

Associations between Circulating Lipids and Fat-Soluble Vitamins and Carotenoids in Healthy Overweight and Obese Men

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ABSTRACT

Inconsistent associations between lipids and circulating markers of fat-soluble vitamin and carotenoid status have been reported. The aim of this hypothesis-generating study was to examine the contribution of the LC-MS-based lipidome, characterized by lipid class, carbon count, and the number of unsaturated bonds, to the interindividual variability in circulating concentrations of retinol, carotenoids, 25-hydroxyvitamin D₃, α -tocopherol, γ -tocopherol, and phyloquinone in 35 overweight and obese, but healthy men. A sparse partial least-squares method was used to accomplish this aim. Highly abundant phospholipids and triglycerides (TGs) contributed to the interindividual variability in phyloquinone, α -tocopherol, and γ -tocopherol. Interindividual variability in lycopene concentrations was driven by concentrations of low-abundant TG. 25-Hydroxyvitamin D₃, retinol, and the other carotenoids were not influenced by lipids. Except for lycopene, evaluation of lipids beyond class does not appear to further explain the interindividual variability in circulating concentrations of fat-soluble vitamins and carotenoids. *Curr Dev Nutr* 2020;4:nzaa089.

Keywords: lipids, fat-soluble vitamins, carotenoids, phyloquinone, tocopherol, retinol, 25-hydroxyvitamin D₃, lipidomics, micronutrients, liquid chromatography-mass spectrometry

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Supplemental Tables 1–4 are available from the “Supplementary data” link in the online posting of the article and from the same link in the online table of contents at <https://academic.oup.com/cdn/>.

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Abbreviations used: ChE, cholesterol ester; DG, diglyceride; FFA, free fatty acid; LPC, lysophosphatidylcholine; PC, phosphatidylcholine; sPLS, sparse partial least-squares; SPM, sphingomyelin; TG, triglyceride.

Introduction

Validated biomarkers of micronutrient status are important tools in observational studies linking micronutrient exposure to disease risk (1). In the case of hydrophobic fat-soluble vitamins and carotenoids, the associations between various lipid classes and the interindividual variability of these circulating biomarkers are not consistent (2–6).

In the fasted state, retinol and 25-hydroxyvitamin D₃ circulate on specific binding proteins, but vitamin E and vitamin K forms and carotenoids are transported with lipids on lipoproteins (2, 7–10). Blood lipid profiles are typically described by triglycerides (TGs), total cholesterol, LDL cholesterol, and HDL cholesterol. However, thousands of distinct lipid molecular species have been identified in the circulation (11). Whereas TGs, free cholesterol, and cholesterol esters (ChEs) are

the most abundant constituents of lipoproteins, glycerophospholipids, diglycerides (DGs), and free fatty acids (FFAs) constitute other important lipid classes (11, 12). Furthermore, esterified fatty acids can differ in carbon-chain length and the number of unsaturated bonds (11). With technological advances in characterization of the lipidome, there is an opportunity to broaden the study of lipids and their influence on these micronutrient biomarkers.

Following work by van den Broek et al. (13), in which micronutrients were correlated with molecules measured across multiple metabolomic and proteomic platforms, the objective of this hypothesis-generating study was to examine the contribution of the lipidome, characterized by lipid class, carbon-chain length, and number of unsaturated bonds, to the interindividual variability in circulating fat-soluble vitamins and carotenoids.

Methods

Study design and participants

This was a secondary analysis of data from the placebo arm of a randomized, double-blind, placebo-controlled, 5-wk intervention trial (NCT00655798) conducted from December 2006 to June 2007, to investigate the anti-inflammatory effects of nutritional interventions in 36 overweight and obese men with low-grade inflammation (14). The men (mean $\pm$ SD age: 47 ± 10 y) were otherwise healthy, with BMI $25.6\text{--}34.7$ kg/m². Exclusion criteria included high fasting total cholesterol; acute inflammation (C-reactive protein >10 mg/L); a chronic disease related to inflammation (e.g., arthritis or inflammatory bowel disease); use of anticoagulant, antiplatelet, or anti-inflammatory medication; smoking; unexplained weight loss; alcohol intake >28 units/wk; following a weight-reduction diet; or use of dietary supplements. Placebo capsules contained 365 mg microcrystalline cellulose (Microz Food Supplements) or 1360 mg soy lecithin (Solgar Vitamin and Herb). The study protocol was approved by the independent Medical Ethics Committee (METOPP) (Tilburg, Netherlands).

Biochemical analyses

At the end of the 5-wk placebo period, blood samples were collected 4 h after a light standard breakfast (1597 kJ, % energy: 8.6 protein, 17.0 fat, 72.4 carbohydrates) preceded by an overnight fast (14). Samples for plasma lipids and micronutrient analysis were collected in lithium-heparinized tubes and tubes containing tri-potassium salts of EDTA (K3-EDTA), respectively. Serum and plasma were centrifuged for 15 min at $2000 \times g$ at 4°C and separated within 30 min of collection and stored at -80°C in the dark until analysis.

