

Supplementary Table 1. Completeness and exclusions of the different metabolites, with limits of detection (LOD) and limits of quantification (LOQ)

Metabolites ¹	Results outside the measurable range (N=599 samples)		CV, % ³
	Samples, %	Criteria for results considered out of the measurable range ²	
Acylcarnitines			
C0	0.0	<LOD	3.5
C2	0.0	<LOD	3.9
C3	0.0	<LOD	4.3
C3-DC (C4-OH)	41.5	<LOD	-
C4	0.0	<LOD	4.7
C5	9.0	<LOD	6.2
C10	62.7	<LOD	-
C10:1	38.8	<LOD	-
C12	37.8	<LOD	-
C12:1	0.2	<LOD	8.5
C14:1	0.0	<LOD	5.0
C14:2	7.5	<LOD	5.1
C16	0.0	<LOD	4.3
C16:1	77.5	<LOD	-
C18	0.0	<LOD	7.0
C18:1	0.0	<LOD	4.8
C18:2	0.0	<LOD	5.9
Amino acids			
Alanine	0.0	<LLOQ (< 20 µmol/L)	4.0
Arginine	1.8	>ULOQ (> 400 µmol/L)	4.4
Asparagine	0.0	<LLOQ (< 5 µmol/L)	4.9
Aspartate	0.0	<LLOQ (< 5 µmol/L)	11.5
Citrulline	0.0	<LLOQ (< 5 µmol/L)	5.2
Glutamine	0.0	<LLOQ (< 20 µmol/L)	5.3
Glutamate	0.0	<LLOQ (< 10 µmol/L)	6.0
Glycine	0.0	<LLOQ (< 25 µmol/L)	7.4
Histidine	0.0	<LLOQ (< 5 µmol/L)	3.9
Isoleucine	0.0	<LLOQ (< 5 µmol/L)	6.2
Leucine	0.0	<LLOQ (< 50 µmol/L)	4.9
Lysine	0.0	<LLOQ (< 10 µmol/L)	7.5
Methionine	0.0	<LLOQ (< 5 µmol/L)	9.6
Ornithine	0.0	<LLOQ (< 5 µmol/L)	7.5
Phenylalanine	0.0	<LLOQ (< 5 µmol/L)	4.1
Proline	0.0	<LLOQ (< 10 µmol/L)	4.7
Serine	0.0	<LLOQ (< 5 µmol/L)	4.0
Threonine	0.0	<LLOQ (< 5 µmol/L)	4.0
Tryptophane	0.0	<LLOQ (< 5 µmol/L)	4.3
Tyrosine	0.0	<LLOQ (< 5 µmol/L)	3.8
Valine	0.0	<LLOQ (< 10 µmol/L)	9.1
Biogenic amines			

ADMA	0.0	<LLOQ (< 0.25 µmol/L)	10.4
Creatinine	0.0	<LLOQ (< 10 µmol/L)	3.7
Kynurenine	0.5	<LLOQ (< 1 µmol/L)	7.5
Putrescine	73.8	<LLOQ (< 0.1 µmol/L)	-
SDMA	0.0	<LLOQ (< 0.1 µmol/L)	7.3
Sarcosine	48.5	<LLOQ (< 1 µmol/L)	-
Serotonin ⁴	1.0	<LLOQ (< 0.1 µmol/L)	-
Spermidine	54.2	<LLOQ (< 0.25 µmol/L)	-
Spermine	94.5	<LLOQ (< 0.25 µmol/L)	-
Taurine	17.3	>ULOQ (> 200 µmol/L)	3.4
alpha-AAA	77.8	<LLOQ (< 1 µmol/L)	-
t4-OH-Pro	0.0		5.8
Glycerophospholipids			
PC aa C28:1	0.0	<LOD	6.1
PC aa C30:0	0.0	<LOD	6.2
PC aa C32:0	0.0	<LOD	6.1
PC aa C32:1	0.0	<LOD	5.9
PC aa C32:2	0.0	<LOD	5.5
PC aa C32:3	0.0	<LOD	6.1
PC aa C34:1	0.0	<LOD	4.6
PC aa C34:2	0.0	<LOD	4.6
PC aa C34:3	0.0	<LOD	5.3
PC aa C34:4	0.0	<LOD	5.1
PC aa C36:0	0.0	<LOD	6.3
PC aa C36:1	0.0	<LOD	5.7
PC aa C36:2	0.0	<LOD	4.6
PC aa C36:3	0.0	<LOD	5.3
PC aa C36:4	0.0	<LOD	4.7
PC aa C36:5	0.0	<LOD	4.8
PC aa C36:6	0.0	<LOD	5.6
PC aa C38:0	0.0	<LOD	7.2
PC aa C38:3	0.0	<LOD	5.3
PC aa C38:4	0.0	<LOD	4.7
PC aa C38:5	0.0	<LOD	4.9
PC aa C38:6	0.0	<LOD	4.9
PC aa C40:1	39.8	<LOD	-
PC aa C40:2	0.0	<LOD	8.2
PC aa C40:3	0.0	<LOD	8.7
PC aa C40:4	0.0	<LOD	5.0
PC aa C40:5	0.0	<LOD	5.3
PC aa C40:6	0.0	<LOD	5.7
PC aa C42:0	0.0	<LOD	7.4
PC aa C42:1	0.0	<LOD	10.9
PC aa C42:2	0.0	<LOD	9.5
PC aa C42:4	0.0	<LOD	13.3
PC aa C42:5	0.0	<LOD	9.0

