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Association between probiotic consumption and serum perfluoroalkyl and polyfluoroalkyl substances (PFAS): results from NHANES, 2003–2018

Yangjie Liao¹, Jiang Chen², Jingbo Li³, Jiayi Wang³, Long Cheng^{1*} and Min Guo^{1*}

Abstract

Background Perfluoroalkyl and polyfluoroalkyl substances (PFAS) represent a category of pervasive and enduring environmental pollutants that present a risk to human health. Although growing evidence suggests that probiotics can potentially alleviate the adverse effects of PFAS, large cross-sectional studies on the relationship between probiotic consumption and PFAS remain lacking.

Objective The objective of this study is to assess the association between the exposure of probiotics and serum levels of PFAS.

Methods This analysis included individuals aged 20 and above who took part in the National Health and Nutrition Examination Survey (NHANES) between 2003 and 2018. Probiotic consumption was considered when a participant reported consuming yogurt during the two 24-h dietary recall or using a probiotic supplement in dietary supplement questionnaires over the past 30 days.

Results This study involved 9469 adults, out of which 1333 had been exposed to probiotics. We found negative associations between probiotic consumption and serum concentrations of perfluorooctanoic acid (PFOA) (β : -0.19 , 95% CI -0.35 to -0.02 ; $P=0.027$), and perfluorooctane sulfonic acid (PFOS) (β : -0.127 , 95% CI -2.23 to -0.32 ; $P=0.010$). The consumption of probiotic supplements alone was associated with reduced perfluorononanoic acid (PFNA) (β : -0.19 , 95% CI -0.28 to -0.10 ; $P<0.001$). No statistically significant association was identified between probiotic consumption and perfluorohexane sulphonic acid (PFHxS).

Conclusion In this cross-sectional, nationally representative study, probiotic ingestion was negatively associated with several serum PFAS compounds. These findings carry substantial implications for designing interventions that target the reduction of accumulated PFAS levels in the body and mitigating the resulting adverse health effects.

Keywords PFAS, Probiotics, Yogurt, NHANES, Nutrients, Epidemiology

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Introduction

Perfluoroalkyl and polyfluoroalkyl substances (PFAS) belong to a group of fluorinated aliphatic chemicals that offer great chemical and thermal stability, along with hydrophobic and oleophobic properties [1]. Thus, they have found wide-ranging applications in diverse industries and everyday consumer products, including industrial detergents, non-stick cookware, food packaging, and textiles [2, 3]. However, the remarkable stability of PFAS poses a challenge as they exhibit resistance to natural degradation. In addition, PFAS can accumulate in water, soil, plants, and organisms throughout the production and usage processes [4]. The persistence and ubiquity of PFAS result in their detection in populations worldwide [5, 6]. PFAS have raised high public health concerns due to their detrimental health effects, which encompass but are not limited to: liver and kidney injury, skeletal disorders, reproductive disorders, immunosuppression, hormonal imbalance, lipid dysregulation, and cancer [7–9].

Recently, the application standards for PFAS have become increasingly stringent in numerous countries [10, 11]. Perfluorooctane sulfonic acid (PFOS) and perfluorooctanoic acid (PFOA) serum levels in the US population witnessed a decline of over 85% and 70% respectively between 1999 and 2018 based on the National Health and Nutrition Examination Survey (NHANES) [12]. Nevertheless, other PFAS may be released as PFOS and PFOA are phased out and replaced. The long-lasting nature of PFAS implies that the bioaccumulation in the human body may still endanger personal well-being. Therefore, the development of effective measures to alleviate individuals' burden to PFAS is of utmost importance. Previous research has indicated that cholestyramine, a bile acid sequestrant, may reduce PFAS reabsorption in bile, leading to an elevated excretion of PFAS through feces [13, 14]. Lifestyle modifications aimed at lowering lipid levels could also potentially influence circulating levels of PFAS [15]. Additionally, numerous studies have suggested a negative association between PFAS and the ingestion of fiber-rich foods, including vegetables, fruits, legumes, and whole grains [16–18]. Interestingly, dietary fiber and folate, which are essential components of these foods, have been found to appear to decrease the concentration of PFAS in the bloodstream [19–21]. The intestinal microbiota has the ability to ferment dietary fiber, resulting in the production of short-chain fatty acids (SCFAs) that exert protective effects on the gut [22]. Consequently, it is worth considering whether the ingestion of probiotics could act as an important confounding factor in the relationship between dietary fiber intake and PFAS levels.

