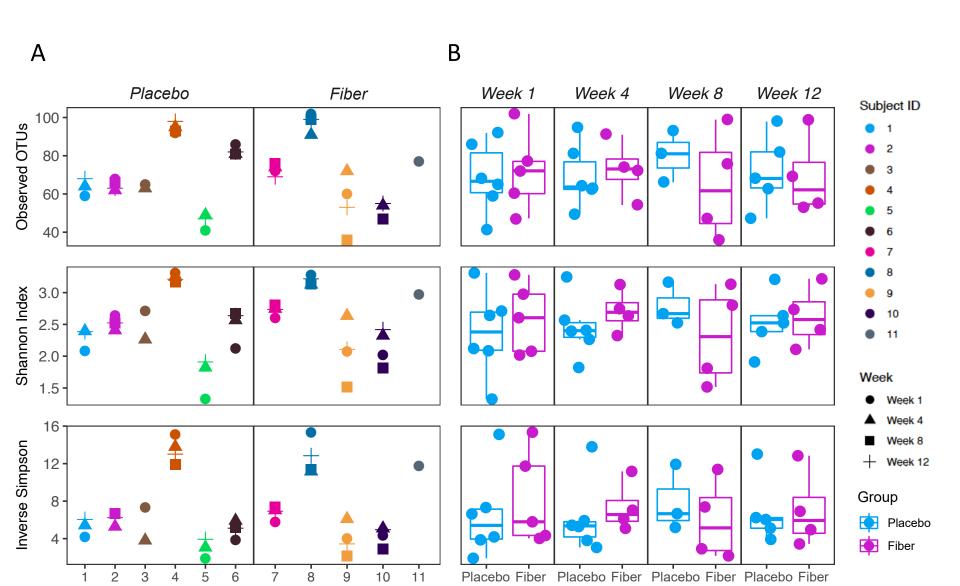
Figures

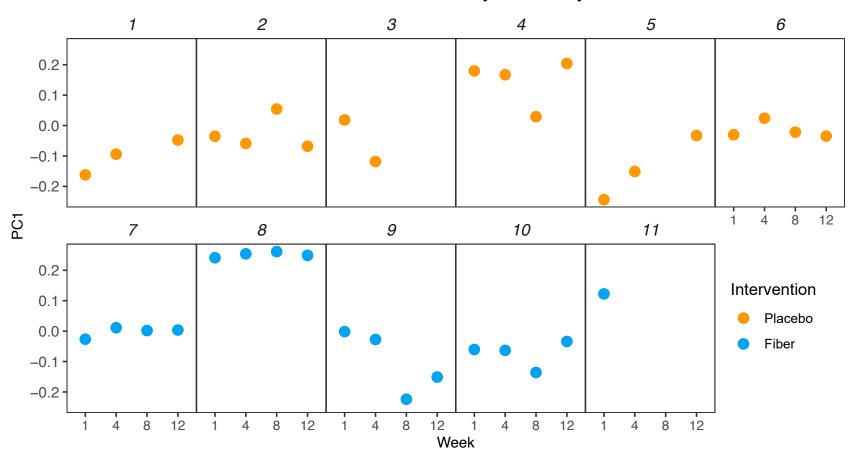
Fiber-Microbiome Study

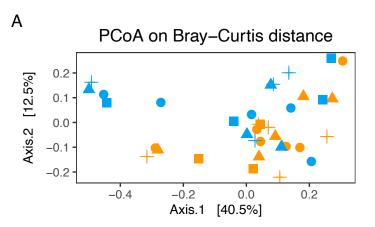
Questions to be addressed

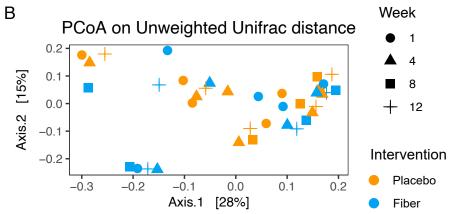
- What is the effect of Prebiotin vs placebo on the gut microbiome (controlling for, age, ethnicity)
 - Stability over time pre vs post intervention (Johnson paper Fig. 6E)
 - Diversity over time pre vs post intervention
 - Alpha FIG 1
 - Beta FIG 2/3
 - Phylogenetic FIG 4
- What is the effect of Prebiotin vs placebo on anthropometrics (controlling for diet, age, ethnicity)- Did the intervention mitigate excess weight gain? NO EFFECTS --> Table 3
 - BMI
 - Lean mass
 - Fat mass
 - Weight, overall
 - Weight change
- What is the effect of Prebiotin vs placebo on measures of stress
 - What is the correlation between the microbiome and stress? Table 3
- What is the effect of Prebiotin vs placebo on clinical variables (blood measures - lipids, PYY) Table 3

