<u>META-ANALYSIS</u>

Potential Value of Probiotics on Lipid Profiles in Hyperlipidemia and Healthy Participants: Systematic Review and Meta-Analysis

Dongdong Su, MD; Yan Liu, MD; Li Zhang, PhD; Shanshan Zhao, PhD; Yifei Wang, MD; Rutao Bian, PhD; Bianling Xu, PhD; Xiaoyang Chen, PhD; Xuegong Xu, PhD

ABSTRACT

Objective • Many randomized controlled trials (RCTs) have reported the effect of probiotics on reducing plasma lipids with inconsistent results. An explicit systematic review and meta-analysis were conducted in this study to evaluate the effect of probiotics on the lipid profile of healthy and hyperlipidemia participants.

Methods • A comprehensive literature search of RCTs was conducted using PubMed, Embase, World Health Organization (WHO) Global Index Medicus, WHO clinical trial registry, and Clinicaltrials.gov. Inclusion criteria included RCTs comparing the use of any strain of a specified probiotic with the placebo control group. The change in lipid profiles was analyzed.

Results • The probiotics can decrease the total cholesterol (TC) level in hyperlipidemia participants but not healthy persons (MD = -0.43, 95% CI -0.60 - -0.25, P < .01; MD = -0.09, 95% CI -0.26 - 0.08, P > .05). Probiotics did not reduce high-density lipoprotein cholesterol (HDL-C) in

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Corresponding author: Xiaoyang Chen, PhD E-mail: doctor_chan@stu.gzucm.edu.cn Corresponding author: Xuegong Xu, PhD E-mail: xuxg1115@126.com patients with hyperlipidemia or healthy people (MD = -0.01, 95% CI -0.09 – 0.07, P > .05; MD = 0.02, 95% CI -0.04 – 0.09, P > .05). Furthermore, probiotics can reduce the low-density lipoprotein cholesterol (LDL-C) level both in hyperlipidemia and healthy persons (MD = -0.34, 95% CI -0.43 – -0.26, P < .01; MD = -0.15, 95% CI -0.28 – -0.02, P < .05). Lastly, the effect of probiotics on reducing triglyceride (TG) levels was significant in hyperlipidemia persons but not in the healthy population (MD = -0.20, 95% CI -0.37 – -0.04, P < .01; MD = -0.01, 95% CI -0.02 – 0.04, P > .05).

Conclusions • Through our analysis, the effect of probiotics on lowering plasma lipid was more obvious in hyperlipidemia participants than healthy population. However, further studies are required to confirm the findings due to pronounced clinical heterogeneity. (*Altern Ther Health Med.* [E-pub ahead of print.])

INTRODUCTION

Hyperlipidemia refers to an imbalance of cholesterol in the blood, including low-density lipoprotein cholesterol (LDL-C) and high-density lipoprotein cholesterol (HDL-C). They regulate cholesterol levels in the blood, and cholesterol imbalances may increase the risk of cardiovascular and cerebrovascular diseases, including myocardial infarction and stroke. Other forms of hyperlipidemia include hypertriglyceridemia as well as mixed hyperlipidemia, in which both cholesterol and triglyceride (TG) levels are elevated.¹ The development of hyperlipidemia is related to many factors, such as high-fat diet, smoking, chronic kidney disease, diabetes, hypertension, and genetic factors.² Taking targeted preventive measures against these risk factors is the key to preventing hyperlipidemia and secondary cardiovascular events.

Animal experiments suggest that intestinal flora plays an important role in regulating lipid metabolism in the host. The disturbance of lipid metabolism could induce changes in the intestinal environment, which could induce dysbiosis of internal microflora. Normal gut microbiota is essential in maintaining lipid-metabolism homeostasis, including mediating satiety in the brain through microglia and vagus nerves, regulating cholesterol metabolism in the liver, promoting lipid oxidation in the muscle and energy storage in the adipose tissue, and maintaining the integrity of the intestinal barrier.³⁻⁵ Some related interventions, such as fecal microbiological transplantation and probiotics, have achieved good results in the treatment of hyperlipidemia.⁶ Probiotics are active microorganisms that are beneficial to the host by regulating the immune function of the host mucosa and the system, and by regulating the balance of intestinal flora.^{6,7} It has been revealed that several probiotic strains are able to improve at least one lipid fraction in both animal and human models.⁸