Probiotics, which are live microorganisms with a beneficial impact on human health [23], can be obtained

from various sources, with yogurt being the most common dietary source [24]. The primary probiotic bacteria are lactic acid bacteria and bifidobacterial [25]. Previous studies have demonstrated a rise in the usage of probiotics among adults and children in the US from 1999 to 2018, which is opposite to the decreasing trend of certain PFAS levels [26]. Probiotics may potentially offer defense against the toxicity of PFAS [27, 28]. However, the precise mechanism by which probiotics mitigate the toxicity of PFAS remains unclear, and there is currently a lack of direct evidence supporting the effect of probiotics on serum PFAS levels. Furthermore, there is a lack of epidemiological studies to evaluate PFAS levels in populations consuming probiotics. This study seeks to examine the association between probiotic consumption and serum PFAS concentrations using NHANES data, providing new insights into the potential role of probiotics in mitigating PFAS toxicity.

Methods

Study design and participants

The public data for this cross-sectional study was obtained from the NHANES 2003–2018 cycle. NHANES is a nationwide survey conducted every 2 years in the US to evaluate the nutritional and health conditions of the general population using questionnaires, physical exams, and biospecimen collection. The study included participants aged 20 and above who had completed probiotics consumption data, PFAS levels measuring, and data for all other covariates. Pregnant or breastfeeding women were excluded from the study because PFAS can be transferred to the fetus through cord blood during pregnancy, or to the baby through breast milk during lactation [29–33]. Participants undergoing dialysis were also excluded due to the potential excretion of PFAS through dialysis [34]. The study population flowchart is depicted in Additional file 1: Fig. S1. All individuals involved in the study provided written informed consent and all protocols were approved by the National Center for Health Statistics's Research Ethics Review Board.

PFAS measurement

Briefly, the online-solid phase extraction combined with high-performance liquid chromatography-Turbo Ion Spray ionization-tandem mass spectrometry was employed to quantify PFAS. This study specifically examined four PFAS substances: PFOA, PFOS, perfluorononanoic acid (PFNA), and perfluorohexane sulfonic acid (PFHxS), which had detection frequencies above 98% in the serum samples. The concentrations below the limit of detection (LOD) were substituted with $LOD/\sqrt{2}$. The LODs for each chemical were as follows: PFOA LOD: 0.1 ng/mL (2003–2018). PFOS LOD: 0.4 ng/mL

(2003–2004), 0.2 ng/mL (2005–2012), and 0.1 ng/mL (2013–2018). PFNA LOD: 0.82 ng/mL (2007–2010), 0.8 ng/mL (2011–2012), and 0.1 ng/mL (2003–2006 and 2013–2018). PFHxS LOD: 0.3 ng/mL (2003–2004) and 0.1 ng/mL (2005–2018). In the 2013–2018 data cycles, separate measurements were conducted for both the linear and branched isomers of PFOA and PFOS. To maintain consistency with previous cycles, the total concentration of PFOA and PFOS was calculated by summing the measured values of their respective linear and branched isomers.

Identifying probiotics

This study assessed probiotic consumption based on participants' consumption of yogurt or probiotic-containing dietary supplements [35, 36]. Yogurt ingestion was evaluated using 24-h dietary recall on the first and second days, with USDA Food Code "114" indicating yogurt. Dietary supplements containing probiotics were assessed using the Dietary Supplement Use 30-day questionnaire, which contained data on dietary supplement and non-prescription antacid. Text mining methodologies were applied to identify probiotics by analyzing data from the dietary supplement products, ingredients, and blends information. The search terms used for identifying probiotics refer to previous research [26], as detailed in Additional file 1: Table S1.

Covariates

Covariates selected a priori for analysis consisted of age, gender, race, body mass index (BMI), highest educational level, marital, poverty income ratio (PIR), smoking status, alcohol drinking, NHANES wave, and the intake of fish, meat, egg, energy, dietary folate equivalent, and dietary fiber. Demographic variables were collected through questionnaires, encompassing age (continuous, in years), gender (categorized as male or female), race (classified as Mexican American, non-Hispanic White, non-Hispanic Black, or other), BMI (continuous, in kg/m²), highest educational level (categorized as ≤high school or >high school), marital (categorized as married or other), and PIR (categorized as <1.30, 1.30 to <3.50, or ≥3.50). Respondents were grouped into current, former, or never smokers in accordance with whether they had smoked 100 cigarettes in their lives and whether they now smoked.

