

<b>INTERPRETATION TABLE</b>					
<b>HLA-Cw*18 SSP subtyping</b>					
<b>Amplification patterns of the Cw*18 alleles</b>					
	<b>Tube</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>395</b>	<b>165</b>
<b>PCR product</b>					
<b>Length of int.</b>	<b>800</b>	<b>1070</b>	<b>1070</b>	<b>800</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>47</b>	<b>412</b>
	5'-Agg <sup>3'</sup>	5'-ggT <sup>3'</sup>	5'-ggA <sup>3'</sup>	5'-Agg <sup>3'</sup>	5'-ATA <sup>3'</sup>
<b>3'-primer<sup>3</sup></b>	<b>302</b>	<b>1034</b>	<b>956</b>	<b>270</b>	<b>538</b>
	5'-ggT <sup>3'</sup>	5'-AgC <sup>3'</sup>	5'-CAg <sup>3'</sup>	5'-TAC <sup>3'</sup>	5'-CCA <sup>3'</sup>
<b>Tube No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>
<b>HLA-Cw allele</b>					
<b>*1801, 0749<sup>4</sup></b>	<b>+</b>	<b>+</b>		<b>+</b>	
<b>*1802</b>	<b>+</b>		<b>+</b>	<b>+</b>	
<b>*1803</b>	<b>+</b>	<b>?</b>	<b>?</b>	<b>+</b>	<b>+</b>
<b>*04010101-040104, 0403, 040401-0421, 0423-0427</b>			<b>+</b>		
<b>*0508, 0609</b>					<b>+</b>
<b>*070101-070107, 0706, 0716, 0718-0722, 0724, 0726, 0728, 0730, 0735, 0736, 0740, 0744</b>		<b>+</b>			
<b>07020101-0705, 0708, 0710- 0715, 0717, 0723, 0725, 0727, 0729, 0731-0734, 0737-0739, 0741-0743, 0745-0748</b>		<b>+</b>		<b>+</b>	
<b>*0707, 0709</b>	<b>+</b>	<b>+</b>			
<b>HLA-Cw allele</b>					
<b>Tube No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>