

101.702-24/06 – including *Taq* pol., IFU-01
101.702-24u/06u – without *Taq* pol., IFU-02

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“Instructions for Use” (IFU)

Lot No.: **3D7**

Lot-specific information

Olerup SSP® HLA-A-B-C SSP Combi Tray

Product number:	101.702-24/06 – including <i>Taq</i> pol. 101.702-24u/06u – without <i>Taq</i> pol.
Lot number:	3D7
Expiry date:	2018-June-01
Number of tests:	24 tests – Product No. 101.702-24/24u 6 tests – Product No. 101.702-06/06u
Number of wells per test:	95+1
Storage - pre-aliquoted primers:	dark at -20°C
- PCR Master Mix:	-20°C
- Adhesive PCR seals	RT
- Product Insert	RT

This Product Description is only valid for Lot No. 3D7.

Complete product documentation consists of generic Instructions for Use (IFU), lot specific Product Insert, Worksheet and Certificate.

CHANGES COMPARED TO THE PREVIOUS OLERUP SSP® HLA-A-B-C SSP COMBI TRAY LOT (48Y)

The format of the Product Insert and Worksheet have been changed.

As of lot series V, the Specificity Table is included in the lot-specific Product Insert, and the Interpretation Table is included in the Worksheet.

The **HLA-A low resolution** specificity and interpretation tables have been updated for the HLA-A alleles described since the previous *Olerup SSP®* HLA-A-B-C SSP Combi Tray lot was made (**Lot No. 48Y**). The kit design is based on IMGT/HLA database 3.21.1.

The primers of the wells detailed below have been exchanged, added or modified compared to the previous lot.

Well	5'-primer	3'-primer	rationale
3	Added	-	5'-primers added for the A*03:204 and for the A*03:01:49 alleles.
22	Added	-	5'-primer added for the A*68:11N allele.

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The **HLA-B low resolution** specificity and interpretation tables have been updated for the HLA-B alleles described since the previous *Olerup* SSP[®] HLA-A-B-C SSP Combi Tray lot was made (**Lot No. 48Y**). The kit design is based on IMGT/HLA database 3.21.1.

The primers of the wells detailed below have been exchanged, added or modified compared to the previous lot.

Well	5'-primer	3'-primer	rationale
28	-	Added	3'-primer added for the B*14:46 allele.
55	-	Added	3'-primer added for the B*15:297 allele.

The **HLA-C low resolution** specificity and interpretation tables have been updated for the HLA-C alleles described since the previous *Olerup* SSP[®] HLA-A-B-C SSP Combi Tray lot was made (**Lot No. 48Y**). The kit design is based on IMGT/HLA database 3.21.1.

The primers of the wells detailed below have been exchanged, added or modified compared to the previous lot.

Well	5'-primer	3'-primer	rationale
80	-	Added	3'-primer added for the C*15:107 allele.
95	Added	-	5'-primer added for the C*16:67 allele.

Changes in revision R01 compared to R00:

1. Primer mix 34 amplifies the C*02:06 and C*15:42 alleles and primer mix 65 amplifies the B*35:291 and the C*15:15 alleles. This has been corrected in the Specificity and Interpretation Tables.
2. Primer mix 21 may generate a false positive band of about 560 base pairs. This band should be disregarded when interpreting HLA-A low resolution typings. A footnote has been added in the Specificity Table.

Change in revision R02 compared to R01:

1. Primer mix 51 does not amplify the B*08:31, 41:06, 41:15 and 42:14 alleles, and primer mix 72 does not amplify the B*08:31 allele. This has been corrected in the Specificity and Interpretation Tables.

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Well **96** contains Negative Control primer pairs, that will amplify more than 95% of the *Olerup* SSP[®] HLA Class I, DRB, DQB1, DPB1 and DQA1 amplicons as well as all the amplicons generated by the control primer pairs matching the human growth hormone gene.

HLA-specific PCR product sizes range from 75 to 200 base pairs.
The PCR product generated by the positive control primer pair is 430 base pairs.

Length of PCR product	105	200	105	80	75	80	85
5'-primer¹	164	340	440	45	45	43	36
	5'-CAC ^{3'}	5'-Agg ^{3'}	5'-TTA ^{3'}	5'-Tgg ^{3'}	5'-Tgg ^{3'}	5'-Tgg ^{3'}	5'-TAC ^{3'}
							36
							5'-TAT ^{3'}
3'-primer²	231	2nd I	507	59	58	57	47
	5'-TgC ^{3'}	5'-AAA ^{3'}	5'-TTg ^{3'}	5'-CTC ^{3'}	5'-ggC ^{3'}	5'-CTC ^{3'}	5'-ACA ^{3'}
							48
							5'-gCA ^{3'}
							48
							5'-gCC ^{3'}
							52
							5'-TgT ^{3'}
A*	+	+	+				
B*	+	+	+				
C*	+	+	+				
DRB1				+	+		
DRB3				+	+		
DRB5				+			
DQB1					+		
DPB1						+	
DQA1							+

¹The nucleotide position for HLA class I genes and the codon for HLA class II genes, in the 2nd or 3rd exon, matching the specificity-determining 3'-end of the primer is given. Nucleotide and codon numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given.

²The nucleotide position for HLA class I genes and the codon for HLA class II genes, in the 2nd or 3rd exon or the 2nd intron, matching the specificity-determining 3'-end of the primer is given in the anti-sense direction. Nucleotide and codon numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given.

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PRODUCT DESCRIPTION

HLA-A-B-C SSP Combi Tray

CONTENT

The primer set contains 5'- and 3'-primers for grouping the HLA-A*01:01 to A*80:03 alleles into the corresponding serological groups A1 to A80.

The primer set contains 5'- and 3'-primers for grouping the B*07:02 to B*83:01 alleles into the corresponding serological groups B7 to B81 as well as primer pairs for recognizing the Bw4 and Bw6 sequence motifs.

The primer set contains 5'- and 3'-primers for grouping the HLA-C*01:02 to C*18:09 alleles into the groups C*01:xx to C*18:xx.

PLATE LAYOUT

Each test consists of 96 PCR reactions in a 96 well PCR plate.

1	2	3	4	5	6	7	8
9	10	11	12	13	14	15	16
17	18	19	20	21	22	23	24
25	26	27	28	29	30	31	32
33	34	35	36	37	38	39	40
41	42	43	44	45	46	47	48
49	50	51	52	53	54	55	56
57	58	59	60	61	62	63	64
65	66	67	68	69	70	71	72
73	74	75	76	77	78	79	80
81	82	83	84	85	86	87	88
89	90	91	92	93	94	95	NC

The 96 well PCR plate is marked with 'A-B-C' in silver/gray ink.

Well No. 1 is marked with the Lot No. '3D7'.

Wells 1 to 24 – HLA-A low resolution primers.

Wells 25 to 72 – HLA-B low resolution primers.

Wells 73 to 95 – HLA-C low resolution primers.

Well 96 – Negative Control.

A faint row of numbers is seen between wells 1 and 2 or wells 7 and 8 of the PCR trays. These stem from the manufacture of the trays, and should be disregarded.

The PCR plates are covered with a PCR-compatible foil.

INTERPRETATION

Only HLA-A alleles will be amplified by the 24 wells of the HLA-A low resolution primer set, **wells 1 to 24**, except that a few HLA-B and HLA-C alleles will be amplified by primer mixes 4 to 6, 9 and 11.

For further details see Specificity Table.

Only HLA-B alleles will be amplified by the 48 wells of the HLA-B low resolution primer set, **wells 25 to 72**, except that a few HLA-A and HLA-C alleles will be amplified by primer mixes 25 to 28, 30 to 32, 37, 38, 41, 43 to 46, 49, 50, 53 to 55, 64 to 67, 69 and 72.

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Lot-specific information

For further details see Specificity Table.

Only HLA-C alleles will be amplified by the 23 wells of the HLA-C low resolution primer set, **wells 73 to 95**, except that a few HLA-A and HLA-B alleles will be amplified by primer mixes 73, 80, 85, 87, 88, 92, 94 and 95.
For further details see Specificity Table.

UNIQUELY IDENTIFIED ALLELES

All the HLA-A alleles, i.e. **A*01:01 to A*80:03**, recognized by the HLA Nomenclature Committee in August 2015^{1,2} will be amplified by the primers in the HLA-A low resolution primer set, **wells 1 to 24**. The HLA-A alleles will be grouped into their corresponding serological specificities^{3,4}.

All the HLA-B alleles, i.e. **B*07:02 to B*83:01**, recognized by the HLA Nomenclature Committee in August 2015^{1,2} will be amplified by the primers in the HLA-B low resolution primer set, **wells 25 to 72**. The HLA-B alleles will be grouped into their corresponding serological specificities³.

All the HLA-C alleles, i.e. **C*01:02 to C*18:09**, recognized by the HLA Nomenclature Committee in August 2015^{1,2} will be amplified by the primers in the HLA-C low resolution primer set, **wells 73 to 95**. The HLA-C alleles will be grouped into the C*01:xx to C*18:xx groups³.

¹HLA-A, HLA-B and HLA-C alleles listed on the IMGT/HLA web page 2015-August-11, release 3.21.1, www.ebi.ac.uk/imgt/hla.

²Alleles that have been deleted from or renamed in the official WHO HLA Nomenclature up to and including the last IMGT/HLA database release can be retrieved from web page <http://hla.alleles.org/alleles/deleted.html>.

³The HLA-A, HLA-B, HLA-C alleles will be grouped into their corresponding serological specificities, except that following alleles give rise to identical amplification patterns.

These alleles can be separated by the full A-B-C primer set or by the respective high resolution SSP primer sets.

Alleles	Alleles
A*03:152, A*24:92	B*41:06, 41:09, 41:29, 41:32, B*45:02-45:03
A*23:14:01-23:14:02, A*24:24, 24:71	B*53:30, 53:39, B*57:45, 57:51, 57:69
A*23:57, 23:66, A*24:94, 24:138, 24:188, 24:228	B*55:04, 55:08, 55:13, 55:27, 55:46, 55:49, B*56:01:05, 56:15, 56:19N, 56:22
A*31:08, A*33:53	B*55:23, 55:32, 56:18, 56:31-56:32
A*32:01:18, B*07:81, B*08:52, B*18:67, B*38:41, B*51:185, B*53:05, 53:16, 53:33	B*57:67:01, B*58:36
B*07:202, B*81:02	C*01:05, 01:07:02, 01:22, 01:35-01:36, 01:79:01-01:79:02, B*39:76, B*54:18
B*08:26:01-08:26:02, 08:50, 08:62, 08:85, 08:94, B*42:07	C*01:67, C*14:58
B*13:21, 13:35, 13:59, 13:71, B*44:135, 44:158, 44:184	C*12:18:02, B*67:02
B*14:08:01-14:08:02, B*39:01:19, 39:25N, 39:30, 39:32-39:34, 39:47, 39:50, 39:74, 39:82, 39:102	C*14:06, 14:08-14:09, 14:28:01-14:28:02, 14:53, 14:63, A*30:62, B*35:252
B*18:29, 18:72, 18:92, 18:102, B*35:32:01-35:32:02, 35:37, 35:53N, 35:64:01-35:64:02, 35:68:01-35:68:02, 35:99, 35:118-35:119, 35:174	

⁴The A*01:136 and the A*11:94, 11:112 and 11:211 alleles may be distinguished by the different sizes of the specific PCR products generated by primer mixes 1 and 11.

The A*03:88 and the A*11:130 alleles may be distinguished by the different sizes of the specific PCR products generated by primer mixes 11.

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The A*11:199:01 and 11:222 and the A*66:23 alleles may be distinguished by the different sizes of the specific PCR products generated by primer mix 11.

The A*24:14-24:15, 24:51-24:53, 24:57, 24:64, 24:114, 24:222N, 24:291 and 24:296 and the C*04:01:03 alleles may be distinguished by the different sizes of the specific PCR products generated by primer mix 5.

