

INTERPRETATION TABLE									
DRB1*01 SSP subtyping									
Amplification patterns of the DRB1*0101 to 0131 alleles									
	Well ⁴								
	1	2	3	4	5	6	7	8	9
Length of spec.	255	235	200	210	220	135	210	175	110
PCR product(s)						210		210	255
Length of int.	515	430	515	430	430	430	430	430	430
pos. control ¹									
5'-primer ²	14 (129)	14 (129)	14 (129)	14 (129)	14 (129)	14 (129)	14 (129)	10 (115)	14 (129)
	5'-gAA 3'	5'-gAA 3'	5'-gAA 3'	5'-gAA 3'	5'-gAA 3'	5'-gAA 3'	5'-gAA 3'	5'-ggg 3'	5'-gAA 3'
								23 (154)	
								5'-Agg 3'	
3'-primer(s) ³	86 (344)	78 (321)	67 (286)	71 (299)	70 (296)	45 (220)	67 (286)	67 (286)	37 (197)
	5'-CAC 3'	5'-CAA 3'	5'-gAg 3'	5'-gCT 3'	5'-TCC 3'	5'-CCT 3'	5'-gAT 3'	5'-gAg 3'	5'-CgT 3'
			67 (286)		77 (317)	48 (230)	67 (286)		85 (341)
			5'-gAg 3'		5'-AAT 3'	5'-CCT 3'	5'-gAT 3'		5'-CAA 3'
						70 (296)	74 (308)		
						5'-TCC 3'	5'-CCT 3'		
Well No.	1	2	3	4	5	6	7	8	9
DRB1 allele									
*010101-010117	1		3						
*010201-010205		2	3						
*0103	1			4			7		
*0104			3		5				9
*0105	1		3			6			
*0106			3						9
*0107	1		3					8	
*0108	1		3						9
*0109	1		3						
*0110	1		3	4					
*0111	1		3		5				
*0112	1		3						
*0113	1		3						
*0114	1		3						
*0115	1						7		
*0116	1		3						
*0117	1		3				7		
*0118	1		3						
*0119	1		3						
*0120			3						9
Well No.	1	2	3	4	5	6	7	8	9

Length of spec.	255	235	200	210	220	135	210	175	110
PCR product(s)						210		210	255
Well No.	1	2	3	4	5	6	7	8	9
*0121	1		3						
*0122	w		w						
*0123		2							
*0124	1								
*0125	1		3						
*0126		2	3						
*0127	1		3					8	
*0128	1		3						
*0129	1		3		5	6			
*0130	1		3						
*0131	1		3			6			
DRB1 allele									
Well No.	1	2	3	4	5	6	7	8	9

			INTERPRETATION TABL						
			DRB1*01 SSP subtyping						
			Amplification patterns of the DRB1*0101 to						
			Well ⁴						
10	11	12	13	14	15	16	17	18	19
210	140	140	85	170	150	200	230	230	125
	210	180	150	210	220	230	260		
		215	215						
430	430	430	430	430	430	430	430	430	430
14 (129)	14 (129)	14 (129)	14 (129)	14 (129)	14 (129)	14 (129)	12 (123)	23 (154)	59 (262)
5'-gAA 3'	5'-gAA 3'	5'-gAA 3'	5'-gAA 3'	5'-gAA 3'	5'-gAA 3'	5'-gAA 3'	5'-AAC 3'	5'-AgT 3'	5'-CCA 3'
							23 (154)		
							5'-Agg 3'		
71 (298)	47 (227)	48 (230)	30 (175)	57 (257)	51 (239)	67 (286)	86 (344)	86 (344)	86 (344)
5'-CgC 3'	5'-ggA 3'	5'-CCT 3'	5'-gAg 3'	5'-CAg 3'	5'-CCA 3'	5'-gAT 3'	5'-CAC 3'	5'-CAC 3'	5'-CAC 3'
	71 (298)	60 (266)	51 (239)	67 (286)	73 (305)	76 (313)			
	5'-CTT 3'	5'-Agg 3'	5'-CCA 3'	5'-gAT 3'	5'-ggC 3'	5'-gTT 3'			
		72 (302)	72 (303)	72 (303)	78 (319)	78 (319)			
		5'-CCA 3'	5'-gCg 3'	5'-gCg 3'	5'-CAC 3'	5'-CAC 3'			
10	11	12	13	14	15	16	17	18	19
10									
10	11								
		12							
			13						
				14					
10									
					15				
	11								
						16			
10	11	12	13	14	15	16	17	18	19

210	140	140	85	170	150	200	230	230	125
	210	180	150	210	220	230	260		
		215	215						
10	11	12	13	14	15	16	17	18	19
					15	16			
							17		
			13	14					
				14		16			
								18	
		12							
							17		
			13		15				
									19
		12							
10	11	12	13	14	15	16	17	18	19

E
0131 alleles
Length of spec. PCR product(s)
Length of int. pos. control¹
5'-primer²
3'-primer(s)³
Well No.
DRB1 allele
*010101-010117
*010201-010205
*0103
*0104
*0105
*0106
*0107
*0108
*0109
*0110
*0111
*0112
*0113
*0114
*0115
*0116
*0117
*0118
*0119
*0120
Well No.

Length of spec.
PCR product(s)
Well No.
*0121
*0122
*0123
*0124
*0125
*0126
*0127
*0128
*0129
*0130
*0131
DRB1 allele
Well No.