

## Probiotic intervention has strain-specific anti-inflammatory effects in healthy adults

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**Supported by** The Research Council for Health of the Academy of Finland, and Valio Research Centre

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Received: December 11, 2007 Revised: February 21, 2008

*freudenreichii* ssp. *shermanii* JS (PJS) or a placebo drink for 3 wk. Venous blood and saliva samples were taken at baseline and on d 1, 7 and 21. Fecal samples were collected at baseline and at the end of intervention.

**RESULTS:** The serum hsCRP expressed as the median AUC<sub>0-21</sub> (minus baseline) was 0.018 mg/L in the placebo group, -0.240 mg/L in the LGG group, 0.090 mg/L in the Bb12 group and -0.085 mg/L in the PJS group ( $P = 0.014$ ). *In vitro* production of TNF- $\alpha$  from *in vitro* cultured peripheral blood mononuclear cells (PBMC) was significantly lower in subjects receiving LGG vs placebo. IL-2 production from PBMC in the Bb12 group was significantly lower compared with the other groups.

**CONCLUSION:** In conclusion, probiotic bacteria have strain-specific anti-inflammatory effects in healthy adults.

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**Key words:** Probiotic; Highly sensitive C-reactive protein; Cytokine; Inflammation; Immune response; Mononuclear cells

**Peer reviewer:** Francesco Costa, Dr, Dipartimento di Medicina Interna-U.O. di Gastroenterologia Università di Pisa-Via Roma, 67-56122-Pisa, Italy

Kekkonen RA, Lummela N, Karjalainen H, Latvala S, Tynkkynen S, Järvenpää S, Kautiainen H, Julkunen I, Vapaatalo H, Korpela R. Probiotic intervention has strain-specific anti-inflammatory effects in healthy adults. *World J Gastroenterol* 2008; 14(13): 2029-2036 Available from: URL: <http://www.wjgnet.com/1007-9327/14/2029.asp> DOI: <http://dx.doi.org/10.3748/wjg.14.2029>

### Abstract

**AIM:** To evaluate the effects of three potentially anti-inflammatory probiotic bacteria from three different genera on immune variables in healthy adults in a clinical setting based on previous *in vitro* characterization of cytokine responses.

**METHODS:** A total of 62 volunteers participated in this randomized, double-blind and placebo-controlled parallel group intervention study. The volunteers were randomized to receive a milk-based drink containing either *Lactobacillus rhamnosus* GG (LGG), *Bifidobacterium animalis* ssp. *lactis* Bb12 (Bb12), or *Propionibacterium*

### INTRODUCTION

Probiotics are defined as living microorganisms that have beneficial effects on human health<sup>[1]</sup>. The immunomodulatory effects of probiotics have mostly been studied in certain disease conditions, such as allergies<sup>[2]</sup> and inflammatory diseases<sup>[3,4]</sup>, though the general, healthy population mostly consumes probiotics. The immunomodulatory effects of probiotics in healthy populations have not been fully established and only a few randomized, double blind, placebo-controlled studies have addressed this question<sup>[5-9]</sup>. Also, there are few studies where the effects of different

probiotic bacteria have been compared within the same clinical setting. Isolauri *et al*<sup>[10]</sup> and Viljanen *et al*<sup>[11]</sup> have compared the effects of two different probiotics or a probiotic mixture with placebo in allergic infants. Schiffrin *et al*<sup>[12]</sup> and Gill *et al*<sup>[13]</sup> evaluated the effects of two different probiotics in healthy adults, but these studies did not have a placebo group. Efforts trying to compare the *in vitro* results of one probiotic to its results in an *in vivo* setting are even more scarce and are at the moment limited to comparisons between *in vitro* and experimental animal studies<sup>[14-16]</sup>.

In our previous studies, we have characterized the capacity of potentially probiotic bacteria to induce cytokine production in human leukocyte cell culture and found that probiotic bacteria direct immune responses to either the Th1 type or the anti-inflammatory direction in a manner specific to the bacterial genera<sup>[17]</sup>. Based on these findings we selected probiotic bacteria from three different genera for the present study and compared their effects on immune variables in healthy adults in a 3-wk intervention trial.

## MATERIALS AND METHODS

### Subjects

The subjects were healthy adults recruited by an advertisement in the Helsinki area. The inclusion criteria were to be healthy (no chronic illnesses), to exercise regularly (at least three times per week), and to not be participating in any other clinical trials. The exclusion criteria was comprised of milk allergies (due to the nature of the study products), use of antibiotics during the two months before the study, acute gastrointestinal disorders during the two months before the study, gastrointestinal diseases and related medications, pregnancy, and lactation. Before entering the study, the subjects gave their written informed consent. The study protocol was approved by the Ethics Committee of the Hospital District of Helsinki and Uusimaa.

