

Probiotics, Prebiotics and Fruit and Vegetable Fiber Improved the Gastrointestinal Tract and Increased Immunity: A Randomized, Single-Blind, Placebo-Controlled Parallel Trial

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Abstract

The intestinal tract is the largest immune organ in the human body. A healthy gut will block bacteria, viruses, molds, parasites, and other attacks. Studies have shown that probiotics can stabilize the intestinal barrier and improve immunity, while prebiotics can help probiotics multiply and grow effectively in the human gut. Dietary fibers increase the frequency of gastrointestinal motility. This study used the combination formula of probiotics, prebiotics, and fruit and vegetable fiber to explore its effects on gastrointestinal health and improving immunity. The study recruited 50 subjects and randomly assigned two groups, the placebo group and the probiotic fruit and vegetable fiber powder group (PF formula). Subjects were measured in body position measurement, blood and stool collection, dietary questionnaire, and somatosensory questionnaire at weeks 0, and 12. The PF formula increased IgG, IgM, and IL-10 and decreased IL-6, IL-12, IFN- γ , and TNF- α compared to baseline (week 0). The PF formula increased the frequency of bowel movements, improved flatulence, stomach pain, hard stools, constipation, and diarrhea. It also reduced headache, muscle pain, and dizziness compared to baseline. Finally, the PF formula increased the abundance of beneficial bacteria *Porphyromonadaceae* and *Parabacteroides* and decreased the abundance of harmful bacteria *Clostridiaceae*. The PF formula had the effect of gastrointestinal health care and enhanced immunity.

Keywords: Gut microbiota; Fruit and vegetable fiber; Immunity; Probiotics; Prebiotics

Introduction

Immunity helps the body resist virus invasion. When the body's immune system is out of balance, it will affect the overall physical condition, also affect the endocrine system, and reduce the function of immune cells, resulting in decreased immunity and cold symptoms or gastrointestinalrelated problems, such as: stomach ulcers, constipation, diarrhea, etc [1]. The gut microbiota is a complex ecosystem that plays a crucial role in the human gastrointestinal tract, affects nutrient metabolism and the immune system, and even defends against pathogens [2]. At the same time, a good gut microbiota can also regulate the function of the gastrointestinal tract [3]. The most common way to improve is to supplement with probiotics to adjust the balance of bacteria in the gut.

In recent years, studies have shown that the consumption of probiotics can adjust the composition of intestinal bacteria, which produce a slightly acidic environment in the intestinal tract and can inhibit the growth of pathogenic bacteria, rebuild the function of the intestinal mucosal barrier and improve the defense capacity of the digestive tract [4]. Probiotics are derived primarily from human intestinal bacteria, mainly Lactobacillus and Bifidobacterium [5]. Probiotics must be resistant to damage caused by strong acid and a high concentration of bile salts in the digestive tract and can colonize the intestinal mucosa of the host [6]. Some studies showed that L. acidophilus can secrete lactic acid and acetic acid, which can enhance immunity, maintain healthy intestinal flora [4], and L. rhamnosus inhibited the growth of vaginal mold [7]. L. paracasei improved allergies and atopic dermatitis [8]. B. longum can improve diarrhea caused by taking antibiotics, improve allergies, and reduce the incidence of colorectal cancer [9]. B. lactis inhibited gastric Helicobacter pylori, improved diarrhea, and constipation [10]. Streptococcus thermophilus improved gastroenteritis, allergic rhinitis, inhibited gastric Helicobacter pylori [11]. In addition, prebiotic supplementation can increase the growth of the gut microbiota. Prebiotics cannot be decomposed by human enzymes, but they can be fermented or metabolized by intestinal bacteria, thus helping to grow the good gut microbiota and inhibiting the growth of the bad gut microbiota [12]. Supplementation with probiotics and prebiotics (fructooligosaccharides) can improve inflammation and intestinal permeability and reduce the burden of diseases [13].

