

WEB MATERIAL.

erythritol			pyroglutamine*		
covariate	β (standard error)	p value	covariate	β (standard error)	p value
age	1.29(0.12)	<0.001	age	0.62(0.12)	<0.001
gender	0.13(0.04)	0.005	gender	1.36(0.07)	<0.001
CHD	0.38(0.13)	0.003	CHD	0.40(0.13)	0.002
smoking	0.02(0.05)	0.635	smoking	0.34(0.05)	<0.001
physical activity	-0.13(0.03)	<0.001	physical activity	-0.01(0.03)	0.782
education	-0.11(0.04)	0.004	education	0.00(0.04)	0.965
BMI	0.44(0.13)	0.001	BMI	-1.09(0.12)	<0.001
SBP	1.95(0.45)	<0.001	SBP	1.57(0.46)	0.001
BP med	0.32(0.05)	<0.001	BP med	-0.04(0.04)	0.420
glucose	0.47(0.05)	<0.001	glucose	-0.32(0.05)	<0.001
diabetes	0.44(0.06)	<0.001	diabetes	-0.33(0.06)	<0.001
TC : HDL ratio	0.28(0.03)	<0.001	TC : HDL ratio	0.17(0.03)	<0.001
lipid med	0.02(0.22)	0.913	lipid med	-0.17(0.22)	0.447
LVH	0.41(0.11)	<0.001	LVH	0.16(0.10)	0.120
eGFR	-6.28(0.36)	<0.001	eGFR	-5.03(0.37)	<0.001

N-acetylalanine			phenylacetylglutamine		
covariate	β (standard error)	p value	covariate	β (standard error)	p value
age	0.93(0.12)	<0.001	age	0.96(0.12)	<0.001
gender	0.10(0.04)	0.020	gender	-0.03(0.04)	0.567
CHD	0.21(0.12)	0.083	CHD	0.40(0.13)	0.002
smoking	0.02(0.05)	0.714	smoking	-0.17(0.05)	<0.001
physical activity	-0.10(0.03)	0.001	physical activity	-0.13(0.03)	<0.001
education	-0.13(0.04)	<0.001	education	-0.16(0.04)	<0.001
BMI	0.53(0.13)	<0.001	BMI	-0.06(0.13)	0.625
SBP	2.13(0.45)	<0.001	SBP	0.74(0.46)	0.104
BP med	0.32(0.05)	<0.001	BP med	0.18(0.05)	<0.001
glucose	-0.22(0.05)	<0.001	glucose	0.17(0.05)	0.001
diabetes	-0.18(0.06)	0.002	diabetes	0.14(0.06)	0.019
TC : HDL ratio	0.13(0.03)	<0.001	TC : HDL ratio	0.01(0.03)	0.698
lipid med	-0.02(0.22)	0.911	lipid med	0.22(0.22)	0.331
LVH	0.35(0.11)	0.001	LVH	0.26(0.10)	0.011
eGFR	-6.83(0.35)	<0.001	eGFR	-4.05(0.37)	<0.001

Metabolite values are in the form of quartile ordinal scores; continuous variables used liner model (e.g, age= intercept+ β ×quartile scores); categorical variables used logistic model (e.g, logit(CHD)= quartile scores).

Abbreviations: CHD, coronary heart disease; BMI, body mass index; SBP, systolic blood pressure; BP med, anti-hypertensive medication use; TC : HDL ratio, serum total cholesterol to high-density-cholesterol lipoprotein ratio; lipid med, lipid-lowering medication use; LVH, left ventricular hypertrophy; eGFR, estimated glomerular filtration rate.

Regression Coefficients with Established Risk Factors for Quartile-score of Sixteen HF Related Metabolites in Model 3

p-cresol sulfate		
covariate	β (standard error)	p value
age	0.59(0.12)	<0.001
gender	-0.01(0.04)	0.841
CHD	0.14(0.12)	0.254
smoking	-0.16(0.05)	0.001
physical activity	-0.11(0.03)	0.001
education	-0.07(0.04)	0.044
BMI	-0.56(0.13)	<0.001
SBP	-0.01(0.46)	0.989
BP med	0.11(0.04)	0.018
glucose	0.03(0.05)	0.623
diabetes	-0.02(0.06)	0.679
TC : HDL ratio	0.02(0.03)	0.508
lipid med	-0.17(0.22)	0.447
LVH	0.17(0.10)	0.098
eGFR	-3.40(0.38)	<0.001