A total of 68 subjects were recruited for the study. Six subjects withdrew from the study during the run-in period and were not included in the analysis. The mean age for the subjects was 44 years (range 23-58) and their mean BMI was 24 kg/m<sup>2</sup> (range 18-30). Of these 62 subjects (45 females, 17 males), one subject withdrew from the study due to a back injury after two study visits and one subject due to an antibiotic treatment after four study visits. These two subjects were included in the statistical analysis.

### Study design and intervention

The study was a randomized, double-blind and placebo-controlled parallel group intervention study. Prior to the intervention period, there was a 3-wk run-in period during which no probiotic-containing products were allowed. Thereafter the subjects received either *Lactobacillus rhamnosus* GG (*n* = 13), *Bifidobacterium animalis* ssp. *lactis* Bb12 (*n* = 16), *Propionibacterium freudenreichii* ssp. *shermanii* JS (*n* = 17) or placebo (*n* = 16) drink for 3 wk. After the intervention period, subjects were followed up for 3 wk without any study drink. A list of probiotic-containing products was given to the subjects, and they were asked

not to consume any other probiotic-containing products at any time during the study.

### Study products

The subjects were advised to consume a 250 mL milk-based fruit drink daily for 3 wk containing either: *L. rhamnosus* GG (ATCC 53103) (LGG) bacteria, on average  $6.2 \times 10^7$  cfu/mL (daily dose of  $1.6 \times 10^{10}$  cfu); *B. animalis* ssp. *lactis* Bb12 (Bb12) bacteria,  $1.4 \times 10^8$  cfu/mL (daily dose of  $3.5 \times 10^{10}$  cfu); *P. freudenreichii* ssp. *shermanii* JS (DSM 7067) (PJS) bacteria,  $1.3 \times 10^8$  cfu/mL (daily dose of  $3.3 \times 10^{10}$  cfu); or a placebo drink without any probiotic bacteria. The subjects consumed the study drinks throughout the 3-wk intervention period after the baseline blood sampling. The amount of probiotic bacteria in the study drinks was measured right after packaging and after 3 wk. The appearance and taste of the study drinks were the same.

### Blood samples

Venous blood samples from the antecubital vein were taken at baseline, on 1, 7 and 21 d, and after the 3-wk follow-up period after an overnight fast. The samples were taken into standard serum tubes and EDTA tubes, centrifuged, and the plasma/serum was collected and stored at -20°C for further analyses. Three EDTA tubes were used in the purification of PBMC.

**Blood cells and immunoglobulins:** Blood cells (leukocytes, monocytes, and lymphocytes) from all time points were determined using an electronic counter (Coulter MAXM Hematology Analyzer, Beckman Coulter, Fullerton, CA, USA). Immunoglobulins (IgA, IgG and IgM) from all time points were measured by immunoturbidimetric method with Tina-quant Roche/Hitachi System reagent using a Roche Hitachi 912 analyzer (Roche Diagnostics GmbH, Mannheim, Germany).

**Highly sensitive C-reactive protein:** Serum levels of C-reactive protein (CRP) were measured at all time points by a highly sensitive particle-enhanced immunoturbidimetric CRP (hsCRP) assay using a Tina-quant C-reactive protein (latex) high sensitive reagent and a Roche Hitachi 912 analyzer (Roche Diagnostics GmbH) with a detection limit of 0.04 mg/L.

**Cytokine levels from serum:** Baseline and 21 d cytokine levels (TNF- $\alpha$ , IL-6, IFN- $\gamma$  and IL-10) in serum were determined using Quantikine HS, Human TNF- $\alpha$ /TNFSF1A (Catalog Number HSTA00D), IL-6 (HS600B), IFN- $\gamma$  (DIF50) and IL-10 (HS100B) immunoassays purchased from R&D Systems (Minneapolis MN, USA). These assays were carried out according to the manufacturer's instructions. The detection limit was 0.5 pg/mL for TNF- $\alpha$ , 0.16 pg/mL for IL-6, 15.6 pg/mL for IFN- $\gamma$  and 0.78 pg/mL for IL-10. For TNF- $\alpha$ , 94% of the samples were over the detection limit, and for IL-6, 89%. For statistical analyses, a detection limit divided by two was given as a value for those samples under the detection limit. None of the IFN- $\gamma$  samples and only 39%

of the IL-10 samples was over the detection limit and were therefore not further analyzed.

### **PBMC cell culture**

**Purification:** Human PBMC were purified by density gradient centrifugation over a Ficoll-Paque gradient (Amersham-Pharmacia Biotech, Uppsala, Sweden), as described previously<sup>[18]</sup>, from freshly collected EDTA blood on the study days (baseline, d 1, 7 and 21 wk and 3 wk after intervention). After washing, the cells were resuspended in RPMI 1640 medium (Sigma, USA) containing 10% heat-inactivated fetal calf serum (FCS) (Integro, Zaandam, Holland) and supplemented with 2 mmol/L L-glutamine (Sigma), 100 U/mL penicillin and 100 µg/mL streptomycin (Gibco BRL, Paisley, Scotland). In stimulation experiments, purified leukocytes ( $2 \times 10^6$  cells/mL) were incubated with stimulants in a final volume of one ml in 24-well plates (Nunc, Roskilde, Denmark) for 24 h in 5% CO<sub>2</sub> at 37°C.

