



Supplementary Figure 3. Quantile-quantile plot for the ImmunoChip association P values of 96,048 markers in 722 cases and 461 controls. $-\log_{10} P$ values corrected for residual inflation by genomic control were plotted against the expected null distribution. Red data points represent the inclusion of SNPs from the MHC locus; black data points represent the exclusion of the MHC and *TNFSF15* loci.