

Electronic Supplementary Material Table 1. Baseline Characteristics in the Analytic Study Population and the Overall ARIC Study

	Analytic Study Population (N=2,939)	ARIC Study Population (N=15,792)
Age, years	53.3 (5.7)	54.2 (5.8)
Female sex	1,755 (59.7)	8,710 (55.2)
African American	1,665 (56.7)	4,266 (27.0)
Study center		
Forsyth County, North Carolina	485 (16.5)	4,035 (25.6)
Jackson, Mississippi	1,566 (53.3)	3,728 (23.6)
Minneapolis, Minnesota	455 (15.5)	4,009 (25.4)
Washington County, Maryland	433 (14.7)	4,020 (25.5)
Batch		
Batch 1	1,275 (43.4)	N/A
Batch 2	1,664 (56.6)	N/A
Education, years		
≤11	835 (28.4)	3,767 (23.9)
12-16	1,029 (35.0)	6,412 (40.6)
17-21	1,075 (36.6)	5,586 (35.4)
Systolic blood pressure, mmHg	122.8 (19.8)	121.4 (19.0)
Diastolic blood pressure, mmHg	76.3 (12.3)	73.7 (11.3)
Body mass index, kg/m ²	28.2 (5.6)	27.7 (5.4)
HDL cholesterol, mg/dL	54.4 (17.4)	51.6 (17.1)
LDL cholesterol, mg/dL	136.8 (39.7)	137.6 (39.4)
Physical activity index	2.3 (0.8)	2.4 (0.8)
eGFR, mL/min/1.73 m ²	106.4 (17.4)	102.5 (15.8)
Current smoking	779 (26.5)	4,132 (26.2)
History of CVD	269 (9.2)	1,365 (8.6)
Fasting glucose, mmol/L	5.5 (0.6)	6.0 (2.3)

CVD, cardiovascular disease; eGFR, estimated glomerular filtration rate; HDL, high-density lipoprotein; LDL, low-density lipoprotein; N/A, not applicable

Data are expressed as means (SD) and *n* (%).

Electronic Supplementary Material Table 2. Heat Map of Hazard Ratios^a for Incident Diabetes Per One Standard Deviation Increase in Log-Transformed Serum Metabolites

Pathway and Sub-Pathway	Metabolites	Model 1	Model 2	Model 3	Color Scale
Amino Acid					
Alanine and Aspartate Metabolism	N-acetylalanine	2.24	1.44	1.59	0
Alanine and Aspartate Metabolism	asparagine	0.67	0.75	0.78	1
Alanine and Aspartate Metabolism	alanine	2.57	1.84	1.26	2
Glutamate Metabolism	glutamate	2.25	2.12	1.86	3
Glutathione Metabolism	5-oxoproline	0.63	0.67	0.76	4
Glycine, Serine and Threonine Metabolism	glycine	0.44	0.60	0.73	5
Glycine, Serine and Threonine Metabolism	N-acetylthreonine	1.50	1.19	1.10	6
Leucine, Isoleucine and Valine Metabolism	alpha-hydroxyisovalerate	1.28	1.30	1.17	7
Leucine, Isoleucine and Valine Metabolism	2-methylbutyrylcarnitine (C5)	1.49	1.25	1.14	8
Leucine, Isoleucine and Valine Metabolism	leucine	4.58	3.12	2.37	9
Leucine, Isoleucine and Valine Metabolism	isoleucine	7.57	4.42	2.96	10
Leucine, Isoleucine and Valine Metabolism	3-hydroxyisobutyrate	1.65	1.53	1.29	11
Leucine, Isoleucine and Valine Metabolism	valine	6.49	3.59	2.41	12
Leucine, Isoleucine and Valine Metabolism	isovalerylcarnitine	1.50	1.26	1.21	13
Leucine, Isoleucine and Valine Metabolism	isovalerate	2.07	2.05	1.41	14
Lysine Metabolism	lysine	2.20	1.62	1.30	
Lysine Metabolism	N6-acetyllysine	1.63	1.21	1.23	
Methionine, Cysteine, SAM and Taurine Metabolism	2-hydroxybutyrate (AHB)	1.61	1.57	1.30	
Phenylalanine and Tyrosine Metabolism	N-acetylphenylalanine	1.36	1.20	1.04	
Phenylalanine and Tyrosine Metabolism	phenyllactate (PLA)	1.57	1.36	1.23	
Phenylalanine and Tyrosine Metabolism	phenylalanine	3.71	2.17	1.69	
Phenylalanine and Tyrosine Metabolism	tyrosine	4.86	2.65	1.73	
Phenylalanine and Tyrosine Metabolism	3-(4-hydroxyphenyl)lactate	2.57	2.00	1.55	
Tryptophan Metabolism	tryptophan	2.33	1.98	1.40	
Tryptophan Metabolism	kynurenine	2.09	1.25	1.14	
Urea cycle; Arginine and Proline Metabolism	ornithine	1.43	1.28	1.25	
Urea cycle; Arginine and Proline Metabolism	pro-hydroxy-pro	0.66	0.74	0.77	
Urea cycle; Arginine and Proline Metabolism	proline	2.10	1.45	1.10	
Carbohydrate					
Disaccharides and Oligosaccharides	trehalose	1.31	1.26	1.16	
Fructose, Mannose and Galactose Metabolism	mannose	4.13	2.53	1.28	
Glycolysis, Gluconeogenesis, and Pyruvate Metabolism	glucose	14.14	11.21	2.24	
Glycolysis, Gluconeogenesis, and Pyruvate Metabolism	lactate	2.94	2.23	1.51	
Cofactors and Vitamins					

