

<b>INTERPRETATION TABLE</b>								
<b>DRB1*09 SSP subtyping</b>								
<b>Amplification patterns of the DRB1*09:01 to 09:17 alleles</b>								
	<b>Well<sup>6</sup></b>							
	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>	<b>7</b>	<b>8</b>
<b>Length of spec.</b>	195	130	185	100	185	105	220	175
<b>PCR product</b>				130	215	150		
				215		250		
				245				
				275				
<b>Length of int.</b>	<b>515</b>	430	430	430	430	430	430	430
<b>pos. control<sup>1</sup></b>								
<b>5'-primer<sup>2</sup></b>	26(164)	28(170)	9(112)	9(112)	9(112)	9(112)	26(165)	12(123)
	5'-gTA 3'	5'-gCA 3'	5'-TgA 3'	5'-TgA 3'	5'-TgA 3'	5'-TgA 3'	5'-TAT 3'	5'-AAC 3'
								13(125)
								5'-gTA 3'
<b>3'-primer(s)<sup>3</sup></b>	78(319)	57(257)	57(257)	29(172)	57(256)	30(176)	86(344)	57(257)
	5'-CAC 3'	5'-CgA 3'	5'-CAT 3'	5'-TCC 3'	5'-gCT 3'	5'-TgT 3'	5'-CCA 3'	5'-CgA 3'
			57(258)	39(203)	67(286)	44(218)		
			5'-gCg 3'	5'-AgA 3'	5'-gAg 3'	5'-CCC 3'		
				67(286)		48(229)		
				5'-gAA 3'		5'-CCC 3'		
				75(311)		78(319)		
				5'-CCC 3'		5'-gTA 3'		
				78(319)				
				5'-CAg 3'				
				87(348)				
				5'-CAT 3'				
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>	<b>7</b>	<b>8</b>
<b>DRB1 allele<sup>4,5</sup></b>								
*09:01:02-09:01:07	1	2						
*09:02:01-09:02:02	1		3					
*09:03, 09:09, 09:13, 09:15, 09:17	1	2		4				
*09:04	1	2			5			
*09:05		2				6		
*09:06	1	2					7	
*09:07, 09:12	1	2						8
*09:08	1				5			
*09:10-09:11, 09:14	1	2				6		
*09:16		2		4				
<b>DRB1 allele<sup>4</sup></b>								
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>	<b>7</b>	<b>8</b>