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SPECIFICITY TABLE

HLA-A low resolution primer set

Specificities and sizes of the PCR products of the 24 primer mixes used for HLA-A low resolution SSP typing

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	HLA-A serology ³	Amplified HLA-A alleles ^{4,5}
1^{6,7}	120 bp, 145 bp, 225 bp	800 bp	A1, A36	*01:01:01:01-01:04N, 01:06-01:33, 01:35-01:175, 01:178N-01:189, 03:18, 03:135, 11:94, 11:112, 11:211, 36:01-36:05
2⁷	175 bp, 215 bp, 255 bp, 365 bp, 415 bp	800 bp	A2, A203, A210, A19, A28	*02:01:01:01-02:01:15, 02:01:17-02:01:19, 02:01:21-02:01:81, 02:01:83-02:06:18, 02:06:20-02:22:02, 02:24:01-02:35:01, 02:35:03-02:47, 02:49-02:77, 02:78 ^w , 02:79:01-02:97:02, 02:99, 02:101:01-02:128, 02:130-02:570, 02:572-02:580
3⁸	205 bp, 235 bp	1070 bp	A1, A3, A11, A32, A34, A36	*01:12, 01:19, 01:21, 01:126, 02:338, 03:01:01:01-03:17:02, 03:19-03:74, 03:76-03:94, 03:96-03:134, 03:136-03:176, 03:178N-03:186, 03:188-03:214, 03:216-03:219, 03:221-03:224, 03:226-03:229, 11:03, 11:20, 11:25:01, 11:60, 11:130, 11:158, 11:175, 11:183, 11:209, 24:92, 32:04, 34:02:01-34:04, 34:07-34:10N, 36:02, 74:23
4	190 bp	800 bp	A1, A3, A11, A30, A36, A68	*01:01:01:01-01:01:22, 01:01:24-01:01:47, 01:01:49-01:01:64, 01:01:67-01:04N, 01:06-01:07, 01:09-01:11N, 01:13, 01:16N-01:18N, 01:20-01:29, 01:31N-01:33, 01:35-01:78, 01:80-01:98, 01:100-01:144, 01:146, 01:148, 01:150-01:158, 01:160N-01:166, 01:168-01:175, 01:180-01:189, 02:78, 02:169, 03:12, 03:18, 03:88, 03:135, 11:01:01:01-11:27, 11:29-11:52Q, 11:54-11:223, 26:19, 26:72, 29:67, 30:08, 32:64, 36:04, 66:23, 68:13:01, 68:66, 68:134, 74:19, C*12:131
5	160 bp, 335 bp, 505 bp	1070 bp	A3, A9, A23, A24, A2403, A29, A31, A32	*03:15, 03:19, 03:30, 03:152, 11:139, 23:01:01-23:69, 24:02:01:01-24:02:92, 24:02:95-24:11N, 24:13:01-24:15, 24:17-24:64, 24:66-24:210, 24:212-24:303N, 24:305-24:314, 29:07, 29:49, 31:08, 31:29, 32:05, 33:19, 33:53, C*04:01:03
6^{9,11}	135 bp, 200 bp	800 bp	A9, A23, A24, A29, A80	*11:166, 23:01:01-23:56, 23:58-23:65, 23:67-23:69, 24:24, 24:71, 29:07, 29:49, 31:29, 31:85, 32:72, 80:01:01:01-80:03, B*18:27
7	175 bp, 205 bp	1070 bp	A2, A3, A9, A23, A24, A2403, A26	*02:17:01 ^w -02:17:03 ^w , 11:139, 23:14:01-23:14:02, 24:02:01:01-24:02:92, 24:02:95-24:11N, 24:13:01-24:13:02, 24:17-24:50, 24:54-24:56, 24:58-24:63, 24:66-24:91, 24:93, 24:95-24:113, 24:115-24:137, 24:139-24:187, 24:189-24:210, 24:212-24:221, 24:223-24:227, 24:229-24:290, 24:292-24:295, 24:297-24:303N, 24:305-24:314, 26:16, 33:19, 68:45, 68:117
8	165 bp, 200 bp	800 bp	A2, A3, A10, A11, A25, A26, A28, A32, A34, A66, A68, A69	*01:51, 02:55, 02:527, 03:24, 03:50, 11:10, 11:183, 11:191, 25:01:01-25:16, 25:18-25:32, 26:01:01:01-26:06, 26:08-26:15, 26:17-26:18, 26:20-26:43:02, 26:45-26:63, 26:65-26:71N, 26:73-26:88,

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				26:90-26:91, 26:93-26:116, 29:28, 32:15, 33:51, 34:01:01-34:12, 66:01:01-66:22, 68:01:01-68:134, 69:01-69:03
9^{6,10}	75 bp	800 bp	A3, A25, A32	*25:01:01-25:32, 32:01:01-32:02, 32:04, 32:06-32:37, 32:39-32:59, 32:61-32:70, 32:72-32:74, B*07:81, B*08:52, B*18:67, B*38:41, B*51:185, B*53:05, B*53:16, B*53:33
10^{6,7,11}	80 bp, 240 bp	800 bp	A10, A26, A43	*01:43, 01:51, 11:17, 11:40, 11:223, 26:01:01-26:02:02, 26:04, 26:07:01-26:20, 26:22-26:29, 26:31-26:43:02, 26:45-26:77, 26:79-26:91, 26:93-26:110, 26:112-26:116, 33:13, 33:48, 43:01, 68:84
11^{6,7,15}	80 bp, 175 bp, 500 bp	1070 bp	A1, A9, A10, A11, A26, A31, A34, A66	*01:13, 01:28, 01:136, 03:63, 03:88, 11:01:01-11:27, 11:29-11:52Q, 11:54-11:223, 24:19, 24:44, 26:03:01, 26:06, 26:21, 26:78, 26:92, 26:111, 34:01:01-34:08, 34:10N-34:12, 66:01:01-66:01:03, 66:04-66:11, 66:13-66:14, 66:17-66:20, 66:22-66:23, 69:02, 80:02, C*07:404
12⁶	125 bp, 195 bp	800 bp	A3, A10, A25, A26, A31, A34, A43, A66	*02:309, 02:454, 03:01:19, 11:11, 25:05-25:06, 26:09, 26:54, 26:91, 31:03-31:04, 34:01:01-34:12, 43:01, 66:02-66:03, 66:16, 66:21, 68:130
13	175 bp, 225 bp	1070 bp	A1, A2, A3, A10, A25, A26, A34, A43, A66	*01:01:56 ^w , 01:13, 02:34-02:35:03, 02:56:01-02:56:02, 02:62, 02:103, 02:135, 02:580, 03:01:01-03:01:22, 03:01:24-03:07, 03:09-03:11N, 03:13-03:31, 03:33-03:35, 03:37-03:40, 03:42-03:56, 03:58, 03:60-03:71, 03:73-03:87, 03:90-03:106, 03:109-03:110, 03:112-03:141, 03:143-03:151, 03:153-03:171, 03:174-03:175, 03:177, 03:179-03:197N, 03:200Q-03:202, 03:204, 03:206-03:210, 03:212-03:218, 03:221-03:229, 11:199:01-11:199:02, 11:222, 25:01:01-25:01:08, 25:02-25:05, 25:07-25:32, 26:01:01-26:01:20, 26:01:22-26:01:38, 26:02:01 ^w -26:02:02 ^w , 26:03:01, 26:05-26:08, 26:10-26:28, 26:29 ^w , 26:30-26:33, 26:35-26:43:02, 26:45-26:48, 26:49 ^w , 26:50-26:72, 26:74-26:77, 26:79-26:90, 26:92-26:116, 30:55, 34:08, 43:01, 66:01:01, 66:01:03, 66:04-66:09, 66:10 ^w , 66:11-66:15, 66:17-66:20, 66:22-66:23, 68:71, 74:13
14^{6,7,11}	80 bp, 115 bp, 200 bp, 240 bp, 470 bp	1070 bp	A26, A29, A31, A34	*02:237, 02:309, 02:454, 03:95, 26:19, 26:22, 29:01:01-29:80, 31:03-31:04, 32:42, 34:04, 66:09
15^{6,7,13,14}	90 bp, 135 bp, 205 bp	1070 bp	A1, A30	*01:13, 01:28, 03:43, 03:82, 03:186, 11:113, 11:162, 30:01:01-30:04:02, 30:06-30:20, 30:22-30:94, 31:35
16	240 bp, 380 bp, 410 bp	1070 bp	A24, A31, A32	*02:237, 03:95, 29:14, 31:01:02-31:94, 31:96-31:98, 32:05, 33:53, 74:13
17	140 bp, 180 bp	1070 bp	A32	*01:95, 03:43, 03:82, 03:186, 24:243, 29:13, 29:39, 30:89, 31:35, 32:01:01-32:01:17, 32:01:19-32:03, 32:05-32:71, 32:73-32:74, 74:07
18	200 bp, 390 bp	1070 bp	A33, A68	*02:243:01-02:243:02, 29:48, 32:15, 33:01:01-33:01:07, 33:03:01-33:37, 33:39-33:52,

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				33:54-33:102, 68:29, 74:04, 74:21
19	340 bp, 375 bp	800 bp	A2, A19, A68, A74	*01:121, 02:65, 02:407, 02:449, 32:62, 68:25, 74:01:01-74:24
20¹¹	210 bp, 240 bp	800 bp	A2, A210, A25, A68	*02:34-02:35:03, 02:46, 02:48, 02:56:01-02:56:02, 02:62, 02:70, 02:78, 02:103, 02:129, 02:580, 11:199:02, 23:01:13, 25:05, 26:54, 34:02:04, 68:01:01:01-68:134
21¹⁶	200 bp, 240 bp, 375 bp	800 bp	A2, A23, A24, A26, A28, A66, A68, A69	*02:17:01-02:17:03, 02:55, 02:108, 02:110, 02:243:01-02:243:02, 02:268, 02:300, 02:303, 23:02, 23:57, 23:66, 24:06, 24:13:01, 24:22, 24:82, 24:94, 24:138, 24:167, 24:188, 24:207, 24:228, 25:30, 26:22, 33:22, 66:06, 66:09, 68:08:02, 68:29, 68:105, 69:01-69:03
22^{6,12}	85 bp, 240 bp, 400 bp	800 bp	A2, A23, A34 A36, A68	*02:34-02:35:03, 02:46, 02:48, 02:56:01-02:56:02, 02:62, 02:70, 02:78, 02:103, 02:129, 02:580, 03:187, 11:155, 11:199:02, 23:01:13, 31:62, 34:02:04, 36:01-36:05, 68:01:32, 68:11N
23^{6,11}	85 bp, 160 bp, 240 bp, 495 bp	800 bp	A2, A3, A24, A26, A28, A36, A68, A80	*02:55, 02:237, 03:41, 03:63, 03:75, 03:88, 03:95, 03:177, 11:130, 24:18, 24:204, 24:213, 26:03:01, 26:05-26:06, 26:21, 26:30, 26:78, 26:111, 33:24, 36:02, 68:05, 68:15, 68:20, 68:109, 80:01:01:01- 80:01:01:02, 80:03
24¹³	360 bp	1070 bp	A10, A26, A31, A66	*02:135, 02:309, 02:454, 03:01:19, 25:13, 26:30, 26:65, 31:04, 34:09, 66:02-66:03, 66:12, 66:16, 66:21

¹Alleles are assigned by the presence of specific PCR product(s). However, the sizes of the specific PCR products may be helpful in the interpretation of HLA-A low resolution SSP typings.

When the primers in a primer mix can give rise to HLA-specific PCR products of more than one length this is indicated if the size difference is more than 20 base pairs. Size differences of 20 base pairs or less are not given. For high resolution SSP kits, the alleles listed are specified according to amplicon length.

Nonspecific amplifications, i.e. a ladder or a smear of bands, may sometimes be seen. GC-rich primers have a higher tendency of giving rise to nonspecific amplifications than other primers.

PCR fragments longer than the control bands may sometimes be observed. Such bands should be disregarded and do not influence the interpretation of the SSP typings.

PCR fragments migrating faster than the control bands, but slower than a 400 bp fragment may be seen in some gel read-outs. Such bands can be disregarded and do not influence the interpretation of the SSP typings.

Some primers may give rise to primer oligomer artifacts. Sometimes this phenomenon is an inherent feature of the primer pair(s) of a primer mix. More often it is due to other factors such as too low amount of DNA in the PCR reactions, taking too long time in setting up the PCR reactions, working at elevated room temperature or using thermal cyclers that are not pre-heated.

²The internal positive control primer pairs amplify segments of the human growth hormone gene. The internal positive control bands are 1070 or 800 base pairs respectively, well distribution as outlined in the table. Well number 1 contains the shorter, 800 bp, internal positive control band. The well distribution of the internal controls can help in orientation of the kit on gel photo, as well as allow for kit identification. In the presence of a specific amplification the intensity of the control band often decreases.

³The serological reactivity of all HLA-A alleles is not known. In this table we use the expert-assigned serological grouping in Tissue Antigens (2009) 73:95-170 and the serological grouping of the sequence-defined allele.

⁴For several HLA Class I alleles 1st and/or 4th exon(s) and beyond, as well as intron nucleotide sequences, are not available. In these instances it is not known whether some of the primers of the SSP sets are completely matched with the target sequences or not. Assumption is made that unknown sequences in these regions are conserved within allelic groups.

⁵Only HLA-A alleles will be amplified by the 24 wells of the HLA-A low resolution primer set, except that a

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few HLA-B and HLA-C alleles will be amplified by primer mixes 4 to 6, 9 and 11.

⁶HLA-specific PCR products shorter than 125 base pairs have a lower intensity and are less sharp than longer PCR products.

⁷The primer pairs in wells 1, 2, 5, 7, 8, 10, 11, 14, 15 and 18 will in many samples give rise to two or three HLA-specific PCR fragments.

