

# Metabolic Signatures Elucidate the Effect of Body Mass Index on Type 2 Diabetes

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## Supplementary Material

### Testing of Assumptions for Multiple Linear Regression and Logistic Regression Model:

$\text{lm}(\text{BMI} \sim \text{metabolite} + \text{age} + \text{sex} + \text{physical activity} + \text{smoking} + \text{systolic blood pressure} + \text{HDL-C} + \text{triglyceride} + \text{fasting glucose})$

$\text{glm}(\text{T2D} \sim \text{metabolite} + \text{age} + \text{sex} + \text{BMI} + \text{physical activity} + \text{smoking} + \text{systolic blood pressure} + \text{HDL-C} + \text{triglyceride} + \text{fasting glucose})$

Basic assumptions of multiple linear regression and logistic regression models were tested for one member of each metabolite specie assuming that a member of one metabolite class sufficiently represented the entire class.

For each metabolite, a separate multiple linear regression model was fitted taking continuous BMI as dependent variable and each of the 146 metabolites as independent variable adjusted for the selected covariates. The basic assumptions of multiple linear regression were tested and satisfied. The diagnostic plots for linear regression were visually inspected. Assumptions of normal distribution of residuals and heteroskedasticity were deemed to be approximately satisfied. **The linearity graphs for representative metabolites are presented below as Figure S4-Figure S11.**

For the logistic regression model, binary diabetes status (yes/no) is the dependent variable and each of the 146 metabolites as an independent variable adjusted for the selected covariates. The basic assumptions of logistic regression were tested and satisfied. The assumption of linearity between continuous predictor variables and the logit of the outcome was tested by analyzing linearity graphs (**Figure S12-Figure S19**). All metabolites seemed to have an approximately linear relationship with the logit of the outcome i.e. Diabetes Status. The assumption of absence of influential points was assumed to have been fulfilled as the data were pre-processed and the natural log-transformed.

For both models, the assumption of absence of multicollinearity was tested using function “vif” of R package “car” employing again one member per class rule. This function gives values of Variance Inflation Factor (VIF) which is a measure of the magnitude of multicollinearity in regression analysis. Multicollinearity exists when there is a correlation between multiple independent variables in a multiple regression model. A value of 1 indicates no collinearity between an independent variable and other predictors of the model whereas infinity suggests perfect collinearity. As a general rule, a VIF value greater than 5 or 10 indicates a serious collinearity problem [1]. This assumption was also deemed to have been satisfied sufficiently as Variance Inflation Factor (VIF) values for nearly all of the predictor variables were close to 1 implying that there is no to very little multicollinearity. **VIF values for multicollinearity between predictors in both models are presented below (Table S16-Table S31).**

Abbreviations: BMI: Body Mass Index; HDL-C: High Density Lipoprotein Cholesterol;

### Diagnostic Plots and Tables for Multiple Linear Regression Model:

Figure S4: The linearity graphs for representative metabolites carnitine (C0) of acylcarnitines:

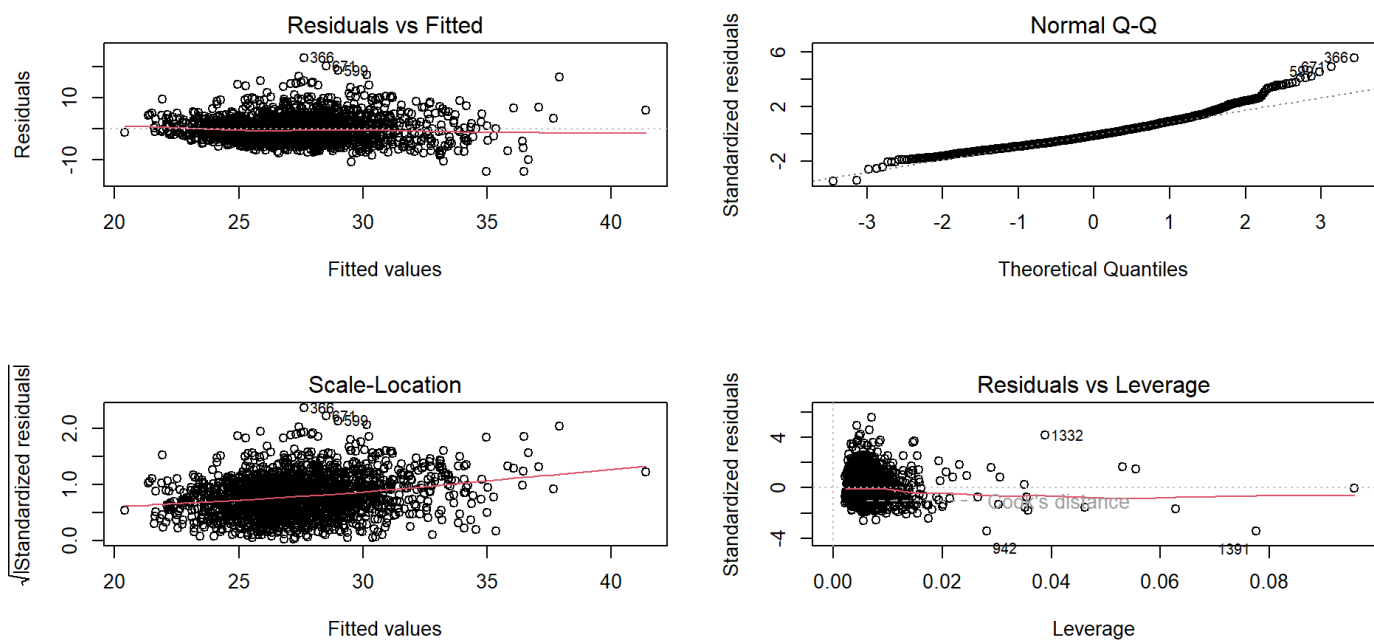


Table S16: VIF values for multicollinearity between predictors in the model for acylcarnitines-carnitine (C0)

Predictors	GVIF	Df	$GVIF^{1/(2 \cdot Df)}$
Carnitine	1.16	1	1.08
Age	1.23	1	1.11
Sex	1.39	1	1.18
Smoking	1.10	2	1.03
Physical activity	1.06	1	1.03
HDL-C	1.59	1	1.26
Systolic Blood Pressure	1.24	1	1.11
Triglycerides	1.40	1	1.18
Fasting serum glucose	1.26	1	1.12

Figure S5: The linearity graphs for representative metabolites lysoPC a C16:0 of lysophosphatidylcholines