Fat-soluble vitamins and carotenoids.

Retinol, α -tocopherol, γ -tocopherol, 25-hydroxyvitamin D₃, α -carotene, β -carotene, β -cryptoxanthin, and lycopene concentrations were measured in 2013, as previously described (13, 15, 16), in a laboratory that adheres to international bioanalytical guidelines for daily and long-term performance and participates in ring-tests. Serum phyloquinone was measured in 2019 from archived serum samples stored for ≤ 12 y at -80°C using reversed-phase HPLC with fluorometric detection in the Vitamin K Laboratory at the Human Nutrition Research Center on Aging at Tufts University, which participates in the vitamin K external quality assurance scheme (17, 18).

Plasma lipids.

Plasma lipids were analyzed directly after study conclusion (2006/2007) with electrospray LC-MS using a Thermo Linear Trap Quadrupole equipped with a Thermo Surveyor HPLC pump, described previously (19). As described, the LC-MS platform performance was assessed by quality control sample (pooled plasma from all participants) analysis, placed after every 10 samples, and method performance was monitored using 5 internal standards and duplicate sample analysis (20). Concentrations of 131 lipids including 6 lysophosphatidylcholines (LPCs), 19 phosphatidylcholines (PCs), 11 sphingomyelins (SPMs), 14 ChEs, 55 TGs, 4 DGs, and 22 FFAs were quantified. Baseline TGs, total cholesterol, and HDL cholesterol were measured with enzymatic techniques. LDL cholesterol was determined by the Friedewald calculation (14, 21).

Statistical analyses

Placebo-phase micronutrient and lipid data were not available for 1 participant so 35 of the 36 participants were included in this analysis. Circulating micronutrient and plasma lipid concentrations were described by means $\pm$ SDs. Given the small sample size and large number of highly correlated lipid exposures, we modeled the variability in circulating concentrations of each micronutrient using sparse partial least-squares (sPLS) regression. This method avoids overfitting when there may be few underlying or latent factors (components) that account for the variability in exposures and response variables (22). Using this technique, components are constructed to account for as much variability in the exposures as possible while maximizing the explained variability of the outcome (micronutrient). Loadings reflect the contribution of exposures (lipids) to the component (22).

The *spls* function from the mixOmics R package (23) in *regression mode* was used to decompose information from the 131 lipids into fewer components. The number of components for each model was determined by maximizing the predictability of the model with the fewest number of components that explained variability in both the predictor and response variables. A component was selected if its Q^2 value, a parameter used to assess the goodness of prediction, was ≥ 0.0975 , a cutoff proposed by Tenenhaus (24). R^2 is regarded as a measure of explained variability and Q^2 is equivalent to R^2 when the training set model is applied to a test set. A Q^2 that approximates R^2 is a qualitative indicator that the sPLS model performance was not unique to the training data set.

Each model was cross-validated using the *tune.spls* function with the *M-fold* method, which resamples the data into n (35 in this analysis) groups, replicated 100 times. With 1 independent outcome per model, we investigated the number of lipids (25, 50, 75, 100, 125, or 131) to retain for ≤ 10 components. Components were added to the model if there was a gain in performance based on a 1-sided t test. All variables were centered and scaled. Phyloquinone, α -tocopherol, γ -tocopherol, lycopene, α -carotene, and β -cryptoxanthin were skewed so these data were ln-transformed.

Results

Table 1 shows descriptive data for micronutrients and plasma lipids. ChEs, TGs, and PCs were the most abundant lipid classes. SPMs, FFAs, and DGs were present in very low quantities. ChEs (18:1) were ~ 4 times more abundant than any other lipid. TGs with 50, 52, and 54 carbons were also highly abundant. These TGs contained 1–3, 1–4, and 3–5 unsaturated bonds, respectively. The top PCs had 34 and 36 carbons.

Table 1 shows the number of components, exposures, R^2 , and Q^2 for each sPLS model. **Figure 1** and **Supplemental Tables 1–4** show the component 1 loadings for phyloquinone, α -tocopherol, γ -tocopherol, and lycopene, the micronutrients for which a robust model was established. Highly abundant LPCs (16:0) and PCs (34 or 36 carbons, 1–3 unsaturated bonds) explained the most variability in circulating phyloquinone. After these, several low-abundant ChEs and high-abundant TGs (50:2, 50:3, 52:1, 52:2) also contributed. The variability in γ -tocopherol was primarily explained by high-abundant TGs (52:4, 54:3–6, and 56:2–5) in addition to PCs (34:0) and few DGs. The

TABLE 1 Descriptive data for the fat-soluble vitamins, carotenoids, and total lipids with sparse partial least-squares model parameters for the micronutrients

