

# 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.		225	220	210	135	220	220	145	210
PCR product(s)			270		150			185	
Length of int.		515	430	430	515	515	515	515	430
pos. control <sup>1</sup>									
5'-primer(s) <sup>2</sup>		26(173)	9(122)	30(185)	21(159)	26(173)	26(173)	38(210)	21(159)
		5'-ggg 3'	5'-gTT 3'	5'-AAg 3'	5'-ACC 3'	5'-TTA 3'	5'-TCT 3'	5'-gCA 3'	5'-ACC 3'
			26(173)		26(173)			48(240)	
			5'-TTA 3'		5'-TCT 3'			5'-CgC 3'	
			26(173)					55(260)	
			5'-TCT 3'					5'-gCC 3'	
								55(260)	
								5'-gCA 3'	
3'-primer(s) <sup>3</sup>		87(356)	86(353)	86(353)	57(266)	86(353)	86(353)	86(353)	77(327)
		5'-ggT 3'	5'-ACg 3'	5'-gCT 3'	5'-Cgg 3'	5'-gCT 3'	5'-gCT 3'	5'-gCT 3'	5'-ACg 3'
			86(353)						
			5'-ACC 3'						
Well No.		88	89	90	91	92	93	94	95
DQB1 allele <sup>4</sup>	ser. <sup>5</sup>								
*050101-0505	5	88							
*060101-0628, 0630-0635	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, 0322, 0324	3, 7, 8					92		94	
*030201-030204, 0307, 0308, 0311, 0318	3, 8				91		93	94	
*030302, 030303, 0306, 0312, 0315, 0320, 0323, 0325	3, 9						93	94	
*030501, 030503	3, 8				91			94	
*030502, 030504, 0317								94	
*040101-0402	4								95
*0403	4						93		95
DQB1 allele <sup>4</sup>	ser. <sup>5</sup>								
Well No.		88	89	90	91	92	93	94	95