

INTERPRETATION TABLE								
DRB1*09 SSP subtyping								
Amplification patterns of the DRB1*09:01 to 09:20 alleles								
	Well⁶							
	1	2	3	4	5	6	7	8
Length of spec.	195	130	135	100	185	105	220	150
PCR product				130	215	150		180
				215		250		
				245				
				275				
Length of int.	515	430	430	430	430	430	430	430
pos. control¹								
5'-primer²	26(164) 5'-gTA ^{3'}	28(170) 5'-gCA ^{3'}	28(170) 5'-gCA ^{3'}	8(112) 5'-TgA ^{3'}	8(112) 5'-TgA ^{3'}	8(112) 5'-TgA ^{3'}	26(165) 5'-TAT ^{3'}	8(112) 5'-TgC ^{3'}
								12(123) 5'-AAC ^{3'}
								21(149) 5'-gAT ^{3'}
3'-primer(s)³	77(319) 5'-CAC ^{3'}	57(257) 5'-CgA ^{3'}	60(266) 5'-AgT ^{3'}	28(172) 5'-TCC ^{3'}	56(256) 5'-gCT ^{3'}	30(176) 5'-TgT ^{3'}	86(344) 5'-CCA ^{3'}	57(257) 5'-CgA ^{3'}
		58(261) 5'-TCC ^{3'}		39(203) 5'-AgA ^{3'}	66(286) 5'-gAg ^{3'}	44(218) 5'-CCC ^{3'}		
		58(261) 5'-TCA ^{3'}		63(276) 5'-TgT ^{3'}		47(229) 5'-CCC ^{3'}		
				66(286) 5'-gAA ^{3'}		77(319) 5'-gTA ^{3'}		
				75(311) 5'-CCC ^{3'}				
				77(319) 5'-CAg ^{3'}				
				87(348) 5'-CAT ^{3'}				
Well No.	1	2	3	4	5	6	7	8
DRB1 allele^{4,5}								
*09:01:02-09:01:09	1	2						
*09:02:01-09:02:02	1		3					
*09:03, 09:09, 09:13, 09:15, 09:17-09:18	1	2		4				
*09:04	1	2			5			
*09:05		2				6		
*09:06	1	2					7	
*09:07, 09:12, 09:19- 09:20	1	2						8
*09:08	1				5			
*09:10-09:11, 09:14	1	2				6		
*09:16		2		4				
DRB1 allele⁴								
Well No.	1	2	3	4	5	6	7	8