



B

| ID | SNP 1 | SNP 2 | SNP 3 | SNP 4 | SNP 5 | SNP 6 | SNP 7 | SNP 8 | SNP 9 | SNP 10 | SNP 11 | SNP 12 | SNP 13 | SNP 14 | SNP 15 | SNP 16 | SNP 17 | Number of accessions |
|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|--------|--------|--------|--------|--------|--------|----------------------|
| Region | Promoter | Promoter | Promoter | Promoter | Promoter | Promoter | Promoter | Promoter | Promoter | Promoter | Promoter | Intron | Intron | Exon | Exon | 3'UTR | 3'UTR | |
| Position | -800 | -766 | -698 | -677 | -671 | -558 | -503 | -396 | -363 | -308 | -233 | 373 | 1689 | 6280 | 6458 | 7226 | 7257 | |
| Allele | T/G | C/T | C/G | G/A | T/C | T/C | G/A | G/A | T/A | C/A | T/C | C/T | C/G | G/A | T/C | G/T | A/T | |
| Ref.seq | T | C | C | G | T | T | G | G | T | C | T | C | C | G | T | G | A | |
| Hap 1 | G | C | C | A | C | C | A | A | A | A | C | T | C | G | T | G | T | |
| Hap 2 | G | C | C | G | T | C | G | G | T | C | T | T | G | G | T | G | A | 3 |
| Hap 3 | G | C | G | A | C | C | A | A | A | A | C | T | C | A | C | G | T | 2 |
| Hap 4 | T | C | C | G | T | T | G | G | T | C | T | T | G | G | T | T | A | 13 |
| Hap 5 | T | C | C | G | T | T | G | G | T | C | T | T | G | G | T | G | A | 39 |
| Hap 6 | T | T | C | G | T | T | G | G | T | C | T | C | G | G | T | T | A | 43 |
| Hap 6 | T | T | C | G | T | T | G | G | T | C | T | C | G | G | T | T | A | 21 |

