



# Potential neuroprotective effects of heat-killed *Lactococcus lactis* KC24 using SH-SY5Y cells against oxidative stress induced by hydrogen peroxide

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**Abstract** The present study was an investigation of the neuroprotective effects of probiotic bacteria in SH-SY5Y neuroblastoma cells experiencing oxidative stress. The bacterial strains were: commercial *Lactobacillus rhamnosus* GG; two isolated bacterial strains (*Lactobacillus delbrueckii* KU200170 and *Lactobacillus plantarum* KU200661); and probiotic *Lactococcus lactis* KC24. To evaluate the neuroprotective effects of the bacteria, a conditioned medium (CM) was prepared using HT-29 cells cultured with the heat-killed probiotic strains. Of the bacterial strains tested, the oxidatively stressed SH-SY5Y cells were most viable when cultured with *L. lactis* KC24-CM. *L. lactis* KC24-CM promoted the expression of brain-derived neurotrophic factor (*BDNF*) in the HT-29 cells. It also significantly increased *BDNF* expression and reduced the apoptosis-related *Bax/Bcl-2* ratio in the oxidatively stressed SH-SY5Y cells. Therefore, *L. lactis* KC24 is a potential psychobiotic for use in the functional food industry.

**Keywords** Probiotics · *Lactococcus lactis* · Gut-brain-axis · Neuroprotective effect · Conditioned medium

## Introduction

Neurobiological investigations into gut–brain crosstalk have revealed two-way communication that ensures gastrointestinal homeostasis, and have multiple effects on the neuronal activities of the brain that are associated with motivation and higher cognitive functions (Mayer, 2011; Foster and Neufeld, 2013). Two-way communication can be explained the autonomic nervous system, the enteric nervous system, the neuroendocrine system, and the immune system. These networks were affected by the gastrointestinal tract. The microbiota of the gut comprises over 5000 bacterial species (Carabotti et al., 2015). The modulatory effect of probiotics on the intestinal microbiota improves the intestinal barrier and mucosal immune response, thereby facilitating the treatment of microbe-related diseases (Holmes et al., 2011).

Reactive oxygen species (ROS) originate from both exogenous and endogenous sources, and can damage proteins, lipids, and other molecules. However, ROS are also essential for cellular function, and can be safely metabolized by an antioxidant mechanism (Phaniendra et al., 2015). The oxidative stress induced by ROS can contribute to pathologies including cancer (Valko et al., 2006), cardiovascular disease (Dhalla et al., 2000), diabetes (Giugliano et al., 1996), and aging (Sohal and Weindruch, 1996). The brain is particularly vulnerable to the effects of ROS because it has a high oxygen demand and an abundance of lipid cells (Leutner et al., 2001).

Several probiotics have been reported to have neuroprotective effects. *Bifidobacterium longum* 1714 had a positive effect on cognition in an anxious mouse (Savignac et al., 2015). Both *Lactobacillus rhamnosus* JB-1 and *Bacteroides fragilis* were able to activate intestinal afferent neurons *ex vivo*, and the polysaccharide A produced by *B.*

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*fragilis* completely represented the neuronal effects (Mao et al., 2013). *Lactobacillus reuteri* enhanced the emotionality of colonic neurons in naive rats by inhibiting  $\text{Ca}^{2+}$ -dependent potassium channels (Kunze et al., 2009). Probiotic mixtures containing *L. rhamnosus* and *Lactobacillus helveticus* ameliorated maternal separation-induced depression by normalizing corticosterone levels (Gareau et al., 2007).

Psychobiotics are of interest to the functional food industry because the gut microbiota affects mental health. Therefore, the aim of this study was to investigate the neuroprotective effects with regard to oxidative stress of lactic acid bacteria isolated from traditional Korean fermented foods.

