

<b>INTERPRETATION TABLE</b>					
<b>HLA-Cw*18 SSP subtyping</b>					
<b>Amplification patterns of the Cw*18 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>425</b>	<b>560</b>	<b>220</b>	<b>530</b>	<b>165</b>
<b>PCR product</b>					
<b>Length of int.</b>	<b>800</b>	<b>1070</b>	<b>1070</b>	<b>1070</b>	<b>1070</b>
<b>pos. control<sup>1</sup></b>					
<b>5'-primer<sup>2</sup></b>	<b>47</b>	<b>956</b>	<b>895</b>	<b>213</b>	<b>412</b>
	5'-Agg3'	5'-ggT3'	5'-ggA3'	5'-CCC3'	5'-ATA3'
<b>3'-primer<sup>3</sup></b>	<b>302</b>	<b>1034</b>	<b>956</b>	<b>459</b>	<b>538</b>
	5'-ggT3'	5'-AgC3'	5'-CAg3'	5'-ACgA3'	5'-CCA3'
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>
<b>HLA-Cw allele</b>					
<b>*1801</b>	<b>1</b>	<b>2</b>		<b>4</b>	
<b>*1802</b>	<b>1</b>		<b>3</b>	<b>4</b>	
<b>*1803</b>	<b>1</b>	<b>?</b>	<b>?</b>	<b>4</b>	<b>5</b>
<b>*04010101-040105, 0403-0420, 0423-0432</b>			<b>3</b>		
<b>*0508, 0609</b>					<b>5</b>
<b>*0617</b>	<b>1</b>				
<b>*070101-0706, 0708, 0710-0733N, 0735-0748, 0750-0754</b>		<b>2</b>			
<b>*0707, 0709, 0749</b>	<b>1</b>	<b>2</b>			
<b>HLA-Cw allele</b>					
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>
<b>B*9537</b>					<b>5</b>
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>