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***Lactobacillus saerimneri* and *Lactobacillus ruminis*: novel human-derived probiotic strains with immunomodulatory activities**

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Abstract

Human-derived lactobacilli were isolated from fecal samples of healthy volunteers. Forty-six isolates from different volunteers were selected and investigated for their immunomodulatory properties. Conditioned medium from each isolate was assessed for its effect on tumor necrosis factor (TNF) production in lipopolysaccharide (LPS) - activated THP-1 monocytes. Of 46 *Lactobacillus* isolates, 12 significantly inhibited TNF production in varying magnitude. *Lactobacillus* strain TH58 displayed the most potent TNF - inhibitory activity (70% inhibition). In contrast, *Lactobacillus* strain TH14 exhibited immunostimulatory property by activating TNF production in THP-1 monocytes. *Lactobacillus* TH14 induced NF- κ B activation in the absence of LPS stimulation, whereas *Lactobacillus* TH58 had no effect on NF- κ B signaling, irrespective of LPS stimulation. Strain TH58 was identified as *Lactobacillus saerimneri* and strain TH14 as *Lactobacillus ruminis* by sequence analysis of the 16S rRNA gene. This is the first report of a human isolate of *L. saerimneri* with TNF inhibitory activity and *L. ruminis*, an indigenous species to humans, with TNF stimulatory activity. Our data suggest the potential use of these two strains as immunoprobiotic candidates.

Keywords

Lactobacillus saerimneri; *Lactobacillus ruminis*; immunoprobiotics; anti-inflammatory; immunostimulatory; tumor necrosis factor

Introduction

Lactobacillus spp. represent sources of beneficial organisms indigenous in the human intestinal tracts and strains of this genus are used as probiotics. Probiotics are defined as

“live microorganisms which when administered in adequate amounts confer a health benefit on the host” (FAO/WHO, 2001). Probiotics are attractive candidates for novel biological therapies because beneficial bacteria may be derived from commensal microorganisms and are generally recognized as safe (GRAS). Several lactobacilli of human origin are currently exploited commercially for various therapeutic applications.

Oral administration of probiotics may modulate cytokine profiles not only locally in the intestines, but also systemically (Solis-Pereyra *et al.*, 1997; Meyer *et al.*, 2007). Results pointing towards stimulation of both T_H1 and T_H2 responses have been reported in animals fed with probiotics (Tejada-Simon *et al.*, 1999; Maassen *et al.*, 2000; Cross, 2002; Perdigón *et al.*, 2002). Ability to modulate cytokine network and innate immunity by probiotic bacteria is strain dependent. Certain probiotics stimulated the production of inflammatory cytokines such as tumor necrosis factor (TNF) (Miettinen *et al.*, 1996; Solis-Pereyra *et al.*, 1997; Morita *et al.*, 2002; Cross *et al.*, 2004) while others effectively inhibit TNF production (Pena & Versalovic, 2003; Pena *et al.*, 2005; Chan Remillard & Ozimek, 2006; Lin *et al.*, 2008).

The objective of this study was to isolate and characterize human-derived *Lactobacillus* strains for potential use as immunoprobiotics. For this purpose, *Lactobacillus* spp. were isolated from fecal samples of healthy human volunteers and investigated for their immunomodulatory properties.

Materials and methods

Subjects and sample collection

Fecal samples were collected from forty-six healthy human volunteers. The volunteers were students at the Thai Red Cross College of Nursing, Chulalongkorn University, which included both males and females in the age group of 17–19 years. No subject had consumed antibiotics or yogurt for at least two months prior to sample collection. Each individual gave informed consent. Fresh fecal samples were obtained by rectal swabbing with sterile cotton swabs, subsequently inserted into a modified Cary-Blair transport medium (Atlas & Snyder, 1995). The samples were processed immediately upon receipt.

