

**Table S1: Results from ordinary least squares regression examining the association of dietary and non-dietary factors with food-based metabolites**

	<b>Hippuric acid</b>	<b>Carnitine</b>
<b>Variable</b>	<b>b (95% CI)</b>	<b>b (95% CI)</b>
<b>Age (years)</b>	0.01 (-0.01, 0.03)	0.00 (-0.00, 0.00)
<b>Gestational age (weeks)</b>	0.01 (-0.01, 0.03)	-0.01** (-0.02, -0.01)
<b>Parity</b>	0.03 (-0.05, 0.11)	0.00 (-0.01, 0.02)
<b>Gestational diabetes (GDM)</b>	0.06 (-0.10, 0.23)	0.02 (-0.02, 0.05)
<b>Pre-pregnancy BMI (kg/m<sup>2</sup>)</b>	-0.01 (-0.03, 0.00)	0.00 (-0.00, 0.00)
<b>Smoking history (ever vs. never smoked)</b>	-0.13 (-0.29, 0.03)	0.06* (0.02, 0.10)
<b>Physical activity (low vs. high)</b>	0.02 (-0.11, 0.15)	-0.01 (-0.04, 0.02)
<b>Social disadvantage index</b>	-0.02 (-0.08, 0.04)	0.00 (-0.01, 0.01)
<b>Fiber intake (g/day)</b>	0.01 (-0.00, 0.02)	0.00 (-0.00, 0.00)
<b>Energy intake (kcal)</b>	0.00* (-0.00, -0.00)	0.00 (-0.00, 0.00)
<b>FFQ before blood draw vs. FFQ at the same time as blood draw</b>	0.10 (-0.07, 0.26)	0.01 (-0.03, 0.05)
<b>FFQ after blood draw vs. FFQ at the same time as blood draw</b>	0.08 (-0.18, 0.34)	0.06 (-0.07, 0.19)
<b>Ethnicity (White European vs. South Asian)</b>	0.04 (-0.18, 0.26)	-0.02 (-0.07, 0.04)
<b>Fruits and vegetables (servings/day)</b>	0.22* (0.08, 0.36)	
<b>Tea (servings/day)</b>	0.01 (-0.01, 0.04)	
<b>Coffee (servings/day)</b>	0.02 (-0.00, 0.04)	
<b>Red meat (servings/day)</b>		0.00 (-0.00, 0.01)

\*p ≤ 0.01, \*\*p ≤ 0.001

**Table S2: Results of model fitting analyses examining the association of dietary and non-dietary factors with food metabolites**

	<b>-2 Log L</b>	<b>BIC</b>	<b>AIC</b>	<b>S<sub>b</sub><sup>2</sup></b>	<b>S<sub>w</sub><sup>2</sup></b>
<b>Proline Betaine</b>					
<b>Model 1</b>	—	—	—	0.14	0.13
<b>Model 2</b>	2100.0	2101.4	2104.0	0.02	0.11
<b>Model 3</b>	1779.0	1780.4	1783.0	0.20	0.12
<b>Hippuric acid</b>					
<b>Model 1</b>	—	—	—	0.00	0.03
<b>Model 2</b>	1394.7	1395.3	1396.7	0.00	0.03
<b>Model 3</b>	1182.1	1182.8	1184.1	0.00	0.03
<b>3-methylhistidine</b>					
<b>Model 1</b>	—	—	—	0.07	0.01
<b>Model 2</b>	515.3	518.8	525.3	0.02	0.01
<b>Model 3</b>	496.0	497.4	500.0	0.02	0.01
<b>Carnitine</b>					
<b>Model 1</b>	—	—	—	0.00	0.00
<b>Model 2</b>	-387.0	-385.6	-383.0	0.00	0.00
<b>Model 3</b>	-298.6	-297.9	-296.6	0.00	0.00
<b>Tryptophan betaine</b>					
<b>Model 1</b>	—	—	—	0.03	0.00
<b>Model 2</b>	-588.1	-586.7	-584.1	0.02	0.00
<b>Model 3</b>	-489.4	-488.0	-485.4	0.02	0.00
<b>TMAO</b>					
<b>Model 1</b>	—	—	—	0.07	0.04
<b>Model 2</b>	1421.5	1422.9	1425.5	0.02	0.03
<b>Model 3</b>	1211.1	1211.8	1213.1	0.00	0.04

AIC = Akaike Information Criterion, BIC = Bayesian Information Criterion, -2 Log L = -2 log likelihood, S<sub>b</sub><sup>2</sup> = Sum of square between, S<sub>w</sub><sup>2</sup> = Sum of square within

Model 1: Unconditional (intercept only) model

Model 2: Random Intercept with Fixed Level-1 Factors (dietary factors)

Model 3: Random Intercept with Fixed Level-1 Factors (dietary and non-dietary factors)

**Table S3: Results from ordinary least squares regression examining the association of dietary and non-dietary factors with food-based metabolites in FAMILY cohort**

	<b>Proline betaine</b>	<b>Hippuric acid</b>	<b>3-Methyl histidine</b>	<b>Carnitine</b>	<b>Tryptophan betaine</b>	<b>TMAO</b>
<b>Variable</b>	<b>b (95% CI)</b>	<b>b (95% CI)</b>	<b>b (95% CI)</b>	<b>b (95% CI)</b>	<b>b (95% CI)</b>	<b>b (95% CI)</b>
<b>Age (years)</b>	0.06** (0.02, 0.10)	0.02 (-0.01, 0.04)	0.00 (-0.01, 0.02)	0.00 (-0.00, 0.01)	0.00 (-0.00, 0.00)	0.02 (-0.01, 0.04)
<b>Gestational age (weeks)</b>	0.02 (-0.02, 0.10)	0.00 (-0.02, 0.02)	0.01 (-0.01, 0.02)	-0.01*** (-0.02, 0.01)	0.00 (-0.00, 0.00)	0.01 (-0.01, 0.04)
<b>Parity</b>	-0.06 (-0.26, 0.15)	-0.02 (-0.11, 0.08)	0.00 (-0.07, 0.07)	-0.01 (-0.03, 0.01)	-0.01 (-0.02, 0.00)	-0.04 (-0.12, 0.05)
<b>Gestational diabetes (GDM)</b>	-0.15 (-0.65, 0.35)	0.01 (-0.23, 0.24)	0.06 (-0.07, 0.19)	0.03 (-0.02, 0.08)	-0.01 (-0.04, 0.02)	-0.07 (-0.29, 0.15)
<b>Pre-pregnancy BMI (kg/m<sup>2</sup>)</b>	-0.03* (-0.06, -0.00)	-0.01 (-0.03, 0.01)	-0.00 (-0.01, 0.00)	-0.00 (-0.01, 0.00)	0.00 (-0.00, 0.00)	-0.01 (-0.02, 0.01)
<b>Smoking history (ever vs. never smoked)</b>	-0.53** (-0.91, 0.14)	-0.10 (-0.26, 0.07)	0.03 (-0.07, 0.13)	0.06** (0.02, 0.10)	0.00 (-0.02, 0.00)	0.01 (-0.17, 0.19)
<b>Physical activity (low vs. high)</b>	0.06 (-0.31, 0.43)	-0.02 (-0.18, 0.14)	-0.05 (-0.15, 0.06)	-0.02 (-0.06, 0.03)	0.00 (-0.03, 0.02)	-0.02 (-0.21, 0.18)
<b>Social disadvantage index</b>	-0.03 (-0.22, 0.17)	-0.01 (-0.10, 0.07)	0.00 (-0.04, 0.05)	-0.00 (-0.02, 0.01)	0.00 (-0.01, 0.01)	0.04 (-0.03, 0.11)
<b>Fiber intake (g/day)</b>	-0.00 (-0.03, 0.02)	0.01 (-0.00, 0.02)	-0.00 (-0.01, 0.01)	0.00 (-0.00, 0.00)	0.00 (-0.00, 0.00)	0.00 (-0.01, 0.02)
<b>Energy intake (kcal)</b>	-0.00 (-0.00, 0.00)	-0.00* (-0.00, -0.00)	-0.00 (0.00, 0.00)	-0.00 (-0.00, 0.00)	0.00 (-0.00, 0.00)	-0.00 (-0.00, 0.00)
<b>FFQ before blood draw vs. FFQ at the same time as blood draw</b>	0.06 (-0.29, 0.40)	0.09 (-0.09, 0.26)	-0.06 (-0.16, 0.05)	0.02 (-0.02, 0.06)	-0.01 (-0.03, 0.02)	0.10 (-0.08, 0.28)
<b>FFQ after blood draw vs. FFQ at the same time as blood draw</b>	0.08 (-0.70, 0.86)	-0.02 (-0.46, 0.41)	0.10 (-0.30, 0.50)	0.05 (-0.10, 0.21)	-0.02 (-0.11, 0.06)	-0.14 (-0.72, 0.45)

