



# The effect of probiotics on inflammatory biomarkers: a meta-analysis of randomized clinical trials

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## Abstract

**Purpose** No study has summarized earlier findings on the effect of probiotic supplementation on inflammatory biomarkers. This systematic review and meta-analysis was conducted to systematically review the available placebo-controlled clinical trials about the effect of probiotic supplementation on several inflammatory biomarkers in adults.

**Methods** Relevant papers published up to March 2018 were searched up through PubMed, MEDLINE, SCOPUS, EMBASE, and Google Scholar, using following suitable keywords. Clinical trials that examined the effect of probiotic supplementation on inflammation in adults were included.

**Results** Overall, 42 randomized clinical trials (1138 participants in intervention and 1120 participants in control groups) were included. Combining findings from included studies, we found a significant reduction in serum hs-CRP [standardized mean difference (SMD)  $-0.46$ ; 95% CI  $-0.73, -0.19$ ], TNF- $\alpha$  ( $-0.21$ ;  $-0.34, -0.08$ ), IL-6 ( $-0.37$ ;  $-0.51, -0.24$ ), IL-12 ( $-0.47$ ;  $-0.67, -0.27$ ), and IL-4 concentrations ( $-0.48$ ;  $-0.76, -0.20$ ) after probiotic supplementation. Pooling effect sizes from 11 studies with 12 effect sizes, a significant increase in IL-10 concentrations was seen ( $0.21$ ;  $0.04, 0.38$ ). We failed to find a significant effect of probiotic supplementation on serum IL-1B ( $-0.17$ ;  $-0.37, 0.02$ ), IL-8 ( $-0.01$ ;  $-0.30, 0.28$ ), and IFN- $\gamma$  ( $-0.08$ ;  $-0.31, 0.15$ ) and IL-17 concentrations ( $0.06$ ;  $-0.34, 0.46$ ).

**Conclusions** Probiotic supplementation significantly reduced serum concentrations of pro-inflammatory cytokines including, hs-CRP, TNF- $\alpha$ , IL-6, IL-12, and IL-4, but it did not influence IL-1B, IL-8, IFN- $\gamma$ , and IL-17 concentrations. A significant increase in serum concentrations of IL-10, as a anti-inflammatory cytokine was also documented after probiotic supplementation.

**Keywords** Diet · Probiotic · Inflammation · Cytokine · Meta-analysis

## Abbreviations

SMD	Standardized mean difference
T2DM	Type 2 diabetes mellitus
MtS	Metabolic syndrome

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CVD	Cardiovascular disease
NAFLD	Non-alcoholic fatty liver disease
SCFAs	Short-chain fatty acids
IBD	Inflammatory bowel disease
IBS	Irritable bowel syndrome
CRP	C-reactive protein
TNF-a	Tumor necrosis factor-a
SD	Standard deviation
SE	Standard error
MD	Mean difference
RCT	Randomized clinical trial
GALT	Gut-associated lymphoid tissue
IL	Interleukin
IL-1B	Interleukin 1 beta
IFN-g	Interferon-gama

## Introduction

Inflammation is a part of innate immunity response to protect the body against injury and harm [1, 2]. However, chronic low-grade systematic inflammation is associated with higher risk of several chronic diseases including type 2 diabetes mellitus (T2DM), obesity, metabolic syndrome (MtS), cardiovascular disease (CVD) [3, 4], and non-alcoholic fatty liver disease (NAFLD) [5].

Adoption of a physically active life-style may attenuate low-grade inflammation [6, 7]. In addition, diet can also play a role in this condition [8]. For instance, higher consumption of processed and fried foods has been linked to increased risk of inflammation [9]. In contrast, adherence to a dietary pattern rich in fibers might decrease inflammatory cytokines which may be due to incremented production of short-chain fatty acids (SCFAs) in the colon [10]. SCFAs are physiologically active byproducts of fermentation by microflora which are suggested to modulate systemic inflammation [11].

Probiotics are non-pathogenic microorganisms which when administered in adequate amounts confer several health benefits in improving conditions associated with inflammation [12]. Probiotic supplementation will change intestinal microbium to increase production of SCFAs [13]. Probiotic supplementation has attenuated inflammation-related diseases such as cardiovascular disease, allergies, inflammatory bowel disease (IBD), irritable bowel syndrome (IBS), autoimmune diseases, and cancers [14, 15]. Studies on the effect of probiotics supplementation on inflammatory cytokines are conflicting. Although some studies have found significant reduction in inflammatory cytokines after supplementation with probiotics compared to placebo [16, 17], some others could not reach the same effect [18, 19]. A recent systematic review and meta-analysis of clinical trials indicated that prebiotics significantly reduce serum

C-reactive protein (CRP), where symbiotic reduced CRP and tumor necrosis factor-a (TNF-a), as compared to placebo [10].

Despite several publications on the effect of probiotics on different inflammatory biomarkers, no further study has summarized the findings from the previous studies in this regard. Therefore, we aimed to systematically review the available placebo-controlled clinical trials about the effect of supplementation with probiotics on several inflammatory biomarkers in adults and to do a meta-analysis, if possible.

## Methods

### Search strategy

Relevant studies published up to March 2018 were searched through PubMed, MEDLINE, SCOPUS, EMBASE, and Google Scholar, using the following suitable MESH and non-MESH keywords: (“Lactobacillus”[All Fields] OR “Streptococcus”[All Fields] OR “Saccharomyces”[All Fields] OR “Enterococcus”[All Fields] OR “Bifidobacterium”[All Fields] OR “Probiotic”[All Fields] OR “Lactococcus”[All Fields]) AND (“Inflammation”[All Fields] OR “inflammatory biomarker”[All Fields] OR “Tumor necrosis factor”[All Fields] OR “C-Reactive protein”[All Fields] OR “Transforming growth factor beta”[All Fields] OR “Cytokine”[All Fields] OR “Acute phase reactant”[All Fields] OR “Matrix metalloproteinase”[All Fields] OR “Intercellular adhesion molecule-1”[All Fields] OR “Monocyte chemotactic protein 1”[All Fields] OR “Inflammation Mediator”[All Fields] OR “Adipokine”[All Fields] OR “Interleukin”[All Fields] OR “Systemic inflammation”[All Fields]) AND (“Clinical Trial” OR “trial”). No restrictions of language or time of publication were used. To avoid missing any publication, we also examined reference lists of all included studies as well as review articles. Unpublished data and grey literatures, including dissertations, congress abstracts, and patents, were not included in the current meta-analysis. In addition, we removed duplicate citations.

### Inclusion criteria

All randomized clinical trials that investigated the effect of probiotic supplementation on inflammatory biomarkers in adults were included. In case of several publications with the same data set, we included only the most complete one. If data for specific subgroups were reported, results for the whole population were used for the meta-analysis, unless it was not available. Moreover, when a study was performed on separate groups of participants, data of each group in

comparison with the control group were considered as an independent study.

### Exclusion criteria

Studies that: (1) were conducted on animal models, pregnant or lactating women, and studies with children or only elderly participants; (2) did not have random allocation; (3) had an observational design; (4) examined the effect of another intervention along with probiotic supplementation; (5) examined postprandial inflammatory responses after an immediate intervention; (6) had no control group; (7) used symbiotic as the intervention; (8) used probiotic enriched foods; (9) examined only gene expression of inflammatory biomarkers; and (10) examined concentrations of inflammatory biomarkers in specific cell lines, were excluded.

### Data extraction

The following data were extracted by two independent reviewers: first author's name, publication year, subjects' health condition, study sample size, participants' sex, number of subjects in each groups, participants' age, type of probiotic microbes, trial design (parallel/cross-over), type of control, duration of intervention, mean and standard deviation (SD) concentrations of inflammatory biomarkers after intervention in each groups, mean (SD) changes in inflammatory biomarkers after intervention in each groups, and covariates. If data were reported as standard errors (SEs) or interquartile ranges, they were converted to SDs using appropriate formulas. When concentration of an inflammatory biomarker was reported in different units, it was converted to the most frequently used one.