Isolation of *Lactobacillus*

To isolate *Lactobacillus*, fecal swabs were suspended in normal saline solution (NSS), serially diluted and applied to deMan-Rogosa-Sharpe (MRS) agar (Becton Dickinson, Sparks, MD) and incubated anaerobically (10% CO₂, 10% H₂ and 80% N₂) at 37°C for 48–72 h in an anaerobic chamber (Concept Plus, Ruskinn Technology, UK). Each isolate was tested for catalase activity using 3% hydrogen peroxide solution and catalase-negative isolates were gram stained (Cappuccino & Sherman, 2005). All catalase-negative, gram-positive rods were tested for vancomycin (VA) resistance using a modified Kirby-Bauer disk diffusion test (Pena *et al.*, 2004). Briefly, isolates were resuspended in NSS to a 0.5 McFarland standard and swabbed onto MRS agar. VA impregnated disks (5 µg; Becton Dickinson) were applied to the bacterial cultures and incubated anaerobically at 37°C for 24–48 h. Isolates displaying an inhibition zone of >15 mm were considered susceptible. All

isolates that were gram-positive, regular rods or short rods, catalase-negative and vancomycin resistant were maintained as frozen cultures in MRS broth with 20% glycerol at -80°C for experimental use.

Preparation of *Lactobacillus* conditioned media

Lactobacillus conditioned media were prepared as previously described (Lin *et al.*, 2008). Human-derived *Lactobacillus* isolates were cultured anaerobically in MRS medium at 37°C for 24 h. Overnight cultures were diluted to an OD_{600} of 0.1 in MRS media and incubated in anaerobic condition for an additional 24 h. Condition media were collected by centrifugation at $4000 \times g$ for 10 min at 4°C , filter-sterilized using a $0.22 \mu\text{m}$ filter (Millipore, Bedford, MA) and concentrated by speed-vacuum drying (Speed vacuum DNA 110, Savant, NY). The residual pellet was resuspended in an equal volume of serum-free RPMI medium 1640 (Gibco-Invitrogen, Carlsbad, CA) and stored at -20°C .

THP-1 monocyte cell culture and bioassay for TNF activity

Cell culture and *in vitro* bioassays were performed as previously described (Lin *et al.*, 2008) with modification. Briefly, THP-1 human monocytic cell lines (ATCC TIB- 202, Manassas, VA) were maintained in RPMI medium 1640 supplemented with 10% heat-inactivated fetal bovine serum (FBS; Gibco-Invitrogen) in 96-well flat-bottomed tissue culture plates (Falcon; Becton Dickinson) and incubated at 37°C in a humidified 5% CO_2 incubator. THP-1 cells (2.5×10^5 cells mL^{-1}) were incubated with conditioned media (5% v/v) alone or in combination with purified LPS (100 ng mL^{-1}) from *Escherichia coli* serotype O127:B8 (Sigma, St. Louis, MO) for 3.5 h at 37°C . The assay condition was previously optimized, i.e. 5% (v/v) of conditioned medium was used to minimize the confounding effects of MRS medium. Supernatants were collected from individual wells by centrifugation at 4°C and assayed for TNF.

Assay for cell viability and apoptosis

Cell viability was determined by Trypan Blue dye exclusion assay (Gibco-Invitrogen). Cell suspension was mixed with trypan blue which was excluded by viable cells but stained dead cells. The cells were counted on a hemocytometer under a phase contrast microscope. Percentage of viable cells was determined from the ratio of viable cells over total cells. In addition to trypan blue staining, TH58 conditioned medium treated cells and controls were stained with Annexin V-FITC and Propidium Iodide (PI) by using BD Pharmingen Annexin V-FITC Apoptosis detection kit I (BD Biosciences Pharmingen, San Diego, CA). Controls included THP-1 cells in RPMI and MRS treated cells. This staining method allowed the detection of viable, apoptotic and necrotic cells. Annexin V binds to cells that expose phosphatidylserine to the external cellular environment, a characteristic of early apoptotic cells. PI stains cells with loss of membrane integrity which accompanies the latest stage of cell death resulting from either apoptotic or necrotic processes. Viable cells were AnnexinV-FITC and PI negative. Early apoptotic cells were AnnexinV-FITC positive and PI negative. Cells that were in late apoptosis or already dead were both AnnexinV-FITC and PI positive. The percentage of viable cells was analyzed on FACSCalibur flow cytometer (BD Biosciences, San Jose, CA) using the Cell Quest Pro software.

Assay for cell proliferation

An increase in the number of THP-1 cells was determined by microscopic count of cells after staining with trypan blue dye. The cells were counted in four 1-mm square grids on a hemocytometer. This assay was performed in TH14 conditioned medium treated cells and controls which included THP-1 cells in RPMI and MRS treated cells. The number of viable cells and total cells in each sample were calculated. The statistical differences were evaluated using the Student's *t* test with a one-tailed distribution. A *p* value of 0.05 was considered statistically significant.