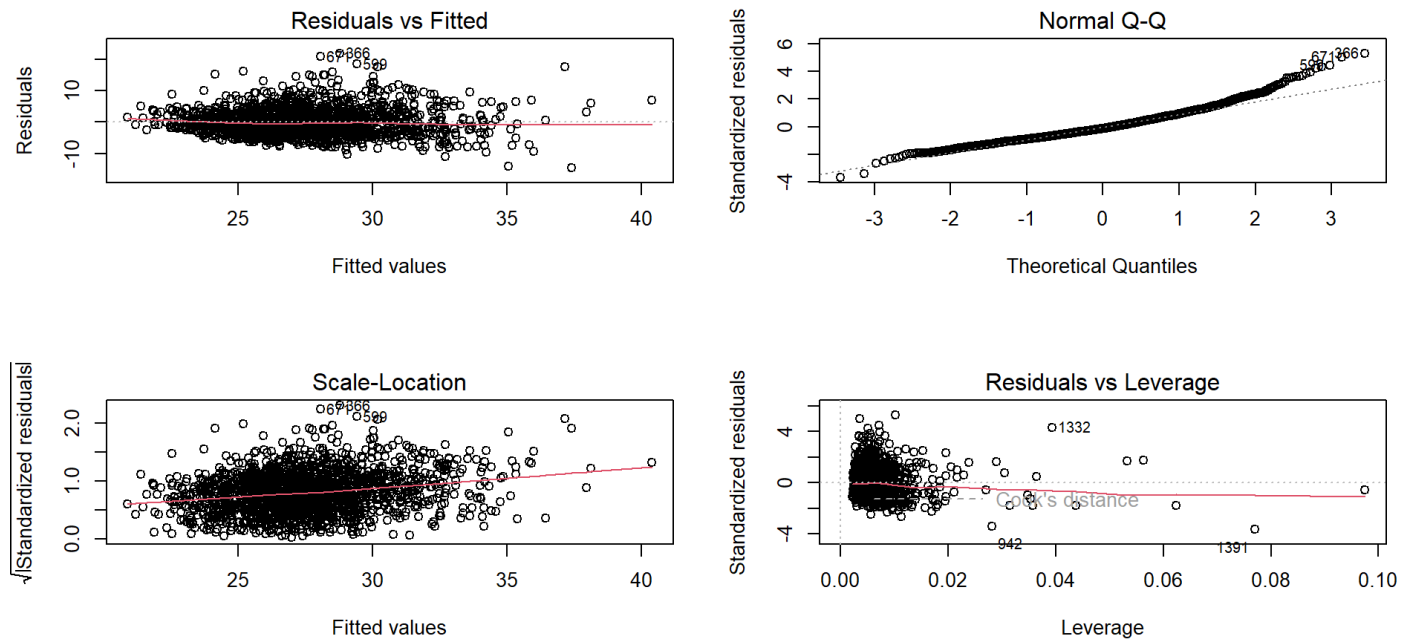


Table S17: VIF values for multicollinearity between predictors in the model for lysophosphatidylcholines - lysoPC a C16:0

Predictors	GVIF	Df	$GVIF^{1/(2 \cdot Df)}$
lysoPC a C16:0	1.12	1	1.06
Age	1.18	1	1.08
Sex	1.42	1	1.19
Smoking	1.09	2	1.02
Physical activity	1.06	1	1.03
HDL-C	1.61	1	1.27
Systolic Blood Pressure	1.24	1	1.11
Triglycerides	1.45	1	1.20
Fasting serum glucose	1.29	1	1.14

Figure S6: The linearity graphs for representative metabolites PC aa C28:1 of diacylphosphatidylcholines

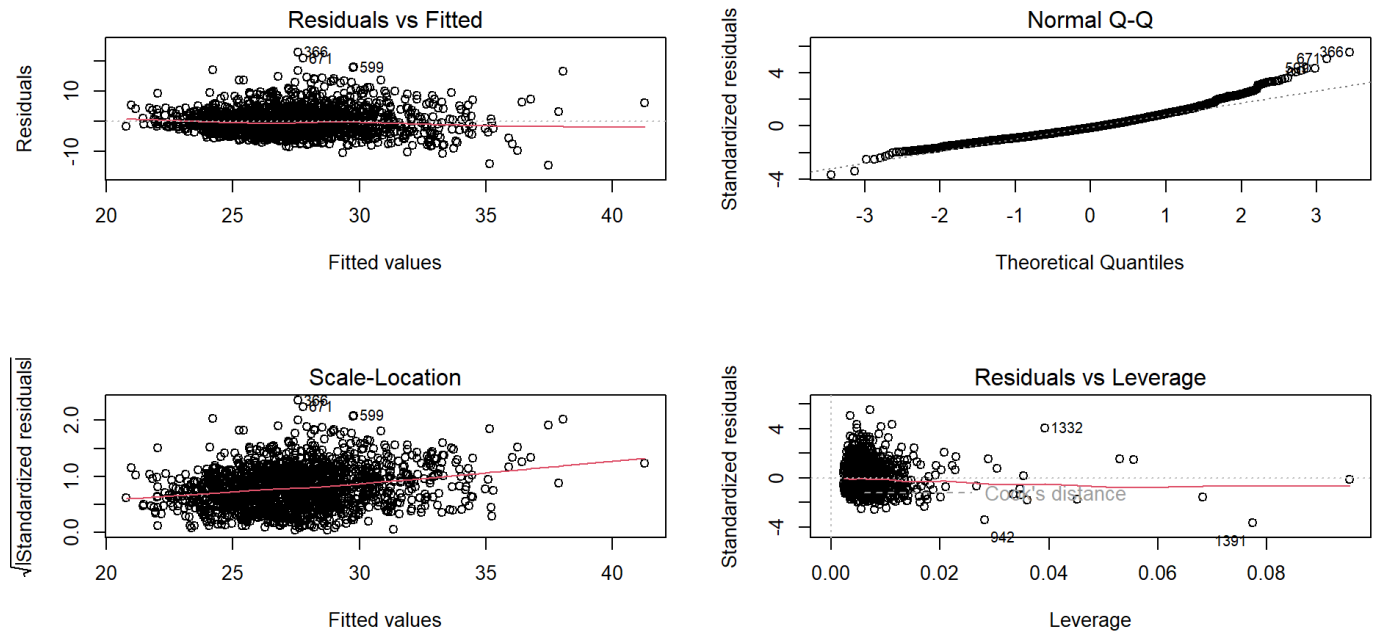


Table S18: VIF values for multicollinearity between predictors in the model for diacylphosphatidylcholines - PC aa C28:1

Predictors	GVIF	Df	GVIF <sup>1/(2*Df)</sup>
PC aa C28:1	1.26	1	1.12
Age	1.18	1	1.09
Sex	1.38	1	1.18
Smoking	1.11	2	1.03
Physical activity	1.06	1	1.03
HDL-C	1.72	1	1.31
Systolic Blood Pressure	1.24	1	1.11
Triglycerides	1.44	1	1.20
Fasting serum glucose	1.27	1	1.13

Figure S7: The linearity graphs for representative metabolites PC ae C30:0 of acylalkylphosphatidylcholine

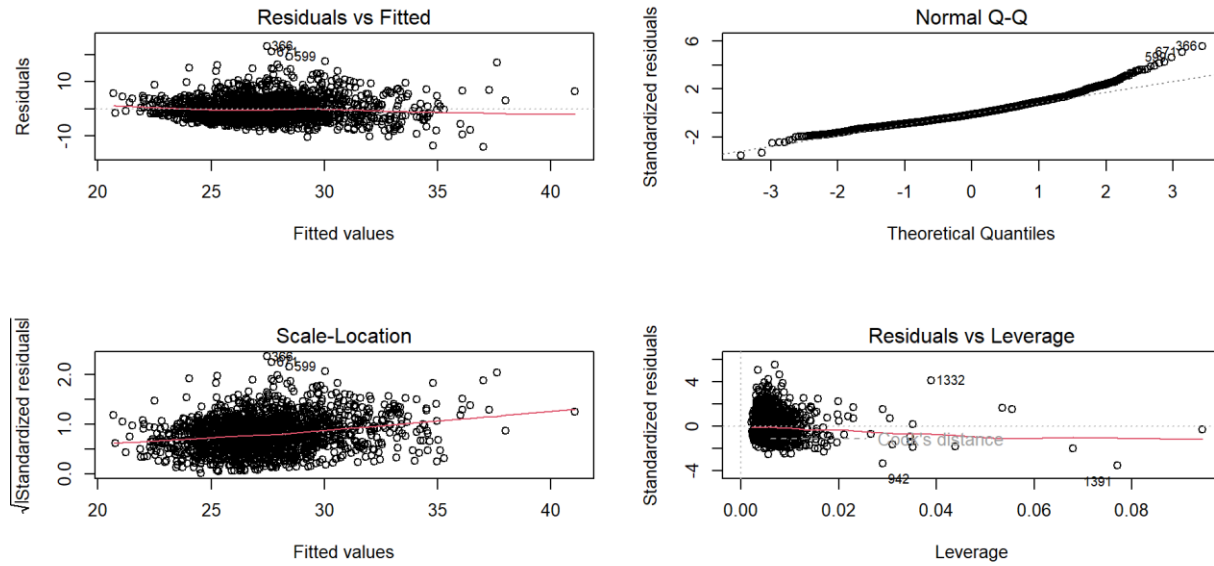


Table S19: VIF values for multicollinearity between predictors in the model for acylalkylphosphatidylcholine - PC ae C30:0

