**Supplementary Table 3. Nine candidate SNPs selected after clump analysis.**

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| **CHR** | **BP** | **Leading SNP** | **Leading SNP located gene** | ***p* for eQTL analysis** | **eQTL gene** |
| 1 | 207279481 | rs12057769 | *C4BPA* | 2.69×10-63 | *C4BPA* |
| 1 | 207505709 | rs6662070 | *CD55* | 4.16×10-29 | *CD55* |
| 6 | 31916951 | rs541862 | *CFB* | 4.90×10-22 | *SKIV2L* |
| 6 | 31914935 | rs1048709 | *CFB* | 2.85×10-16 | *HLA-DRB5* |
| 1 | 207535246 | rs1507765 | *CD55* | 3.86×10-9 | *CD55* |
| 19 | 862912 | rs3826945 | *CFD* | 7.05×10-7 | *CFD* |
| 1 | 207305670 | rs4266889 | *C4BPA* | 5.59×10-6 | *CD55* |
| 1 | 207496896 | rs28371597 | *CD55* | 5.89×10-6 | *CD55* |
| 11 | 57370742 | rs78364821 | *SERPING1* | 1.99×10-5 | *SERPING1* |

BP, base pair; CHR, chromosome; eQTL, expression quantitative trait locus; SNP, single-nucleotide polymorphism.