

<b>INTERPRETATION TABLE</b>				
<b>DQB1*04 SSP subtyping</b>				
<b>Amplification patterns of the DQB1*04 alleles</b>				
	<b>Well</b>			
	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>
<b>Length of spec.</b>	<b>210</b>	<b>205</b>	<b>205</b>	<b>195</b>
<b>PCR product</b>				
<b>Length of int.</b>	<b>515</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>21(159)</b>	<b>23(164)</b>	<b>23(164)</b>	<b>26(173)</b>
	5'-ACC 3'	5'-gCT 3'	5'-gCg 3'	5'-TCT 3'
<b>3'-primer<sup>3</sup></b>	<b>77(327)</b>	<b>77(327)</b>	<b>77(327)</b>	<b>77(327)</b>
	5'-ACg 3'	5'-ACg 3'	5'-ACg 3'	5'-ACg 3'
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>
<b>DQB1 allele</b>				
<b>*0401</b>	<b>1</b>	<b>2</b>		
<b>*0402</b>	<b>1</b>		<b>3</b>	
<b>*0403</b>	<b>1</b>		<b>3</b>	<b>4</b>
<b>*0306</b>				<b>4</b>
<b>DQB1 allele</b>				
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>