Predictors	GVIF	Df	GVIF <sup>1/(2*Df)</sup>
PC ae C30:0	1.24	1	1.11
Age	1.17	1	1.08
Sex	1.37	1	1.17
Smoking	1.11	2	1.03
Physical activity	1.06	1	1.03
HDL-C	1.68	1	1.30
Systolic Blood Pressure	1.24	1	1.11
Triglycerides	1.40	1	1.18
Fasting serum glucose	1.28	1	1.13

Figure S8: The linearity graphs for representative metabolite SM (OH) C14:1 of sphingomyelins

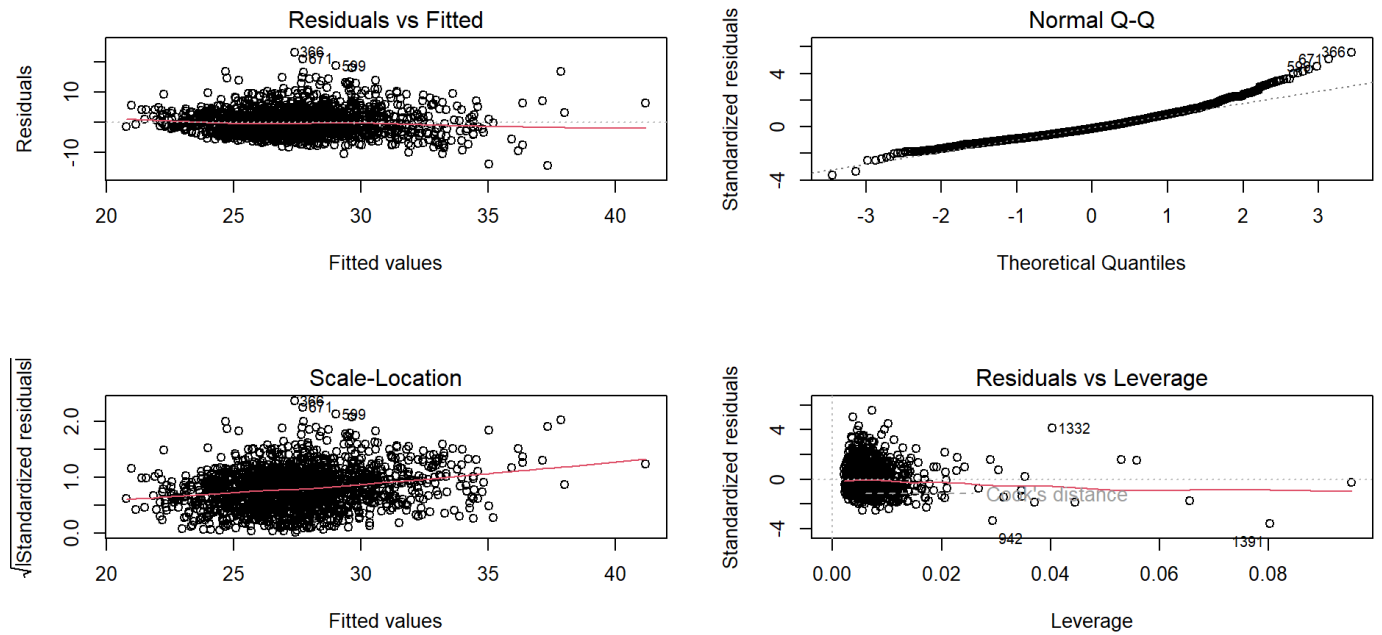


Table S20: VIF values for multicollinearity between predictors in the model for sphingomyelins - SM (OH) C14:1

Predictors	GVIF	Df	$GVIF^{1/(2 \cdot Df)}$
SM (OH) C14:1	1.31	1	1.15
Age	1.20	1	1.09
Sex	1.41	1	1.19
Smoking	1.12	2	1.03
Physical activity	1.06	1	1.03
HDL-C	1.62	1	1.27
Systolic Blood Pressure	1.24	1	1.11
Triglycerides	1.39	1	1.18
Fasting serum glucose	1.28	1	1.13

Figure S9: The linearity graphs for representative metabolite Hexoses (H1)

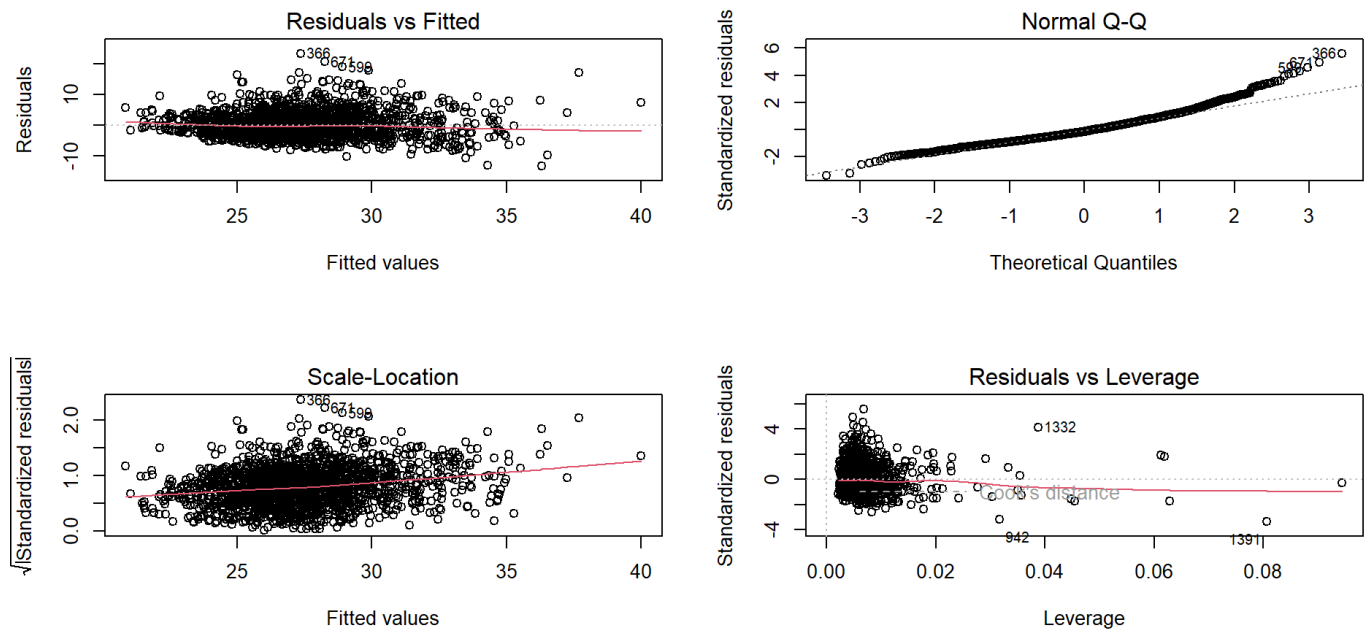


Table S21: VIF values for multicollinearity between predictors in the model for Hexoses (H1)