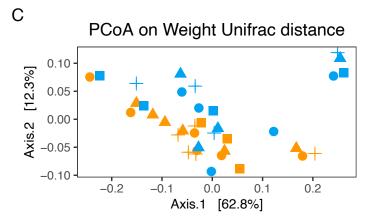


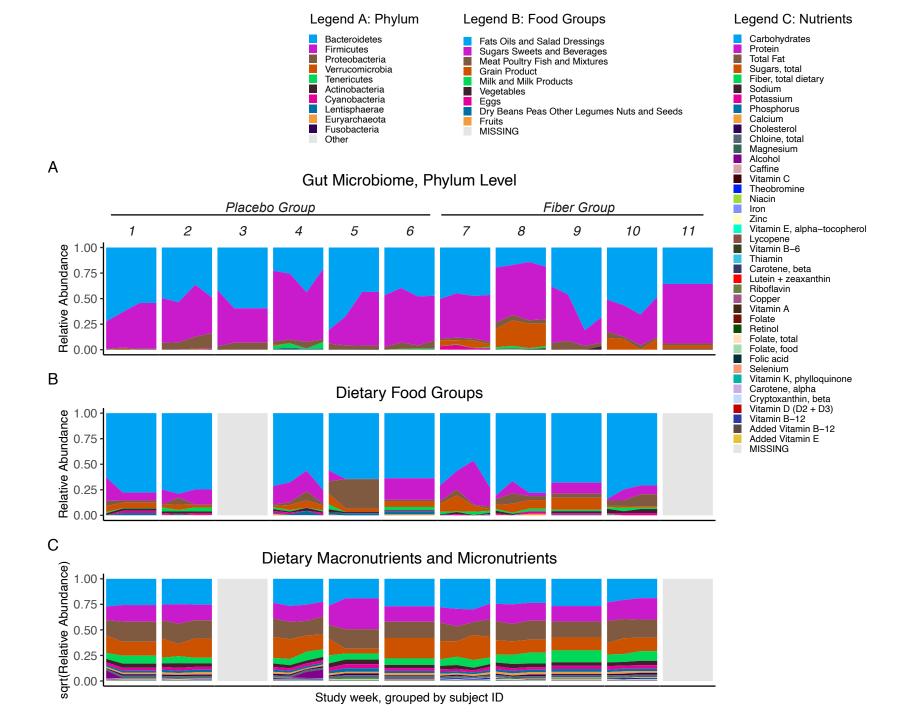
Individualized beta-diversity over study duration