### Statistical analysis

The overall effect sizes were calculated as mean differences (MDs) and SEs of concentrations of inflammatory biomarkers between probiotic and control groups. If mean (SDs) changes were not reported, we took end-of-trial means (SDs) of biomarkers in each group. Then, we used fixed-effects model to calculate overall effect size, because random-effects model gives larger weights to small extreme studies [20]. Between-study heterogeneity was examined by the Cochran's  $Q$  test and  $I^2$  statistic. To find probable sources of between-study heterogeneity, we conducted subgroup analyses based on participants' health condition (healthy/gastrointestinal disease/skeletal disorders/metabolic diseases/allergy and autoimmune diseases/renal diseases/critically ill/other diseases), sex (male/female/both genders), age (adult/adult + elderly), study design (parallel/cross-over), supplement dosage ( $< 1/1-10/10-100/\geq 100$  CFU/day), duration of intervention ( $< 10$  weeks/ $\geq 10$  weeks), outcome assessment

method (immunoassay/electrochemiluminescence/bead assay), and probiotic type (lactobacillus/bifidobacter/saccharomyces/different types), using a fixed-effects model. All statistical analyses were done using Stata software, version 11.2 (Stata Corp, College Station, TX, USA).  $P < 0.05$  was considered as statistically significant.

## Results

### Study characteristics

Overall, 42 publications with 46 effect sizes were included in this meta-analysis [16–19, 21–58]. The flow diagram of study selection is shown in (Supplementary Figure 1). All studies were randomized clinical trials (RCTs), published between 2003 and 2018. Duration of intervention was varied from 1 to 52 weeks. A total of 1138 participants in intervention group and 1120 participants in control group were enrolled in these studies (43.22% male and 56.78% female). Characteristics of included studies are summarized in (Supplementary Table 1).

Among publications included, 40 studies were parallel RCTs [16–19, 21–29, 31–41, 43–58], whereas two remaining studies had cross-over design [30, 42]. These 46 effect sizes were related to studies in healthy subjects [31, 37, 39, 45, 46, 51], patients with gastrointestinal diseases [17, 19, 25, 32, 33, 44, 47, 49], patients with skeletal disorders [18, 22, 32, 34–36, 53, 58], those with metabolic diseases [23, 24, 29, 40, 50], those with allergy and autoimmune diseases [32, 38, 41, 43, 56, 57], patients with renal disease [26, 42, 48, 52, 55], critically ill patients [16, 28], and the remaining among patients with HIV [54], heart failure [27], edetulous [30], and depression [21]. Two effect sizes were from studies were done in men [23, 43], seven effect sizes from studies on women [22, 31, 32, 35, 41, 50, 53], and 37 effect sizes from studies on both genders [16–19, 24–34, 36–40, 42, 44–49, 51, 52, 54–58].

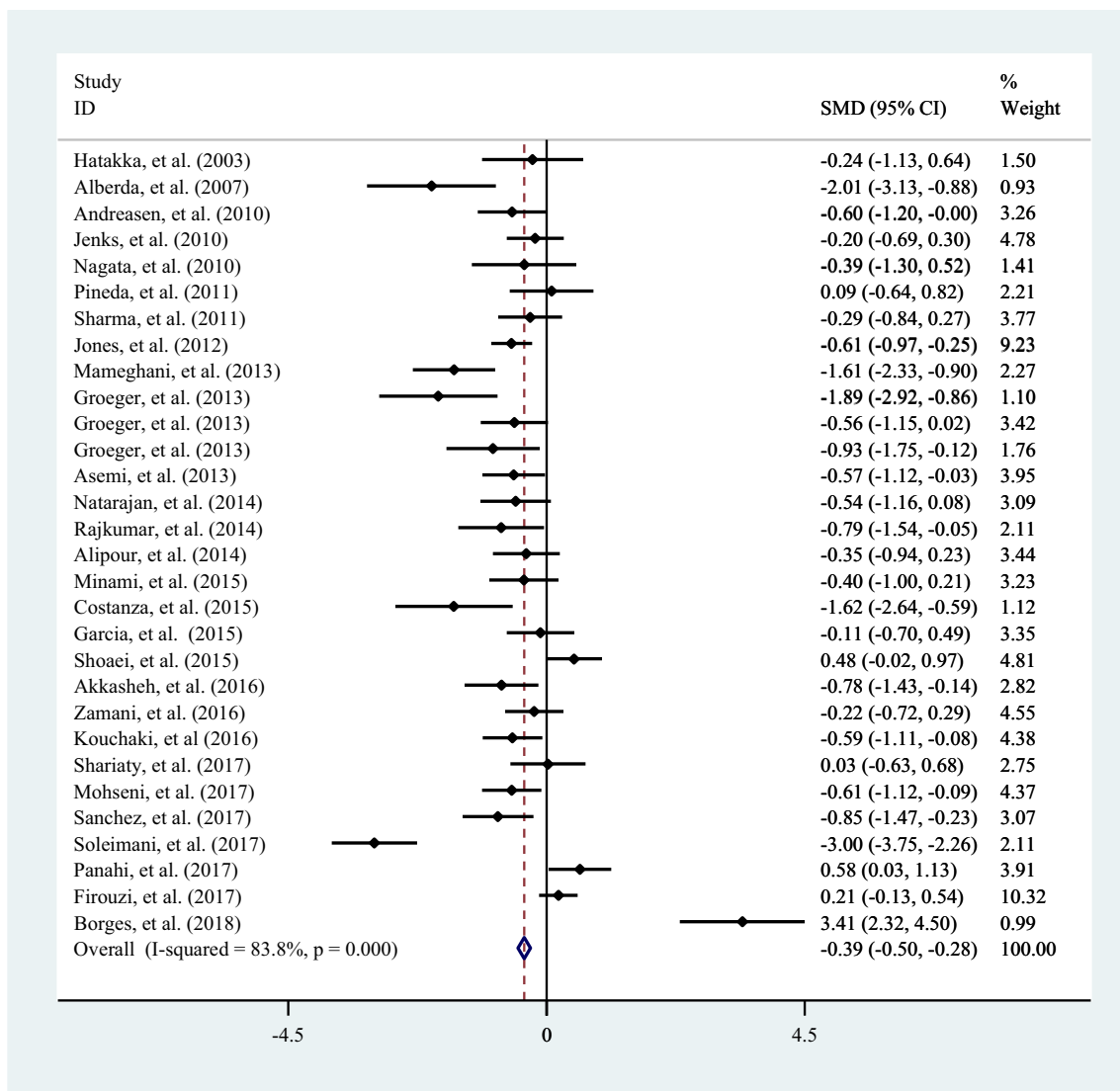
Probiotics administered were lactobacillus in 12 studies [18, 19, 22, 23, 30, 34, 37, 41, 46, 51, 53, 57], bifidobacter in 4 studies [32, 39, 44, 56], and saccharomyces in two studies [27, 54], whereas 24 studies used more than 1 type of probiotic [16, 17, 21, 24–26, 28, 29, 31, 33, 35, 36, 38, 40, 42, 43, 45, 47–50, 52, 55, 58]. Daily dose of probiotic bacteria was ranged from 0.06 to 1800 billion. Moreover, three studies did not report daily dose of consumed bacteria [17, 27, 57]. Placebo was used in all studies. Four studies had a third arm intervention [16, 25, 33, 45], which was not entered in the current meta-analysis. Adjustment for the baseline measures of inflammatory biomarkers was conducted in 14 studies [21, 22, 25, 31, 37–40, 44, 45, 50, 52, 53, 58], while 4 studies did not report any adjustment [16, 23, 33, 42].

Biochemical analyses were conducted using immunoassay [17–19, 21–26, 29, 33, 35, 37, 38, 40, 41, 44, 45, 47, 49–58], electrochemiluminescence [32], and bead assay [30, 31, 46]. In addition, nine studies did not report methods used to quantify inflammatory biomarkers [16, 27, 28, 34, 36, 39, 42, 43, 48].