	Mean $\pm$ SD	Components, <i>n</i>	Explained variability, R^2			Predictability, Q^2
			Exposures, <i>n</i>	Vitamin or carotenoid	Lipids	
Phylloquinone, ¹ nM	1.7 $\pm$ 1.7	1	131	1.00	0.33	0.33
Retinol, μ M	1.97 $\pm$ 0.31	1	131	1.00	0.34	0.03
α -Tocopherol, ¹ μ M	29.0 $\pm$ 6.2	1	75	1.00	0.30	0.27
γ -Tocopherol, ¹ μ M	1.74 $\pm$ 0.68	2	50, 131	1.00, 0.60	0.24, 0.21	0.20, 0.18
25-OH-vitamin D ₃ , nM	63.2 $\pm$ 32.8	9	50, 25, 131, 131, 131, 131, 100, 25, 50	1.00, 0.54, 0.45, 0.33, 0.26, 0.16, 0.11, 0.6, 0.5	0.10, 0.13, 0.10, 0.25, 0.11, 0.7, 0.3, 0.4, 0.2	0.17, -0.28, -0.38, -0.15, 0.26, 0.10, 0.21, -0.20, -0.34
Lycopene, ¹ μ M	0.62 $\pm$ 0.30	1	25	1.00	0.26	0.20
α -Carotene, ¹ μ M	0.06 $\pm$ 0.05	1	50	1.00	0.19	-0.17
β -Carotene, μ M	0.40 $\pm$ 0.17	1	100	1.00	0.22	0.03
β -Cryptoxanthin, ¹ μ M	0.22 $\pm$ 0.20	1	25	1.00	0.25	-0.15
Total cholesterol, ² mg/dL	231 $\pm$ 37					
LDL cholesterol, ² mg/dL	152 $\pm$ 31					
HDL cholesterol, ² mg/dL	47 $\pm$ 9					
Total triglycerides, ² mg/dL	147 $\pm$ 72					

¹Skewed distribution.²Plasma lipids were measured at the baseline study visit.

variability in α -tocopherol was also explained by TGs and select PCs and DGs, with an overlap in carbon count and the number of unsaturated bonds. The variability in circulating lycopene was explained by TGs with 40–52 carbons. All loadings were negative, indicating an inverse relation. The top contributors had 40–46 carbons and 0–2 unsaturated bonds. Q^2 was not comparable with R^2 for retinol, α -carotene, β -carotene, and β -cryptoxanthin so these models were not robust. For 25-hydroxyvitamin D₃, the model appeared overfitted, requiring 9 components, none of which sufficiently accounted for variability in the exposures and response variables with adequate performance. For example, component 1 maximized the variability in circulating 25-hydroxyvitamin D₃ with $Q^2 > 0.0975$, but only explained 10% of the variance in lipids. In contrast, component 4 explained more variance in lipids (25%), but limited variability in 25-hydroxyvitamin D₃ (33%) and had poor performance ($Q^2 = -0.15$).

Discussion

Using a sPLS technique, the relations between lipids (characterized by class, carbon-chain length, and the number of unsaturated bonds) and the interindividual variability in blood-based markers of fat-soluble vitamin and carotenoid status were assessed in healthy overweight and obese adult males. We found that lipids contributed to the variability in circulating phyloquinone, α -tocopherol, γ -tocopherol, and lycopene, but not 25-hydroxyvitamin D₃, retinol, α -carotene, β -carotene, or β -cryptoxanthin.

The largest interindividual variability in circulating concentrations was observed for phyloquinone. We observed that high relative abundance TGs contributed to the variability in serum phyloquinone, but the contribution of highly abundant PC and LPC was unexpectedly greater. Phyloquinone concentrates in the TG-rich lipoprotein fraction (10), but it is plausible phyloquinone and TGs share

transport mechanisms without a causal relation between the molecules. That high-abundant lipids explained the most variability in circulating phyloquinone suggests that total PCs and TGs sufficiently account for the variability in circulating phyloquinone and differentiation by number of carbons and unsaturated bonds does not explain more. These data also suggest that variances in PC/LPC may be more indicative of circulating phyloquinone than variances in TGs. The association between circulating phyloquinone and PC/LPC is not well-studied. However, there is biological plausibility to our findings. PC is critical to VLDL synthesis and higher circulating phyloquinone has been associated with higher VLDL particle number, independently of TGs (25) (JM Kelly, unpublished results). In addition, based on the chemical structure of phyloquinone, we posit that phyloquinone resides at the surface of the lipoprotein with PC rather than in the core with TGs (26, 27). Further research is required to confirm this. Importantly, these findings should not be extrapolated to other forms of vitamin K because the transport of menaquinones in the body is not well understood.

Consistent with the report by van den Broek et al. (13), the interindividual variability in α -tocopherol concentrations was also sensitive to variances in highly abundant PC/LPC. We also observed contributions of TGs for both vitamin E vitamers. That vitamin K and vitamin E vitamers were similarly influenced by lipids is in line with evidence that suggests there is overlap in the metabolic pathways of these 2 fat-soluble vitamins (26, 28, 29).