<b>Citrus food (servings/day)</b>	0.40*** (0.24, 0.57)					
<b>Fruits and vegetables (servings/day)</b>		0.30*** (0.16, 0.45)				
<b>Tea (servings/day)</b>		0.00 (-0.02, 0.03)				
<b>Coffee (servings/day)</b>		0.01 (-0.01, 0.04)				
<b>Chicken (servings/day)</b>			0.03 (-0.00, 0.06)			
<b>Red meat (servings/day)</b>			0.01 (-0.03, 0.05)	0.02* (0.00, 0.04)		0.04 (-0.02, 0.10)
<b>Eggs (servings/day)</b>			0.03 (-0.00, 0.06)			0.03 (-0.04, 0.10)
<b>Nuts and legumes (servings/day)</b>			0.06* (0.00, 0.12)		0.02* (0.00, 0.04)	
<b>Canned fish (servings/day)</b>						0.02 (-0.01, 0.06)
<b>Fried fish (servings/day)</b>						0.02 (-0.02, 0.07)
<b>Seafood (servings/day)</b>						0.07** (0.02, 0.11)

\*p ≤ 0.05, \*\*p ≤ 0.01, \*\*\*p ≤ 0.001

**Table S4: Results from ordinary least squares regression examining the association of dietary and non-dietary factors with food-based metabolites in START cohort**

	<b>Proline betaine</b>	<b>Hippuric acid</b>	<b>3-Methyl histidine</b>	<b>Carnitine</b>	<b>Tryptophan betaine</b>	<b>TMAO</b>
<b>Variable</b>	<b>b (95% CI)</b>	<b>b (95% CI)</b>	<b>b (95% CI)</b>	<b>b (95% CI)</b>	<b>b (95% CI)</b>	<b>b (95% CI)</b>
<b>Age (years)</b>	-0.01 (-0.07, 0.04)	-0.01 (-0.04, 0.02)	0.00 (-0.01, 0.01)	-0.00 (-0.01, 0.00)	0.00 (-0.01, 0.01)	0.03 (-0.01, 0.06)
<b>Gestational age (weeks)</b>	0.02 (-0.08, 0.13)	0.05 (-0.01, 0.11)	-0.00 (-0.03, 0.02)	-0.02*** (-0.03, -0.01)	-0.01 (-0.02, 0.00)	0.02 (-0.03, 0.07)
<b>Parity</b>	-0.18 (-0.45, 0.10)	0.11 (-0.02, 0.24)	-0.03 (-0.09, 0.04)	0.03 (-0.00, 0.06)	-0.01 (-0.04, 0.03)	0.07 (-0.06, 0.21)
<b>Gestational diabetes (GDM)</b>	0.18 (-0.18, 0.55)	0.09 (-0.14, 0.32)	-0.02 (-0.11, 0.07)	0.01 (-0.03, 0.06)	0.04 (-0.00, 0.09)	0.05 (-0.18, 0.27)
<b>Pre-pregnancy BMI (kg/m<sup>2</sup>)</b>	0.01 (-0.04, 0.05)	-0.02 (-0.05, 0.00)	-0.01 (-0.02, 0.00)	0.00 (-0.00, 0.01)	0.00 (-0.00, 0.01)	-0.02 (-0.04, 0.01)
<b>Physical activity (low vs. high)</b>	-0.57* (-0.12, 0.03)	-0.05 (-0.27, 0.17)	-0.01 (-0.11, 0.09)	-0.00 (-0.05, 0.05)	0.03 (-0.02, 0.08)	-0.04 (-0.36, 0.28)
<b>Social disadvantage index</b>	-0.05 (-0.18, 0.09)	-0.03 (-0.10, 0.05)	-0.00 (-0.04, 0.03)	0.01 (-0.01, 0.02)	-0.00 (-0.02, 0.02)	-0.07 (-0.15, 0.01)
<b>Fiber intake (g/day)</b>	0.01 (-0.02, 0.04)	0.01 (-0.01, 0.03)	-0.00 (-0.01, 0.00)	-0.00 (-0.01, 0.00)	0.00** (0.00, 0.01)	-0.00 (-0.02, 0.01)
<b>Energy intake (kcal)</b>	-0.00 (-0.00, 0.00)	0.00* (0.00, 0.00)	0.00 (-0.00, 0.00)	0.00** (0.00, 0.00)	-0.00* (-0.00, -0.00)	0.00 (-0.00, 0.00)
<b>FFQ before blood draw vs. FFQ at the same time as blood draw</b>	-0.76 (-1.58, 0.07)	0.24 (-0.25, 0.72)	0.00 (-0.18, 0.19)	0.05 (-0.14, 0.05)	-0.01 (-0.08, 0.05)	0.02 (-0.37, 0.41)
<b>FFQ after blood draw vs. FFQ at the same time as blood draw</b>	0.94** (0.29, 1.59)	0.14 (-0.23, 0.50)	-0.07 (-0.28, 0.13)	0.06 (-0.14, 0.27)	-0.06 (-0.24, 0.12)	-0.19 (-0.64, 0.26)
<b>Citrus food (servings/day)</b>	0.18*** (0.17, 0.28)					

