaired function of D₁R in hypertension is not due to decreased renal dopamine production, mutation of the D₁R gene, or decreased expression of D₁R, but is caused by increased D₁R phosphorylation and a defect in the coupling of the D₁R to its G protein/effector complex in renal proximal tubules (Yu et al., 2006), which is ascribed to increased GRK4 expression or activity (Felder et al., 2002; Jose et al., 2010; Lu et al., 2018; Sun, Chen, Wang, Zhou, & Zeng, 2020) (Fig. 2). SHRs have higher renal GRK4 expression than WKY rats (Sanada et al., 2006a, b). The intrarenal infusion of GRK4 antisense oligodeoxynucleotides or ultrasound-microbubble destruction-targeted GRK4 siRNA delivery to the kidney effectively decreases GRK4 expression, attenuates the augmented D₁R phosphorylation, normalizes the sodium balance, and reduces the blood pressure of SHRs (Huang et al., 2016; Sanada et al., 2006a, b). Besides the SHR, GRK4-mediated inhibition of D₁R is also reported in other hypertensive animal models (Lu et al., 2018; Sun et al., 2020; Wang et al., 2014; Ye et al., 2018). GRK4 γ 142 V transgenic mice have higher blood pressure than GRK4 γ wild-type mice, which is related to renal D₁R hyperphosphorylation and impaired D₁R-mediated natriuresis and diuresis (Felder et al., 2002; Wang et al., 2016; Yang et al., 2020).

The regulatory effect of GRK4 on D₁R in vivo is confirmed by in vitro studies. GRK4 activity is enhanced in human renal proximal tubule cells from hypertensive subjects, which constitutively phosphorylates D₁R even in the absence of D₁R agonist (Felder et al., 2002). Inhibition of GRK4 activity with heparin or depletion of GRK4 with antisense oligodeoxynucleotides blunts the D₁-like receptor agonist fenoldopam-induced D₁R desensitization in human renal proximal tubule cells (Watanabe, Xu, Bengra, Jose, & Felder, 2002). In Chinese hamster ovary cells transfected with GRK4 γ variants, including A142V, R65L and A486V, GRK4 activity is also increased that is associated with an increase in basal D₁R phosphorylation and impairment of D₁R-mediated cAMP production (Felder et al., 2002).

4.4.2. GRK4 regulates AT₁R function by affecting its expression and phosphorylation—Ang II, the principal renin-angiotensin system (RAS) effector peptide, exerts its physiological functions via its receptors, type 1 (AT₁R) and type 2 (AT₂R) (Fatima, Patel, & Hussain, 2021; Ziaja, Urbanek, Kowalska, & Piastowska-Ciesielska, 2021). The vast majority of the actions of Ang II are transduced via AT₁R, which increases renal sodium reabsorption and induces vasoconstriction (Paz et al., 2020).

The renal expression of GRK4 and AT₁R is higher in kidneys of SHRs than WKY rats (Yatabe et al., 2008). The selective intrarenal cortical infusion of GRK4 antisense oligodeoxynucleotides increases sodium excretion and decreases arterial blood pressure in SHRs (Sanada et al., 2006a, b); these effects are increased with the combined inhibition of renal GRK4 and AT₁R (Yatabe et al., 2008). Another study also found that the increased

renal GRK4 expression correlates with increased renal AT₁R expression and function in old Fischer 344 × Brown Norway rats, suggesting that GRK4-mediated increase in renal AT₁R expression may be necessary for the development of age-associated hypertension (Chugh, Lokhandwala, & Asghar, 2012).

AT₁R expression and function are regulated by GRK4 variants, as related to regulation of blood pressure (Fig. 2). The hypertension in hGRK4 γ 142 V transgenic mice is associated with increased AT₁R expression and function in the kidneys and arteries. GRK4 γ 142 V transgenic mice that are deficient of AT₁R have normal blood pressure (Wang et al., 2016). We have reported that both AT₁R expression and AT₁R-mediated vasoconstriction are higher in the aorta of hGRK4 γ 142 V than hGRK4 γ wild type transgenic mice. Moreover, Ang II causes a greater increase in blood pressure while infusion of the AT₁R antagonist candesartan causes a greater decrease in blood pressure in hGRK4 γ 142 V than hGRK4 γ WT transgenic mice (Chen et al., 2014). *Agtr1* mRNA and AT₁R protein expression and function are greater in VSMCs expressing hGRK4 γ 142 V than in cells expressing hGRK4 γ WT. In those cells expressing GRK4 γ 142 V, the activity of NF- κ B, a regulator of AT₁R promoter activity, is increased, accompanied by an increase in its binding to the AT₁R. The opposite is true for AT₁R protein degradation, indicating that the regulation of AT₁R expression by hGRK4 γ occurs at both transcriptional and post-translational levels (Chen et al., 2014).

Histone deacetylase type 1 (HDAC1) is involved into the regulation of *Agtr1* mRNA expression by GRK4. The phosphorylation of HDAC1 by GRK4 promotes the export of HDAC1 from the nucleus to the cytoplasm, which causes an increase in AT₁R promoter activity (Wang et al., 2016). The phosphorylation of AT₁R is also regulated by GRK4. The phosphorylation of AT₁R protein is lower in hGRK4 γ 142 V-transduced VSMCs than in hGRK4 WT-transduced cells, which seems to be inconsistent with GRK4 function as a phosphorylating enzyme. However, our further study showed that the interaction between GRK4 and AT₁R in VSMCs is reduced in hypertension, as determined by co-immunoprecipitation (Chen et al., 2014). This may be, at least in part, explained by the lower level of AT₁R phosphorylation, although the underlying mechanism leading to the reduced GRK4/AT₁R vascular interaction is not known. Additional studies are needed to elucidate this GRK4/AT₁R vascular interaction.

Similar to hGRK4 γ 142 V transgenic mice, hGRK4 γ 486 V transgenic mice also have higher blood pressure and greater renal AT₁R expression than hGRK4 γ WT mice. However, there is a distinct difference that hGRK4 γ 142 V transgenic mice are hypertensive on a normal salt diet. By contrast, hGRK4 γ 486 V transgenic mice develop hypertension only on a high salt diet (Diao et al., 2017; Wang et al., 2006a, b). The contribution of AT₁R in the salt sensitivity of blood pressure in hGRK4 γ 486 V transgenic mice remains to be determined.