Fruits and vegetables are important sources of dietary fiber with a balanced ratio of insoluble and soluble fractions. Adding fruit and vegetable fiber can expand the volume of feces and speed up defecation. The fermentation process of dietary fiber favors bacteria with carbohydrate degrading

Food Science & Nutrition Technology

ability, leading to the production of different metabolites, such as short-chain fatty acids (SCFAs) [14]. Increasing the levels of SCFAs in the intestine helps to lower its pH, creating a suitable environment for helpful bacteria, and hence, reducing the proportion of pathogenic bacteria. Papaya contains a special digestive enzyme called papain [15]. Studies have shown that this digestive enzyme can break down protein, which is easy for the body to absorb. Dioscorea opposita contains a high molecular mucin, which has a protective effect on digestive organs, can repair the gastric mucosa, buffer gastric acid, and avoid gastric and duodenal ulcers [16]. Pumpkin is rich in vitamins A and D, which can protect the gastrointestinal mucosa and prevent gastritis and gastric ulcers [17]. However, there were still few clinical studies on the combined use of probiotics, prebiotics, and fruit and vegetable fiber powder.

In this study, we used a complex formula containing probiotics, prebiotics, and fruit and vegetable fiber powder (PF formula) to explore whether the PF formula had an effect on gastrointestinal health care, immune regulation, and gut microbiota. 50 subjects were recruited and randomly assigned to two groups, namely the placebo group and the PF formula group. Subjects were examined for measurement of body position, blood analysis, stool analysis, and questionnaires at week 0 and week 12.

Materials and Methods

Clinical Trial Design

The clinical study had been approved by the Institutional Review Board of Chung Shan Medical University (CS2-21043) and the study had been registered on ClinicalTrials.gov ID: NCT04903600. In this trial, 50 adult subjects (20-60 years of age), who have had at least 4 colds in the past year or who have suffered from gastric Helicobacter pylori infection and gastric ulcer, were recruited in this trial. Informed consent was obtained from all subjects before the study. Subjects were divided into a placebo group (n=25) and a group of probiotics fruit vegetable fiber powder (PF formula) (n=25). Each subject was informed about taking placebo or PF twice a day for 12 weeks, and measurements of body position, blood analysis, stool analysis, and questionnaires were examined at week 0, week 4, week 8, and week 12. The exclusion criteria included: i) Subjects diagnosed with cancer and still on active treatment; ii) Subjects diagnosed with heart disease and still on active treatment; iii) Subjects with systemic infection requiring systemic antibiotics; iv) Who taking supplements containing probiotics, prebiotics or fruit and vegetable fiber; v) Subjects with dairy allergy or gastrointestinal distress.

Test Sample

The YIBENEF probiotics fruit & vegetable fiber solid powder (PF formula) (SHENZHEN YIBAIFEN INDUSTRY Co. LTD.) contained *Lactobacillus fermentum* TCI275, *Bifidobacterium longum* TCI068, *Streptococcus thermophilus* TCI25& TCI633, *Lactobacillus acidophilus* TCI800, *Lactobacillus paracasei* TCI058, *Lactobacillus rhamnosus* TCI366, *Bifidobacterium lactis* TCI604, fructooligosaccharides, papaya extract, dioscorea opposita extract, pumpkin extract, maltodextrin. The dose was 4.48×10^{11} CFU/day. The placebo powder contained skim milk powder and maltodextrin.

Clinical Measurements

Body weight, body mass index (BMI), body fat, and visceral fat were measured using the OMRON body fat calculator HBF-370. Tape measure was used to measure waist and hips. Systolic and diastolic blood pressures were measured with the NISSEI DS-G10J blood pressure machine. Blood was collected in serum separator tubes (BD Vacutainer, Plymouth, UK), and serum was removed by centrifugation at $3000 \times g$ for 15 minutes, aliquoted in microcentrifuge tubes and stored at -80°C in a freezer for analysis. Biochemical tests include aspartate transaminase (AST), alanine transaminase (ALT), creatinine, blood urea nitrogen (BUN), triglyceride, total cholesterol, low-density lipoprotein (LDL), high-density lipoprotein (HDL), sodium (Na⁺), potassium (K⁺), calcium (Ca²⁺), phosphorus (P⁻), chloride (Cl⁻), immunoglobulin A (IgA), immunoglobulin G (IgG), immunoglobulin M (IgM), interleukin-4 (IL-4), interleukin-8 (IL-8), interleukin-10 (IL-10), interleukin-12 (IL-12), interferon-y (IFN-y), tumor necrosis factor- α (TNF- α), high-sensitivity C-reactive protein (hsCRP) by an automatic modular biochemical immunoassay analyzer (C501, Roche, Basel, Switzerland).