TNF measurement

TNF production in THP-1 cell culture supernatants was measured with human TNF-specific sandwich quantitative enzyme-linked immunosorbent assay (DuoSet, R&D Systems, Minneapolis, MN) according to the manufacturer's instructions. Recombinant human TNF was used as standard. Absorbance was measured at 450 nm using a Spectramax 340PC (Molecular Devices Corporation, CA). Cytokine concentrations were quantified from standard curve and expressed as pg mL^{-1} of culture medium. Results were reported as mean value with standard deviations (SD) of three different experiments in triplicate. The statistical differences were evaluated using the Student's *t* test with a one-tailed distribution. A *p* value of 0.05 was considered statistically significant.

Nuclear extraction from THP-1 monocytes and NF- κ B measurement

THP-1 cells (2.5×10^5 cells mL^{-1}) were incubated with conditioned media (5% v/v) in the presence or absence of purified LPS (100 ng mL^{-1}) for 30 min. The THP-1 cells were collected and stored on ice before nuclear extraction. Nuclear extracts were isolated using the Nuclear Extract kit (Active Motif, Carlsbad CA, USA) according to the manufacturer's instructions. Briefly, THP-1 cells were washed in ice-cold PBS, resuspended in 200 μL of hypotonic buffer and incubated on ice for 15 min. Detergent (25 μL) was added and each sample was vortexed. The nuclear pellets were collected by centrifugation and resuspended in ice-cold lysis buffer (20 μL) containing protease inhibitor. The samples were vortexed 8 times, each for 30 s, at highest setting and subjected to centrifugation at 14,000 \times g at 4°C for 10 min. The nuclear extracts were stored at -80°C. The protein content of nuclear extracts were quantified using the microplate procedure of BCA™ Protein Assay kit (Pierce Biotechnology, Rockford, IL) as per the manufacturer's instruction. The levels of nuclear factor NF- κ B in THP-1 nuclear extracts were measured by ELISA with TransAM™ NF- κ B p65 Transcription Factor Assay kit (Active Motif, Carlsbad, CA) according to the manufacturer's instruction.

16S rRNA gene sequencing

The species of selected bacterial isolates was determined using 16S rRNA gene sequencing as done previously (Pena *et al.*, 2004). Genomic DNA was extracted from overnight cultures using the Wizard® Genomic DNA Purification kit (Promega Corporation Madison, WI, USA) following the manufacturer's instructions. The 16S rRNA gene was amplified by PCR with primers 16S-8F (5'-AGA GTT TGA TCY TGG YTY AG-3') and 16S-1541R (5'-AAG GAG GTG WTC CAR CC-3') (Pena *et al.*, 2004) in a reaction containing 200 ng DNA, 50

pmol of each primer, 1.25 U AmpliTaq DNA polymerase (Applied Biosystems, Foster City, CA), 2.5 mM dNTPs, 1X AmpliTaq Reaction Buffer and 75 mM MgCl₂. The following thermal cycling conditions were used: an initial denaturation of 94°C for 2 min; 35 cycles of denaturation at 94°C for 30 s, annealing at 57°C for 30 s, extension at 72°C for 1 min; and a final extension at 72°C for 5 min. PCR products were purified by Qiagen MinElute PCR Purification kit (Qiagen Inc., Valencia, CA) and sequenced by SeqWright DNA Technology Services (Houston, TX) using primers 16S 8F and 16S 1541R. Approximately 1,400 nucleotides were analyzed by using the sequence match program at the Ribosomal Database Project –II (<http://rdp.cme.msu.edu/>).

Results

Isolation of *Lactobacillus* isolates

Bacterial colonies isolated from 46 healthy human volunteers were selected for genus *Lactobacillus* by presumptive tests including gram stain, catalase production and vancomycin susceptibility. Four hundred and thirty-seven isolates were gram-positive, catalase-negative and vancomycin resistant, making them candidates for *Lactobacillus* spp. Cell morphology in each isolate varied from long and slender rods, straight rods to bent rods, sometimes shot rods to coccobacilli; arranged in single, in pairs or short chains. Some isolates exhibited bipolar staining or internal granulations. The most frequently found colonies were small to medium sizes (1–2 mm) with white, circular, smooth and convex colonial morphologies. Most isolates grew well under anaerobic condition. Most *Lactobacillus* isolates were obligate anaerobes while some isolates were facultative anaerobes. One isolate from each volunteer was randomly selected and characterized further for immunomodulatory effects on THP-1 human monocytes.