The studies suggest that GRK4 regulates AT₁R expression and its actions in kidneys and arteries. In the kidney, the increased expression of GRK4 or presence of GRK4 variants increases the transcription of AT₁R, leading to an increase in AT₁R protein expression and renal sodium reabsorption, and subsequently hypertension. In the artery,

the increased expression of GRK4 or presence of GRK4 variants increases AT₁R protein expression, via both transcriptional and post-translational levels and aggravates AT₁R-induced vasoconstriction and subsequently hypertension. However, it should be noted that not only AT₁R but also other receptors, such as dopamine receptors, are targets of GRK4. Thus, the effect of GRK4 expression or variants on blood pressure could be ascribed, at least in part, to their regulation of AT₁R and other GPCRs.

The activation of AT₂R, the other receptor of Ang II, induces natriuresis and lowers blood pressure (Kemp et al., 2020). Our preliminary data indicate that hGRK4 γ 142 V transgenic mice have increased phosphorylation of renal AT₂R with impaired AT₂R-mediated natriuresis, compared with hGRK4 γ WT transgenic mice. This is supported by an in vitro study in which overexpression with hGRK4 γ 142 V in renal proximal tubule cells impairs AT₂R-mediated inhibition of Na⁺-K⁺-ATPase activity (Zhang and Yang, unpublished data). Accumulating pieces of evidence indicate that GRK4 regulates blood pressure by affecting several GPCRs, which affect sodium retention, arterial function, consequently leading to elevated blood pressure level.

4.4.3. GRK4 regulation of other GPCRs—More and more GPCRs are reported to be regulated by GRK4. These include endothelin receptor type B (ETBR), thromboxane receptors, and dopamine D₃ receptor (D₃R).

Endothelin, via its receptor, ETBR, decreases renal tubular sodium transport. However, in hypertensive states, the renal ETBR is hyperphosphorylated and ETBR-mediated sodium excretion is impaired (Dhaun & Webb, 2019). GRK4 is believed to play an important role in this process because down-regulation of renal GRK4 by siRNA normalizes the phosphorylation of ETBR and ameliorates the impaired renal ETBR function in the SHR (Yang et al., 2020). Moreover, hGRK4 γ 142 V transgenic mice, which are hypertensive, have increased phosphorylation of renal ETBR and impaired ETBR-mediated natriuresis and diuresis (Yang et al., 2020) (Fig. 2).

Thromboxane receptors may be also regulated by GRK4. GRK4 expression in aortic smooth muscles is higher in SHR than in WKY rats (Zhao et al., 2015). More importantly, in the SHR but not the WKY aorta, α_1 -AR activation desensitizes thromboxane receptors through activation of PKC- ϵ (Zhao et al., 2015), which is a positive regulator of GRK4 (Gildea et al., 2013). These findings suggest that GRK4 may be involved in the regulation of aortic thromboxane receptors in hypertension. However, the direct relationships between GRK4 and those receptors have not been fully elucidated.

In addition to D₁R, GRK4 also regulates other dopamine receptor subtypes, including D₃R. GRK4 is required for D₃R-mediated mitogenesis and activation of the mitogen activated protein kinase pathway in human renal proximal tubule cells (Villar et al., 2009). This effect of GRK4 on the regulation of D₃R is isoform-specific because even though all four GRK4 isoforms are expressed in human renal proximal tubule cells, only GRK4 α and GRK4 γ (GRK4 γ > GRK4 α) isoforms, not GRK4 β and GRK4 δ , modulate the phosphorylation of D₃R (Villar et al., 2009). However, the role of GRK4-mediated D₃R regulation in the development of hypertension has not been determined.

4.4.4. GRK4 regulation of non-GPCRs—A few studies also reported that non-GPCRs, such as adiponectin receptor 1 (AdipoR1), are regulated by GRK4. AdipoR1, as with GPCRs, has 7 transmembrane domains. However, the orientation of AdipoR1 is opposite that of GPCRs; its N-terminus is intracellular and its C-terminus is external (Tanabe et al., 2015).

Obesity, characterized by excess body fat, is associated with impaired natriuresis and increased risk of hypertension (Hall et al., 2021; Powell-Wiley et al., 2021). As an endocrine organ, adipose tissue produces and releases adipokines, such as adiponectin (Zhao, Kusminski, & Scherer, 2021). Adiponectin, an adipocytokine produced by adipose tissue can also be produced by renal proximal tubules (Perri et al., 2013). In addition to adiponectin receptor 1 (AdipoR1), there are two other adiponectin receptors, AdipoR2 and T-cadherin; AdipoR1 mRNA is 20 times higher than AdipoR2 in human renal proximal tubule cells (Shen, Hughes, Charlesworth, Kelly, & Peake, 2008). Adiponectin, via AdipoR1, decreases renal sodium transport. However, adiponectin-mediated natriuresis and diuresis are impaired in SHR. The impaired adiponectin function in hypertension may be due, in part, to the hyperphosphorylation of adiponectin receptor (Zhang et al., 2020). GRK4 is thought to play a role in the impairment of adiponectin receptor function in hypertension (Fig. 2) because hGRK4 γ 142 V transgenic mice have increased phosphorylation of renal AdipoR1 and impaired diuretic and natriuretic response to adiponectin, relative to hGRK4 γ WT mice. Moreover, renal-selective GRK4 knockdown via renal ultrasound-directed siRNA restores the adiponectin-mediated increase in sodium excretion and reduces blood pressure in SHR (Zhang et al., 2020). These observations indicate that GRK4 causes the hyperphosphorylation and impaired function of renal AdipoR1 in hypertension. These studies also show that GRK4 can regulate the function of GPCRs and non-GPCRs.

4.4.5. GRK4 regulation of cellular senescence—Cellular senescence, an age-related physiological process, is recognized as a vital contributor to the development of cardiovascular diseases, including hypertension (Gorgoulis et al., 2019). The heterologous expression of GRK4 in HEK293 cells causes cell cycle G1/G0 phase arrest, accompanied by an increase in senescence-associated β -galactosidase activity, indicating that GRK4 halts cell proliferation and induces cellular senescence (Luo et al., 2019; Xiao et al., 2017). Therefore, GRK4 may be associated with age-related hypertension, by impairing cellular growth and promoting senescence. However, the causal role of GRK4-mediated regulation of cellular proliferation or senescence on the pathogenesis of hypertension needs to be demonstrated.