Fecal Nucleic Acid Extraction

Nucleic acid was extracted from fecal bacteria with the QIA amp Fast DNA Stool Mini Kit. According to the standard operating procedure, the feces samples were removed from the preservation solution at 13200 rpm for 10 min, Inhibit EX buffer was added to dissolve the feces by shaking, and subsequent reagents were added in sequence. The concentration and OD values were determined by NanoDrop 2000.

Sequencing

According to the standard operation procedure of the original preparation of the Illumina 16S Metagenomic Sequencing Library preparation, amplify the V3 and V4 gene fragments of bacterial 16S rRNA. After the PCR products

Food Science & Nutrition Technology

are cleaned up, index PCR is performed, and the two ends of the DNA fragments are connected by two-stage PCR. The Illumina adapter sequence and index sequence, followed by the second PCR clean up, the two library calibrations were analyzed by capillary electrophoresis with Fragment Analyzer and quantified by Qubit 3.0 Fluorometer, and then DNA was denatured into single strands with sodium hydroxide and Illumina MiSeq was used for the second time. The generation is determined to perform bridge sub amplification, and the amplification is completed to pull into a single strand, add different bases, calibrate dNTPs, and reaction reagents that can remove fluorescent molecules, and repeat fluorescent label removal and detection to achieve rapid sequencing results. Sequencing was unified with the MiSeq Reagent Kit, V3 (600 cycles). Sequencing quality Q30≥80%, single sample≥100000 reads.

Statistical Analysis

The experimental data were analyzed using statistical product and service solutions (SPSS) version 22.0 computer statistical software. First, the paired t-test was used to analyze whether there was a significant difference between the same groups, and then the t-test was used to analyze the differences between the groups. Data from the above studies were expressed as mean and standard deviation (mean \pm SD), and p<0.05 was considered a significant difference.

Results

The PF formula Does Not Affect Liver Function, Kidney Function, or Blood Lipids

Figure 1 presents the recruitment flow chart for the subject. A total of 92 subjects were recruited, 42 subjects who did not meet the conditions were excluded, and the final total number of subjects was 50 subjects. Subjects were randomly assigned to the placebo (n=25) and PF formula groups (n=25). Table 1 shows the basic information on the subjects. The results showed that there were no differences between the placebo and PF formula group in terms of age, height, weight, BMI, waist circumference, hip circumference, waist-hip ratio, body fat, systolic blood pressure, and diastolic blood pressure. Table 2 showed the effect of serum biochemical parameters of the subjects. From the results, after taking the PF formula for 12 weeks, liver function (aspartate aminotransferase, alanine aminotransferase), kidney function (urea nitrogen, creatinine), uric acid, fasting blood glucose, blood lipids (cholesterol, triglycerides, highdensity lipoprotein cholesterol and low-density lipoprotein cholesterol) and electrolytes (sodium ion, potassium ion, calcium ion, and phosphorus ion) were not significantly different from the placebo group. However, taking Placebo for 12 weeks decreased albumin, sodium, and chloride

compared to week 0; taking PF formula for 12 weeks decreased albumin, chloride, and calcium compared to week 0. Although the results of the above serum biochemical

values were different, the serum biochemical values are still within the physiological range of the human body.