### Findings on the effect of probiotics on hs-CRP

Combining findings from 31 studies with 33 effect sizes, we found a significant reduction in serum hs-CRP concentrations after probiotic supplementation, as compared to placebo [standardized mean difference (SMD)  $-0.39$ ; 95% CI  $(-0.50, -0.28)$ ,  $I^2 = 83.8\%$ ] (Fig. 1). However, a significant

between-study heterogeneity was found. Subgroup analysis by the participants' age and sex, study duration and outcome assessment method did not provide any explanation for between-study heterogeneity (Table 1). When the subgroup analysis was done based on health status of participants, between-study heterogeneity was disappeared. In this analysis, the effect of probiotic supplementation on hs-CRP was significant in participants with all health conditions except for patients with metabolic diseases and those with allergy and autoimmune diseases. In addition, subgroup analysis based on the type and dose of supplemented bacteria resulted in the disappearance of between-study heterogeneity for studies used lactobacillus and bifidobacter and those used  $\geq 100$  CFU/day of bacteria, respectively. In addition,



**Fig. 1** Forest plot for the effect of probiotics supplementation on serum hs-CRP concentrations, expressed as the mean differences between the intervention and the control diets. The area of each

square is proportional to the inverse of the variance of the SMD. Horizontal lines represent 95% CIs. Diamonds represent pooled estimates from fixed-effects analysis

**Table 1** Subgroup analysis for the effect of probiotics supplementation on serum concentrations of CRP and IL-6

Subgroup	Number of effect sizes		Participants, <i>N</i>		Effect size (95% CI)		<i>f</i> <sup>2</sup>		<i>P</i> within <sup>a</sup>		<i>P</i> between <sup>a</sup>	
	CRP	IL-6	CRP	IL-6	CRP	IL-6	CRP	IL-6	CRP	IL-6	CRP	IL-6
Overall	31	20	1579	840	-0.33 (-0.44, -0.23)	-0.37 (-0.51, -0.24)	84.2	69.7	<0.001	<0.001	-	-
Participants' age												
Adult	14	13	521	585	-0.31 (-0.48, -0.14)	-0.34 (-0.50, -0.17)	85.4	74.5	<0.001	<0.001	0.04	0.40
Adult + elderly	16	7	1038	255	-0.33 (-0.46, -0.20)	-0.47 (-0.72, -0.21)	84.1	59.6	<0.001	0.02	-	-
NR	1	-	20	-	-1.62 (-2.64, -0.59)	-	-	-	-	-	-	-
Supplement dosage												
< 10 CFU/day	12	10	668	93	-0.45 (-0.61, -0.29)	-0.49 (-0.68, -0.31)	83.5	55.4	<0.001	0.01	<0.001	0.07
10–100 CFU/day	15	8	802	282	-0.12 (-0.26, 0.02)	-0.14 (-0.39, 0.10)	83.4	77.4	<0.001	<0.001	-	-
≥ 100 CFU/day	3	2	89	465	-1.36 (-1.83, -0.89)	-0.47 (-0.89, -0.05)	49.8	83.9	0.13	0.01	-	-
NR	1	-	20	-	-1.62 (-2.64, -0.23)	-	-	-	-	-	-	-
Study duration												
< 10 week	18	15	837	674	-0.48 (-0.62, -0.34)	-0.39 (-0.55, -0.24)	71.4	70.2	<0.001	<0.001	<0.01	0.60
≥ 10 week	13	5	742	166	-0.16 (-0.32, -0.01)	-0.30 (-0.62, 0.01)	90.1	74.1	<0.001	<0.01	-	-
Outcome assessment												
Immunoassay	18	3	1129	96	-0.27 (-0.39, -0.15)	-0.50 (-0.93, -0.08)	87.7	81.6	<0.001	<0.01	0.02	0.18
Electrochemiluminescence	3	10	86	419	-0.90 (-1.33, -0.47)	-0.44 (-0.64, -0.24)	58.4	64.6	0.09	<0.01	-	-
Bead assay	-	5	-	274	-	-0.34 (-0.59, -0.10)	-	81.1	-	<0.001	-	-
NR	10	2	364	51	-0.39 (-0.61, -0.18)	0.20 (-0.36, 0.75)	77.0	0.0	<0.001	0.57	-	-
Health condition												
Healthy	3	3	201	206	-0.59 (-0.87, -0.31)	-0.33 (-0.61, -0.05)	0.0	90.0	0.70	<0.001	<0.001	0.71
Gastrointestinal disease	3	6	116	218	-0.73 (-1.11, -0.34)	-0.41 (-0.69, -0.14)	73.0	79.0	0.02	<0.001	-	-
Skeletal disorders	6	4	267	161	-0.26 (-0.51, -0.02)	-0.44 (-0.76, -0.12)	0.0	0.0	0.83	0.44	-	-
Metabolic diseases	6	1	496	45	-0.01 (-0.19, 0.17)	-0.17 (-0.76, 0.42)	74.8	-	<0.01	-	-	-
Allergy an autoimmune diseases	3	1	105	26	0.01 (-0.40, 0.42)	0.25 (-0.52, 1.02)	80.2	-	<0.01	-	-	-
Renal diseases	4	2	171	72	-0.51 (-0.87, -0.15)	-0.52 (-1.01, -0.03)	96.9	90.9	<0.001	<0.01	-	-
Critical illness	2	-	59	-	-1.73 (-2.33, -1.12)	-	0.0	-	0.56	-	-	-
Other diseases	4	3	164	112	-0.60 (-0.91, -0.28)	-0.43 (-0.81, -0.06)	55.9	0.0	0.07	0.53	-	-
Participants' sex												
Male	3	1	124	45	-0.03 (-0.40, 0.34)	-0.17 (-0.76, 0.42)	77.4	-	0.01	-	0.03	0.24
Female	3	4	159	183	-0.07 (-0.39, 0.24)	-0.18 (-0.48, 0.12)	76.0	82.5	0.01	<0.01	-	-
Both genders	25	15	1296	612	-0.40 (-0.51, -0.28)	-0.45 (-0.61, -0.28)	85.5	67.2	<0.001	<0.001	-	-
Probiotic type												
Lactobacillus	5	10	268	448	-0.46 (-0.70, -0.21)	-0.37 (-0.56, -0.18)	0.0	39.6	0.49	0.09	0.01	0.34
Bifidobacter	6	4	203	140	-0.72 (-1.01, -0.43)	-0.60 (-0.95, -0.25)	30.9	74.0	0.20	<0.01	-	-
Saccharomyces	2	1	64	44	-0.48 (-1.00, 0.03)	-0.52 (-1.12, 0.08)	84.1	-	0.01	-	-	-
Different types	18	5	1044	208	-0.22 (-0.34, -0.09)	-0.20 (-0.49, 0.08)	89.5	87.8	<0.001	<0.001	-	-

<sup>a</sup>*P* values were obtained by fixed-effect analysis

the effect of probiotic supplementation was seen only in studies that were performed in both gender and it was not significant in studies that confined the participants to males or females.