5. Regulation of GRK4 expression and activity

GRK4 expression and activity are higher in the hypertensive than normotensive state. It is known that the presence of GRK4 variants leads to increased activity of GRK4, although the mechanisms are still unclear. To our knowledge, until today, there is only one article that reported the possible mechanism on how GRK4 polymorphism affects enzymatic activity (Allen et al., 2015). There are differences in the crystal structure of GRK4 α A486V and wild-type GRK4 α and other GRKs, e.g., GRK6. Allen et al found that there is a lag in the autophosphorylation of wild-type GRK4 α , which is required for full kinase activity. However, this lag is not observed in GRK4 α A486V, which has an increased rate of

autophosphorylation of a number of residues. These kinetic differences between wild-type GRK4 α and GRK4 α A486V may be related to their structural differences. The precise effect of GRK4 polymorphisms on its enzymatic activity needs further studies.

In addition to the inherent increase in GRK4 expression in the kidneys and arteries in primary hypertension, environmental factors such as cold stress, fine particulate matter exposure, and infection also increase the expression of GRK4 (Lu et al., 2018; Sun et al., 2020; Wang et al., 2014; Ye et al., 2018). Reactive oxygen species (ROS), may be the vital link between environmental factors and GRK4. For example, long-term exposure of fine particulate matter (PM_{2.5}) in SD rats increases ROS production, renal GRK4 expression, and blood pressure (Lu et al., 2018). The elevated GRK4 expression with long-term exposure to PM_{2.5} is due to oxidative stress because inhibition of ROS production by tempol, an antioxidant, decreases renal GRK4 expression, alleviates the hyperphosphorylation of renal D₁R, increases sodium excretion, and lowers the blood pressure of PM_{2.5}-exposed SD rats (Lu et al., 2018). In utero exposure to PM_{2.5} and other stressors, such as cold stress and infection, plays an important role in early life-induced hypertension in offspring that may be related to up-regulation of GRK4 expression, at least in arteries and kidneys (Sun et al., 2020; Wang et al., 2014; Ye et al., 2018).

Besides ROS, the increase in GRK4 expression caused by environmental and disease factors may be also a consequence of alterations of some transcription factors and signaling molecules. The GRK4 core promoter resides in the first 1851 bp upstream of its transcription start site (Hasenkamp et al., 2008), suggesting that the DNA-protein and/or protein-protein interaction patterns at this region may affect the transcription and expression of GRK4. The regulatory regions of the GRK4 promoter are independent of cell type and differentiation state (Hasenkamp et al., 2008). The transcription factor c-Myc, by binding to the promoter of GRK4, positively regulates GRK4 protein expression in human renal proximal tubule cells (Gildea et al., 2013). In the PM_{2.5} regulation of GRK4 expression, c-Myc is the key signal between ROS and increased GRK4 transcription (Lu et al., 2018; Ye et al., 2018). *trans*-Activator C/EBP family members, including CCAAT/enhancer-binding protein (C/EBP) α , C/EBP β , or C/EBP δ , also increase GRK4 promoter activity in renal cell lines HEK293T and COS7 cells and osteoblast-like osteosarcoma cell line SaOs-2 (Hasenkamp et al., 2008).

In addition to the regulation of the promoter of GRK4, other regions are also regulated by certain proteins and microRNAs. For example, in the cerebellum, FMRP, an RNA-binding protein, decreases the translation of GRK4 by an interaction between its C-terminal region and the G4RIF domain of GRK4 mRNA (Maurin et al., 2015). A ubiquitous calcium-binding protein calmodulin (CaM) interacts with the GRK4 subfamily, including GRK4, but not the GRK1-like- and the GRK2-like subfamilies, by binding to the C-terminal-or amino-terminal domain (Pronin, Satpaev, Slepak, & Benovic, 1997; Sallese et al., 1997; Sallese et al., 2000). Some miRNAs, such as miR430a and miR218a, have also been implicated in the regulation of GRK4 by targeting the 3' untranslated region of GRK4 (Guo et al., 2019), suggesting that miRNAs, a class of endogenously-initiated non-coding RNAs, may post-transcriptionally control GRK4 expression via either translational repression or mRNA degradation.

The process of intracellular trafficking may be also involved in the regulation of GRK4. Our previous studies showed that GRK4 is regulated by sorting nexin (SNX), an intracellular transport protein, which is involved in D₁R endocytosis and trafficking through the endosomes (Li et al., 2015a, b; Villar et al., 2013). SNX5 directly interacts with GRK4 and prevents GRK4 from targeting the phosphorylation sites of the D₁R, which is enhanced after D₁R activation (Villar et al., 2013). By contrast, depletion of SNX5 markedly increases the ability of GRK4 to phosphorylate constitutively D₁R in human renal proximal tubule cells, consistent with the in vivo studies showing that renal SNX5 depletion increases blood pressure, causes insulin resistance, and decreases D₁R-mediated sodium excretion (Li et al., 2015a, b; Li et al., 2018; Villar et al., 2013).

GRKs may be regulated by sex steroids, estrogens, progestins, and androgens. Estrogen, via estrogen receptor α and β , has beneficial effects on the regulation of blood pressure (Colafella & Denton, 2018; Mercurio et al., 2010; Somani, Pawelczyk, De Souza, Kris-Etherton, & Proctor, 2019; Zheng, Ji, Maric, Wu, & Sandberg, 2008). A few studies have found that GRKs, such as GRK2 and GRK6, are regulated by estrogens (Abraham et al., 2018; Ansonoff & Etgen, 2001; Miyoshi, Otsuka, & Shimasaki, 2013) but there is no study on the ability of GRK4 to regulate estrogens or their receptors. In fact, previous studies have shown that there is no difference on GRK4-mediated regulation of blood pressure between male and female hGRK4 γ WT mice and hGRK4 γ 142 V or hGRK4 γ 486 V transgenic mice, or male and female WKY rats and SHR (Diao et al., 2017; Felder et al., 2002; Wang et al., 2007; Wang et al., 2016).

6. GRK4 and human essential hypertension

6.1. GRK4 polymorphisms and hypertension in humans

Loci in chromosome 4 are associated with hypertension. The *GRK4* locus on human chromosome 4p16.3, correlates with essential hypertension and salt sensitivity (Allayee et al., 2001; Casari et al., 1995; Zeng et al., 2008).

