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ABSTRACT

Objective: The purpose of this study was to compare equivalent amounts of two choline sources on plasma choline and its derivatives including trimethylamine N-oxide (TMAO), inflammatory markers and changes in fecal microbiota in men and women aged 32 to 70 years old with metabolic syndrome (MetS).

Methods: Twenty-three subjects with MetS were included in this randomized, crossover clinical trial. Participants underwent an initial period of 2 weeks without consuming any eggs, which was followed by a random allocation to either 3 eggs/day or a choline-supplement for 4 weeks (both diets had a choline equivalent of 400 mg per day). Following a 3-week washout period, participants were allocated to the alternate diet. We measured plasma choline and plasma TMAO as well as C-reactive protein (CRP), inflammatory markers, insulin, liver enzymes and analyzed fecal microbiota.

Results: Although there was an overall significant increase in plasma choline after egg intake, compared to baseline ($P < 0.01$), there were no significant differences between egg and supplement at the end of the respective interventions ($P > 0.05$). Baseline values were 7.9 ± 2.1 nmol/ml compared to 9.9 ± 2.2 and 9.5 ± 2.1 nmol/ml for the egg and supplement, respectively ($P < 0.01$). In addition, plasma TMAO was not different between baseline, or at the end of the egg and supplement periods ($P > 0.1$). In regards to inflammatory and insulin resistance markers, C-reactive protein, plasma insulin, and insulin resistance (HOMA-IR) were each lower after the egg period compared to baseline, while no differences in these parameters were observed after the supplement period when compared to baseline. In contrast, plasma IL-6 concentrations were lower by both dietary treatments compared to baseline. There were no significant differences in fecal bacterial diversity when baseline was compared to either the egg or the choline supplement treatment by using the Shannon diversity index or when time and treatment were compared. Some interesting observations were that the family of *Lachnospiraceae* (Phylum Bacteroidetes) appeared to increase after choline treatment (both after the supplement and the egg period). Also *Akkermansia* (Phylum Verrucomicrobia) was higher after both treatments, yet more so during after the choline period.

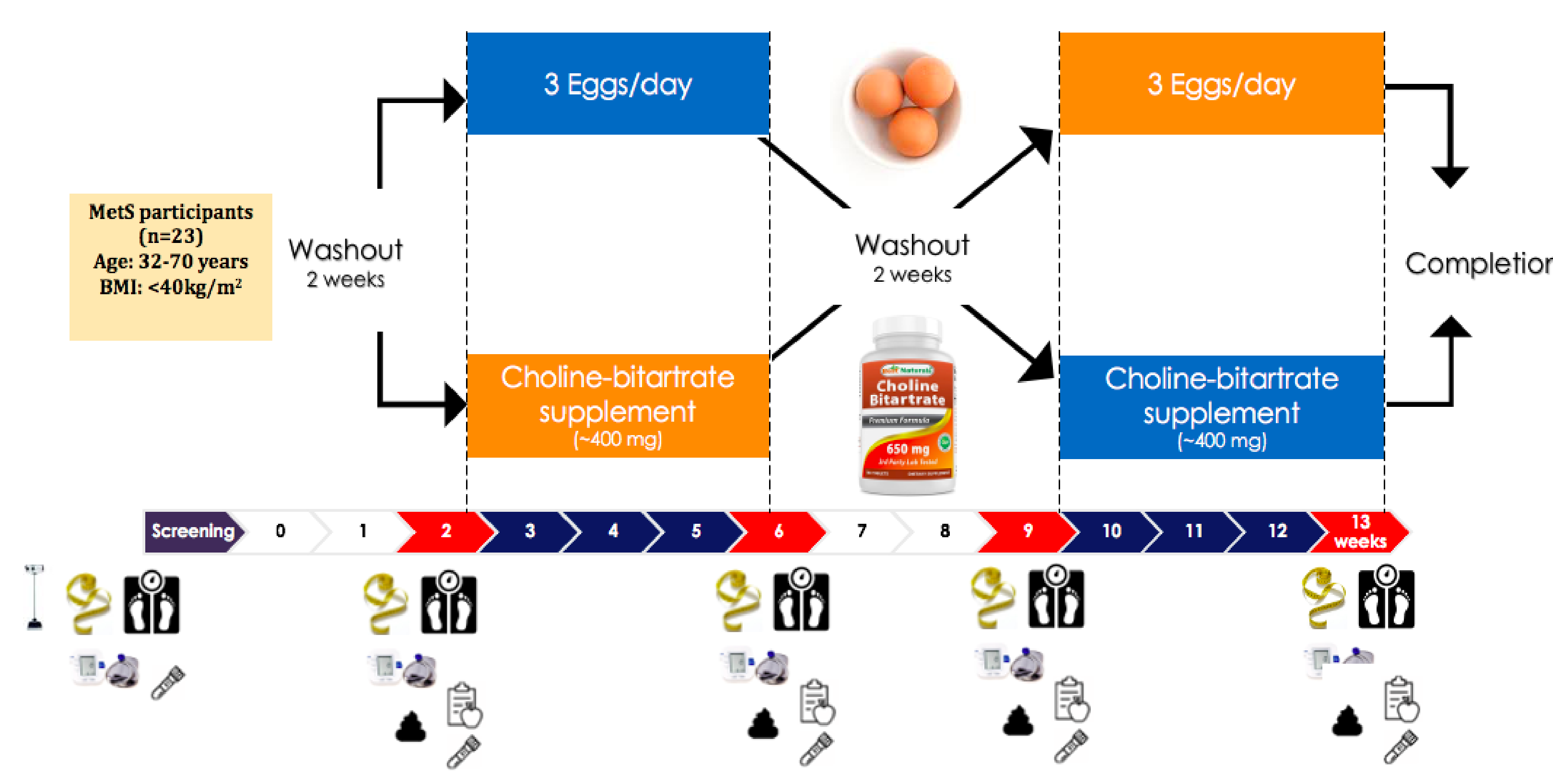
Conclusions: These studies indicate that plasma choline response appears to be similar in MetS participants, independent of its source or chemical composition. Analyzing the fecal microbiota of these subjects also indicated no significant differences between treatments indicating that both bioavailability of choline, TMAO production and microbiota composition are not affected by choline source in MetS patients.