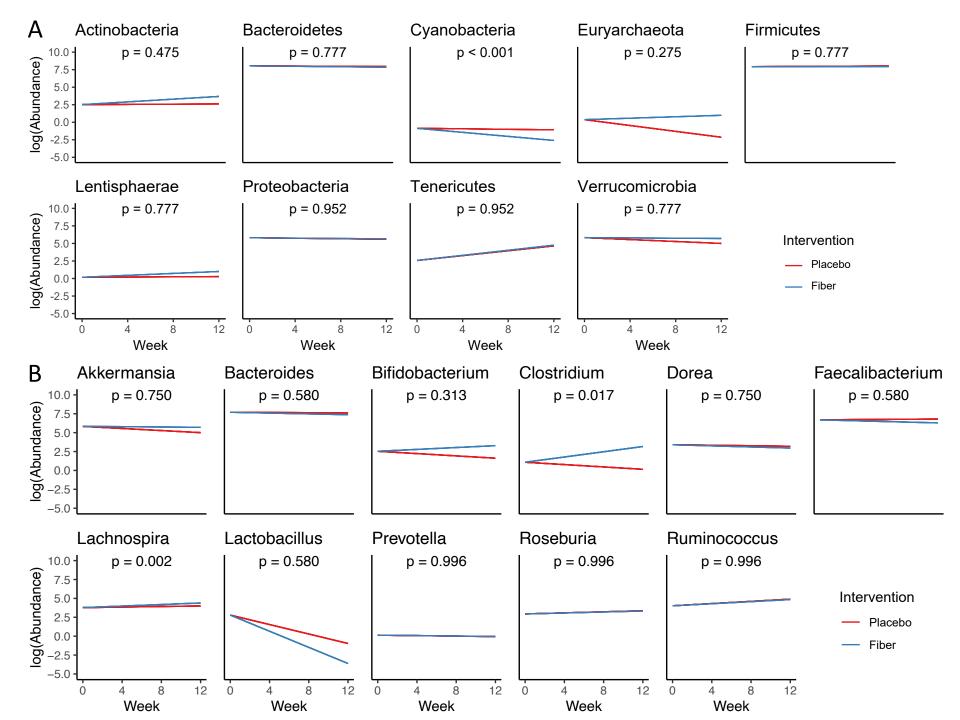


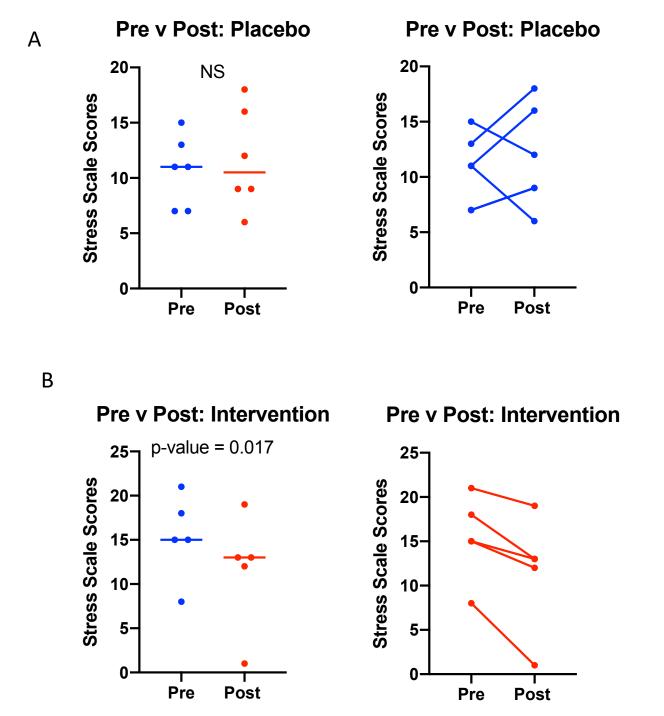












Tables

Tables 1 & S1

Table 1. Population Demographics

	Placebo mean (SD)	Intervention mean (SD)	P-value
N	6	5.00	
% Male	33.33	40.00	
Age	27.33(1.5)	28.4(2.6)	0.638
Weight (kg)	68.96(12.19)	73.88(9.23)	0.537
Height (cm)	164.26(10.06)	163.58(6.3)	0.927
BMI	25.45(3.12)	27.51(1.72)	0.247
VFL	8.83(4.3)	10.2(2.94)	0.405
% Body fat	29.75(8.55)	32.34(4.99)	0.464
% Lean fat	26.86(6.9)	27.86(3.99)	0.407
HEI score	89.36(0.44)	89.81(0.36)	0.201

P-values calculted from Student's t-tests

VFL=visceral fat level

Supplemental Table 1						
Week	ASA24	FFQ*	Stool Samples	Blood Samples*		
Baseline (Pre)	9	9	11	11		
4	7	-	10	-		
8	4	-	7	-		
12 (Post)	1	-	9	11		

^{*}FFQ was only given once at baseline

^{*}Blood Samples only collected Pre-Post

Table S2 - Intervention Effect of Change in Alpha Diversity

Table S2: Intervention effect on changes in alpha diversity							
Metric	Estimate	Std. Error	df		t value	Pr(> t)	p.adjust
Observed	-2.291	2.097		24.504	-1.093	0.285	0.569
Shannon	-0.065	0.072		24.78	-0.905	0.374	0.569
InvSimpson	-0.309	0.382		24.367	-0.807	0.427	0.569

