

TABLE S4: Complete adjusted negative binomial models stratified by HDP status

	Coefficient	Standard Error	Z Value	p-value	IRR (95% CI)
(Intercept)	1.296	0.493	2.629	0.009	3.653 (1.390, 9.598)
T Alleles = 1	0.013	0.260	0.052	0.959	1.014 (0.608, 1.689)
T Alleles = 2	0.080	0.302	0.264	0.792	1.083 (0.600, 1.955)
AFB	0.022	0.021	1.022	0.307	1.022 (0.980, 1.065)
Age	-0.043	0.010	-4.302	0.000	0.958 (0.940, 0.977)
Family History	0.441	0.229	1.924	0.054	1.554 (0.992, 2.435)
BMI	-0.020	0.014	-1.369	0.171	0.98 (0.953, 1.009)
Parity	0.398	0.131	3.028	0.002	1.489 (1.151, 1.927)
Menarche	-0.041	0.073	-0.556	0.578	0.96 (0.831, 1.109)

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