PC aa C42:6	0.0	<LOD	6.3
PC ae C30:0	0.0	<LOD	8.5
PC ae C30:2	0.0	<LOD	11.7
PC ae C32:1	0.0	<LOD	6.2
PC ae C32:2	0.0	<LOD	6.7
PC ae C34:0	0.0	<LOD	6.8
PC ae C34:1	0.0	<LOD	5.5
PC ae C34:2	0.0	<LOD	6.2
PC ae C34:3	0.0	<LOD	5.6
PC ae C36:0	3.2	<LOD	7.6
PC ae C36:1	0.0	<LOD	7.0
PC ae C36:2	0.0	<LOD	5.2
PC ae C36:3	0.0	<LOD	4.8
PC ae C36:4	0.0	<LOD	5.5
PC ae C36:5	0.0	<LOD	5.4
PC ae C38:0	0.0	<LOD	7.6
PC ae C38:2	0.0	<LOD	7.6
PC ae C38:3	0.0	<LOD	7.1
PC ae C38:4	0.0	<LOD	5.1
PC ae C38:5	0.0	<LOD	5.5
PC ae C38:6	0.0	<LOD	4.9
PC ae C40:1	0.0	<LOD	8.3
PC ae C40:2	0.0	<LOD	5.7
PC ae C40:3	0.0	<LOD	5.6
PC ae C40:4	0.0	<LOD	7.4
PC ae C40:5	0.0	<LOD	6.7
PC ae C40:6	0.0	<LOD	5.8
PC ae C42:1	0.0	<LOD	10.1
PC ae C42:2	0.0	<LOD	6.4
PC ae C42:3	0.0	<LOD	7.4
PC ae C42:4	0.5	<LOD	7.8
PC ae C42:5	0.0	<LOD	5.2
PC ae C44:3	0.0	<LOD	11.8
PC ae C44:4	0.0	<LOD	7.1
PC ae C44:5	0.0	<LOD	6.5
PC ae C44:6	0.0	<LOD	5.9
lysoPC a C16:0	0.0	<LOD	5.7
lysoPC a C16:1	0.0	<LOD	6.4
lysoPC a C17:0	0.0	<LOD	6.3
lysoPC a C18:0	0.0	<LOD	5.8
lysoPC a C18:1	0.0	<LOD	6.1
lysoPC a C18:2	0.0	<LOD	6.8
lysoPC a C20:3	0.0	<LOD	5.2
lysoPC a C20:4	0.0	<LOD	5.9
lysoPC a C24:0	0.0	<LOD	9.9
Sphingolipids			

SM C16:0	0.0	<LOD	6.1
SM C16:1	0.0	<LOD	6.3
SM C18:0	0.0	<LOD	7.2
SM C18:1	0.0	<LOD	6.4
SM C20:2	0.0	<LOD	9.7
SM C24:0	0.0	<LOD	6.0
SM C24:1	0.0	<LOD	6.5
SM C26:0	9.0	<LOD	11.7
SM C26:1	1.0	<LOD	10.7
SM (OH) C14:1	0.0	<LOD	6.5
SM (OH) C16:1	0.0	<LOD	6.2
SM (OH) C22:1	0.0	<LOD	7.0
SM (OH) C22:2	0.0	<LOD	5.8
SM (OH) C24:1	0.0	<LOD	9.9
Hexose			
Hexose	0.0	<LLOQ (< 1500 µmol/L)	3.4
¹ Metabolites in gray were excluded from the statistical analysis because more than 20% of the measures were outside the measurable range. ² Limit of detection (LOD) is applicable to acylcarnitines, glycerophospholipids, sphingolipids. Lower limit of quantification (LLOQ) or upper limit of quantification (ULOQ) is applicable to amino acids, biogenic amines and hexose. No metabolites had values both under LLOQ and over ULOQ. ³ Mean intra- and inter-batch coefficients of variation (CVs) based on two quality control samples measured in duplicate in each batch. ⁴ CV for serotonin not calculated because measures in quality controls were lower than LLOQ.			

