

Hierarchical Bayesian Model Selection

The Role of Nutritional Supplementation on Betaine Concentration in Pregnant Pakistani Women

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Introduction

- Hierarchical Bayesian Models** occur when the parameter(s) of a prior distribution has/have a prior distribution
 - Common in correlated responses
- Betaine** has been connected to pregnancy outcomes such as infant length at birth.
 - Nutritional supplements for malnourished pregnant women may have an impact on the concentration of Betaine in the mother after 34 weeks gestation.

The Data

- 31 pregnant Pakistani women
 - 13 on NS since preconception (Arm 1)
 - 18 on NS since 12 Weeks gestation (Arm 2)
- Women from 13 different locations in Pakistan (clusters, denoted with subscript i)
 - Implies **correlated responses**

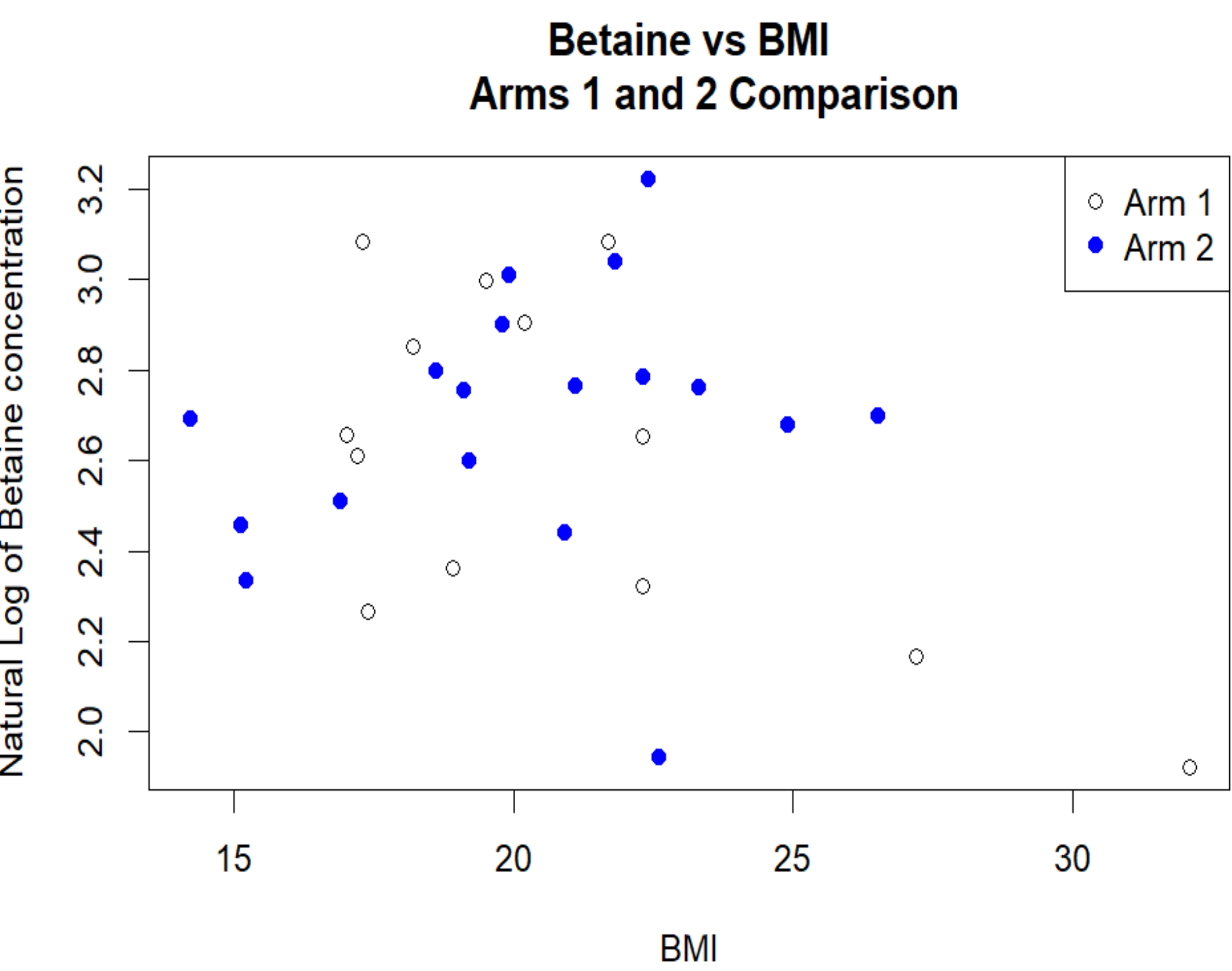


Figure 1. Betaine Scatterplot by BMI: The natural log of betaine concentration for the observed Pakistani women. Values from Arm 1 and Arm 2 are signified by color (white and blue, respectively). Future work will consider the role of BMI (the x-axis).