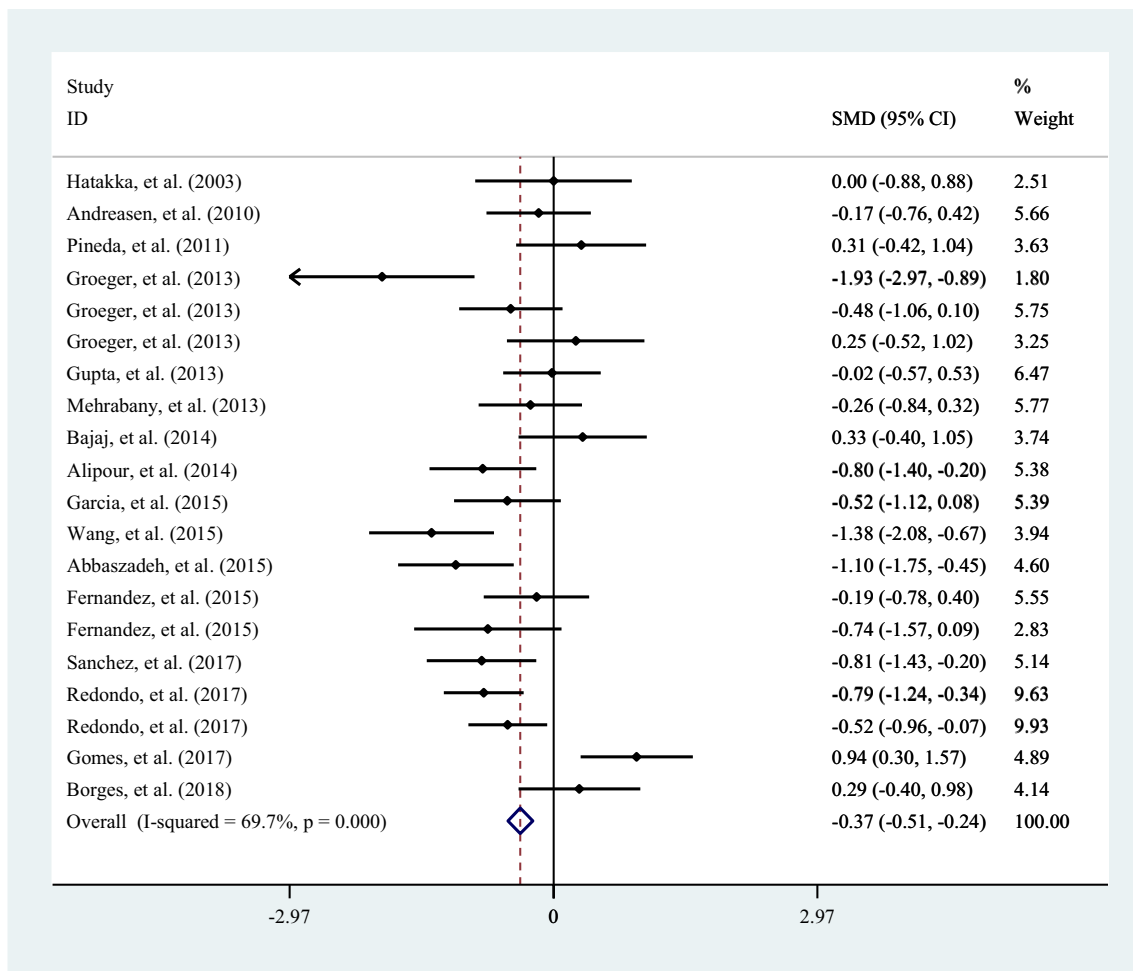
### Findings on the effect of probiotics on serum IL-6

Based on 16 studies with 20 effect sizes, a significant reduction in serum IL-6 was found after probiotic supplementation [SMD  $-0.37$ ; 95% CI  $(-0.51, -0.24)$ ,  $I^2 = 69.7%$ ] (Fig. 2). A significant between-study heterogeneity was seen; however, participants' age and sex, supplement dose, study duration, and method of outcome assessment did not explain the heterogeneity (Table 1). Health status of participants as well as type of bacteria used made the between-study heterogeneity disappeared. In all above-mentioned subgroups, the significant effect of probiotics on serum IL-6 concentrations was still significant; however, the effect was not significant

among studies that used a dosage of 10–100 CFU/day, those that were performed on women as well as those that used multiple types of bacteria in one study.

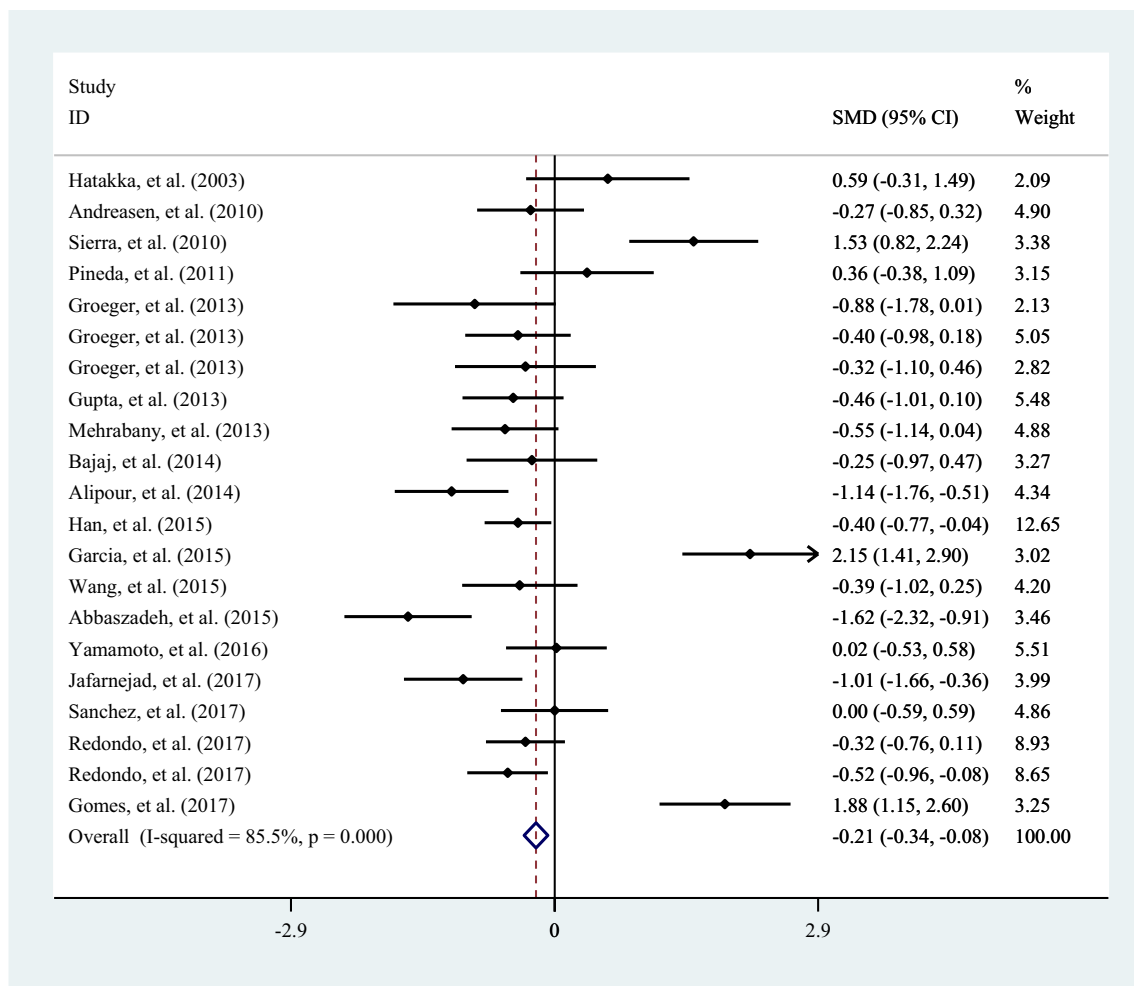
### Findings on the effect of probiotics on TNF-a

Pooling effect sizes from 18 studies with 21 effect sizes, we found a significant reduction in serum TNF-a concentrations after probiotic supplementation [SMD  $-0.21$ ; 95% CI  $(-0.34, -0.08)$ ,  $I^2 = 85.5%$ ] (Fig. 3). Due to heterogeneity, we conducted subgroup analysis to find possible sources (Table 2). Participants' age and sex as well as study duration did not explain the between-study heterogeneity. In all above-mentioned subgroups, except for studies with an intervention duration of  $\geq 10$  weeks, the effect of probiotic supplementation on TNF-a was still significant. Dosages of probiotic supplements, method of outcome assessment, health status of participants, and type of bacteria could



**Fig. 2** Forest plot for the effect of probiotics supplementation on serum IL-6 concentrations, expressed as the mean differences between the intervention and the control diets. The area of each

square is proportional to the inverse of the variance of the SMD. Horizontal lines represent 95% CIs. Diamonds represent pooled estimates from fixed-effects analysis



**Fig. 3** Forest plot for the effect of probiotics supplementation on serum TNF- $\alpha$  concentrations, expressed as the mean differences between the intervention and the control diets. The area of each

square is proportional to the inverse of the variance of the SMD. Horizontal lines represent 95% CIs. Diamonds represent pooled estimates from fixed-effects analysis

provide some explanations for the heterogeneity. In these analyses, we found that the effect of probiotics on TNF- $\alpha$  was not significant among studies that used probiotics at the dosages of < 1 and 10–100 CFU/day. In addition, the effect of supplementation was not significant among studies that were done on healthy participants and those with allergy and autoimmune diseases.