Predictors	GVIF	Df	GVIF <sup>1/(2*Df)</sup>
Hexoses	3.92	1	1.98
Age	1.18	1	1.09
Sex	1.35	1	1.16
Smoking	1.10	2	1.02
Physical activity	1.06	1	1.03
HDL-C	1.60	1	1.26
Systolic Blood Pressure	1.24	1	1.11
Triglycerides	1.39	1	1.18
Fasting serum glucose	3.94	1	1.98



Figure S10: The linearity graphs for representative metabolite alanine (Ala) for amino acids

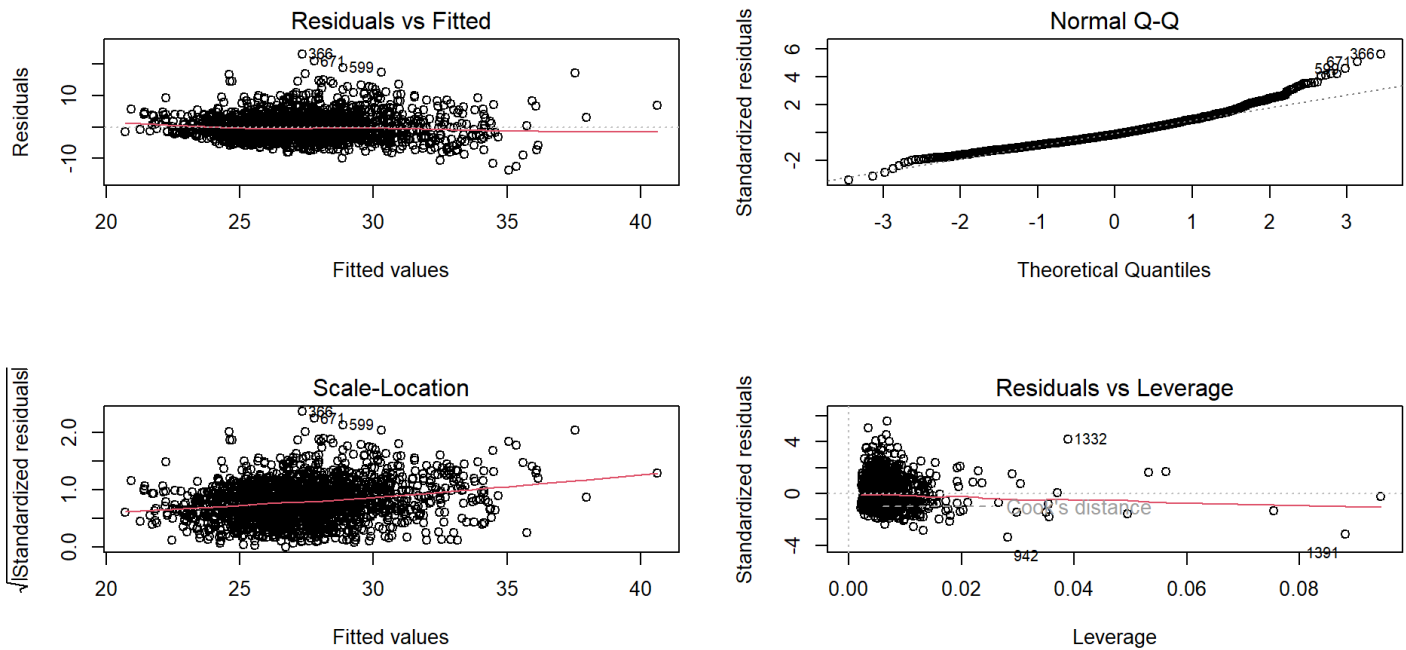


Table S22: VIF values for multicollinearity between predictors in the model for amino acids - alanine (Ala)

Predictors	GVIF	Df	GVIF <sup>1/(2*Df)</sup>
Alanine	1.19	1	1.09
Age	1.19	1	1.09
Sex	1.34	1	1.16
Smoking	1.10	2	1.02
Physical activity	1.06	1	1.03
HDL-C	1.59	1	1.26
Systolic Blood Pressure	1.24	1	1.11
Triglycerides	1.41	1	1.19
Fasting serum glucose	1.32	1	1.15

Figure S11: The linearity graphs for representative metabolite acetylnornithine (Ac-Orn) for biogenic amines

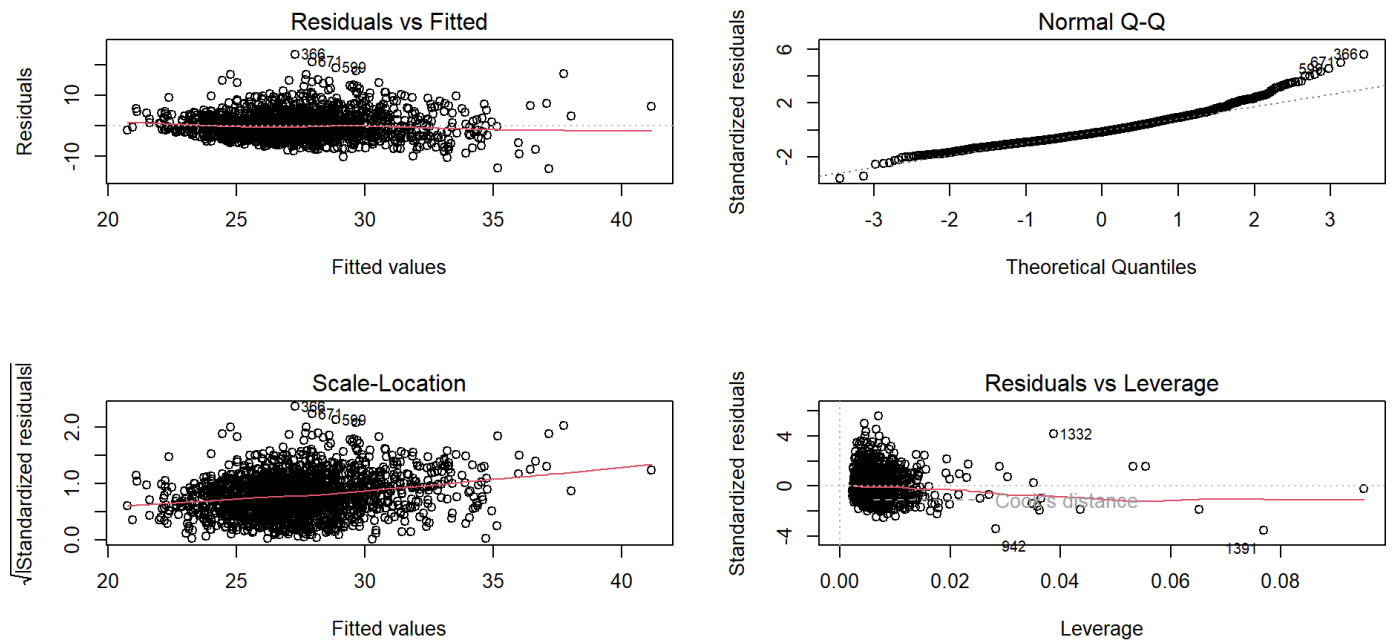


Table S23: VIF values for multicollinearity between predictors in the model for biogenic amines - acetylnornithine

