

INTERPRETATION TABLE								
HLA-Cw*16 SSP subtyping								
Amplification patterns of the HLA-Cw*1601 to 1612 alleles								
	Well⁵							
	1	2	3	4	5	6	7	8
Length of spec.	205	340	85	140	160	130	100	135
PCR product			220			160	210	
						210		
Length of int.	800	1070	800	1070	800	1070	1070	1070
pos. control¹								
5'-primer(s)²	361	1st 	256	201	419	113	244	126
	5'-AgT ^{3'}	5'-CgA ^{3'}	5'-ACg ^{3'}	5'-CCA ^{3'}	5'-gTC ^{3'}	5'-CCA ^{3'}	5'-CgC ^{3'}	5'-ggA ^{3'}
			361			368	369	
			5'-AgT ^{3'}			5'-gTC ^{3'}	5'-TAC ^{3'}	
						418		
						5'-Agg ^{3'}		
3'-primer(s)³	527	302	302	302	539	201	302	220
	5'-CCg ^{3'}	5'-ggT ^{3'}	5'-ggT ^{3'}	5'-ggC ^{3'}	5'-TCT ^{3'}	5'-CTT ^{3'}	5'-ggT ^{3'}	5'-CgA ^{3'}
			538			539	539	
			5'-CCA ^{3'}			5'-TCT ^{3'}	5'-TCT ^{3'}	
Well No.	1	2	3	4	5	6	7	8
HLA-Cw allele⁴								
*160101, 160103	1			4	5			
*160102	1				5			
*1602	1	2			5			
*160401	1		3	4				
*1606				4	5	6		
*1607				4	5		7	
*1608	1			4	5			8
*1609	1	2			5		7	
*1610	1			4		6		
*1611	1			4	5	6		
*1612	1	2	3		5			
*0104, 0109, 0611, 120304			3					
Well No.	1	2	3	4	5	6	7	8

Length of spec.	205	340	85	140	160	130	100	135
PCR product			220			160	210	
						210		
Well No.	1	2	3	4	5	6	7	8
*0114, 020201-020203, 020205-0204, 0206-0211, 0213-0216, 0218-0220, 0222, 0223, 0307, 0315, 0345, 04010101-040106, 0403-0410, 0412-0420, 0423-0428, 0430-0435, 0437, 050101-050104, 0503-0523, 0604-0606, 0614, 0707, 0709, 0749, 0810, 1209, 1221, 1404, 150201-1506, 1508-1513, 1515-1520, 1701-1705, 1801-1803	2							
*0121, 0411, 0429, 0436, 080101-0809, 0811-0819, 120201-120203, 1208, 1210, 1214, 1216-1220, 140203, 1403, 1408, 1410, 1507				4				
*0205, 0217, 06020101-0603, 0607-0610, 0612, 0613, 0615-0619, 120401-1205	2		3					
*0212, 1521				w				
*0221	2			6				
*0753					5			
*12030101-120303, 120305-120306, 1206, 1207, 1211-1213, 1215			3	4				
HLA-Cw allele ⁴								
Well No.	1	2	3	4	5	6	7	8
B*350802, B*6702				4				
Well No.	1	2	3	4	5	6	7	8