Immunomodulatory effects of *Lactobacillus* isolates

Forty-six *Lactobacillus* isolates were tested for their ability to modulate TNF production in LPS-activated THP-1 monocytic cells. Conditioned media from 12 *Lactobacillus* isolates significantly inhibited TNF production in LPS-activated THP-1 human monocytes with different magnitude. *Lactobacillus* TH58 showed the strongest TNF inhibitory activity of 70% ($p < 0.001$) in LPS-activated THP-1 monocytes (Fig. 1). In contrast, conditioned medium from *Lactobacillus* TH14 stimulated TNF production ($p < 0.01$) (Fig. 1). Ability of *Lactobacillus* TH 58 and TH14 to modulate TNF production in the presence or absence of LPS was investigated. The results confirmed the TNF inhibitory effect of *Lactobacillus* TH58 and TNF stimulatory effect of *Lactobacillus* TH14 conditioned medium (Fig. 2). Results also indicated that in the absence of LPS activation, *Lactobacillus* TH14 conditioned medium stimulated TNF production ($p < 0.001$) in THP-1 human monocytes, while *Lactobacillus* TH58 conditioned medium did not (Fig. 2).

The inhibition of TNF production by TH58 conditioned medium did not have cytotoxic effects on the THP-1 cells as determined by Trypan Blue dye exclusion assay and Annexin V- Propidium Iodide staining and analysis by flow cytometer (Fig. 3). The percentage of viable cells in each sample was more than 90% when determined by both methods. The effect of TH14 conditioned medium on proliferation of THP-1 cells was examined and it

was found that the stimulation of TNF production by TH14 conditioned medium did not result from the proliferation of THP-1 cells. The total number of TH14 conditioned medium treated cells was not significantly different from untreated controls and the percentage of viable cells in each sample was found to be more than 90%. As shown in Fig. 4, the number of viable cells in TH14 conditioned medium treated samples was not significantly different from the untreated controls.

Pro-inflammatory mediators such as LPS induce TNF production via activation of the NF- κ B signaling pathway (Ulivitch & Tobia, 1999; Aderem & Ulevitch, 2000). We speculated that *Lactobacillus* TH14 and TH58 mediated their pro- and anti-inflammatory effects by modulating the NF- κ B signaling pathway. As expected, *Lactobacillus* TH14 conditioned medium stimulated phosphorylation of NF- κ B in THP-1 cells in the absence of LPS stimulation ($p < 0.001$) whereas *Lactobacillus* TH58 conditioned medium did not. However, *Lactobacillus* TH14 and TH58 conditioned media did not affect the phosphorylation of NF- κ B in LPS-activated THP-1 cells ($p > 0.05$) (Fig. 5).

Genotypic identification of *Lactobacillus* isolates with immunomodulatory properties

Sequence analysis of the amplified 16S rRNA gene products allowed the identification of *Lactobacillus* isolates (Table 1). Seven out of 12 TNF-inhibitory isolates were identified as *Lactobacillus plantarum* or *Lactobacillus pentosus*, three as *Lactobacillus plantarum*, one as *Lactobacillus salivarius* and TH58 as *Lactobacillus saerimneri*. *Lactobacillus* TH14 was identified as *Lactobacillus ruminis*.

Discussion

Lactobacillus isolates derived from humans are optimal candidates for exhibiting beneficial effects in human beings (FAO/WHO, 2001). Since probiotic strains have been reported to possess anti-inflammatory and immunostimulatory effects, an effort was made in this study to isolate human- derived *Lactobacillus* strains with these potential probiotic properties. Several *Lactobacillus* species produce immunoregulatory factors, known as immunomodulins, which may modulate cytokine production (Guarner *et al.*, 2006). Among the inflammatory cytokines, TNF is important in mediating inflammation and immune function. In the current study, human-derived *Lactobacillus* isolates modulated TNF production via a contact- independent manner thereby suggesting the possibility that these isolates may secrete immunomodulins. Most of the immunomodulatory *Lactobacillus* isolated in this study inhibited TNF production in varying magnitude and *Lactobacillus* TH58 had the strongest TNF-inhibitory activity.

