

TABLE S4: Complete adjusted negative binomial models stratified by HDP status with alleles treated linearly for trend

For HDP+ women only					
	Coefficient	Standard Error	Z Value	p-value	CR (95% CI)
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					
	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)

T alleles are treated linearly. The reference for T alleles is no T alleles. The reference for family history is no family history.