|  |  |  |  |
| --- | --- | --- | --- |
| **DBSNP** | **Genotype** | **Overall Likelihood Ratio** | **Asian Likelihood Ratio** |
| rs1470579 | AA | 0.89 | 0.90 |
| AC | 1.07 | 1.06 |
| CC | 1.28 | 1.31 |
| rs2237892 | TT | 0.61 | 0.63 |
| CT | 0.92 | 0.92 |
| CC | 1.21 | 1.21 |
| rs2383208 | GG | 0.71 | 0.71\* |
| AG | 0.95 | 0.95\* |
| AA | 1.27 | 1.27\* |
| rs4402960 | GG | 0.90 | 0.89 |
| TG | 1.06 | 1.08 |
| TT | 1.27 | 1.34 |
| rs4712523 | AA | 0.75 | 0.75\* |
| AG | 1.03 | 1.03\* |
| GG | 1.43 | 1.43\* |
| rs5015480 | TT | 0.89 | 0.89 |
| TC | 1.22 | 1.23 |
| CC | 1.63 | 1.65 |
| rs7172432 | GG | 0.89 | 0.90 |
| AG | 0.98 | 0.98 |
| AA | 1.09 | 1.07 |
| rs7578597 | CC | 0.67 | 0.31 |
| TC | 0.83 | 0.56 |
| TT | 1.02 | 1.01 |
| rs7756992 | AA | 0.83 | 0.79 |
| AG | 1.04 | 0.98 |
| GG | 1.35 | 1.32 |
| rs7903146 | CC | 0.70 | 0.96 |
| TC | 1.17 | 1.16 |
| TT | 1.97 | 1.66 |
| rs8050136 | CC | 0.88 | 0.89 |
| AC | 1.05 | 0.95 |
| AA | 1.26 | 1.00 |
| rs9295475 | AA | 0.89 | 0.89\* |
| AG | 0.94 | 0.94\* |
| GG | 1.39 | 1.39\* |
| rs9300039 | AA | 0.61 | **NA** |
| AC | 0.80 | **NA** |
| CC | 1.05 | **NA** |
| rs10906115 | GG | 0.85 | 0.85\* |
| AG | 0.97 | 0.97\* |
| AA | 1.10 | 1.10\* |
| rs12255372 | GG | 0.81 | 0.97 |
| TG | 1.12 | 1.75 |
| TT | 1.66 | 1.01 |
| rs13266634 | TT | 0.83 | 0.87 |
| TC | 0.95 | 0.98 |
| CC | 1.11 | 1.11 |

**NA: The SNP was never verified in an Asian based GWAS and was excluded**

**\* no likelihood ratio difference is possible since only Asian-based studies exist**