Several studies have shown that the three GRK4 gene variants (R65L, A142V, and A486V) are positively associated with essential hypertension in several ethnic groups, including African-Brazilian-, Caucasian and Black American-, Caucasian-Australian-, Chinese-, Italian-, and Japanese-populations (Table 3) (Bengra et al., 2002; Gu et al., 2006; Kimura et al., 2012; Sanada et al., 2016; Speirs et al., 2004; Wang et al., 2006a, b). A meta-analysis confirmed the association between the GRK4 SNPs and hypertension risk among different populations (Liu & Xi, 2012; Zeng et al., 2008; Zhang, Sun, Liu, & Yang, 2015). Not included in the meta-analysis are recent studies of the association of new GRK4 SNPs and hypertension (GRK4 rs1644731) (Jiang et al., 2021), risk of both hypertension and diabetes (GRK4rs1557213) (Du et al., 2021), or cardiovascular disease risk and diabetes (GRK4rs60314379) (Cheng et al., 2021) in Han Chinese population.

About 50% of subjects with hypertension are salt-sensitive (Elijovich et al., 2016). The association between GRK4 variants and salt sensitivity or impaired urinary sodium excretion has been studied in both hypertensive patients and normotensive subjects (Table 3).

The positive association of GRK4 gene variants and hypertension are contrasted by other reports showing no such association (Martinez Cantarin et al., 2010; Rana et al., 2007; Staessen et al., 2008). For example, a study in a family-based random sample of a white population found that GRK4 A142V polymorphism did not contribute to increased blood pressure but the genetic variation in the DRD1 promoter associated with impaired renal sodium handling and blood pressure; no other GRK4 genotypes were studied (Staessen et al., 2008). Another study based on a large, community-based sample in White Americans also showed no association between GRK4 A142V or GRK4 A486V and hypertension (Rana et al., 2007). A study in African Americans aged 18–49 years even showed a negative association between GRK4 variants and hypertension, in which the GRK4 A486V variant was negatively associated with hypertension in the non-obese group (Martinez Cantarin et al., 2010). The reasons for these diverse outcomes are still unknown. Some factors, including ethnicity, age, salt sensitivity, failure to study all the GRK4 variants that are associated with hypertension, and gene-gene interaction should be taken into account.

The role of gene-gene interaction in phenotypes has attracted increasing attention. For example, Pereira et al found that there is an epistatic interaction between GRK4 variants R65L and A142V, and angiotensinogen that affects blood pressure and cardiovascular risk (Pereira Da Silva et al., 2020). In an African population from Ghana, the combination, that is most predictive of hypertension, is GRK4 R65L and angiotensin-converting enzyme (ACE), with an estimated prediction success of 70.5% (Williams et al., 2004). Moreover, the multi-gene interaction is also involved in the salt sensitivity of blood pressure. Sanada et al reported that among Japanese participants, GRK4 R65L, A142V, or A486V impaired a dopaminergic agonist-induced natriuretic effect, and a genetic model of the three GRK4 variants predicted the presence of salt-sensitive hypertension in 94.4% of cases. By contrast, the single-locus model with only GRK4 A142V was 78.4% predictive and a 2-locus model of GRK4 A142V and CYP11B2 C-344 T was 77.8% predictive of low-renin hypertension (Sanada et al., 2006a, b). A study in Korean children aged between 8 and 9 years also showed that the risk of obesity, a well-known pathogenic factor in hypertension, increased with GRK4 A486V, ACE, and SLC12A3 variants in boys, whereas it increased with GRK4 A486V and CYP11B2 variants in girls, as sodium intake increased (Lee et al., 2015).

6.2. Potential role of GRK4 in pharmacogenomics of hypertension

Recently, precision medicine, defined as personalized medicine enhanced by different technologies such as genome-wide association studies, has attracted great attention in the treatment of hypertension (Padmanabhan & Dominiczak, 2021). The presence or absence of GRK4 gene variants has been shown to be valuable in guiding therapeutic antihypertensive strategies. Common variants of GRK4 can predict the blood pressure response to antihypertensive medicines, such as angiotensin receptor blockers (ARBs), β -adrenergic blockers, low salt diet, and diuretics. In addition, studies have also shown the association between GRK4 variants and the blood pressure response to lifestyle modification, such as reduction of dietary salt intake. The role of GRK4 variants in hypertension and response to antihypertensive treatment in humans is summarized in Table 3.

7. Conclusions

In summary, genetic studies have shown a correlation between hypertension and GRK4 variants. Increasing evidence shows that GRK4, via several molecular mechanisms, plays a vital role in regulating the expression and function of blood pressure-related receptors, consequently affecting renal sodium handling, arterial function, and blood pressure (Fig. 2). The downregulation of increased GRK4 activity restores the normal blood pressure-related receptor function. Moreover, pharmacogenomics studies show that GRK4 variants can predict the blood pressure response to antihypertensive medicines. Thus, further studies targeting GRK4 or identifying additional GRK4 variants may provide new therapeutic antihypertensive strategies in the future.

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Abbreviations:

Ang II	angiotensin II
AT₁R	angiotensin II receptor type 1
AT₂R	angiotensin II receptor type 2
AdipoR1	adiponectin receptor 1
AR	adrenergic receptor
D₁R	dopamine D ₁ receptor
D₃R	dopamine D ₃ receptor
ETBR	endothelin receptor type B
GPCRs	G protein-coupled receptors
GRK4	G protein-coupled receptor kinase 4
hGRK4γ	human GRK4 γ
HDAC1	Histone deacetylase type 1
L-NAME	L-NG-Nitro arginine methyl ester
ROS	reactive oxygen species
SD	Sprague-Dawley
SHRs	spontaneously hypertensive rats
SNPs	single nucleotide polymorphisms

