

INTERPRETATION TABLE									
DQ low resolution SSP typing									
Amplification patterns of the DQB1*0501 to DQB1*0403 alleles									
		Well							
		88	89	90	91	92	93	94	95
Length of spec.		220	220	205	130	220	220	135	210
PCR product(s)			270		145				
Length of int.		515	430	430	515	515	515	515	430
pos. control ¹									
5'-primer(s) ²		26	9	30	21	26	26	55	21
		5'-g gg ^{3'}	5'-g TT ^{3'}	5'-A Ag ^{3'}	5'-ACC ^{3'}	5'-T TA ^{3'}	5'-T CT ^{3'}	5'-g CC ^{3'}	5'-ACC ^{3'}
			26		26			55	
			5'-T TA ^{3'}		5'-T CT ^{3'}			5'-g CA ^{3'}	
			26						
			5'-T CT ^{3'}						
3'-primer(s) ³		87	86	86	57	86	86	86	77
		5'-g gT ^{3'}	5'-A Cg ^{3'}	5'-g CT ^{3'}	5'-C gg ^{3'}	5'-g CT ^{3'}	5'-g CT ^{3'}	5'-g CT ^{3'}	5'-AC g ^{3'}
			86						
			5'-A CC ^{3'}						
Well No.		88	89	90	91	92	93	94	95
DOB1 allele ⁴	ser. ⁵								
*050101-0505	5	88							
*060101-0628, 0630-0634	6		89						
*0629	6		89		91				
*020101-0202, 0204, 0205	2			90	91		93		
*0203	2			90			93		
*030101-030104, 0304, 0309, 0310, 0313, 0314, 0316, 0319, 0321	3, 7, 8					92		94	
*030201-030204, 0307, 0308, 0311, 0318	3, 8				91		93	94	
*030302-030303, 0306, 0312, 0315, 0320	3, 9						93	94	
*030501, 030503	3, 8				91			94	
*030502, 030504, 0317								94	
*0401, 0402	4								95
*0403	4						93		95
DOB1 allele ⁴	ser. ⁵								
Well No.		88	89	90	91	92	93	94	95