

INTERPRETATION TABLE								
DQB1*04 SSP subtyping								
Amplification patterns of the DQB1*04 alleles								
	Well⁴							
	1	2	3	4	5	6	7	8
Length of spec.	210	205	205	195	245	110	160	95
PCR product(s)	245							
Length of int.	515	515	430	430	430	430	430	430
pos. control¹								
5'-primer²	9(122)	23(164)	23(164)	26(173)	9(122)	55(259)	145(529)	167(596)
	⁵ -gTT ³	⁵ -gCT ³	⁵ -gCg ³	⁵ -TCT ³	⁵ -gTA ³	⁵ -ggT ³	⁵ -CCg ³	⁵ -gCA ³
	21(159)		23(164)					
	⁵ -ACC ³		⁵ -gCg ³					
3'-primer³	77(327)	77(327)	77(327)	77(327)	77(327)	77(327)	185(650)	185(650)
	⁵ -ACg ³	⁵ -ACg ³	⁵ -ACg ³	⁵ -ACg ³	⁵ -ACg ³	⁵ -ACg ³	⁵ -CgA ³	⁵ -CgA ³
Well No.	1	2	3	4	5	5	5	5
DQB1 allele								
*04:01:01-04:01:02	1	2						
*04:02:01-04:02:02	1		3					
*04:03:01	1		3	4				
*04:03:02	1			4				
*04:04	1		3		5			
*04:05	1	2			5			
*04:06	1	2				6		
*04:07	1	2					7	
*04:08	1	2						8
*03:06				4	5			?
*03:07-03:08, 03:11-03:18, 03:20, 03:23, 03:26, 03:37								?
*03:19								8
*03:25				4	5			
DQB1 allele								
Well No.	1	2	3	4	5	6	7	8