Acknowledgments

- Dr. Joshua French and his course MATH 7393
- Dr. Audrey Hendricks, Dr. Stephanie Gilley, and Dr. Sarah Borengasser for guiding this study
- The Women First Research Group led by Dr. Michael Hambidge and Dr. Nancy Krebs for providing the data and producing the overall study

Model Designs

$Y_{i,j}$: Betaine concentration for the j^{th} person from the i^{th} cluster.
 β_0 : **parameter** for the mean metabolite concentration of Women from Arm 2

β_1 : **parameter** for the difference in mean metabolite concentration between Arm1 and Arm 2 Women

Model with No Random Effect (NRE)

Sampling Distribution:

$$\ln(Y_{i,j}) \sim N(\mu_{i,j}, \sigma^2), \mu_{i,j} = \beta_0 + \beta_1 X_{i,j} + \beta_i$$

Prior Distributions:

$$\beta_k \sim N(0, 1,000), k = 0, 1, \dots, 13$$

$$\sigma^2 \sim IG(0.001, 0.001)$$

Model with Random Effect for Site (RES)

Sampling Distribution:

$$\ln(Y_{i,j}) \sim N(\mu_{i,j} + \alpha_i, \sigma_0^2), \mu_{i,j} = \beta_0 + \beta_1 X_{i,j}$$

Prior Distributions:

$$\beta_k \sim N(0, 1,000), k = 0, 1$$

$$\alpha_i \sim N(0, \sigma_1^2)$$

$$\sigma_l^2 \sim IG(0.001, 0.001), l = 0, 1$$

Model with Random Effect for Site and Woman (RESW)

Sampling Distribution:

$$\ln(Y_{i,j,k}) \sim N(\mu_{i,j} + \alpha_i + \gamma_k, \sigma_0^2), \mu_{i,j} = \beta_0 + \beta_1 X_{i,j}$$

Prior Distributions:

$$\beta_m \sim N(0, 1,000), m = 0, 1$$

$$\alpha_i \sim N(0, \sigma_1^2)$$

$$\gamma_k \sim N(0, \sigma_2^2)$$

$$\sigma_l^2 \sim IG(0.001, 0.001), l = 0, 1, 2$$

Where k is the k^{th} observation

Model Selection

Model	LOOIC
NRE	43.382
RES	23.951
RESW	11.678

Table 2. Selection by LOOIC: Leave-one-out-information-criterion (LOOIC) recommends the model with the smallest value (RESW).

Results

Parameter	Posterior Mean	95% CPI
β_1	-0.106	(-0.349, 0.138)

Table 3. Arm Parameter Results: The posterior mean and 95% central posterior interval for the distribution of the β_1 parameter. Results are from the RESW model, as determined through model selection.

Model Analysis

Model	Variance Breakdown	Mean	SD	2.5 Percentile	97.5 Percentile
NRE	σ^2	0.1094	0.0428	0.0544609	0.2171397
	Total Variance	0.1094	0.0428	0.0544609	0.2171397
RES	σ^2	0.0993	0.0316	0.0535146	0.1758947
	σ_1^2	0.0219	0.0259	0.0008010	0.0903016
RESW	Total Variance	0.1212	0.0375	0.0696655	0.2118633
	σ^2	0.0510	0.0431	0.0009505	0.1449823
	σ_1^2	0.0216	0.0260	0.0007846	0.0897476
	σ_2^2	0.0508	0.0431	0.0008985	0.1452992
	Total Variance	0.1234	0.0384	0.0706971	0.2162895

Table 1. Random Effects: Posterior values for distributions of the variance components for each model.