### Findings on the effect of probiotics on serum Interleukin-10 (IL-10)

Pooling effect sizes from 11 studies with 12 effect sizes, a significant increase in IL-10 concentrations was seen following probiotic supplementation [SMD 0.21; 95% CI (0.04, 0.38),  $I^2 = 48.5%$ ] (Fig. 4). However, in the subgroup analysis, the effect was not significant among studies conducted exclusively on adults as well as those performed on both genders, studies that used bifidobacter, those that used the

dosage of  $\geq 1$  CFU/day of probiotic supplement, and those that used bead assay for outcome assessment (Table 2). In addition, no significant effect was seen among studies that were performed on healthy participants or subjects with gastrointestinal diseases.

### Findings on the effect of probiotics on serum interleukin-1 beta (IL-1B)

Combining ten effect sizes from nine studies, we failed to find a significant effect of probiotic supplementation on IL-1B concentrations [SMD -0.17; 95% CI (-0.37, 0.02),  $I^2 = 80.7%$ ] (Fig. 5). A significant between-study heterogeneity was seen. Subgroup analysis based on participants' age and gender, duration of study, and health status of participants did not explain the heterogeneity (Table 3). When the analysis was done stratified by supplement dosage, the between-study heterogeneity was disappeared for studies

**Table 2** Subgroup analysis for the effect of probiotics supplementation on serum concentrations of TNF- $\alpha$  and IL-10

Subgroup	Number of effect sizes		Participants, <i>N</i>		Effect size (95% CI)		<i>I</i> <sup>2</sup>		<i>P</i> within <sup>a</sup>		<i>P</i> between <sup>a</sup>	
	TNF- $\alpha$	IL-10	TNF- $\alpha$	IL-10	TNF- $\alpha$	IL-10	TNF- $\alpha$	IL-10	TNF- $\alpha$	IL-10	TNF- $\alpha$	IL-10
Overall	21	12	987	545	-0.21 (-0.34, -0.08)	0.21 (0.04, 0.38)	85.5	48.5	<0.001	0.03	-	-
Participants' age												
Adult	13	9	592	424	-0.16 (-0.33, 0.01)	0.16 (-0.03, 0.35)	84.5	57.2	<0.001	0.01	0.40	0.27
Adult + elderly	7	3	395	121	-0.27 (-0.48, -0.07)	0.39 (0.03, 0.75)	88.2	0.0	<0.001	0.47	-	-
Supplement dosage												
< 1 CFU/day	4	3	176	132	0.26 (-0.07, 0.59)	0.67 (0.31, 1.02)	95.3	22.7	<0.001	0.27	<0.001	0.01
1–10 CFU/day	6	5	302	261	-0.39 (-0.62, -0.17)	0.11 (-0.14, 0.35)	37.5	49.0	0.15	0.09	-	-
10–100 ( $\geq$ 10) CFU/day	7	4	249	152	0.05 (-0.21, 0.31)	0.02 (-0.30, 0.34)	82.6	0.0	<0.001	0.47	-	-
$\geq$ 100 CFU/day	2	-	93	-	-0.91 (-1.34, -0.47)	-	84.4	-	0.01	-	-	-
NR	2	-	167	-	-0.28 (-0.58, 0.03)	-	36.3	-	0.21	-	-	-
Study duration												
< 10 week	15	8	763	412	-0.31 (-0.45, -0.16)	0.18 (-0.01, 0.38)	83.7	54.8	<0.001	0.03	<0.01	0.52
$\geq$ 10 week	6	4	224	133	0.15 (-0.13, 0.43)	0.31 (-0.04, 0.66)	88.6	45.2	<0.001	0.14	-	-
Outcome assessment												
Immunoassay	13		634	7	-0.23 (-0.40, -0.07)	0.39 (0.16, 0.63)	87.7	61.0	<0.001	0.01	0.28	0.09
Electrochemiluminescence	3		96	-	-0.48 (-0.89, -0.07)	-	0.0	-	0.60	-	-	-
Bead assay	3		206	3	-0.06 (-0.35, 0.22)	0.03 (-0.25, 0.30)	94.0	0.0	<0.001	0.52	-	-
NR	2		51	2	0.08 (-0.49, 0.64)	-0.02 (-0.58, 0.53)	50.7	0.0	0.15	0.92	-	-
Health condition												
Healthy	4	4	246	246	0.16 (-0.10, 0.43)	0.18 (-0.07, 0.43)	94.0	69.0	<0.001	0.02	<0.001	0.09
Gastrointestinal disease	6	2	306	74	-0.50 (-0.73, -0.27)	-0.20 (-0.66, 0.26)	64.1	0.0	0.01	0.47	-	-
Skeletal disorders	6	3	231	113	-0.48 (-0.75, -0.21)	0.40 (0.02, 0.77)	71.0	0.0	<0.01	0.44	-	-
Metabolic diseases	1	-	45	-	-0.27 (-0.85, 0.32)	-	-	-	-	-	-	-
Allergy an autoimmune diseases	2	1	76	44	-0.09 (-0.55, 0.36)	0.12 (-0.48, 0.71)	0.0	-	0.48	-	-	-
Renal diseases	1	1	39	39	-0.39 (-1.02, 0.25)	0.99 (0.32, 1.65)	-	-	-	-	-	-
Other diseases	1	1	44	29	2.15 (1.41, 2.90)	0.05 (-0.68, 0.78)	-	-	-	-	-	-
Participants' sex												
Male	1	-	45	-	-0.27 (-0.85, 0.32)	-	-	-	-	-	0.48	0.12
Female	5	3	224	135	-0.35 (-0.63, -0.07)	0.44 (0.10, 0.79)	91.5	0.0	<0.001	0.73	-	-
Both genders	15	9	718	410	-0.16 (-0.31, -0.01)	0.14 (-0.06, 0.33)	84.3	56.6	<0.001	0.01	-	-
Probiotic type												
Lactobacillus	10	7	470	346	-0.19 (-0.38, -0.01)	0.22 (0.00, 0.43)	78.0	52.0	<0.001	0.05	<0.001	0.12
Bifidobacter	4	2	140	88	-0.32 (-0.66, 0.01)	0.11 (-0.53, 0.31)	0.0	10.9	0.43	0.28	-	-
Saccharomyces	1	1	44	29	2.15 (1.41, 2.90)	0.05 (-0.68, 0.78)	-	-	-	-	-	-



Table 2 (continued)

Subgroup	Number of effect sizes		Participants, N		Effect size (95% CI)		$I^2$	$P$ within <sup>a</sup>		$P$ between <sup>a</sup>	
	TNF-a	IL-10	TNF-a	IL-10	TNF-a	IL-10		TNF-a	IL-10	TNF-a	IL-10
Different types	6	2	333	82	-0.39 (-0.61, -0.16)	0.62 (0.17, 1.07)	90.6	51.5	<0.001	0.15	

<sup>a</sup> $P$  values were obtained by fixed-effect analysis

that used < 1 CFU/day of bacteria ( $I^2 = 19.2\%$ ). The effect of supplementation was non-significant in all subgroups, except for studies that were performed on adults and on females, in which a significant reduction in serum IL-1B concentrations was found after supplementation. In addition, probiotic supplementation at the dosages of < 1 or  $\geq 1$  CFU/day resulted in a significant reduced concentrations of serum IL-1B.

**Findings on the effect of probiotics on serum IL-12**

When we combined data from eight studies with nine effect sizes, a significant reduction in serum IL-12 concentrations was found after probiotic supplementation [SMD - 0.47; 95% CI (- 0.67, - 0.27),  $I^2 = 85.2\%$ ] (Fig. 6). To find possible sources of between-study heterogeneity, we did subgroup analysis, but participants’ age, gender, and health status and supplement dosage did not provide any explanation for this heterogeneity (Table 3). However, heterogeneity was disappeared in studies with a duration of  $\geq 10$  weeks.

