**Table S1.** Details of the genotyping procedures used in the present study

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| ***SNP identifier*** | **Consequence** | **NAT2 allele cluster** | **TaqMan Assay Id.** | **PCR conditions** | **Allelic discrimination** |
| rs1801280 (T/C) | I114T | *NAT2\*5* | C\_\_\_1204093\_20 | 92ºC 15 sec  60ºC 90 sec  45 cycles | Fluorescent dyes as indicated by the manufacturer |
| rs1799930 (G/A) | R197Q | *NAT2\*6* | C\_\_\_1204091\_10 | 92ºC 15 sec  60ºC 90 sec  45 cycles | Fluorescent dyes as indicated by the manufacturer |
| rs1799931 (G/A) | G286E | *NAT2\*7* | C\_\_\_\_572770\_20 | 92ºC 15 sec  60ºC 90 sec  45 cycles | Fluorescent dyes as indicated by the manufacturer |
| rs1801279 (G/A) | R64Q | *NAT2\*14* | C\_\_\_\_572771\_10 | 92ºC 15 sec  60ºC 90 sec  45 cycles | Fluorescent dyes as indicated by the manufacturer |

According to standard procedures, all alleles lacking these SNPs are classified as *NAT2\*4* for this analysis.