Many randomized controlled trials (RCTs) have reported the effect of probiotics on reducing plasma cholesterol levels with inconsistent results,⁹⁻²⁶ most likely due to variations in experiment design and methodological measurements. Some recent studies have been published to demonstrate the benefits of probiotics. However, they did not compare the cholesterol-lowering effects of probiotics in healthy individuals and those with hyperlipidemia,²⁷ and just considered the effect of probiotics on total cholesterol.²⁸ Therefore, an explicit systematic review and meta-analysis are needed to evaluate the effect of probiotics on reducing plasma cholesterol levels in healthy and hyperlipidemia participants.

MATERIALS AND METHODS

Study Selection

All RCTs evaluating the effect of probiotics on lipid profiles in healthy and hyperlipidemia persons were searched using PubMed (1966-2022), Embase (1980-2022), and World Health Organization (WHO) Global Index Medicus. Unpublished or ongoing studies were identified by checking clinical trial registers through Clinicaltrials.gov and WHO clinical trial registry. Literature in all languages was included in the search. Meta-analyses and systematic reviews were also hand-searched to find relevant literature that might have been missed by the initial search. Logical combinations of "probiotics" and "hyperlipidemia" were used as keywords to search for relevant literature. RCTs of any kind of probiotics and dose were accepted, either in healthy persons or hyperlipidemia patients, who did not have underlying diseases such as cardiovascular disease, kidney disease, and diabetes. Pregnant women were excluded from the study. Control groups were those that did not receive any probiotics.

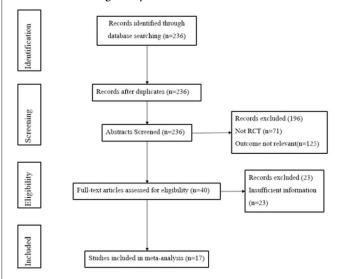
Data Extraction

Articles retrieved from the searches were evaluated independently by 2 reviewers (Bianling Xu and Shanshan Zhao) using predefined standardized data extraction forms, and then data were evaluated by a third reviewer (Li Zhang) independently based on the US National Institute of Health National Heart, Lung, and Blood Institute (NHLBI) study quality assessment tool for controlled intervention studies.²⁹ Clinical outcome of interest was the changes in lipid profiles after administration of probiotics compared to the control. Data pertaining to the participants, the kinds of probiotics used, control groups, and methodology were extracted (Table 1).

Table 1. Characteristics of All Randomized Control Trials Evaluating the Use of Probiotics on Lipid Profiles in Healthy and Hyperlipidemia Participants (1966–2022)

			No. of parti (Male/Fer	Follow- up period		
Study name	Participants	Probiotics	Intervention			
Agerbaek 1995	Healthy	S. thermophilus, E. faecium	29/0	28/0	6	
Ataie 2009	Hyperlipidemia	L. acidophilus, B. lactis	4/10	4/10	6	
Bertolami 1999	Hyperlipidemia	E. faecium, S. thermophilus	11/21	11/21	8	
Greany 2008	Healthy	L. acidophilus, B. longum	15/22	7/11	8	
Simons 2006	Hyperlipidemia	L. fermentum	8/15	8/13	10	
Sadrzadeh 2010	Healthy	L. bulgaricus, L. acidophilus, S. thermophilius, B. lactis	0/30	0/30	6	
Jones 2012	Hyperlipidemia	L. reuteri	28/39	27/34	9	
Anderson 1999	Hyperlipidemia	L. acidophilus	9/12	9/10	4	
Fabian 2006	Healthy	S. thermophilus, L. bulgaricus, L. casei	0/17	0/16	4	
Fuentes 2013	Hyperlipidemia	L. plantarum	30	30	12	
Xiao 2003	Healthy	B. longum	16/0	16/0	4	
Valentini 2015	Healthy	VSL#3	31	31	8	
Rerksuppaphol 2015	Hyperlipidemia	L. acidophilus, B. bifidum	12/19	10/23	6	
Cavallini 2016	Hyperlipidemia	E. faecium, L. helveticus	17/0	15/0	6	
Chiu 2021	Hyperlipidemia	L. acidophilus, L. casei, B. lactis	20	20	12	
Guerrero 2021	Hyperlipidemia	L. plantarum	18	18	12	
Lee 2017	Healthy	B. animalis, B. lactis	26	25	4	