Table S3: Intervention Effect on Change in Clinical Biomarkers

Table S3. Effect of intervention on change in clinical biomarkers						
Outcome	Estimate*	Std. Error	t value	Pr(> t)	p.adj	
Glucose	13.47	7.29	1.85	0.082	0.489	
BUN	-3.60	1.83	-1.97	0.065	0.474	
Creatinine	-0.07	0.08	-0.91	0.376	0.786	
Sodium	-0.70	1.45	-0.48	0.636	0.898	
Potassium	0.12	0.34	0.34	0.736	0.903	
Chloride	1.47	1.67	0.88	0.393	0.786	
arbon_Dioxid	-1.00	2.02	-0.49	0.628	0.898	
Calcium	-0.30	0.23	-1.32	0.206	0.609	
Cholesterol	-2.47	30.98	-0.08	0.937	0.957	
Triglycerides	38.77	72.27	0.54	0.599	0.898	
HDL	2.43	13.11	0.19	0.855	0.929	
LDL	-10.13	26.73	-0.38	0.709	0.903	
.DL_HDL_Rati	-0.50	0.63	-0.80	0.436	0.804	
hsCRP	2.90	2.21	1.32	0.206	0.609	
Insulin	-0.27	24.02	-0.01	0.991	0.991	
PYY	26.90	160.88	0.17	0.869	0.929	

^{*}multivariable model adjusted for age, ethnicity, and stress scores

Table S4

Table S4. Effect of intervention on phylum level relative abundance						
Phylum	Estimate	Std. Error	z value	Pr(> z)	p.adjust	
Actinobacteria	0.356	0.285	1.250	0.211	0.475	
Bacteroidetes	-0.032	0.066	-0.486	0.627	0.777	
Cyanobacteria	-0.504	0.069	-7.302	< .001	< .001	
Euryarchaeota	1.043	0.618	1.687	0.092	0.275	
Firmicutes	-0.026	0.047	-0.564	0.573	0.777	
Lentisphaerae	0.243	0.509	0.477	0.633	0.777	
Proteobacteria	0.018	0.131	0.134	0.893	0.952	
Tenericutes	0.035	0.579	0.060	0.952	0.952	
Verrucomicrobia	0.236	0.468	0.504	0.614	0.777	

Table S5

Table S5. Effect of intervention on changes in relative abundance							
Genus	Estimate	Std. Error	z value	Pr(> z)	p.adjust		
Akkermansia	0.236	0.468	0.504	0.614	0.750		
Bacteroides	-0.076	0.085	-0.898	0.369	0.580		
Bifidobacterium	0.553	0.368	1.504	0.133	0.313		
Clostridium	1.004	0.363	2.766	0.006	0.017		
Dorea	-0.078	0.149	-0.524	0.601	0.750		
Faecalibacterium	-0.170	0.183	-0.932	0.351	0.580		
Lachnospira	0.131	0.037	3.500	0.000	0.002		
Lactobacillus	-0.883	0.976	-0.906	0.365	0.580		
Prevotella	0.003	0.613	0.005	0.996	0.996		
Roseburia	-0.011	0.270	-0.043	0.966	0.996		
Ruminococcus	-0.018	0.307	-0.059	0.953	0.996		

Supplemental Figures

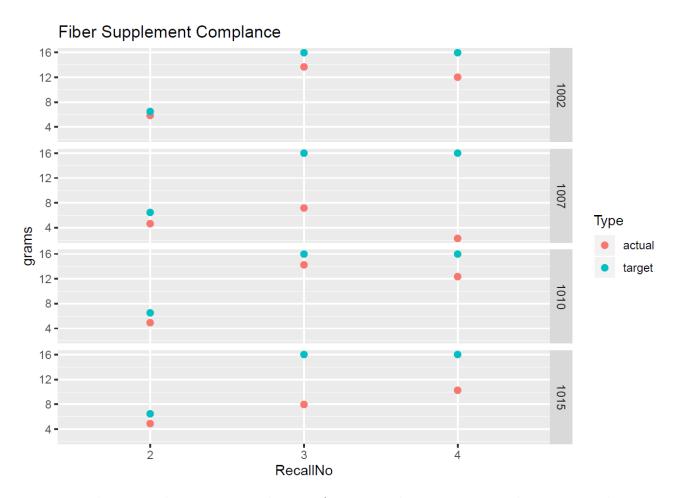


Fig. S1. Fiber Supplement Compliance. *Note: Only participants that received actual supplements are displayed in this chart. Participants receiving placebo were not included.