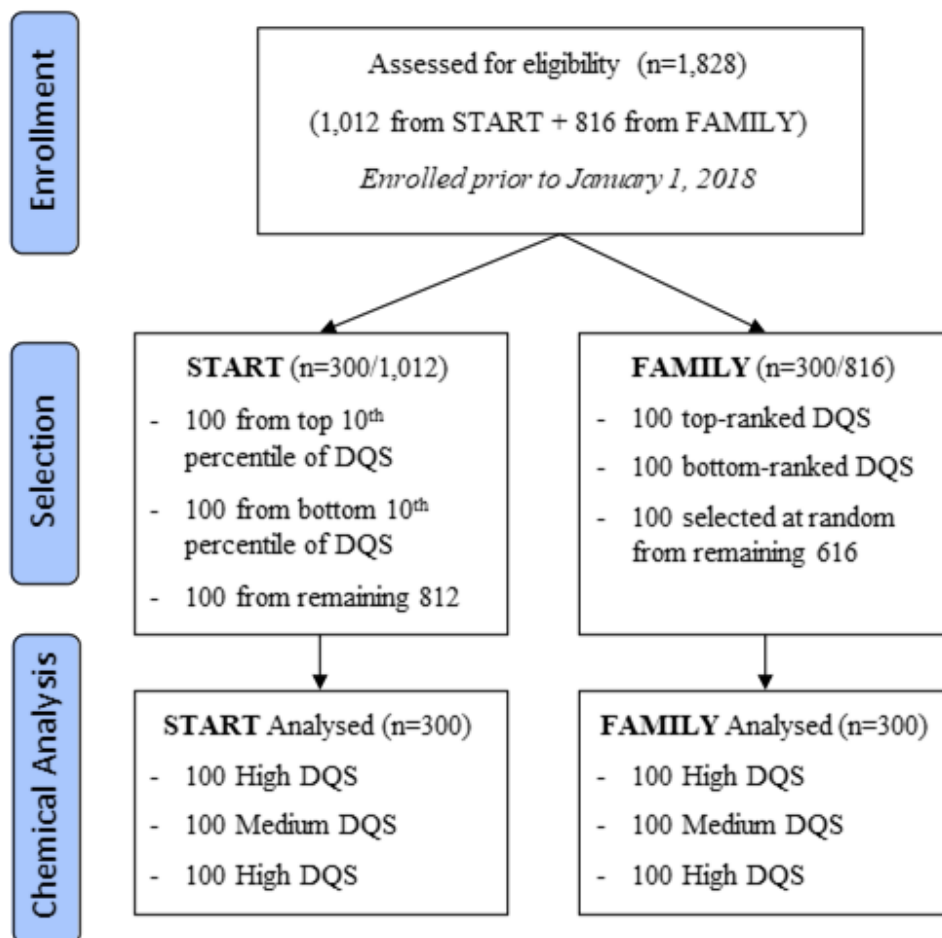
<b>Fruits and vegetables (servings/day)</b>		0.11 (-0.09, 0.30)				
<b>Tea (servings/day)</b>		0.03 (-0.02, 0.08)				
<b>Coffee (servings/day)</b>		0.02 (-0.04, 0.07)				
<b>Chicken (servings/day)</b>			-0.00 (-0.03, 0.03)			
<b>Red meat (servings/day)</b>			0.06** (0.02, 0.10)	-0.00 (-0.01, 0.01)		0.00 (-0.06, 0.06)
<b>Eggs (servings/day)</b>			-0.01 (-0.02, 0.01)			0.00 (-0.03, 0.07)
<b>Nuts and legumes (servings/day)</b>			-0.05 (-0.11, 0.01)		0.02 (-0.01, 0.05)	
<b>Canned fish (servings/day)</b>						-0.08 (-0.24, 0.07)
<b>Fried fish (servings/day)</b>						-0.00 (-0.08, 0.07)
<b>Seafood (servings/day)</b>						0.11* (0.01, 0.21)

\*p ≤ 0.05, \*\*p ≤ 0.01, \*\*\*p ≤ 0.001

**Table S5: Results from ordinary least squares regression examining the association of dietary and non-dietary factors with serum non-esterified fatty acid (NEFA) in FAMILY cohort**