**Stimulations:** During the stimulation experiments, the PBMC were maintained in RPMI-1640 medium containing 10% FCS. PBMC were left unstimulated or were stimulated with one of three different stimulants, simulating Gram-positive bacteria, a Gram-negative bacteria or a virus. Live Group A streptococci *S. pyogenes* serotype T1M1 obtained from the National Public Health Institute, Helsinki, Finland, grown as previously described<sup>[19]</sup>, was used as a Gram-positive bacteria at 1:1 host-cell:bacteria ratio; lipopolysaccharide (LPS) from *E. coli* serotype 0111:B4 (L-3024, Sigma) was used as a model for Gram-negative bacteria at a concentration of 100 ng/mL; and Influenza A H3N2 virus (A/Beijing/353/89) was used to infect cells at a multiplicity of infection of 5. Cell culture supernatants were collected individually at the 24 h time point and stored at -20°C before analysis.

**Cytokine levels from cell culture supernatants of stimulated PBMC:** Cytokine levels (TNF-α, IFN-γ, IL-1β, IL-2, IL-4, IL-5, IL-6, IL-8, IL-10 and IL-12p70) in cell culture supernatants from each time point (baseline, 1 d, 7 d, 21 d and 3 wk after intervention) were determined using the FlowCytomix human Th1/Th2 10 plex kit II (BMS716FFCE) from Bender MedSystems (Vienna, Austria) according to manufacturer's instructions. The detection limit was 4.5 pg/mL for IL-1β, 8.9 pg/mL for IL-2, 6.4 pg/mL for IL-4, 5.3 pg/mL for IL-5, 4.7 pg/mL for IL-6, 6.4 pg/mL for IL-8, 6.9 pg/mL for IL-10, 7.9 pg/mL for TNF-α, 9.7 pg/mL for IL-12p70 and 7.0 pg/mL for IFN-γ. Only those cytokines from which over 80% of the samples were above the detection limit were statistically analyzed. Therefore, all unstimulated samples, IL-4 and IL-5 in all stimulated samples, and IFN-γ in LPS stimulated samples were not included in further analyses. For statistical analyses, samples under the detection limit were replaced by the values obtained by dividing the detection limit by two.

### **Saliva samples and secretory IgA**

An unstimulated saliva sample was taken at every visit (at baseline, d 1, 7 and 21 and 3 wk after the intervention)

after the blood sampling. The saliva samples were placed in Eppendorf tubes, chilled, and stored at -20°C until secretory IgA was analyzed. SIgA from saliva was determined with an ELISA assay (catalog number K8870) purchased from Gentaur (Brussels, Belgium) according to the manufacturers' instructions.

### **Fecal samples and microbiological analyses**

The fecal samples were collected at home at baseline and at the end of the 3-wk intervention period. Immediately after the collection the subjects were asked to deep-freeze (-20°C) the samples at home. They were subsequently transported to the study center on the morning of the study day and the samples were immediately put on dry ice and stored at -70°C until analysis. The amounts of the probiotic strains *L. rhamnosus* GG, *B. animalis* ssp. *lactis* Bb12 and *P. freudenreichii* ssp. *shermanii* JS in the fecal samples were analyzed with a previously described real-time quantitative PCR method<sup>[20]</sup>.

### **Study diary**

Subjects were asked to fill in a structured study diary throughout the study. The study diary included questions about the use of the study product, the presence of any symptoms of respiratory infection, gastrointestinal symptoms or any other symptoms, the amount of exercise, and the use of any medication. No respiratory tract infections or major symptoms were recorded by the subjects during the study. The amount of weekly exercise carried out by the study subjects remained the same throughout the study.

### **Outcome measures and statistical analysis**

The intention-to-treat population (all randomized patients who took at least one dose of the study product) was included in the analysis. The last-observation-carried-forward (LOCF) approach was used for missing data and for subjects who withdrew early.