	Placebo (n=25)			PF (n=25)			
	Week 0	Week12	Δ	Week 0	Week12	Δ	
Subject (n)	25		-	25		-	
Male (n)	11		-	12		-	
Age (years)	37.64±12.86		-	37.76±12.33		-	
Height (cm)	165.44±9.06		-	165.84±10.04		-	
Weight (kg)	63.10±11.72	63.48±11.48	0.38±2.02	64.01±11.83	64.48±11.98	0.47±1.13	
BMI (kg/m ²)	22.77±2.66	23.02±2.74	0.26±0.55	23.12±2.58	23.32±2.52	0.20±0.48	
Waist (cm)	77.16±9.84	75.96±7.80	-1.20±5.32	79.32±10.70	79.20±8.23	-0.12±6.08	
Hip (cm)	95.26±6.20	94.28±5.18	-0.98±4.14	95.68±6.65	94.96±5.60	-0.72±4.36	
Waist-hip ratio	0.81±0.08	0.81±0.07	0.00±0.05	0.83±0.10	0.83±0.06	0.00±0.09	
SBP (mmHg)	126.32±21.34	119.08±13.85	7.24±13.17	122.52±19.02	114.84±14.25	-7.68±16.06	
DBP (mmHg)	78.08±14.99	74.80±10.07	-3.28±11.98	77.68±10.09	74.48±11.60	-3.20±7.66	
Body Fat (%)	25.47±0.06	26.59±5.96	1.12±1.30	27.37±0.04	27.54±4.32	0.17±1.51	
The reported values are the mean ± SD. All data between two groups do not have significant differences. PF means probiotics fruit vegetable fiber powder. BMI means body mass index. SBP means systolic blood pressure. DBP means diastolic blood							

pressure. Δ Mean change between week 0 and week 12 (week 12-week 0).

Table 1: Differences between week 0 and week 12 of the subject's characteristics after the Placebo and PF intervention.

	Placebo (n=25)		PF (n=25)		Normal
	Week 0	Week 12	Week 0	Week 12	range1
Glucose(mg/dL)	89.08±13.46	86.72±8.48	87.32±12.85	92.76±27.05	60~100
BUN (mg/dL)	11.92±2.89	13.12±3.68	13.12±3.80	14.16±3.91	5~25
Creatinine(mg/dL)	0.80±0.18	0.77±0.20	0.79±0.17	0.80±0.19	0.1~1.4
Uric Acid (mg/dL)	5.12±1.12	5.06±1.07	5.52±1.34	5.18±1.13	2.0~7.0
AST (U/L)	21.68±9.20	25.56±22.00	20.76±7.93	20.36±5.34	0~40
ALT (U/L)	19.20±14.51	21.04±13.93	20.56±20.09	19.44±8.48	0~40
Albumin (g/dL)	4.84±0.26	4.56±0.25**	5.00±0.20#	4.72±0.20**	3.5~5.5
Triglyceride(mg/dL)	80.00±40.69	80.08±46.46	88.36±43.35	94.36±62.99	30~150
Cholesterol (mg/dL)	182.36±32.51	184.28±36.53	174.00±26.28	183.64±32.42	130~200
HDL-C (mg/dL)	57.28±12.53	59.72±15.90	58.16±13.84	58.80±15.0	>40
LDL-C (mg/dL)	117.36±31.09	109.80±30.27	107.60±30.00	107.88±33.12	<130
LDL-C /HDL-C ratio	2.14±0.65	1.95±0.64	2.01±0.88	2.01±0.91	<3.5
Na⁺ (mmol/L)	140.12±1.76	138.32±1.35**	139.84±2.13	138.88±2.05	130~145
K⁺ (mmol/L)	4.67±0.66	4.50±0.44	4.72±0.48	4.49±0.36	3.5~5.0
Cl ⁻ (mmol/L)	103.49±1.27	100.29±1.41**	101.78±1.51#	100.17±2.18**	90~110
Ca2+ (mmol/L)	9.19±0.35	9.06±0.34	9.35±0.25	9.20±0.28**	8.5~11.5
P⁻ (mmol/L)	3.80±0.43	3.83±0.35	3.83±0.45	3.89±0.60	2.6~5.0

The reported values are the mean ± SD. #Mean significant between two groups at the same time point, 0.01≦p<0.05 usinig the student's t-test. **Mean significant between week 0 and week 12 within the same group, p<0.01 using the paired t-test. 1 means provided from the medical laboratory. PF means probiotics fruit vegetable fiber powder. AST means aspartate aminotransferase. ALT means alanine aminotransferase. BUN means blood urea nitrogen. HDL-C means high density lipoprotein-cholesterol. LDL-C means low density lipoprotein-cholesterol.