## Materials and methods

### Bacterial strains and culture conditions

*Lactobacillus delbrueckii* KU200170 (Lim et al., 2020), *Lactobacillus plantarum* KU200661 (Lim et al., 2020), *Lactococcus lactis* KC24 (Lee et al., 2015), and *Lactobacillus rhamnosus* GG (Cell Biotech., Ltd., Gimpo, Korea) were used in the present study. The lactic acid bacteria were propagated and maintained in lactobacilli MRS medium (MRS; BD Biosciences, Franklin Lakes, NJ, USA) at 37 °C, and were then cultured in MRS broth for 24 h. The bacterial cells were centrifuged ( $12,000\times g$  at 4 °C for 10 min), and washed in phosphate-buffered saline (PBS; HyClone Laboratories Inc., Logan, UT, USA). The harvested bacteria were killed by heating at 121 °C for 15 min, and stored at – 80 °C until required.

### Cell line culture

HT-29 human intestinal cells (KCLB 30038) and SH-SY5Y human neuroblastoma cells (KCLB 22266) were obtained from the Korean Cell Line Bank (Seoul, Korea). The HT-29 and SH-SY5Y cells were grown in Roswell Park Memorial Institute medium 1640 (RPMI 1640; Gibco, Grand Island, NY, USA) and Dulbecco's modified Eagle's medium (DMEM; HyClone Laboratories Inc.), respectively. The cells were kept at 37 °C in 5%  $\text{CO}_2$ . All media were supplemented with 5% fetal bovine serum (HyClone) with 1% (v/v) penicillin–streptomycin (HyClone).

### Preparation of conditioned medium (CM) from HT-29 cells

The conditioned medium (CM) was prepared using HT-29 cells cultured with heat-killed probiotics (Park et al., 2017). The probiotics culture was centrifuged at  $14,000\times g$ , 4 °C

for 10 min, and the collected cell was washed twice with PBS. The harvested bacteria were then heated at 121 °C for 15 min to produce heat-killed lactic acid bacteria (LAB), which were stored at – 80 °C until required. The monolayered HT-29 cells were treated with two concentrations ( $1.0 \times 10^7$  and  $1.0 \times 10^8$  CFU/well) of the heat-killed LAB for 24 h. After incubation, the supernatant was collected by centrifugation ( $12,000\times g$  for 10 min) and filtered by syringe filtration (0.45- $\mu\text{m}$  membrane filter). This CM was stored at –80 °C until required.

### Protective effect of LAB-CM on SH-SY5Y neuroblastoma cells experiencing hydrogen peroxide-induced oxidative stress

The neuroprotective effect on the oxidatively stressed SH-SY5Y cells was determined by the following methods (Park et al., 2017). The SH-SY5Y cells ( $1.0 \times 10^5$  cells/well) were incubated to form a confluent monolayer on a 96-well plate. After pretreatment with 80  $\mu\text{L}$  of LAB-CM for 4 h, the SH-SY5Y cells in each well were treated with 20  $\mu\text{L}$  of 50  $\mu\text{M}$  hydrogen peroxide ( $\text{H}_2\text{O}_2$ ; Junsei Chemical Co. Ltd., Tokyo, Japan) for 20 h. After incubation, the media were removed, and the cells were incubated with 100  $\mu\text{L}$  of 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT; 5 mg/mL) for 1 h. The absorbance was measured using a microplate reader at 540 nm. The cell viability (%) was calculated using the Eq. (1):

$$\text{Cell viability (\%)} = \frac{\text{Absorbance of sample}}{\text{Absorbance of control}} \times 100 \quad (1)$$

### BDNF gene expression in the HT-29 cells

BDNF gene expression was determined using the following modified methods (Park et al., 2017). The HT-29 cells were inoculated into 6-well plates at a density of  $1.0 \times 10^6$  cells/well, and incubated to form a confluent monolayer. After incubation, the cells were treated with heat-killed LAB for 24 h. The total RNA of the HT-29 cells was isolated using a total RNA isolation kit (Qiagen, Hilden, Germany), and converted to complementary DNA (cDNA) using a cDNA synthesis kit (Thermo Fisher Scientific, Waltham, MA, USA). Brain-derived neurotrophic factor (BDNF) gene expression was confirmed by real-time polymerase chain reaction (PCR) analysis.

