TABLE 2: Results of adjusted negative binomial model with interaction terms.

	Coefficient	Standard Error	Z Value	p-value	CR (95% CI)
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)
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)

HDP compares HDP-positive women to HDP-negative women. T alleles are treated as a factor variable. The reference for T alleles is no T alleles (T alleles = 0). These results are adjusted for family history, age at biopsy, parity, age at menarche, age at first birth, and BMI. Full model covariates can be found in the supplemental material, Table S2.