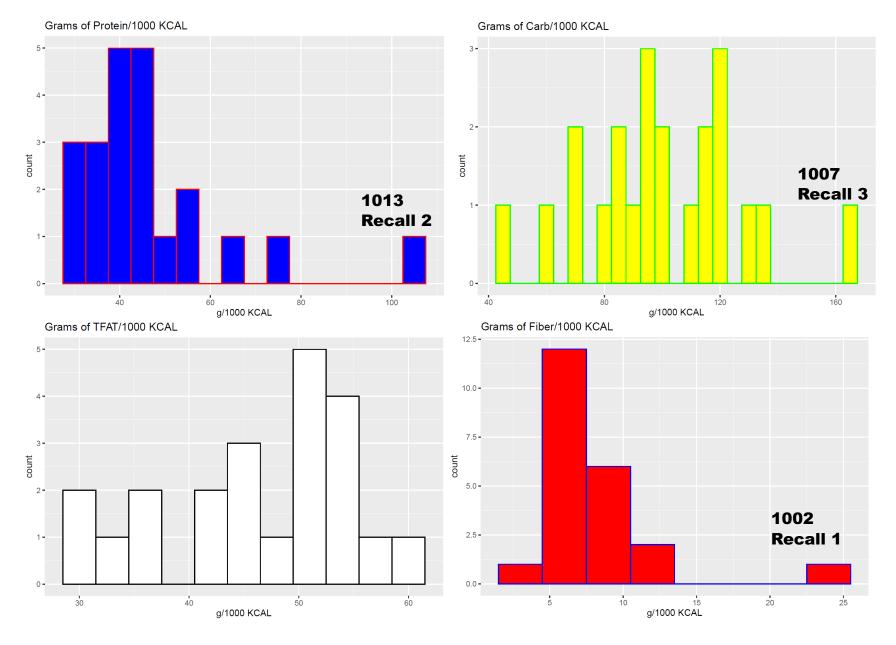


Fig. S2. Assessment of outliers from ASA24 dietary recalls. NHANES interquartile range was used to evaluate outliers along with visual inspection of dietary food reports.

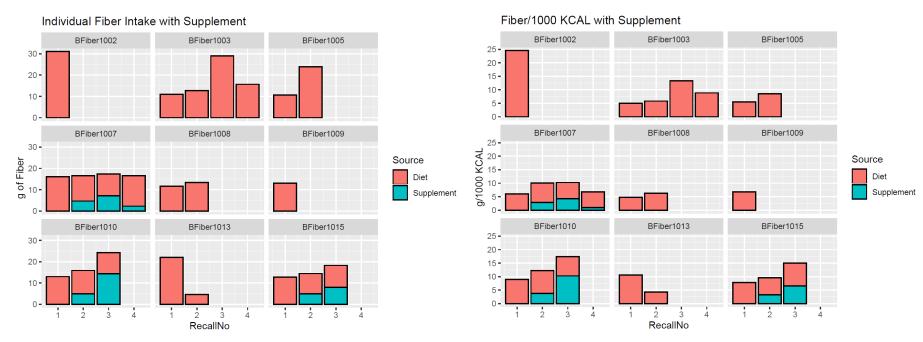


Fig. S3. Individual dietary and fiber supplement intake. Left panel are values by grams of dietary fiber; Right panel is fiber intake per 1000 kcals. *Note: Fiber1002 did not complete ASA24 after Recall 1, thus fiber supplement could not be added. 1007, 1010, and 1015 show supplemental fiber intake. 1003, 1005, 1008, 1009, 1013 received placebo, thus no supplemental fiber was added to their data.