Similar results have been demonstrated in previous investigations. Report indicated that certain lactic acid bacteria secreted anti-inflammatory metabolites that inhibited TNF production and suppressed NF- κ B activation in LPS-activated THP-1 cells (Menard *et al.*, 2004). Conditioned media from *Lactobacillus rhamnosus* GG (Pena & Versalovic, 2003) and other murine lactobacilli significantly inhibited TNF production in LPS-activated murine macrophages (Pena *et al.*, 2004). The supernatants of nine strains of probiotic *Lactobacillus* were able to inhibit TNF secretion in murine macrophages (Chan Remillard & Ozimek, 2006). *Lactobacillus reuteri* strain ATCC PTA 6475 was able to inhibit TNF

production by LPS-activated THP-1 cells and human monocyte-derived macrophages from pediatric patients with Crohn's disease by a contact-independent mechanism (Lin *et al.*, 2008). In addition, the anti-inflammatory property of *Lactobacillus reuteri* ATCC PTA 6475 was shown to be strain-specific as the conditioned media of *Lactobacillus reuteri* strains CF48-3A and ATCC 55730 did not inhibit TNF production in those monocytic cells. Our data in a separate study of Panpetch, 2008 (Thesis, Graduate school, Chulalongkorn university) also supported the strain-specific nature of *Lactobacillus* sp. We found that condition media of some isolates of *Lactobacillus gasseri* and *Lactobacillus salivarius* cultivated from gastric biopsies of peptic ulcer patients could inhibit TNF production in LPS-activated THP-1 cells. On the contrary, some isolates of these species could not inhibit TNF production. These data supported the possibility of the secretion of immunomodulin(s) by *Lactobacillus* TH58 since it is unlikely that specific strains of *Lactobacillus reuteri*, some isolates of *Lactobacillus gasseri* and *Lactobacillus salivarius* metabolize MRS media to yield products that can inhibit TNF production, whereas specific strain or isolates of the same species do not metabolize MRS by the same pathway.

In this study, we also found that the anti-inflammatory activity (TNF inhibition) of *Lactobacillus* TH58 was most potent in conditioned media collected at 48 h (80% inhibition) compared to conditioned media collected at 24 h (70% inhibition) (data not shown). Based on growth characteristics of TH58, 24–28 h time point represented late logarithmic phase, > 28–40 h represented stationary phase and > 48 h time point represented early decline growth phase. It was possible that during the time point of stationary phase until the early decline phase, *Lactobacillus* TH58 increased secretion of putative immunomodulin(s) into the supernatant. Similar results by Chan Remillard and Ozimek indicated that supernatants from milk fermented by nine different strains of *Lactobacillus* showed increased TNF inhibitory activity over time (Chan Remillard & Ozimek, 2006).

Previous studies demonstrated that probiotic bacteria may stimulate the production of certain pro-inflammatory cytokines including TNF (Miettinen *et al.*, 1996; Solis-Pereyra *et al.*, 1997; Morita *et al.*, 2002; Cross *et al.*, 2004; Meyer *et al.*, 2007). Results from the current study indicated that the conditioned medium of *Lactobacillus* TH14 exhibited immunostimulatory activity and stimulated TNF production in the absence of LPS. Similar immunostimulatory capabilities were also demonstrated in *Lactobacillus reuteri* strains CF48-3A and ATCC 55730 (Lin *et al.*, 2008).

NF- κ B transcription factor is known to regulate inflammatory mediators and cytokines (Bibiloni *et al.*, 2005) including TNF, IL-12, IL-1, IL-6 and IL-8 (Aderem & Ulevitch, 2000; Egan & Toruner, 2006). Several reports indicated that certain *Lactobacillus* species diminish pro-inflammatory cytokine production via inhibiting NF- κ B activation (Bai *et al.*, 2004; Ma *et al.*, 2004; Menard, 2004; Guarner *et al.*, 2006; Iyer *et al.*, 2008). As expected, *Lactobacillus* TH14 (TNF stimulatory strain) induced NF- κ B activation in the absence of LPS stimulation. Whereas, *Lactobacillus* TH58 (TNF inhibitory strain) had no effect on NF- κ B signaling pathway irrespective of LPS stimulation.