⁸Primer mix 3 may faintly amplify the A*30:04:01-30:04:02, 30:06, 30:17, 30:29, 30:46 and 30:77 alleles.

⁹Primer mix 6 may give rise to a lower yield of A*23 alleles than the other A low primer mixes.

¹⁰Primer mix 9 may weakly amplify the A*34 alleles.

¹¹Primer mixes 6, 10, 14, 20 and 23 have a tendency to giving rise to primer oligomer formation.

¹²Primer mix 22 might faintly amplify most A*11 alleles.

¹³Primer mixes 15 and 24 may give rise to a lower yield of HLA-specific PCR product than the other HLA-A low primer mixes.

¹⁴Primer mix 15 may have tendency of unspecific amplification.

¹⁵In primer mix 11 the positive control band may be weaker than for other HLA-A low primer mixes.

¹⁶Primer mix 21 may generate a false positive band of about 560 base pairs. This band should be disregarded when interpreting HLA-A low resolution typings.

‘w’, might be weakly amplified.

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SPECIFICITY TABLE

HLA-B low resolution primer set

Specificities and sizes of the PCR products of the 48 primer mixes used for
HLA-B low resolution SSP typing

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	HLA-B serol. ³	Amplified HLA-B alleles ^{4,5}
25^{6,10}	110 bp	800 bp	7, 703, 40, 41, 42, 48, 61	*07:02:01-07:18:02, 07:20-07:32, 07:34-07:39, 07:41-07:47, 07:49N-07:50, 07:52, 07:54-07:59, 07:61-07:99, 07:101-07:121, 07:123-07:138, 07:140-07:154, 07:156-07:163, 07:166-07:201N, 07:203-07:208, 07:210-07:219, 07:221-07:222, 07:224-07:225, 07:227-07:234, 07:236-07:249, 07:251N-07:256, 08:79, 15:138, 15:230, 15:241, 15:297, 37:07, 40:15-40:16, 40:23, 40:32, 40:98, 40:136, 40:158, 40:183, 40:198, 40:231, 40:279, 41:08, 42:05:01-42:05:02, 44:150, 48:05, 48:08, 48:15, C*02:23, C*03:260, C*04:77
26	215 bp	1070 bp	8, 44	*08:01:01-08:05, 08:07-08:25, 08:27-08:49, 08:51-08:61, 08:63-08:64, 08:66-08:84, 08:86N-08:93, 08:95-08:122, 08:124-08:131, 08:133-08:145, 08:147-08:151, 15:142, 15:180, 35:218, 35:256, 44:49, 44:156, 49:25, 51:68, 51:176, C*06:147
27¹⁰	140 bp, 235 bp	1070 bp	7, 8, 13, 15, 35, 4005, 44, 49, 50, 61, 62, 77	*07:20, 07:24, 07:60, 07:100, 07:131, 07:220, 07:223, 08:21, 08:25, 13:01:01-13:04, 13:06-13:08, 13:10-13:23, 13:25-13:38, 13:40-13:57, 13:59-13:66, 13:68-13:84, 15:07:01-15:07:03, 15:36 ^w , 15:45, 15:55, 15:68, 15:89 ^w , 15:126, 15:207, 15:324, 15:329, 15:331, 15:360, 35:05:01-35:05:03, 35:16-35:17:02, 35:22, 35:30, 35:51, 35:58, 35:72, 35:89, 35:97, 35:113-35:114, 35:125, 35:164, 35:199, 35:232, 35:242, 40:05, 40:71, 40:174, 44:08 ^w , 44:54, 44:57 ^w , 44:60, 44:106, 44:110, 44:135, 44:158, 44:184, 44:211, 44:213, 46:12, 51:64, 51:148, 53:14, 55:51, 58:18, C*03:05, C*03:25, C*03:27, C*03:143, C*03:167, C*03:296
28⁷	130 bp, 265 bp	800 bp	12, 13, 14, 15, 17, 21, 35, 40, 41, 44, 45, 47, 49, 50, 57, 58, 60, 61, 64, 65	*13:01:01-13:01:09, 13:02:01-13:04, 13:06-13:13:02, 13:15-13:23, 13:25-13:57, 13:59-13:82, 13:84, 14:01:01-14:02:08, 14:02:10-14:04, 14:07N, 14:09, 14:11-14:12, 14:14-14:36, 14:38-14:49, 15:46, 15:53, 15:106, 15:143, 15:212, 15:336, 18:44:01-18:44:02, 18:107, 35:19, 35:46-35:47, 35:63, 35:154, 35:207, 35:217, 39:79, 40:01:01-40:01:39, 40:01:42-40:01:43, 40:07, 40:10:01:01-40:10:02, 40:14:01-40:16, 40:22N-40:23, 40:25, 40:30-40:34, 40:36, 40:38, 40:42-40:43, 40:45, 40:47-40:49, 40:51-40:55, 40:58-40:63, 40:65-40:67, 40:69, 40:72:01-40:73, 40:76-40:77, 40:79-40:81, 40:84, 40:87:01-40:88, 40:92, 40:100-40:102, 40:106, 40:108, 40:110, 40:112-40:114:02, 40:116-40:118N, 40:121, 40:123-40:126, 40:128-40:130:02, 40:132, 40:134-40:136, 40:139-40:141, 40:146-40:147, 40:150-40:156, 40:158, 40:160:01-40:160:02, 40:163-40:164, 40:166, 40:168, 40:170-40:172, 40:175, 40:178-40:179, 40:182-40:183, 40:186-40:188, 40:191-40:199, 40:204, 40:207-40:208, 40:212-40:213, 40:215-40:219, 40:221-40:223, 40:227-40:228, 40:231, 40:233-40:242, 40:245, 40:247, 40:249-40:250, 40:252-40:253, 40:257-40:264, 40:272-40:273, 40:277-40:282, 40:285-40:286N, 40:288, 40:298-40:299, 40:301, 40:308, 41:01:01-41:39, 44:02:01:01-44:02:31, 44:02:33-44:03:06,

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29⁸	185 bp, 235 bp	800 bp	7, 16, 17, 27, 2708, 37, 38, 39, 3901, 3902, 64, 67	*07:02:32, 07:197, 14:01:01-14:01:05, 14:07N-14:08:02, 14:10, 14:12, 14:14, 14:19, 14:26, 14:32, 14:40, 14:46-14:47, 14:49, 27:01-27:05:15, 27:05:17-27:05:19, 27:05:21-27:06, 27:07:04-27:10, 27:12-27:13, 27:16-27:18, 27:20, 27:23, 27:26-27:27, 27:29, 27:31, 27:35-27:37, 27:39-27:42, 27:44-27:46, 27:48-27:61, 27:64N-27:69, 27:72-27:75, 27:77-27:80, 27:82-27:124, 27:126, 27:128-27:129, 27:131-27:137, 27:140-27:142, 37:02, 38:01:01-38:01:07, 38:01:09-38:02:02, 38:02:04-38:03, 38:07-38:24, 38:26-38:32, 38:34N-38:57, 39:01:01-39:01:01:03, 39:01:03-39:01:08, 39:01:10-39:02:01, 39:03, 39:05:01-39:06:02, 39:06:04-39:09:03, 39:11, 39:14-39:15, 39:18, 39:19:02, 39:22, 39:24:01-39:39:01, 39:40:01N-39:44, 39:46-39:48, 39:50-39:62, 39:64-39:71, 39:73-39:75, 39:77, 39:79-39:82, 39:84-39:88, 39:90-39:91, 39:93-39:95N, 39:97N-39:98, 39:100, 39:102, 39:104, 47:04-47:05, 48:21, 48:26, 67:01:01, 67:03-67:04
30	190 bp	800 bp	14, 35, 38, 39, 65	*07:28, 08:87, 14:02:01:01-14:02:02, 14:02:04-14:02:05, 14:02:07-14:06:02, 14:09, 14:11, 14:13, 14:15-14:18, 14:20, 14:22-14:23, 14:25, 14:27, 14:29-14:31, 14:33-14:39, 14:41N, 14:43-14:45, 14:48, 15:77, 15:189, 15:233, 35:03:17, 35:26, 38:05, 38:33, 39:04, 44:16, 44:37:01-44:37:02, 44:64:01-44:64:02, 44:91, 44:132, 44:150, 44:190, 52:26, 57:04:01, A*23:31, A*24:106, C*02:97, C*07:231, C*16:10
31	290 bp	1070 bp	15, 22, 46, 62, 63, 71, 72, 75, 76, 77	*15:01:01:01-15:01:03, 15:01:06-15:02:09, 15:03:03, 15:04:01-15:08:02, 15:10:02, 15:11:01-15:11:03, 15:11:05-15:15, 15:17:01:01-15:17:03, 15:19-15:21, 15:24:01-15:28, 15:30-15:36, 15:38:01-15:40, 15:42-15:46, 15:48, 15:50, 15:55-15:58, 15:60, 15:63, 15:65-15:66, 15:70-15:71, 15:73, 15:75-15:79N, 15:81-15:89, 15:92, 15:94N, 15:96-15:97, 15:101-15:102, 15:104-15:107, 15:109-15:113, 15:116-15:118, 15:120-15:122, 15:125-15:126, 15:128-15:129, 15:135-15:150, 15:152, 15:154-15:155, 15:157, 15:159-15:160, 15:162-15:172, 15:174-15:175, 15:177-15:179:02, 15:181N-15:185, 15:187-15:196, 15:199, 15:201-15:209N, 15:211, 15:213-15:219, 15:223-15:225, 15:227-15:228, 15:230-15:234, 15:236-15:237, 15:239-15:241, 15:244-15:251, 15:256-15:262N, 15:264-15:265, 15:267-15:273, 15:276-15:280, 15:283-15:289, 15:291, 15:295-15:305, 15:308-15:310, 15:315-15:322, 15:324-15:328, 15:330-15:334, 15:339-15:346, 15:353, 15:355-15:361, 15:363, 46:01:01-46:64, C*01:73, C*05:01:28, C*07:02:30, C*08:16:02
32	165 bp, 225 bp, 285 bp, 330 bp	1070 bp	5, 8, 12, 21, 22, 37, 41, 42, 44, 45, 48, 51, 56, 57, 60, 62, 70, 71,	*08:01:01-08:01:22, 08:01:24-08:05, 08:08N-08:13, 08:15-08:27, 08:29-08:36, 08:38-08:48, 08:50-08:73, 08:75-08:76, 08:78, 08:80-08:86N, 08:88, 08:90-08:106, 08:109-08:110, 08:112-08:151, 13:46, 15:03:01:01-15:03:05, 15:09-15:10:04, 15:18:01-15:18:06, 15:23, 15:29, 15:37, 15:46-15:47:02, 15:49, 15:51-15:54, 15:61-15:62, 15:64:01-15:64:02, 15:69, 15:72,

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34	180 bp	800 bp	13, 22, 55, 60	*07:78, 13:01:01-13:02:18, 13:07N-13:09, 13:11, 13:14-13:20, 13:22:01-13:23, 13:25, 13:27-13:34, 13:36-13:45, 13:47, 13:49N-13:50, 13:52, 13:54-13:58, 13:60-13:61, 13:63N-13:70, 13:73-13:78, 13:80-13:84, 40:48, 45:10, 49:07, 50:31, 54:26, 55:09, 55:22, 55:24, 59:09, C*02:06, C*15:42
35⁶	100 bp, 195 bp	1070 bp	8, 12, 27, 38, 39, 3902, 44, 45, 48, 60, 70, 71, 72, 82	*07:27, 07:50, 07:219, 08:04, 08:17, 08:54, 08:110, 15:03:01:01-15:03:05, 15:47:01-15:47:02, 15:49, 15:54, 15:61-15:62, 15:64:01-15:64:02, 15:68-15:69, 15:91, 15:98, 15:103, 15:123, 15:127, 15:131-15:132, 15:151, 15:156, 15:158, 15:173, 15:210, 15:220, 15:235, 15:242, 15:253, 15:259, 15:266, 15:274, 15:281-15:282, 18:37, 27:18, 27:29, 37:28, 38:03, 39:02:01-39:02:02, 39:08, 39:13:01-39:13:02, 39:23, 39:39:01-39:39:02, 39:49, 39:88, 39:101, 39:105, 40:12, 40:46, 40:93, 42:11, 44:10, 44:15, 44:18, 44:40, 44:44, 44:130, 44:140, 44:156, 44:210, 45:01:01-45:01:02, 45:05-45:07, 45:11-45:16, 48:01:01-48:05, 48:07-48:30, 48:32-48:37, 49:20, 50:02, 52:16, 55:18, 56:44, 58:44, 82:01-82:03
36	280 bp	1070 bp	5, 7, 8, 13, 15, 16, 17, 18, 22, 35, 40, 42, 44, 46, 48, 49, 51, 53, 55, 56, 57, 58, 60, 61, 62, 63, 70, 71,	*07:09, 07:11, 07:17, 07:162, 07:235, 07:237, 07:255, 08:28, 08:35, 08:37, 08:89, 08:107, 13:04, 13:10, 13:26:01-13:26:02, 13:72, 15:01:01:01-15:01:04, 15:01:06-15:08:02, 15:11:01-15:16:03, 15:18:01-15:21, 15:23-15:29, 15:31-15:36, 15:38:01-15:40, 15:43-15:44, 15:46-15:47:02, 15:49-15:57, 15:60-15:62, 15:64:01-15:72, 15:74-15:76, 15:78:01-15:82, 15:84-15:85, 15:87-15:89, 15:91-15:98, 15:101-15:129, 15:131-15:132, 15:134-15:136, 15:138-15:149N, 15:151-15:161, 15:163-15:167, 15:169-15:176, 15:178-15:187, 15:189-15:195,