Figure 1. PRISMA Flow Diagram Showing the Process of Literature Screening, Study Selection, and Reasons for Exclusion



Meta Analysis

The Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) statement methodology³⁰ was adhered to. The mean difference (MD) with a 95% CI for lipid profiles of each trial was calculated to estimate treatment effects. Meta-analysis of the pooled data was performed using the fixed-effect model or random-effect model, depending upon the heterogeneity of the included studies. If clinical heterogeneity was observed, data was analyzed using a random-effect model. Heterogeneity was quantified using the Cochrane's Q statistic and statistics, with the values of 25%, 50%, and 75% signifying the limits of low, moderate, and high statistical heterogeneity, respectively.³¹ A funnel plot was used to explore publication bias for the studies.³² A two-tailed P < .05 was considered statistically significant. All statistical analyses were performed using the R package meta (R version 4.0.1). The risk of bias was evaluated using the

Cochrane risk of bias tool. It was used to evaluate the selection bias, performance bias, detection bias, attrition bias, reporting bias, and other biases.

RESULTS

Demographic Characteristics of the Studies

The literature search process, shown in Figure 1, identified 236 potential studies for full analyses. Following the application of exclusion criteria, 17 studies were identified as appropriate for further quantitative meta-analyses (Table 1). Of 17 studies included in the final analysis, 7 studies included healthy participants, while 10 studies were conducted in hyperlipidemia patients. The probiotic strains used were *L. acidophillus, L. casei, L. paracasei, L. bulgaricus, L. fermentum, L. helveticus, L. paracasei, L. bulgaricus, S. thermophilus, E. faecium.* The follow-up period ranged from 4 weeks to 12 weeks. The present study compared lipid changes at the end of follow-up between the intervention and treatment groups.

Effects of probiotics on total cholesterol (TC) in healthy and hyperlipidemia participants

Overall, probiotics can lower the level of TC as compared to the control group who did not receive probiotics (MD = -0.28, 95 % CI -0.43 - -0.13, P < .01; Figure 2). There was significantly high heterogeneity between the trials ($I^2 = 86\%$). Of 17 studies included in the final analysis, 7 studies involved healthy participants, while 10 studies were conducted in hyperlipidemia patients. In the healthy subgroup, no significant difference was observed in the level of TC compared to the control group (MD = -0.09, 95% CI -0.26 – 0.08, P > .05; Figure 2). High heterogeneity was observed between the studies ($I^2 = 82\%$). In the hyperlipidemia subgroup, the meta-analysis showed a significantly lower level of TC compared to the control group (MD = -0.43, 95%CI -0.60 – -0.25, P < .01; Figure 2). There is moderate heterogeneity between the studies ($I^2 = 69\%$). The subgroup difference between healthy and hyperlipidemia participants was significant (P < .01).

Effects of probiotics on HDL-C in healthy and hyperlipidemia participants

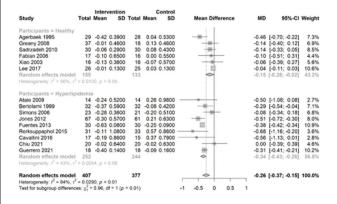
Overall, probiotics cannot lower the level of HDL-C as compared to the control group who did not receive probiotics (MD = 0.00, 95 %CI -0.05 – 0.05, P > .05; Figure 3). There was high heterogeneity between the trials ($I^2 = 88\%$). Of 15 studies included in the final analysis, 6 studies involved healthy participants, while 9 studies were conducted in hyperlipidemia patients. In the healthy subgroup, the level of HDL-C showed no significant difference compared to the control group (MD = 0.02, 95% CI -0.04 – 0.09, P > .05; Figure 3). Moderate heterogeneity was observed between the studies ($I^2 = 71\%$). In the hyperlipidemia subgroup, the metaanalysis showed no significant difference in the HDL level compared to the control group (MD = -0.01, 95% CI -0.09 – **Figure 2.** Forest Plot of Probiotics from Randomized Controlled Trials Demonstrating the Effect on the Mean Difference of Total Cholesterol in Healthy and Hyperlipidemia Participants