Predictors	GVIF	Df	$GVIF^{1/(2 \cdot Df)}$
Acetylnornithine	1.06	1	1.03
Age	1.23	1	1.11
Sex	1.35	1	1.16
Smoking	1.10	2	1.02
Physical activity	1.06	1	1.03
HDL-C	1.58	1	1.26
Systolic Blood Pressure	1.24	1	1.11
Triglycerides	1.39	1	1.18
Fasting serum glucose	1.26	1	1.12

### Diagnostic Plots and Tables for Multiple Linear Regression Model:

Figure S12: The linearity graphs for representative metabolite carnitine (C0) for acylcarnitines

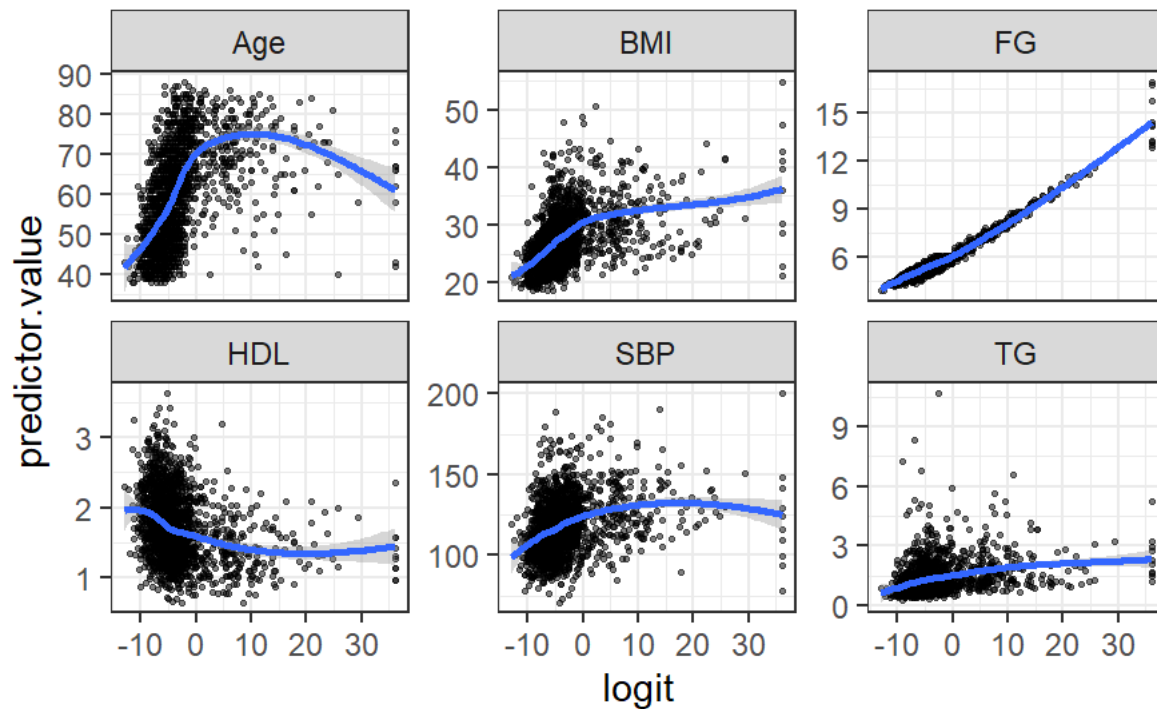


Table S24: VIF values for multicollinearity between predictors in the model for acylcarnitines - carnitine (C0)

Predictors	GVIF	Df	$GVIF^{1/(2 \cdot Df)}$
Carnitine	1.13	1	1.06
Age	1.15	1	1.07
Sex	1.52	1	1.23
BMI	1.19	1	1.09
Smoking	1.37	2	1.08
Physical activity	1.07	1	1.04
HDL-C	1.70	1	1.30
Systolic Blood Pressure	1.13	1	1.06
Triglycerides	1.39	1	1.18
Fasting serum glucose	1.10	1	1.05

Figure S13: The linearity graphs for representative metabolite lysoPC a C16:0 for lysophosphatidylcholines

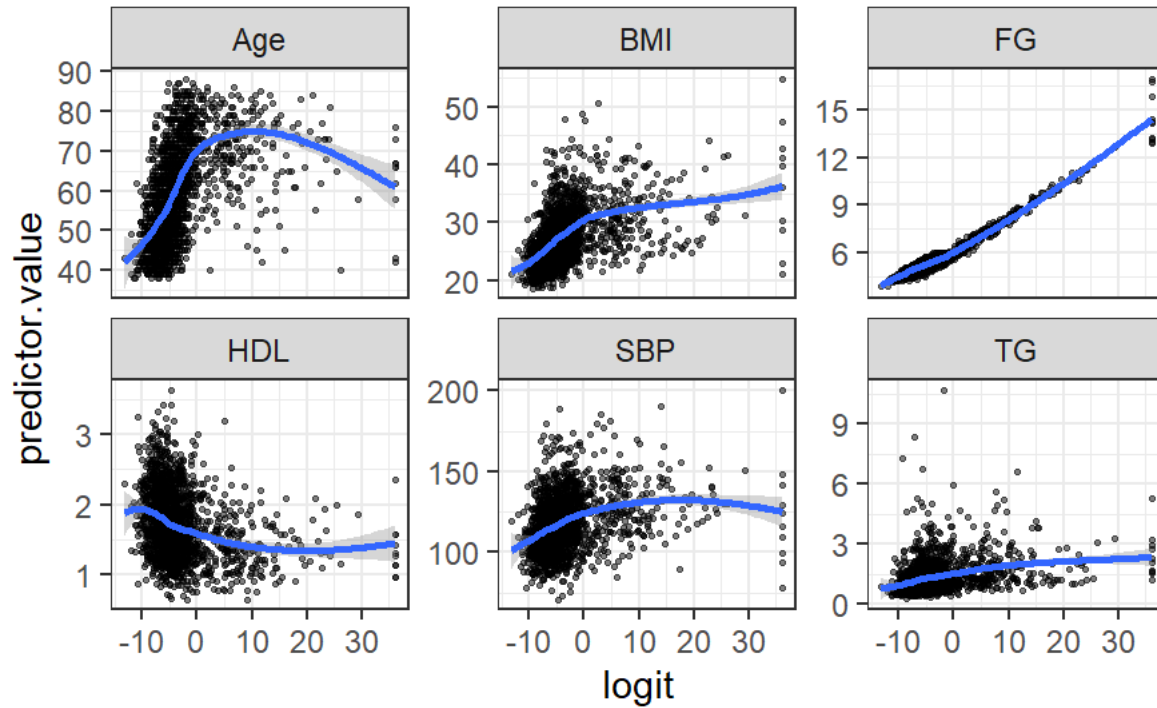


Table S25: VIF values for multicollinearity between predictors in the model for lysophosphatidylcholines - lysoPC a C16:0

Predictors	GVIF	Df	GVIF <sup>1/(2*Df)</sup>
lysoPC a C16:0	1.15	1	1.07
Age	1.14	1	1.07
Sex	1.53	1	1.24
BMI	1.22	1	1.10
Smoking	1.38	2	1.08
Physical activity	1.07	1	1.03
HDL-C	1.71	1	1.31
Systolic Blood Pressure	1.12	1	1.06
Triglycerides	1.43	1	1.20
Fasting serum glucose	1.10	1	1.05

Figure S14: The linearity graphs for representative metabolite PC aa C28:1 for diacylphosphatidylcholines