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68⁶	95 bp	1070 bp	8, 18, 22, 35, 37, 38, 39, 44, 51, 5102, 5103, 53, 78	*07:65 ^w , 07:134 ^w , 08:32, 08:133, 15:308, 18:01:01-18:11, 18:13-18:15, 18:17N-18:36, 18:38-18:47, 18:49-18:65, 18:67-18:106, 18:108-18:113, 35:01:01-35:08:04, 35:08:06-35:09:03, 35:11:01-35:12:03, 35:14:01-35:15:02, 35:17:01-35:18, 35:20:01-35:24:02, 35:27, 35:29:01-35:45, 35:48, 35:50-35:62, 35:64:01-35:68:02, 35:70-35:72, 35:74-35:75, 35:76 ^w , 35:77-35:79, 35:81-35:153, 35:155-35:185, 35:187-35:190, 35:192-35:197, 35:199-35:206, 35:208-35:209, 35:211-35:216N, 35:218-35:225, 35:227-35:253, 35:255-35:269, 35:272-35:273, 35:275-35:282, 35:285-35:296, 37:08, 38:06-38:07, 39:19:01-39:19:02, 44:06, 51:01:01-51:24:05, 51:26-51:46, 51:48-51:103, 51:105-51:111, 51:113-51:117, 51:119-51:146, 51:148-51:190, 53:01:01-53:16, 53:18-53:27, 53:29, 53:31-53:37, 56:06 ^w , 56:45 ^w , 78:01:01-78:04, 78:07-78:08
69^{6,10}	115 bp, 150 bp	1070 bp	15, 18, 22, 27, 35, 39, 46, 55, 62, 72, 73, 75, 76	*07:100, 13:31, 13:41, 15:01:01-15:01:04, 15:01:06-15:01:16, 15:01:18-15:01:30, 15:01:32-15:01:39, 15:04:01-15:04:02, 15:07:01-15:08:02, 15:11:01-15:12, 15:14-15:15, 15:19, 15:24:01-15:24:02, 15:26N-15:28, 15:30, 15:32:01-15:32:02, 15:34-15:35, 15:38:01-15:38:02, 15:43, 15:45-15:46, 15:50, 15:53-15:54, 15:56-15:58, 15:60, 15:63, 15:66, 15:68, 15:70-15:71, 15:73, 15:75-15:77, 15:79N, 15:81-15:82, 15:85, 15:87, 15:92, 15:94N, 15:96-15:97, 15:101-15:102, 15:104-15:105, 15:109-15:111N, 15:113, 15:117-15:118, 15:120, 15:122, 15:125-15:126, 15:128-15:129, 15:135, 15:137, 15:140, 15:142-15:149N, 15:152, 15:154, 15:157, 15:159-15:160, 15:163-15:167, 15:169, 15:171-15:172, 15:174-15:175, 15:178, 15:180-15:184, 15:187, 15:189-15:193, 15:201-15:203, 15:205-15:207, 15:209N, 15:211-15:212, 15:215, 15:217, 15:225, 15:227-15:228, 15:231-15:234, 15:236, 15:239, 15:241, 15:244-15:247, 15:251, 15:256-15:262N, 15:264, 15:267, 15:269-15:270, 15:272N, 15:276-15:280, 15:284-15:286, 15:296, 15:298-15:299, 15:303-15:305, 15:309, 15:315-15:318, 15:320-15:321Q, 15:324, 15:326-15:327, 15:331, 15:333-15:334, 15:336, 15:340, 15:342-15:344, 15:346-15:347, 15:353, 15:355, 15:359, 15:363, 18:19, 27:25, 27:75, 27:115, 35:14:01-35:14:02, 35:43:01-35:44, 35:62, 35:67, 35:79, 35:86, 35:102, 35:117, 35:135, 35:163, 35:185, 35:213, 35:265, 39:18, 39:36, 44:146, 46:01:01-46:02, 46:04-46:05, 46:07N, 46:09-46:10, 46:12, 46:14-46:17, 46:20, 46:22-46:24, 46:26-46:32, 46:34-46:42, 46:44, 46:46-46:53, 46:55N-46:57, 46:60-46:61, 46:63-46:64, 51:61:01-51:61:02, 52:21, 54:06, 54:33, 55:21, 56:03, 56:43, 73:01-73:02, A*26:68, A*26:100, A*68:56, C*02:56, C*06:20, C*12:50
70¹⁵	360 bp	1070 bp	Bw4	
71	350 bp	1070 bp	Bw6	
72	285 bp	800 bp	7, 703, 8, 35, 40, 41, 42, 48, 53, 60, 61, 81	*07:02:01-07:02:38, 07:02:40-07:06, 07:08-07:18:02, 07:20-07:32, 07:34-07:39, 07:41-07:52, 07:54-07:59, 07:61-07:76, 07:79-07:83, 07:85-07:99, 07:101-07:121, 07:123-07:138, 07:140-07:164, 07:166-07:192, 07:194-07:195, 07:199-07:213,

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08:01:24, 08:01:26-08:05, 08:07-08:08N, 08:10-08:11, 08:13-
08:15, 08:17-08:30N, 08:32-08:83, 08:85-08:88, 08:90-08:102,
08:104-08:106, 08:108-08:112, 08:114-08:115, 08:117-08:133,
08:135-08:151, 15:297, 35:66, 35:87, 37:07, 40:15-40:16,
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42:15-42:21, 44:150, 44:166, 48:01:01-48:01:06, 48:05-48:12,
48:14-48:20, 48:22, 48:27-48:37, 53:15, 81:01-81:07, **C*03:260**

¹Alleles are assigned by the presence of specific PCR product(s). However, the sizes of the specific PCR products may be helpful in the interpretation of HLA-B low resolution SSP typings. When the primers in a primer mix can give rise to HLA-specific PCR products of more than one length this is indicated if the size difference is more than 20 base pairs. Size differences of 20 base pairs or less are not given. For high resolution SSP kits, the alleles listed are specified according to amplicon length. Nonspecific amplifications, i.e. a ladder or a smear of bands, may sometimes be seen. GC-rich primers have a higher tendency of giving rise to nonspecific amplifications than other primers. PCR fragments longer than the control bands may sometimes be observed. Such bands should be disregarded and do not influence the interpretation of the SSP typings. PCR fragments migrating faster than the control bands, but slower than a 400 bp fragment may be seen in some gel read-outs. Such bands can be disregarded and do not influence the interpretation of the SSP typings. Some primers may give rise to primer oligomer artifacts. Sometimes this phenomenon is an inherent feature of the primer pair(s) of a primer mix. More often it is due to other factors such as too low amount of DNA in the PCR reactions, taking too long time in setting up the PCR reactions, working at elevated room temperature or using thermal cyclers that are not pre-heated.

²The internal positive control primer pairs amplify segments of the human growth hormone gene. The internal positive control bands are 1070 or 800 base pairs respectively, well distribution as outlined in the table. Well number 25 contains the shorter, 800 bp, internal positive control band. The well distribution of the internal controls can help in orientation of the kit on gel photo, as well as allow for kit identification. In the presence of a specific amplification the intensity of the control band often decreases.

³The serological reactivity of all HLA-B alleles is not known. In this table we use the expert-assigned serological grouping in Tissue Antigens (2009) 73:95-170 and the serological grouping of the sequence-defined allele.

⁴For several HLA Class I alleles 1st and/or 4th exon(s) and beyond, as well as intron nucleotide sequences, are not available. In these instances it is not known whether some of the primers of the SSP sets are completely matched with the target sequences or not. Assumption is made that unknown sequences in these regions are conserved within allelic groups.

⁵Only HLA-B alleles will be amplified by the 48 wells of the HLA-B low resolution, primer set, except that a few HLA-A and HLA-C alleles will be amplified by primer mixes 25 to 28, 30 to 32, 37, 38, 41, 43 to 46, 49, 50, 53 to 55, 64 to 67, 69 and 72.

⁶HLA-specific PCR products shorter than 125 base pairs have a lower intensity and are less sharp than longer PCR products.

⁷Primer mix 28 may give rise to a lower yield of HLA-specific PCR product than the other HLA-B low resolution primer mixes in B*40, B*41, B*45, B*49 and B*50 alleles.

⁸Primer mixes 29, 33 and 63 may give rise to a lower yield of HLA-specific PCR product than the other HLA-B low resolution primer mixes.

⁹Primer mixes 33, 52 and 62 may have tendencies of unspecific amplifications, most pronounced in primer mix 33.

¹⁰Primer mixes 25, 27, 38, 41, 43, 44, 50, 58, 67 and 69 have a tendency to giving rise to primer oligomer formation.

¹¹The B*57 and B*58 alleles might be faintly amplified by primer mix 33.

¹²The C*17:01 to C*17:04 alleles might be faintly amplified by primer mix 49.

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¹³Primer mixes 56 and 67 may generate a false positive band of about 800 base pairs. This band should be disregarded when interpreting HLA-B low resolution typings.

¹⁴Primer mix 63 may give rise to a lower yield of B*54 alleles than the other B low primer mixes.

¹⁵The Bw4-associated HLA-A specificities A9, A23, A24, A2403, A25 and A32 are not amplified by the primer pair in primer mix 70.

‘w’, might be weakly amplified.

‘?’, nucleotide sequence information not available for the primer matching sequence.

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SPECIFICITY TABLE

HLA-C low resolution SSP typing

Specificities and sizes of the PCR products of the 23+1 primer mixes used for
HLA-C low resolution SSP typing

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	Amplified HLA class I ^{3,4} alleles
73⁵	90 bp, 150 bp	800 bp	*01:02:01-01:107, 01:109N-01:111N, 04:71, 07:316, 07:338, 14:58, 15:104, B*39:76, B*54:18
74⁶	130 bp, 200 bp, 270 bp, 300 bp	800 bp	*01:10, 01:43, 02:02:01-02:02:03, 02:02:05-02:40:02, 02:42-02:57, 02:59-02:106, 04:32, 04:77, 05:105, 06:08, 07:101, 07:148, 07:161, 08:31, 12:119, 14:25, 15:42, 15:111, 16:29, 16:50, 17:01:01-01:17:06, 17:08-17:16, 17:18-17:22, 17:24-17:30, 18:03
75¹³	280 bp	800 bp	*02:02:01-02:02:03, 02:02:05-02:02:29, 02:03, 02:04 ^w , 02:05:01-02:13, 02:14:01 ^w , 02:15-02:25Q, 02:26:02-02:40:02, 02:42-02:86, 02:88-02:106, 03:02:01-03:02:09, 03:02:11-03:02:14, 03:04:01-01-03:04:25, 03:04:27-03:10, 03:14-03:17, 03:19, 03:23-03:29, 03:32-03:38:02, 03:40:01-03:42, 03:44-03:48, 03:51, 03:54, 03:57, 03:60, 03:63-03:64:01, 03:65, 03:70-03:74, 03:77-03:78, 03:80, 03:82, 03:84, 03:87:01-03:87:02, 03:89-03:95, 03:98, 03:100-03:101, 03:104-03:111, 03:113-03:115, 03:117-03:118, 03:121N, 03:123, 03:125, 03:128-03:131, 03:134-03:136, 03:138-03:140, 03:143, 03:145-03:149, 03:153-03:155, 03:157, 03:159, 03:162-03:164, 03:169Q, 03:172-03:174, 03:178-03:181, 03:183-03:184:02, 03:186:01-03:186:02, 03:190-03:191, 03:193-03:194, 03:197-03:201N, 03:208N-03:213, 03:215-03:216, 03:218-03:219, 03:221-03:222, 03:224N-03:226, 03:232-03:236, 03:238-03:240, 03:244Q-03:250, 03:252, 03:255-03:261, 03:263, 03:265N-03:266, 03:269-03:270, 03:277N, 03:278 ^w , 03:279-03:280, 03:282-03:283, 03:286-03:287, 03:292-03:294, 03:296-03:297, 04:03:01-04:03:02, 04:06, 04:16, 04:80, 04:107, 04:147, 04:160, 04:171, 05:58:01, 06:03:01, 06:132:01-06:132:02, 07:96:01-07:96:02, 07:314:02, 12:03:23, 15:02:01-01-15:02:20, 15:02:22-15:05:08, 15:05:10-15:09, 15:10:02-15:11, 15:13, 15:15-15:19, 15:21-15:22, 15:24-15:35, 15:37-15:60, 15:62, 15:64-15:106, 15:108-15:117, 16:34, 16:70
76⁷	170 bp, 275 bp	1070 bp	*03:02:01-03:02:12, 03:02:14-03:04:06, 03:04:08-03:11:02, 03:13:01-03:15, 03:17-03:40:04, 03:42-03:57, 03:59-03:79, 03:81-03:85, 03:87:01-03:93, 03:95-03:98, 03:100-03:109, 03:111-03:112, 03:114-03:150, 03:152-03:164, 03:166-03:250, 03:252-03:259, 03:261-03:266, 03:268-03:280, 03:282-03:285, 03:287-03:292, 03:294-03:297, 07:242
77	280 bp	800 bp	*03:03:01-03:03:20, 03:03:22-03:03:29, 03:11:01-03:11:02, 03:13:01-03:13:02, 03:18:02, 03:20N-03:22Q, 03:30-03:31, 03:43:01-03:43:02, 03:49-03:50, 03:52-03:53, 03:55-03:56, 03:58-03:59, 03:61-03:62, 03:66, 03:67 ^w , 03:68-03:69, 03:75-03:76, 03:79, 03:81, 03:83, 03:85-03:86, 03:88, 03:96-03:97, 03:102-03:103, 03:112, 03:116:01-03:116:02, 03:119-03:120, 03:122, 03:124, 03:126-03:127, 03:132-03:133, 03:141-03:142, 03:150-03:152, 03:158, 03:160-03:161, 03:165, 03:167-03:168, 03:171, 03:175-03:177, 03:185, 03:187-03:189N, 03:192, 03:196, 03:202-03:207, 03:214, 03:217, 03:220, 03:223, 03:227-03:230,

