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| --- | --- | --- | --- | --- | --- | --- | --- |
| Supplementary Table S1 SNP information | | | |  |  |  |  |
| NCBI SNP ID | Positiona | Alleles | No. of SNPs tagged | HWE-pval | Genotyping rate(%) | MAF in CHBb | MAF in study |
|
| rs303218 | 89392836 | A:G | 1 | 0.36 | 98.2 | 0.47 | 0.45 |
| rs303215 | 89397074 | T:C | 2 | 0.71 | 99.1 | 0.46 | 0.49 |
| rs11203109 | 89397402 | T:C | 1 | 0.04 | 94.6 | 0.21 | 0.19 |
| rs303212 | 89401598 | C:T | 1 | 0.93 | 96.9 | 0.33 | 0.34 |
| Abbreviations: CHB: Han Chinese in Beijing, China; MAF: Minor Allele Frequency; NCBI: National Center for Biotechnology Information; HWE: Hardy-Weinberg equilibrium; SNP: Single nucleotide polymorphism | | | | | | | |
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| a. Position was referred to GRCh38.p2 from NCBI. | | | | | | | |
| b. MAF was referred to the HapMap Project. | | | | | | | |