

Description of Additional Supplementary Files

Supplementary Data 1 Summary association statistics of reported SLE-associated variants

Supplementary Data 2 Summary association statistics of 38 novel SLE susceptibility loci

Supplementary Data 3 Transcription factors whose binding sites are enriched in the SLE-associated loci

Supplementary Data 4 Putative disease genes identified at each associated locus

Supplementary Data 5 Summary of fine-mapping results based on ancestry-dependent and trans-ancestral analyses