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Running Title (within 10 words)	LpK79 regulates vasodilatory factor activation and inflammation
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Abstract

The reduction of nitric oxide bioavailability in the endothelium induces endothelial dysfunction, contributing to the development of hypertension. Although *Lactobacillus* consumption decreases blood pressure, intracellular signaling pathways related to hypertension have not been well elucidated. Thus, this study examined the effect of spray-dried Lactiplantibacillus plantarum K79 (LpK79) on nitric oxide (NO) production, intracellular signaling pathways, and inflammatory responses related to vascular function and hypertension. NO production was assessed in human umbilical vein endothelial cells (HUVECs) treated with LpK79. Endothelial NO synthase (eNOS) and intracellular signaling molecules were determined using Western blot analysis. LpK79 dose-dependently increased NO production and activated eNOS via the PI3K/Akt signaling pathway HUVECs. Moreover, LpK79 mitigated the activation of crucial factors pivotal for vascular contraction in smooth muscle cells, such as PLCy, MYPT-1, and ROCK2. When HUVECs were treated with LpL79 in the presence of Escherichia coli lipopolysaccharide (LPS), LpK79 effectively suppressed mRNA and protein expression of proinflammatory mediators induced by E. coli LPS. These results suggest that LpK79 provided a beneficial effect on the regulation of vascular endothelial function.

Keywords: *Lactiplantibacillus plantarum*; hypertension; vascular function; signaling pathway; inflammation

Introduction

Hypertension is strongly associated with substantial cardiovascular disease risks, such as heart attack and stroke (Khalesi et al., 2014). Recent findings revealed that the increasing global prevalence of hypertension is a pressing health concern worldwide, affecting nearly one in two Americans (Ritchey et al., 2018). Furthermore, an estimated 1.3 billion adults are affected by hypertension worldwide (Zhou et al., 2021). Nitrovasodilators, such as organic nitrates and sodium nitroprusside, have proven effective in managing hypertensive crises because nitric oxide (NO) is a vasodilator directly linked to blood pressure regulation (Bryan, 2022). Therefore, NO is pivotal in regulating hypertension, and the reduction of NO bioavailability, otherwise known as endothelial dysfunction, is a characteristic feature of arterial hypertension (Hermann et al., 2006).

Additionally, inadequate NO production impairs blood flow regulation and circulation, subsequently increasing inflammation, oxidative stress, and immune dysfunction (Cervantes-Gracia et al., 2020). Vascular smooth muscle cells (VSMCs), which constitute the vascular wall, are crucial for regulating hypertension due to their plastic and dynamic features (Touyz et al., 2018). Extensive studies have reported that the Rho-associated kinase (ROCK) signaling pathway and its downstream effectors, such as the myosin light chain (MLC), promote inflammation and remodeling in VSMCs, resulting in the pathogenesis of cardiovascular disease (Zhang et al., 2015). Moreover, enhanced Rho-associated kinase signaling via MLC phosphorylation contributes to VSMC contraction (Ma et al., 2023).

Probiotic lactobacilli, including *Lactiplantibacillus plantarum*, have emerged as a novel therapeutic approach for regulating blood pressure. For instance, *L. plantarum* supplementation significantly reduced blood pressure in hypertensive patients (Sharafedtinov et al., 2013). Additional studies have indicated that probiotic supplementation had positive effects on lowering blood pressure (Dong et al., 2013; He et al., 2017; Khalesi et al., 2014). Malik et al. (Malik et al.,

2018) demonstrated that *L. plantarum* 299v supplementation improved vascular endothelial function and decreased systemic inflammation in patients with coronary artery disease. Furthermore, conditioned media derived from *L. acidophilus* exhibited anti-apoptotic and anti-inflammatory activities in human umbilical vein endothelial cells treated with lipopolysaccharide (Kalani et al., 2016). However, although extensive clinical trials reported positive effects on lowering blood pressure, the cellular mechanism by which probiotic lactobacilli accomplishes this attenuation is not well documented. Thus, this study investigated the impact of spray-dried *L. plantarum* K79 on NO production and elucidated its intracellular signaling pathway in human umbilical vein endothelial cells (HUVECs). In addition, the signaling pathway in vascular function and hypertension was also examined.

Materials and methods

Bacterial sample preparation

Spray-dried *L. plantarum* K79 (LpK79) was kindly provided from Novalacto Co. Ltd. (Daejon, Korea). Briefly, pasteurized skim milk containing glucose was aseptically inoculated with *L. plantarum* K79 and incubated at 37°C until pH reached 4.2. Next, the bacterial culture was subjected to heat treatment at 85°C for 20 min, cooled at 50°C, and subsequently underwent spray drying.