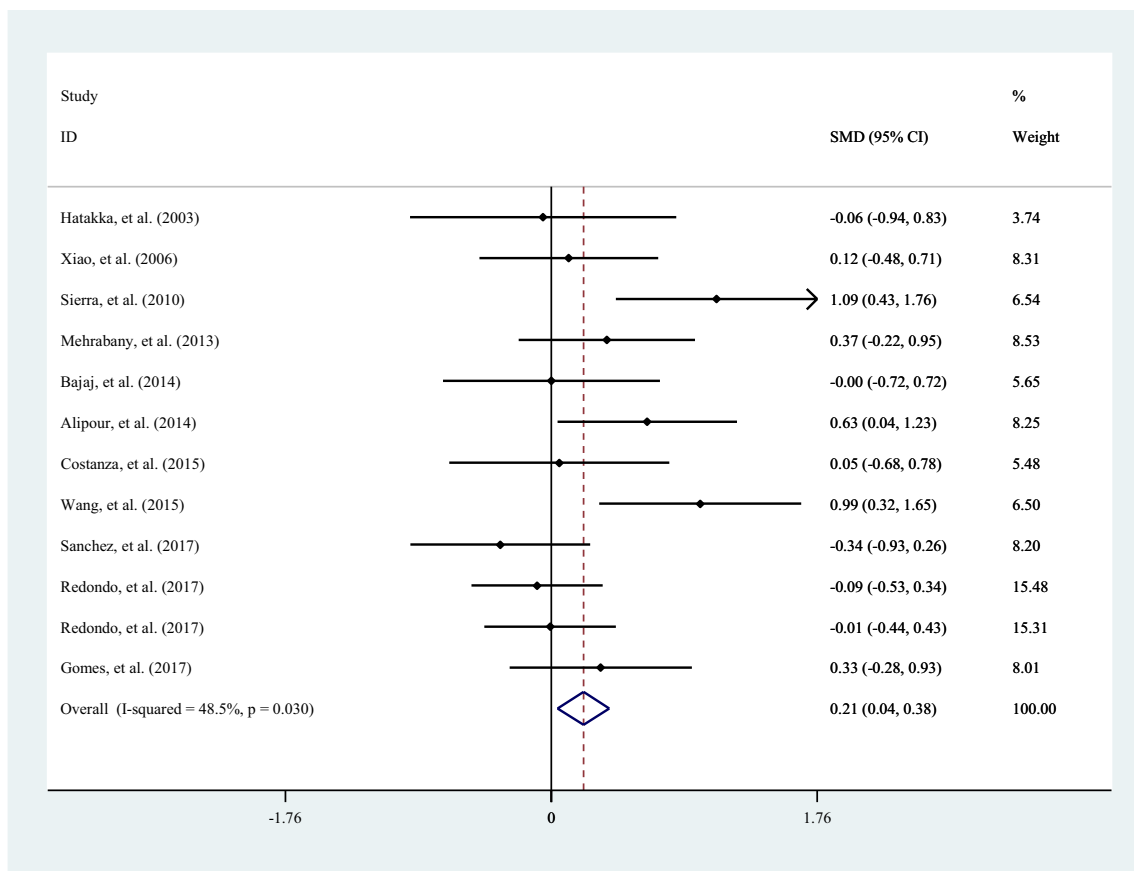
**Findings on the effect of probiotics on other cytokines**

Pooling five effect sizes from four studies, no significant effect of probiotic supplementation on serum IL-8 [SMD - 0.01; 95% CI (- 0.30, 0.28),  $I^2 = 73.0\%$ ] and interferon-gama (IFN-g) concentrations was found [SMD - 0.08; 95% CI (- 0.31, 0.15),  $I^2 = 0.0\%$ ] (Fig. 7a, b). In addition, we failed to find a significant effect of probiotic supplementation on serum levels of IL-17 based on three studies [SMD 0.06; 95% CI (- 0.34, 0.46),  $I^2 = 0.0\%$ ] (Fig. 7c). However, probiotic supplementation resulted in a lower concentration on IL-4 concentrations compared to placebo [SMD - 0.48; 95% CI (- 0.76, - 0.20),  $I^2 = 0.0\%$ ] (Fig. 7d).

**Discussion**

In the current study, we found a significant reduction in serum hs-CRP, TNF-a, IL-6, IL-12, and IL-4 concentrations after probiotic supplementation. In addition, serum concentrations of IL-10 were significantly increased following probiotic supplementation. However, no significant effects of probiotic supplementation on IL-1B, IL-8, IFN-g, and IL-17 concentrations were found in our meta-analysis.

We found a significant reduction in some inflammatory cytokines including hs-CRP, TNF-a, IL-6, IL-12, and IL-4 after probiotic supplementation. A meta-analysis of RCTs on the effect of probiotic supplementation on serum CRP concentrations, published in 2017, showed a significant reduction in serum levels after probiotic intake [59]. Probiotic supplementation resulted in reduced TNF-a concentrations in patients with non-alcoholic fatty liver disease, as found in



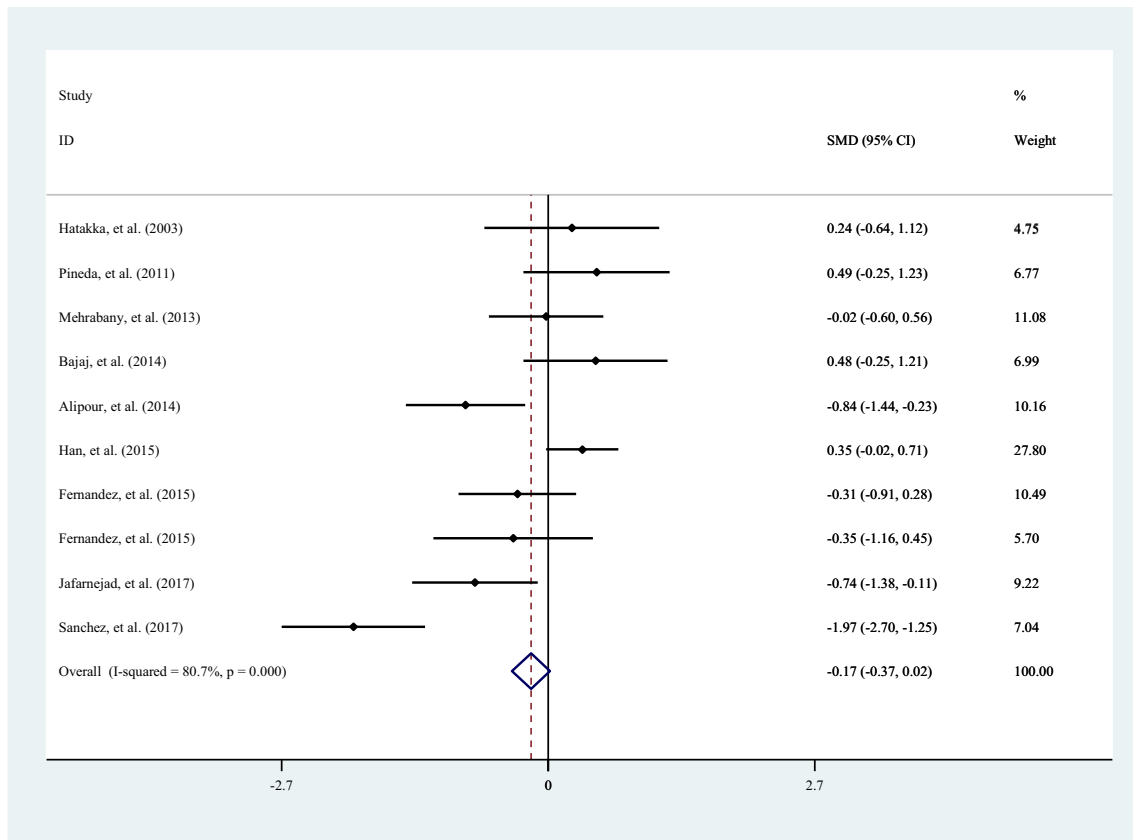
**Fig. 4** Forest plot for the effect of probiotics supplementation on serum IL-10 concentrations, expressed as the mean differences between the intervention and the control diets. The area of each

square is proportional to the inverse of the variance of the SMD. Horizontal lines represent 95% CIs. Diamonds represent pooled estimates from fixed-effects analysis

a meta-analysis [60]. Among patients with colorectal cancer, probiotic supplementation led to reduced serum concentrations of CRP, but changes in serum levels of IL-6 were not significant [61]. However, sample sizes of included studies in that meta-analysis were small. In addition, probiotic supplementation did not change serum concentrations of inflammatory cytokines including TNF- $\alpha$ , IL-1 $\beta$ , IL-6, IL-10, and IL-12 in patients with rheumatoid arthritis [62]. However, only four studies were included in that study. Although a significant reduction in IL-4 concentrations after probiotic supplementation was found in our study, the number of included studies in this area was low. Further studies are needed to reach a firm conclusion.