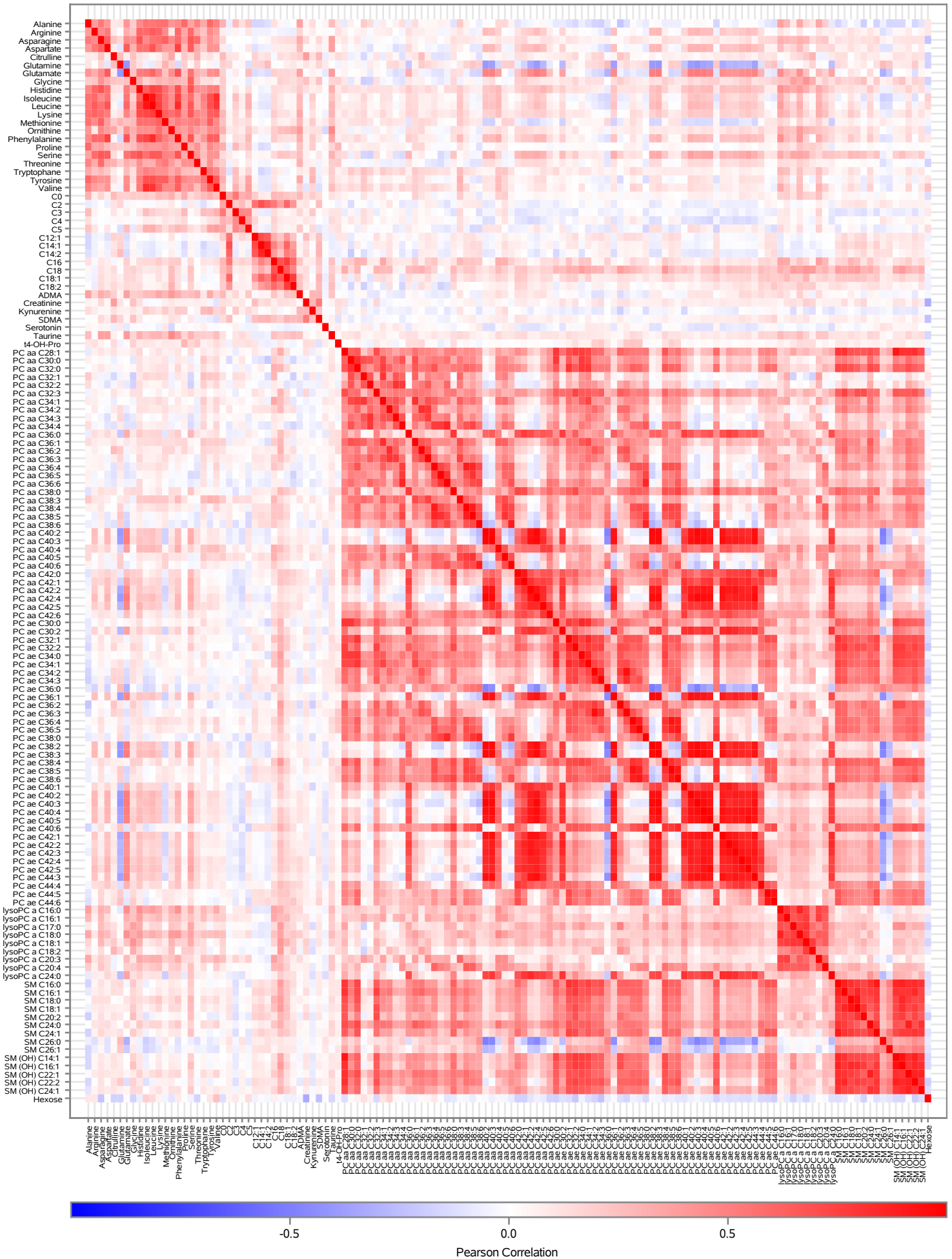
Supplementary Table 2. Geometric means of metabolites concentrations

