

TABLE S5: Complete adjusted negative binomial models stratified by HDP status with alleles treated linearly for trend

	Coefficient	Standard Error	Z Value	p-value	IRR (95% CI)
(Intercept)	4.047	0.749	5.401	0.000	57.204 (13.174, 248.392)
T Alleles	-0.380	0.194	-1.960	0.050	0.684 (0.468, 1.000)
AFB	-0.046	0.028	-1.645	0.100	0.955 (0.904, 1.009)
Age	-0.022	0.013	-1.767	0.077	0.978 (0.954, 1.002)
Family History	0.103	0.300	0.342	0.733	1.108 (0.615, 1.996)
BMI	-0.028	0.018	-1.560	0.119	0.973 (0.939, 1.007)
Parity	-0.032	0.178	-0.182	0.856	0.968 (0.684, 1.371)
Menarche	-0.220	0.093	-2.375	0.018	0.802 (0.669, 0.962)

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.