In contrast to the observations for phyloquinone, α -tocopherol, and γ -tocopherol, lycopene was inversely related to variances in a few low-abundant TGs. That low-abundant TGs accounted for the most variability in lycopene suggests there is something unique about TGs with 40–46 carbons and few unsaturated bonds with respect to circulating lycopene. Interestingly, a robust sPLS model was established for lycopene, but not the other carotenoids. Variances in circulating retinol and 25-hydroxyvitamin D₃ were not explained by lipids. This was not

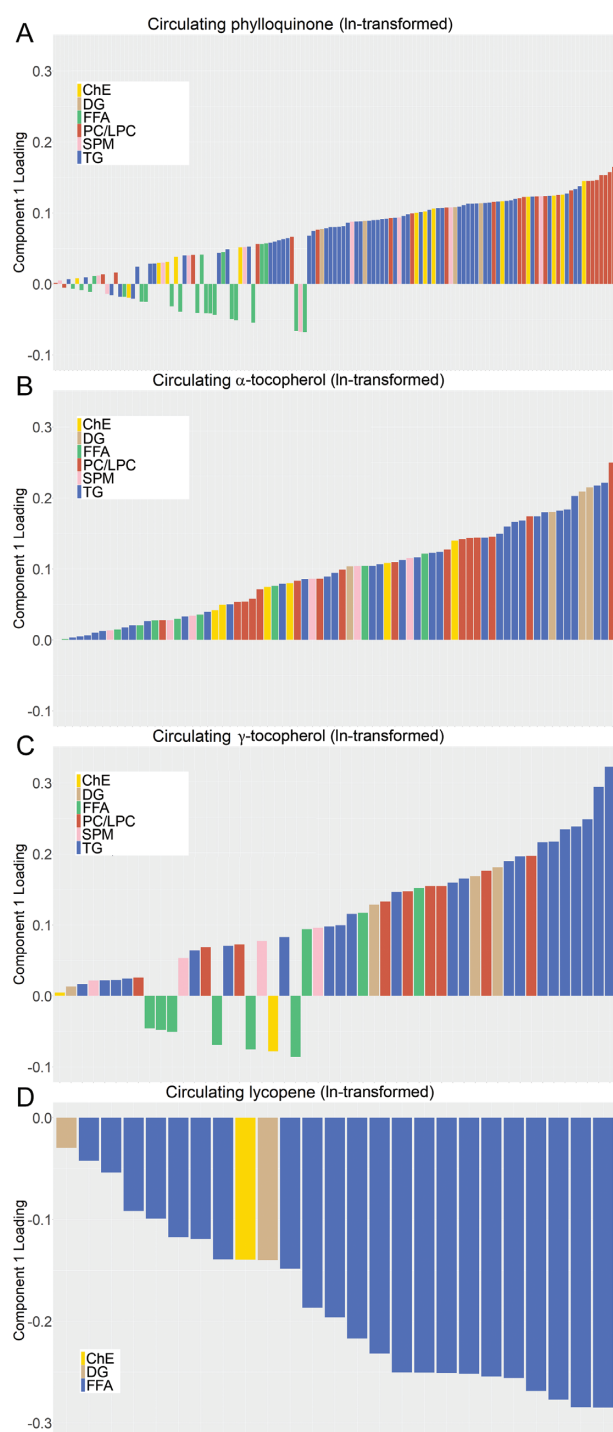


FIGURE 1 Plasma lipid loading plots of component 1 from cross-sectional sparse partial least-squares regression models for circulating concentrations of (A) phyloquinone (131 lipids), (B) α -tocopherol (75 lipids), (C) γ -tocopherol (50 lipids), and (D) lycopene (25 lipids). ChE, cholesterol ester; DG, diglyceride; FFA, free fatty acid; LPC, lysophosphatidylcholine; PC, phosphatidylcholine; SPM, sphingomyelin; TG, triglyceride.

unexpected given that both fat-soluble vitamins are transported on specific binding proteins (7, 8).

Total lipids and the relative abundance of lipids according to the number of carbons and unsaturated bonds for these participants are consistent with previous reports (11) and micronutrient concentrations were within expected ranges for this population (13, 30). However, there are several limitations of this study. First, this was a cross-sectional analysis so it cannot be determined whether lipids influence micronutrients or vice versa. Second, the cohort was a small sample of healthy overweight and obese males and there was no control group of normal-weight males, which limits generalizability. Finally, for this pilot study, the cross-validation procedure was conducted in subsets of the same cohort rather than in separate training and validation data sets, so replication in larger and more diverse cohorts is needed.

This was a study to investigate the associations of lipids, categorized by lipid class, the number of carbons, and the number of unsaturated bonds, with circulating concentrations of fat-soluble vitamins and carotenoids. Except for lycopene, refinement of the lipidome beyond class did not further characterize the interindividual variability in these micronutrients. However, based on these analyses, future investigations of the role of PC/LPC in vitamin E and K metabolism are merited.

