

Inherited variants affecting RNA editing may contribute to ovarian cancer susceptibility: results from a large-scale collaboration

Supplementary Material

Supplementary Table 1: Description of individual OCAC studies and case-control sets included in the analysis of RNA editing SNPs and EOC susceptibility¹.

For Table S1, please see the attached file

Supplementary Table 2: HaploReg view of top *ADAR* eQTL SNP rs1127313 and proxies.

For Table S2, please see the attached Excel file