OBJECTIVE AND AIMS

Objective: To evaluate the effect of two sources of choline: phosphatidyl choline (from eggs) vs choline bitartrate from supplement on choline metabolism and microbiota diversity in participants with MetS.

- AIM 1:** Measure plasma choline and metabolites including betaine, dimethylglycine (DGM), and TMAO
- AIM 2:** Evaluate whether inflammatory markers were reduced in these subjects
- AIM 3:** Evaluate microbiota diversity and potential changes due to dietary treatments

EXPERIMENTAL DESIGN



MATERIALS AND METHODS

Study Population	Metabolic Syndrome Participants (3/5 of NCEP-ATP III criteria)				
N = 23	Waist circumference	Blood pressure	HDL-C	Triglycerides	Fasting blood glucose
	≥ 102 cm for men or ≥ 88 cm for women	$\geq 130/85$ or $Sys \geq 130$ or $Diast \geq 85$	< 40 mg/dL for men or < 50 mg/dL for women	≥ 150 mg/dL	≥ 100 mg/dL
Diet Assessment	Fasting blood samples		Stool samples		
	3 day-diet record pre- and post egg or supplement treatment		Fasting blood collected from antecubital vein at the beginning and at the end of each intervention, centrifuged at 2000 x g for 20 minutes, and plasma collected and stored at -80°C for analyses		Feces collected in OMNigen OM 200 stool kits Aliquoted
Diets	Blood Biomarkers		Gut microbiome analysis		
	Evaluated using NDSR software		Plasma choline, Betaine, DGM and TMAO evaluated using LC-MS/MS IL-6, Mtp-1, TNF- α measured using Luminex Multiplex CRP measured using Cobas c-111 analyzer Plasma Insulin was measured using ELISA immunoassay (R&D systems)		16S rRNA gene sequencing for microbiome analysis Processed in Mothur v. 1.39.1

Statistics were performed in SPSS version 20. Data are represented as mean \pm SD in all figures; $p < 0.05$ was considered significant

RESULTS

- Dietary choline significantly increased to meet the DRI (DRI: 550 mg/day of choline for men and 425 mg/day for women)

Plasma concentration of Choline and metabolites at baseline and after consumption of 3 eggs or choline bitartrate supplementation for 4 weeks

Parameter	Baseline	Eggs	Choline supplement
Choline (nmol/mL) ²	7.9 ± 2.0^a	9.9 ± 2.2^b	9.6 ± 2.1^b
TMAO (nmol/mL)	4.1 ± 3.0	5.6 ± 3.5	5.2 ± 5.3
Methionine (nmol/mL)	29.4 ± 4.0	29.8 ± 4.0	29.9 ± 5.2
Betaine (nmol/mL)	37.6 ± 14.8^a	43.0 ± 14.7^b	43.5 ± 18.7^b
DGM (nmol/mL)	2.4 ± 0.6^a	2.8 ± 0.7^b	2.8 ± 0.6^b