Nicotinate and Nicotinamide Metabolism	N1-Methyl-2-pyridone-5-carboxamide	1.27	1.10	1.14
Energy				
TCA Cycle	succinate	2.73	2.11	1.46
Lipid				
Carnitine Metabolism	carnitine	2.63	1.63	1.15
Fatty Acid Metabolism(Acyl Carnitine)	hydroxybutyrylcarnitine	1.21	1.13	1.09
Long Chain Fatty Acid	myristate (14:0)	1.46	1.50	1.36
Long Chain Fatty Acid	palmitate (16:0)	2.10	1.99	1.71
Long Chain Fatty Acid	10-heptadecenoate (17:1n7)	1.46	1.36	1.24
Long Chain Fatty Acid	palmitoleate (16:1n7)	1.41	1.32	1.26
Long Chain Fatty Acid	oleate (18:1n9)	1.49	1.47	1.41
Long Chain Fatty Acid	stearate (18:0)	1.76	2.01	1.72
Long Chain Fatty Acid	10-nonadecenoate (19:1n9)	1.38	1.36	1.26
Lysolipid	1-arachidonoylglycerophosphoinositol	1.53	1.65	1.48
Lysolipid	1-oleoylglycerophosphocholine (18:1)	0.64	0.78	0.81
Monoacylglycerol	1-oleoylglycerol (1-monoolein)	1.60	1.29	1.21
Phospholipid Metabolism	glycerophosphorylcholine (GPC)	0.69	0.87	0.93
Phospholipid Metabolism	choline	0.43	0.58	0.62
Polyunsaturated Fatty Acid (n3 and n6)	dihomo-linolenate (20:3n3 or n6)	2.12	1.80	1.57
Polyunsaturated Fatty Acid (n3 and n6)	dihomo-linoleate (20:2n6)	1.40	1.44	1.32
Polyunsaturated Fatty Acid (n3 and n6)	stearidonate (18:4n3)	1.28	1.32	1.18
Polyunsaturated Fatty Acid (n3 and n6)	arachidonate (20:4n6)	1.63	1.76	1.47
Polyunsaturated Fatty Acid (n3 and n6)	adrenate (22:4n6)	1.59	1.51	1.33
Polyunsaturated Fatty Acid (n3 and n6)	eicosapentaenoate (EPA; 20:5n3)	1.31	1.36	1.17
Secondary Bile Acid Metabolism	glycocholate sulfate	1.46	1.30	1.22
Steroid	4-androsten-3beta,17beta-diol disulfate (1)	1.24	1.23	1.09
Steroid	4-androsten-3beta,17beta-diol disulfate (2)	1.31	1.20	1.05
Nucleotide				
Purine Metabolism, (Hypo)Xanthine/Inosine containing	urate	2.79	1.78	1.42
Pyrimidine Metabolism, Uracil containing	N-acetyl-beta-alanine	1.50	1.33	1.32
Peptide				
Dipeptide	phenylalanylleucine	1.36	1.22	1.17
Dipeptide	phenylalanylphenylalanine	1.43	1.34	1.19
Dipeptide	pyroglutamylglycine	1.23	1.17	1.09
Dipeptide	leucylleucine	1.28	1.18	1.06
Dipeptide Derivative	N-acetylcarnosine	1.38	1.20	1.12
Gamma-glutamyl Amino Acid	gamma-glutamylglutamate	0.82	0.85	0.90
Gamma-glutamyl Amino Acid	gamma-glutamylthreonine	0.71	0.76	0.84
Polypeptide	HWESASLLR	1.12	1.08	1.07
Xenobiotics				

Food Component/Plant	erythritol	2.41	1.93	1.66
Food Component/Plant	gluconate	1.35	1.30	1.16
Food Component/Plant	piperine	1.11	1.07	1.02
Xanthine Metabolism	caffeine	1.11	1.08	1.06
Xanthine Metabolism	theobromine	1.13	1.11	1.09
Xanthine Metabolism	theophylline	1.11	1.09	1.06

^a Hazard ratios are bolded if the association is statistically significant at the Bonferroni-corrected level ($0.05/245=2.04 \times 10^{-4}$).