### Confirmation of *BDNF*, *Bax*, and *Bcl-2* gene expression in the SH-SY5Y cells experiencing $H_2O_2$ -induced oxidative stress by reverse transcription polymerase chain reaction (RT-PCR) analysis

The SH-SY5Y cells ( $1.0 \times 10^6$  cells/well) were seeded into a 6-well plate and incubated to form a confluent monolayer. After incubation, the cells were treated with 800  $\mu$ L of CM for 4 h. Subsequently, 200  $\mu$ L of  $H_2O_2$  (50  $\mu$ M) was added to induce oxidative stress for 20 h (Cheon et al., 2020; Park et al., 2017). The total RNA was isolated using an RNeasy Mini total RNA isolation kit and primers. The cDNA was manufactured using a cDNA synthesis kit (Thermo Fisher Scientific, Waltham, MA). The primers were those described by Cheon et al. (2020). The PCR data were evaluated using the  $2^{-\Delta\Delta CT}$  method, and are expressed as fold changes.

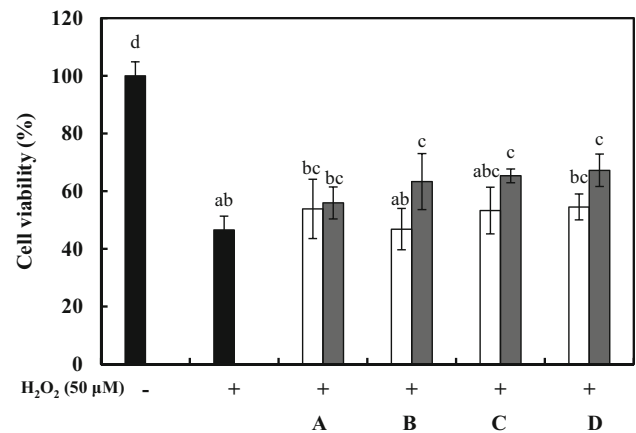
### Statistical analysis

All data are represented as the mean and standard deviation by three replicates. Analysis of variance (ANOVA) was used to determine significant differences. The mean values were utilized for the Duncan's multiple range test for post hoc verification ( $p < 0.05$ ).

## Results and discussion

### Protective effects of LAB-CM in SH-SY5Y cells experiencing $H_2O_2$ -induced oxidative stress

*L. delbrueckii* KU200170, *L. plantarum* KU200661, and *L. lactis* KC24 was reported on their probiotic properties included antimicrobial effect (Lee et al., 2015; Lim et al., 2020). Therefore, this study was dealt on neuroprotective effects using heat-killed probiotics. Cell viability was determined to evaluate the neuroprotective effects of LAB-CM in oxidatively stressed SH-SY5Y cells (Fig. 1). The positive control treated with  $H_2O_2$  exhibited  $47.46 \pm 1.38\%$  cell viability. The LAB-CM ( $1.0 \times 10^8$  CFU/mL) using *L. rhamnosus* GG, *L. delbrueckii* KU200170, *L. plantarum* KU200661, and *L. lactis* KC24 exhibited  $55.93 \pm 5.53\%$ ,  $63.28 \pm 5.08\%$ ,  $65.00 \pm 2.73\%$ , and  $67.23 \pm 5.59\%$  cell viability, respectively. LAB-CM concentrations of  $1.0 \times 10^7$  CFU/mL and  $1.0 \times 10^8$  CFU/mL resulted in similar or higher cell viability values to that in the positive control. These results confirm that in *Ruminococcus albus*-CM, the viability of oxidatively stressed SH-SY5Y cells depends on the concentration of colony-forming units ( $10^6$ – $10^8$  CFU/mL) (Park et al., 2017). The antioxidant effects of *L. rhamnosus* GG, and of *L. lactis* KC24 and LGG have been

