TABLE S6: Complete adjusted negative binomial models stratified by HDP status with alleles treated linearly for trend

|  | Coefficient | Standard Error | Z Value | p-value | CR (95% CI) |
|---|---|---|---|---|---|
| T Alleles | 0.041 | 0.151 | 0.270 | 0.787 | 1.042 (0.775, 1.400) |
| AFB | 0.022 | 0.021 | 1.031 | 0.303 | 1.022 (0.981, 1.065) |
| Age | -0.043 | 0.010 | -4.308 | 0.000 | 0.958 (0.940, 0.977) |
| Family History | 0.436 | 0.228 | 1.911 | 0.056 | 1.547 (0.989, 2.419) |
| BMI | -0.020 | 0.014 | -1.348 | 0.178 | 0.981 (0.953, 1.009) |
| Parity | 0.402 | 0.131 | 3.071 | 0.002 | 1.495 (1.157, 1.933) |
| Menarche | -0.039 | 0.072 | -0.550 | 0.582 | 0.961 (0.836, 1.106) |

For HDP- women only. T alleles are treated linearly. The reference for T alleles is no T alleles (T alleles = 0). The reference for family history is no family history.