Model 1: Adjusted for age, sex, race-center, and batch

Model 2: Adjusted for covariates in Model 1 as well as education level, systolic blood pressure, diastolic blood pressure, body mass index, high density lipoprotein cholesterol, low density lipoprotein cholesterol, smoking status, physical activity level, history of cardiovascular disease, and estimated glomerular filtration rate

Model 3: Adjusted for covariates in Model 2 as well as fasting glucose

Electronic Supplementary Material Table 3. Serum Metabolites Significantly Associated with Incident Diabetes According to Metabolic Pathway and Stratified by Race

Pathway and Sub-Pathway	Metabolites	White		Black		<i>p</i> value for interaction
		HR ^a (95% CI)	<i>p</i> value	HR ^a (95% CI)	<i>p</i> value	
Amino Acid						
Leucine, Isoleucine and Valine Metabolism	isoleucine	2.73 (1.48, 5.02)	1.26×10 ⁻³	2.93 (1.78, 4.83)	2.48×10 ⁻⁵	0.94
Alanine and Aspartate Metabolism	asparagine	0.69 (0.60, 0.80)	3.70×10 ⁻⁷	0.83 (0.73, 0.93)	1.26×10 ⁻³	0.05
Leucine, Isoleucine and Valine Metabolism	leucine	2.81 (1.51, 5.22)	1.06×10 ⁻³	2.06 (1.28, 3.33)	3.06×10 ⁻³	0.49
Phenylalanine and Tyrosine Metabolism	3-(4-hydroxyphenyl)lactate	1.69 (1.18, 2.41)	4.05×10 ⁻³	1.48 (1.13, 1.92)	4.08×10 ⁻³	0.90
Leucine, Isoleucine and Valine Metabolism	valine	2.61 (1.30, 5.23)	7.03×10 ⁻³	2.12 (1.21, 3.72)	8.48×10 ⁻³	0.43
Carbohydrate						
Disaccharides and Oligosaccharides	trehalose	1.13 (1.02, 1.25)	2.08×10 ⁻²	1.20 (1.10, 1.32)	1.23×10 ⁻⁴	0.24
Xenobiotic						
Food Component/Plant	erythritol	2.91 (1.72, 4.92)	6.89×10 ⁻⁵	1.42 (1.05, 1.93)	2.50×10 ⁻²	0.12

CI, confidence interval; HR, hazard ratio

^a Hazard ratio per one standard deviation increase in the log-transformed metabolite in multivariable regression models adjusted for age, sex, center, batch, education level, systolic blood pressure, diastolic blood pressure, body mass index, high density lipoprotein cholesterol, low density lipoprotein cholesterol, smoking status, physical activity level, history of cardiovascular disease, estimated glomerular filtration rate, and fasting glucose.

Electronic Supplementary Material Table 4. Pearson's Correlation Coefficients for Serum Metabolites Significantly Associated with Incident Diabetes

	Isoleucine	Asparagine	Leucine	Trehalose	3-(4-hydroxyphenyl) lactate	Valine	Erythritol
Isoleucine	1						
Asparagine	-0.07 <i>p</i> <0.001	1					
Leucine	0.86 <i>p</i> <0.001	0.009 <i>p</i> =0.64	1				
Trehalose	0.11 <i>p</i> <0.001	-0.11 <i>p</i> <0.001	0.11 <i>p</i> <0.001	1			
3-(4-hydroxyphenyl) lactate	0.50 <i>p</i> <0.001	-0.10 <i>p</i> <0.001	0.45 <i>p</i> <0.001	0.10 <i>p</i> <0.001	1		
Valine	0.83 <i>p</i> <0.001	-0.06 <i>p</i> <0.001	0.86 <i>p</i> <0.001	0.11 <i>p</i> <0.001	0.42 <i>p</i> <0.001	1	
Erythritol	0.28 <i>p</i> <0.001	-0.009 <i>p</i> =0.64	0.24 <i>p</i> <0.001	0.03 <i>p</i> =0.06	0.31 <i>p</i> <0.001	0.23 <i>p</i> <0.001	1

Color rules for abs(corr)	0	0.2	0.4	0.6	0.8	1