**2017 Gene x Environment Summary Statistics**

**Summary Statistics for Models Adjusting for Smoking Status**

**Column headers:**

1. chromosome

2. rs\_id: dbSNP ID

3. markername: chr:pos

4. allele\_1: effect allele

5. allele\_2: other allele

6. freq\_Allele1\_HapMapCEU: The allele frequency of Allele1 in the HapMap CEU population

7. effect: beta

8. stderr: standard error

9. p\_value: p-value after correction for inflation of test statistics using genomic control both at the individual study level and again after meta-analysis

10. N: sample size