NO production in HUVECs

HUVECs were purchased from the American Type Culture Collection (ATCC; Manassas, VA, USA) and maintained in Kaighn's modification of Ham's F-12 medium (Gibco, Burlington, ON, Canada) supplemented with 10% fetal bovine serum (Gibco) and antibiotics (HyClone, Logan, UT, USA). The cells were seeded on a 96-well culture plate and treated with various

concentrations of LpK79 (0, 2.5, 5, and 10 mg/mL) at 37°C for 24 and 48 h. After obtaining spent culture supernatants, NO production was measured using a nitrate/nitrite fluorometric assay kit (Cayman Chemical, Ann Arbor, MI, USA).

Western blot analysis

HUVECs were seeded onto a 100 mm tissue culture dish and incubated at 37°C for two days. Then, cells were treated with various concentrations of LpK79 (0, 2.5, 5, and 10 mg/mL) at 37°C for 2 h, and cell lysates were obtained using a lysis buffer. In a separate experiment, VSMCs were purchased from ATCC and maintained in Kaighn's modification of Ham's F-12 medium (Gibco) under culture conditions as previously described. VSMCs were seeded and incubated at 37°C for two days. Afterward, cells were treated with the different concentrations of LpK79 (0, 2.5, 5, and 10 mg/mL) at 37°C for 2 h, and cell lysates were obtained using a lysis buffer to determine ROCK2 and MLC activation. Lysate concentrations were determined using a bicinchonic acid protein assay to perform western blot analysis. Equal volumes of lysate samples were loaded on a 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis, electrotransferred onto polyvinylidene difluoride membranes (Millipore, Bedford, MA, USA), and blocked with 5% skim milk. Next, the membranes were incubated with primary antibodies specific for phosphoinositide 3-kinase (PI3K), phosphorylated PI3K, Akt, phosphorylated Akt, endothelial nitric oxide synthase (eNOS), phosphorylated eNOS, MLC, phosphorylated MLC, ROCK2 (Cell Signaling Technology, Danvers, MA, USA), or β-actin (Santa Cruz Biotechnology, Santa Cruz, CA, USA) overnight at 4°C. After washing the membranes with Tris-buffered saline containing 0.1% Tween 20, membranes were incubated with horseradish peroxidase-conjugated IgG at room temperature for 2 h. Immunoreactive bands on the membranes were detected using

an enhanced chemiluminescence reagent (Dyne Bio, Seongnam, Korea) and visualized with a C-Digit blot scanner (Li-Cor Bioscience, Lincoln, NE, USA).

Quantitative reverse transcription polymerase chain reaction (qRT-PCR)

HUVECs were seeded in a 6-well culture plate and incubated at 37°C for two days. The cells were then stimulated with Escherichia coli lipopolysaccharide (LPS) (1 µg/mL) with or without LpK79 (0, 2.5, 5, and 10 mg/mL) at 37°C for 3 h. Total RNA was extracted, and complementary DNA (cDNA) was synthesized using random hexamers and M-MLV reverse transcriptase (Promega, Madison, WI, USA). Pro-inflammatory cytokines, interleukin (IL)-6, and IL-8 genes were amplified through polymerase chain reaction (PCR) using the SYBR green real-time PCR master mix (Toyobo, Osaka, Japan) and a StepOnePlusTM Real-Time PCR System (Applied Biosystems, Foster City, CA, USA) according to the following conditions: initial denaturation at 95°C for 10 min, 40 cycles of denaturation at 95°C for 15 s, and annealing/extension at 60°C for 1 min. The specific primers used in this study were as follows: IL-6, forward 5'-TGCAATAACCACCCTGACC-3' and reverse 5'-GTGCCCATGCTACATTTGCC-3'; IL-8, forward 5'-GTGCAGTTTTGCCAAGGAGT-3' and reverse 5'-CTCTGCACCCAGTTTTCCTT-3'; and GAPDH, forward 5'-AAGGTGAAGGTCGGAGTCAA-3' and reverse 5'-ATGACAAGCTTCCCGTTCTC-3'. The relative mRNA expression of IL-6 and IL-8 was normalized to that of GAPDH using the $2^{-\Delta\Delta CT}$ method.