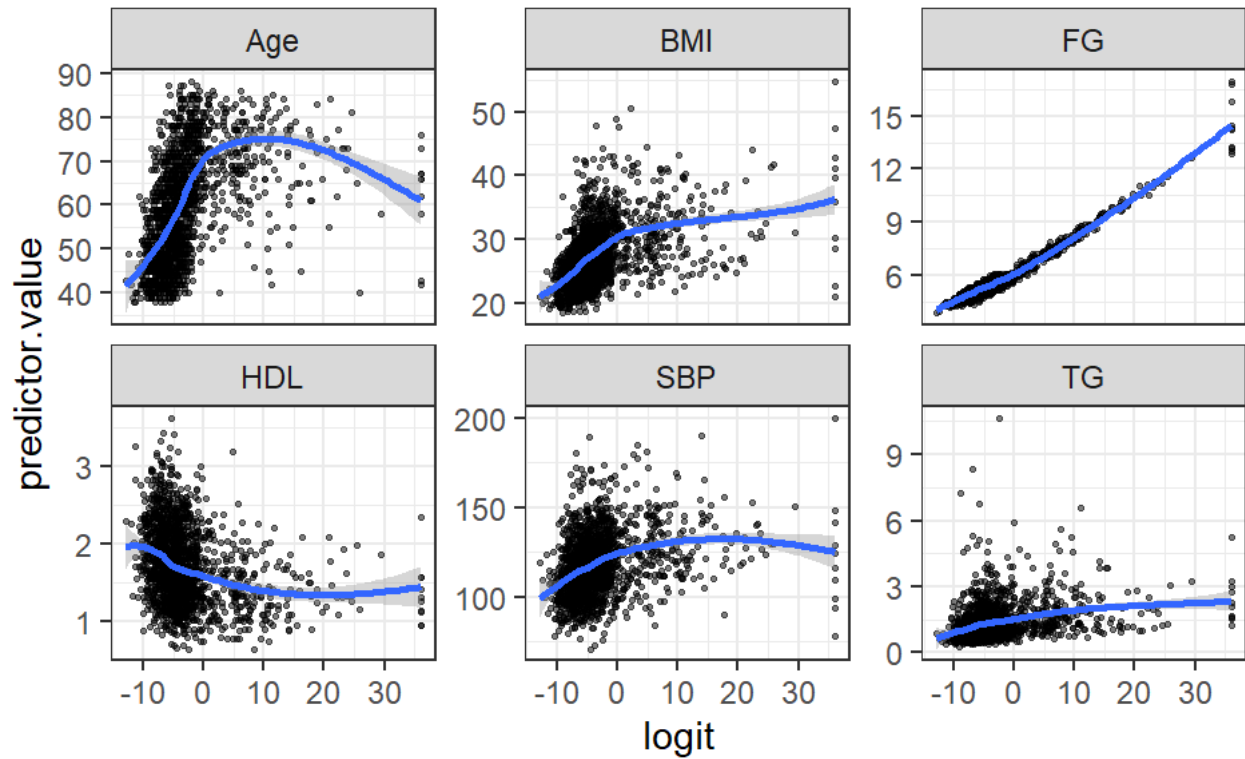


Table S26: VIF values for multicollinearity between predictors in the model for diacylphosphatidylcholines - PC aa C28:1

Predictors	GVIF	Df	GVIF <sup>1/(2*Df)</sup>
PC aa C28:1	1.25	1	1.12
Age	1.14	1	1.07
Sex	1.54	1	1.24
BMI	1.18	1	1.09
Smoking	1.38	2	1.08
Physical activity	1.07	1	1.03
HDL-C	1.80	1	1.34
Systolic Blood Pressure	1.12	1	1.06
Triglycerides	1.43	1	1.19
Fasting serum glucose	1.10	1	1.05

Figure S15: The linearity graphs for representative metabolite PC ae C30:0 for acylalkylphosphatidylcholine

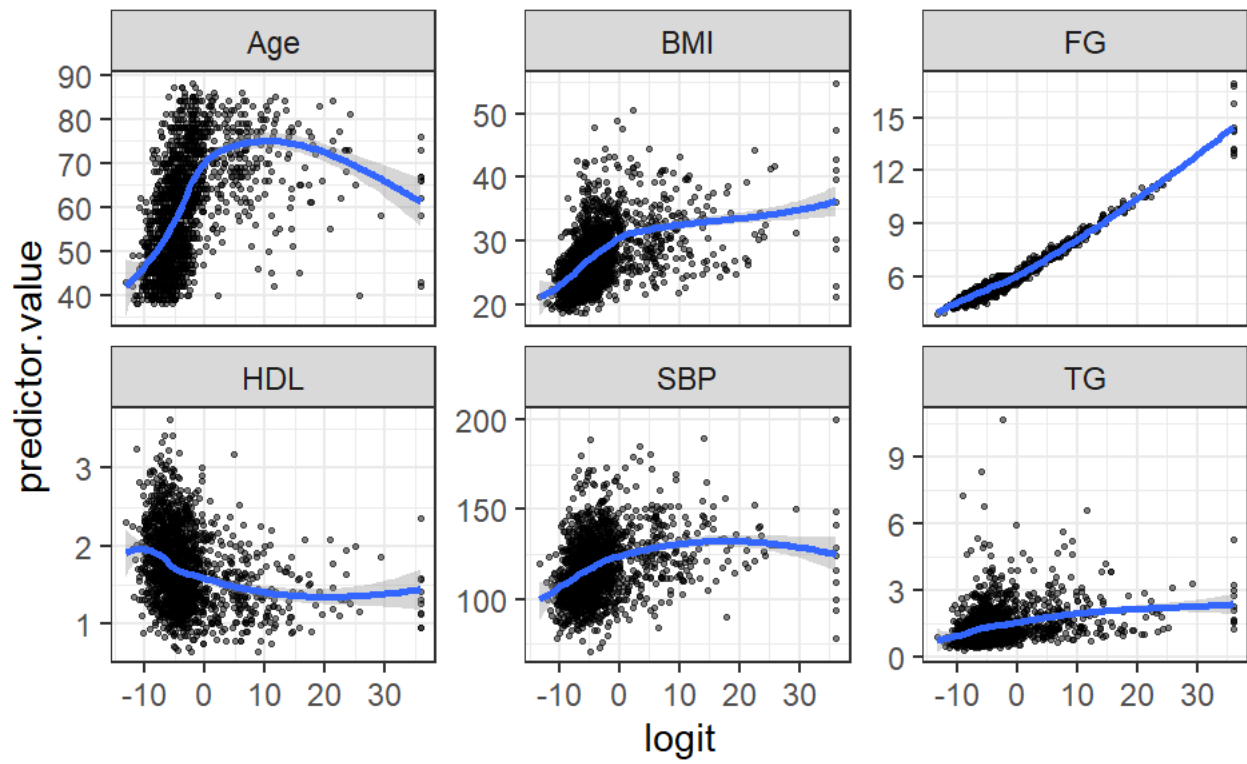


Table S27: VIF values for multicollinearity between predictors in the model for acylalkylphosphatidylcholine - PC ae C30:0

Predictors	GVIF	Df	GVIF <sup>1/(2*Df)</sup>
PC ae C30:0	1.18	1	1.09
Age	1.16	1	1.08
Sex	1.48	1	1.22
BMI	1.19	1	1.09
Smoking	1.42	2	1.09
Physical activity	1.07	1	1.03
HDL-C	1.74	1	1.32
Systolic Blood Pressure	1.12	1	1.06
Triglycerides	1.39	1	1.18
Fasting serum glucose	1.11	1	1.05

Figure S16: The linearity graphs for representative metabolite SM (OH) C14:1 for sphingomyelins

