

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: Supporting evidence of 13 novel loci identified by C-GWAS

File Name: Supplementary Data 2

Description: Study-wide suggestive significant results of C-GWAS and MinGWAS

File Name: Supplementary Data 3

Description: 56 C-GWAS suggestively significant regional lead SNPs and their replication results

File Name: Supplementary Data 4

Description: Study-wide suggestive significant results of C-GWAS and MTAG

File Name: Supplementary Data 5

Description: Significant results of C-GWAS and MinGWAS in gene ontology analysis

File Name: Supplementary Data 6

Description: Colocalizations between C-GWAS findings and eQTL in 22 tissues

File Name: Supplementary Data 7

Description: CNCC regulatory subnetwork based on C-GWAS and MinGWAS findings