¹Data are presented as mean \pm SD for n = 23 subjects. ² Numbers in the same row with different superscripts are significantly different ($P < 0.01$)

Concentration of inflammatory markers IL-6, CRP and Insulin at baseline and after consumption of 3 eggs or choline bitartrate supplementation for 4 weeks

Parameter	Baseline	Egg	Choline supplement
IL-6 (pg/ml) ²	5.6 ± 0.9^a	4.6 ± 1.7^b	4.4 ± 1.3^b
CRP	0.48 ± 0.57^a	0.32 ± 0.31^b	0.36 ± 0.31^{ab}
Insulin (pg/mL)	152.4 ± 87.2^a	97.7 ± 66.5^b	111.3 ± 100^{ab}

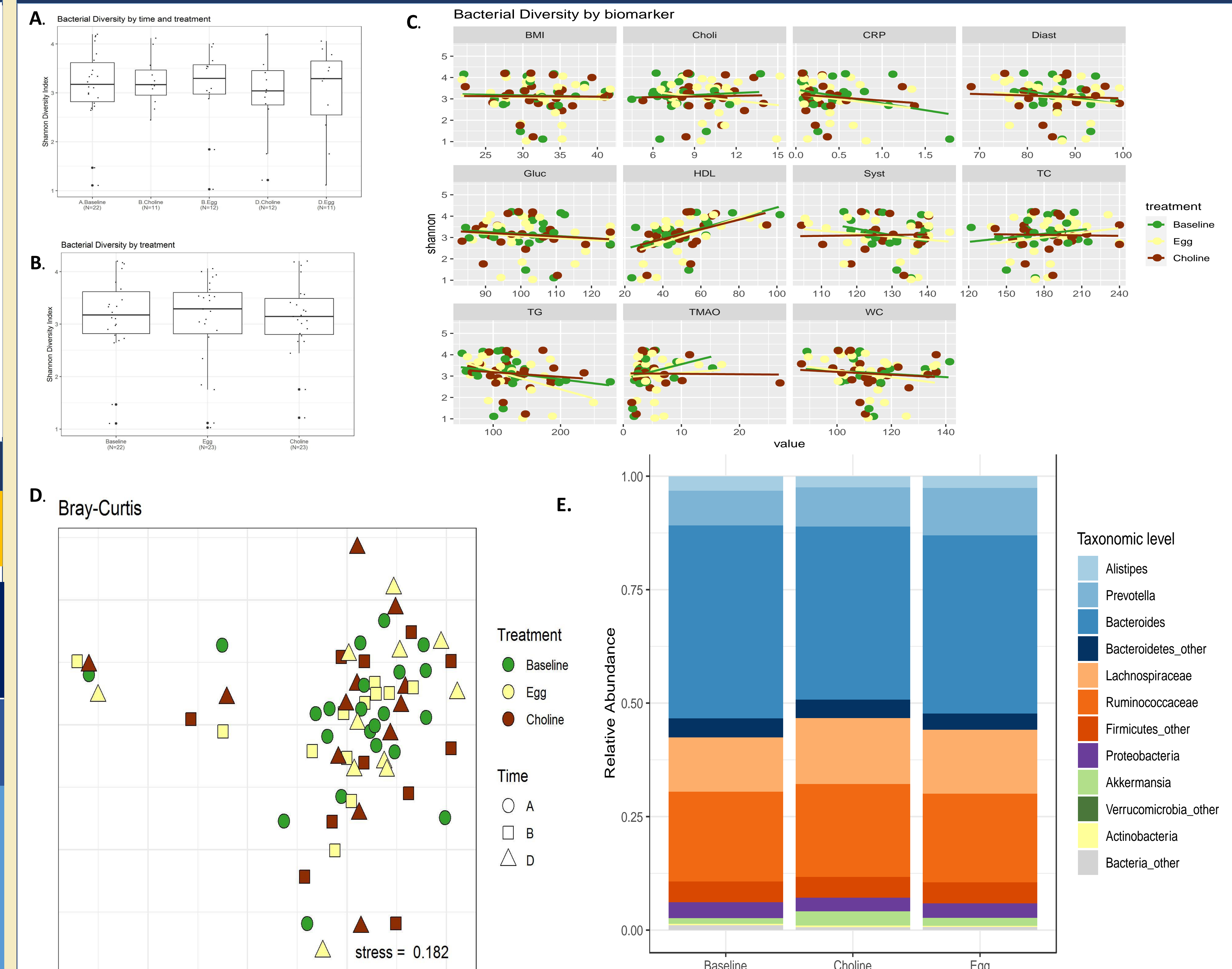
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FUNDING SOURCE