Variable	N	Geometric mean (95% CI) (μmol/L)
C0	573	32.18 (31.69-32.69)
C2	573	5.31 (5.19-5.44)
C3	573	0.30 (0.29-0.31)
C4	573	0.15 (0.14-0.15)
C5	573	0.11 (0.11-0.12)
C12:1	573	0.21 (0.21-0.21)
C14:1	573	0.06 (0.06-0.06)
C14:2	573	0.03 (0.03-0.03)
C16	573	0.11 (0.11-0.11)
C18	573	0.05 (0.04-0.05)
C18:1	573	0.12 (0.12-0.13)
C18:2	573	0.06 (0.06-0.06)
Alanine	573	469.21 (461.08-477.48)
Arginine	573	157.23 (153.14-161.43)
Asparagine	573	51.05 (50.19-51.92)
Aspartate	573	51.50 (50.28-52.75)
Citrulline	573	25.86 (25.32-26.41)
Glutamine	573	491.29 (483.10-499.62)
Glutamate	573	153.86 (148.98-158.90)
Glycine	573	348.92 (341.28-356.73)
Histidine	573	99.11 (97.83-100.41)
Isoleucine	573	76.22 (74.96-77.51)
Leucine	573	155.26 (152.62-157.94)
Lysine	573	247.69 (243.53-251.93)
Methionine	573	23.47 (22.99-23.95)
Ornithine	573	93.29 (90.95-95.68)
Phenylalanine	573	99.09 (97.54-100.66)
Proline	573	201.05 (196.96-205.23)
Serine	573	179.38 (176.39-182.41)
Threonine	573	133.40 (130.66-136.19)
Tryptophane	573	63.24 (62.40-64.09)
Tyrosine	573	73.30 (72.02-74.61)
Valine	573	231.29 (227.26-235.39)
ADMA	573	0.47 (0.47-0.48)
Creatinine	573	53.09 (52.37-53.83)
Kynurenine	573	2.08 (2.04-2.12)
SDMA	573	0.42 (0.41-0.42)
Serotonin	573	0.67 (0.64-0.70)
Taurine	573	147.71 (144.49-151.00)
t4-OH-Pro	573	10.59 (10.18-11.01)
PC aa C28:1	573	1.78 (1.73-1.82)
PC aa C30:0	573	2.31 (2.25-2.37)
PC aa C32:0	573	17.31 (16.97-17.65)

PC aa C32:1	573	18.59 (17.95-19.24)
PC aa C32:2	573	6.75 (6.58-6.94)
PC aa C32:3	573	0.99 (0.97-1.01)
PC aa C34:1	573	238.47 (234.24-242.78)
PC aa C34:2	573	525.26 (518.64-531.96)
PC aa C34:3	573	25.11 (24.62-25.61)
PC aa C34:4	573	2.36 (2.29-2.42)
PC aa C36:0	573	2.64 (2.58-2.70)
PC aa C36:1	573	44.09 (43.19-45.00)
PC aa C36:2	573	288.26 (283.94-292.64)
PC aa C36:3	573	177.98 (175.08-180.92)
PC aa C36:4	573	197.37 (193.37-201.45)
PC aa C36:5	573	16.77 (16.21-17.34)
PC aa C36:6	573	0.94 (0.91-0.97)
PC aa C38:0	573	2.30 (2.25-2.35)
PC aa C38:3	573	49.27 (48.27-50.29)
PC aa C38:4	573	99.18 (97.06-101.34)
PC aa C38:5	573	48.30 (47.22-49.41)
PC aa C38:6	573	59.60 (58.12-61.12)
PC aa C40:2	573	0.48 (0.45-0.52)
PC aa C40:3	573	0.62 (0.59-0.65)
PC aa C40:4	573	3.43 (3.35-3.51)
PC aa C40:5	573	7.31 (7.14-7.48)
PC aa C40:6	573	18.15 (17.70-18.61)
PC aa C42:0	573	0.29 (0.28-0.29)
PC aa C42:1	573	0.16 (0.16-0.16)
PC aa C42:2	573	0.16 (0.15-0.17)
PC aa C42:4	573	0.18 (0.17-0.19)
PC aa C42:5	573	0.26 (0.25-0.26)
PC aa C42:6	573	0.28 (0.28-0.29)
PC ae C30:0	573	0.21 (0.20-0.21)
PC ae C30:2	573	0.08 (0.08-0.09)
PC ae C32:1	573	3.41 (3.34-3.49)
PC ae C32:2	573	1.04 (1.02-1.06)
PC ae C34:0	573	1.64 (1.60-1.68)
PC ae C34:1	573	12.64 (12.41-12.88)
PC ae C34:2	573	16.59 (16.24-16.94)
PC ae C34:3	573	10.13 (9.88-10.39)
PC ae C36:0	573	0.61 (0.58-0.63)
PC ae C36:1	573	14.13 (13.54-14.74)
PC ae C36:2	573	19.10 (18.72-19.49)
PC ae C36:3	573	10.47 (10.26-10.68)
PC ae C36:4	573	20.80 (20.34-21.28)
PC ae C36:5	573	11.87 (11.57-12.18)
PC ae C38:0	573	1.70 (1.66-1.73)
PC ae C38:2	573	4.49 (4.25-4.76)