		Inten	tervention			Control				
Study	Total	Mean	SD	Total	Mean	SD	Mean Difference	MD	95%-CI	Weight
Participants = Healthy							1.1			
Agerbaek 1995	29	-0.37	0.4100	28	-0.30	0.6300		-0.07	[-0.35; 0.21]	6.2%
Greany 2008	37	-0.07	0.5200	18	0.07	0.5200		-0.14	[-0.43; 0.15]	6.1%
Sadrzadeh 2010	30	-0.20	0.3300	30	0.18	0.5100		-0.38	[-0.60; -0.16]	6.8%
abian 2006	17	0.10	0.8000	16	0.00	0.7000		0.10	[-0.41; 0.61]	4.1%
Gao 2003	16	-0.17	0,4000	16	-0.02	0.4600		-0.15	[-0.45: 0.15]	6.0%
/alentini 2015	31	-0.16	0.2100	31	-0.34	0.1900	10	0.18	[0.08; 0.28]	7.7%
.00 2017	26	-0.08	0.1600	25	0.05	0.1600	-	-0.13	[-0.22; -0.04]	7.7%
Random effects model	186			164			~	-0.09	[-0.26: 0.08]	44.5%
Heterogeneity: $l^2 = 82\%$, τ^2	= 0.03	65, p <	0.01							
Participants = Hyperlip	idemia	1								
Ataie 2009	14	-0.32	0.0300	14	0.36	1.0200		-0.68	[-1.21; -0.15]	3.9%
Bertolami 1999	32	-0.32	0.6700	32	0.03	0.4400		-0.35	[-0.63; -0.07]	6.2%
Simons 2006	23	-0.38	0.5000	21	-0.31	0.4400	+==	-0.07	[-0.35; 0.21]	6.2%
Jones 2012	67	-0.42	0.6300	61	0.16	0.6700		-0.58	[-0.81; -0.35]	6.7%
Anderson 1999	21	-0.27	0.5000	19	-0.05	0.6100		-0.22	[-0.57; 0.13]	5.5%
Fuentes 2013	30	-0.87	0.7000	30	-0.28	0.3400		-0.59	[-0.87; -0.31]	6.2%
Rerksuppaphol 2015	31	-0.64	0.9500	33	0.57	0.7700		-1.21	[-1.64; -0.78]	4.8%
Cavallini 2016	17	-0.35	0.9400	15	-0.10	0.7700		-0.25	[-0.84; 0.34]	3.5%
Chiu 2021	20	-0.04	0.5500	20	-0.01	0.7100	+	-0.03	[-0.42: 0.36]	5.1%
Suerrero 2021	18	-0.47	0,1600	18	-0.07	0.2400		-0.40	[-0.53: -0.27]	7.5%
Random effects model	273			263			\$	-0.43	[-0.60; -0.25]	55.5%
Heterogeneity: $l^2 = 69\%$, τ^2	= 0.04	74, p <	0.01							
Random effects model	459			427			\$	-0.28	[-0.43; -0.13]	100.0%
leterogeneity: 12 = 86%, r2	= 0.07	21. p <	0.01							
Test for subgroup difference	es: y2 =	7.53. 0	f = 1 (p	< 0.01)			1.5 -1 -0.5 0 0.5 1	1.5		

Figure 3. Forest Plot of Probiotics from Randomized Controlled Trials Demonstrating the Effect on the Mean Difference of High-Density Lipoprotein in Healthy and Hyperlipidemia Participants

