

<b>INTERPRETATION TABLE</b>							
<b>HLA-A*74 SSP subtyping</b>							
<b>Amplification patterns of the A*7401 to 7414N alleles</b>							
	<b>Well</b>						
	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>	<b>7</b>
<b>Length of spec.</b>	<b>160</b>	<b>285</b>	<b>285</b>	<b>135</b>	<b>230</b>	<b>155</b>	<b>150</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>800</b>
<b>pos. control<sup>1</sup></b>							
<b>5'-primer<sup>2</sup></b>	<b>180</b>	<b>67</b>	<b>67</b>	<b>144</b>	<b>98</b>	<b>180</b>	<b>180</b>
	5' -TTT 3'	5' -CCA 3'	5' -CCT 3'	5' -gCC 3'	5' -CTT 3'	5' -TTT 3'	5' -TTT 3'
<b>3'-primer(s)<sup>3</sup></b>	<b>299</b>	<b>180</b>	<b>180</b>	<b>240</b>	<b>289</b>	<b>292</b>	<b>290</b>
	5' -CCA 3'	5' -TCA 3'	5' -TCA 3'	5' -ggA 3'	5' -AgC 3'	5' -gTg 3'	5' -CAA 3'
	<b>299</b>						
	5' -CCA 3'						
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>	<b>7</b>
<b>HLA-A allele</b>							
<b>*7401</b>	<b>1</b>	<b>2</b>					
<b>*7402</b>	<b>1</b>		<b>3</b>				
<b>*7403</b>	<b>1</b>	<b>2</b>					
<b>*7404</b>	<b>1</b>	<b>?</b>	<b>?</b>	<b>4</b>			
<b>*7405</b>	<b>1</b>	<b>?</b>	<b>?</b>		<b>5</b>		
<b>*7406</b>	<b>1</b>	<b>?</b>	<b>?</b>			<b>6</b>	
<b>*7407</b>	<b>1</b>		<b>3</b>				<b>7</b>
<b>*7408</b>	<b>1</b>	<b>?</b>	<b>?</b>				
<b>*7409</b>	<b>1</b>	<b>?</b>	<b>?</b>				
<b>*7410</b>		<b>?</b>	<b>?</b>				
<b>*7411</b>	<b>1</b>	<b>?</b>	<b>?</b>				
<b>*7412N</b>	<b>1</b>	<b>?</b>	<b>?</b>				
<b>*7413</b>	<b>1</b>	<b>?</b>	<b>?</b>				
<b>*7414N</b>	<b>1</b>	<b>?</b>	<b>?</b>				
<b>*020109, 023503, 0323, 260702, 3308, 3309, 6830</b>				<b>4</b>			
<b>*030103</b>				<b>w</b>			
<b>*29010101-2903, 2911, 320101-3202, 3205, 3217, 3219N</b>			<b>3</b>				
<b>*2904-2910, 2912-2918, 3203, 3206-3216, 3218</b>			<b>?</b>				
<b>*2919</b>	<b>1</b>		<b>?</b>			<b>6</b>	
<b>HLA-A allele</b>							
<b>Well No.</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>	<b>7</b>

<b>INTERPRETATION</b>							
<b>HLA-A*74 SSP s</b>							
<b>Amplification patterns of the A*</b>							
<b>Well</b>							
<b>8</b>	<b>9</b>	<b>10</b>	<b>11</b>	<b>12</b>	<b>13</b>	<b>14</b>	<b>15</b>
<b>115</b>	<b>170</b>	<b>225</b>	<b>200</b>	<b>130</b>	<b>215</b>	<b>145</b>	<b>85</b>
<b>1070</b>	<b>800</b>	<b>1070</b>	<b>800</b>	<b>1070</b>	<b>1070</b>	<b>1070</b>	<b>1070</b>
<b>448</b>	<b>180</b>	<b>113</b>	<b>180</b>	<b>211</b>	<b>364</b>	<b>180</b>	<b>180</b>
5'-CCT 3'	5'-TTT 3'	5'-CCC 3'	5'-TTT 3'	5'-AgT 3'	5'-gA 3'	5'-TTT 3'	5'-TTT 3'
<b>523</b>	<b>308</b>	<b>299</b>	<b>341</b>	<b>299</b>	<b>538</b>	<b>282</b>	<b>225</b>
5'-ACT 3'	5'-TCg 3'	5'-CCA 3'	5'-CgT 3'	5'-CCA 3'	5'-CAA 3'	5'-gAC 3'	5'-ATT 3'
<b>8</b>	<b>9</b>	<b>10</b>	<b>11</b>	<b>12</b>	<b>13</b>	<b>14</b>	<b>15</b>
	<b>9</b>						
<b>8</b>							
		<b>10</b>					
			<b>11</b>				
				<b>12</b>			
					<b>13</b>		
						<b>14</b>	
	<b>9</b>						<b>15</b>
<b>8</b>	<b>9</b>	<b>10</b>	<b>11</b>	<b>12</b>	<b>13</b>	<b>14</b>	<b>15</b>

<b>N TABLE</b>	
<b>Subtyping</b>	
<b>7401 to 7414N alleles</b>	
	<b>Length of spec.</b>
	<b>PCR product</b>
	<b>Length of int.</b>
	<b>pos. control<sup>1</sup></b>
	<b>5'-primer<sup>2</sup></b>
	<b>3'-primer(s)<sup>3</sup></b>
	<b>Well No.</b>
	<b>HLA-A allele</b>
	<b>*7401</b>
	<b>*7402</b>
	<b>*7403</b>
	<b>*7404</b>
	<b>*7405</b>
	<b>*7406</b>
	<b>*7407</b>
	<b>*7408</b>
	<b>*7409</b>
	<b>*7410</b>
	<b>*7411</b>
	<b>*7412N</b>
	<b>*7413</b>
	<b>*7414N</b>
	<b>*020109, 023503, 0323, 260702, 3308, 3309, 6830</b>
	<b>*030103</b>
	<b>*29010101-2903, 2911, 320101- 3202, 3205, 3217, 3219N</b>
	<b>*2904-2910, 2912-2918, 3203, 3206-3216, 3218</b>
	<b>*2919</b>
	<b>HLA-A allele</b>
	<b>Well No.</b>