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78¹³	135 bp, 330 bp	800 bp	*02:02:01-02:02:03, 02:02:05-02:02:12, 02:02:14-02:02:25, 02:02:27-02:02:28, 02:02:30-02:20, 02:22-02:25Q, 02:27:01-02:38N, 02:40:01-02:40:02, 02:42-02:44, 02:46-02:86, 02:88-02:100, 02:101 ^w , 02:102-02:106, 03:231, 04:01:01:01-04:01:27, 04:01:29-04:01:60, 04:01:62, 04:01:64, 04:01:66-04:01:68, 04:03:01-04:15:03, 04:17-04:20, 04:23-04:41, 04:43-04:102, 04:104-04:139, 04:141-04:195, 04:197-04:206, 05:26, 05:78, 07:02:09, 07:125, 07:356, 15:11, 15:36, 16:34
79	165 bp, 390 bp, 445 bp	1070 bp	*02:94, 04:129, 05:01:01:01-05:01:31, 05:03-05:122, 06:05, 06:67, 08:10, 12:21, 12:33, 15:107, 16:40, 16:53, 17:05
80	130 bp, 355 bp	800 bp	*01:90, 01:101-01:102, 02:06, 02:23, 02:36, 02:47, 04:108 ^w , 04:178, 06:02:01:01-06:02:01:03, 06:02:03-06:02:38, 06:02:40-06:16N, 06:18-06:31, 06:33-06:76:01, 06:77-06:93, 06:96-06:98, 06:100-06:132:01, 06:133-06:164, 08:113, 12:02:11, 12:03:09, 12:03:26, 12:15, 12:113, 15:02:01:01-15:03, 15:05:01 ^w -15:05:11 ^w , 15:06:01-15:13, 15:15-15:19, 15:21, 15:22 ^w -15:23 ^w , 15:26-15:28, 15:29 ^w , 15:31-15:35, 15:36 ^w , 15:37-15:39, 15:41-15:45, 15:46 ^w , 15:47-15:53, 15:54 ^w , 15:55-15:58, 15:59 ^w , 15:60-15:64, 15:67-15:68, 15:69 ^w -15:70 ^w , 15:71-15:75, 15:78-15:89, 15:90 ^w , 15:91-15:101, 15:103, 15:104 ^w -15:105Q ^w , 15:106-15:107, 15:108 ^w , 15:109, 15:110 ^w -15:111 ^w , 15:112-15:114, 15:115N ^w -15:117 ^w , 16:01:16, 16:20, B*58:02
81⁸	245 bp, 425 bp	800 bp	*07:01:01:01-07:33N, 07:35-07:294, 07:296-07:446
82^{5,9,13}	110 bp, 165 bp, 390 bp	800 bp	*01:43, 02:87, 03:280, 07:101, 07:148, 07:161, 08:01:01-08:63, 08:65-08:117, 08:119-08:123, 12:127
83	340 bp	1070 bp	*01:14, 01:59, 02:02:01-02:02:03, 02:02:05-02:11, 02:13-02:26:03, 02:28-02:40:02, 02:42-02:86, 02:88-02:106, 03:07, 03:15, 03:45, 03:130, 03:140, 03:163, 03:243, 03:268, 03:297, 04:01:01:01-04:01:68, 04:03:01-04:10, 04:12-04:20, 04:23-04:28, 04:30-04:35, 04:37-04:54, 04:56-04:171, 04:173N-04:206, 05:01:01:01-05:01:31, 05:03-05:122, 06:02:01:01-06:02:01:03, 06:02:03-06:02:11, 06:02:13-06:10, 06:12-06:51, 06:53:01-06:121, 06:123, 06:125-06:146, 06:148-06:164, 07:07, 07:09, 07:49, 07:76:01-07:76:02, 07:210, 07:238, 07:247, 07:315, 07:328, 07:403, 07:406, 08:10, 12:04:01-12:05, 12:09, 12:21, 12:33, 12:41, 12:54, 12:60, 12:146, 14:04, 14:12, 14:49, 14:64, 15:02:01:01-15:05:10, 15:06:01-15:06:03, 15:08-15:13, 15:15-15:19, 15:22-15:24, 15:26-15:42, 15:44-15:70, 15:72-15:115N, 15:117, 16:02:01-16:02:13, 16:09, 16:12, 16:19, 16:25, 16:46-16:48, 16:57, 16:60, 16:63, 16:69-16:70, 16:74, 16:77N, 16:84, 17:01:01:01-17:21, 17:23-17:30, 18:01-18:09
84^{5,11}	100 bp, 155 bp, 220 bp	800 bp	*01:02:18, 01:04, 01:21, 06:02:38, 06:76:02, 12:02:01-12:03:01:02, 12:03:03-12:03:07, 12:03:09-12:03:15, 12:03:17-12:03:19, 12:03:21-12:03:29, 12:03:31-12:03:33, 12:03:35-12:03:37, 12:04:02-12:08, 12:10:01-12:13, 12:14:02-12:18:01, 12:19-12:25, 12:27-12:32, 12:34-12:58, 12:60-12:143, 12:145-12:162, 14:02:08, 16:01:17, 16:02:13, 16:15:02
85^{5,11}	120 bp, 250 bp	800 bp	*01:21, 02:12, 02:49, 02:55:01, 04:01:01:01-04:01:68, 04:03:01-04:09N, 04:12-04:20, 04:23-04:35, 04:37-04:54, 04:56-04:152,

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			04:154-04:168, 04:170N-04:191N, 04:193, 04:195-04:206, 05:42, 05:46, 05:112, 06:76:02, 07:02:09, 07:125, 07:356, 08:05, 08:21, 08:25, 12:02:01-12:02:10, 12:02:12-12:03:03, 12:03:05-12:03:08, 12:03:10-12:03:12, 12:03:13 ^w , 12:03:14-12:03:23, 12:03:24 ^w , 12:03:25-12:03:33, 12:03:35-12:03:37, 12:04:02, 12:06-12:08, 12:10:01-12:20, 12:22-12:32, 12:34-12:48, 12:50-12:97, 12:99-12:122, 12:124-12:162, 15:03, 15:16, 15:25, 15:75, 16:01:01-16:02:13, 16:06-16:28, 16:30N-16:32, 16:34, 16:36-16:39:02, 16:41, 16:43-16:47, 16:49-16:52, 16:54, 16:56-16:60, 16:62-16:65, 16:67, 16:69-16:77N, 16:79-16:81, 16:83-16:86, 17:01:04, B*67:02
86¹⁴	160 bp, 210 bp	800 bp	*01:04, 01:09, 02:05:01-02:05:03, 02:17, 04:42:01-04:42:02, 06:02:01:01-06:02:01:03, 06:02:03-06:02:15, 06:02:17-06:02:42, 06:02:44-06:03:02, 06:07-06:13, 06:15-06:34:01, 06:36-06:39, 06:41-06:71, 06:73-06:78, 06:80, 06:82-06:100, 06:102-06:122, 06:124-06:126, 06:128N-06:135, 06:137-06:142, 06:145-06:152N, 06:154-06:164, 07:31:01-07:31:02, 07:125, 07:177, 07:356, 12:03:01:01-12:07, 12:11-12:13, 12:15, 12:19, 12:23, 12:25-12:26, 12:28-12:29, 12:31-12:35, 12:37-12:39N, 12:42Q-12:43, 12:45-12:48, 12:50-12:55, 12:57-12:63, 12:65-12:66, 12:70-12:71, 12:75-12:79, 12:81-12:82, 12:87-12:95, 12:97-12:102, 12:107-12:111, 12:113, 12:115-12:116, 12:119-12:122, 12:125, 12:129, 12:131, 12:133, 12:135, 12:138-12:141, 12:143-12:144, 12:147, 12:149-12:150, 12:152, 12:154, 12:156-12:160, 14:15-14:16, 16:04:01, 16:04:03, 16:29, 16:33, 16:35, 16:42, 16:48, 16:55, 16:61, 16:66, 16:78, 16:82
87¹³	130 bp, 255 bp, 555 bp	1070 bp	*02:02:01 ^w , 02:02:02:01-02:02:03, 02:02:05-02:13, 02:15-02:26:03, 02:28-02:40:02, 02:42-02:64, 02:66-02:86, 02:88-02:106, 03:07, 03:10, 03:15, 03:29, 03:45, 03:58, 03:86, 03:94, 03:99, 03:163, 03:209, 03:268, 03:297, 04:03:01-04:03:02, 04:06, 04:16, 04:37, 04:80, 04:103, 04:107, 04:147, 04:160, 04:171, 04:190, 05:01:01:01-05:01:31, 05:03-05:19, 05:21-05:42, 05:44:01-05:77, 05:79-05:112, 05:114-05:122, 06:03:01-06:03:02, 06:76:01-06:76:02, 06:132:01-06:132:02, 08:10, 12:04:01-12:05, 12:09, 12:21, 12:33, 12:41, 12:60, 12:72, 12:135, 12:146, 12:154, 14:45, 15:02:01:01-15:06:03, 15:08-15:13, 15:15-15:19, 15:21-15:22, 15:24, 15:26-15:35, 15:37-15:42, 15:44-15:62, 15:64-15:84Q, 15:86-15:117, 16:02:01-16:02:13, 16:09, 16:12, 16:18-16:19, 16:25, 16:37, 16:46-16:48, 16:60, 16:63, 16:69-16:70, 16:74, 16:77N, 16:84, 17:01:01:01-17:19, 17:21, 17:23-17:30, 18:09, B*15:277, B*35:222
88¹²	160 bp, 260 bp, 445 bp	1070 bp	*01:67, 03:155, 03:231, 04:11, 04:29, 04:36, 04:55, 04:114, 04:172, 07:64, 07:402, 08:95, 12:55, 14:02:01-14:11, 14:13-14:48, 14:50-14:69, 14:71, 16:40, 16:53, A*30:62, B*35:252
89^{5,10}	110 bp, 325 bp	800 bp	*01:90, 01:101-01:102, 02:06, 02:23, 02:36, 02:68, 03:81, 03:175, 03:199, 03:245, 04:108, 04:112, 04:169, 04:178, 05:36, 06:89, 07:123, 07:173, 07:294, 08:113, 12:08, 12:15, 12:81, 12:113, 15:02:01:01-15:13, 15:15-15:19, 15:21-15:24, 15:26-15:117, 16:20, 16:64, 16:70
90¹²	180 bp, 210 bp, 240 bp	1070 bp	*02:13, 02:18, 02:33, 02:49, 02:75, 04:01:01:01-04:01:22, 04:01:24-04:01:68, 04:03:01-04:10, 04:12-04:20, 04:23-04:32, 04:34-04:106, 04:108-04:115N, 04:117-04:129, 04:131-04:168, 04:170N-04:171, 04:173N-04:206, 05:17, 05:25, 05:42, 05:55, 05:68, 05:76, 05:79, 05:115, 06:05, 06:31, 06:76:02, 06:118, 07:02:09, 07:31:01-07:31:02, 07:154, 07:177, 08:01:01-08:01:17,

