

<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</b>	<b>26</b>	<b>30</b>	<b>30</b>	<b>30</b>	<b>38</b>
	5'-g gg <sup>3'</sup>	5'-g gg <sup>3'</sup>	5'-gA C <sup>3'</sup>	5'-gA C <sup>3'</sup>	5'-gA T <sup>3'</sup>	5'-C gC <sup>3'</sup>
<b>3'-primer<sup>3</sup></b>	<b>87</b>	<b>57</b>	<b>57</b>	<b>47</b>	<b>57</b>	<b>86</b>
	5'-g gT <sup>3'</sup>	5'-C AA <sup>3'</sup>	5'-gCT <sup>3'</sup>	5'-Cg A <sup>3'</sup>	5'-gCT <sup>3'</sup>	5'-A Cg <sup>3'</sup>
<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>