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Online Supplementary Material

Supplemental Table 1: Component 1 Loadings for circulating phylloquinone (nM)

#	Lipid	Component 1 Loading
1	C16_0_LPC	0.165
2	C36_1_PC	0.157
3	C36_3_PC	0.153
4	C34_3_PC	0.153
5	C34_2_PC	0.146
6	C34_1_PC	0.145
7	C36_2_PC	0.145
8	C18_3_ChE	0.145
9	C52_2_TG	0.138
10	C52_1_TG	0.134
11	C18_0_LPC	0.132
12	C50_2_TG	0.127
13	C14_1_ChE	0.126
14	C38_3_PC	0.125
15	C16_2_ChE	0.124
16	C50_3_TG	0.124
17	C32_1_PC	0.124
18	C24_0_SPM	0.123
19	C36_0_PC	0.123
20	C50_1_TG	0.123
21	C18_4_ChE	0.123
22	C34_0_PC	0.122
23	C18_1_LPC	0.121
24	C48_2_TG	0.120
25	C50_4_TG	0.118
26	C52_6_TG	0.117
27	C18_0_Che	0.117
28	C50_0_TG	0.116
29	C16_1_LPC	0.116
30	C52_5_TG	0.114
31	C56_4_TG	0.114
32	C36_1_DG	0.114
33	C54_2_TG	0.113
34	C48_1_TG	0.113
35	C50_5_TG	0.113
36	C54_3_TG	0.111
37	C52_3_TG	0.109
38	C36_2_DG	0.108
39	C22_0_SPM	0.108
40	C38_7_PC	0.108
41	C48_3_TG	0.107
42	C56_3_TG	0.107

#	Lipid	Component 1 Loading
43	C14_0_ChE	0.106
44	C52_7_TG	0.104
45	C16_1_ChE	0.102
46	C48_0_TG	0.101
47	C18_1_ChE	0.100
48	C32_2_PC	0.099
49	C56_2_TG	0.098
50	C54_7_TG	0.095
51	C16_1_SPM	0.094
52	C46_1_TG	0.093
53	C34_4_PC	0.093
54	C56_5_TG	0.092
55	C54_1_TG	0.091
56	C46_0_TG	0.090
57	C48_4_TG	0.090
58	C52_4_TG	0.089
59	C36_3_DG	0.089
60	C50_6_TG	0.088
61	C54_4_TG	0.088
62	C14_0_SPM	0.087
63	C54_6_TG	0.086
64	C46_2_TG	0.082
65	C54_5_TG	0.081
66	C40_2_TG	0.080
67	C46_3_TG	0.080
68	C58_6_TG	0.078
69	C36_4_DG	0.077
70	C14_0_LPC	0.076
71	C40_1_TG	0.074
72	C44_0_TG	0.068
73	C36_6_PC	0.066
74	C44_1_TG	0.064
75	C44_2_TG	0.063
76	C56_6_TG	0.062
77	C56_9_TG	0.060
78	C42_2_TG	0.058
79	C20_0_FA	0.057
80	C22_0_FA	0.056
81	C18_2_LPC	0.056
82	C56_10_TG	0.052
83	C22_1_SPM	0.052
84	C18_2_ChE	0.052
85	C42_1_TG	0.049
86	C24_0_FA	0.045
87	C42_0_TG	0.044
88	C20_3_FA	0.041
89	C32_0_PC	0.041
90	C16_0_SPM	0.040

#	Lipid	Component 1 Loading
91	C40_0_TG	0.040
92	C20_3_ChE	0.038
93	C16_0_ChE	0.031
94	C14_1_SPM	0.030
95	C20_5_ChE	0.030
96	C58_7_TG	0.029
97	C56_7_TG	0.028
98	C56_8_TG	0.024
99	C36_5_PC	0.016
100	C36_4_PC	0.014
101	C24_2_SPM	0.012
102	C18_0_FA	0.011
103	C58_10_TG	0.009
104	C22_6_ChE	0.008
105	C58_11_TG	0.006
106	C24_1_SPM	0.005
107	C38_4_PC	0.001
108	C38_6_PC	-0.005
109	C20_4_FA	-0.007
110	C20_5_FA	-0.009
111	C24_1_FA	-0.011
112	C18_1_SPM	-0.014
113	C58_9_TG	-0.016
114	C58_8_TG	-0.018
115	C22_6_FA	-0.018
116	C20_4_ChE	-0.020
117	C58_12_TG	-0.021
118	C22_5_FA	-0.025
119	C16_0_FA	-0.025
120	C18_3_FA	-0.032
121	C15_0_FA	-0.039
122	C20_2_FA	-0.041
123	C22_4_FA	-0.041
124	C12_0_FA	-0.042
125	C14_0_FA	-0.044
126	C17_1_FA	-0.050
127	C18_1_FA	-0.051
128	C18_2_FA	-0.055
129	C16_1_FA	-0.066
130	C18_0_SPM	-0.067
131	C14_1_FA	-0.068

