

*15:40	1	2	3	4												
*15:41	1	2	3	4												
*15:42	1	2	3	4												
*15:43	1	2	3	4												
*15:44	1	2	3		5											
*15:45	1	2	3	4												
*15:46	1	2	3	4												
*15:47	1	2	3		5											
*15:48		2	3	4								12				
*15:49	1			4												
*15:50N	1			4												
*15:54, 15:66	1			4				8								
*15:55	1	2	3	4												
*15:57	1	2		4		6										
*15:58	1	2			5											
*15:59		2	3	4												
*15:63	1				5			8								
*15:65	1	2	3	4												
*15:68	1	2	3		5											
*15:69			3							9						
*15:70, 15:89, 15:91	1			4												
*15:74	1	2	3													16
*15:78	1	2			5	6										
*15:80N	1				5											
*15:84	1			4											13	
*15:85	1	2	3													
*15:88				4			7			10						
Well No.	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
Length of spec. PCR product(s)	140	200	150	260	260	150	140	200	90	200	170	95	110	200	200	200
						180		245	215				215			260
Well No.	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
*04:73, 04:105:01-04:105:02, 11:82, 13:09, 13:129, 14:37			3													
*11:88												12				
*16:01:01-16:03, 16:08, 16:11, 16:13N-16:14, 16:16-16:17, 16:19-16:21N					5		7									
*16:04, 16:18					5		7									
*16:05:01-16:05:02, 16:07					5		7								15	
*16:09-16:10	1				5											
*16:15				4			7									
DRB1 allele																
Well No.	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16

INTERPRETATION TABLE

DRB1*15 SSP subtyping

Amplification patterns of the DRB1*15:01 to 15:92 alleles

Well ⁶																
17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	
160	205	175	165	165	165	125	200	70	75	175	135	200	165	150	155	Length of spec.
		225	200		200	170		195			255		220	215	200	PCR product(s)
						220										
430	515	430	430	430	430	430	430	430	430	430	430	430	430	515	430	Length of int. pos. control ¹
																5'primer(s) ²
5'-gAT ^{3'}	5'-gAT ^{3'}	5'-Agg ^{3'}	5'-Agg ^{3'}	5'-gAC ^{3'}	5'-AAG ^{3'}	5'-Agg ^{3'}	5'-gTA ^{3'}	5'-AgA ^{3'}	5'-gTC ^{3'}	5'-TgC ^{3'}	5'-Agg ^{3'}	5'-Agg ^{3'}	5'-Agg ^{3'}	5'-Agg ^{3'}	5'-Agg ^{3'}	
					5'-gTA ^{3'}		5'-gTA ^{3'}	5'-ACA ^{3'}		5'-gCA ^{3'}						
					5'-gCA ^{3'}		5'-ggT ^{3'}									
5'-CTC ^{3'}	5'-CCT ^{3'}	5'-gCC ^{3'}	5'-gCA ^{3'}	5'-CgC ^{3'}	5'-gAT ^{3'}	5'-gTg ^{3'}	5'-CgC ^{3'}	5'-gAT ^{3'}	5'-CgC ^{3'}	5'-gAT ^{3'}	5'-CCT ^{3'}	5'-CTC ^{3'}	5'-CCA ^{3'}	5'-CCT ^{3'}	5'-CCC ^{3'}	3'primer(s) ³
17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	Well No.
160	205	175	165	165	165	125	200	70	75	175	135	200	165	150	155	Length of spec.
		225	200		200	170		195			255		220	215	200	PCR product(s)
						220										
17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	Well No.
																DRB1 allele ⁴