TNF production is regulated at different levels such as transcription, translation and secretion (Watkins *et al.*, 1999; Papadakis & Targan, 2000). Kim *et al.* demonstrated that *L.*

rhamnosus GG- and *L. rhamnosus* GR-1 suppressed LPS- induced TNF production in THP-1 cells via paracrine route involving granulocyte-colony stimulating factor (G-CSF). The suppression of TNF production by G-CSF was mediated through activation of signal transducer and activator of transcription (STAT)-3, subsequently inhibiting activation of c-Jun-N-terminal kinases (JNKs) in THP-1 cells (Kim *et al.*, 2006). In another report, Lin *et al.* demonstrated the suppression of TNF in Mono-Mac-6 cells and THP-1 cells via down-regulation of activator protein-1 (AP-1) pathway without affecting NF- κ B signaling (Lin *et al.*, 2008). Results of these investigations suggested that *Lactobacillus* TH58-mediated suppression of TNF production may not be mediated via NF- κ B signaling pathway. It is possible that *Lactobacillus* TH58 may mediate its anti-inflammatory effects via inhibition of c-Jun-N-terminal kinase (JNK) activation (Kim *et al.*, 2006) or suppression of AP-1 transcription factor (Lin *et al.*, 2008). However, these speculations require further investigation.

Lactobacillus TH58 (TNF-inhibitory strain) was identified as *Lactobacillus saerimneri* which was first isolated from a pig in 2004 (Pedersen & Roos, 2004), and until now not identified in humans. *Lactobacillus* TH14, TNF-stimulatory strain, was identified as *Lactobacillus ruminis* which is known to be a dominant indigenous species in the digestive tracts of humans and pigs (Reuter, 2001; Yin & Zheng, 2005).

In conclusion, the results of the present study demonstrated that *Lactobacillus* isolates exhibited different immunomodulatory properties. This is the first report of human-derived *L. saerimneri* strain with anti-inflammatory property and *L. ruminis*, the indigenous gut microbiota, with immunostimulatory property.

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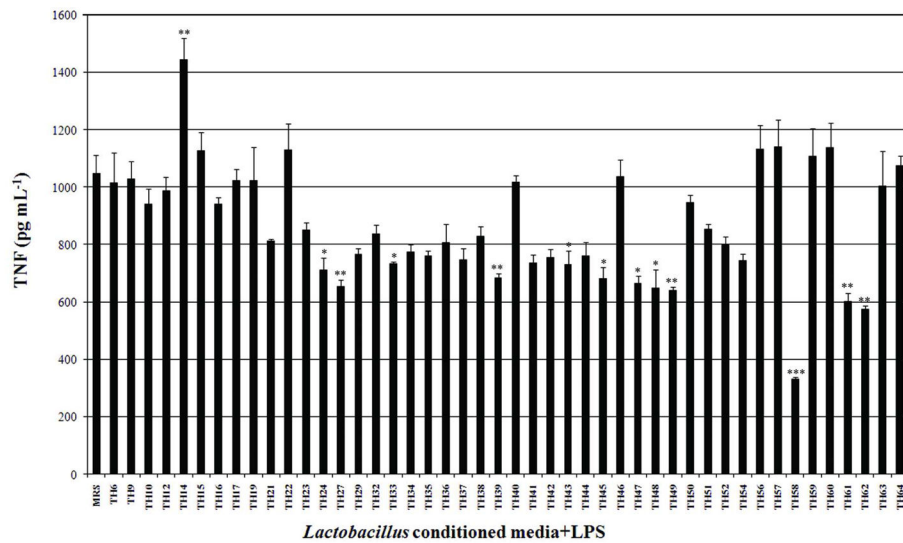


Fig. 1. Effects of *Lactobacillus* conditioned media on TNF production in LPS-activated THP-1 cells. THP-1 cells were coincubated with *Lactobacillus* conditioned media and *E. coli* - derived lipopolysaccharide. TNF secretions were determined by human TNF ELISA. The results were from three independent experiments in triplicate. MRS: deMan-Rogosa-Sharpe, media control; Asterisks denote significantly different from media control; ***, $p < 0.001$; **, $p < 0.01$; *, $p < 0.05$; Error bars indicate standard deviations.