Supplemental Table 2: Component 1 Loadings for circulating α -tocopherol (μM)

#	Lipid	Component 1 Loading
1	C34_0_PC	0.250
2	C54_3_TG	0.221
3	C54_4_TG	0.217
4	C36_3_DG	0.215
5	C36_2_DG	0.209
6	C52_3_TG	0.203
7	C56_4_TG	0.184
8	C52_4_TG	0.182
9	C36_4_DG	0.180
10	C52_2_TG	0.180
11	C54_2_TG	0.174
12	C34_1_PC	0.174
13	C56_5_TG	0.168
14	C56_3_TG	0.166
15	C54_5_TG	0.160
16	C52_5_TG	0.150
17	C18_0_LPC	0.145
18	C56_2_TG	0.144
19	C16_0_LPC	0.144
20	C36_0_PC	0.144
21	C18_1_LPC	0.142
22	C18_0_Che	0.140
23	C34_2_PC	0.127
24	C54_6_TG	0.124
25	C50_4_TG	0.123
26	C18_0_FA	0.121
27	C54_1_TG	0.116
28	C22_0_SPM	0.115
29	C58_6_TG	0.113
30	C36_2_PC	0.110
31	C18_1_ChE	0.108
32	C50_3_TG	0.107
33	C56_6_TG	0.104
34	C20_2_FA	0.104
35	C16_0_SPM	0.104
36	C36_1_DG	0.104
37	C34_3_PC	0.099
38	C52_1_TG	0.094
39	C50_5_TG	0.089
40	C38_4_PC	0.086
41	C24_0_SPM	0.086
42	C52_6_TG	0.086
43	C36_4_PC	0.083
44	C18_3_ChE	0.080
45	C54_7_TG	0.079
46	C20_4_FA	0.076
47	C16_2_ChE	0.075

#	Lipid	Component 1 Loading
48	C34_4_PC	0.071
49	C36_1_PC	0.058
50	C32_0_PC	0.054
51	C36_3_PC	0.054
52	C48_3_TG	0.050
53	C20_4_ChE	0.049
54	C16_0_ChE	0.042
55	C52_7_TG	0.040
56	C24_1_FA	0.036
57	C16_1_SPM	0.034
58	C48_4_TG	0.033
59	C22_4_FA	0.030
60	C22_1_SPM	0.028
61	C16_1_LPC	0.028
62	C22_0_FA	0.027
63	C50_2_TG	0.027
64	C20_3_FA	0.021
65	C48_2_TG	0.021
66	C56_7_TG	0.018
67	C18_2_FA	0.015
68	C14_0_SPM	0.013
69	C50_6_TG	0.013
70	C40_2_TG	0.010
71	C58_7_TG	0.006
72	C46_3_TG	0.005
73	C56_9_TG	0.003
74	C18_1_FA	0.001
75	C14_1_ChE	0.000
76	C12_0_FA	0.000
77	C14_0_FA	0.000
78	C14_1_FA	0.000
79	C15_0_FA	0.000
80	C16_0_FA	0.000
81	C16_1_FA	0.000
82	C17_1_FA	0.000
83	C18_3_FA	0.000
84	C20_0_FA	0.000
85	C20_5_FA	0.000
86	C22_5_FA	0.000
87	C22_6_FA	0.000
88	C24_0_FA	0.000
89	C14_0_ChE	0.000
90	C16_1_ChE	0.000
91	C18_2_ChE	0.000
92	C18_4_ChE	0.000
93	C20_3_ChE	0.000
94	C20_5_ChE	0.000
95	C22_6_ChE	0.000

#	Lipid	Component 1 Loading
96	C14_0_LPC	0.000
97	C18_2_LPC	0.000
98	C14_1_SPM	0.000
99	C18_0_SPM	0.000
100	C18_1_SPM	0.000
101	C24_1_SPM	0.000
102	C24_2_SPM	0.000
103	C32_1_PC	0.000
104	C32_2_PC	0.000
105	C36_5_PC	0.000
106	C36_6_PC	0.000
107	C38_3_PC	0.000
108	C38_6_PC	0.000
109	C38_7_PC	0.000
110	C40_0_TG	0.000
111	C40_1_TG	0.000
112	C42_0_TG	0.000
113	C42_1_TG	0.000
114	C42_2_TG	0.000
115	C44_0_TG	0.000
116	C44_1_TG	0.000
117	C44_2_TG	0.000
118	C46_0_TG	0.000
119	C46_1_TG	0.000
120	C46_2_TG	0.000
121	C48_0_TG	0.000
122	C48_1_TG	0.000
123	C50_0_TG	0.000
124	C50_1_TG	0.000
125	C56_10_TG	0.000
126	C56_8_TG	0.000
127	C58_10_TG	0.000
128	C58_11_TG	0.000
129	C58_12_TG	0.000
130	C58_8_TG	0.000
131	C58_9_TG	0.000

