

| INTERPRETATION TABLE | | | | | |
|--|-------------------------|----------------|----------------|----------------|----------------|
| DQB1*04 SSP subtyping | | | | | |
| Amplification patterns of the DQB1*04 alleles | | | | | |
| | Well⁴ | | | | |
| | 1 | 2 | 3 | 4 | 5 |
| Length of spec. | 210 | 205 | 205 | 195 | 245 |
| PCR product(s) | 245 | | | | |
| Length of int. | 515 | 515 | 430 | 430 | 430 |
| pos. control¹ | | | | | |
| 5'-primer² | 9(122) | 23(164) | 23(164) | 26(173) | 9(122) |
| | 5' -gTT 3' | 5' -gCT 3' | 5' -gCg 3' | 5' -TCT 3' | 5' -gTA 3' |
| | 21(159) | | | | |
| | 5' -ACC 3' | | | | |
| | | | | | |
| 3'-primer³ | 77(327) | 77(327) | 77(327) | 77(327) | 77(327) |
| | 5' -ACg 3' | 5' -ACg 3' | 5' -ACg 3' | 5' -ACg 3' | 5' -ACg 3' |
| Well No. | 1 | 2 | 3 | 4 | 5 |
| DQB1 allele | | | | | |
| *04:01:01-04:01:02 | 1 | 2 | | | |
| *04:02 | 1 | | 3 | | |
| *04:03:01 | 1 | | 3 | 4 | |
| *04:03:02 | 1 | | | 4 | |
| *04:04 | 1 | | 3 | | 5 |
| *04:05 | 1 | 2 | | | 5 |
| *03:06, 03:25 | | | | 4 | 5 |
| DQB1 allele | | | | | |
| Well No. | 1 | 2 | 3 | 4 | 5 |