

INTERPRETATION TABLE						
DQB1*02 PCR-SSP subtyping						
Allele-specific amplification patterns of the 02:01 to 02:05 alleles						
	Well					
	1	2	3	4	5	6
Length of spec.	120	85	145	140	145	180
PCR product						
Length of int.	515	430	515	430	430	430
pos. control¹						
5'-primer²	30(185)	57(266)	135(500)	102(400)	136(502)	30(185)
	5'-AAg ^{3'}	5'-TgA ^{3'}	5'-TgA ^{3'}	5'-TCT ^{3'}	5'-ACg ^{3'}	5'-AAg ^{3'}
3'-primer³	57(266)	71(309)	170(604)	135(500)	170(604)	77(326)
	5'-Cgg ^{3'}	5'-CgT ^{3'}	5'-gAC ^{3'}	5'-ggC ^{3'}	5'-gAC ^{3'}	5'-CCg ^{3'}
Well No.	1	2	3	4	5	6
DQB1 allele⁴						
*02:01:01-02:01:03	1		3			
*02:02	1			4		
*02:03		2		4		
*02:04	1		3		5	
*02:05	1		3			6
*03:01:01-03:23, 03:25-03:31, 04:01:01-04:06, 05:01:01-05:08, 06:01:01-06:37, 06:39-06:40			3			
DQB1 allele⁴						
Well No.	1	2	3	4	5	6