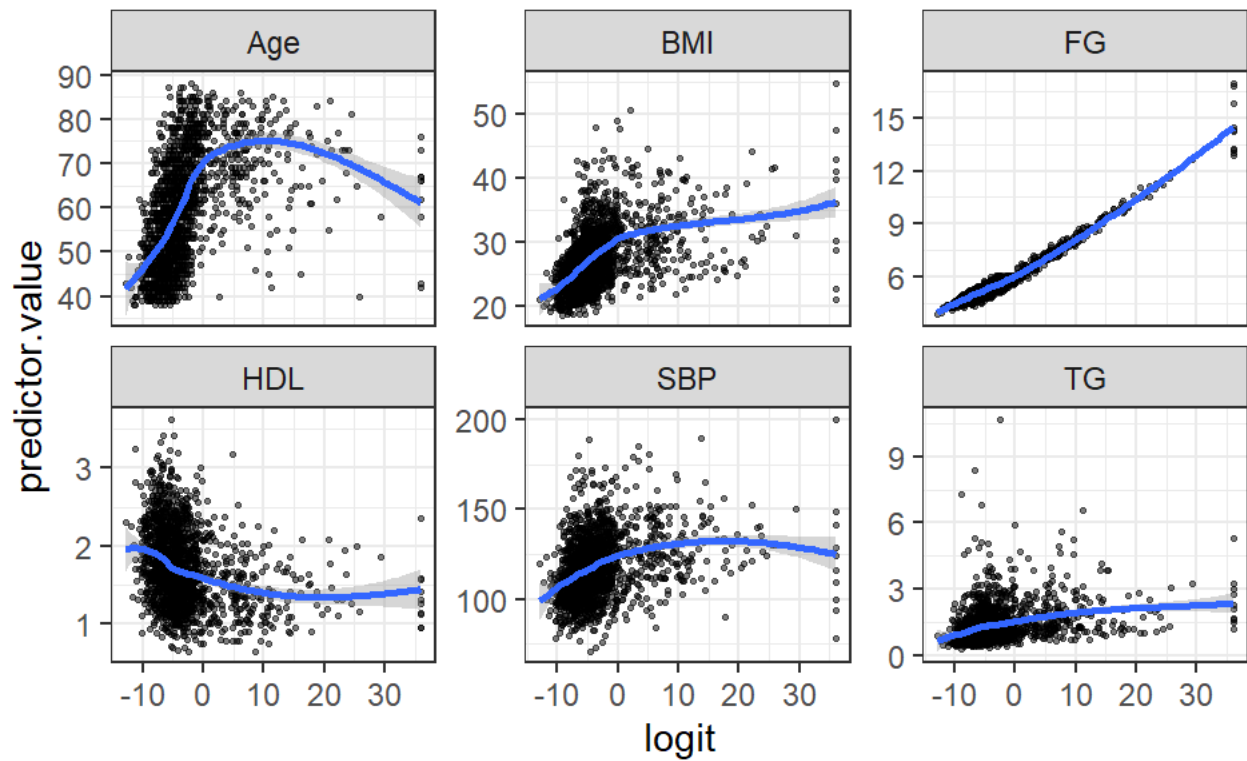


Table S28: VIF values for multicollinearity between predictors in the model for sphingomyelins - SM (OH) C14:1

Predictors	GVIF	Df	$GVIF^{1/(2 \cdot Df)}$
SM (OH) C14:1	1.28	1	1.13
Age	1.20	1	1.10
Sex	1.47	1	1.21
Smoking	1.12	2	1.03
Physical activity	1.04	1	1.02
HDL-C	1.58	1	1.26
Systolic Blood Pressure	1.22	1	1.10
Triglycerides	1.33	1	1.15
Fasting serum glucose	1.25	1	1.12

Figure S17: The linearity graphs for representative metabolite Hexoses (H1)

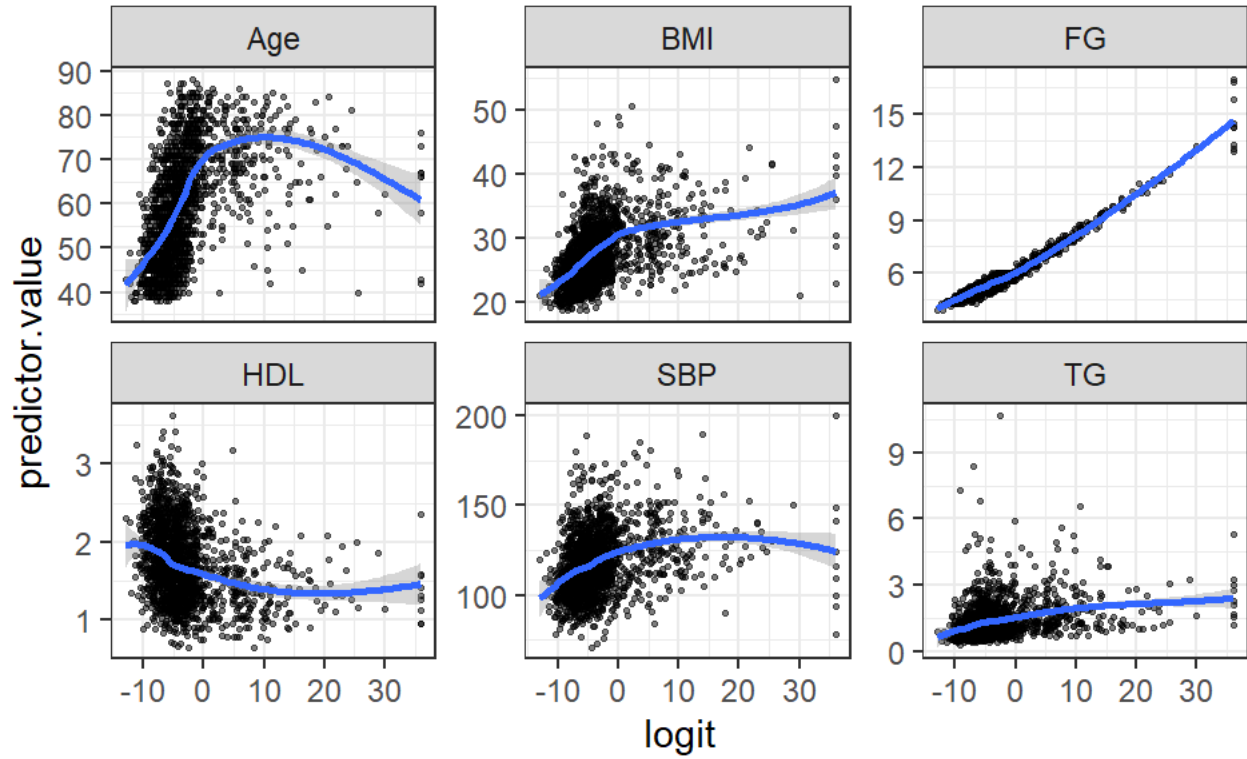


Table S29: VIF values for multicollinearity between predictors in the model for Hexoses (H1)

Predictors	GVIF	Df	GVIF <sup>1/(2*Df)</sup>
Hexoses	1.34	1	1.16
Age	1.14	1	1.07
Sex	1.47	1	1.21
BMI	1.18	1	1.09
Smoking	1.37	2	1.08
Physical activity	1.07	1	1.03
HDL-C	1.71	1	1.31
Systolic Blood Pressure	1.12	1	1.06
Triglycerides	1.38	1	1.17
Fasting serum glucose	1.40	1	1.18



Figure S18: The linearity graphs for representative metabolite alanine (Ala) of amino acids