X - 11429		
covariate	β (standard error)	p value
age	1.00(0.12)	<0.001
gender	0.02(0.04)	0.672
CHD	0.12(0.12)	0.307
smoking	0.20(0.05)	<0.001
physical activity	-0.16(0.03)	<0.001
education	-0.12(0.04)	0.001
BMI	0.28(0.13)	0.026
SBP	1.54(0.46)	0.001
BP med	0.23(0.05)	<0.001
glucose	-0.36(0.05)	<0.001
diabetes	-0.34(0.06)	<0.001
TC : HDL ratio	0.02(0.03)	0.448
lipid med	-0.54(0.24)	0.028
LVH	0.30(0.10)	0.005
eGFE	-5.56(0.36)	<0.001

pro-hydroxy-pro		
covariate	β (standard error)	p value
age	0.69(0.12)	<0.001
gender	0.04(0.04)	0.370
CHD	0.15(0.12)	0.209
smoking	0.20(0.05)	<0.001
physical activity	-0.05(0.03)	0.131
education	-0.17(0.04)	<0.001
BMI	-0.64(0.13)	<0.001
SBP	1.44(0.46)	0.002
BP med	-0.10(0.04)	0.031
glucose	-0.13(0.05)	0.020
diabetes	-0.08(0.06)	0.190
TC : HDL ratio	-0.06(0.03)	0.064
lipid med	-0.02(0.22)	0.912
LVH	0.14(0.10)	0.177
eGFR	-3.20(0.38)	<0.001

X - 11687_200		
covariate	β (standard error)	p value
age	0.61(0.11)	<0.001
gender	0.06(0.04)	0.144
CHD	0.03(0.11)	0.810
smoking	0.19(0.04)	<0.001
physical activity	-0.11(0.03)	<0.001
education	-0.06(0.03)	0.098
BMI	0.00(0.12)	0.993
SBP	1.67(0.42)	<0.001
BP med	0.27(0.04)	<0.001
glucose	0.07(0.05)	0.132
diabetes	0.08(0.05)	0.132
TC : HDL ratio	0.12(0.03)	<0.001
lipid med	-0.19(0.21)	0.368
LVH	0.18(0.09)	0.050
eGFE	-3.64(0.35)	<0.001

Metabolite values are in the form of quartile ordinal scores; continuous variables used liner model (e.g, age= intercept+ β ×quartile scores); categorical variables used logistic model (e.g, logit(CHD)= quartile scores).

Abbreviations: CHD, coronary heart disease; BMI, body mass index; SBP, systolic blood pressure; BP med, anti-hypertensive medication use; TC : HDL ratio, serum total cholesterol to high-density-cholesterol lipoprotein ratio; lipid med, lipid-lowering medication use; LVH, left ventricular hypertrophy; eGFR, estimated glomerular filtration rate.

Regression Coefficients with Established Risk Factors for Quartile-score of Sixteen HF Related Metabolites in Model 3

X - 12096			X - 11334		
covariate	β (standard error)	p value	covariate	β (standard error)	p value
age	0.92(0.12)	<0.001	age	0.95(0.12)	<0.001
gender	-0.06(0.04)	0.161	gender	0.31(0.05)	<0.001
CHD	0.20(0.12)	0.106	CHD	0.23(0.12)	0.064
smoking	0.02(0.05)	0.635	smoking	-0.13(0.05)	0.006
physical activity	-0.10(0.03)	0.001	physical activity	-0.04(0.03)	0.156
education	-0.15(0.04)	<0.001	education	-0.08(0.04)	0.041
BMI	1.57(0.12)	<0.001	BMI	0.05(0.13)	0.701
SBP	1.05(0.46)	0.021	SBP	0.96(0.46)	0.036
BP med	0.35(0.05)	<0.001	BP med	0.17(0.05)	<0.001
glucose	0.22(0.05)	<0.001	glucose	-0.04(0.05)	0.406
diabetes	0.28(0.06)	<0.001	diabetes	-0.02(0.06)	0.767
TC : HDL ratio	0.33(0.03)	<0.001	TC : HDL ratio	0.11(0.03)	<0.001
lipid med	-0.22(0.22)	0.330	lipid med	0.02(0.22)	0.913
LVH	0.37(0.11)	<0.001	LVH	0.18(0.10)	0.080
eGFR	-4.51(0.37)	<0.001	eGFR	-5.87(0.36)	<0.001