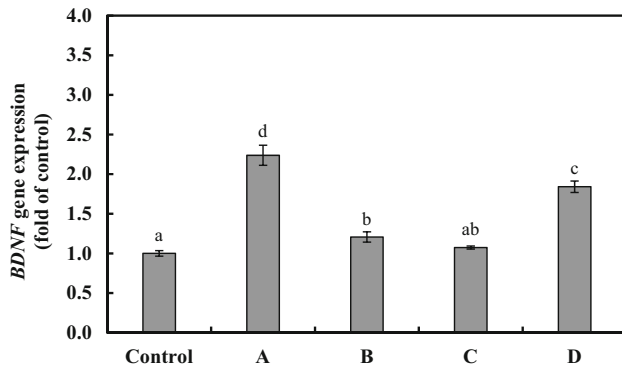


**Fig. 1** Protective effects of heat-killed lactic acid bacteria-conditioned medium (LAB-CM) on cell viability in SH-SY5Y cells experiencing oxidative stress induced by  $H_2O_2$  (50  $\mu$ M). Cell viability was evaluated using a 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay. A, *Lactobacillus rhamnosus* GG; B, *Lactobacillus delbrueckii* KU200170; C, *Lactobacillus plantarum* KU200661; D, *Lactobacillus lactis* KC24. The error bars indicate standard deviations from three independent experiments. The letters on each bar exemplify significant differences ( $p < 0.05$ ). open square,  $1.0 \times 10^7$  CFU/mL; filled square,  $1.0 \times 10^8$  CFU/mL

evaluated by determining ferric reducing ability of plasma (FRAP) values, and the inhibition of  $\beta$ -carotene and linoleic acid (Lee et al., 2015; Mishra et al., 2015). Therefore, the neuroprotective effects of *L. rhamnosus* GG and *L. lactis* KC24 may be influenced by their antioxidant effects.

### *BDNF* gene expression in HT-29 cells following treatment with heat-killed LAB

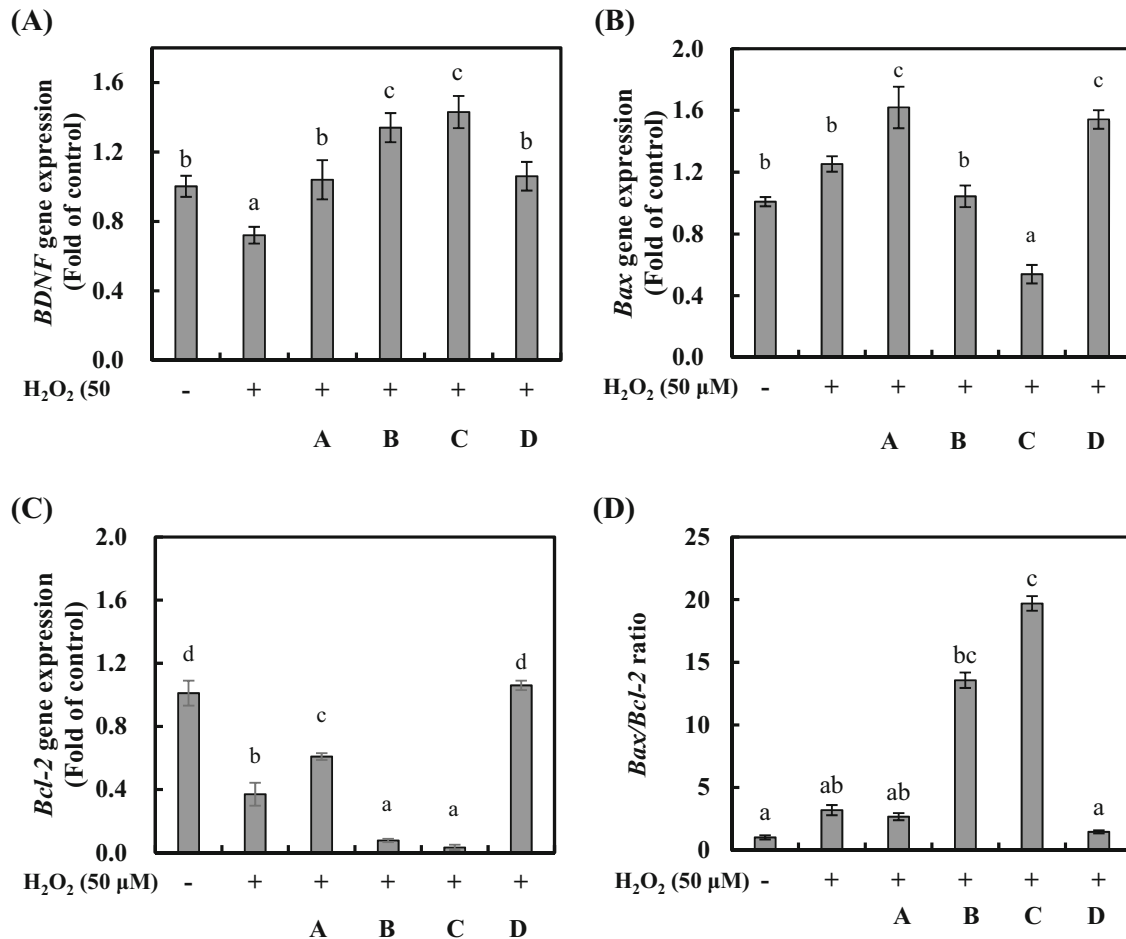
Microbial dysbiosis can reduce BDNF expression in the brain and gut (Bistoletti et al., 2019). BDNF expression is also involved in the homeostatic regulation of intestinal barrier integrity (Li et al., 2018; Yu et al., 2017). Therefore, probiotic bacterial strains may increase the expression of BDNF by modulating intestinal cells. The HT-29 cells were treated with  $1.0 \times 10^8$  CFU/mL of heat-killed LAB, and RT-PCR analysis was used to determine the level of *BDNF* gene expression from the extracted RNA (Fig. 2). The levels of *BDNF* gene expression in the HT-29 cells treated with heat-killed *L. delbrueckii* KU200170 and *L. plantarum* KU200661 were not significantly different to those in the PBS-treated controls. The levels of *BDNF* gene expression in the HT-29 cells treated with heat-killed *L. rhamnosus* GG and with *L. lactis* KC24 were  $2.23 \pm 0.13$ - and  $1.84 \pm 0.07$ -fold, respectively. Therefore, heat-killed *L. rhamnosus* GG and *L. lactis* KC24 may be useful for the treatment of intestinal conditions.



