Table S7 Correlation between *NAT2* SNPs and acetylator status

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| SNP | | Acetylator status | | | Concordance (%) |
| SA | IA | RA |
| rs1041983 | |  |  |  | 90.3 |
|  | AA | 25 | 0 | 0 |  |
|  | AG | 9 | 37 | 0 |  |
|  | GG | 1 | 5 | 26 |  |
| rs1495741 | |  |  |  | 98.1 |
|  | AA | 34 | 1 | 0 |  |
|  | AG | 1 | 39 | 1 |  |
|  | GG | 0 | 2 | 25 |  |

Concordance was calculated as the proportion of patients with AA genotypes and SA status and AG/GG genotypes and IA/RA status for both SNPs.

IA: intermediate acetylator, RA: rapid acetylator, SA: slow acetylator