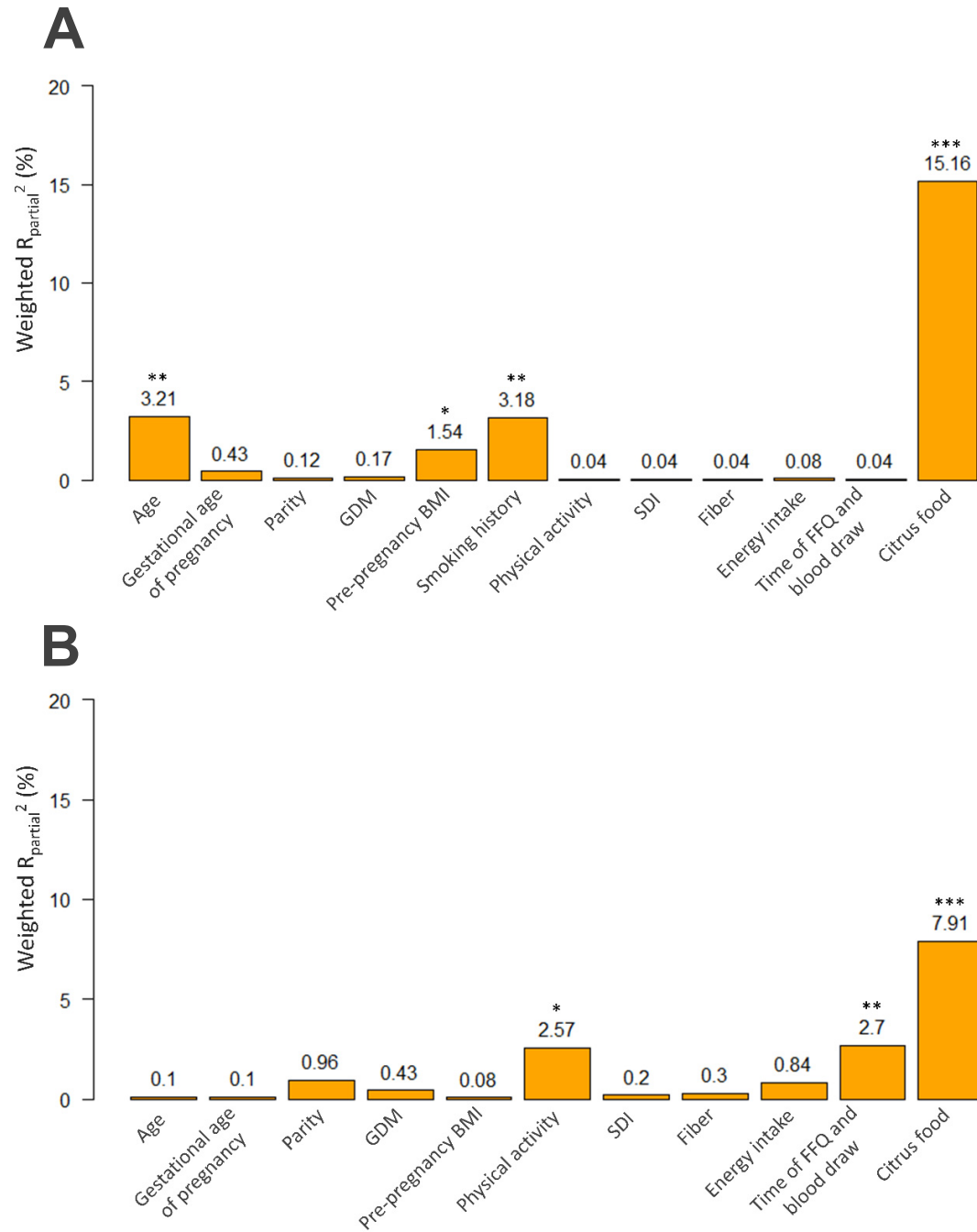
	<b>ω-3 PUFA</b>		
	<b>EPA</b>	<b>DHA</b>	<b>EPA + DHA</b>
<b>Variable</b>	<b>b (95% CI)</b>	<b>b (95% CI)</b>	<b>b (95% CI)</b>
<b>Age (years)</b>	-0.01 (-0.02, 0.00)	-2.92×10 <sup>-3</sup> (-0.01, 0.01)	-0.01 (-0.02, 0.00)
<b>Gestational age (weeks)</b>	-0.01 (-0.03, 0.00)	-0.01* (-0.02, -0.00)	-0.01* (-0.02, -0.00)
<b>Parity</b>	-0.01 (-0.07, 0.06)	-0.05* (-0.09, -0.00)	-0.03 (-0.08, 0.02)
<b>Gestational diabetes (GDM)</b>	-0.06 (-0.22, 0.10)	-0.06 (-0.17, 0.05)	-0.07 (-0.18, 0.05)
<b>Pre-pregnancy BMI (kg/m<sup>2</sup>)</b>	-3.34×10 <sup>-3</sup> (-0.01, 0.01)	-0.01** (-0.02, -0.00)	-0.01 (-0.01, 0.00)
<b>Smoking history (ever vs. never smoked)</b>	-0.01 (-0.13, 0.11)	-0.04 (-0.13, 0.04)	-0.03 (-0.12, 0.05)
<b>Physical activity (low vs. high)</b>	-0.03 (-0.18, 0.12)	-0.02 (-0.12, 0.08)	-0.02 (-0.13, 0.09)
<b>Social disadvantage index</b>	0.03 (-0.04, 0.10)	0.03 (-0.01, 0.07)	0.03 (-0.02, 0.08)
<b>Fiber intake (g/day)</b>	0.01 (-0.00, 0.01)	3.71×10 <sup>-3</sup> (-0.00, 0.01)	4.88×10 <sup>-3</sup> (-0.00, 0.01)
<b>Energy intake (kcal)</b>	-7.71×10 <sup>-5</sup> (-0.00, 0.00)	-3.76×10 <sup>-5</sup> (-0.00, 0.00)	-5.39×10 <sup>-5</sup> (-0.00, 0.00)
<b>FFQ before blood draw vs. FFQ at the same time as blood draw</b>	-0.01 (-0.15, 0.13)	0.04 (-0.05, 0.13)	0.02 (-0.08, 0.11)
<b>FFQ after blood draw vs. FFQ at the same time as blood draw</b>	0.06 (-0.26, 0.39)	0.25 (-0.04, 0.54)	0.17 (-0.11, 0.44)
<b>Fish (servings/day)</b>	0.02 (-0.01, 0.05)	0.04** (0.01, 0.06)	0.03* (0.01, 0.05)

\*p ≤ 0.05, \*\*p ≤ 0.01



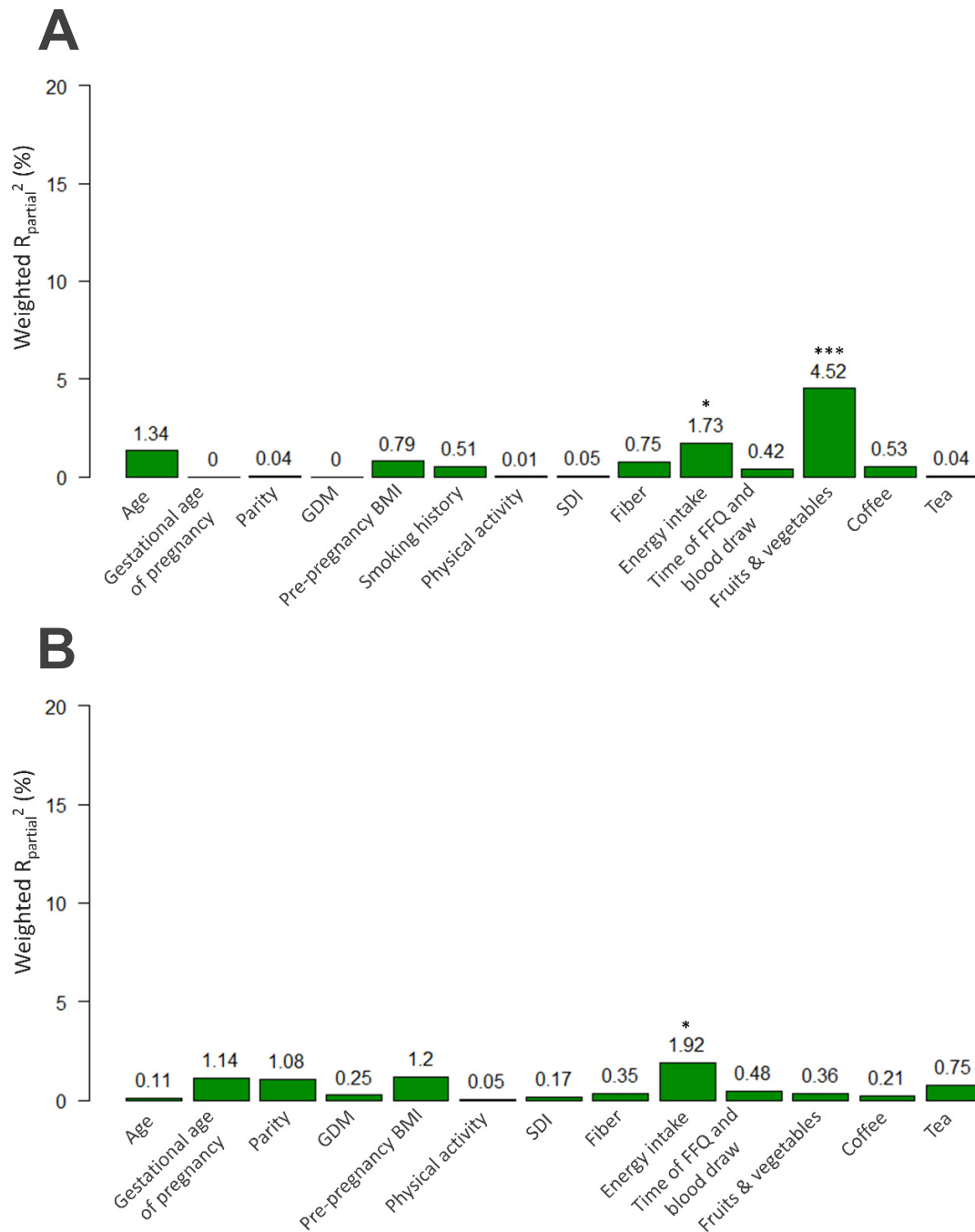
**Figure S1. Consort flow diagram outlining selection criteria used in a cross-sectional study involving participants from the FAMILY and START birth cohorts**