		Interventio Mean SI	n D Total	Control Mean SD	Mean Difference	MD	95%-CI	
Study	Total	Mean S	lotal	Mean SD	Mean Difference	MD	95%-CI	weign
Participants = Healthy								
Agerbaek 1995	29	0.02 0.130	28	-0.01 0.1600		0.03 [-	0.05; 0.11]	8.6%
Greany 2008	37	-0.03 0.180	0 18	0.02 0.1800		-0.05 [-	0.15; 0.05]	7.5%
Sadrzadeh 2010	30	0.11 0.150	0 30	-0.01 0.1300		0.12 [0.05; 0.19]	8.89
Fabian 2006	17	0.20 0.400	0 16	0.10 0.3000		0.10 [-	0.14: 0.34]	3.19
Xiao 2003	16	-0.07 0.350	0 16	-0.07 0.4000		0.00 [-	0.26; 0.26]	2.89
Lee 2017	26	0.01 0.050	0 25	0.04 0.0500		-0.03 [-	0.06: 0.001	10.4%
Random effects model	155		133		\diamond	0.02 [-	0.04: 0.091	41.39
Heterogeneity: $l^2 = 71\%$, τ^2	= 0.00	39, p < 0.01						
Participants = Hyperlip	idemia	1						
Ataie 2009	14	0.01 0.140	0 14	0.15 0.2000		-0.14 [-1	0.27; -0.01]	6.49
Simons 2006	23	-0.05 0.110	0 21	-0.07 0.1800	-10-	0.02 [-	0.07; 0.11]	8.09
Jones 2012	67	0.00 0.240	0 61	0.01 0.2900		-0.01 [-	0.10; 0.08]	7.99
Anderson 1999	21	0.00 0.340	0 19	0.03 0.4100		-0.03 [-	0.26; 0.20]	3.29
Fuentes 2013	30	-0.08 0.030	0 30	0.01 0.0500		-0.09 [-1	0.11: -0.07]	10.59
Rerksuppaphol 2015	31	-0.12 0.350	0 33	0.04 0.4400		-0.16 [-	0.35; 0.03]	4.29
Cavallini 2016	17	-0.10 0.220	0 15	-0.37 0.3300		- 0.27 [0.07; 0.47]	4.19
Chiu 2021	20	-0.00 0.350	0 20	0.02 0.2800		-0.02 [-	0.22; 0.17]	4.19
Guerrero 2021	18	0.08 0.030	0 18	0.01 0.0500			0.04; 0.10]	10.49
Random effects model	241		231		-	-0.01 [-0	0.09; 0.07]	58.7%
Heterogeneity: $I^2 = 92\%$, τ^2	= 0.01	03, <i>p</i> < 0.01						
Random effects model	396		364		4	0.00 [-4	0.05; 0.05]	100.05
Heterogeneity: $l^2 = 88\%$, τ^2	= 0.00	63. p < 0.01						
Test for subgroup difference	os: x2 =	0.43, df = 1 (r	= 0.51)		-0.4 -0.2 0 0.2 0	.4		

Figure 4. Forest Plot of Probiotics from Randomized Controlled Trials Demonstrating the Effect on the Mean Difference of Low-Density Lipoprotein in Healthy and Hyperlipidemia Participants

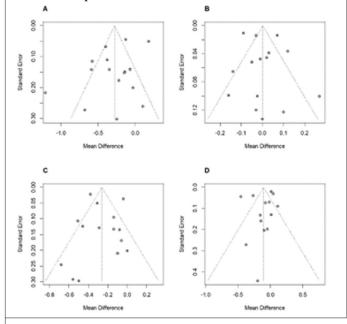


0.07, P > .05; Figure 3). There is high heterogeneity between the studies ($I^2 = 92\%$). The subgroup difference between healthy and hyperlipidemia participants was not significant (P = .51).

Figure 5. Forest Plot of Probiotics from Randomized Controlled Trials Demonstrating the Effect on the Mean Difference of Triglyceride in Healthy and Hyperlipidemia Participants