Enzyme-linked immunosorbent assay (ELISA)

To determine the protein secretion of *E. coli* LPS-stimulated IL-6 and IL-8 in HUVECs with LpK79, HUVECs were seeded as previously described, and cells were stimulated with *E. coli*

LPS (1 μ g/mL) with or without LpK79 (10 mg/mL) at 37° for 24 h. After obtaining supernatants, IL-6 and IL-8 were quantified using ELISA kits (R&D Systems, Minneapolis, MN, USA).

Statistical analysis

Statistical significance between groups was determined through one-way analysis of variance (ANOVA) using IBM SPSS Statistics 23 software (IBM, Armonk, NY, USA) or unpaired two-tailed t-test using GraphPad Prism 5 (GraphPad Software Inc., La Jolla, CA, USA). P < 0.05 was considered statistically significant.

Results and discussion

LpK79 significantly increases NO production in HUVECs

When HUVECs were incubated with LpK79 at different concentrations (2.5, 5, and 10 mg/mL) for 24 h, LpK79 dose-dependently increased NO production (Fig. 1A). HUVECs were further incubated with LpK79 for 48 h and LpK79 also significantly increased NO production in a dose-dependent manner (Fig. 1B). Previous studies demonstrated that probiotic lactobacilli exert antihypertensive effects in animal models and human clinical trials (Gómez-Guzmán et al., 2015; Jauhiainen et al., 2010; Malik et al., 2018). Our findings corroborate the previous reports that LpK79 promotes NO production in HUVECs, subsequently reducing hypertension. NO released by endothelial cells are critical regulators for vasculature relaxation. Moreover, NO diffused by smooth muscle cells promoted relaxation by increasing cyclic guanosine monophosphate levels (Gu et al., 2020). Although the impact of particular compounds from probiotics on the vascular functions has not been thoroughly explained, a study has demonstrated that short chain fatty acids reduce the blood pressure through G protein-coupled receptor 41, present in vascular

endothelial cells (Natarajan et al., 2016). Thus, it may be speculated that metabolic short chain fatty acids derived from probiotics play a role in the regulation of vascular functions.

LpK79 activates eNOS through the PI3K/Akt pathway in HUVECs

Endothelial NO synthase (eNOS) primarily facilitated NO production in vascular endothelium. Numerous studies discovered that the phosphatidylinositol 3-kinase (PI3K) pathway triggers the activation of serine-threonine protein kinase B (Akt), leading to the direct phosphorylation of eNOS (Ahmad et al., 2018; Gu et al., 2020). HUVECs were exposed to LpK79 at concentrations of 2.5, 5, and 10 mg/mL for 2 h to investigate the impact of LpK79 on PI3K, Akt, and eNOS phosphorylation. As shown in Figure 2A, LpK79 significantly enhanced PI3K phosphorylation at 5 and 10 mg/mL and Akt phosphorylation at 2.5, 5, and 10 mg/mL. Furthermore, eNOS phosphorylation increased in the presence of LpK79. A recent report determined that while the probiotic Bifidobacterium longum ZL0210 did not substantially affect PI3K phosphorylation, it significantly upregulated Akt and eNOS phosphorylation in cardiomyocytes (Qi et al., 2023). Additional studies found that the PI3K/Akt pathway is closely linked to various diseases, including metabolic syndrome, gastrointestinal disorders, and allergies. Probiotic lactobacilli, such as L. rhamnosus GG, have been shown to modulate the PI3K/Akt pathway (Mohseni et al., 2021). However, probiotic lactobacilli increasing eNOS phosphorylation via PI3K/Akt pathway activation in human endothelial cells has yet to be established. In this study, we demonstrate that LpK79 activates the PI3K/Akt pathway, directly affecting eNOS activation in HUVECs.

LpK79 decreases factors for VSMC contraction

In addition to endothelial dysfunction, VSMC contractions contribute to hypertension (Yang and Hori, 2021). The ROCK pathway is crucial, and the myosin phosphatase target subunit (MYPT-1), an essential subset of ROCK targets, participates in VSMC contraction. Additionally,

