INTERPRETATION TABLE

DQ low resolution SSP typing

Amplification patterns of the DQB1*0501 to DQB1*0403 alleles

		Well							
		1	2	3	4	5	6	7	8
Length of spec.		225	220	210	220	130	135	145	210
PCR product(s)			270					185	
Length of int.		515	430	430	515	515	515	515	430
pos. control[1]									
5'-primer(s)[2]		26 (173)	9 (122)	30 (185)	26 (173)	28 (179)	26 (173)	38 (210)	21 (159)
		5' -ggg 3'	5' -gTT 3'	5' -AAg 3'	5' -TTA 3'	5' -gAC 3'	5' -TCT 3'	5' -gCA 3'	5' -ACC 3'
			26 (173)			28 (179)		48 (240)	
			5' -TTA 3'			5' -gAC 3'		5' -CgC 3'	
			26 (173)					55 (260)	
			5' -TCT 3'					5' -gCC 3'	
								55 (260)	
								5' -gCA 3'	
3'-primer(s)[3]		87 (356)	86 (353)	86 (353)	86 (353)	57 (266)	57 (266)	86 (353)	77 (327)
		5' -ggT 3'	5' -ACg 3'	5' -gCT 3'	5' -gCT 3'	5' -Cgg 3'	5' -CgT 3'	5' -gCT 3'	5' -ACg 3'
			86 (353)						
			5' -ACC 3'						
Well No.		1	2	3	4	5	6	7	8
DQB1 allele[4]	ser.[5]								
*050101-0505	5	1							
*060101-0628, 0630-0639	1, 5, 6, null		2						
*0629	6		2			5			
*020101-0202, 0204, 0205	2			3					
*0203	2			3			6		
*030101-030104, 0309, 0310, 0313, 0316, 0319, 0321, 0322, 0324	3, 7				4			7	
*030201-030204, 030501-030504, 0307, 0308, 0311, 0318	8					5		7	
*030302, 030303, 0306, 0312, 0315, 0320, 0325, 0326	3, 9						6	7	
*0304, 0314	3, 7				4	w		7	
*0317, 0323	9							7	
*040101-0402, 0404	4								8
*040301	-						6		8
*040302	-						6		
DQB1 allele[4]	ser.[5]								

Well No.		1	2	3	4	5	6	7	8