To take into potential confounding factors account, food items that have been identified as having high levels of PFAS or being associated with higher PFAS levels, such as fish, meat, and egg, were included in the analysis [37–40]. Food and nutrient calculations were based on data from two 24-h recalls, using the USDA Food Patterns Equivalents Database to determine

food consumption. The fish variable (continuous, in ounce/d) encompassed fish, shellfish, and other seafood. The meat variable (continuous, in ounce/d) encompassed beef, pork, veal, lamb, and game, organ meats, frankfurters, sausages, luncheon meat, and poultry (chicken, turkey, and other). The egg variable (continuous, in ounce/d) encompassed eggs and egg substitutes. Nutrient intake covariates included dietary fiber (continuous, in g/d), energy (continuous, in kcal/d), dietary folate equivalent (continuous, µg/d), and alcohol consumption. Dietary fiber and folate have been found to be negatively associated with PFAS levels in adult serum [19, 20]. Adjusting for total energy intake allows for a relative rather than an absolute representation of dietary intake, considering the variation in energy intake among individuals engaged in different physical activities or with varying body weights [41]. Subjects were classified as heavy alcohol consumption if their daily ethanol consumption exceeded 10 g for females or 20 g for males; otherwise, they were classified as no heavy alcohol consumption.

Statistical analysis

Baseline characteristics were presented using median and quartiles for continuous variables, while percentages were utilized for categorical variables. To investigate the relationship between probiotic consumption and various PFAS compounds (PFOA, PFNA, PFOS, and PFHxS), a generalized linear regression model was applied. Model 1 was unadjusted, Model 2 adjusted for age, gender, and BMI, and Model 3 adjusted for all covariates. To investigate whether the associations differed depending on the source of probiotics (probiotics supplement or yogurt), we conducted further analysis. Subgroup analyses were performed for different factors, including age (<54 y, ≥54 y), gender, race, BMI (<20, 20–<25, 25–<30, 30+), educational level, marital status, PIR (<1.30, 1.30–<3.50, 3.50+), smoking status, alcohol drinking, NHANES wave (2003–2005, 2007–2009, 2011–2013, 2015–2017), and the intake of fish (Yes or No), meat (<2.19, 2.19–<3.71, 3.71–<5.74, 5.74+), egg (<0.035, 0.035–<0.230, 0.230–<0.885, 0.885+), energy (<1545, 1545–<1972.5, 1972.5–<2559, 2559+), dietary folate equivalent (<333, 333–<466, 466–<657, 657+), and dietary fiber (<10.75, 10.75–<15.25, 15.25–<21.35, 21.35+). Likelihood ratio tests were conducted to assess the presence of interaction effects by comparing models with and without interaction terms. The statistical analysis was conducted utilizing R version 4.3.1 and the "survey" package. Statistical significance was determined based on a significance level of $P < 0.05$.

Results

The baseline population characteristics regarding probiotic consumption are outlined in Table 1. This study included 9469 adults aged 20 years or above, of whom 1333 had been exposed to probiotics. The median age of the probiotic exposure group, 50 years, was higher than that of the non-exposure group, which had a median age of 46 years. In the probiotic exposure group, there was a higher proportion of female participants (62.28%) compared to the non-exposure group (49.28%). The individuals in the non-exposure group had a median BMI of 28.10 kg/m², whereas those in the probiotic consumption group had a median BMI of 27.49 kg/m². The probiotic exposure group had higher proportions of non-Hispanic White individuals (79.71%) and married participants (61.41%). Probiotic consumption also had higher socioeconomic status, as indicated by both PIR and education. For instance, 76.52% of individuals with an education level beyond high school reported probiotic consumption, compared to 58.66% of those who did not. In the non-probiotic exposure group, the daily intake of meat, dietary fiber, and dietary folate equivalents was 3.77 oz, 14.70 g and 462.00 mcg, respectively, while in the probiotic exposure group, it was 3.30 oz, 17.95 g and 480.00 mcg, respectively. In the group exposed to probiotics, participants demonstrated a higher preference for dietary fiber and dietary folate equivalents rather than meat. The ingestion of energy, egg, and the proportion of heavy alcohol consumption were similar between the two groups. Individuals exposed to probiotics had higher percentages of never-smoker statuses (62.68%) and lower proportions of current smokers (9.70%) compared to those not exposed to probiotics. Among the PFAS compounds studied, the serum levels of PFOA, PFNA, and PFOS showed significant differences between the two groups, with lower median concentrations by 0.30 ng/mL, 0.09 ng/mL, and 2.30 ng/mL respectively, compared to the non-exposure group. However, the concentrations of PFHxS did not differ significantly between the two groups.

