

<b>INTERPRETATION TABLE</b>						
<b>DQB1*05 SSP subtyping</b>						
<b>Amplification patterns of the DQB1*05 alleles</b>						
	<b>Well</b>					
	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>
<b>Length of spec.</b>	<b>225</b>	<b>135</b>	<b>120</b>	<b>95</b>	<b>120</b>	<b>185</b>
<b>PCR product</b>						
<b>Length of int.</b>	<b>515</b>	<b>430</b>	<b>430</b>	<b>515</b>	<b>430</b>	<b>430</b>
<b>pos. control<sup>1</sup></b>						
<b>5'-primer<sup>2</sup></b>	<b>26(173)</b>	<b>26(173)</b>	<b>30(184)</b>	<b>30(184)</b>	<b>30(184)</b>	<b>38(209)</b>
	5'-ggg 3'	5'-ggg 3'	5'-gAC 3'	5'-gAC 3'	5'-gAT 3'	5'-CgC 3'
<b>3'-primer<sup>3</sup></b>	<b>87(356)</b>	<b>57(266)</b>	<b>57(265)</b>	<b>47(237)</b>	<b>57(265)</b>	<b>86(353)</b>
	5'-ggT 3'	5'-CAA 3'	5'-gCT 3'	5'-CgA 3'	5'-gCT 3'	5'-ACg 3'
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>
<b>DQB1 allele</b>						
<b>*050101, 050102</b>	<b>1</b>	<b>2</b>				
<b>*050201</b>	<b>1</b>		<b>3</b>			
<b>*050202</b>	<b>1</b>		<b>3</b>	<b>4</b>		
<b>*050301, 050302</b>	<b>1</b>			<b>4</b>		
<b>*0504</b>	<b>1</b>				<b>5</b>	
<b>*0505</b>	<b>1</b>		<b>3</b>			<b>6</b>
<b>DQB1 allele</b>						
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>