Supplemental Table 3: Component 1 Loadings for circulating γ -tocopherol (μM)

#	Lipid	Component 1 Loading
1	C54_4_TG	0.323
2	C54_5_TG	0.295
3	C54_3_TG	0.249
4	C56_3_TG	0.239
5	C56_4_TG	0.234
6	C52_4_TG	0.218
7	C54_6_TG	0.216
8	C34_0_PC	0.197
9	C56_2_TG	0.196
10	C56_5_TG	0.190
11	C36_3_DG	0.181
12	C34_2_PC	0.176
13	C36_4_DG	0.169
14	C54_2_TG	0.165
15	C52_3_TG	0.160
16	C36_2_PC	0.155
17	C18_1_LPC	0.155
18	C22_0_FA	0.152
19	C38_7_PC	0.148
20	C56_6_TG	0.146
21	C18_2_LPC	0.133
22	C36_2_DG	0.129
23	C24_1_FA	0.117
24	C52_5_TG	0.116
25	C58_6_TG	0.100
26	C54_1_TG	0.098
27	C22_1_SPM	0.096
28	C24_0_FA	0.094
29	C56_10_TG	0.083
30	C22_0_SPM	0.078
31	C18_0_LPC	0.073
32	C54_7_TG	0.070
33	C36_0_PC	0.069
34	C52_2_TG	0.064
35	C16_1_SPM	0.053
36	C16_0_LPC	0.026
37	C50_4_TG	0.024
38	C56_7_TG	0.022
39	C56_9_TG	0.022
40	C24_0_SPM	0.022
41	C58_7_TG	0.017
42	C36_1_DG	0.013
43	C18_1_ChE	0.005
44	C16_0_FA	0.000
45	C18_0_FA	0.000
46	C18_1_FA	0.000
47	C18_2_FA	0.000

#	Lipid	Component 1 Loading
48	C18_3_FA	0.000
49	C20_0_FA	0.000
50	C20_2_FA	0.000
51	C20_3_FA	0.000
52	C20_4_FA	0.000
53	C20_5_FA	0.000
54	C22_4_FA	0.000
55	C22_5_FA	0.000
56	C22_6_FA	0.000
57	C14_0_ChE	0.000
58	C14_1_ChE	0.000
59	C16_0_ChE	0.000
60	C16_1_ChE	0.000
61	C16_2_ChE	0.000
62	C18_0_Che	0.000
63	C18_2_ChE	0.000
64	C18_3_ChE	0.000
65	C18_4_ChE	0.000
66	C20_4_ChE	0.000
67	C20_5_ChE	0.000
68	C22_6_ChE	0.000
69	C14_0_LPC	0.000
70	C16_1_LPC	0.000
71	C14_0_SPM	0.000
72	C14_1_SPM	0.000
73	C16_0_SPM	0.000
74	C18_0_SPM	0.000
75	C18_1_SPM	0.000
76	C24_1_SPM	0.000
77	C24_2_SPM	0.000
78	C32_0_PC	0.000
79	C32_1_PC	0.000
80	C32_2_PC	0.000
81	C34_1_PC	0.000
82	C34_3_PC	0.000
83	C34_4_PC	0.000
84	C36_1_PC	0.000
85	C36_3_PC	0.000
86	C36_4_PC	0.000
87	C36_5_PC	0.000
88	C36_6_PC	0.000
89	C38_3_PC	0.000
90	C38_4_PC	0.000
91	C38_6_PC	0.000
92	C40_0_TG	0.000
93	C40_1_TG	0.000
94	C40_2_TG	0.000
95	C42_0_TG	0.000

#	Lipid	Component 1 Loading
96	C42_1_TG	0.000
97	C42_2_TG	0.000
98	C44_0_TG	0.000
99	C44_1_TG	0.000
100	C44_2_TG	0.000
101	C46_0_TG	0.000
102	C46_1_TG	0.000
103	C46_2_TG	0.000
104	C46_3_TG	0.000
105	C48_0_TG	0.000
106	C48_1_TG	0.000
107	C48_2_TG	0.000
108	C48_3_TG	0.000
109	C48_4_TG	0.000
110	C50_0_TG	0.000
111	C50_1_TG	0.000
112	C50_2_TG	0.000
113	C50_3_TG	0.000
114	C50_5_TG	0.000
115	C50_6_TG	0.000
116	C52_1_TG	0.000
117	C52_6_TG	0.000
118	C52_7_TG	0.000
119	C56_8_TG	0.000
120	C58_10_TG	0.000
121	C58_11_TG	0.000
122	C58_12_TG	0.000
123	C58_8_TG	0.000
124	C58_9_TG	0.000
125	C12_0_FA	-0.046
126	C14_0_FA	-0.048
127	C16_1_FA	-0.051
128	C14_1_FA	-0.069
129	C15_0_FA	-0.075
130	C20_3_ChE	-0.078
131	C17_1_FA	-0.086

