

INTERPRETATION TABLE								
HLA-A*29 SSP subtyping								
Amplification patterns of the A*2901 to 2916 alleles								
	Tube⁴							
	1	2	3	4	5	6	7	8
Length of spec.	475	130	440	165	130	130	210	85
PCR product								
Length of int.	800	1070	800	800	1070	800	1070	1070
pos. control¹								
5'-primer²	180	808	219	448	180	448	448	368
	5'-TTT ^{3'}	5'-CgT ^{3'}	5'-gCA ^{3'}	5'-CCT ^{3'}	5'-TTT ^{3'}	5'-CCT ^{3'}	5'-CCT ^{3'}	5'-gTT ^{3'}
3'-primer³	376	895	376	570	268	539	616	413
	5'-gTg ^{3'}	5'-CTC ^{3'}	5'-gTC ^{3'}	5'-CCg ^{3'}	5'-ATg ^{3'}	5'-TCT ^{3'}	5'-CgC ^{3'}	5'-gCC ^{3'}
Tube No.	1	2	3	4	5	6	7	8
HLA-A allele								
*29010101	+							
*29010102N	+	+						
*290201-290203			+					
*2903			+	+				
*2904			+		+			
*2905			+			+		
*2906			+				+	
*2907			+					+
*2908N			+					
*2909			+					
*2910			+					
*2911			+					
*2912	+							
*2913			+					
*2914			+					
*2915	+							
*2916	+							
*0327								
*2303, 3105, 3213, 3310				+				
*2417, 2441								+
*3107, 3108, 3110								
*3116								
*3202						+		
*3212							+	

Tube No.	1	2	3	4	5	6	7	8
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9	10	11	12	13	14	15	16	Tube No.
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