WKY	Wistar Kyoto
WT	wild-type
VSMCs	vascular smooth muscle cells

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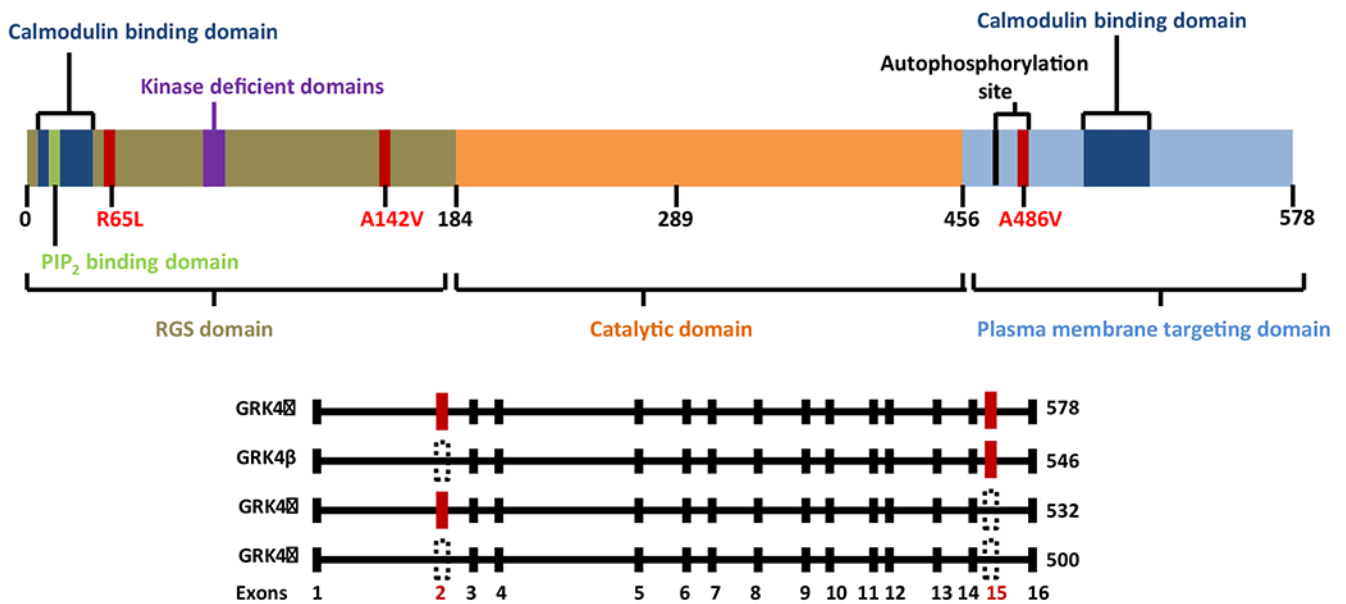


Fig. 1. Schematic representation of GRK4 domain architecture. Top: GRK4 has three primary domains: RGS domain; catalytic domain; and plasma membrane targeting domain. The numbers below the schematic diagram represent relative amino acid residues in GRK4. The positions of the GRK4 gene variants (amino acid substitutions R65L, A142V, and A486V) associated with hypertension are shown in red. The established and purported functional domains of GRK4 are also depicted in different colors. Bottom: The four GRK4 splice variants (GRK4 α , GRK4 β , GRK4 γ , and GRK4 δ) are distinguished by the presence or in-frame deletion of exon 2 (GRK4 β), exon 15 (GRK4 γ), or both (GRK4 δ). The red colored rectangles show the presence of exon 2 and/or exon 15 while the black-dotted triangles show the deletion of exon 2 and/or exon 15. Abbreviations: GPCR, G protein-coupled receptor; GRK4, G protein-coupled receptor kinase 4; PIP₂, phosphatidylinositol bisphosphate; RGS, regulator of G-protein signaling. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

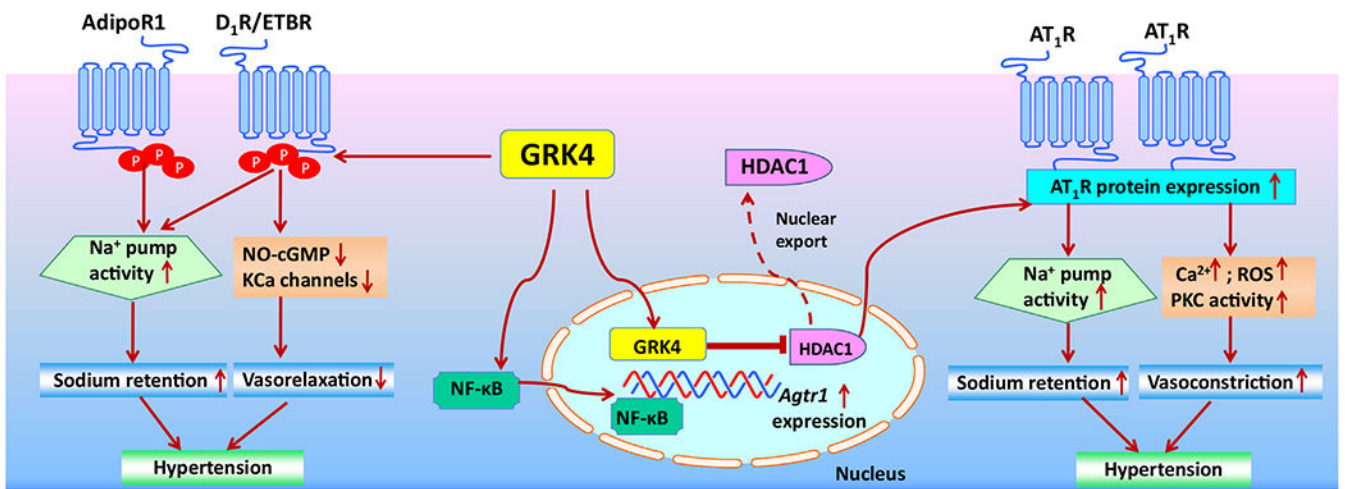


Fig. 2.

GRK4-mediated regulation of blood pressure via the kidneys and arteries.

GRK4 regulates blood pressure by modulating GPCRs and non-GPCRs in the kidneys and arteries by several mechanisms: 1) renal or arterial GRK4 increases the phosphorylation of D₁R, AdipoR1, and ETBR impairs receptor-mediated inhibition of the sodium pump (Na⁺/K⁺-ATPase) and attenuates receptor-induced vasorelaxation, subsequently leading to hypertension; 2) renal GRK4 promotes HDAC1 egress from the nucleus into the cytoplasm, which up-regulates *Agtr1* expression and the ability of AT₁R to increase the activity of the sodium pump (Na⁺/K⁺-ATPase), which results in an increase in renal sodium reabsorption and extracellular fluid volume, and consequently hypertension; 3) in the arteries, GRK4 increases NF-κB activity with more NF-κB binding to the AT₁R promoter, which increases *Agtr1* expression and AT₁R protein abundance, intracellular calcium concentration, ROS production, and PKC activity, enhancing vasoconstriction and consequently hypertension. Abbreviations: AT₁R, angiotensin II receptor type 1; AdipoR1, adiponectin receptor 1; cGMP, cyclic guanosine monophosphate; D₁R, dopamine D₁ receptor; ETBR, endothelin receptor type B; GRK4, G protein-coupled receptor kinase 4; HDAC1, histone deacetylase type 1; KCa: calcium-activated K⁺ channel; NF-κB, nuclear factor kappa B; NO: nitric oxide; PKC: protein kinase C; ROS, reactive oxygen species.