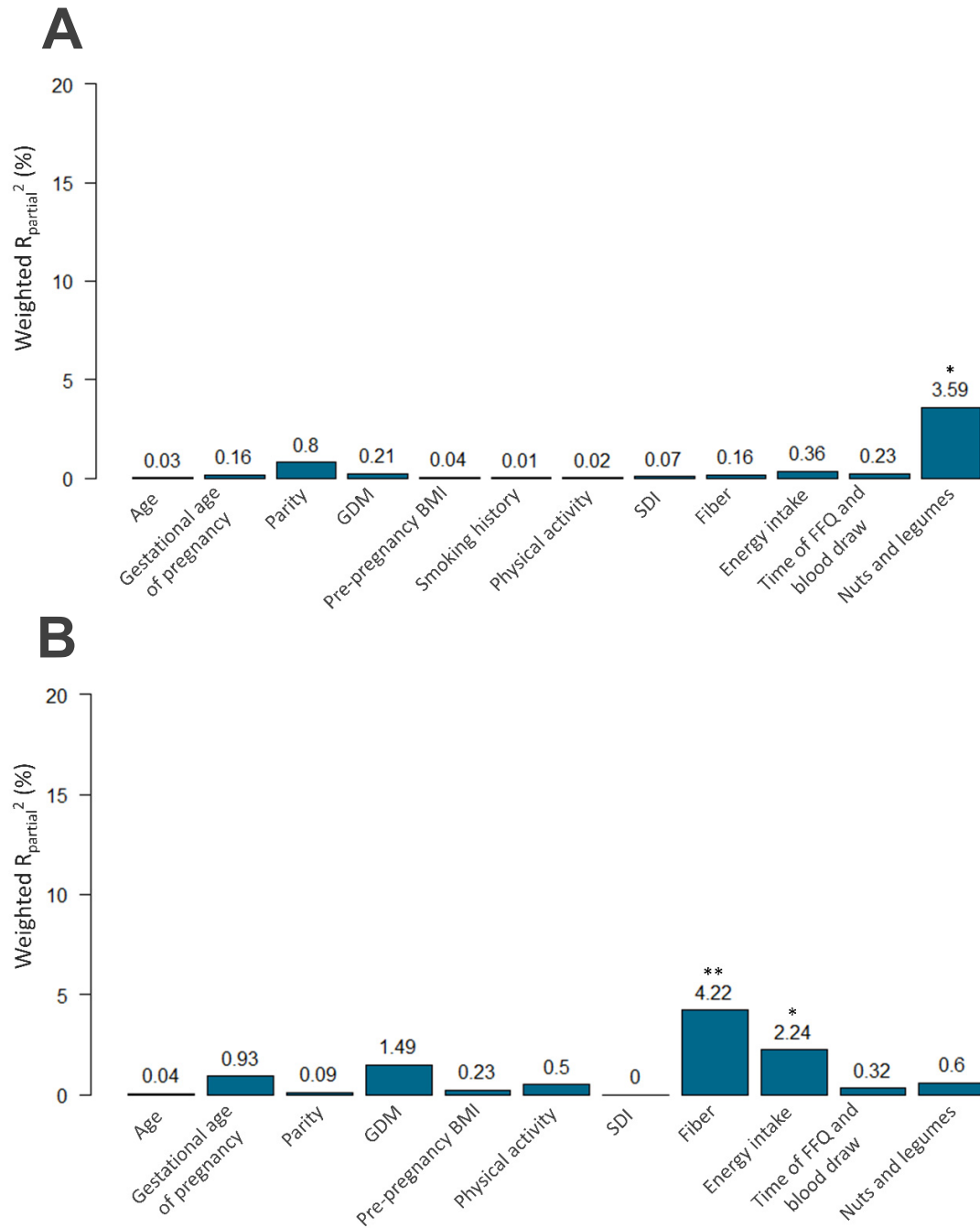
**Figure S2: Weighted  $R_{\text{partial}}^2$  for each factor showing the percentage of explained variability in Proline betaine in (A) FAMILY and (B) START cohort**

Statistical significance was based on hierarchical linear models. \*  $p \leq 0.05$ , \*\* $p \leq 0.01$ , \*\*\* $p \leq 0.001$