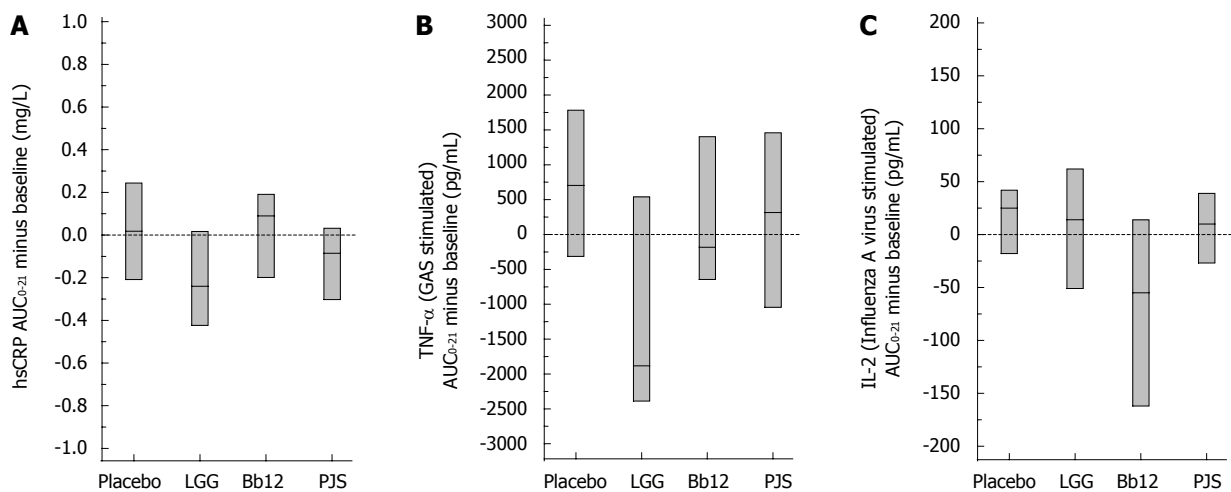
The main outcome measures were the serum hsCRP levels and the cytokines produced by PBMCs. The responses for these outcomes were calculated as the area under the curve from the 0, 1 d, 7 d and 21 d, subtracted by the baseline value (AUC<sub>0-21</sub> minus baseline).

Data is presented as mean with standard deviation (SD) or as median with interquartile range (IQR). The differences between the groups were tested using the Kruskal-Wallis test or median regression analysis with Holm's adjustment for pair wise comparisons. A *P*-value below 0.05 was regarded as statistically significant, but no adjustment was made for multiple testing.

## **RESULTS**

### **Highly sensitive CRP**

In order to study the effect of probiotic bacteria on inflammatory markers, we determined serum CRP levels at different time points during the intervention. The median AUC<sub>0-21</sub> minus baseline (IQR) for hsCRP was 0.018 (-0.209-0.244) mg/L in the placebo group, -0.240 (-0.424-0.017) mg/L in the LGG group, 0.090 (-0.199-0.191) mg/L in the Bb12 group and -0.085 (-0.303-0.032) mg/L



**Figure 1** The median AUC<sub>0-21</sub> (minus baseline) with IQR for serum highly sensitive CRP (hsCRP) levels (A), for *Streptococcus pyogenes* (GAS)-stimulated TNF- $\alpha$  production from peripheral blood mononuclear cells (B) and for Influenza A virus-stimulated IL-2 production from peripheral blood mononuclear cells (C) during the 3-wk intervention period in healthy adults ( $n = 62$ ). LGG: *Lactobacillus rhamnosus* GG; Bb12: *Bifidobacterium animalis* ssp. lactis Bb12; PJS: *Propionibacterium freudenreichii* ssp. *shermanii* JS.

**Table 1** Counts of cells of innate and adaptive immunity ( $10^9/L$ ) and levels of immunoglobulins (g/L) in serum and secretory IgA (g/mL) in saliva in healthy adults ( $n = 62$ ) at baseline presented as median (IQR)

	Placebo ( $n = 16$ )	LGG ( $n = 13$ )	Bb12 ( $n = 16$ )	PJS ( $n = 17$ )	$P$ value <sup>1</sup>
Leukocytes	4.90 (3.90-7.05)	5.20 (4.90-6.40)	5.25 (4.60-6.00)	4.90 (4.35-5.70)	0.55
Monocytes	5.00 (4.25-6.75)	5.00 (5.00-6.00)	6.00 (5.00-6.00)	6.00 (4.50-6.50)	0.84
Neutrophils	2.05 (1.67-3.72)	3.10 (2.50-3.30)	2.95 (2.12-3.47)	3.10 (2.15-3.45)	0.39
Basophils	0.05 (0.00-0.10)	0.10 (0.00-0.10)	0.05 (0.00-0.10)	0.00 (0.00-0.10)	0.73
Eosinophils	4.00 (3.00-5.75)	3.00 (2.00-6.00)	3.00 (2.25-4.75)	2.00 (1.50-3.00)	0.077
Lymphocytes	38.0 (33.5-48.7)	35.0 (31.5-37.5)	34.0 (30.0-40.5)	31.0 (26.0-40.5)	0.24
IgM	1.28 (0.97-1.65)	0.87 (0.69-1.32)	1.10 (0.73-1.66)	1.44 (0.80-1.77)	0.31
IgG	10.7 (9.5-12.2)	10.3 (9.3-11.7)	10.6 (9.1-12.1)	10.4 (8.4-11.7)	0.72
IgA	2.65 (2.45-3.21)	2.42 (2.04-3.44)	2.20 (1.73-2.94)	2.44 (1.53-2.85)	0.22
sIgA	0.23 (0.15-0.34)	0.27 (0.14-0.42)	0.40 (0.27-0.88)	0.28 (0.17-0.49)	0.065

<sup>1</sup>Kruskal-Wallis test with Monte Carlo  $P$  values. IQR: Interquartile range.

in the PJS group ( $P = 0.014$ ); a statistically significant difference was observed between LGG and Bb12 group by pair wise comparisons. In the LGG and PJS groups, hsCRP appeared to be at a lower level during the 3-wk intervention period compared with the Bb12 and placebo groups (Figure 1A).