Table 2: Differences between week 0 and week 12 of serum biochemical parameters after the Placebo and PF intervention.

The PF Formula Does Not Alter The Dietary Intake and has an Anti-Inflammatory Effect

Table 3 shows that there were no significant differences between the Placebo and PF groups in terms of energy, major nutrients (protein, fat, carbohydrate), dietary fiber, sodium, and water intake. Figure 2 shows the values of the immunerelated biochemical parameters of the subjects. After taking the PF formula for 12 weeks, IL-4, IFN- γ , and TNF- α were significantly decreased compared to the placebo group. Furthermore, after taking the PF formula for 12 weeks, IgG, IgM, IL-10 increased significantly compared to baseline (week 0), and IL-4, IL-6, IL-12, IFN- γ and TNF- α significantly decreased compared to baseline (week 0).

	Placebo	(n=25)	PF (n=25)				
	Week 0	Week12	Week 0	Week12			
Energy (Kcal/day)	1487.41±205.57	1433.32±274.68	1438.55±236.88	1405.87±269.92			
	Protein						
(g/day)	56.26±20.86	56.30±15.84	59.23±16.04	60.55±16.51			
(% of total energy)	16.58±0.04	17.30±0.06	16.38±0.03	17.14±0.03			
Fat							
(g/day)	58.97±18.03	52.05±14.67	55.81±16.26	58.04±15.68			
(% of total energy)	35.33±0.09	34.68±0.12	35.05±0.08	36.82±0.07			
Carbohydrate							
(g/day)	184.21±48.00	178.02±78.30	174.40±40.81	163.05±42.27			
(% of total energy)	49.70±0.13	48.01±0.16	48.58±0.09	46.04±0.09			

Dietary fiber (g/ day)	10.61±3.78	9.64±4.22	11.44±5.12	10.62±4.00	
Sodium (mg/day)	1857.75±668.86	1966.69±331.37	2053.81±544.39	2012.60±674.89	
Water (mL/day)	2146.21±915.74	2331.57±700.71	2022.21±783.63	2130.08±738.08	
The reported values are the mean ± SD. Analyzed with Ekitchen plus3. All data between two groups do not have significant					
differences. PF means probiotics fruit vegetable fiber powder.					

Table 3: Nutrition intakes of subjects.



Figure 2: Differences between week 0 and week 12 of the subject's serum immune after Placebo and PF intervention. (A) Immunoglobulin A (IgA; mg/dL); (B) Immunoglobulin G (IgG; mg/dL); (C) Immunoglobulin M (IgM; mg/dL); (D) Interleukin-4 (IL-4; pg/mL); (E) Interleukin-6 (IL-6; pg/mL); (F) Interleukin-10 (IL-10; pg/mL); (G) Interleukin-12 (IL-12; pg/mL); (H) Interferon gamma (IFN- γ ; pg/mL); (I) Tumor Necrosis Factor- α (TNF- α ; pg/mL), and (J) High-sensitivity CRP (hsCRP; mg/dL). The reported values are the mean ± SD. n=25. *Mean significant between two groups at the same time point, p<0.01 using the student's *t*-test. **Mean significant between two groups at the same time point, p<0.01 using the student's *t*-test. **Mean significant between week 0 and week 12 within in same group, p<0.01 using the paired *t*-test. PF means probiotics fruit vegetable fiber powder.

