

INTERPRETATION TABLE									
DQ low resolution SSP typing									
Amplification patterns of the DQB1*05:01 to DQB1*04:08 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	245
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)	9 (122)
		5'-ggg 3'	5'-gTT 3'	5'-AAG 3'	5'-TTA 3'	5'-gAC 3'	5'-TCT 3'	5'-gCA 3'	5'-gTT 3'
			26 (173)			28 (179)		48 (240)	21 (159)
			5'-TTA 3'			5'-gAC 3'		5'-CgC 3'	5'-ACC 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								
*05:01:01:01-05:14	5, –	1							
*06:01:01-06:28, 06:30-06:44, 06:47	1, 5, 6, Null, –		2						
*06:29	6		2			5			
*02:01:01-02:02, 02:04-02:06	2, –			3					
*02:03	2			3			6		
Well No.		1	2	3	4	5	6	7	8
Well No.		1	2	3	4	5	6	7	8
DQB1 allele ⁴	ser. ⁵								
*03:01:01:01-03:01:06, 03:04, 03:09-03:10, 03:13-03:14, 03:16, 03:19, 03:21-03:22, 03:24, 03:27-03:29, 03:35-03:36	3, 7, –				4			7	
*03:02:01-03:02:05, 03:05:01-03:05:04, 03:07-03:08, 03:11, 03:18, 03:32, 03:37	8, –					5		7	

*03:03:02:01-03:03:04, 03:06, 03:12, 03:15, 03:20, 03:25-03:26, 03:30- 03:31, 03:33-03:34, 03:38-03:39	3, 9, –						6	7	
*03:17, 03:23	9, –							7	
*04:01:01-04:02:02, 04:04-04:08	4, –								8
*04:03:01-04:03:02	–						6		8
DQB1 allele ⁴	ser. ⁵								
Well No.		1	2	3	4	5	6	7	8