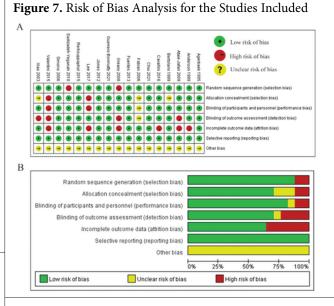
		Interv	ention			Control						
Study	Total	Mean	SD	Total	Mean	SD		Mean Difference	MC	9	5%-CI	Weight
Participants = Hyperlip	idemia							11				
Ataie 2009	14	-0.22	0.6800	14	-0.12	0.3400			-0.10	[-0.50;	0.30]	4.6%
Simons 2006	23	-0.03	0.4500	21	0.13	0.4200			-0.16	1-0.42	0,101	6.8%
Jones 2012	67	-0.02	0.7100	61	0.13	1.0400			-0.15	1-0.46	0.161	5.8%
Fuentes 2013	30	-0.33	0.1400	30	-0.06	0.1600				[-0.35;		
Rerksuppaphol 2015	31	-0.32	1.3900	33	0.06	0.6100				1-0.91		
Cavallini 2016	17	-0.02	0.9100	15	0.18	1.4800			-0.20	[-1.06;	0.661	1.5%
Chiu 2021	20	0.08	0.2400	20	-0.03	0.3200				1-0.07		
Guerrero 2021	18	-0.34	0.1300	18	0.12	0.1300			-0.46	1-0.54	-0.381	9.8%
Random effects model	220			212						[-0.37:		
Heterogeneity: $I^2 = 82\%$, τ^2	= 0.03	29, p < 0	0.01									
Participants = Healthy												
Greany 2008	37		0.4500	18		0.4500				[-0.24;		
Sadrzadeh 2010	30		0.3000	30		0.2600				3 [-0.22;		
Fabian 2006	17		0.2000	16		0.2000		*		2 [-0.16;		
Xiao 2003	16		0.5100	16		0.6000			-0.05	5 [-0.44;	0.34]	
Valentini 2015	31	-0.01	0.1000	31	-0.05	0.1300		1.00	0.04	[-0.02;	0.10]	10.1%
Lee 2017	26	-0.01	0.0700	25	-0.02	0.0700		10	0.01	[-0.03;	0.05]	10.3%
Random effects model	157			136				•	0.0	[-0.02;	0.04]	50.0%
Heterogeneity: $I^2 = 0\%$, τ^2	= 0, p =	0.73										
Random effects model				348				\$	-0.1	[-0.22;	0.01]	100.0%
Heterogeneity: I2 = 91%, r2												
Test for subgroup difference	es: y2 =	6.67, df	= 1 (p	< 0.01)			-1 -4	0.5 0 0.5	1			

Figure 6. Funnel Plots of Included Randomized Controlled Trials Demonstrating the Treatment Effect Relative to Study Size in (A) Total Cholesterol, (B) High-Density Lipoprotein, (C) Low-Density Lipoprotein, and (D) Triglyceride Solid black dots represent the true effect values.



Effects of probiotics on LDL in healthy and hyperlipidemia participants

Overall, the probiotics can lower the level of LDL-C as compared to the control group who did not receive probiotics (MD = -0.26, 95% CI -0.37 – -0.15, P < .05; Figure 4). There was high heterogeneity between the trials ($I^2 = 84\%$). Of 15 studies included in the final analysis, 6 studies involved healthy participants, while 9 studies were conducted in hyperlipidemia patients. In the healthy subgroup, the level of LDL-C showed significantly lower differences compared to the control group (MD = -0.15, 95% CI -0.28 – -0.02, P < .05; Figure 4). Moderate heterogeneity was observed between the studies ($I^2 = 56\%$). In the hyperlipidemia subgroup, the meta-analysis showed a significantly lower level of LDL-C compared to the control group (MD = -0.34, 95% CI -0.43 – -0.26, P < 1000



.01; Figure 4). There is low heterogeneity between studies $(I^2 = 43\%)$. The subgroup difference between healthy and hyperlipidemia participants was significant (P = .01).

Effects of probiotics on TG in healthy and hyperlipidemia participants

Overall, probiotics cannot lower the level of TG in hyperlipidemia participants as compared to the control group who did not receive probiotics (MD = -0.11, 95% CI -0.22 - 0.01, P > .05; Figure 5). There was high heterogeneity between the trials ($I^2 = 91\%$). Of 14 studies included in the final analysis, 6 studies were performed involving healthy participants, while 8 studies were conducted in hyperlipidemia patients. In the healthy subgroup, no significant difference was observed in the level of TG compared to the control group (MD = -0.01, 95% CI -0.02 – 0.04, P > .05; Figure 5). No heterogeneity was observed between the studies ($I^2 = 0\%$). In the hyperlipidemia subgroup, the meta-analysis showed a significantly lower level of TG compared to the control group (MD = -0.20, 95% CI - 0.37 - -0.04, P < .01; Figure 5). There is a high heterogeneity between the studies ($I^2 = 82\%$). The subgroup difference between healthy and hyperlipidemia participants was significant (P < .01).