PC ae C38:3	573	8.33 (7.93-8.75)
PC ae C38:4	573	14.14 (13.86-14.41)
PC ae C38:5	573	17.94 (17.56-18.32)
PC ae C38:6	573	6.55 (6.39-6.70)
PC ae C40:1	573	1.02 (0.99-1.05)
PC ae C40:2	573	1.77 (1.70-1.84)
PC ae C40:3	573	2.20 (2.05-2.36)
PC ae C40:4	573	3.12 (3.00-3.26)
PC ae C40:5	573	4.60 (4.43-4.77)
PC ae C40:6	573	3.54 (3.47-3.62)
PC ae C42:1	573	0.42 (0.40-0.43)
PC ae C42:2	573	0.42 (0.41-0.43)
PC ae C42:3	573	0.61 (0.59-0.63)
PC ae C42:4	573	0.74 (0.71-0.76)
PC ae C42:5	573	1.74 (1.69-1.79)
PC ae C44:3	573	0.09 (0.08-0.09)
PC ae C44:4	573	0.20 (0.19-0.20)
PC ae C44:5	573	0.68 (0.66-0.69)
PC ae C44:6	573	0.57 (0.56-0.59)
lysoPC a C16:0	573	156.23 (152.58-159.96)
lysoPC a C16:1	573	4.30 (4.20-4.41)
lysoPC a C17:0	573	2.96 (2.88-3.04)
lysoPC a C18:0	573	47.71 (46.49-48.96)
lysoPC a C18:1	573	25.45 (24.89-26.02)
lysoPC a C18:2	573	25.22 (24.68-25.77)
lysoPC a C20:3	573	2.08 (2.03-2.13)
lysoPC a C20:4	573	5.91 (5.75-6.07)
lysoPC a C24:0	573	0.11 (0.11-0.12)
SM C16:0	573	67.14 (65.77-68.54)
SM C16:1	573	10.49 (10.29-10.69)
SM C18:0	573	12.19 (11.92-12.46)
SM C18:1	573	6.78 (6.63-6.93)
SM C20:2	573	0.28 (0.27-0.28)
SM C24:0	573	6.60 (6.47-6.73)
SM C24:1	573	13.11 (12.82-13.40)
SM C26:0	573	0.03 (0.03-0.03)
SM C26:1	573	0.06 (0.06-0.07)
SM (OH) C14:1	573	3.84 (3.74-3.94)
SM (OH) C16:1	573	2.02 (1.97-2.07)
SM (OH) C22:1	573	5.16 (5.05-5.27)
SM (OH) C22:2	573	3.64 (3.56-3.73)
SM (OH) C24:1	573	0.39 (0.38-0.40)
Hexose	573	4423.7 (4334.1-4515.2)

Supplementary Figure 1. Pearson's partial correlations between metabolites adjusted for age, state, and batch



Supplementary Figure 2. Associations between metabolites and non-dense area. A: Raw P-values. B: Adjusted P-values. Estimates per standard deviation increase in residuals of metabolites on batch were obtained from linear regression adjusted for age, BMI, age at menarche, family history of cancer, history of benign breast disease, use of oral contraceptive, number of full-term pregnancies, age at first full-term pregnancy, breastfeeding, alcohol intake, smoking status, socioeconomic status, and physical activity. Dotted lines represent statistical significance thresholds for raw P-values (A) and for P-values adjusted by permutation-based stepdown *minP* (B). Abbreviations: ae: acyl-alkyl; aa: acyl-acyl; PC: phosphatidylcholine; SM: sphingomyelin; SD: standard deviation.

