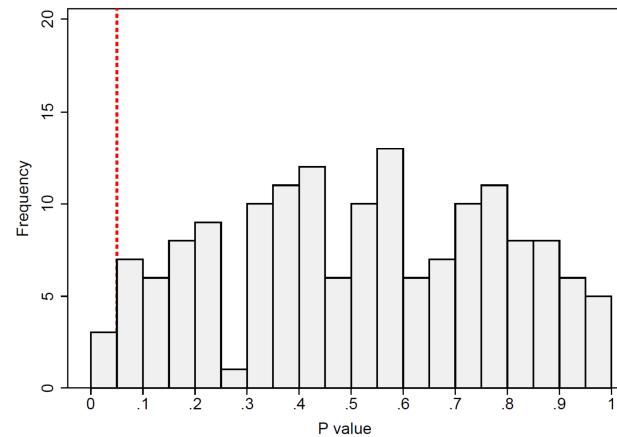
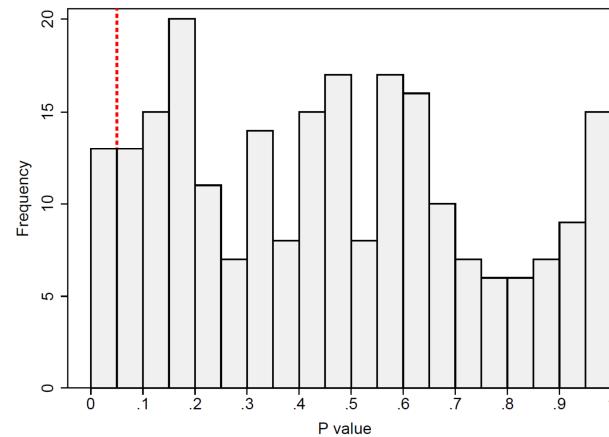


S4 Fig. Histograms of P-values obtained from logistic regression analyses assessing associations between SNPs in the indicated category and risk of EA. (A) miRNA biogenesis genes, (B) miRNA genes, (C) miRNA target genes, (D) all SNPs.

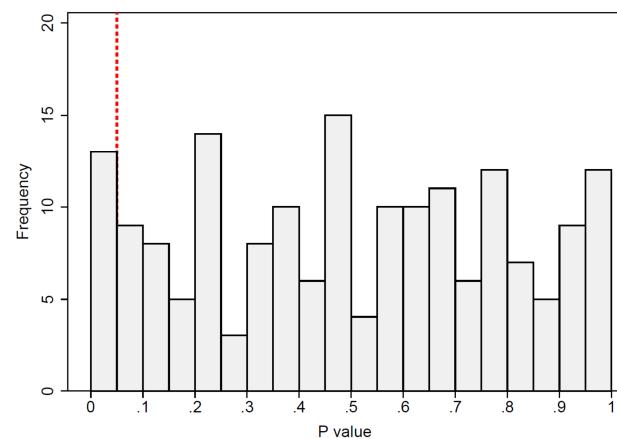
A.



B.



C.



D.