The study utilized baseline and multivariate design-based survey linear regression models to analyze the association between probiotic consumption and individual PFAS, as shown in Table 2. In the unadjusted analyses (Model 1), it was observed that probiotic consumption exhibited a statistically significant association with the reductions in PFOA levels, PFNA levels, and PFOS levels. After controlling for age, gender, and BMI (Model 2), the statistical significance of the association between probiotic consumption and decreased levels of PFOA (β : -0.43, 95% CI -0.63 to -0.23; $P < 0.001$), PFNA (β : -0.13, 95% CI -0.21 to -0.05; $P = 0.002$), and PFOS (β : -3.21, 95% CI -4.23 to -2.19; $P < 0.001$) was maintained.

After full adjustment (Model 3), although the association between probiotic consumption and PFNA disappeared, probiotic consumption remained negatively associated with PFOA and PFOS. Notably, no significant association was found between probiotic consumption and PFHxS across all models.

In Model 3, further analysis was undertaken to assess the association between probiotic supplements, yogurt consumption, and serum PFAS concentrations (Fig. 1). Among the participants who were exposed to probiotics, a total of 1194 individuals were exposed to yogurt, 139 individuals were exposed to probiotic supplements, and 50 participants were exposed to both simultaneously. Any type of probiotics and probiotic supplements were associated with reduced PFOA concentration. Serum PFNA concentration was not associated with exposure to any type of probiotics or yogurt but showed a significant association with exposure to probiotic supplements alone (β : -0.19, 95% CI -0.28 to -0.10; $P < 0.001$). Irrespective of the type of probiotics, a negative association was observed with serum PFOS levels.

The stratified analysis based on population characteristics (Table 3 and Additional file 1: Table S2) revealed significant interactions between probiotic consumption and serum PFOA levels by race (P -interaction = 0.009), education (P -interaction = 0.012), PIR (P -interaction = 0.021), and meat intake (P -interaction = 0.046). Additionally, significant interactions were observed between probiotic consumption and PFNA by race (P -interaction = 0.038), as well as between probiotic consumption and PFOS by PIR (P -interaction = 0.022) and heavy alcohol consumption (P -interaction = 0.036). Other associations did not yield statistically significant differences. Notably, the negative association between probiotic consumption and PFOA levels was more apparent among non-Hispanic White individuals, those with an education level beyond high school, high household income, and the highest quartile of meat intake. A statistically significant association between probiotic consumption and decreased PFNA concentration was also observed among non-Hispanic White individuals. In terms of PFOS, participants with moderate family incomes or heavy alcohol consumption were more susceptible to the association.

Discussion

This representative large-scale cross-sectional study suggested a strong negative association between the use of probiotics and serum concentrations of PFOA and PFOS. While there were no associations found between exposure to any type of probiotics and serum PFNA concentration, exposure to probiotic supplements alone was significantly associated with reduced PFNA levels. The negative associations between probiotic consumption

Table 1 Population characteristics at baseline categorized by probiotic consumption (n = 9469)