### Serum cytokines

The baseline values for pro-inflammatory cytokine TNF- $\alpha$  in serum were 1.2 pg/mL in the placebo group, 1.0 pg/mL in the LGG, 1.0 pg/mL in the Bb12 and 0.8 pg/mL in the PJS. The change (median with IQR) from baseline to the end of 3-wk intervention for TNF- $\alpha$  in these study groups was 0.1 (-0.1-0.3) pg/mL, 0.1 (-0.02-0.2) pg/mL, 0.3 (-0.04-0.4) pg/mL and 0.0 (-0.1-0.3) pg/mL, respectively ( $P = 0.44$ ).

The baseline values for pro-inflammatory cytokine IL-6 were 0.3 pg/mL in the placebo group, 0.6 pg/mL in the LGG, 0.3 pg/mL in the Bb12 and 0.4 pg/mL in the PJS. The change (median with IQR) from baseline to the end of 3-wk intervention for IL-6 in these study groups was -0.5 (-0.6-0.0) pg/mL, -0.2 (-0.3-0.2) pg/mL, 0.1 (-0.3-0.3)

pg/mL and -0.04 (-0.3-0.1) pg/mL, respectively ( $P = 0.26$ ). There were no statistically significant differences between the study groups with respect to serum cytokine levels.

### Blood cells and immunoglobulins

Baseline values for leukocytes, monocytes, neutrophils, basophils, lymphocytes and immunoglobulins are presented in Table 1. There were no differences in these variables between the groups during the intervention.

### Cytokines produced by PBMC

We also determined whether the use of probiotic bacteria has an effect on the overall responsiveness of PBMC to various microbial stimuli in *in vitro* cultured cells. The microbe-induced cytokine production by PBMC is presented in Table 2. *S. pyogenes*-stimulated production of pro-inflammatory cytokine TNF- $\alpha$  was significantly different between the groups ( $P = 0.025$ ); a statistically significant difference was observed between LGG and placebo groups by pair wise comparisons (Figure 1B). Influenza A virus-stimulated production of Th1 cytokine IL-2 was significantly different between the groups ( $P <$

**Table 2** The effect of a 3-wk probiotic intervention on *in vitro* cytokine production (pg/mL) in peripheral blood mononuclear cells stimulated with *Streptococcus pyogenes*, lipopolysaccharide (LPS) from *E. coli* and Influenza A H3N2 virus of healthy adults ( $n = 62$ ) presented as median AUC<sub>0-21</sub> minus baseline (IQR)

	Placebo ( $n = 16$ )	LGG ( $n = 13$ )	Bb12 ( $n = 16$ )	PJS ( $n = 17$ )	<i>P</i> value <sup>1</sup>	Localization
TNF- $\alpha$						
<i>Streptococcus</i>	703 (-315-1784)	-1883 (-2389-540)	-645 (-1843-1403)	315 (-1045-1460)	0.025	LGG vs placebo
Influenza	31 (-3-66)	6 (-83-84)	-27 (-101-37)	29 (-39-83)	0.32	
LPS	11 (-16-38)	-15 (-31-10)	14 (-36-48)	-6 (-78-36)	0.53	
IFN- $\gamma$						
<i>Streptococcus</i>	-19 (-221-97)	-10 (-227-284)	71 (-161-360)	-23 (-223-155)	0.6	
Influenza	117 (29-284)	-72 (-221-194)	-7 (-436-189)	102 (-59-218)	0.25	
LPS	NA	NA	NA	NA		
IL-1 $\beta$						
<i>Streptococcus</i>	2308 (45-5222)	-1324 (-5609-352)	444 (-6152-5000)	649 (-3412-3747)	0.49	
Influenza	166 (13-478)	83 (-183-273)	-15 (-817-170)	156 (-75-791)	0.69	
LPS	67 (-29-138)	5 (-147-114)	51 (-40-122)	17 (-43-199)	0.66	
IL-2						
<i>Streptococcus</i>	5 (-23-79)	-46 (-176-0)	1 (-135-97)	39 (-105-213)	0.33	Bb12 vs others
Influenza	25 (-18-42)	14 (-51-62)	-55 (-162-14)	10 (-27-39)	< 0.001	
LPS	2 (-18-35)	27 (-25-57)	1 (-42-51)	6 (-35-25)	0.54	
IL-6						
<i>Streptococcus</i>	793 (-691-4829)	-379 (-3955-397)	205 (-2757-1553)	175 (-4344-909)	0.82	
Influenza	1530 (166-4632)	1288 (-4329-4073)	550 (-7178-1905)	1644 (-327-3344)	0.74	
LPS	947 (-418-2527)	-2189 (-3675-4053)	329 (-1949-3950)	639 (-1217-1607)	0.13	
IL-8						
<i>Streptococcus</i>	240 (-1585-3914)	34 (-3066-3143)	400 (-3638-2891)	132 (-2667-2976)	0.96	
Influenza	-455 (-1966-1670)	-193 (-2953-1520)	-1675 (-4245 to -601)	-1148 (-2550-1707)	0.31	
LPS	-742 (-3384-180)	-149 (-1608-1517)	-334 (-2692-739)	-1111 (-2457-836)	0.78	
IL-10						
<i>Streptococcus</i>	907 (263-2149)	-4 (-881-2420)	452 (-2982-1735)	226 (-86-950)	0.29	
Influenza	95 (48-284)	-57 (-159-233)	4 (-301-130)	95 (13-350)	0.25	
LPS	381 (19-602)	78 (-298-656)	347 (-403-796)	187 (28-1440)	0.51	
IL-12						
<i>Streptococcus</i>	26 (-18-75)	-32 (-99-36)	22 (-36-67)	35 (-73-172)	0.46	
Influenza	0 (-4-16)	7 (-9-30)	8 (-19-51)	5 (-25-37)	0.88	
LPS	15 (1-36)	3 (-7-28)	0 (-6-29)	0 (-62-20)	0.23	

