

INTERPRETATION TABLE								
DRB1*09 SSP subtyping								
Amplification patterns of the DRB1*09 alleles								
	Well⁵							
	1	2	3	4	5	6	7	8
Length of spec.	195	130	185	215	185	250	220	175
PCR product					215			
Length of int.	515	430	430	430	430	430	430	430
pos. control¹								
5'-primer²	26(164)	28(170)	9(112)	9(112)	9(112)	9(112)	26(165)	13(125)
	5'-gTA 3'	5'-gCA 3'	5'-TgA 3'	5'-TgA 3'	5'-TgA 3'	5'-TgA 3'	5'-TAT 3'	5'-gTA 3'
3'-primer(s)³	78(319)	57(257)	57(257)	67(286)	57(256)	78(319)	86(344)	57(257)
	5'-CAC 3'	5'-CgA 3'	5'-CAT 3'	5'-gAA 3'	5'-gCT 3'	5'-gTA 3'	5'-CCA 3'	5'-CgA 3'
			57(258)		67(286)			
			5'-gCg 3'		5'-gAg 3'			
Well No.	1	2	3	4	5	6	7	8
DRB1 allele⁴								
*090102-090104	1	2						
*090201, 090202	1		3					
*0903	1	2		4				
*0904	1	2			5			
*0905		2				6		
*0906	1	2					7	
*0907	1	2						8
*0908	1				5			
DRB1 allele⁴								
Well No.	1	2	3	4	5	6	7	8