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101.702-24u/06u – without *Taq* pol., IFU-02

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Lot-specific information

			08:01:19, 08:03:01-08:03:03, 08:06, 08:08:01-08:11, 08:14, 08:16:01-08:16:02, 08:20-08:22, 08:24, 08:26N, 08:28, 08:36N, 08:38, 08:40-08:42, 08:44, 08:46-08:47, 08:50, 08:56-08:61, 08:65, 08:72:01-08:72:02, 08:78-08:89N, 08:91, 08:95-08:99, 08:101-08:102, 08:105-08:106, 08:109, 08:117, 08:119, 08:121N-08:122, 12:14:01-12:14:02, 12:28, 12:58, 12:132, 12:135, 12:146, 14:10, 14:15, 14:17, 15:12, 15:25, 15:31, 15:62, 15:65, 15:75, 16:01:01-16:02:13, 16:04:01, 16:04:03, 16:06-16:39:02, 16:41-16:42, 16:44-16:52, 16:54-16:86
91^{11,15}	225 bp, 250 bp	800 bp	*01:60, 04:58, 04:160, 05:23, 05:62, 06:118, 08:07, 08:47, 08:104, 12:14:01-12:14:02, 14:17, 15:25, 15:65, 17:01:01-17:30
92	210 bp, 425 bp	800 bp	*01:02:01-01:03, 01:06-01:07:01, 01:08, 01:10-01:20, 01:23-01:34, 01:37N-01:48, 01:51-01:54, 01:56N-01:78, 01:80-01:107, 01:109N-01:111N, 03:58, 03:86, 03:94, 03:99, 04:37, 05:16, 05:85, 05:107, 06:05-06:06, 06:17, 06:31, 07:07, 07:09, 07:49, 07:76:01-07:76:02, 07:210, 07:238, 07:247, 07:315, 07:328, 07:403, 07:406, 08:12, 12:09, 12:24, 14:02:01-14:05, 14:07N, 14:10-14:14, 14:17-14:27, 14:29-14:52, 14:55-14:62, 14:64-14:69, 14:71, 15:102, 16:04:01, 16:04:03, 16:29, 16:33, 16:42, 16:53, 16:55, 16:61, 16:66, 16:68, 16:78, 16:82, 18:01-18:09, B*07:239, B*14:03
93	325 bp, 380 bp	1070 bp	*01:03, 01:24, 01:34, 01:49:01-01:50, 01:55, 01:78, 02:22, 02:62, 03:03:01-03:04:48, 03:06:01-03:11:02, 03:14, 03:18:01-03:24, 03:26, 03:28-03:32, 03:34, 03:37:01-03:59, 03:61-03:70, 03:72-03:83, 03:85, 03:87:01-03:88, 03:90-03:93, 03:96, 03:98, 03:100-03:107, 03:109, 03:111-03:120, 03:122-03:131, 03:133-03:134, 03:136-03:138, 03:140-03:145, 03:147-03:153, 03:155-03:166, 03:168, 03:170-03:176, 03:179-03:189N, 03:191-03:193, 03:195-03:196, 03:202-03:215, 03:217-03:220, 03:223, 03:227-03:237, 03:239-03:244Q, 03:246-03:247, 03:250-03:257, 03:259-03:261, 03:262 [?] , 03:263, 03:265N-03:270, 03:272-03:280, 03:282-03:291, 03:293-03:295, 03:297, 04:01:01-04:01:15, 04:01:17-04:01:68, 04:03:01-04:20, 04:24-04:53, 04:55-04:71, 04:73-04:107, 04:109-04:206, 05:01:01-05:01:31, 05:03, 05:05-05:21, 05:23-05:102, 05:104-05:106, 05:108-05:122, 06:09, 06:14, 06:35, 06:72, 06:143-06:144, 07:10, 07:28, 07:41, 07:43, 07:184, 07:196, 07:367, 08:01:01-08:08:02, 08:10, 08:12-08:47, 08:49-08:63, 08:65-08:82, 08:84-08:107, 08:109-08:112, 08:114-08:117, 08:119-08:123, 12:31, 12:44, 12:144, 14:54, 15:02:01-15:13, 15:15-15:19, 15:21, 15:23-15:36, 15:38-15:54, 15:56-15:57, 15:59-15:64, 15:66-15:71, 15:73-15:101, 15:103-15:117, 16:45, 17:01:01-17:30, 18:01-18:09
94¹³	135 bp	800 bp	*02:02:29, 03:02:01-03:02:09, 03:02:11-03:03:20, 03:03:22-03:04:24, 03:04:27-03:11:02, 03:13:01-03:17, 03:18:02-03:38:02, 03:40:01-03:64:01, 03:65-03:66, 03:67 ^w , 03:68-03:98, 03:100-03:136, 03:138-03:143, 03:146-03:155, 03:157-03:165, 03:167-03:169Q, 03:171, 03:173-03:181, 03:183-03:194, 03:196-03:230, 03:232-03:247, 03:249-03:263, 03:265N-03:277N, 03:278 ^w , 03:279-03:280, 03:282-03:294, 03:295 ^w , 03:296-03:297, 04:32, 04:77, 06:03:01, 06:132:01-06:132:02, 07:96:01-07:96:02, 07:314:02, 14:25, 15:02:10, 15:02:17, 15:43, 18:03, B*13:13:02
95¹³	155 bp, 235 bp	1070 bp	*04:14, 04:42:01-04:42:02, 04:68, 05:112, 06:02:01-01-06:02:01:03, 06:02:03-06:02:44, 06:04:01-06:75, 06:78-06:131, 06:133-06:164, 07:01:01-07:01:40, 07:01:41 ^w , 07:01:42-07:02:07, 07:02:09-07:02:28, 07:02:30-07:02:40, 07:02:41 ^w ,

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101.702-24u/06u – without *Taq* pol., IFU-02

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Lot-specific information

	07:02:42-07:25, 07:27:01-07:32N, 07:35-07:38:02, 07:41-07:63, 07:65-07:91, 07:93-07:95, 07:97-07:138, 07:140-07:151, 07:153- 07:155, 07:157-07:176, 07:178-07:209, 07:211-07:222, 07:223 ^w , 07:224-07:237, 07:239-07:245, 07:247-07:266, 07:268-07:294, 07:297-07:313, 07:315-07:316, 07:318-07:321, 07:323-07:334, 07:336-07:350N, 07:352-07:401, 07:403-07:404, 07:406-07:414, 07:415 ^w -07:416 ^w , 07:417-07:446, 12:16, 12:147, 15:75, 16:01:01:01-16:02:13, 16:06-16:28, 16:30N-16:32, 16:34, 16:36- 16:39:02, 16:41, 16:43-16:47, 16:49-16:52, 16:54, 16:56-16:60, 16:62-16:65, 16:67, 16:69-16:77N, 16:79-16:81, 16:83-16:86, 17:20, 18:01-18:08, A*24:64, A*24:96, A*24:106, A*24:174, B*08:16, B*08:123, B*08:145		
96 ¹⁶	-	-	Negative control

¹Alleles are assigned by the presence of specific PCR product(s). However, the sizes of the specific PCR products may be helpful in the interpretation of HLA-C low resolution SSP typings.

When the primers in a primer mix can give rise to HLA-specific PCR products of more than one length this is indicated if the size difference is more than 20 base pairs. Size differences of 20 base pairs or less are not given. For high resolution SSP kits, the alleles listed are specified according to amplicon length.

Nonspecific amplifications, i.e. a ladder or a smear of bands, may sometimes be seen. GC-rich primers have a higher tendency of giving rise to nonspecific amplifications than other primers.

PCR fragments longer than the control bands may sometimes be observed. Such bands should be disregarded and do not influence the interpretation of the SSP typings.

PCR fragments migrating faster than the control bands, but slower than a 400 bp fragment may be seen in some gel read-outs. Such bands can be disregarded and do not influence the interpretation of the SSP typings.

Some primers may give rise to primer oligomer artifacts. Sometimes this phenomenon is an inherent feature of the primer pair(s) of a primer mix. More often it is due to other factors such as too low amount of DNA in the PCR reactions, taking too long time in setting up the PCR reactions, working at elevated room temperature or using thermal cyclers that are not pre-heated.

²The internal positive control primer pairs amplify segments of the human growth hormone gene. The internal positive control bands are 1070 or 800 base pairs respectively, well distribution as outlined in the table. Well number 73 contains the shorter, 800 bp, internal positive control band. The well distribution of the internal controls can help in orientation of the kit on gel photo, as well as allow for kit identification. In the presence of a specific amplification the intensity of the control band often decreases.

³For several HLA Class I alleles 1st and/or 4th exon(s) and beyond, as well as intron nucleotide sequences, are not available. In these instances it is not known whether some of the primers of the SSP sets are completely matched with the target sequences or not. Assumption is made that unknown sequences in these regions are conserved within allelic groups.

⁴Only HLA-C alleles will be amplified by the 23 wells of the HLA-C low resolution primer set, except that a few HLA-A and HLA-B alleles will be amplified by primer mixes 73, 80, 85, 87, 88, 92, 94 and 95.

⁵HLA-specific PCR products shorter than 125 base pairs have a lower intensity and are less sharp than longer PCR products.

⁶Primer mix 74 will for most C*02 alleles give rise to two specific PCR fragments.

⁷Primer mix 76 will for most C*03 alleles give rise to two specific PCR fragments.

⁸Primer mix 81 will for most C*07 alleles give rise to two specific PCR fragments.

⁹Primer mix 82 will for most C*08 alleles give rise to multiple specific PCR fragments.

¹⁰Primer mix 89 will for most C*15 alleles give rise to two specific PCR fragments.

¹¹Primer mixes 84, 85 and 91 may have tendencies of unspecific amplifications.

¹²Primer mixes 88 and 90 have a tendency to giving rise to primer oligomer formation.

¹³Primer mixes 75, 78, 82, 87, 94 and 95 may yield less HLA-specific PCR fragments than the other HLA-C low resolution primer mixes.

¹⁴Primer mix 86 might faintly amplify most C*01 and the C*14 alleles.

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Lot-specific information

¹⁵Primer mix 91 might generate a false band of about 500 base pairs. This band should be disregarded when interpreting HLA-C low resolution typings.

¹⁶Primer mix 96 contains a negative control, which will amplify more than 95% of HLA amplicons as well as the amplicons generated by the control primer pairs matching the human growth hormone gene. HLA-specific PCR product sizes range from 75 to 200 base pairs and the PCR product generated by the HGH positive control primer pair is 430 base pairs.

‘w’, might be weakly amplified.

‘?’, nucleotide sequence information not available for the primer matching sequence.

101.702-24/06 – including *Taq* pol., IFU-01
101.702-24u/06u – without *Taq* pol., IFU-02

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Lot No.: **3D7**

Lot-specific information

HLA-A LOW PRIMER SPECIFICATION

Well No.	1	2	3	4	5	6	7	8	9	10	11	12
Length of spec. PCR product	120	175	205	190	160	135	175	165	75	80	80	125
	145	215	235		335	200	205	200		240	175	195
	225	255			505						500	
		365										
		415										
Length of int. pos. control ¹	800	800	1070	800	1070	800	1070	800	800	800	1070	800
5'-primer(s) ²	98	48	357	98	144	176	98	98	261	98	301	103
	5'-CTT 3'	5'-gCT 3'	5'-ATg 3'	5'-CTA 3'	5'-gCC 3'	5'-gCA 3'	5'-CTC 3'	5'-CTA 3'	5'-AAC 3'	5'-CTA 3'	5'-Cgg 3'	5'-CCT 3'
	103	78	363	413	317	368	368	102	266	261	302	415
	5'-CCT 3'	5'-TCT 3'	5'-ATA 3'	5'-CCg 3'	5'-gCT 3'	5'-gTT 3'	5'-gTT 3'	5'-ACA 3'	5'-ACg 3'	5'-AAC 3'	5'-ggA 3'	5'-ggT 3'
	123	106	363					413			385	423
	5'-AgT 3'	5'-CCA 3'	5'-ATA 3'					5'-CCg 3'			5'-ggC 3'	5'-gCT 3'
	363											
	5'-ATA 3'											
3'-primer(s) ³	203	240	527	256	265	270	259	259	302	299	341	257
	5'-TCT 3'	5'-ggA 3'	5'-CCA 3'	5'-CTg 3'	5'-CCC 3'	5'-ACA 3'	5'-gTT 3'	5'-gTT 3'	5'-ggC 3'	5'-TCg 3'	5'-CgT 3'	5'-gCA 3'
	545	292	527	559	368	521	502	259	303		521	506
	5'-AgA 3'	5'-gTg 3'	5'-CCT 3'	5'-CCg 3'	5'-CAA 3'	5'-ggg 3'	5'-CTT 3'	5'-gTT 3'	5'-AgA 3'		5'-ggg 3'	5'-TgT 3'
			527			534	539	538				559
			5'-CCT 3'			5'-CgT 3'	5'-TCT 3'	5'-CCA 3'				5'-CTC 3'
			555									559
			5'-CCA 3'									5'-CgT 3'
			555									559
			5'-gCA 3'									5'-CgC 3'