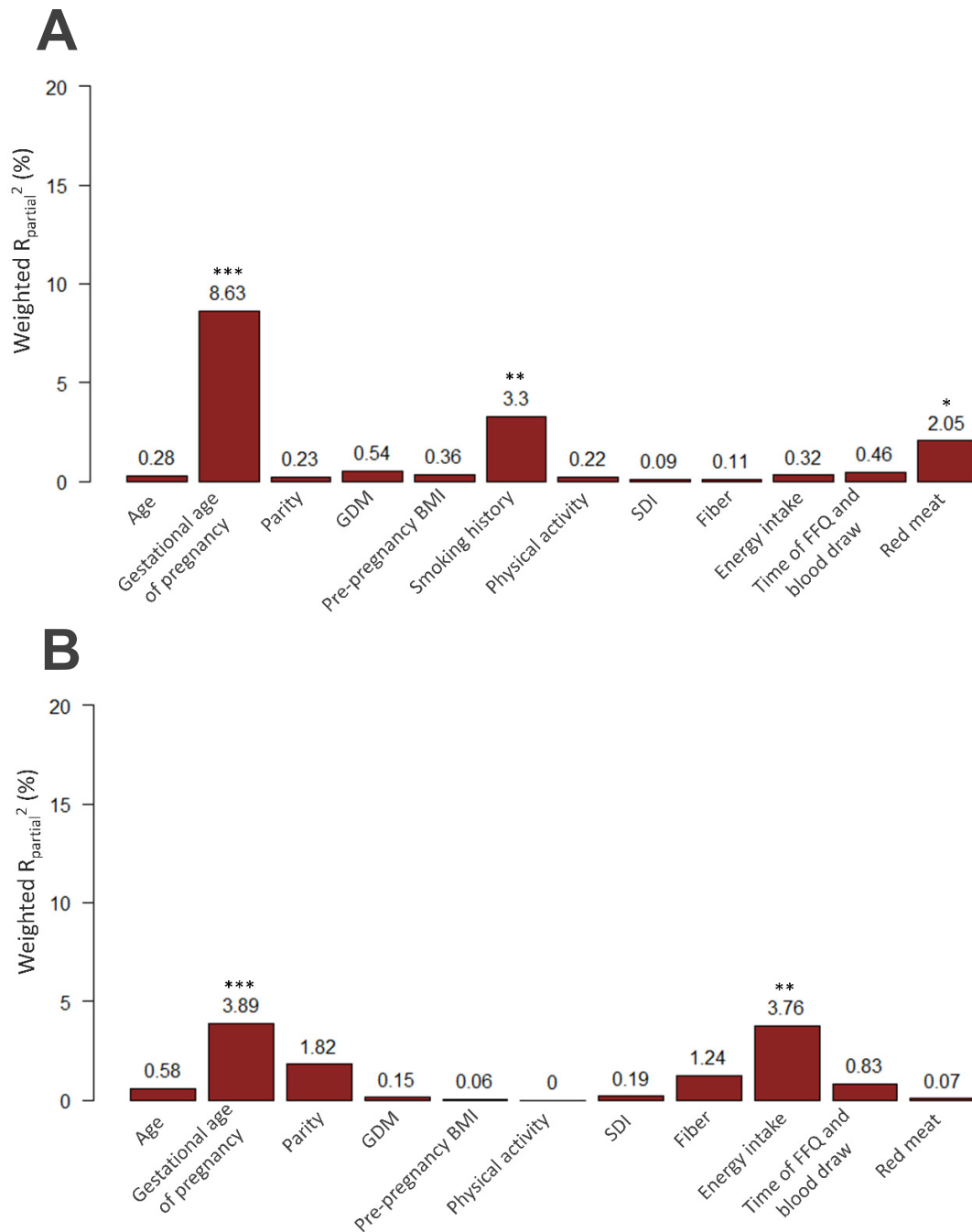
**Figure S3: Weighted  $R_{\text{partial}}^2$  for each factor showing the percentage of explained variability in Hippuric acid in (A) FAMILY and (B) START cohort**

Statistical significance was based on hierarchical linear models. \*  $p \leq 0.05$ , \*\*\* $p \leq 0.001$



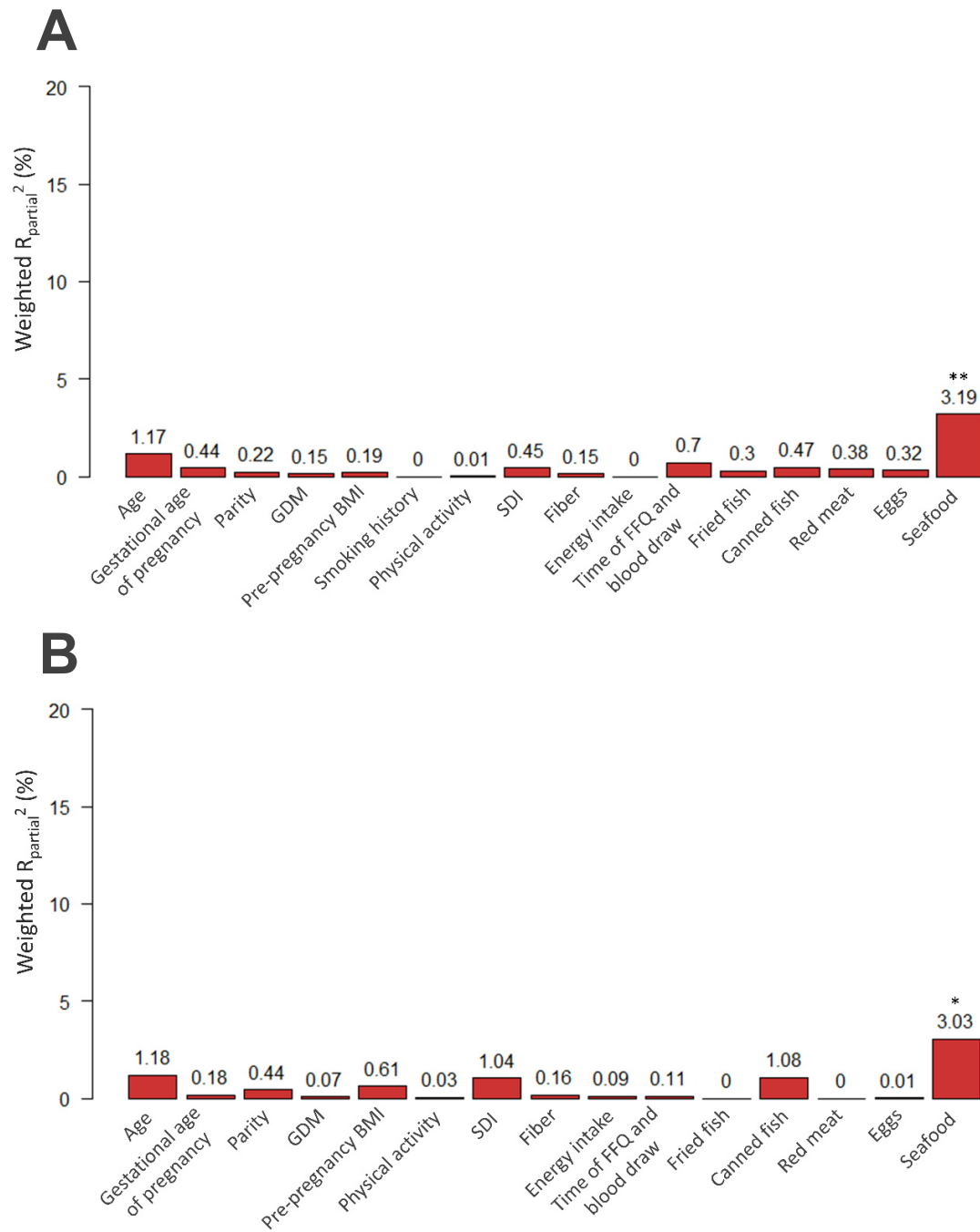
**Figure S4: Weighted  $R_{\text{partial}}^2$  for each factor showing the percentage of explained variability in Tryptophan betaine in (A) FAMILY and (B) START cohort**

Statistical significance was based on hierarchical linear models. \*  $p \leq 0.05$ , \*\* $p \leq 0.01$



