TABLE S2: Complete adjusted negative binomial model with interaction terms

	Coefficient	Standard Error	Z Value	p-value	IRR (95% CI)
(Intercept)	1.849	0.494	3.741	0.000	6.352 (2.411, 16.735)
HDP	0.208	0.345	0.604	0.546	1.231 (0.627, 2.419)
T Alleles = 1	-0.027	0.268	-0.102	0.918	0.973 (0.575, 1.645)
T Alleles = 2	0.099	0.307	0.322	0.747	1.104 (0.605, 2.016)
AFB	0.008	0.017	0.498	0.619	1.008 (0.975, 1.043)
Age	-0.041	0.008	-5.232	0.000	0.96 (0.945, 0.975)
Family History	0.372	0.185	2.008	0.045	1.451 (1.009, 2.086)
BMI	-0.022	0.011	-1.918	0.055	0.979 (0.957, 1.000)
Parity	0.334	0.107	3.132	0.002	1.396 (1.133, 1.720)
Menarche	-0.102	0.058	-1.763	0.078	0.903 (0.806, 1.011)
HDP×T alleles = 1	-0.310	0.417	-0.743	0.457	0.734 (0.324, 1.661)
HDP×T alleles = 2	-0.740	0.482	-1.536	0.124	0.477 (0.185, 1.227)

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