Well No.	13	14	15	16	17	18	19	20	21	22	23	24
Length of spec. PCR product	175	80	90	240	140	200	340	210	200	85	85	360
	225	115	135	380	180	390	375	240	240	240	160	
		200	205	410					375	400	240	
		240							545		495	
		470										
Length of int. pos. control ¹	1070	1070	1070	1070	1070	1070	800	800	800	800	800	1070
5'-primer(s) ²	98	98	203	41	180	41	302	78	28	47	176	341
	5'-CTT 3'	5'-CAC 3'	5'-gAA 3'	5'-CTT 3'	5'-TTT 3'	5'-CTT 3'	5'-ggA 3'	5'-TCT 3'	5'-TCg 3'	5'-g.T 3'	5'-gCA 3'	5'-ggC 3'
	423	219	362	355	203	98	302	106	261	78	261	
	5'-gCT 3'	5'-gCA 3'	5'-ggT 3'	5'-CCg 3'	5'-gAA 3'	5'-CAC 3'	5'-ggA 3'	5'-CCA 3'	5'-AAC 3'	5'-TCT 3'	5'-AAC 3'	
		238	363		418		341	2 nd I	368	527	341	
		5'-AgA 3'	5'-ATA 3'		5'-AgC 3'		5'-ggA 3'	5'-CCT 3'	5'-gTT 3'	5'-TgC 3'	5'-ggA 3'	
		355	363								355	
		5'-CCg 3'	5'-ATA 3'								5'-CCC 3'	
			363								362	
			5'-ATA 3'								5'-gAg 3'	
			369								362	
			5'-TAC 3'								5'-gAg 3'	
3'-primer(s) ³	282	180	299	238	290	256	397	265	97	265	292	418
	5'-gAC 3'	5'-TCA 3'	5'-CCA 3'	5'-CCT 3'	5'-CAA 3'	5'-CCC 3'	5'-gAg 3'	5'-CCC 3'	5'-ggT 3'	5'-CCC 3'	5'-gTg 3'	5'-gTC 3'
	282	257	411	238	317	256		282	355	282	292	
	5'-gAC 3'	5'-gCA 3'	5'-TCA 3'	5'-CCT 3'	5'-ggA 3'	5'-CTC 3'		5'-gAC 3'	5'-gAC 3'	5'-gAC 3'	5'-gTT 3'	
	559	418	526	243	555	259		282	524	282	299	
	5'-CCC 3'	5'-gTC 3'	5'-CCA 3'	5'-TCA 3'	5'-CCA 3'	5'-gTT 3'		5'-gAC 3'	5'-CAT 3'	5'-gAC 3'	5'-TCT 3'	
	559	555		265				502	538	570	555	
	5'-CCg 3'	5'-CCA 3'		5'-CCC 3'				5'-CTT 3'	5'-CCA 3'	5'-CAC 3'	5'-CCA 3'	
				282				506				
				5'-gAC 3'				5'-TgT 3'				
				555								
				5'-CCA 3'								

101.702-24/06 – including *Taq* pol., IFU-01
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Lot No.: **3D7**

Lot-specific information

¹The internal positive control primer pairs amplify segments of the human growth hormone gene. The internal positive control bands are 1070 or 800 base pairs respectively, well distribution as outlined in the table. Well number 1 contains the shorter, 800 bp, internal positive control band. The well distribution of the internal controls can help in orientation of the kit on gel photo, as well as allow for kit identification. In the presence of a specific amplification the intensity of the control band often decreases.

²The nucleotide position matching the specificity-determining 3'-end of the primer is given. Nucleotide numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given.

³The nucleotide position matching the specificity-determining 3'-end of the primer is given in the anti-sense direction. Nucleotide numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given.

HLA-B LOW PRIMER SPECIFICATION

Well No.	25	26	27	28	29	30	31	32	33	34	35	36
Length of spec. PCR product	110	215	140	130	185	190	290	165	165	180	100	280
			235	265	235			225	190		195	
								285	390			
								330				
Length of int. pos. control ¹	800	1070	1070	800	800	800	1070	1070	1070	800	1070	1070
5'-primer(s) ²	527	97	209	103	103	103	45	45	45	420	206	419
	5'-TgA 3'	5'-TCg 3'	5'-ggC 3'	5'-CCg 3'	5'-CCT 3'	5'-CCg 3'	5'-ggA 3'	5'-ggA 3'	5'-ggA 3'	5'-TTA 3'	5'-AgA 3'	5'-gTC 3'
	540		363	103	103	418		357	435		420	
	5'-gAC 3'		5'-AgC 3'	5'-CCg 3'	5'-CCT 3'	5'-Agg 3'		5'-Tgg 3'	5'-AAA 3'		5'-TTA 3'	
			363	361	363			412				
			5'-AgC 3'	5'-AgT 3'	5'-AAT 3'			5'-ATA 3'				
3'-primer(s) ³	605	272	309	193	246	246	165	206	266	559	256	3 rd I
	5'-gCT 3'	5'-Tgg 3'	5'-gTg 3'	5'-CgT 3'	5'-TAT 3'	5'-TAT 3'	5'-Tgg 3'	5'-CCT 3'	5'-TCC 3'	5'-CTC 3'	5'-CCC 3'	5'-TAT 3'
		272	312	193	246	572		538	559		272	
		5'-TgA 3'	5'-gCC 3'	5'-CgT 3'	5'-TAT 3'	5'-gCC 3'		5'-gTC 3'	5'-CAg 3'		5'-Tgg 3'	
			559	463	559			603	583		572	
			5'-CAg 3'	5'-gCT 3'	5'-CTC 3'			5'-gTg 3'	5'-gTg 3'		5'-gCg 3'	
				583								
				5'-gTg 3'								
Well No.	25	26	27	28	29	30	31	32	33	34	35	36

101.702-24/06 – including *Taq* pol., IFU-01
101.702-24u/06u – without *Taq* pol., IFU-02

Visit www.olerup.com for
“Instructions for Use” (IFU)

Lot No.: **3D7**

Lot-specific information

Well No.	37	38	39	40	41	42	43	44	45	46	47	48
Length of spec. PCR product	195	105 395 435	115 150	80	150	140	55 245 390 415	210	170	110	395	160 425
Length of int. pos. control ¹	1070	1070	1070	1070	800	1070	1070	800	800	1070	1070	1070
5'-primer(s) ²	142 5'-TCT 3' 419 5'-gTC 3'	45 5'-ggA 3' 540 5'-gAC 3'	161 5'-Cgg 3'	167 5'-gCT 3'	355 5'-TCA 3'	206 5'-gAC 3'	45 5'-ggA 3' 368 5'-gTT 3' 557 5'-ggA 3'	142 5'-TCT 3' 368 5'-gTC 3'	409 5'-ggC 3' 420 5'-TTA 3'	246 5'-gAA 3'	44 5'-ggC 3'	44 5'-ggC 3' 357 5'-Tgg 3'
3'-primer(s) ³	301 5'-gTC 3' 301 5'-gTC 3' 570 5'-CCg 3' 572 5'-gCg 3'	272 5'-TgC 3' 272 5'-TgT 3' 309 5'-ATC 3' 605 5'-gCT 3'	234 5'-TCT 3'	204 5'-TCT 3'	463 5'-gCT 3'	302 5'-ggC 3' 312 5'-AgT 3'	259 5'-gTT 3' 259 5'-gTT 3' 266 5'-TCC 3' 272 5'-Tgg 3' 292 5'-gTA 3' 572 5'-gCg 3'	311 5'-ggg 3' 311 5'-ggg 3' 538 5'-gTC 3'	544 5'-ggT 3'	317 5'-ggA 3'	272 5'-TgC 3'	302 5'-ggC 3' 302 5'-ggT 3' 477 5'-gCg 3'
Well No.	37	38	39	40	41	42	43	44	45	46	47	48

Well No.	49	50	51	52	53	54	55	56	57	58	59	60
Length of spec. PCR product	180	290 465	105	325	80 115 160 195 225 260	215	130 270	90 410	90 175	145 430	120 210	430
Length of int. pos. control ¹	1070	1070	1070	800	1070	800	800	1070	1070	1070	800	1070
5'-primer(s) ²	355 5'-TCA 3' 363 5'-Agg 3'	72 5'-gCT 3' 209 5'-ggC 3'	540 5'-gAC 3'	1 st 5'-CAg 3'	355 5'-TCA 3' 418 5'-Agg 3' 499 5'-TCT 3'	141 5'-ATT 3' 357 5'-Tgg 3'	209 5'-ggC 3' 209 5'-ggg 3' 355 5'-TCA 3'	41 5'-CTg 3' 368 5'-gTT 3'	209 5'-ggC 3' 363 5'-AgC 3'	48 5'-gCC 3'	357 5'-Tgg 3'	49 5'-CAg 3'
3'-primer(s) ³	499 5'-ggA 3'	193 5'-CgT 3' 193 5'-TgT 3' 387 5'-TCg 3'	603 5'-gTg 3' 605 5'-gCT 3'	282 5'-gCC 3'	538 5'-gTC 3' 572 5'-gCg 3' 573 5'-AgT 3'	309 5'-ATC 3' 538 5'-gTC 3'	299 5'-TCA 3' 583 5'-gTg 3' 605 5'-gCT 3'	282 5'-gCC 3' 418 5'-gTC 3'	259 5'-gTT 3' 499 5'-ggA 3'	309 5'-ATC 3'	435 5'-TCT 3'	309 5'-ATC 3' 527 5'-CCT 3'
Well No.	49	50	51	52	53	54	55	56	57	58	59	60

101.702-24/06 – including *Taq* pol., IFU-01
101.702-24u/06u – without *Taq* pol., IFU-02

Visit www.olerup.com for
“Instructions for Use” (IFU)

Lot No.: **3D7**

Lot-specific information

Well No.	61	62	63	64	65	66	67	68	69	70	71	72
Length of spec.	145	300	160	180	180	90	90	95	115	360	350	285
PCR product			370	210	205	240			150			
Length of int. pos. control ¹	1070	1070	1070	1070	1070	800	1070	1070	1070	1070	1070	800
5'-primer(s) ²	206	1 st I	15	141	97	209	208	206	165	1 st I	1 st I	355
	5'-gAA 3'	5'-CAg 3'	5'-gCA 3'	5'-ATT 3'	5'-TCg 3'	5'-ggC 3'	5'-CgT 3'	5'-gAC 3'	5'-ACC 3'	5'-CAg 3'	5'-CAg 3'	5'-TCA 3'
			418	420	420	209	209		463			363
			5'-Agg 3'	5'-TTA 3'	5'-TTA 3'	5'-ggA 3'	5'-ggA 3'		5'-TgA 3'			5'-AgC 3'
						362	369					
						5'-ggT 3'	5'-TAC 3'					
3'-primer(s) ³	311	259	207	311	259	256	256	259	272	317	311	603
	5'-ggT 3'	5'-CTC 3'	5'-TCC 3'	5'-ggT 3'	5'-gTT 3'	5'-CCC 3'	5'-CTC 3'	5'-gTT 3'	5'-TgC 3'	5'-ggA 3'	5'-ggT 3'	5'-gTg 3'
		259	226	559	559	259	256		538			605
		5'-CTC 3'	5'-CAC 3'	5'-CgT 3'	5'-CAg 3'	5'-CTT 3'	5'-CCC 3'		5'-CCA 3'			5'-gCT 3'
		262	538			263	420					
		5'-TgC 3'	5'-gTC 3'			5'-gTT 3'	5'-gCT 3'					
						559						
						5'-CgT 3'						
Well No.	61	62	63	64	65	66	67	68	69	70	71	72

¹The internal positive control primer pairs amplify segments of the human growth hormone gene. The internal positive control bands are 1070 or 800 base pairs respectively, well distribution as outlined in the table. Well number 25 contains the shorter, 800 bp, internal positive control band. The well distribution of the internal controls can help in orientation of the kit on gel photo, as well as allow for kit identification. In the presence of a specific amplification the intensity of the control band often decreases.

²The nucleotide position matching the specificity-determining 3'-end of the primer is given. Nucleotide numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given.

³The nucleotide position matching the specificity-determining 3'-end of the primer is given in the anti-sense direction. Nucleotide numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given.