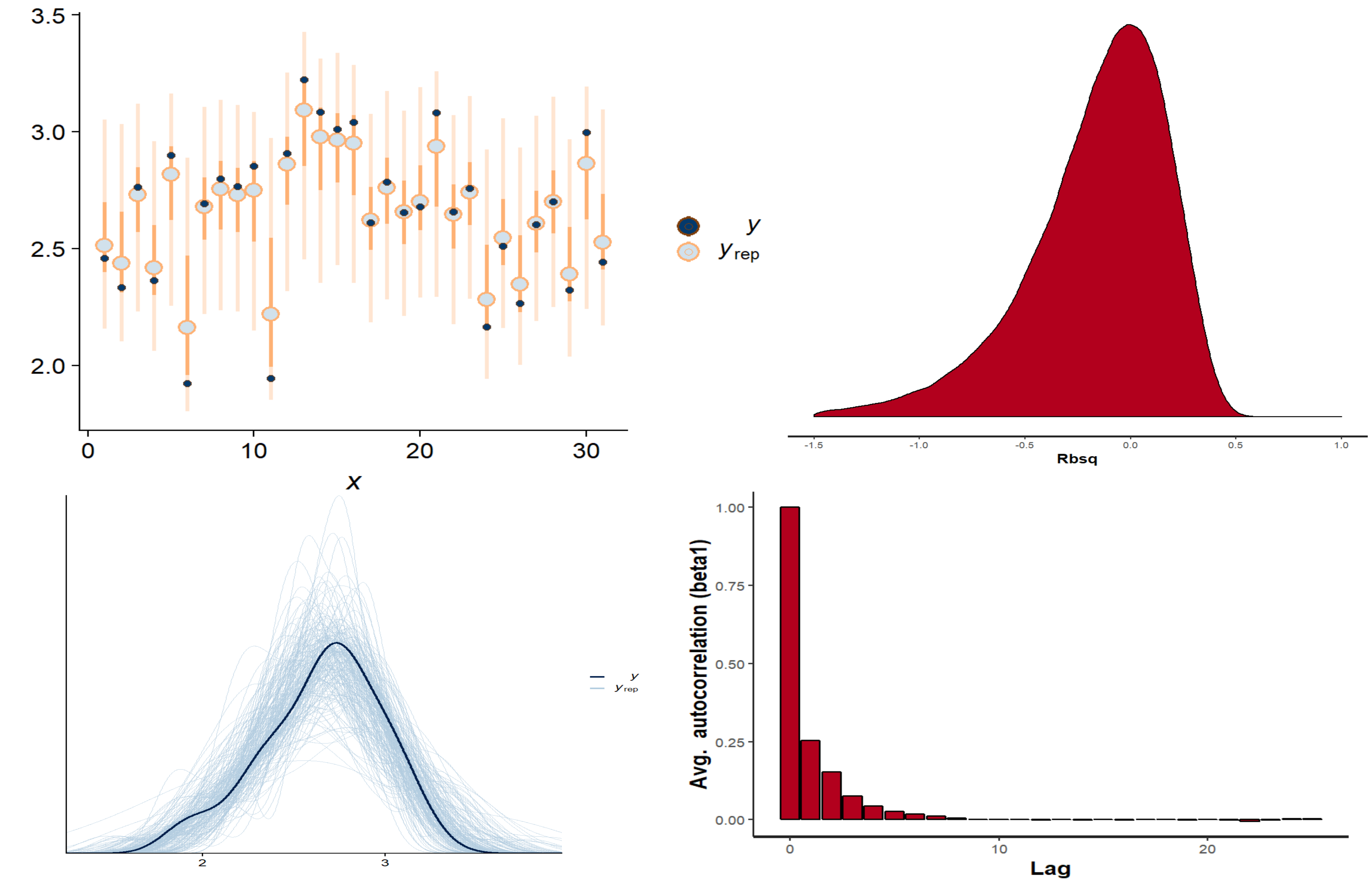


Figure 3. Model verification: (TL) 50% (dark orange), 90% (light orange) bands and mean estimate (gray circle) for the replicated response value at each observed response (dark blue circle). Calculated from 10,000 replicated data sets. (TR) Density of the Bayesian R-squared value for the RESW model with mean estimate - 0.171. (BL) Overlay of 200 replicated data densities generated from the posterior distributions of the RESW model. (BR) Average autocorrelation for β_1 in RESW model.

Conclusion/Future Work

- Arm 1 and Arm 2 do not appear to have significant differences in terms of metabolic concentration among the Pakistani women.
- The final model (RESW) has poor predictive ability
- Consider the interaction of BMI and Arm type to potentially strengthen the model