X - 11787			X - 11423		
covariate	β (standard error)	p value	covariate	β (standard error)	p value
age	1.07(0.12)	<0.001	age	0.78(0.12)	<0.001
gender	0.24(0.05)	<0.001	gender	0.23(0.05)	<0.001
CHD	0.49(0.13)	<0.001	CHD	0.24(0.12)	0.050
smoking	0.05(0.05)	0.255	smoking	0.09(0.05)	0.053
physical activity	-0.02(0.03)	0.412	physical activity	-0.09(0.03)	0.002
education	-0.15(0.04)	<0.001	education	0.04(0.04)	0.312
BMI	0.30(0.13)	0.019	BMI	-0.14(0.13)	0.283
SBP	0.91(0.46)	0.047	SBP	1.74(0.45)	<0.001
BP med	0.13(0.05)	0.003	BP med	0.20(0.05)	<0.001
glucose	0.29(0.05)	<0.001	glucose	-0.18(0.05)	0.001
diabetes	0.29(0.06)	<0.001	diabetes	-0.19(0.06)	0.002
TC : HDL ratio	0.15(0.03)	<0.001	TC : HDL ratio	0.16(0.03)	<0.001
lipid med	0.60(0.25)	0.017	lipid med	-0.07(0.22)	0.737
LVH	0.35(0.11)	0.001	LVH	0.25(0.10)	0.015
eGFR	-3.76(0.38)	<0.001	eGFR	-6.20(0.36)	<0.001

Metabolite values are in the form of quartile ordinal scores; continuous variables used liner model (e.g, age= intercept+ β ×quartile scores); categorical variables used logistic model (e.g, logit(CHD)= quartile scores).

Abbreviations: CHD, coronary heart disease; BMI, body mass index; SBP, systolic blood pressure; BP med, anti-hypertensive medication use; TC : HDL ratio, serum total cholesterol to high-density-cholesterol lipoprotein ratio; lipid med, lipid-lowering medication use; LVH, left ventricular hypertrophy; eGFR, estimated glomerular filtration rate.

Regression Coefficients with Established Risk Factors for Quartile-score of Sixteen HF Related Metabolites in Model 3

X - 04499			X - 11333		
covariate	β (standard error)	p value	covariate	β (standard error)	p value
age	0.39(0.12)	0.001	age	0.63(0.12)	<0.001
gender	0.12(0.04)	0.004	gender	0.32(0.05)	<0.001
CHD	0.07(0.12)	0.531	CHD	0.27(0.12)	0.027
smoking	0.21(0.05)	<0.001	smoking	0.22(0.05)	<0.001
physical activity	-0.05(0.03)	0.078	physical activity	-0.04(0.03)	0.223
education	-0.07(0.04)	0.065	education	-0.10(0.04)	0.007
BMI	-0.16(0.12)	0.197	BMI	-0.02(0.13)	0.861
SBP	1.04(0.44)	0.018	SBP	1.39(0.46)	0.002
BP med	0.16(0.04)	<0.001	BP med	0.23(0.05)	<0.001
glucose	0.12(0.05)	0.020	glucose	-0.02(0.05)	0.748
diabetes	0.06(0.06)	0.285	diabetes	0.00(0.06)	0.953
TC : HDL ratio	0.00(0.03)	0.936	TC : HDL ratio	0.16(0.03)	<0.001
lipid med	-0.03(0.21)	0.894	lipid med	-0.07(0.22)	0.744
LVH	0.21(0.10)	0.037	LVH	0.20(0.10)	0.051
eGFR	-2.82(0.37)	<0.001	eGFR	-4.76(0.37)	<0.001