<sup>1</sup>Median regression analysis. AUC: Area under curve (calculated from baseline, 1, 7 and 21 d minus baseline); IQR: Interquartile range; LGG: *Lactobacillus rhammosus* GG; Bb12: *Bifidobacterium animalis* ssp. *lactis* Bb12; PJS: *Propionibacterium freudenreichii* ssp. *shermanii* JS; NA: Not analyzed.

0.001); the statistically significant difference was observed between Bb12 and other groups (Figure 1C). There were no significant differences between the study groups with respect to the other cytokines produced by PBMC.

### Detection of probiotic strains from feces

In order to determine whether the ingested bacteria could also be found in the fecal samples, the bacterial DNA levels were determined at the baseline and after the 3-wk intervention. The baseline levels for all three studied probiotics were low in fecal samples (Table 3). Despite the 3-wk run-in with probiotic restriction, a detectable level of the probiotic strains, especially LGG, was harbored in some of the subjects at baseline before the probiotic ingestion (Table 3). The amount of studied probiotic in feces in a given probiotic intervention group increased significantly from the baseline values during the intervention ( $P < 0.001$ ). In the placebo group, the levels of different probiotics in feces remained low during the whole intervention period.

### Follow-up samples

Three weeks after the intervention period, follow-up samples were taken. The levels for blood cells, immunoglobulins,

hsCRP and cytokines produced by PBMC were at the baseline levels.

## DISCUSSION

In the present study, we studied the *in vivo* effects of three probiotic bacteria from three different genera on immune variables in healthy adults in a randomized, double-blind, placebo-controlled setting. The selection of these probiotics was based on our previous findings showing that, in human leukocyte cell cultures, probiotic bacteria readily induce cytokine production in PBMCs, but different bacteria are able to direct immune responses to either the Th1 type or the anti-inflammatory side in a genera-specific manner<sup>[17]</sup>. Based on the cell culture results, two potentially anti-inflammatory strains (a *Bifidobacterium* and a *Propionibacterium* strain) and a well-studied *L. rhammosus* GG strain<sup>[21]</sup> as a reference probiotic were selected. Our data indicates that *in vivo* probiotics differ in their ability to induce anti-inflammatory and cytokine responses and may have a weak, genera-specific anti-inflammatory effect reflected as a decrease in serum hsCRP levels in healthy adults. In addition, we observed

**Table 3** Detection of individual probiotic genomic DNA from fecal samples by quantitative PCR at baseline and after the 3-wk probiotic intervention in healthy adults ( $n = 62$ )

Strain	Fecal samples							
	Placebo ( $n = 16$ )		LGG ( $n = 13$ )		Bb12 ( $n = 16$ )		PJS ( $n = 17$ )	
	Baseline	After intervention	Baseline	After intervention	Baseline	After intervention	Baseline	After intervention
<i>L. rhamnosus</i> GG								
Number of subjects <sup>1</sup>	7	10	7	13	10	5	6	9
Mean (SD) <sup>2</sup>	4.7 (1.2)	5.1 (1.1)	5.1 (1.4)	8.6 (0.6)	5.2 (1.3)	4.6 (1.5)	4.5 (1.2)	5.0 (1.4)
<i>B. animalis</i> ssp. <i>lactis</i> Bb12								
Number of subjects <sup>1</sup>	6	5	5	2	7	16	2	4
Mean (SD) <sup>2</sup>	5.4 (1.6)	5.3 (1.7)	5.3 (1.4)	4.9 (1.3)	5.4 (1.6)	8.6 (0.5)	4.7 (1.1)	5.1 (1.6)
<i>P. freudenreichii</i> ssp. <i>shermanii</i> JS								
Number of subjects <sup>1</sup>	4	2	2	2	4	4	1	16
Mean (SD) <sup>2</sup>	4.5 (1.4)	4.0 (0.7)	4.1 (0.8)	4.0 (0.6)	4.4 (1.6)	4.2 (0.9)	3.8 (0.3)	8.3 (1.0)