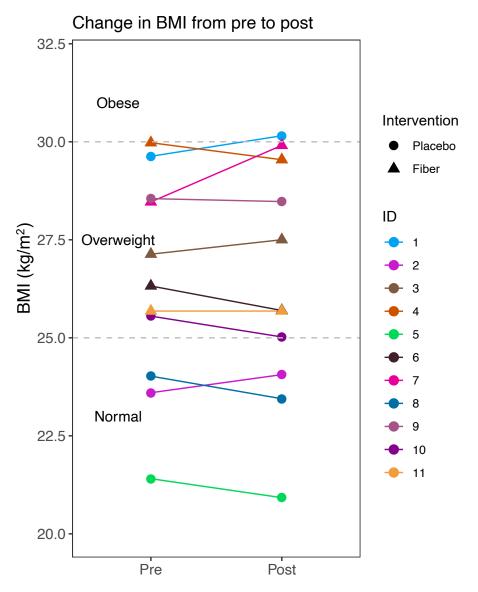
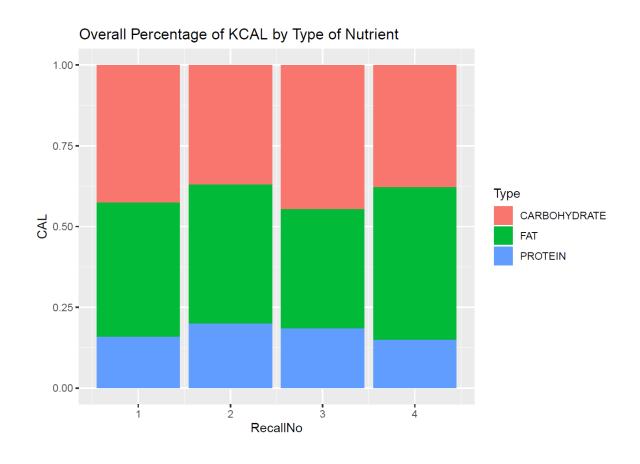


Fig. S4. Individual changes in BMI at baseline at end of the intervention. Red lines are the WHO guidelines for differentiating normal, overweight, and obese BMI.

Discussion

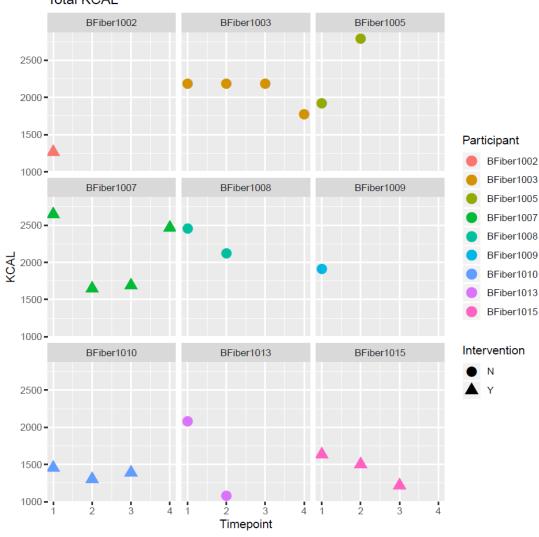
- Cookies were not accounted for
 - Some participants logged the cookies. Others did not
 - Those that logged the cookies logged them all differently
 - How would we like to handle this?
- Most participants given supplement were very compliant with the exception of 1007 over the last 4 weeks
- Most of placebo group appeared to be fairly noncompliant during the last 4 weeks with the exception of 1003
- Supplement did appear to make a difference in overall fiber consumption in test participants with the exception of 1007, who was noncompliant during the last 4 weeks, and 1002, who did not complete enough ASA_24 to analyze.

Overall Macronutrient Percentages



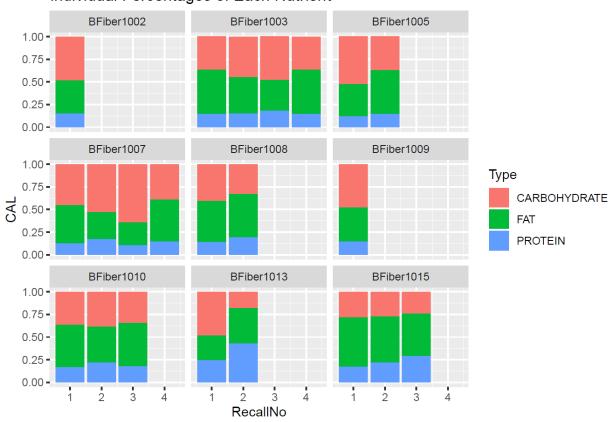
Total KCAL Intake (ASA24)