The effect of probiotic supplementation on hs-CRP concentrations was not significant in studies conducted in patients with metabolic diseases. Chronic low-grade inflammation plays a pivotal role in the pathogenesis of these conditions [63]. Most studies performed on these patients did not consider baseline status of inflammation in their analyses. Moreover, due to high concentrations of inflammatory cytokines in these patients [63], it seems that

interventions with higher doses of probiotics for longer periods would be required to observe the affect on inflammation. In this meta-analysis, the effect of probiotics on serum CRP and TNF- $\alpha$  concentrations was not significant among patients with allergy and autoimmune diseases. In addition, no significant effect of probiotic supplementation was seen on TNF- $\alpha$  concentrations among studies conducted on healthy participants. However, studies in these areas were relatively rare and between-study heterogeneity was also high. Probiotic supplementation at a dose of 10–100 CFU/day did not significantly change serum concentrations of CRP, TNF- $\alpha$ , and IL-6. This was also the case with a dosage of < 1 CFU/day for CRP concentrations. Different responses to probiotic supplementation in different health conditions might be due to changes in gut microbiota, in particular due to the severity and direction of immune system stimulation through the gut-associated lymphoid tissue (GALT) [64, 65]. Data for the effects of each strain of probiotics on serum concentrations of cytokines were insufficient, and a considerable number of included studies had used multiple strains. However,



**Fig. 5** Forest plot for the effect of probiotics supplementation on serum IL-1B concentrations, expressed as the mean differences between the intervention and the control diets. The area of each

square is proportional to the inverse of the variance of the SMD. Horizontal lines represent 95% CIs. Diamonds represent pooled estimates from fixed-effects analysis

our subgroup analysis showed that bifidobacter had higher impact on hs-CRP and IL-6 levels compared to lactobacillus, while saccharomyces did not have a significant effect. Due to the limited number of studies in this regard, further studies are needed to confirm these findings and to explore probable mechanisms of action.

A significant increase in serum IL-10 concentrations was seen after probiotic supplementation. In a recent meta-analysis on the effect of probiotic administration on hs-CRP concentrations, the investigators did not find a significant effect of supplementation on IL-10 levels [59]. However, the main aim of that meta-analysis was focusing on serum hs-CRP levels and they did not include all studies that examined the effect on IL-10 concentrations. In addition, in another meta-analysis on 4 published papers on the effect of probiotic supplementation on serum concentrations of IL-10 in patients with rheumatoid arthritis, no significant effect was reported [62]. IL-10 has anti-inflammatory properties [66]. Several studies in animal models have shown that IL-10 concentrations are increased after probiotic consumption [67, 68]. However, as the significant effect of probiotic supplementation on IL-10 concentrations disappeared in subgroup

analyses, it seems that further studies are needed to shed light in this issue.

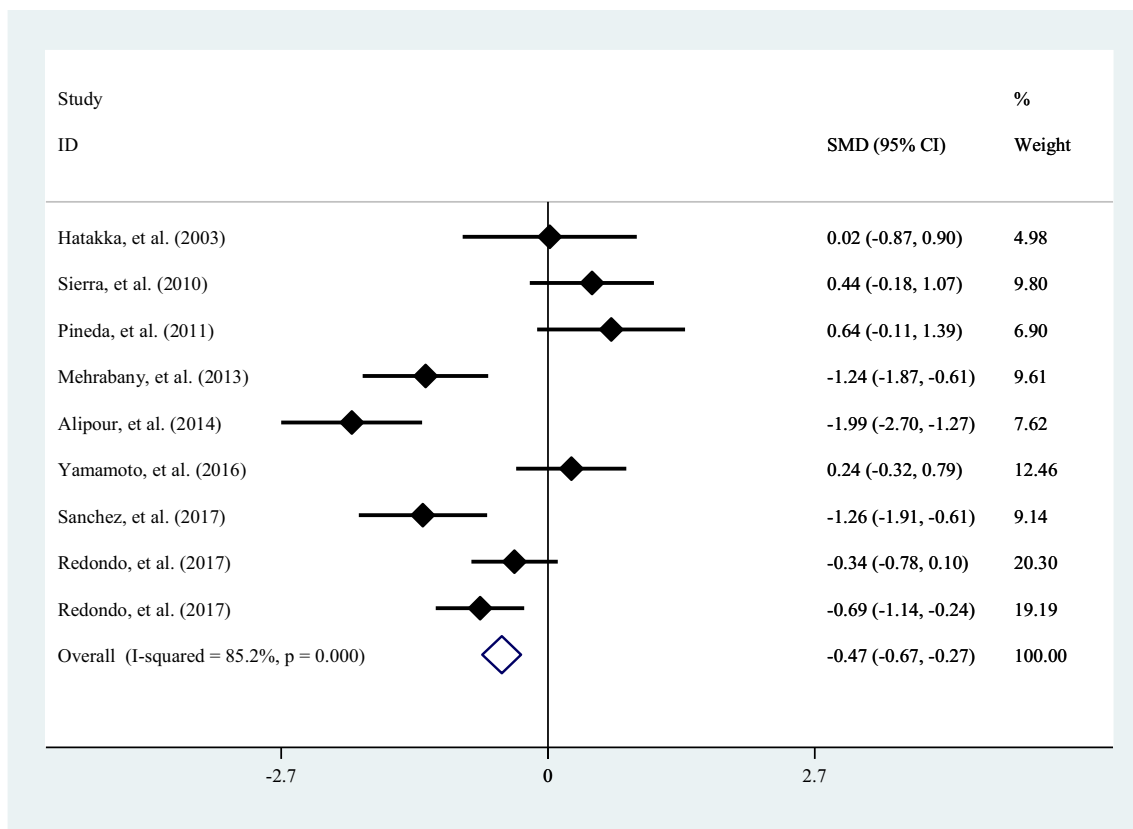
The effect of probiotic supplementation on IL-1B, IL-8, IFN-g, and IL-17 concentrations was not significant in our meta-analysis. Findings from a meta-analysis on the effect of probiotics on cytokines in patients with rheumatoid arthritis revealed no significant effect on serum levels of IL-1B [62]. Further studies using different doses of probiotic supplements are recommended for the future investigations.

