INTERPRETATION TA	
DQB1*06:02 SSP	
	7
Length of spec.	210
PCR product	
Length of int.	515
pos. control[1]	
5'-primer[2]	29(184)
	5' -gAT 3'
	29(184)
	5' -gAT 3'
3'-primer[3]	86(353)
	5' -ACg 3'
Well No.	7
DQB1 allele	
DQB1*06:02:01-06:02:02	7
DQB1*06:01:01, 06:01:03-06:01:06, 06:10-06:11:02, 06:13, 06:18, 06:29, 06:35, 06:43	7
DQB1*06:01:02	7
DQB1*06:03:01-06:05:01, 06:07:01-06:09, 06:12, 06:17, 06:21, 06:25-06:28, 06:30, 06:32, 06:34, 06:36, 06:38-06:39, 06:41-06:42, 06:44, 06:52, *DQB1*03:38*	
DQB1*06:05:02	?
DQB1*06:06	?
DQB1*06:14:01-06:14:02, 06:46	
DQB1*06:15	
DQB1*06:16, 06:19, 06:48, 06:50	7

DQB1*06:20, 06:33, 06:49	**7**
DQB1*06:22	
DQB1*06:23	
DQB1*06:24, 06:47	**7**
DQB1*06:31	
DQB1*06:37	**7**
DQB1*06:45	**7**
DQB1*06:51	**7**
DQB1*03:30, 04:01:01-04:03:02, 04:06-04:08	
DQB1 allele	
Well No.	**7**

BLE				
Well[4]				
8	9	10	11	12
185	155	115	50	
	195	225	100	
	230	265	175	
			220	
430	430	430	430	
9(122)	9(122)	11(129)	26(173)	
5' -gTT 3'	5' -gTT 3'	5' -TTA 3'	5' -TCT 3'	
		13(134)	154(558)	
		5' -ggT 3'	5' -ACT 3'	
		13(136)		
		5' -gCC 3'		
		26(173)		
		5' -ggg 3'		
		62(282)		
		5' -AAg 3'		
57(266)	47(237)	87(356)	29(184)	
5' -CAT 3'	5' -CgA 3'	5' -ggA 3'	5' -gTg 3'	
	59(274)		70(307)	
	5' -gTT 3'		5' -ggC 3'	
	69(304)		86(353)	
	5' -CCT 3'		5' -ACC 3'	
	75(322)		174(618)	
	5' -gTg 3'		5' -ACT 3'	
8	9	10	11	12
8				
		W		
			11	
	9		?	
			?	
8			11	
8	9		11	
8	9			

Negative Control

ative Control

8		10		Nega
	9		11	
8		10		
8			11	
		10	11	
8	9	10		
		10		
	9			
	9			
8	9	10	11	12