Characteristic ^a	Overall (n = 9469)	No exposure to probiotics (n = 8136)	Exposure to probiotics (n = 1333)	P value ^b
Age (years)	47.00 (34.00, 60.00)	46.00 (33.00, 60.00)	50.00 (37.00, 62.00)	< 0.001
Gender				< 0.001
Male	4614 (48.50)	4136 (50.72)	478 (37.72)	
Female	4855 (51.50)	4000 (49.28)	855 (62.28)	
BMI (kg/m ²)	27.95 (24.26, 32.50)	28.10 (24.40, 32.70)	27.49 (23.89, 31.60)	0.001
Race				< 0.001
Mexican American	1425 (7.50)	1254 (7.95)	171 (5.30)	
Non-Hispanic Black	1967 (10.24)	1804 (11.31)	163 (5.02)	
Non-Hispanic White	4445 (70.89)	3700 (69.08)	745 (79.71)	
Other race	1632 (11.37)	1378 (11.66)	254 (9.97)	
Marital				0.007
Married	5004 (56.83)	4243 (55.89)	761 (61.41)	
Other	4465 (43.17)	3893 (44.11)	572 (38.59)	
Education				< 0.001
≤ High school	4421 (38.29)	4007 (41.34)	414 (23.48)	
> High school	5048 (61.71)	4129 (58.66)	919 (76.52)	
PIR				< 0.001
< 1.30	2796 (19.54)	2527 (21.23)	269 (11.33)	
1.30 to < 3.50	3619 (35.93)	3175 (37.16)	444 (29.97)	
≥ 3.50	3054 (44.53)	2434 (41.61)	620 (58.70)	
Fish (ounce/d)	0.00 (0.00, 0.43)	0.00 (0.00, 0.37)	0.00 (0.00, 0.73)	0.020
Meat (ounce/d)	3.71 (2.19, 5.74)	3.77 (2.27, 5.87)	3.30 (1.92, 5.19)	< 0.001
Egg (ounce/d)	0.23 (0.04, 0.89)	0.23 (0.04, 0.89)	0.22 (0.04, 0.88)	0.900
Energy (kcal/d)	1972.44 (1544.63, 2559.00)	1975.00 (1533.71, 2571.50)	1960.00 (1603.00, 2470.80)	0.800
Dietary fiber (g/d)	15.25 (10.75, 21.35)	14.70 (10.45, 20.70)	17.95 (13.00, 23.60)	< 0.001
Dietary folate equivalent (μg/d)	466.00 (333.00, 657.00)	462.00 (329.50, 652.50)	480.00 (346.00, 679.02)	0.047
Smoking status				< 0.001
Never	5180 (55.10)	4333 (53.56)	847 (62.58)	
Former	2400 (24.83)	2040 (24.24)	360 (27.72)	
Current	1889 (20.06)	1763 (22.20)	126 (9.70)	
Heavy alcohol consumption				0.100
No	7903 (79.68)	6791 (80.19)	1112 (77.20)	
Yes	1566 (20.32)	1345 (19.81)	221 (22.80)	
PFOA (ng/mL)	2.70 (1.67, 4.40)	2.77 (1.67, 4.60)	2.47 (1.50, 3.78)	< 0.001
PFNA (ng/mL)	0.90 (0.55, 1.39)	0.90 (0.57, 1.40)	0.81 (0.50, 1.30)	0.002
PFOS (ng/mL)	9.40 (4.90, 17.30)	9.80 (5.10, 18.00)	7.50 (4.02, 13.48)	< 0.001
PFHxS (ng/mL)	1.50 (0.90, 2.70)	1.53 (0.90, 2.70)	1.40 (0.80, 2.60)	0.064
NHANES wave				< 0.001
2003–2004	1121 (11.89)	1020 (12.96)	101 (6.73)	
2005–2006	1126 (12.84)	1001 (13.48)	125 (9.73)	
2007–2008	1284 (11.93)	1133 (12.31)	151 (10.10)	
2009–2010	1400 (12.34)	1165 (11.97)	235 (14.14)	
2011–2012	1128 (12.21)	953 (11.98)	175 (13.33)	
2013–2014	1176 (13.07)	967 (12.62)	209 (15.29)	
2015–2016	1125 (12.90)	929 (11.77)	196 (18.37)	
2017–2018	1109 (12.81)	968 (12.92)	141 (12.31)	

BMI body mass index, NHANES National Health and Nutrition Examination Survey, PFHxS perfluorohexane sulfonic acid, PFNA perfluorononanoic acid, PFOA perfluorooctanoic acid, PFOS perfluorooctane sulfonic acid, PIR poverty income ratio

Bold type denotes $P \leq 0.05$

^a Median (IQR)—continuous variables; n (%)—categorical variables

^b Wilcoxon rank-sum test for complex survey samples; chi-squared test with Rao & Scott's second-order correction

Table 2 Association between probiotic consumption and PFAS serum concentrations

PFAS	Model 1, β (95% CI)	P value	Model 2, β (95% CI)	P value	Model 3, β (95% CI)	P value
PFOA	-0.46 (-0.67, -0.25)	<0.001	-0.43 (-0.63, -0.23)	<0.001	-0.19 (-0.35, -0.02)	0.027
PFNA	-0.12 (-0.20, -0.04)	0.003	-0.13 (-0.21, -0.05)	0.002	-0.07 (-0.14, 0.01)	0.072
PFOS	-3.28 (-4.31, -2.26)	<0.001	-3.21 (-4.23, -2.19)	<0.001	-1.27 (-2.23, -0.32)	0.010
PFHxS	-0.14 (-0.36, 0.07)	0.181	-0.07 (-0.27, 0.13)	0.491	-0.02 (-0.22, 0.18)	0.865

PFHxS perfluorohexane sulfonic acid, PFNA perfluorononanoic acid, PFOA perfluorooctanoic acid, PFOS perfluorooctane sulfonic acid