Table 1

Abnormal GRK4 expression and function in hypertensive states.

Experimental animal models of hypertension or hypertensive humans	Abnormal GRK4 expression and function in the hypertensive state	Blood pressure and targets regulated by GRK4 in the hypertensive state	References
SHRs	Higher mRNA and protein expressions of GRK4 in the renal cortex and aorta but not in the heart of SHRs, compared with WKY rats	Higher systolic blood pressure and renal cortical D ₁ R and ETBR phosphorylation, and increased sodium retention in SHRs compared with WKY rats; impaired ETBR-mediated diuresis and natriuresis in both male and female SHRs; inhibition of contractions to PGE ₂ induced by previous phenylephrine exposure, and desensitization of TP receptors induced by α ₁ -adrenoceptor activation in the aortae of SHRs, not WKY rats; GRK2 and GRK4 expressions increased in the aortae of SHRs, relative to WKY rats	Sanada et al., 2006; Yang et al., 2020; Zhao et al., 2015
Hypertensive humans	Higher basal (ligand-independent) and D ₁ R-stimulated GRK activities in RPT cells from hypertensive than normotensive subjects	Higher serine phosphorylation of D ₁ R and lower D ₁ R-mediated cAMP accumulation in RPT cells from hypertensive than normotensive subjects	Felder et al., 2002
Long-term exposure to fine particulate matter (PM _{2.5})-mediated hypertension	Increased mRNA and protein expressions of GRK4 in the kidney of PM _{2.5} -induced hypertension in SD rats and PM _{2.5} -treated RPT cells from WKY rats	Higher systolic and diastolic blood pressures; and renal D ₁ R phosphorylation, and decreased D ₁ R expression and receptor-induced natriuresis and diuresis in PM _{2.5} -treated SD rats. Decreased D ₁ R expression, increased D ₁ R phosphorylation and decreased D ₁ R-mediated inhibition of NKA activity in PM _{2.5} -treated RPT cells	Lu et al., 2018
In utero PM _{2.5} exposure mediated-hypertension in offspring	Increased GRK4 protein expression in renal cortex of offspring of PM _{2.5} -treated dams	Higher systolic blood pressure, lower 24 h urine volume and sodium excretion, decreased renal D ₁ R expression, increased renal D ₁ R phosphorylation, and decreased D ₁ -like receptor-induced natriuresis and diuresis in the offspring of PM _{2.5} -treated dams than the offspring of vehicle-treated dams	Ye et al., 2018
Prenatal LPS exposure-mediated hypertension in offspring	Increased renal GRK4 and GRK2 protein expressions in offspring of LPS-treated dams	Higher systolic blood pressure, lower 24 h urine volume and sodium excretion, decreased renal D ₁ R expression, increased D ₁ R phosphorylation, and decreased D ₁ -like receptor-induced natriuresis and diuresis in the offspring of LPS-treated than vehicle-treated dams	Wang et al., 2014
Prenatal cold stress exposure-mediated hypertension in offspring	Increased arterial GRK4 expression in offspring of prenatal cold stress-treated dams	Higher systolic-, diastolic- and mean blood pressures, impaired D ₁ R-mediated vasodilation, increased D ₁ R phosphorylation and cell internalization in mesenteric artery from offspring of prenatal cold stress-treated than control dams	Sun et al., 2020
Old FBN rats with hypertension	Increased renal GRK4 protein expression	Higher systolic, diastolic, and mean arterial blood pressures, decreased D ₁ R expression, impaired renal D ₁ R-mediated natriuresis, and no change in AT ₁ R expression in RPTs, and increased renal AT ₁ R function and oxidative stress in old rats; GRK4-mediated increase in HEK293 senescence	Chugh et al., 2011; Chugh et al., 2012; Xiao et al., 2017
High salt-induced hypertension	Renal GRK4 expression is lower on normal salt diet but higher on high salt diet in C57BL/6 J than in SJL/J mice	Higher systolic and diastolic arterial pressures, increased renal D ₅ R expression, impaired renal D ₁ R-like receptor-mediated natriuresis, shift to the right of the blood pressure-natriuresis plot in C57BL/6J mice, relative to SJL/J mice fed high salt diet	Escano et al., 2009

Abbreviations: AT₁R, angiotensin II receptor type 1; cAMP, cyclic adenosine monophosphate; D₁R, dopamine D₁ receptor; D₅R, dopamine D₅ receptor; ETBR, endothelin receptor type B; FBN rats, Fischer 344 × Brown Norway rats; GRK4, G protein-coupled receptor kinase 4; LPS, lipopolysaccharides; NKA, Na⁺/K⁺-ATPase; PGE₂, prostaglandin E₂; RPT, renal proximal tubule; SD, Sprague-Dawley; SHRs, spontaneously hypertensive rats; TP, thromboxane prostanoid; WKY, Wistar-Kyoto.

Table 2

In vivo and in vitro studies of GRK4 modification or variants.