The PF Formula Improved The Gastrointestinal Tract and Immune Function

Table 4 shows the results of the somatosensory questionnaire. The content of the questionnaire was

mainly for the gastrointestinal tract and immune function. After taking the PF formula for 12 weeks, symptoms of the gastrointestinal tract (defecation frequency, frequency of bowel movements, rectal tenesmus, defecation time, abdominal pain, gastralgia, flatulence, belching) and

gastroesophageal reflux (nausea, appetite loss, vomit, constipation, hard stool, and diarrhea) and cold symptoms (fever, headache, cough, muscle pain, dizziness, acne) were not significantly different compared to the placebo group. In particular, after 12 weeks of taking the PF formula, there was an increase in the frequency of bowel movements and

a decrease in gastralgia, flatulence, constipation, hard stool, diarrhea, headache, muscle pain, dizziness compared to baseline (week 0). From the above results, it was suggested that the administration of the PF formula can improve the gastrointestinal tract and immune function.

Sumptoma	Placebo	PF (n=	PF (n=25)		
Symptoms	Week 0	Week12	Week 0	Week12	
	Gast	rointestinal tract symp	toms	•	
Defecation frequency	1.44±0.82	1.20±0.76	1.32±1.07		1.12±0.97
Bowel movement frequency	1.56±1.16	1.80±0.91	1.52±0.87		2.04±0.89*
Rectal tenesmus	1.56±1.00	1.48±0.87	1.72±1.17		1.40±0.65
Defecate time	0.92±0.76	0.72±0.54	0.88±1.01		0.64±0.28
Abdominal pain	0.44±0.71	0.20±0.50	0.24±0.44		0.32±0.75
Gastralgia	0.24±0.60	0.16±0.47	0.44±0.65		0.12±0.33*
Flatulence	0.88±1.20	0.32±0.63*	1.28±1.14		0.48±0.82**
Belching	0.60±1.00	0.32±0.63	0.36±0.64		0.20±0.50
Gastroesophageal reflux	0.44±0.65	0.28±0.54	0.52±1.08		0.24±0.44
Nausea	0.36±0.70	0.08±0.28*	0.32±0.69		0.16±0.47
Appetite loss	0.32±0.63	0.04±0.20*	0.36±0.76		0.16 ± 0.47
Vomit	0.16±0.47	0.00±0.00	0.08±0.28		0.04±0.20
Constipation	0.52±0.77	0.28±0.54	0.68±1.11		0.16±0.37*
Hard stools	0.40±0.91	0.24±0.52	1.16±1.31		0.20±0.50**
Diarrhea	0.48±0.65	0.08±0.28	0.56±1.00		0.12±0.33*
		Other symptoms			
Fever	0.04±0.20	0.04±0.20	0.04±0.20		0.04 ± 0.20
Headache	0.76±1.13	0.16±0.37**	0.44±0.65		0.24±0.44**
Cough	0.12±0.33	0.12±0.33	0.28±0.46		0.16±0.37
Muscle pain	0.64±1.08	0.16±0.37*	0.92±1.35		0.52±1.00**
Dizziness	0.52±0.71	0.24±0.44	0.80±0.91		0.24±0.44**
Acne	0.52±0.92	0.16±0.37	0.84±1.18		0.32±0.90

ous). PF means probiotics fruit vegetable fiber powder. *Mean significant between week 0 and week 12 within the same group,0.01≦p<0.05 using the paired t-test. **Mean significant between week 0 and week 12 within the same group, p<0.01 using the paired t-test.

Table 4: Differences between week 0 and week 12 of subject's gastrointestinal and immune function after the Placebo and PF intervention.

The PF Formula Increased The Abundance of Good Bacteria *Porphyromonadaceae* and *Parabacteroides* and Decreased The Abundance of Bad Bacteria *Clostridiaceae*.

To analyze whether the PF formula affected the gut microbiota, the feces of the subject were collected for analysis.

Figure 3 represents the principal coordinate analysis (PCoA), and each point represents one subject. The distance between two points indicates the similarity of the clusters, and the smaller the distance, the more similar the clusters are. The results showed that the PF formula did not alter the fecal microbiota compared to placebo at week 0 and week 12.