Model 1: unadjusted

Model 2: adjusted for age, gender, and body mass index

Model 3: adjusted for age, gender, race, body mass index, highest educational level, marital status, poverty income ratio, smoking status, alcohol drinking, NHANES wave, and the intake of fish, meat, egg, energy, dietary folate equivalent, and dietary fiber

Bold type denotes $P \leq 0.05$

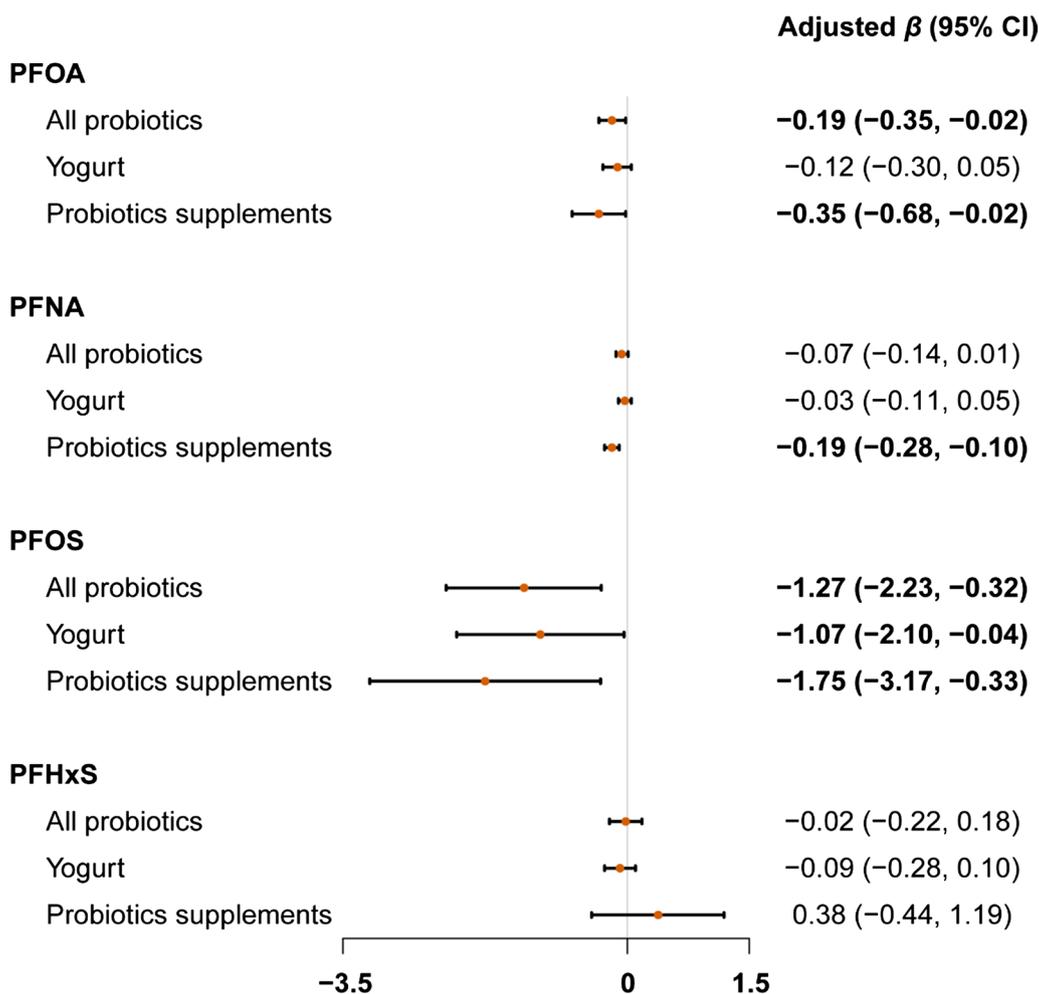


Fig. 1 Association between different sources of probiotics and serum PFAS concentration. Adjusted for age, gender, race, body mass index, highest educational level, marital status, poverty income ratio, smoking status, alcohol drinking, NHANES wave, and the intake of fish, meat, egg, energy, dietary folate equivalent, and dietary fiber

