

Genetic Determinants Of Anti-Malarial Acquired Immunity In A Large Multi-Centre Study

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This file contains **Additional Figure SF5: Plot for 178 SNPs with logged anti-malarial antibody levels***. Values of $-\log_{10}$ p-values plotted against chromosomal positions; only the lowest meta-analysis p-value for each SNP-antibody association is plotted. The red dotted lines indicates a Bonferroni threshold p-value of 6×10^{-5} (equivalent to a Bonferroni correction). Each colour represents a different anti-malarial antibody whilst shape of point represents the genetic model of best fit for the SNP-antibody association: (circle) for additive, (triangle) for dominant, and (square) for heterozygote and (plus) for recessive.

*Adjusted for age, gender, parasite density, village (>20), ethnicity (>20), sample month (>20) and study.

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