<sup>1</sup>Number of subjects harboring a detectable level of the strain. <sup>2</sup>Mean ( $\log_{10}$ ) genome copies/g (SD). Detection limits for LGG and PJS is  $3.7 \log_{10}$  genome copies/g and for Bb12  $4.3 \log_{10}$  genome copies/g.

that, during the intervention, *S. pyogenes*-induced TNF- $\alpha$  responses and influenza A virus-induced IL-2 responses in *in vitro* cultured PBMC were reduced, indicating a clear anti-inflammatory potential of some probiotic bacteria.

To our knowledge, this is the first study to show that probiotics may reduce serum hsCRP levels in healthy adults in a randomized, double-blind, placebo-controlled setting. It appeared that in the *L. rhamnosus* GG and *P. freudenreichii* ssp. *shermanii* JS treated groups, the hsCRP level tended to be lower during the intervention, whereas in *B. animalis* ssp. *lactis* Bb12 and the placebo groups, serum hsCRP levels remained unchanged. CRP is a sensitive marker of inflammation<sup>[22]</sup> and provides an easy way to measure the anti-inflammatory potential of probiotics and other biological or pharmacological substances. This result was somewhat contradictory to our previous findings in leukocyte cell culture<sup>[17]</sup>, where *B. animalis* ssp. *lactis* Bb12 and *P. freudenreichii* ssp. *shermanii* JS were both good inducers of anti-inflammatory cytokines, whereas *L. rhamnosus* GG was a rather poor inducer of any cytokine. Previously, the effect of probiotics on CRP has only been studied in immunocompromised patients<sup>[23-27]</sup>, allergic children<sup>[28]</sup> and patients suffering from rheumatoid arthritis<sup>[29]</sup>. In immunocompromised patients, a combination of *L. casei*, *B. breve* and prebiotic galactooligosaccharides<sup>[26]</sup> and *B. longum*<sup>[30]</sup> have reduced serum CRP levels and also resulted in improvement in the overall clinical appearance of chronic inflammation<sup>[30]</sup>. In contrast to the studies above and to our results in the present study, *Lactobacillus rhamnosus* GG increased serum hsCRP levels compared to placebo in infants with IgE-associated atopic eczema dermatitis syndrome<sup>[28]</sup>. However, *L. rhamnosus* GG had no effect on serum CRP levels in patients with rheumatoid arthritis<sup>[29]</sup>. It is of interest that a combination of four probiotic bacteria (*L. rhamnosus* GG, *L. rhamnosus* Lc705, *B. breve* 99, *P. freudenreichii* ssp. *shermanii* JS) did not have an effect on sensitive CRP<sup>[28]</sup> in the same clinical setting with allergic children. In immunocompromised patients undergoing surgical procedures, *L. plantarum* 299V<sup>[23,25]</sup> or a combination of *L. acidophilus* La5, *B. animalis* ssp. *lactis* Bb12, *S. thermophilus* and *L. bulgaricus*<sup>[24,27]</sup> did not change serum CRP concentrations, either. It appears that the

effect of probiotics on CRP is controversial, and it is very difficult to compare the effects due to the differences in the measurement technique (highly sensitive *vs* normal CRP measurement), the different patient materials (healthy *vs* various diseases) and the different probiotic strains that have been used. It seems that age, the immunological status of the individual and the probiotic strain used in the study has a great impact on the immunomodulatory effects. Probiotics may have a strain-specific ability to lower serum CRP levels, thus having anti-inflammatory effects in apparently healthy adults and in patients suffering from different inflammatory conditions. In allergic patients, however, probiotics seem to induce a low-grade inflammatory response, as evidenced by increased serum CRP levels, and thus the treatment may have a beneficial effect on the host Th1/Th2 balance.