Table 3 P-interaction values with various subgroups

	P values for the interaction			
	PFOA	PFNA	PFOS	PFHxS
Age	0.656	0.414	0.245	0.144
Gender	0.496	0.482	0.268	0.589
BMI	0.506	0.297	0.251	0.462
Race	0.009	0.038	0.092	0.330
Marital	0.593	0.662	0.207	0.631
Education	0.012	0.277	0.154	0.223
PIR	0.021	0.173	0.022	0.411
Fish	0.220	0.844	0.415	0.702
Meat	0.046	0.198	0.106	0.817
Egg	0.305	0.272	0.454	0.451
Energy	0.539	0.866	0.464	0.822
Dietary fiber	0.658	0.550	0.934	0.326
Dietary folate equivalent	0.097	0.989	0.801	0.730
Smoking status	0.620	0.541	0.633	0.167
Heavy alcohol consumption	0.335	0.118	0.036	0.235
NHANES wave	0.315	0.212	0.467	0.448

BMI body mass index, NHANES National Health and Nutrition Examination Survey, PFHxS perfluorohexane sulfonic acid, PFNA perfluorononanoic acid, PFOA perfluorooctanoic acid, PFOS perfluorooctane sulfonic acid, PIR poverty income ratio

Meat (ounce/d)—Q1: < 2.19, Q2: 2.19–< 3.71, Q3: 3.71–< 5.74, Q4: 5.74+; Egg (ounce/d)—Q1: < 0.035, Q2: 0.035–< 0.230, Q3: 0.230–< 0.885, Q4: 0.885+; Energy (kcal/d)—Q1: < 1545, Q2: 1545–< 1972.5, Q3: 1972.5–< 2559, Q4: 2559+; Dietary fiber (g/d)—Q1: < 10.75, Q2: 10.75–< 15.25, Q3: 15.25–< 21.35, Q4: 21.35+; dietary folate equivalent (µg/d)—Q1: < 333, Q2: 333–< 466, Q3: 466–< 657, Q4: 657+

Adjusted for age, gender, race, body mass index, highest educational level, marital status, poverty income ratio, smoking status, alcohol drinking, NHANES wave, and the intake of fish, meat, eggs, energy, dietary folate equivalent, and dietary fiber

Bold type denotes P-interaction ≤ 0.05

and PFOA (among non-Hispanic White individuals, those with an education level beyond high school, high household income, and the highest quartile of meat intake), PFNA (among non-Hispanic White individuals), and PFOS (among moderate family incomes and heavy alcohol consumption) were more pronounced.

As far as we know, this study is the first large cross-sectional analysis investigating the association between probiotic consumption (through supplements or yogurt) and serum PFAS levels. Previous research on the association between probiotics and PFAS has been limited to animal and cell-based studies, which have found that probiotics can mitigate the toxicity of PFAS. Specifically, exposure to PFOA toxicity in mouse models has been shown to induce liver damage, reduces the abundance of beneficial probiotics, and lowers levels of SCFAs in feces, but these effects can be alleviated through the administration of probiotics or synbiotics [42–44]. Probiotics may alleviate PFOA toxicity by absorbing the compound,

exerting antioxidant effects, and regulating gut microbiota [43]. Additionally, an in vitro study assessed the binding capacity of 25 Lactobacillus strains to PFOA and found that *Lactobacillus plantarum* CCFM738 exhibited the highest binding capability. This strain also demonstrated excellent cellular antioxidative properties, tolerance to acidic and bile salt conditions, and adhesion to intestinal epithelial cells, which are important factors in assessing the functional properties of probiotics [45]. In mice, PFOS has been demonstrated to cause liver damage, alter lipid and glucose metabolism, and disturb intestinal homeostasis [46, 47]. Lactic acid bacteria can help mitigate PFOS toxicity by binding to PFOS, exerting antioxidant effects, and restoring the intestinal environment through up-regulating the content of SCFAs in the cecum and enhancing the expression of tight junction proteins in the intestines [48]. Our study reveals a negative association between probiotic consumption and serum PFAS concentrations, suggesting that probiotics may have the potential to alleviate the burden of PFAS in the human body and mitigate their detrimental effects.

According to the analysis of probiotic sources, the consumption of probiotic supplements, as opposed to yogurt, was observed to be associated with a decrease in serum levels of PFOA and PFNA. Both types of probiotics were significantly associated with a reduction in PFOS concentration. In this study, the consumption of a dietary supplement containing probiotics within the past 30 days was considered probiotic supplement consumption, which reflected a relatively long-term pattern of probiotic consumption. In contrast, yogurt consumption referred to consuming yogurt 1 day between 2 days, which had a certain degree of chance and did not accurately reflect a long-term habit of consuming yogurt. The quantity of probiotics ingested by individuals through yogurt consumption may vary depending on the brand, type, and serving size. Considering the persistence of PFAS compounds, this finding suggested that long-term probiotic consumption may have a stronger association with the reduction of serum PFAS concentrations.