Total KCAL

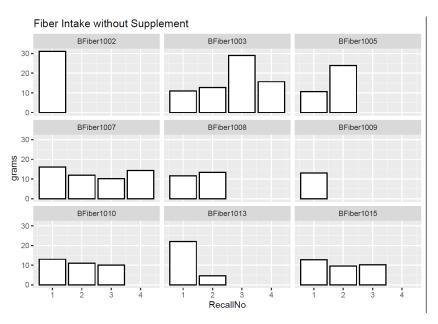


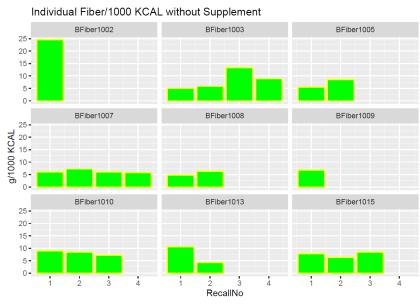
Individual Percentages of Each Nutrient

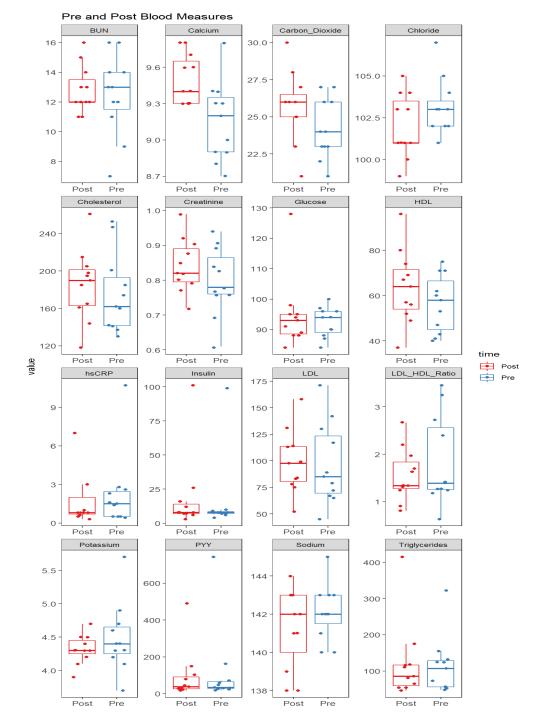




Individual Fiber Intake Without Supplement







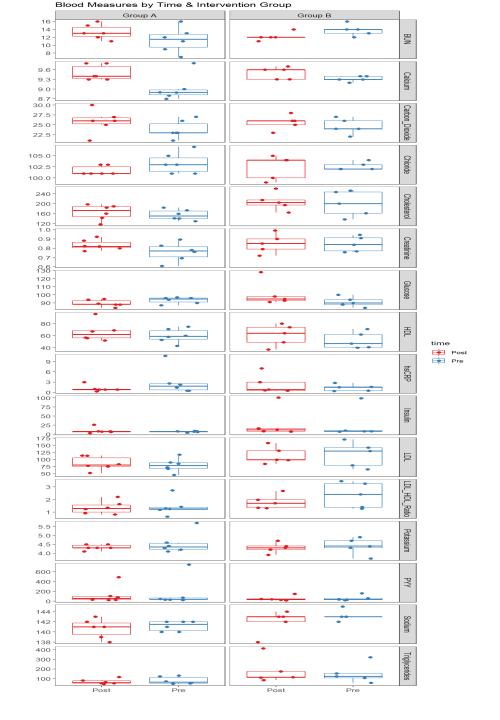


Table S3

Supplemental Table 1							
	Baseline		1.5 Year Follow Up				
Variables	Mean	Standard Deviation	Mean	Standard Deviation			
Weight	76.92	17.61	3.02	4.99			
Waist	34.90	4.00	0.87	2.95			
DXA	26.33	8.79	3.35	2.51			
Fat weight	44.69	16.62	9.27	6.45			
Lean weight	127.52	32.82	-3.50	9.95			
Visceral fat	328.24	144.84	54.67	65.35			
Total cholesterol	183.04	29.28	8.00	27.95			
LDLC	111.21	24.94	7.60	23.51			
HDLC	55.88	14.80	-5.82	5.77			
Triglycerides	95.92	52.53	17.30	59.64			
Glucose	89.92	7.58	9.95	13.74			

Preliminary Findings: Findings from an on-going study by co-investigator, Dr. Grandjean, indicate that over a short 18-month period, residents experience an increase in body weight, fat mass, waist circumference and loss of lean mass. Also of concern is that their lipid profiles change unfavorably and glucose levels increase. Please see table 1 for details.