Gut is the main reservoir of antigens [69]. It is involved in immune function [61]. The role of gut in inflammation might be partially due to the presence of GALT, which is a lymphoid organ containing the majority of total lymphocytes in body [70]. Previous studies have shown that administration of probiotics can modulate gut immune function by restoring normal function of mucosal barrier [71]. In addition, probiotics may skew immune responses towards immunoregulation by inducing T-reg cells [72]. Lack of microbial antigens-induced immune stimulation in gut deviates cytokine profile from Th1- to Th2-related types [73]. Therefore, it is assumed that consumption of probiotics will alter microbial population in gut and subsequently regulate

**Table 3** Subgroup analysis for the effect of probiotics supplementation on serum concentrations of IL-1B and IL-12

Subgroup	Number of effect sizes		Participants, N		Effect size (95% CI)		$I^2$		$P$ within <sup>a</sup>		$P$ between <sup>2</sup>	
	IL-1B	IL-12	IL-1B	IL-12	IL-1B	IL-12	IL-1B	IL-12	IL-1B	IL-12	IL-1B	IL-12
Overall	10	9	442	439	-0.17 (-0.37, 0.02)	-0.47 (-0.67, -0.27)	80.7	85.2	<0.001	<0.001	-	-
Participants' age												
Adult	3	5	95	268	-0.50 (-0.94, -0.06)	-0.43 (-0.68, -0.18)	92.1	75.3	<0.001	<0.01	0.11	0.60
Adult + elderly	7	4	347	171	-0.10 (-0.31, 0.12)	-0.54 (-0.86, -0.21)	67.9	92.0	<0.01	<0.001		
Supplement dosage												
< 1 CFU/day	4	3	160	132	-0.37 (-0.69, -0.06)	-0.84 (-1.22, -0.46)	19.2	92.7	0.29	<0.001	<0.01	<0.01
≥ 1 CFU/day	5	5	165	257	-0.37 (-0.70, -0.04)	-0.45 (-0.70, -0.20)	87.6	75.9	<0.001	<0.01		
NR	1	1	117	50	0.35 (-0.02, 0.71)	0.24 (-0.32, 0.79)	-	-	-	-		
Study duration												
< 10 week	7	6	351	339	-0.19 (-0.40, 0.03)	-0.72 (-0.94, -0.49)	84.9	85.1	<0.001	<0.001	0.76	<0.001
≥ 10 week	3	3	91	100	-0.12 (-0.54, 0.31)	0.31 (-0.09, 0.70)	71.2	0.0	0.03	0.53		
Health condition												
Healthy	-	3	-	203	-	-0.32 (-0.60, -0.04)	-	75.8	-	0.01	0.41	<0.01
Gastrointestinal disease	3	1	191	44	-0.02 (-0.32, 0.28)	-1.26 (-1.91, -0.61)	65.3	-	<0.001	-		
Skeletal disorders	5	4	183	142	-0.26 (-0.56, 0.03)	-0.77 (-1.14, -0.41)	94.0	90.0	0.02	<0.001		
Allergy and autoimmune diseases	-	1	-	50	-	0.24 (-0.32, 0.79)	-	-	-	-		
Other diseases	2	-	68	-	-0.33 (-0.81, 0.15)	-	0.0	-	0.93	-		
Participants' sex												
Female	3	2	133	94	-0.51 (-0.86, -0.16)	-1.57 (-2.04, -1.10)	54.2	57.5	0.11	0.12	0.02	<0.001
Both genders	7	7	309	345	-0.03 (-0.26, 0.21)	-0.24 (-0.46, -0.02)	83.8	77.4	<0.001	<0.001		

<sup>a</sup> $P$  values were obtained by fixed-effect analysis



**Fig. 6** Forest plot for the effect of probiotics supplementation on serum IL-12 concentrations, expressed as the mean differences between the intervention and the control diets. The area of each

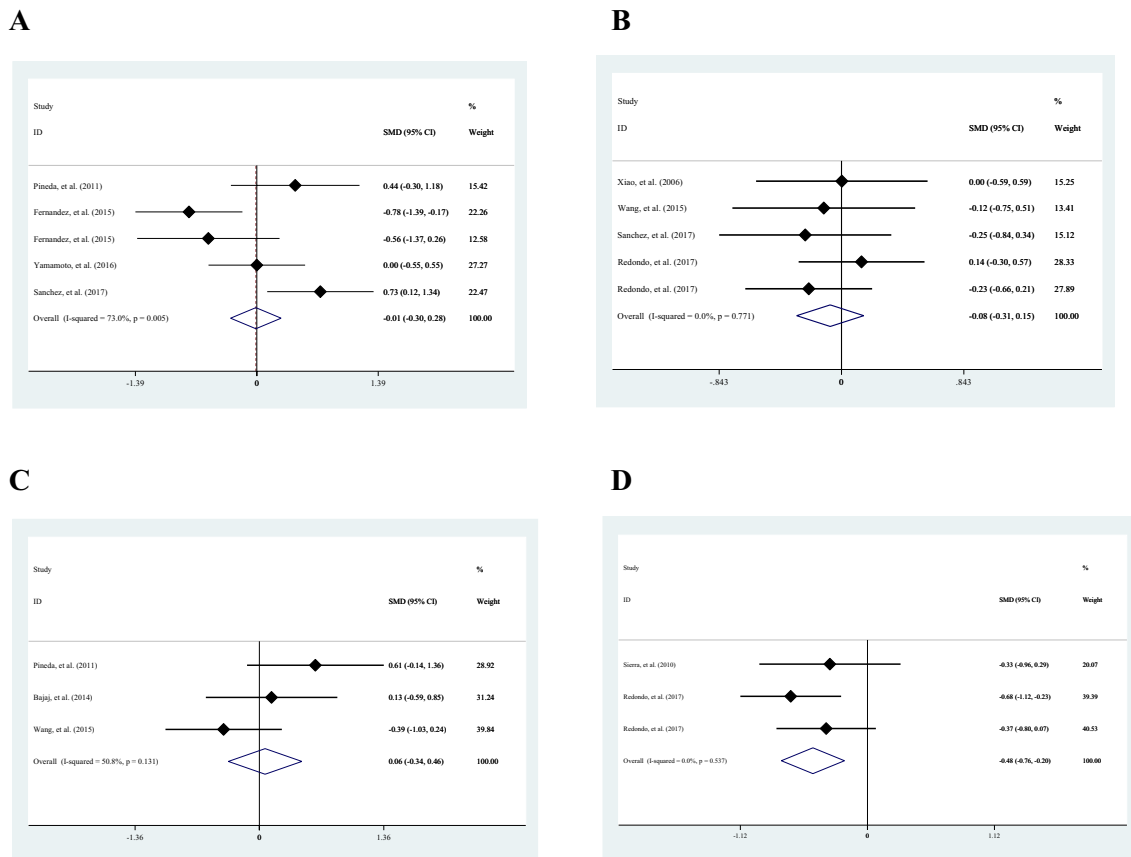
square is proportional to the inverse of the variance of the SMD. Horizontal lines represent 95% CIs. Diamonds represent pooled estimates from fixed-effects analysis

secretion of inflammatory cytokines towards the reduced body inflammation.

This is the first comprehensive meta-analysis on the effect of probiotics on various inflammatory cytokines. All studies we included were RCTs. Some limitations should be taken into account. Probiotics were used in different dosages in included studies. Moreover, different strains of probiotics were used. We tried to take in account this problem using subgroup analysis based on probiotic type. The wide ranges of intervention periods were another possible source of bias in this meta-analysis. In addition, participants in included studies were in different physiological status. Therefore, future studies should be separately conducted in participants with different health conditions. Furthermore, most included studies did not adjust their findings for the baseline levels of cytokines.

## Conclusion

In conclusion, we found a significant reduction in serum concentrations of pro-inflammatory cytokines, including hs-CRP, TNF- $\alpha$ , IL-6, IL-12, and IL-4 after probiotic supplementation. Moreover, serum concentrations of IL-10, as an anti-inflammatory agent, were significantly increased. The effects of probiotic supplementation on serum IL-1B, IL-8, IFN-g, and IL-17 concentrations were not significant. Therefore, it seems that intake of probiotics might considerably reduce inflammation in humans. Further studies measuring the effects of probiotics on serum concentrations of other inflammatory biomarkers are needed to confirm our conclusion.



**Fig. 7** Forest plot for the effect of probiotics supplementation on serum IL-8 (a), IFN-g (b), IL-17 (c), and IL-4 (d) concentrations, expressed as the mean differences between the intervention and the

control diets. The area of each square is proportional to the inverse of the variance of the SMD. Horizontal lines represent 95% CIs. Diamonds represent pooled estimates from fixed-effects analysis

**Author contributions** AM, BL, and AE designed research; AM, SMM, AS, ASM, and MP conducted research; AM analyzed data; AM and AE wrote the paper; and AE had primary responsibility for final content. All authors read and approved the final manuscript.

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### Compliance with ethical standards

**Conflict of interest** The authors have no conflicts of interest to declare.

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