GRK4 modification or variants	Animal or cell	Type of GRK4 modification	Effects of GRK4 modification on blood pressure and related functions	References
In vivo studies of GRK4 modification or variants				
GRK4 knockout	Knockout mice	Global knock mice	Decreases systolic and diastolic blood pressures	Wang et al., 2016
GRK4 siRNA	SHRs	Silencing renal GRK4 expression via UTMd-delivered GRK4 siRNA	Decreases blood pressure; increases sodium excretion and urine volume; reduces D ₁ R phosphorylation and improves D ₁ R-mediated sodium excretion; restores the adiponectin- or ETBR-mediated increase in sodium excretion	Huang et al., 2016; Yang et al., 2020; Zhang et al., 2020
GRK4 As-Odns	SHRs and WKY rats	Renal GRK4 depletion via the chronic renal cortical interstitial infusion of GRK4 As-Odns	Decreases D ₁ R phosphorylation to a greater extent in SHRs than in WKY rats; increases sodium excretion and urine volume, attenuates the increase in arterial blood pressure with age, and decreases protein excretion in SHRs, but not WKY rats	Sanada et al., 2006; Yatabe et al., 2008
GRK4 and AT ₁ R As-Odns	SHRs and WKY rats	Silencing both renal GRK4 and AT ₁ R via intrarenal cortical infusion of both GRK4 and AT ₁ R As-Odns	Decreases blood pressure and increases sodium excretion to a greater extent in SHRs than WKY rats; decreases circulating levels of renin, ang II, and aldosterone, reduces urine protein excretion and improves the GSI in SHRs and WKY rats; greater natriuresis and amelioration of hypertension in SHRs than the rats treated with either GRK4 or AT ₁ R As-Odn	Yatabe et al., 2008
Overexpression of hGRK4 γ 142V	hGRK4 γ WT and hGRK4 γ 142 V transgenic mice	Global transgenic mice	Higher blood pressure and renal D ₁ R phosphorylation, impaired D ₁ R-mediated sodium excretion, increased renal AT ₁ R expression and pressor response to Ang II in hGRK4 γ 142 V transgenic mice than those in hGRK4 γ WT transgenic mice; higher AT ₁ R protein expression in the aorta of GRK4 γ 142 V transgenic mice than hGRK4 γ WT transgenic mice; hyperphosphorylation of renal ETBR and AdipoR1, impairment of ETBR or AdipoR1 function, less natriuresis and diuresis induced by endothelin or adiponectin in SHRs than WKY rats and GRK4 γ 142 V transgenic mice than hGRK4 γ WT transgenic mice	Felder et al., 2002; Yang et al., 2020; Wang et al., 2016; Wang et al., 2007; Zhang et al., 2020.
Overexpression of hGRK4 γ 486 V	hGRK4 γ WT and hGRK4 γ 486 V transgenic mice	Global transgenic mice	Impairs sodium excretion, increases renal oxidative stress, and increases blood pressure with high salt diet in hGRK4 γ 486 V transgenic mice, relative to hGRK4 γ WT transgenic mice	Diao et al., 2017; Wang et al., 2006
Overexpression of hGRK4 γ 65 L	hGRK4 γ WT and hGRK4 γ 65 L transgenic mice	Global transgenic mice	Increases blood pressure with high salt diet in hGRK4 γ 65 L transgenic mice, relative to hGRK4 γ WT transgenic mice	Asico and Jose, unpublished data
In vitro studies of GRK4 modification or variants				
Inhibition of GRK4 activity	RPT cells from normotensive humans	Treatment with heparin, an inhibitor of GRK activity	Decreases the expression of GRK2 and GRK4 and attenuates the desensitization of D ₁ R, blunts p44/42 phosphorylation and mitogenesis induced by D ₃ R stimulation in hRPTCs	Watanabe et al., 2002; Villar et al., 2009
Inhibition of GRK4 expression	RPT cells from hypertensive humans	Treatment with antibody recognizing GRK4 γ / δ isoforms	Blocks the stimulatory effect of the D ₁ -like receptor agonist, fenoldopam, on GRK activity	Felder et al., 2002
GRK4 As-Odns	RPT cells from hypertensive humans	GRK4 knockdown with its specific As-Odns	Blocks the D ₁ R phosphorylation, blunts fenoldopam-induced D ₁ R desensitization, and restores the D ₁ -like receptor-mediated cAMP production	Felder et al., 2002

GRK4 modification or variants	Animal or cell	Type of GRK4 modification	Effects of GRK4 modification on blood pressure and related functions	References
GRK4 siRNA	RPT cells from SHR	GRK4 knockdown with its specific siRNA	Recovers the impaired inhibitory effect of ETBR on Na ⁺ /K ⁺ -ATPase activity	Yang et al., 2020
GRK4 siRNA	RPT cells from normotensive humans	RPT cells transfected with GRK4 siRNA	Blunts the D ₃ R-mediated mitogenesis	Villar et al., 2009
Inhibition of GRK4 activity	Rat intestinal epithelial cells	Treatment with heparin, an inhibitor of GRK activity	Heparin prevents the loss of inhibition of NHE activity caused by 25 min exposure to SKF-38393, a D ₁ -like receptor agonist	Fraga et al., 2004
Inhibition of GRK4 expression	Rat intestinal epithelial cells	Treatment with the anti-GRK4-6 antibody	Restores the loss of inhibition of NHE activity and stimulation of AC activity caused by 25 min exposure to SKF-38393, a D ₁ -like agonist	Fraga et al., 2004
Overexpression of GRK4 variants	CHO cells	Transfected with GRK4γ with SNPs (A142V, R65L, A486V or combined R65L and A486V)	Increases basal D ₁ R phosphorylation and impairs D ₁ R-mediated cAMP accumulation	Felder et al., 2002
Overexpression of GRK4 splice variants	CHO cells	Transfected with GRK4α, GRK4β, GRK4γ or GRK4δ	GRK4γ and GRK4α increase the phosphorylation of D ₃ R compared with untreated control cells; activation of D ₃ R results in a 3-fold and 2-fold increase in D ₃ R phosphorylation in cells expressing GRK4γ and GRK4α, respectively	Villar et al., 2009
Overexpression of hGRK4γ 142 V	RPT cells from normotensive humans	RPT cells transfected with hGRK4γ 142 V and GRK4γ WT plasmids	Increases activity of <i>Agtr1</i> and AT ₁ R protein expression	Wang et al., 2016
Overexpression of hGRK4γ 142 V	VSMCs	Transfected with hGRK4γ 142 V and GRK4γ WT plasmids	Higher AT ₁ R protein and mRNA expressions, lower AT ₁ R phosphorylation and protein degradation, higher AT ₁ R-mediated intracellular calcium concentration after stimulation with Ang II in hGRK4γ 142 V than in GRK4γ WT cells	Chen et al., 2014
Overexpression of GRK4	HEK293 cells	Transfected with full-length GRK4 plasmid	Induces cellular senescence, halts cell proliferation	Xiao et al., 2017; Luo et al., 2019

Abbreviations: AC, adenylyl cyclase; AdipoR1, adiponectin receptor 1; Ang II, angiotensin II; As-Odms, antisense oligodeoxynucleotides; AT₁R, angiotensin II receptor type 1; cAMP, cyclic adenosine monophosphate; CHO, Chinese hamster ovary; D₁R, dopamine D₁ receptor; D₃R, dopamine D₃ receptor; ETBR, endothelin receptor type B; GRK4, G protein-coupled receptor kinase 4; GSI, glomerular sclerosis index; hRPTCs: human renal proximal tubules cells; NHE: sodium-hydrogen exchanger; RPT, renal proximal tubule; SHR, spontaneously hypertensive rats; siRNA, small interfering RNA; UTMD, ultrasound-targeted microbubble destruction; VSMCs, vascular smooth muscle cells; WKY, Wistar-Kyoto; WT, wild-type.