X - 11308			X - 11564		
covariate	β (standard error)	p value	covariate	β (standard error)	p value
age	-0.29(0.12)	0.020	age	0.94(0.12)	<0.001
gender	0.17(0.04)	<0.001	gender	0.37(0.05)	<0.001
CHD	-0.09(0.12)	0.434	CHD	0.42(0.13)	0.001
smoking	-0.13(0.05)	0.007	smoking	0.18(0.05)	<0.001
physical activity	0.18(0.03)	<0.001	physical activity	-0.15(0.03)	<0.001
education	0.21(0.04)	<0.001	education	-0.09(0.04)	0.017
BMI	-0.71(0.13)	<0.001	BMI	0.15(0.13)	0.234
SBP	-2.02(0.45)	<0.001	SBP	2.36(0.45)	<0.001
BP med	-0.13(0.05)	0.003	BP med	0.26(0.05)	<0.001
glucose	-0.11(0.05)	0.049	glucose	-0.03(0.05)	0.599
diabetes	-0.07(0.06)	0.237	diabetes	0.02(0.06)	0.722
TC : HDL ratio	-0.02(0.03)	0.450	TC : HDL ratio	0.09(0.03)	0.006
lipid med	-0.02(0.22)	0.913	lipid med	-0.22(0.22)	0.330
LVH	-0.01(0.10)	0.960	LVH	0.27(0.10)	0.008
eGFR	-0.07(0.39)	0.866	eGFR	-5.10(0.37)	<0.001

Metabolite values are in the form of quartile ordinal scores; continuous variables used liner model (e.g, age= intercept+ β ×quartile scores); categorical variables used logistic model (e.g, logit(CHD)= quartile scores).

Abbreviations: CHD, coronary heart disease; BMI, body mass index; SBP, systolic blood pressure; BP med, anti-hypertensive medication use; TC : HDL ratio, serum total cholesterol to high-density-cholesterol lipoprotein ratio; lipid med, lipid-lowering medication use; LVH, left ventricular hypertrophy; eGFR, estimated glomerular filtration rate.

The β regression coefficient (standard error as well as the p value testing the association of the ordinal quartile score of each of the 16 metabolites identified in Model 3 and the covariates adjusted in Model 4 are presented in the above tables. Consistently, most of the metabolites were significantly related to eGFR (except for X-11308), while few of them were related to lipid-lowering medication use (except for X-11787 and X-11429). Their associations with the other established risk factors (covariates) vary among different metabolites.

Figure S1. Cumulative Hazards Curves of Incident Hospitalized Heart Failure across Quartiles of the Three Candidate Metabolites Identified in Model 4 among 1,744 African Americans in Jackson, MS field center of the Atherosclerosis Risk in Communities Study over the Period from 1987 to 2008.

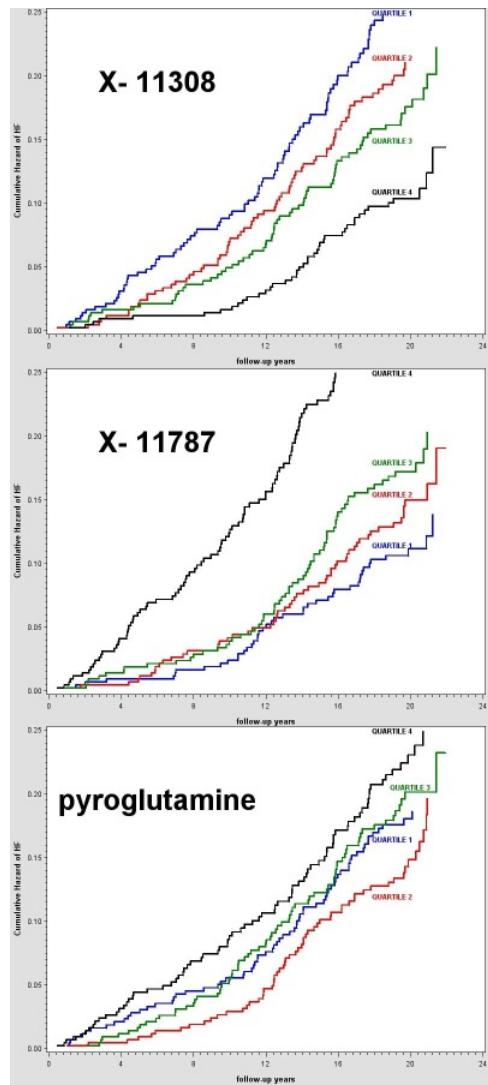


Figure S2. Heat map of Metabolite-Metabolite Correlations among All 187 Metabolites (Treated as Continuous Variables, Figure A) and among the 16 Metabolites Significant in Model 3 (Figure B) collected at 1986-1987 among 1,744 African Americans in Jackson, MS field center of the Atherosclerosis Risk in Communities Study. Metabolites were grouped by pathway and the three metabolites identified in Model 4 are labeled on the diagonal axis in Figure

A.