**Fig. 2** mRNA expression of the *BDNF* gene in HT-29 cells treated with heat-killed lactic acid bacteria (LAB). A, *Lactobacillus rhamnosus* GG; B, *Lactobacillus delbrueckii* KU200170; C, *Lactobacillus plantarum* KU200661; D, *Lactobacillus lactis* KC24. The control was treated with phosphate-buffered saline (PBS). The error bars indicate standard deviations from three independent experiments. The letters on each bar exemplify significant differences ( $p < 0.05$ )

### *BDNF* gene expression and anti-apoptosis in heat-killed LAB-CM-treated SH-SY5Y neuroblastoma cells experiencing $H_2O_2$ -induced oxidative stress

Oxidative stress was induced in the SH-SY5Y cells by treatment with  $H_2O_2$ , and BDNF production was determined using heat-killed LAB-CM (Fig. 3A). *BDNF* gene expression in the  $H_2O_2$ -treated positive control was reduced by  $0.72 \pm 0.05$ -fold compared to that in the negative control. The *BDNF* gene expression rates were higher in the CMs than in the negative control group. The *BDNF* gene expression rates in the SH-SY5Y cells treated with the LAB-CM ( $1.0 \times 10^8$  CFU/mL) using heat-killed *L. rhamnosus* GG, *L. delbrueckii* KU200170, *L. plantarum* KU200661, and *L. lactis* KC24 were  $1.04 \pm 0.11$ -,  $1.34 \pm 0.08$ -,  $1.43 \pm 0.09$ -, and  $1.05 \pm 0.08$ -fold, respectively. There was no difference in the level of BDNF expression between the intestinal cells and the