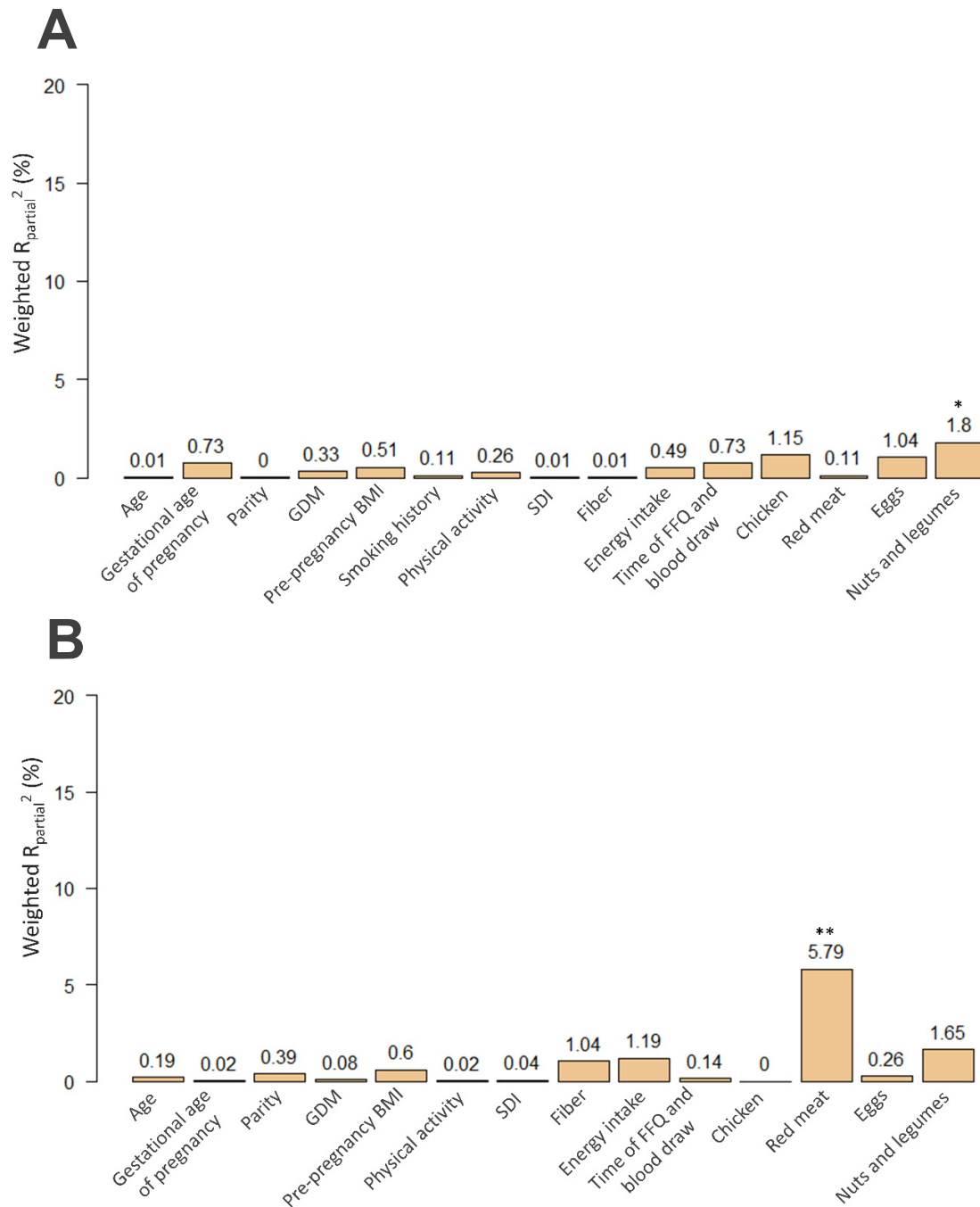
**Figure S5: Weighted  $R^2_{\text{partial}}$  for each factor showing the percentage of explained variability in Carnitine in (A) FAMILY and (B) START cohort**

Statistical significance was based on hierarchical linear models. \*  $p \leq 0.05$ , \*\* $p \leq 0.01$ , \*\*\* $p \leq 0.001$



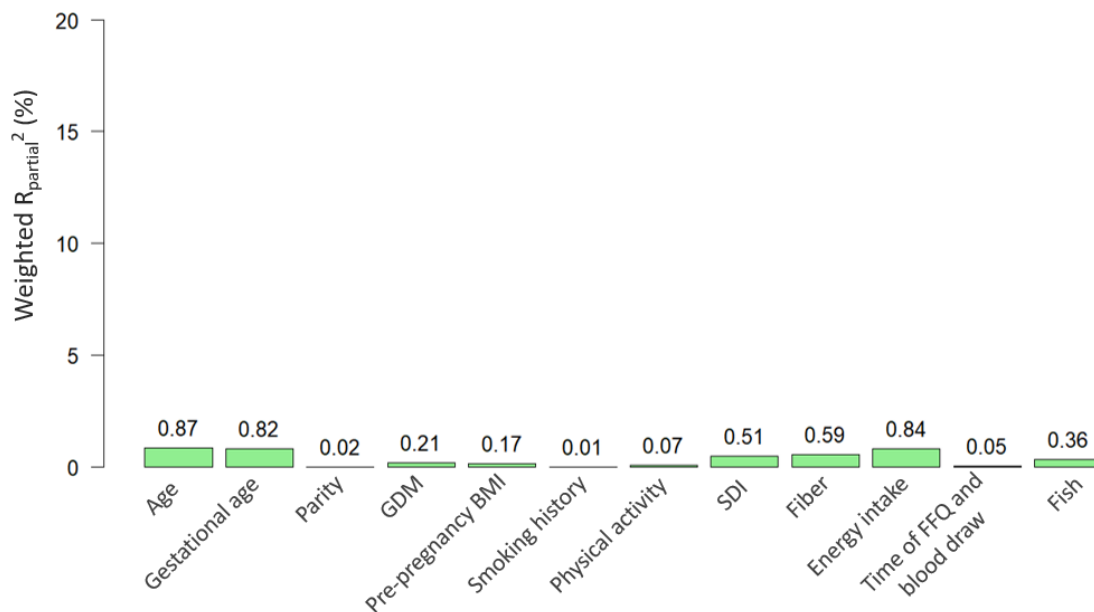
**Figure S6: Weighted  $R_{\text{partial}}^2$  for each factor showing the percentage of explained variability in trimethylamine *N*-oxide (TMAO) in (A) FAMILY and (B) START cohort**

Statistical significance was based on hierarchical linear models. \*  $p \leq 0.05$ , \*\* $p \leq 0.01$



