

<b>INTERPRETATION TABLE</b>	
<b>HLA-C*18 SSP subtyping</b>	
<b>Amplification patterns of the C*18:01 to 18:04 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>425</b>	<b>560</b>	<b>225</b>	<b>535</b>	<b>165</b>	<b>265</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>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>	<b>47</b>
	5'-Agg 3'	5'-ggT 3'	5'-ggA 3'	5'-CCC 3'	5'-ATA 3'	5'-Agg 3'
<b>3'-primer<sup>3</sup></b>	<b>302</b>	<b>1034</b>	<b>956</b>	<b>459</b>	<b>538</b>	<b>142</b>
	5'-ggT 3'	5'-AgC 3'	5'-CAg 3'	5'-AgA 3'	5'-CCA 3'	5'-TgC 3'
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>
<b>HLA-C allele</b>						
<b>*18:01</b>	<b>1</b>	<b>2</b>		<b>4</b>		
<b>*18:02</b>	<b>1</b>		<b>3</b>	<b>4</b>		
<b>*18:03</b>	<b>1</b>	<b>?</b>	<b>?</b>	<b>4</b>	<b>5</b>	
<b>*18:04</b>	<b>1</b>	<b>?</b>	<b>?</b>	<b>4</b>		<b>6</b>
<b>*02:22, 05:08, 06:09, 08:27, 08:29, 08:31, 12:31, B*15:137</b>					<b>5</b>	
<b>*04:01:01-04:01:26, 04:03-04:20, 04:23-04:81</b>			<b>3</b>			
<b>*06:02:08</b>				<b>4</b>		
<b>*06:17</b>	<b>1</b>					
<b>*07:01:01-07:06, 07:08, 07:10-07:19, 07:21-07:33N, 07:35-07:48, 07:50-07:63, 07:65-07:72, 07:74-07:75, 07:77-07:91, 07:93-07:95, 07:97-07:160</b>		<b>2</b>				
<b>*07:07, 07:09, 07:49, 07:76</b>	<b>1</b>	<b>2</b>				
<b>*07:20, 07:64, 07:73, 07:92, 07:96</b>		<b>2</b>				<b>6</b>
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>