Publication Bias

Funnel plots were used to visually assess for publication bias in the treatment effect of probiotics on TC, HDL-C, LDL-C, and TG in healthy and hyperlipidemia participants (Figure 6). There was minor asymmetry on the funnel plot suggesting that studies are less likely to be published if positive outcomes are demonstrated.

Risk of Bias Analysis

The risk of bias in the studies included is summarized in Figure 7. The Cochrane risk of bias tool was used to evaluate the selection bias, performance bias, detection bias, attrition bias, and other biases.

DISCUSSION

In vitro and *in vivo* experiments as well as cohort studies have shown that probiotics play an important role in lowering blood lipids by regulating intestinal flora.^{33,34} In the present study, we evaluated the effect of probiotics on lipid profiles in healthy and hyperlipidemia participants explicitly and systematically.

Through our analysis, the effect of probiotics on lowering blood lipids is different in hyperlipidemia participants and healthy populations. Probiotics can decrease the TC level in hyperlipidemia participants but not healthy persons as compared to the control group who received placebo (MD = -0.43, 95% CI -0.60 – -0.25, *P* < .01; MD = -0.09, 95% CI -0.26 - 0.08, P > .05, respectively). The subgroup difference between them was significant. The lowering effect of probiotics on HDL-C was not detected either in hyperlipidemia participants or in healthy persons (MD = -0.01, 95% CI -0.09 - 0.07, P > .05; MD = 0.02, 95% CI -0.04 -0.09, P > .05, respectively). It can be concluded that the probiotics can reduce the LDL-C level both in hyperlipidemia and healthy persons (MD = -0.34, 95% CI -0.43 - -0.26, P < .01; MD = -0.15, 95% CI -0.28 – -0.02, *P* < .05, respectively). The subgroup difference between them was significant. Therefore, the effect of probiotics on lowering LDL-C in participants with hyperlipidemia was more obvious than that in healthy population. The effect of probiotics on reducing TG level was significant in hyperlipidemia persons but not in healthy population (MD = -0.20, 95% CI -0.37 - -0.04, P <.01; MD = -0.01, 95% CI -0.02 - 0.04, P > .05, respectively).

However, although probiotic use has been greatly popularized among the general public, there are conflicting clinical results for many probiotic strains and formulations.³⁵ Theoretical risks have been described in case reports, clinical trial results, and experimental models, including systemic infections, deleterious metabolic activities, excessive immune stimulation in susceptible individuals, gene transfer, and gastrointestinal side effects.^{36,37} The included studies did not report the incidence of side effects and mortality after administration of probiotics. Therefore, probiotics should be used in consideration of their possible side effects, which need to be confirmed by more studies.

Some limitations of this study should be acknowledged. First, the composition of probiotics used in each study varied, and different probiotic strains or antibiotics may present different values in lowering plasma lipids. Second, the duration of administration of probiotics was different in the studies included, which should be important for the outcomes and could account for the high clinical heterogeneity. Third, we did not distinguish the difference in the lowering lipid effect of probiotics in different genders. Moreover, there was an imbalance in the ratio of men to women between the treatment and control groups in the included studies. Despite these limitations, our findings support the fact that the effects of probiotics on lowering plasma lipids were more obvious in hyperlipidemia participants than in healthy populations. However, further studies are required to be conducted to confirm the findings due to pronounced clinical heterogeneity.

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AUTHOR DISCLOSURE STATEMENT

We declare that we do not have any commercial or associative interest that represents a conflict of interest in connection with the work submitted.

AUTHOR CONTRIBUTIONS

Dongdong Su and Yan Liu contributed equally to this work and should be considered co-first authors. We thank every participant who contributed to the study. The authors' responsibilities were as follows—Xuegong Xu and Xiaoyang Chen: research design; Rutao Bian, Dongdong Su, Yifei Wang, and Yan Liu: data collection; Yan Liu and Dongdong Su: data analysis; Bianling Xu, Shanshan Zhao, and Li Zhang: data assessment; Dongdong Su: Manuscript writing.

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