

INTERPRETATION TABLE

DQB1*02 SSP subtyping

Amplification patterns of the 02:01 to 02:06 alleles

	Well						
	1	2	3	4	5	6	7
Length of spec.	120	85	145	140	145	180	245
PCR product							
Length of int.	515	430	515	430	430	430	430
pos. control ¹							
5'-primer ²	30(185)	57(266)	135(500)	101(400)	135(502)	30(185)	101(400)
	5' -AAg 3'	5' -TgA 3'	5' -TgA 3'	5' -TCT 3'	5' -ACg 3'	5' -AAg 3'	5' -TCT 3'
3'-primer ³	57(266)	71(309)	169(604)	135(500)	169(604)	77(326)	169(604)
	5' -Cgg 3'	5' -CgT 3'	5' -gAC 3'	5' -ggC 3'	5' -gAC 3'	5' -CCg 3'	5' -gAT 3'
Well No.	1	2	3	4	5	6	7
DQB1 allele							
*02:01:01-02:01:04	1		3				
*02:02	1			4			
*02:03		2		4			
*02:04	1		3		5		
*02:05	1		3			6	
*02:06	1			4			7
*03:01:01:01-03:23, 03:25-03:40, 04:01:01-04:08, 05:01:01:01-05:13, 06:01:01-06:37, 06:39-06:49			3				
*03:24							7
DQB1 allele							
Well No.	1	2	3	4	5	6	7