

Figure S1A. The cell enrichment of SLE genes implicated by 63 SNPs in Eastern Asian population within 249 cell types expression matrix for mus musculus. *The bottom indicates the log transformed p value. The vertical line indicated the Bonferroni-corrected significance criteria. The cell types are listed in the right.*

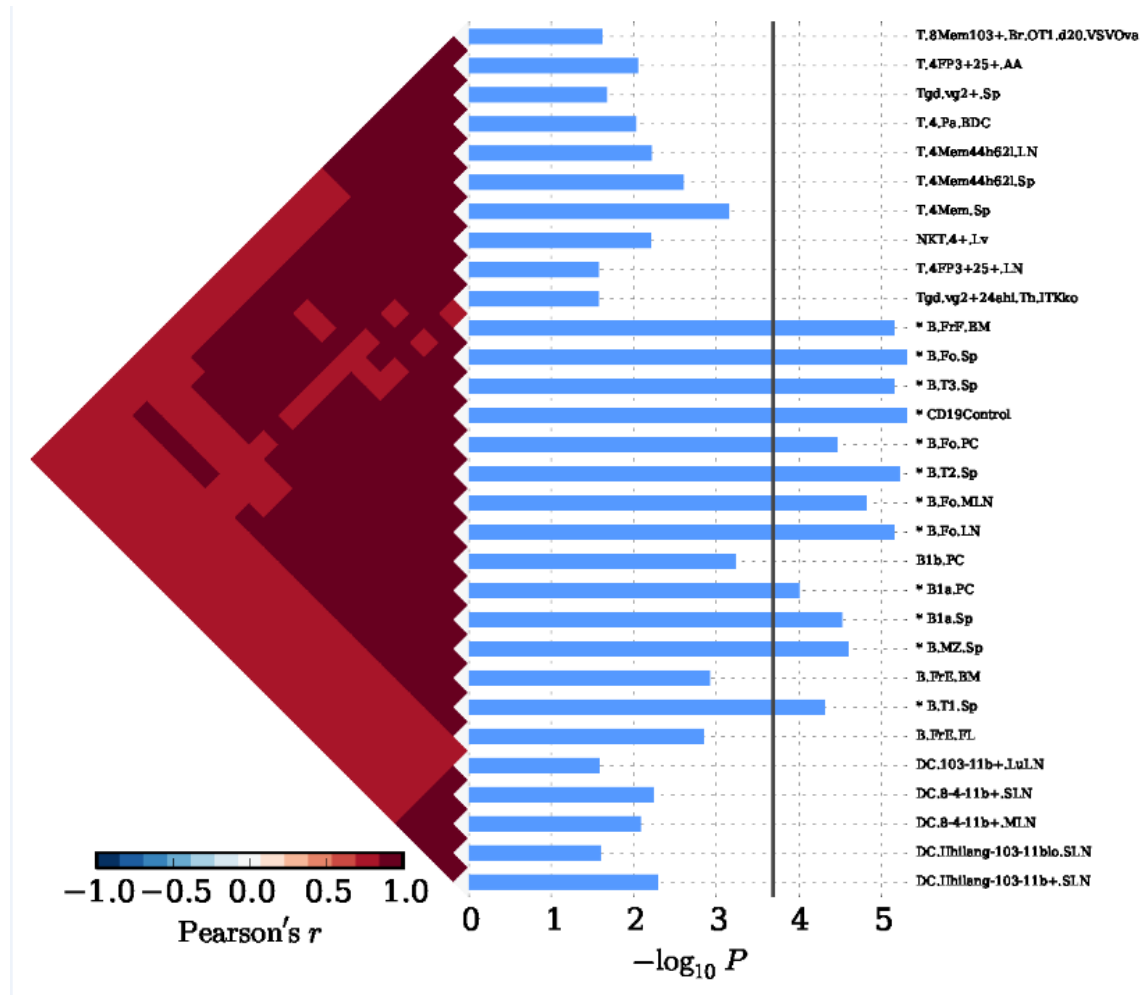


Figure S1B. The cell enrichment of SLE implicated genes by 118 SNPs in Caucasian population within 249 cell types expression matrix for mus musculus. *The bottom indicates the log transformed p value. The vertical line indicated the Bonferroni-corrected significance criteria. The cell types are listed in the right.*

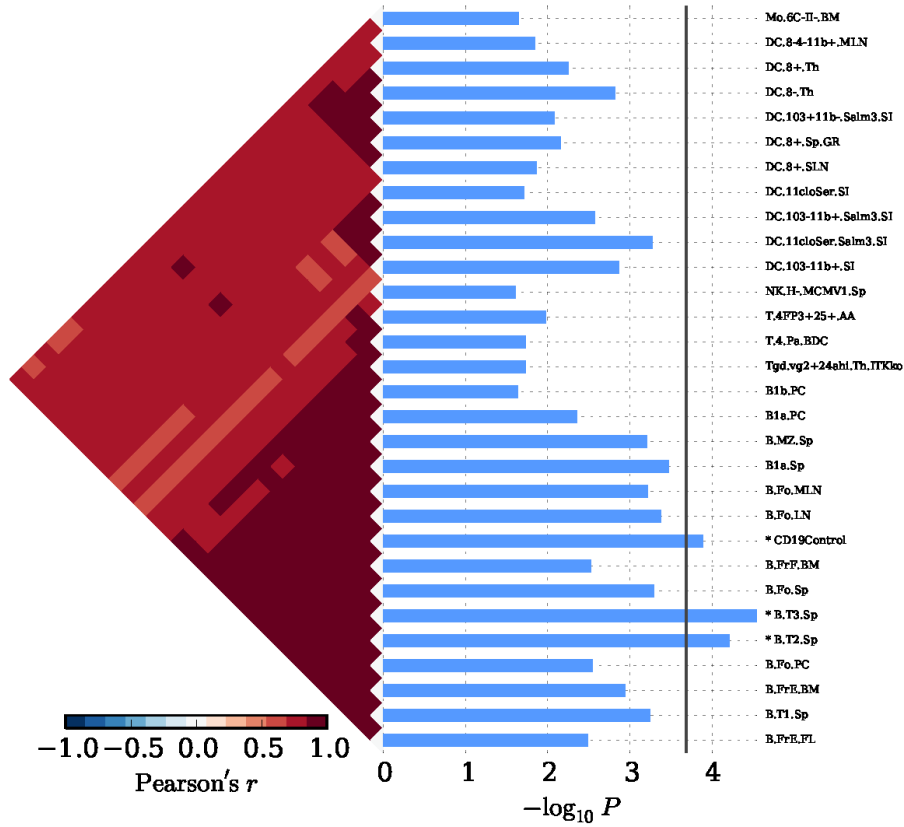


Figure S1C. The cell enrichment of SLE implicated gene by 105 SNPs without HLA region SNPs in Caucasian population within 249 cell types expression matrix for mus musculus. *The bottom indicates the log transformed p value. The vertical line indicated the Bonferroni-corrected significance criteria. The cell types are listed in the right.*