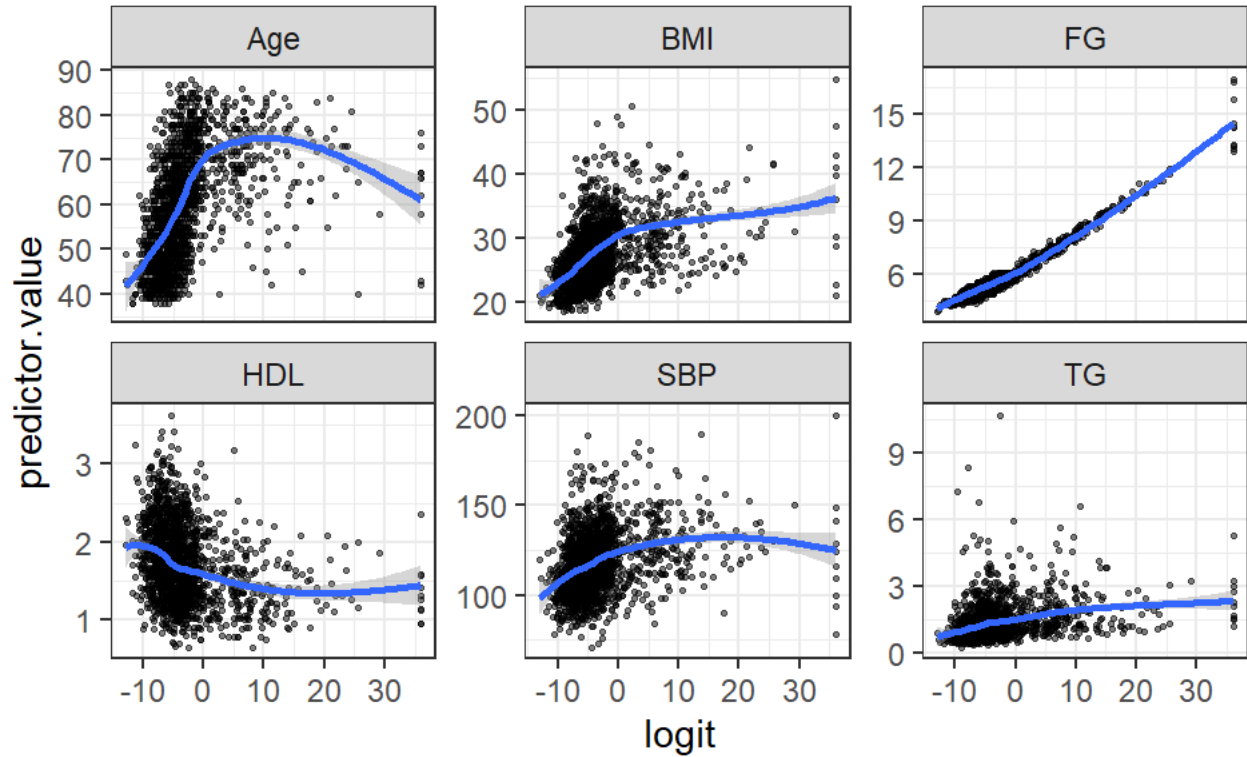


Table S30: VIF values for multicollinearity between predictors in the model for amino acids - alanine (Ala)

Predictors	GVIF	Df	$GVIF^{1/(2*Df)}$
Alanine	1.11	1	1.05
Age	1.15	1	1.07
Sex	1.47	1	1.21
BMI	1.18	1	1.09
Smoking	1.39	2	1.09
Physical activity	1.07	1	1.03
HDL-C	1.70	1	1.31
Systolic Blood Pressure	1.12	1	1.06
Triglycerides	1.43	1	1.20
Fasting serum glucose	1.10	1	1.05

Figure S19: The linearity graphs for representative metabolite acetylmethionine (Ac-Met) of biogenic amines

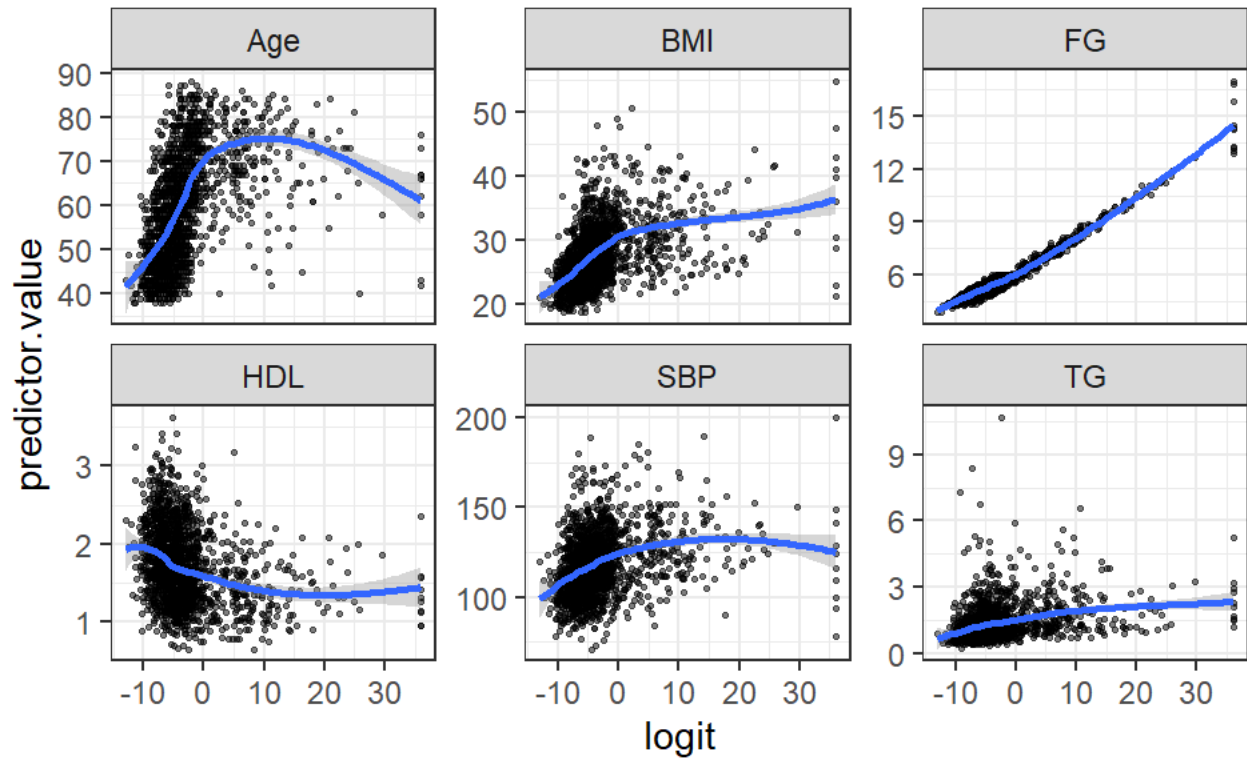


Table S31: VIF values for multicollinearity between predictors in the model for biogenic amines - acetylmethionine

Predictors	GVIF	Df	$GVIF^{1/(2 \cdot Df)}$
Acetylmethionine	1.08	1	1.04
Age	1.16	1	1.08
Sex	1.49	1	1.22
BMI	1.18	1	1.09
Smoking	1.37	2	1.08
Physical activity	1.07	1	1.03
HDL-C	1.70	1	1.30
Systolic Blood Pressure	1.13	1	1.06
Triglycerides	1.39	1	1.18
Fasting serum glucose	1.10	1	1.05

## Reference

1. James, G.; James, G.; Witten, D.; Hastie, T.; Tibshirani, R.; Ebscohost. *An introduction to statistical learning: with applications in R*, Uncorrected edition. ed.; Springer: New York, 2013.