Identification of new susceptibility loci for gastric noncardia adenocarcinoma: Pooled results from two Chinese genome-wide association studies

Supp. Figure 1  QQ Plot of the meta analysis $p$ values for 3,996,998 SNPs after filtering out those with $P_{het} < 1.0 \times 10^{-4}$ or $I^2 > 75\%$. The genomic control lambda value is 1.02.
Supp Figure 2  Manhattan plot of the meta-analysis $p$ values (stage1)