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Figure 4A shows the composition of the fecal microbiota phylum. The most abundant microbiota in two groups (placebo and PF formula) were *Firmicutes* and *Bacteroridetes*. Figure 4B shows the composition of the class of fecal microbiota. The most abundant microbiota were *Bacterodia* and *Clostridia*. Figure 4C shows the composition of the fecal microbiota in order. The most abundant microbiota were *Bacteroidales* and *Clostridiales*. Figure 4D shows the composition of the fecal microbiota family. The most abundant microbiota was *Bacteroidaceae*. Figure 4E shows the composition of the fecal microbiota genus. The most abundant microbiota was *Bacteroides*. Figure 5 shows the compositional differences of phylum, order, family, and genus after taking the PF formula for 12 weeks. The PF formula slightly increased the relative abundance of *Bacteroidetes* (phylum), slightly decreased *Actinobacteria*, and *Cyanobacteria* (phylum) compared to the placebo group. Furthermore, the PF formula slightly decreased *Sphingobacteriales* (order), *Clostridiaceae* (family), and significantly increased *Porphyromonadaceae* (family), *Parabacteroides* (genus) compared to the placebo group.



Figure 5: Effect of Placebo and PF on fecal microbiota phylum, order, family, and genus level composition in week 12. (A) *Bacteroidetes* (phylum), (B) *Actinobacteria* (phylum), (C) *Cyanobacteria* (phylum), (D) *Sphingobacteriales* (order), (E) *Clostridiaceae* (family), (F) *Porphyromonadaceae* (family), and (G) *Parabacteroides* (genus). The reported values are the mean ± SD (n=25).

[#]Mean significant between two groups at the same time point, p<0.05 usinig the student's *t*-test. PF means probiotics fruit vegetable fiber powder.

Discussion

The intestines are the largest immune organ in the human body, with the intestinal mucosa as the first line of defense and immune cells as the second line of defense [18]. The intestinal mucosa as the first line of defense can block undigested protein molecules, toxins, bacteria, molds, etc. from entering the body and excreted through feces; the immune cells of the second line of defense can fight against bacterial invasion [19]. A decrease in gastrointestinal tract health can lead to an overly sensitive inflammatory response from the immune system. In order to reduce the frequency of gastrointestinal discomfort and improve immunity, many healthy foods have been developed on the market recently, such as fruits and vegetables, probiotics, and prebiotics. In this study, we found that the PF formula (combine probiotics, prebiotics with fruit and vegetable fiber) has a beneficial effect on gastrointestinal health and immune enhancement and can also regulate the intestinal microbiota. The PF formula does not affect basic body position, blood pressure, and dietary

intake. However, the PF formula group had a higher albumin and a lower chloride ion compared to the placebo group in week 0. However, it is still within the normal physiological range of the human body. Human albumin range is 3.5~5.5 g/dL, and chloride ions range 90-110 mmol/L. Therefore, from the above results, the PF formula does not have any negative effects on safety indicators (liver, kidney indicators, and electrolyte concentration), and the PF formula is a safe health food.

Studies have shown that lactobacilli and bifidobacteria can increase IL-10, which in turn improves type II helper T cell production and inhibits pro-inflammatory cytokines TNF- α , IFN- γ and IL-6 [20,21]. Supplementation of 2 types of lactobacilli can increase IL-10 and IL-4, thus promoting T cell production [22]. Studies have shown that supplementation with 2×10¹⁰ CFU/day of Bifidobacterium lactis or Lactobacillus acidophilus can significantly increase IgG concentration [23]; Supplementation with 1×10¹⁰ CFU/day Lactobacillus fermentum can enhance IgM concentration and reduce the chance of influenza [24]. Consistent with our results, the PF formula for 12 weeks can increase IgG and IgM compared to baseline (week 0). Studies have shown that supplementation with 1×10^9 CFU/day of Lactobacillus bizakii for 18 weeks can increase IL-10 and reduce TNF- α [25]. Supplementation with Lactobacillus plantarum 1×109 CFU/day for 12 weeks can reduce IFN- γ and TNF- α and increase IL-4 and IL-10 [26]. Supplementation with 1×10⁹ CFU/day of Bacillus coagulans reduced IL-6 [27]. Consistent with our results, the PF formula for 12 weeks increased IL-10 and decreased IL-4, IL-6, IL-12, IFN- γ , and TNF- α compared to baseline (week 0). Since no IgE was measured in the study, it is speculated that the PF formula may have an allergy modifying effect [28], suggesting that the PF formula has an immunomodulatory effect. Supplementation with Bifidobacterium lactis can increase the frequency of bowel movements, reduce flatulence, and constipation [29]. Taking Lactobacillus plantarum can reduce flatulence, abdominal pain, diarrhea, and irritable bowel syndrome (IBS), so it can be known that probiotics intervention is beneficial for gastrointestinal health. Taking Lactobacillus and Bizobacteria can improve antibioticassociated diarrhea [30]. In addition, taking 3×10¹⁰ CFU/day of Lactobacillus plantarum can improve muscle pain after exercise [31]. Consistent with our results, the PF formula for 12 weeks can significantly increase the frequency of bowel movement, reduce stomach pain, flatulence, constipation, hard stool, diarrhea, and reduce the incidence of headache, muscle pain, and dizziness.

