

Supplementary Data 3. Forest plots of 10 variants that have been previously reported to have association with migraine but have *P*-value above genome-wide significance threshold ($P > 5 \times 10^{-8}$) in our study (N = 873,341; 102,084 migraine cases and 771,257 controls). For each variant, we plot the log-odds-ratio estimate (BETA) with its 95%-confidence intervals (green) from each of the five studies included in the meta-analysis and a combined beta estimate from the fixed-effect meta-analysis (blue diamond). Grey squares indicate the sample sizes of each study. We annotate each plot with the lead variant and effect allele, uncorrected two-sided *P*-value by the inverse-variance weighted fixed-effect meta-analysis and the heterogeneity index (I^2).



