**Fig. 3** mRNA expression of the *BDNF* gene (A), *Bax* gene (B), *Bcl-2* gene (C), and the *Bax/Bcl-2* ratio (D) in oxidatively stressed SH-SY5Y cells treated with heat-killed lactic acid bacteria-conditioned medium (LAB-CM). Oxidative stress was induced by  $H_2O_2$  (50  $\mu$ M). A, *Lactobacillus rhamnosus* GG; B, *Lactobacillus delbrueckii*

KU200170; C, *Lactobacillus plantarum* KU200661; D, *Lactobacillus lactis* KC24. The error bars indicate standard deviations from three independent experiments. The letters on each bar exemplify significant differences ( $p < 0.05$ )

neuroblastoma cells. The administration of *L. rhamnosus* modulates the gut microbiota of zebrafish, and the expression of *BDNF* in the gut and brain (Borrelli et al., 2016).

*Bax* is a pro-apoptotic Bcl-2-family protein in cytosol and translocated to mitochondria upon induction of apoptosis (Hsu et al., 1997). The *Bax/Bcl-2* ratio is more important than the expression for each gene in determining susceptibility to apoptosis (Raisova et al., 2001).

The mRNA levels of the pro-apoptotic factor *Bax* was increased 1.25-fold under  $H_2O_2$  treatment compared with the negative control (Fig. 3B). CM using *L. rhamnosus* GG and *L. lactis* KC24 increased by 1.62- and 1.54-fold. While *L. delbrueckii* KU200170 and *L. plantarum* KU200661 was 1.04- and 0.53-fold. The mRNA levels of anti-apoptotic factor, *Bcl-2* was decreased 0.37-fold under  $H_2O_2$  treatment compared with the negative control (Fig. 3C). The LAB-CM using *L. rhamnosus* GG, *L. delbrueckii* KU200170, and *L. plantarum* KU200661 reduced 0.37-, 0.08-, and 0.04-fold, respectively. However, *L. lactis* KC24 maintained by 1.06-fold.

A decrease in apoptosis can be explained by a reduced *Bax/Bcl-2* ratio. The *Bax/Bcl-2* ratio improved significantly in the SH-SY5Y cells that had been co-cultured with  $H_2O_2$ , resulting in apoptosis. The *Bax/Bcl-2* ratios in the negative control and in the positive control that had received  $H_2O_2$  treatment were  $1.01 \pm 0.17$  and  $3.19 \pm 0.41$ , respectively (Fig. 3D). The *Bax/Bcl-2* ratios in the SH-SY5Y cells that had been treated with the LAB-CM ( $1.0 \times 10^8$  CFU/mL) using *L. rhamnosus* GG, *L. delbrueckii* KU200170, *L. plantarum* KU200661, and *L. lactis* KC24 were  $2.67 \pm 0.28$ ,  $13.56 \pm 0.62$ ,  $19.69 \pm 0.58$ , and  $1.45 \pm 0.58$ , respectively. Therefore, the *L. lactis* KC24 ratio was similar to that in the negative control. *L. lactis* KC24-CM exhibited the highest neuroprotective effect of all the samples in the oxidatively stressed SH-SY5Y cells. A probiotic cocktail comprising *L. rhamnosus* GG, *Bifidobacterium animalis* subsp. *lactis*, and *Lactobacillus acidophilus* prevented neurodegenerative diseases in mice (Srivastav et al., 2019). *R. albus*-CM significantly affected the *Bax/Bcl-2* ratio (by 1.3–1.5) in oxidatively stressed SH-SY5Y cells (Park et al., 2017). Therefore, *L. lactis* KC24 has the potential to prevent neurodegenerative diseases.

In conclusion, the antimicrobial, antioxidant, and anti-cancer effects of the probiotic *L. lactis* KC24 have been reported; the present study evaluated its neuroprotective effects in oxidatively stressed SH-SY5Y neuroblastoma cells. Treatment with *L. lactis* KC24 increased cell viability, increased *BDNF* gene expression, and reduced the *Bax/Bcl-2* ratio in oxidatively stressed SH-SY5Y cells. These results suggest it has potential as a prophylactic therapy for neuroprotection.

## Compliance with ethical standards

**Conflict of interest** The authors declare no conflict of interest.

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