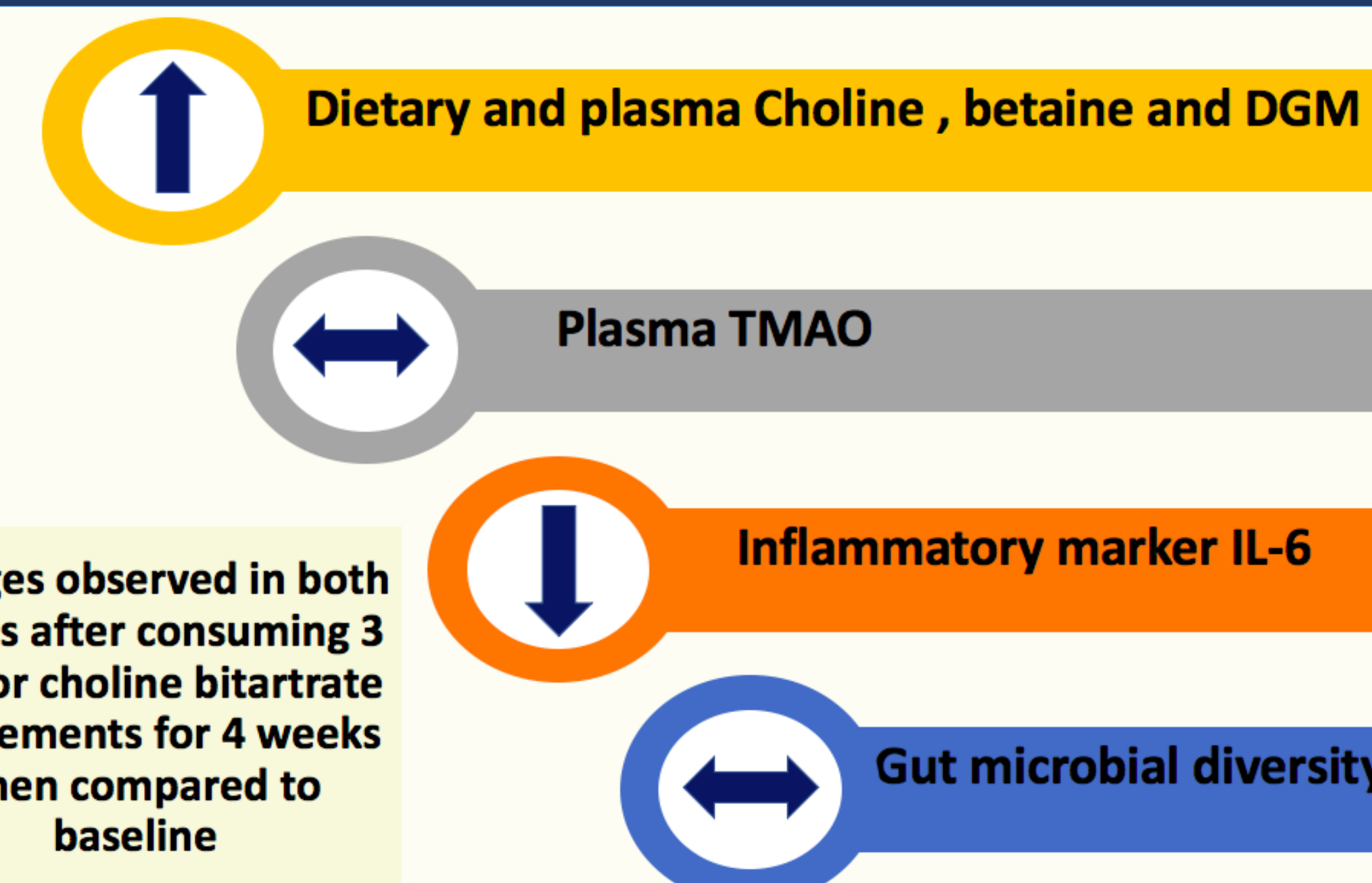
This study was funded Egg Nutrition Center

RESULT



(A,B) Alpha diversity measures by time and treatment (C) Correlations between plasma biomarkers and bacterial diversity based on treatments (D,E) Nonmetric multidimensional scaling of the Bray-Curtis dissimilarity index. This shows the relationship between individual samples based on the species and their abundances. Both ordinations are of the same data, but the ordination on the right also includes an overlay of the OTUs that are significantly correlated with this ordination colored by Taxonomic level.

SUMMARY



CONCLUSION

Choline provided by two distinct sources: Phosphatidyl choline from eggs or choline bitartrate resulted in significant increases in plasma choline without changes in TMAO or microbiota diversity in participants with metabolic syndrome.