Table 3

Summary of the role of GRK4 variants in hypertension and response to antihypertensive treatment in humans.

Ethnic Group	Subjects	Single- or multi-locus analyses	Association between GRK4 variants and hypertension	Response to antihypertensive medicines or dietary salt intake	Reference
Italian	Hypertensive: 60 Normotensive: 60	Single-locus	GRK4 486 V correlates with mild hypertension		Bengra et al., 2002
Ghanaian	Hypertensive: 126 Normotensive: 51	Multi-locus	The combination of GRK4 R65L and ACE predicts the hypertensive phenotype with an estimated success of 70.5%		Williams et al., 2004
Caucasian Australian	Hypertensive: 168 Normotensive: 312	Single-locus	GRK4 486 V is associated with hypertension		Speiris et al., 2004
Chinese	Hypertensive: 503 Normotensive: 490	Single-locus	GRK4 A486V is associated with hypertension		Gu et al., 2006
Chinese	Hypertensive: 503 Normotensive: 490	Single-locus	A486 allele is associated with hypertension		Wang et al., 2006
American	Normotensive adolescents: 664	Single-locus	GRK4 65 L is associated with stress-induced reduction of urinary sodium excretion in the blacks		Zhu et al., 2006
Japanese	Newly diagnosed, untreated hypertensives: 184	Single-locus	GRK4 R65L, A142V, and A486V, are associated with salt-sensitive hypertension; the GRK4 A142V or combination of GRK4 A142V and CYP11B2 is associated with low-renin hypertension; sodium excretion is inversely related to the number of GRK4 variants in hypertension	GRK4 variants (R65L, A142V, and A486V) impair renal dopamine-induced natriuresis, even in the absence of hypertension	Sanada et al., 2006
African Americans	AASK study Hypertensives randomized to metoprolol: 328	Single-locus		GRK4 A142V is associated with blood pressure response among men. Men with a GRK4 A142 are less responsive to metoprolol if they have the GRK4 L65 variant	Bhatnagar et al., 2009
African Americans	Hypertensive: 173 Normotensive: 239	Single-locus	GRK4 A486V variant is negatively associated with HBP in the nonobese group		Martinez et al., 2010
European ancestry	Hypertensive: 55; Normotensive: 130	Single-locus	GRK4 486 V is associated with salt sensitivity of blood pressure		Carey et al., 2012
African-derived Brazilian	Participants: 652	Multi-locus	Combination of NOS3-rs1799983 and GRK A486V is associated with DBP levels		Kimura et al., 2012
Black subjects	Mild to moderate hypertensive: 40	Single-locus		GRK4 variants, A142V and R65L, predict blood pressure response to reduction of dietary salt intake	Rayner et al., 2012
American	Mild-to-moderate hypertensive: 768	Single-locus	All three GRK4 variants (65 L, 142 V, and 486 V) correlate with increased risk for the primary outcome.	Increasing number of copies of the GRK4 variant 65 L-142 V haplotype is associated with	Vandell et al., 2012

Ethnic Group	Subjects	Single- or multi-locus analyses	Association between GRK4 variants and hypertension	Response to antihypertensive medicines or dietary salt intake	Reference
American	Healthy males: 24	Multi-locus		reduced response to the β -blocker monotherapy with atenolol. Subjects with at least three GRK4 allele variants have impaired natriuretic response to diuretics than those with less than three GRK4 allele variants	Wagner et al., 2012
Chinese	Participants: 3025	Single-locus	GRK4 variant rs2488815 correlates with lower values of eGFR		Montasser et al., 2014
Swiss	Hypertensive: 100	Single-locus		Subjects with GRK4 65 L and 142 V need more antihypertensive treatment and especially diuretic therapy (particularly for thiazide and thiazide-like diuretics) to reach the same MBP	Muskalla et al., 2014
Japanese	Hypertensive: 588 Normotensive: 486	Single-locus	GRK4 R65L, A142V, or A486V is associated with hypertension	Subjects with GRK4 142 V have a greater decrease in SBP in response to ARBs than non-carrier hypertensive patients	Sanada et al., 2016
Korean	Children: 2163	Multi-locus		A high sodium intake increases the obesity risk in children with GRK4 A486V; the combination of GRK4 A486V, ACE, and SLC12A3 variants increases the obesity risk in boys, whereas the combination of GRK4 A486V and CYP11B2 variants increases it in girls	Lee et al., 2015
Black Africans	Hypertensive: 105	Multi-locus	A high frequency of four NSVs of GRK4 (R65L, A116T, A142V, V486A) in patients with low renin-resistant hypertension		Jones et al., 2017
American	Patients with breast, lung, ovarian, or other cancers, 38 developed BIH: 114	Single-locus	GRK4 variant rs1419044 is associated with BIH		Frey et al., 2017
Korean	Adults: 15034	Single-locus	GRK4 65 L TT genotype is inversely associated with hypertension risk		Jeong et al., 2020
Chinese	cardiovascular disease patients: 326; noncardiovascular disease patients: 1209	Multi-locus	GRK4 rs60314379 is associated with cardiovascular disease risk in T2DM		Cheng et al., 2020
Chinese	Healthy: 1152; T2DM:1152; Hypertensive: 1152	Single-locus	GRK4 rs1557213 contributes to the risk of hypertension and T2DM		Du et al., 2021
Chinese	Hypertensive: 1239	Single-locus	Patients with GRK4 variants (A142V, A486V or R65L) are more likely to be non-dippers	Patients with GRK4 variants (A142V, A486V or R65L), relative to those without these variants, have a better antihypertensive response to candesartan	Cao et al., 2021
Chinese	Healthy: 65; Hypertensive: 151	Multi-locus	GRK4 rs1644731 and RDH8 rs1801058 are associated with hypertension.		Jiang et al., 2021

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Abbreviations: ACE, angiotensin-converting enzyme; ARBs, angiotensin receptor blockers; BIH, bevacizumab-induced hypertension; DBP, diastolic blood pressure; eGFR, estimated glomerular filtration rate; GRK4, G protein-coupled receptor kinase 4; HBP, high blood pressure; MBP, mean arterial blood pressure; NSVs, nonsynonymous variants; SBP, systolic blood pressure; T2DM, type 2 diabetes mellitus.