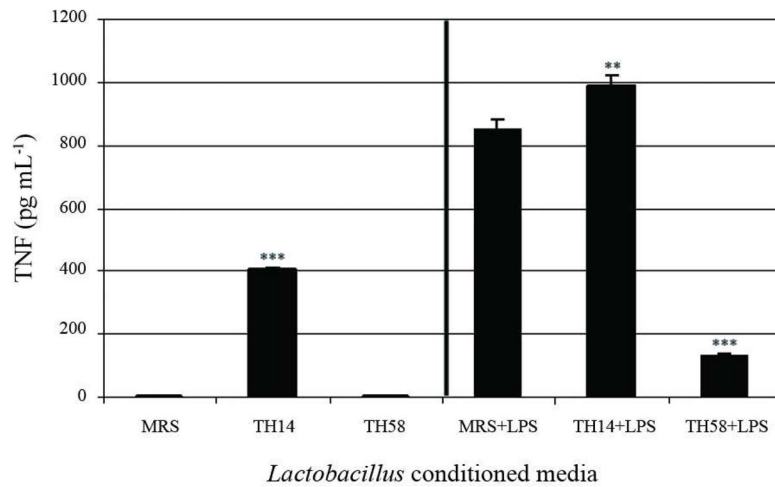


Fig. 2. Immunomodulatory effects of *Lactobacillus* TH14 and TH58 conditioned media on TNF production in THP-1 cells. Bioassays for TNF secretion were performed without or with LPS activation. The results represent three independent experiments. MRS: deMan-Rogosa-Sharpe, media control; Asterisks denote significantly different from media control; ***, $p < 0.001$; **, $p < 0.01$; Error bars indicate standard deviations.

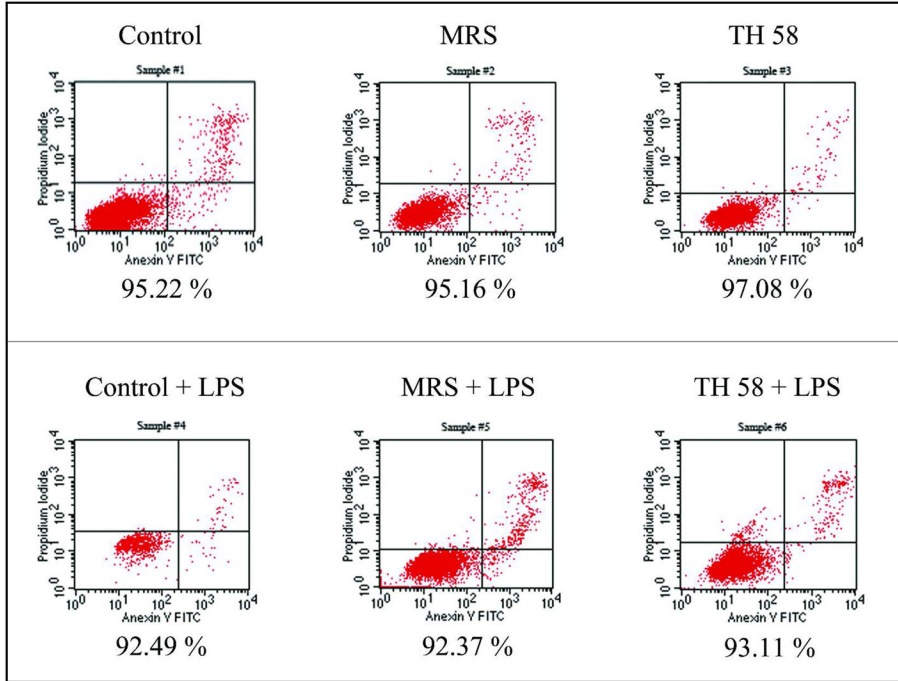


Fig. 3. The viability of THP-1 cells determined by staining with Annexin V-FITC and Propidium Iodide (PI) followed by flow cytometric analysis. Viable cells are negative for both Annexin V-FITC and PI. Cells that are in early apoptosis are Annexin V-FITC positive and PI negative. Cells that are in late apoptosis or already dead are both Annexin V-FITC and PI positive. The percentages of cell viability under each plot are representative of three experiments. Control: RPMI cell culture medium; MRS: deMan-Rogosa-Sharpe, bacterial media control; TH58: *Lactobacillus* TH58 conditioned medium.