phospholipase Cy (PLCy) is integral for promoting VSMC contractions by increasing the intracellular calcium concentration (Touyz et al., 2018). Therefore, PLCy and MYPT1 phosphorylation and ROCK2 activation in VSMCs were exposed to LpK79 to explore its impact on VSMC contraction. Notably, LpK79 treatment significantly decreased PLCy and MYPT1 phosphorylation and ROCK2 expression (Fig. 2B). Calcium-independent MLC kinase (MLCK) and calcium-dependent MLC phosphatase (MLCP) pathways regulate myosin light chain (MLC) phosphorylation in smooth muscle cell contraction. The RhoA/ROCK signaling pathway, a calcium-independent MLCK pathway, activates MYPT1 phosphorylation, prompting MLC activation (Zhao et al., 2021). Additionally, PLC phosphorylation in the calcium-dependent MLCP pathway increases intracellular calcium ions, enhancing MLC phosphorylation and promoting VSMC contraction (Ma et al., 2023). Lactobacillus fermentum CECT5716 demonstrated antihypertensive effects in rats with NO synthase inhibition through vasodilation (Gómez-Guzmán et al., 2015). However, the intracellular mechanisms underlying VSMC contraction associated with probiotic lactobacilli remain unclear. Our study indicates that LpK79 is essential for regulating essential factors linked to VSMC contraction.

LpK79 suppresses inflammatory responses in HUVECs

Impaired eNOS activation reduces NO production in HUVECs, resulting in endothelial dysfunction, such as inflammation. Thus, we examined whether LpK79 attenuated lipopolysaccharide (LPS)-induced inflammatory responses in HUVECs. As illustrated in Figure 3A, *Escherichia coli* LPS augmented IL-6 and IL-8 mRNA expression. In contrast, LpK79 treatment significantly reduced IL-8 mRNA expression but not IL-6 mRNA. However, LpK79 significantly diminished IL-6 secretion in *E. coli* LPS-induced HUVECs and IL-8 secretion. These results suggest that LpK79 suppresses endothelial dysfunction-related inflammatory

responses. IL-6 impairs endothelial barrier function and elevates angiotensin II type 1 receptor expression (Wassmann et al., 2004). Angiotensin II is an endocrine ligand that contributes to hypertension by interacting with its angiotensin II type 1 receptor (Eguchi et al., 2018). Therefore, a high IL-6 level in HUVECs has a positive influence on the interaction between angiotensin II and angiotensin II type 1 receptors, increasing hypertension risk. While our observations indicated that IL-6 mRNA expression in HUVECs was not significantly reduced when exposed to LpK79, there was an effective suppression of IL-6 protein levels. This discrepancy may be because mRNA expression was assessed in HUVECs for an extended duration of 24 h. In contrast, IL-8 mRNA and protein levels were significantly suppressed in HUVECs with LpK79 at 24 h. IL-8 is also closely related to the vascular inflammatory process, and excessive IL-8 production was observed in atherosclerotic lesions (Papadopoulou et al., 2008). Thus, inhibiting IL-8 and other inflammatory mediators in activated HUVECs could be beneficial in maintaining endothelial homeostasis and mitigating the risk of vascular diseases.

Conclusions

This study demonstrated that LpK79 increased NO production by activating eNOS through the PI3K/Akt pathway in HUVECs while reducing the expression of factors associated with VSMC contraction. Additionally, LpK79 suppressed inflammatory mediator expressions in HUVECs. These results suggest that LpK79 can provide a beneficial effect on the regulation of vascular endothelial function.

Conflicts of interest

The authors declare that they have no conflicts of interest.

Acknowledgments

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Author contributions

Conceptualization: Hwang Y, Kang SS. Formal analysis: Kim KH, Kang SS. Methodology: Kim KH, Kang SS. Validation: Kim KH, Hwang Y, Kang SS. Investigation: Kim KH, Kang SS. Writing - original draft: Kim KH, Kang SS. Writing - review & editing: Kim KH, Hwang Y, Kang SS.

Ethics Approval

This article does not require IRB/IACUC approval because there were no human or animal participants.

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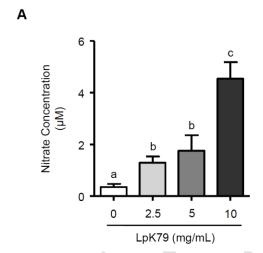
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Figure legends

Fig. 1. Effect of LpK79 on nitric oxide (NO) production in human umbilical vein endothelial cells (HUVECs). HUVECs were incubated with LpK79 (2.5, 5, and 10 mg/mL) at 37°C for 24 h (A) and 48 h (B). NO production was measured using a nitrate/nitrite fluorometric assay kit. *P* <

0.05 was considered statistically significant. Different letters (a–d) indicate statistical differences between groups.