Probiotics improve the balance of the bacterial phase balance of intestinal microorganisms and promote host health, while prebiotics are probiotic growth promoting substances, fruit and vegetable fiber increase the survival of beneficial flora and regulate the diversity of the intestinal microbiota [32]. However, taking the PF formula for 12 weeks, PCoA was used to analyze the diversity of the gut microbiota, but no bacterial phase differences were observed, only the distribution of the gut microbiota was changed. At the phylum level of the microbiota, the Bacteroidetes were positively correlated with serum IgA, IgM, and IgG [33]. Consistent with our results, the relative abundance of Bacteroidetes was higher than placebo after 12 weeks of administration of the PF formula. Studies have shown that the proportion of Actinobacteria and Proteobacteria in the stool of patients with ulcerative colitis is higher than that of healthy people [34]. Cyanobacteria increase the production of proinflammatory cytokines [35]. Consistent with our results, the relative abundance of Actinobacteria, Cyanobacteria, was lower than placebo after 12 weeks of administration of the PF formula. At the order level of the microbiota, Lactobacillus rhamnosus increases the concentrations of IL-4 and IL-10 in the blood and reduces the abundance of Sphingobacteriales [36]. Consistent with our results, the relative abundance of Sphingobacteriales was lower than placebo after 12 weeks of administration of the PF formula. At the family level of the microbiota, Clostridiaceae increase when the body is in an inflammatory state [37]. Porphyromonadaceae regulate short-chain fatty acid production and improve gastrointestinal function [38]. Consistent with our results, the relative abundance of Clostridiaceae was lower and Porphyromonadaceae was higher than placebo after 12 weeks of administration of the PF formula. At the genus level of the microbiota, *Parabacteroides* inhibit TNF- α and pro-inflammatory cytokines (IL-12, IFN-y), with potential for immunomodulation [39]. Consistent with our results, the relative abundance of Parabacteroides was higher than placebo after 12 weeks of PF formula administration.

Fruits and vegetables, dietary fibers, are an essential component of a healthy diet pattern, which is suggested to play an important role in maintaining the balance of the gut microbiota and improving intestinal ecology [40]. High intake of fruits and vegetables can increase good gut bacteria, such as Faecalibacterium prausnitzii, Akkermansia muciniphila, Ruminococcaceae, Clostridiales, and Acidaminococcus, which indicated that high intake of dietary fiber was potentially beneficial to human health by increasing short-chain fatty acid production, maintaining intestinal mucosal integrity, improving insulin sensitivity, and anti-inflammatory properties [41]. Some studies showed that papaya, opposite dioscorea, pumpkin, which are fermented dietary fibers, are high in butyrate, which is associated with maintaining intestinal cell integrity and reducing intestinal inflammation, and increased Bacteroides or decreased Roseburia results in an increase in butyrate [42]. In conclusion, the PF formula can regulate the immune inflammatory response and improve

gastrointestinal function after 12 weeks of administration.

Conflict of Interest

The authors declare that they have no conflict of interest.

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