TABLE S3: Complete adjusted negative binomial models stratified by HDP status

	For HDP+ women only				
	Coefficient	Standard Error	Z Value	p-value	CR (95% CI)
T Alleles = 1	-0.500	0.326	-1.536	0.124	0.606 (0.320, 1.148)
T Alleles = 2	-0.760	0.386	-1.967	0.049	0.468 (0.219, 0.997)
AFB	-0.045	0.028	-1.628	0.104	0.956 (0.905, 1.009)
Age	-0.023	0.013	-1.793	0.073	0.978 (0.954, 1.002)
Family History	0.071	0.303	0.235	0.814	1.074 (0.593, 1.946)
BMI	-0.027	0.018	-1.522	0.128	0.973 (0.940, 1.008)
Parity	-0.048	0.179	-0.267	0.790	0.953 (0.671, 1.354)
Menarche	-0.224	0.095	-2.360	0.018	0.799 (0.663, 0.963)

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

T alleles are treated as factor variables. The reference for T alleles is no T alleles. The reference for family history is no family history.