- **Fig. 2.** Effect of LpK79 on PI3K/Akt phosphorylation in human umbilical vein endothelial cells (HUVECs) and MYPT1, PLCγ, and ROCK2 activation in vascular smooth muscle cells (VSMCs). HUVECs and VSMCs were incubated with LpK79 (2.5, 5, and 10 mg/mL) at 37°C for 2 h. PI3K and Akt phosphorylation in HUVECs (A), MYPT1 and PLCγ phosphorylation, and ROCK2 expression in VSMCs (B) were measured using Western blot analysis. *P* < 0.05 was considered statistically significant. Different letters (a–c) indicate statistical differences between groups.
- **Fig. 3.** The inhibitory effect of LpK79 on IL-6 and IL-8 expression in human umbilical vein endothelial cells (HUVECs). HUVECs were incubated with E. coli LPS (1 μ g/mL) with or without LpK79 (10 mg/mL) at 37°C for 24 h. IL-6 (A) and IL-8 mRNA expressions (B) were determined using qRT-PCR. IL-6 (C) and IL-8 (D) protein secretions were measured using ELISA. P < 0.05 was considered statistically significant. Different letters (a–c) indicate statistical differences between groups.
- **Fig. 4.** Schematic illustration of LpK79 regulating factors related to vascular functions in endothelial and smooth muscle cells.



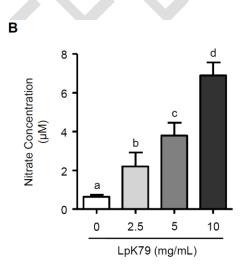
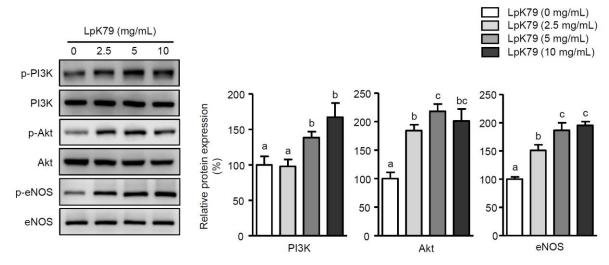


Figure 1





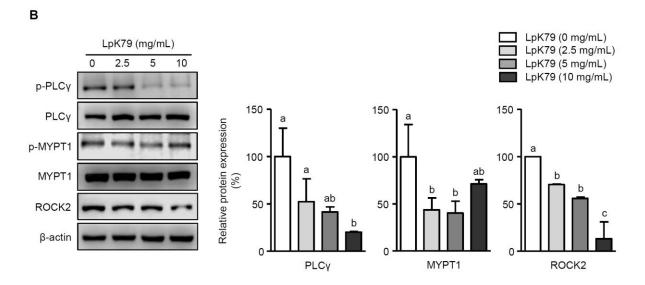
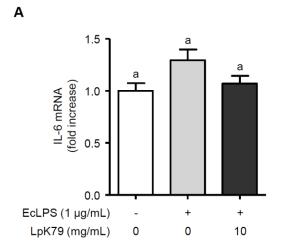
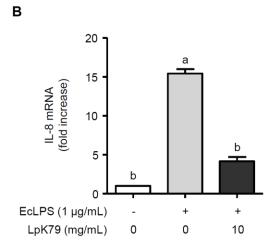
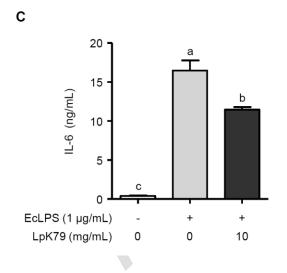


Figure 2







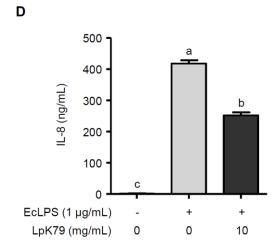


Figure 3

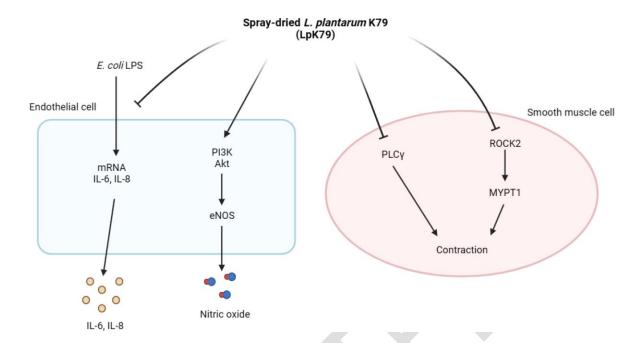


Figure 4