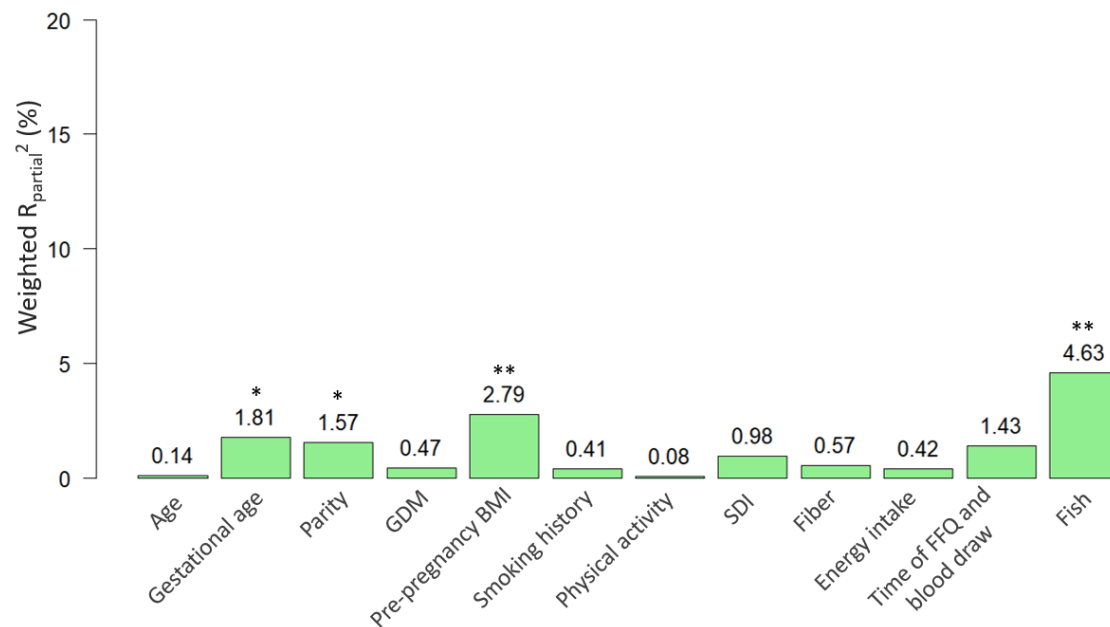
**Figure S7: Weighted  $R_{\text{partial}}^2$  for each factor showing the percentage of explained variability in 3-methylhistidine in (A) FAMILY and (B) START cohort**

Statistical significance was based on hierarchical linear models. \*  $p \leq 0.05$ , \*\* $p \leq 0.01$



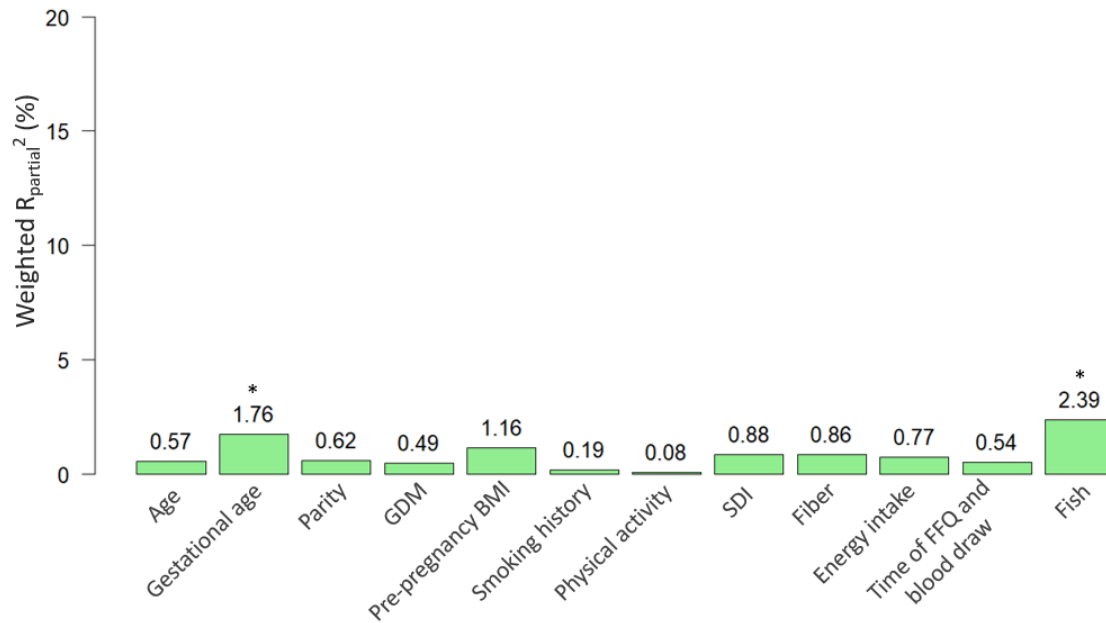
**Figure S8: Weighted  $R_{\text{partial}}^2$  for each factor showing the percentage of explained variability in Eicosapentaenoic acid (EPA, 20:5n-3) in FAMILY cohort**

Statistical significance was based on ordinary least squares regression.



**Figure S9: Weighted  $R_{\text{partial}}^2$  for each factor showing the percentage of explained variability in Docosahexaenoic acid (DHA; 22:6n-3) in FAMILY cohort**

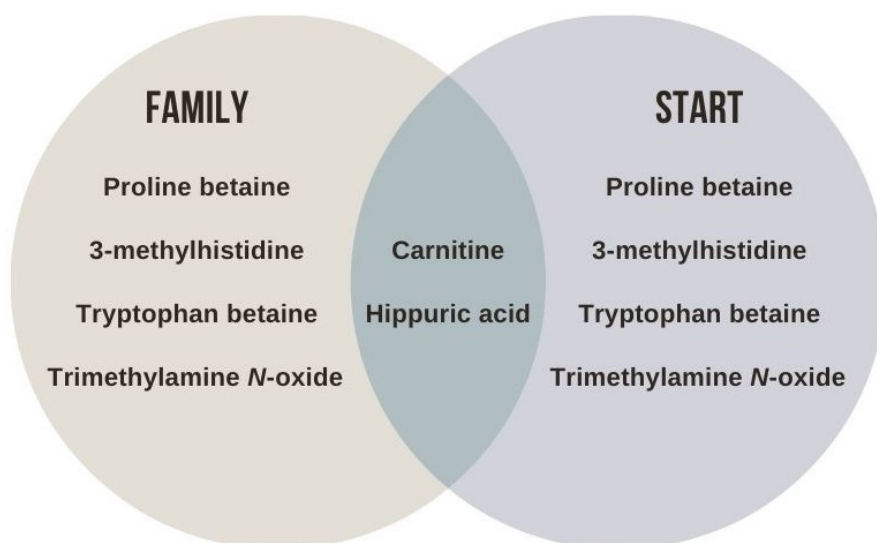
Statistical significance was based on ordinary least squares regression. \*  $p \leq 0.05$ , \*\* $p \leq 0.01$



**Figure S10: Weighted  $R_{\text{partial}}^2$  for each factor showing the percentage of explained variability in EPA + DHA in FAMILY cohort**

Statistical significance was based on ordinary least squares regression. \*  $p \leq 0.05$





**Figure S11: Venn diagram showing overlap of serum metabolites based on the cluster effect by ethnicity/cohort**