

<b>INTERPRETATION TABLE</b>					
<b>DRB1*10 SSP subtyping</b>					
<b>Amplification patterns of the DRB1*10 alleles</b>					
	<b>Well</b>				
	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>
<b>Length of spec.</b>	<b>210</b>	<b>205</b>	<b>205</b>	<b>70</b>	<b>160</b>
<b>PCR product</b>					
<b>Length of int.</b>	<b>515</b>	<b>515</b>	<b>430</b>	<b>430</b>	<b>430</b>
<b>pos. control<sup>1</sup></b>					
<b>5'-primer<sup>2</sup></b>	<b>30(178)</b>	<b>30(178)</b>	<b>30(178)</b>	<b>30(178)</b>	<b>30(178)</b>
	5' -gCg 3'	5' -gCg 3'	5' -gCg 3'	5' -gCg 3'	5' -gCg 3'
<b>3'-primer<sup>3</sup></b>	<b>86(346)</b>	<b>86(344)</b>	<b>86(344)</b>	<b>40(206)</b>	<b>70(296)</b>
	5' -CTC 3'	5' -CAC 3'	5' -CCA 3'	5' -CgA 3'	5' -TCT 3'
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>
<b>DRB1 allele<sup>4</sup></b>					
<b>*10:01:01-10:01:03</b>	<b>1</b>	<b>2</b>			
<b>*10:02</b>	<b>1</b>		<b>3</b>		
<b>*10:03</b>	<b>1</b>	<b>2</b>		<b>4</b>	
<b>*10:04</b>	<b>1</b>	<b>2</b>			<b>5</b>
<b>DRB1 allele</b>					
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>