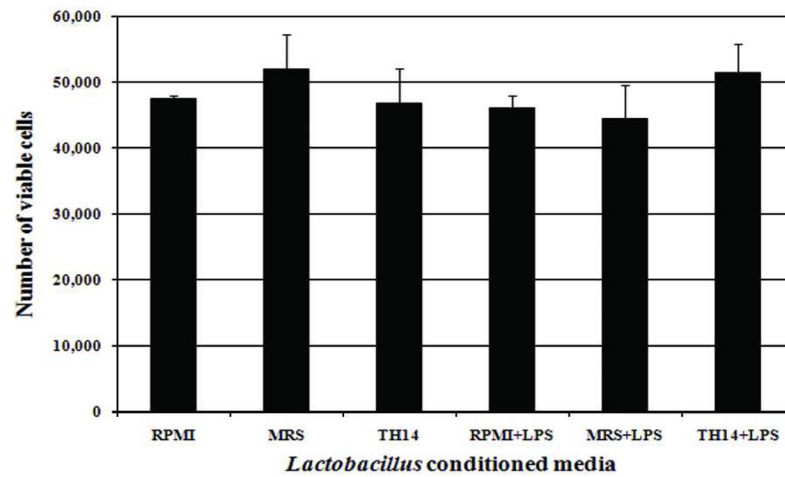


Fig. 4. Effect of *Lactobacillus* TH 14 conditioned medium on the proliferation of THP-1 cells. THP-1 cells were incubated with TH14 conditioned medium in the absence or presence of LPS. Cell suspensions were stained with trypan blue, counted on a hemocytometer and the number of viable cells in each sample was calculated. RPMI: cell culture medium; MRS: deMan-Rogosa-Sharpe, bacterial media control; TH14: *Lactobacillus* TH14 conditioned medium. The number of viable cells is shown as mean value with standard deviation from three different experiments.

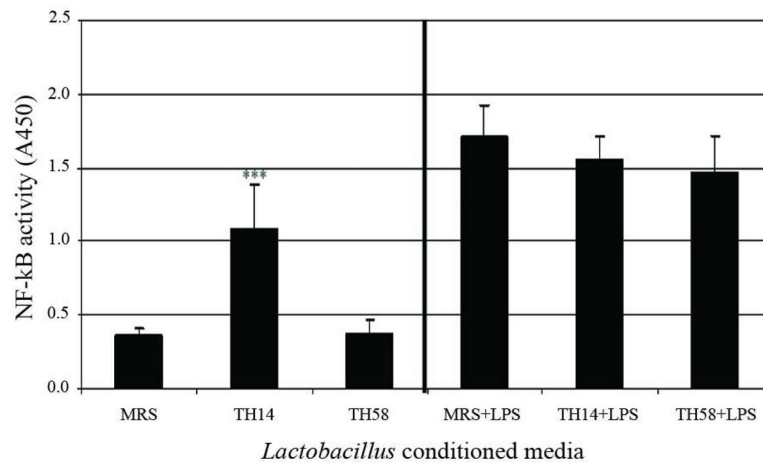


Fig. 5. Effect on NF- κ B activation by *Lactobacillus* TH14 and TH58 in THP-1 cells. THP-1 cells were incubated with TH14 or TH58 conditioned media in the absence or presence of LPS. Nuclear extracts were prepared and assayed for NF- κ B levels. The results were from two independent experiments in triplicate. MRS: deMan-Rogosa-Sharpe, media control; Asterisk denotes significantly different from media control; ***, $p < 0.001$; Error bars indicate standard deviations.

Table 1Identification of *Lactobacillus* isolates with immunomodulatory properties by 16 S rRNA gene sequencing

Isolate	Match organism	Identity (%)
TH14	<i>Lactobacillus ruminis</i>	99.5
TH24	<i>Lactobacillus plantarum</i> or <i>Lactobacillus pentosus</i>	100
TH27	<i>Lactobacillus plantarum</i> or <i>Lactobacillus pentosus</i>	100
TH33	<i>Lactobacillus salivarius</i>	100
TH39	<i>Lactobacillus plantarum</i> or <i>Lactobacillus pentosus</i>	100
TH43	<i>Lactobacillus plantarum</i> or <i>Lactobacillus pentosus</i>	100
TH45	<i>Lactobacillus plantarum</i> or <i>Lactobacillus pentosus</i>	100
TH47	<i>Lactobacillus plantarum</i>	100
TH48	<i>Lactobacillus plantarum</i>	100
TH49	<i>Lactobacillus plantarum</i>	100
TH58	<i>Lactobacillus saerimneri</i>	98
TH61	<i>Lactobacillus plantarum</i> or <i>Lactobacillus pentosus</i>	100
TH62	<i>Lactobacillus plantarum</i> or <i>Lactobacillus pentosus</i>	99.5