Supplemental Table 4: Component 1 Loadings for circulating lycopene (μM)

#	Lipid	Component 1 Loading
1	C14_0_FA	0.000
2	C14_1_FA	0.000
3	C15_0_FA	0.000
4	C16_0_FA	0.000
5	C16_1_FA	0.000
6	C17_1_FA	0.000
7	C18_0_FA	0.000
8	C18_1_FA	0.000
9	C18_2_FA	0.000
10	C18_3_FA	0.000
11	C20_0_FA	0.000
12	C20_2_FA	0.000
13	C20_3_FA	0.000
14	C20_4_FA	0.000
15	C20_5_FA	0.000
16	C22_0_FA	0.000
17	C22_4_FA	0.000
18	C22_5_FA	0.000
19	C22_6_FA	0.000
20	C24_0_FA	0.000
21	C24_1_FA	0.000
22	C14_0_ChE	0.000
23	C14_1_ChE	0.000
24	C16_0_ChE	0.000
25	C16_1_ChE	0.000
26	C16_2_ChE	0.000
27	C18_0_Che	0.000
28	C18_1_ChE	0.000
29	C18_2_ChE	0.000
30	C18_3_ChE	0.000
31	C18_4_ChE	0.000
32	C20_3_ChE	0.000
33	C20_4_ChE	0.000
34	C20_5_ChE	0.000
35	C22_6_ChE	0.000
36	C16_0_LPC	0.000
37	C16_1_LPC	0.000
38	C18_0_LPC	0.000
39	C18_1_LPC	0.000
40	C18_2_LPC	0.000
41	C14_0_SPM	0.000
42	C14_1_SPM	0.000
43	C16_0_SPM	0.000
44	C16_1_SPM	0.000
45	C18_0_SPM	0.000
46	C18_1_SPM	0.000
47	C22_0_SPM	0.000

#	Lipid	Component 1 Loading
48	C22_1_SPM	0.000
49	C24_0_SPM	0.000
50	C24_1_SPM	0.000
51	C24_2_SPM	0.000
52	C32_0_PC	0.000
53	C32_1_PC	0.000
54	C34_0_PC	0.000
55	C34_1_PC	0.000
56	C34_2_PC	0.000
57	C34_3_PC	0.000
58	C34_4_PC	0.000
59	C36_0_PC	0.000
60	C36_1_PC	0.000
61	C36_2_PC	0.000
62	C36_3_PC	0.000
63	C36_4_PC	0.000
64	C36_5_PC	0.000
65	C36_6_PC	0.000
66	C38_3_PC	0.000
67	C38_4_PC	0.000
68	C38_6_PC	0.000
69	C38_7_PC	0.000
70	C36_1_DG	0.000
71	C36_2_DG	0.000
72	C36_3_DG	0.000
73	C36_4_DG	0.000
74	C50_1_TG	0.000
75	C50_2_TG	0.000
76	C50_3_TG	0.000
77	C50_4_TG	0.000
78	C52_1_TG	0.000
79	C52_2_TG	0.000
80	C52_3_TG	0.000
81	C52_4_TG	0.000
82	C52_5_TG	0.000
83	C52_6_TG	0.000
84	C54_1_TG	0.000
85	C54_2_TG	0.000
86	C54_3_TG	0.000
87	C54_4_TG	0.000
88	C54_5_TG	0.000
89	C54_6_TG	0.000
90	C54_7_TG	0.000
91	C56_10_TG	0.000
92	C56_2_TG	0.000
93	C56_3_TG	0.000
94	C56_4_TG	0.000
95	C56_5_TG	0.000

#	Lipid	Component 1 Loading
96	C56_6_TG	0.000
97	C56_7_TG	0.000
98	C56_8_TG	0.000
99	C56_9_TG	0.000
100	C58_10_TG	0.000
101	C58_11_TG	0.000
102	C58_12_TG	0.000
103	C58_6_TG	0.000
104	C58_7_TG	0.000
105	C58_8_TG	0.000
106	C58_9_TG	0.000
107	C32_2_PC	-0.029
108	C50_0_TG	-0.042
109	C50_5_TG	-0.054
110	C48_0_TG	-0.092
111	C40_2_TG	-0.099
112	C48_2_TG	-0.118
113	C48_1_TG	-0.119
114	C52_7_TG	-0.139
115	C12_0_FA	-0.140
116	C14_0_LPC	-0.140
117	C48_3_TG	-0.148
118	C46_0_TG	-0.187
119	C48_4_TG	-0.196
120	C46_3_TG	-0.217
121	C50_6_TG	-0.232
122	C46_1_TG	-0.250
123	C42_2_TG	-0.250
124	C40_1_TG	-0.251
125	C46_2_TG	-0.252
126	C44_2_TG	-0.254
127	C40_0_TG	-0.256
128	C42_1_TG	-0.269
129	C44_1_TG	-0.277
130	C44_0_TG	-0.285
131	C42_0_TG	-0.285