Race, education, income, meat intake, and alcohol abuse may influence the relationship between probiotic consumption and specific PFAS concentrations. Compared to other races, a stronger negative association was observed between the consumption of probiotics and serum levels of PFOA and PFNA in non-Hispanic White individuals. This could be due to genetic, environment, and dietary differences among different races, as well as variations in their gut microbiota composition [49–51]. It is worth noting that the sample size of non-Hispanic White individuals was approximately 2–3 times larger than that of other races. In a study utilizing NHANES data from 1999 to 2018 to investigate the

trends in probiotic usage among adults and children in the US, it was observed that individuals with higher levels of education and income exhibited greater prevalence of probiotic use [26]. The effect heterogeneity by education and income in our study possibly because individuals with higher levels of education and income tend to have a greater interest in health-related knowledge and more disposable income. As a result, they are more likely to develop habits of consuming probiotic-rich foods or foods that promote the growth of beneficial gut bacteria. However, these possible confounding factors cannot be comprehensively addressed in our study. Previous studies have found a link between meat consumption and higher levels of PFAS [39, 40]. The observation of a stronger negative association between probiotic consumption and serum PFAS levels in populations with higher meat intake requires further investigation and careful evaluation. Heavy alcohol consumption has been found to potentially interact with PFAS in relation to liver function biomarkers [52]. Our finding that associations were more apparent among individuals with heavy alcohol consumption should be interpreted with caution, given the small sample size of the population with heavy alcohol consumption. One possible explanation is that the antagonistic effect of probiotics on PFAS levels could be influenced by impaired liver function due to alcohol abuse. Alternatively, unmeasured confounding variables may contribute to difference by groups.

When interpreting the results, it is important to take into account several limitations of the present study. Firstly, the cross-sectional study design hinders the establishment of temporal relationships between probiotic exposure and PFAS concentrations in serum, and therefore, causality cannot be determined. It is not possible to infer whether probiotic consumption leads to a decrease in PFAS concentration. Although we made efforts to control for potential confounding factors, there may still be unaccounted or residual confounding. It is possible that other dietary sources of probiotics not considered in this study could also influence the association between yogurt consumption and serum PFAS levels. Future studies could examine the effects of other fermented foods to expand on our findings. Additionally, the study population was categorized based on whether they consumed probiotics (either through supplements or yogurt), but information regarding the specific types, quantity and quality of probiotics consumed was unavailable. Therefore, we were unable to investigate the potential relationship between the dosage and types of probiotics ingestion and PFAS levels. Finally, the available information regarding the exposure of other PFAS

compounds released into the environment as PFOS and PFOA are phased out and replaced remains unclear. It is important to recognize the need for further monitoring to understand the potential implications of exposure to these emerging PFAS compounds.

Conclusion

In summary, this large nationally representative study observed negative associations between probiotic consumption and several serum PFAS levels. Considering the widespread exposure, persistent nature, and harmfulness of PFAS, these findings present a promising approach to mitigating the body burden of PFAS. Further prospective and experimental studies are needed to confirm or refute the association between probiotic consumption and PFAS concentration, and to explore the underlying mechanisms involved.

Abbreviations

BMI	Body mass index
LOD	Limit of detection
NHANES	National Health and Nutrition Examination Survey
PFHxS	Perfluorohexane sulfonic acid
PFNA	Perfluorononanoic acid
PFOA	Perfluorooctanoic acid
PFOS	Perfluorooctane sulfonic acid
PIR	Poverty income ratio
SCFAs	Short-chain fatty acids

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s12302-023-00808-2>.

Additional file 1: Figure S1. A schematic flow chart of study design.

Table S1. Probiotic search terms. **Table S2.** Association between probiotic consumption and PFAS serum concentrations according to different subgroups.

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Author contributions

YL designed research, analyzed data and drafted the manuscript; JC analyzed data and interpreted data; JL and JW revised the manuscript; LC and MG conceived research, revised the manuscript and had primary responsibility for final content. All authors have read and approved the final manuscript.

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Availability of data and materials

The datasets generated during and/or analyzed during the current study are available in the NHANES repository, <https://www.cdc.gov/nchs/nhanes/> [53]. Further information is available from the corresponding author upon request.

Declarations

Ethics approval and consent to participate

All individuals provided written informed consent and all protocols were approved by the Research Ethics Review Board of the National Center for Health Statistics.

Consent for publication

Not applicable.

Competing interests

The authors have no relevant financial or non-financial interests to disclose.

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