We found that *L. rhamnosus* GG was also able to reduce pro-inflammatory TNF- $\alpha$  production in the Gram-positive bacteria-stimulated PBMC. TNF- $\alpha$  is secreted by the monocytes, and it acts as an inflammatory mediator activating many types of cells. In our previous work with leukocyte cell culture, *L. rhamnosus* GG was found to be a relatively poor inducer of TNF- $\alpha$ , IL-12, IFN- $\gamma$  and IL-10<sup>[17]</sup>. Our present findings are supported by another clinical study carried out in healthy adults showing that *L. rhamnosus* GG treatment leads to decreased TNF- $\alpha$  production in PBMC<sup>[31]</sup>. In addition, when the cytokine expression pattern in the small bowel mucosa was studied, it was found that *L. rhamnosus* GG induced the expression of genes involved in immune response and inflammation (TGF-beta and TNF family members, cytokines, nitric oxide synthase 1, defensin alpha 1)<sup>[32]</sup>. Schultz and coworkers<sup>[31]</sup> observed a decreased IL-6 and IFN- $\gamma$  and an increased IL-10 and IL-4 production in PBMC obtained from *L. rhamnosus* GG treated individuals. We, however, did not find any significant changes in bacteria-induced production of cytokines apart from the TNF- $\alpha$  in the PBMC cultures of our study subjects after *L. rhamnosus* GG treatment. In another study with healthy adults and with patients with Crohn's disease, *L. rhamnosus* GG decreased the production of IL-2, IL-10 and IL-4 from PBMCs sorted as naive and memory T cells<sup>[33]</sup>. It seems that *L. rhamnosus* GG has a role in modulating the cytokine responses and may possess an

anti-inflammatory potential in healthy individuals.

In the present study, we also found that *B. animalis* ssp. *lactis* Bb-12 decreased the T lymphocyte growth factor IL-2 in the influenza-virus-stimulated PBMC, indicating an anti-inflammatory effect, which is consistent with our previous findings in human leukocyte cell culture<sup>[17]</sup>. Our finding is a new one since, in healthy adults, a combination of *B. animalis* ssp. *lactis* Bb-12 and *L. paracasei* ssp. *paracasei* CRL-431 had no effect in *in vitro*-stimulated blood cytokine production<sup>[7]</sup>. IL-2 is a very important cytokine in viral infections and inflammatory responses since it activates NK cells and induces activation and proliferation of T lymphocytes. Therefore, IL-2 production might be an important factor for a probiotic fighting against respiratory tract infections. Based on our present results, the *Bifidobacterium* strain might not be the most optimal strain against respiratory infections. Indeed, it is mainly probiotic strains from *Lactobacillus* genera-*L. rhamnosus* GG<sup>[34]</sup>, *L. casei* DN-114001<sup>[35]</sup>, a combination of *L. gasseri* PA 16/8, *B. longum* SP 07/3 and *B. bifidum* MF 20/5<sup>[5-6]</sup>, and *L. reuteri*<sup>[36]</sup>-that have reduced the incidence or symptoms of common cold or respiratory tract infections. However, the immunomodulatory effects underlying the results observed in these studies have not been fully elucidated.

In conclusion, it appears that probiotics have an anti-inflammatory potential seen as a decrease in serum CRP levels and as a reduction in bacteria-induced production of pro-inflammatory cytokines in PBMC in healthy adults. However, all of the markers were in the normal range, and therefore the real impact of probiotics as anti-inflammatory substances warrants further evaluation in studies during inflammatory processes and with individuals suffering from various types of inflammatory or autoimmune diseases.

## ACKNOWLEDGMENTS

We would like to thank Hanna Valtonen and Mari Aaltonen for their expert technical assistance in the PBMC isolations, Juha Laukonmaa for technical assistance in qPCR analysis and Sirkka Kokkonen for manufacturing the study products.

## COMMENTS

### Background

Probiotics have been mostly studied in the prevention and treatment of different gastrointestinal diseases and allergy. Probiotic products, however, are usually consumed by the general, healthy population but not much is known what kind of effects they have on immune system in healthy adults.

### Research frontiers

It is not fully clarified how probiotics exert their health effects, but one of the most probable action mechanisms is the modulation of immune responses *via* gut mucosal immune system.

### Innovations and breakthroughs

In the present study the immunomodulatory effects of probiotics were studied in healthy adults. Probiotic bacteria had strain-specific anti-inflammatory effects reflected in reduced sensitive C-reactive protein, which is a new finding, and decreased proinflammatory cytokine production in peripheral blood mononuclear cells (PBMC).

### Applications

Understanding of the specific immunomodulatory effects of probiotics may help in

designing future probiotics for targeted purposes. As the effects in the present study were investigated in healthy adults, the real impact of probiotics on inflammatory variables warrants further evaluation during inflammatory processes and with individuals suffering from various types of inflammatory or autoimmune diseases.

### Peer review

The paper by Kekkonen and co-workers investigated the effects of three probiotic bacteria on immune variables in healthy adults. They observed strain-specific anti-inflammatory effects for distinct bacteria. Overall this paper is interesting and it has clearly stated aims, the sample size and the overall designs of the study are fair, the results adequate to provide experimental evidence and to support valid conclusions. As placebo per se could cause effects on immune response, a further control group, formed by healthy subjects, would be advisable in order to analyze the basic fluctuation of all the parameters studied.

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