





*26:62-26:63 <sup>13</sup>	1	2																							
*26:64	1	2																							
*26:65	1	2																							
*26:66	1	2																							
*26:68	1	2																							
*26:71N	1	2																							
*26:72	1																								
*26:76	1																								
*26:78				4	5																				
*26:82	1	2																					21	23	
*01:01:13, 01:83:01-01:83:02, 02:146									10																
*01:02, 24:04, 24:109, 29:37, 30:57										12															
*01:03, 36:03, 74:10, C*06:72																								24	
*01:13, 01:17, 03:63, 03:88, 11:01:01-11:05, 11:07-11:11, 11:13-11:16, 11:20-11:25, 11:27, 11:29-11:39, 11:41-11:52Q, 11:54-11:90, 11:92-11:95, 11:97, 11:99N-11:105, 11:107-11:115N, 11:117, 11:120, 11:122-11:133, 11:135-11:139														15											
*01:20										12								w							
*01:51, 33:48		2																							
*01:60, C*06:71																		18							
*01:66																		w							
*01:104, 02:346, 80:01:01:01																									
*02:34, 02:56:01-02:56:02, 02:62, 02:103																		w							
*02:35:01-02:35:03, 02:78, 24:19, 68:01:01:01-68:02:05, 68:06-68:14, 68:16-68:19, 68:21:01-68:30, 68:32-68:35, 68:37-68:56, 68:58-68:60, 68:62-68:83, 68:85, 68:87-68:88, 68:91-68:93, 68:95-68:99, 69:01																			16						
<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>8</b>	<b>9</b>	<b>10</b>	<b>11</b>	<b>12</b>	<b>13</b>	<b>14</b>	<b>15</b>	<b>16</b>	<b>17</b>	<b>18</b>	<b>19</b>	<b>20</b>	<b>21</b>	<b>22</b>	<b>23</b>	<b>24</b>	
<b>Length of spec.</b>	<b>##</b>	<b>80</b>	<b>##</b>	<b>##</b>	<b>##</b>	<b>80</b>	<b>##</b>	<b>##</b>	<b>##</b>	<b>85</b>	<b>##</b>	<b>##</b>	<b>##</b>	<b>##</b>	<b>##</b>	<b>##</b>	<b>##</b>	<b>##</b>	<b>##</b>	<b>##</b>	<b>##</b>	<b>##</b>	<b>##</b>	<b>##</b>	
<b>PCR product(s)</b>						<b>##</b>			<b>##</b>	<b>##</b>		<b>##</b>	<b>##</b>					<b>##</b>	<b>##</b>		<b>##</b>	<b>##</b>	<b>##</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>8</b>	<b>9</b>	<b>10</b>	<b>11</b>	<b>12</b>	<b>13</b>	<b>14</b>	<b>15</b>	<b>16</b>	<b>17</b>	<b>18</b>	<b>19</b>	<b>20</b>	<b>21</b>	<b>22</b>	<b>23</b>	<b>24</b>	
*02:38, 02:101:01													14					19							





# INTERPRETATION TABLE

## HLA-A\*26 SSP subtyping

Amplification patterns of the A\*26:01 to 26:84 alleles

Well<sup>14</sup>

		25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	
		360	150	125	90	330	125	210	90	105	135	110	125	245	190	220	90	140	460	135	105	365	155	425	Length of spec.
		###	###	###	###	###	###	800	###	###	###	###	###	800	###	###	###	###	###	###	###	###	###	###	PCR product(s)
		###	###	###	###	###	###	###	###	###	###	###	###	###	###	###	###	###	###	###	###	###	###	###	Length of int.
		5'-ggC <sup>3'</sup>	5'-CTA <sup>3'</sup>	5'-CCA <sup>3'</sup>	5'-CTC <sup>3'</sup>	5'-TCg <sup>3'</sup>	5'-CTA <sup>3'</sup>	5'-ACC <sup>3'</sup>	5'-gCT <sup>3'</sup>	5'-CCA <sup>3'</sup>	5'-gCT <sup>3'</sup>	5'-gCT <sup>3'</sup>	5'-gCT <sup>3'</sup>	5'-TCg <sup>3'</sup>	5'-CgA <sup>3'</sup>	5'-TCg <sup>3'</sup>	5'-AAC <sup>3'</sup>	5'-CCT <sup>3'</sup>	5'-TCg <sup>3'</sup>	5'-gCT <sup>3'</sup>	5'-gCT <sup>3'</sup>	5'-TCg <sup>3'</sup>	5'-gCT <sup>3'</sup>	5'-TCg <sup>3'</sup>	pos. control <sup>1</sup>
		341	98	106	292	28	98	391	423	401	423	136	423	28	191	28	261	2 <sup>nd</sup>	28	423	423	28	423	28	5'-primer(s) <sup>2</sup>
		5'-gCT <sup>3'</sup>	5'-gCT <sup>3'</sup>	5'-gAg <sup>3'</sup>			5'-CTg <sup>3'</sup>			5'-CAC <sup>3'</sup>		5'-AAA <sup>3'</sup>		5'-CAG <sup>3'</sup>											
		418	208	341	341	186	180	559	472	559	518	341	506	265	341	77	311	426	319	517	486	224	538	283	3'-primer(s) <sup>3</sup>
		5'-gTC <sup>3'</sup>	5'-CCA <sup>3'</sup>	5'-CgT <sup>3'</sup>	5'-CgT <sup>3'</sup>	5'-TCC <sup>3'</sup>	5'-TCC <sup>3'</sup>	5'-CCg <sup>3'</sup>	5'-ggC <sup>3'</sup>	5'-CCg <sup>3'</sup>	5'-CCA <sup>3'</sup>	5'-CgT <sup>3'</sup>	5'-TgT <sup>3'</sup>	5'-CCA <sup>3'</sup>	5'-CgT <sup>3'</sup>	5'-ggC <sup>3'</sup>	5'-ggA <sup>3'</sup>	5'-TCC <sup>3'</sup>	5'-gCg <sup>3'</sup>	5'-CgC <sup>3'</sup>	5'-gCg <sup>3'</sup>	5'-TCT <sup>3'</sup>	5'-CAG <sup>3'</sup>	5'-TgC <sup>3'</sup>	
		559					743		538		570	595	559			271									
		5'-CCA <sup>3'</sup>					5'-TCC <sup>3'</sup>		5'-CAA <sup>3'</sup>		5'-CCg <sup>3'</sup>	5'-CCA <sup>3'</sup>	5'-CCg <sup>3'</sup>	5'-CCg <sup>3'</sup>	5'-CAT <sup>3'</sup>	5'-CCA <sup>3'</sup>									
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47		Well No.	
																								HLA-A allele <sup>4,5</sup>	
																								*26:01:01-26:01:27, 26:27, 26:74-26:75, 26:77, 26:79-26:80, 26:83- 26:84 <sup>15</sup>	
			28																					*26:02	
																								*26:03:01-26:03:02 *26:04	
			28																					*26:05 *26:06	
																								*26:07:01-26:07:02 *26:08, 26:67, 26:81	
																								*26:09 *26:10 *26:11N	
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47		Well No.	
##	##	##	90	##	##	##	90	##	##	##	##	##	##	##	90	##	##	##	##	##	##	##	##	Length of spec.	









