

INTERPRETATION TABLE								
DQB1*04 SSP subtyping								
Amplification patterns of the DQB1*04 alleles								
	Well ⁴							
	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. control ¹								
5'-primer ²	9(122)	23(164)	23(164)	26(173)	9(122)	54(259)	144(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'-primer ³	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	6	7	8
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, 03:40, 03:44								?
*03:19								8
*03:25				4	5			
DQB1 allele								
Well No.	1	2	3	4	5	6	7	8