101.702-24/06 – including *Taq* pol., IFU-01
101.702-24u/06u – without *Taq* pol., IFU-02

Visit www.olerup.com for
“Instructions for Use” (IFU)

Lot No.: **3D7**

Lot-specific information

HLA-C LOW PRIMER SPECIFICATION

Well No.	73	74	75	76	77	78	79	80	81	82	83	84
Length of spec. PCR product	90	130	280	170	280	135	165	130	245	110	340	100
	150	200		275		330	390	355	425	165		155
		270					445			390		220
		300										
Length of int. pos. control ¹	800	800	800	1070	800	800	1070	800	800	800	1070	800
5'-primer(s) ²	89	47	105	355	105	47	1 st	28	47	1 st	1 st	361
	5'-gAA 3'	5'-Agg 3'	5'-gCT 3'	5'-TCA 3'	5'-gCT 3'	5'-Agg 3'	5'-CgA 3'	5'-TCA 3'	5'-Agg 3'	5'-CgA 3'	5'-CgA 3'	5'-AgT 3'
	89	89		459		112	176	2 nd	648	176		419
	5'-gAA 3'	5'-gAA 3'		5'-gAT 3'		5'-CCT 3'	5'-gCA 3'	5'-CCA 3'	5'-CAC 3'	5'-gCA 3'		5'-gTC 3'
	98	2 nd				118	527			527		
	5'-CTT 3'	5'-CCA 3'				5'-CCA 3'	5'-TgA 3'			5'-TAC 3'		
		703										
		5'-CTA 3'										
3'-primer(s) ³	142	176	343	589	343	201	302	213	302	302	302	474
	5'-TgA 3'	5'-ACT 3'	5'-C 3'	5'-CTT 3'	5'-T 3'	5'-CTT 3'	5'-ggT 3'	5'-Cgg 3'	5'-ggC 3'	5'-ggC 3'	5'-ggT 3'	5'-gCA 3'
	201	559	343			218	3 rd	409	853	595	304	477
	5'-CTT 3'	5'-CTC 3'	5'-g 3'			5'-gCT 3'	5'-gCA 3'	5'-ATg 3'	5'-CAT 3'	5'-CCC 3'	5'-CAA 3'	5'-gCA 3'
	201	861						420				538
	5'-CTC 3'	5'-TCg 3'						5'-gCT 3'				5'-gCA 3'

Well No.	85	86	87	88	89	90	91	92	93	94	95
Length of spec. PCR product	120	160	130	160	110	180	225	210	325	135	155
	250	210	255	260	325	210	250	425	380		235
			555	445		240					
Length of int. pos. control ¹	800	800	1070	1070	800	1070	800	800	1070	800	1070
5'-primer(s) ²	201	97	98	98	201	201	2 nd	47	355	105	97
	5'-CCA 3'	5'-TCg 3'	5'-CTA 3'	5'-CTC 3'	5'-CCA 3'	5'-CCA 3'	5'-CCA 3'	5'-Agg 3'	5'-TCA 3'	5'-gCT 3'	5'-TCg 3'
	218	361	368	98	409	2 nd		361	412	459	97
	5'-ggA 3'	5'-AgT 3'	5'-gTg 3'	5'-CTT 3'	5'-ggC 3'	5'-CCA 3'		5'-AgT 3'	5'-ATA 3'	5'-gAT 3'	5'-TTg 3'
	2 nd			194		361					361
	5'-CCA 3'			5'-CgT 3'		5'-AgT 3'					5'-AgT 3'
				527							418
				5'-TgA 3'							5'-Agg 3'
											419
											5'-gTC 3'
											419
											5'-gTT 3'
3'-primer(s) ³	289	218	312	311	270	341	512	302	3 rd	201	213
	5'-AgC 3'	5'-gCT 3'	5'-AgT 3'	5'-ggT 3'	5'-TAG 3'	5'-CgT 3'	5'-CCA 3'	5'-ggT 3'	5'-CTC 3'	5'-CTC 3'	5'-Cgg 3'
	289	527	361	317	3 rd	343	538	527		559	289
	5'-AgC 3'	5'-CCg 3'	5'-CCA 3'	5'-CgT 3'	5'-CTC 3'	5'-T 3'	5'-gTC 3'	5'-CCg 3'		5'-CTC 3'	5'-AgC 3'
	291	538	459	3 rd		343		538			289
	5'-TCg 3'	5'-CCA 3'	5'-AgA 3'	5'-gCA 3'		5'-g 3'		5'-CCg 3'			5'-AgC 3'
	539	538				527					539
	5'-TCT 3'	5'-gCA 3'				5'-CCg 3'					5'-TCT 3'
						527					
						5'-CCg 3'					
						527					
						5'-CCg 3'					
						527					
						5'-CCg 3'					
						530					
						5'-CCA 3'					
Well No.	85	86	87	88	89	90	91	92	93	94	95

101.702-24/06 – including *Taq* pol., IFU-01
 101.702-24u/06u – without *Taq* pol., IFU-02

Visit www.olerup.com for
 “Instructions for Use” (IFU)

Lot No.: 3D7

Lot-specific information

¹The internal positive control primer pairs amplify segments of the human growth hormone gene. The internal positive control bands are 1070 or 800 base pairs respectively, well distribution as outlined in the table. Well number 73 contains the shorter, 800 bp, internal positive control band. The well distribution of the internal controls can help in orientation of the kit on gel photo, as well as allow for kit identification. In the presence of a specific amplification the intensity of the control band often decreases.

²The nucleotide position matching the specificity-determining 3'-end of the primer is given. Nucleotide numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given.

³The nucleotide position matching the specificity-determining 3'-end of the primer is given in the anti-sense direction. Nucleotide numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given.

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Lot-specific information

CE
0088

101.702-24/06 – including *Taq* pol., IFU-01
101.702-24u/06u – without *Taq* pol., IFU-02

Visit www.olerup.com for
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Lot No.: **3D7**

Lot-specific information

CELL LINE VALIDATION SHEET													
HLA-A low resolution primer set ²													
					Prod. No.:	Well							
						17	18	19	20	21	22	23	24
						201553417	201553418	201553419	201553420	201561121	201559022	201553423	201552924
	IHCW cell line ¹		A*	A*									
1	9001	SA	*24:02			-	-	-	-	-	-	-	-
2	9280	LK707	*02:01			-	-	-	-	-	-	-	-
3	9011	E4181324	*01:01			-	-	-	-	-	-	-	-
4	9275	GU373	*30:01			-	-	-	-	-	-	-	-
5	9009	KAS011	*01:01			-	-	-	-	-	-	-	-
6	9353	SM	*02:01	*26:03		-	-	-	-	-	-	+	-
7	9020	QBL	*26:01			-	-	-	-	-	-	-	-
8	9025	DEU	*31:01			-	-	-	-	-	-	-	-
9	9026	YAR	*26:01			-	-	-	-	-	-	-	-
10	9107	LKT3	*24:02			-	-	-	-	-	-	-	-
11	9051	PITOUT	*29:02			-	-	-	-	-	-	-	-
12	9052	DBB	*02:01			-	-	-	-	-	-	-	-
13	9004	JESTHOM	*02:01			-	-	-	-	-	-	-	-
14	9071	OLGA	*31:01			-	-	-	-	-	-	-	-
15	9075	DKB	*24:02			-	-	-	-	-	-	-	-
16	9037	SWEIG007	*29:02			-	-	-	-	-	-	-	-
17	9282	CTM3953540	*03:01	*80:01		-	-	-	-	-	-	+	-
18	9257	32367	*33:03	*74:01		-	+	+	-	-	-	-	-
19	9038	BM16	*02:01			-	-	-	-	-	-	-	-
20	9059	SLE005	*02:01			-	-	-	-	-	-	-	-
21	9064	AMALA	*02:17			-	-	-	-	+	-	-	-
22	9056	KOSE	*02:01			-	-	-	-	-	-	-	-
23	9124	IHL	*02:01	*34:01		-	-	-	-	-	-	-	-
24	9035	JBUSH	*32:01			+	-	-	-	-	-	-	-
25	9049	IBW9	*33:01			-	+	-	-	-	-	-	-
26	9285	WT49	*02:05			-	-	-	-	-	-	-	-
27	9191	CH1007	*24:10	*29:01		-	-	-	-	-	-	-	-
28	9320	BEL5GB	*02:01	*29:02		-	-	-	-	-	-	-	-
29	9050	MOU	*29:02			-	-	-	-	-	-	-	-
30	9021	RSH	*30:01	*68:02		-	-	-	+	-	-	-	-
31	9019	DUCAF	*30:02			-	-	-	-	-	-	-	-
32	9297	HAG	*02:01			-	-	-	-	-	-	-	-
33	9098	MT14B	*31:01			-	-	-	-	-	-	-	-
34	9104	DHIF	*31:01			-	-	-	-	-	-	-	-
35	9302	SSTO	*32:01			+	-	-	-	-	-	-	-
36	9024	KT17	*02:06	*11:01		-	-	-	-	-	-	-	-
37	9065	HHKB	*03:01			-	-	-	-	-	-	-	-
38	9099	LZL	*02:17			-	-	-	-	-	-	-	-
39	9315	CML	*01:01	*03:01		-	-	-	-	-	-	-	-
40	9134	WHONP199	*02:07	*30:01		-	-	-	-	-	-	-	-
41	9055	H0301	*03:01			-	-	-	-	-	-	-	-
42	9066	TAB089	*02:07			-	-	-	-	-	-	-	-
43	9076	T7526	*02:06	*02:07		-	-	-	-	-	-	-	-
44	9057	TEM	*66:01			-	-	-	-	-	-	-	-
45	9239	SHJO	*23:01	*24:02		-	-	-	-	-	-	-	-
46	9013	SCHU	*03:01			-	-	-	-	-	-	-	-
47	9045	TUBO	*02:16	*03:01		-	-	-	-	-	-	-	-
48	9303	TER-ND	*02:01	*11:01		-	-	-	-	-	-	-	-

101.702-24/06 – including *Taq* pol., IFU-01
 101.702-24u/06u – without *Taq* pol., IFU-02

Visit www.olerup.com for
 “Instructions for Use” (IFU)

Lot No.: 3D7

Lot-specific information

¹The provided cell line HLA specificities are retrieved from the <http://www.ihwg.org/hla> web site. The specificity of an individual cell line may thus be subject to change.

²The specificity of each primer solution in the kit has been tested against 48 well characterized cell line DNAs and where applicable, additional cell line DNAs.

Additional 5'- and 3'-primers in primer solutions 4, 5, 7, 8, 12, 14 to 17 and 20 to 23 were tested by separately adding one 3'-primer, respectively one 5'-primer. Additional 5'-primers in primer solution 1 and 19 were tested by separately adding one 3'-primer. Additional 3'-primers in primer solutions 3 and 18 were tested by separately adding one 5'-primer.

In primer solutions 2, 3, 11, 12, 15 and 22 one or two 5'-primers were not possible to test, and in primer solutions 3, 6, 8, 9, 12, 13, 16 and 18 one 3'-primer was not possible to test.

101.702-24/06 – including *Taq* pol., IFU-01
101.702-24u/06u – without *Taq* pol., IFU-02

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Lot No.: **3D7**

Lot-specific information

¹The provided cell line HLA specificities are retrieved from the <http://www.ihwg.org/hla> web site. The specificity of an individual cell line may thus be subject to change.

²The specificity of each primer solution in the kit has been tested against 48 well characterized cell line DNAs and where applicable, additional cell line DNAs.

Additional 5'-primers and 3'-primers in primer solutions 27, 30, 38, 43, 53, 55, 64 and 65 were tested by separately adding one additional 3'-primer, respectively one additional 5'-primer. Additional 3'-primers in primer solutions 26, 32, 35, 37, 39 and 42 were tested by separately adding one additional 5'-primer. Additional 5'-primers in primer solutions 25, 45, 49 and 72 were tested by separately adding one additional 3'-primer.

In primer mixes 55, 66 and 67 one 5'-primer was not possible to test, and in primer mixes 27, 53, 62, 63 and 66 one or two 3'-primer was not possible to test.

101.702-24/06 – including *Taq* pol., IFU-01
101.702-24u/06u – without *Taq* pol., IFU-02

Visit www.olerup.com for
“Instructions for Use” (IFU)

Lot No.: **3D7**

Lot-specific information

CELL LINE VALIDATION SHEET									
HLA-C low resolution SSP primer set ²									
					Well				
					89	90	91	92	93
					94	95			
					Prod. No.:	201553117	201553118	201553119	201553120
						201553121	201553122	201553123	201550823
IHC cell line ¹					C*				
1	9001	SA	*07:02			-	-	-	-
2	9280	LK707	*07:01	*15:05		+	-	-	+
3	9011	E4181324	*12:02			-	-	-	-
4	9275	GU373	*03:04	*04:01		-	+	-	+
5	9009	KAS011	*06:02			-	-	-	-
6	9353	SM	*03:04	*07:02		-	-	-	+
7	9020	QBL	*05:01			-	-	-	+
8	9007	DEM	*04:01			-	+	-	-
9	9026	YAR	*12:03			-	-	-	-
10	9107	LKT3	*01:02			-	-	-	+
11	9051	PITOUT	*16:01			-	+	-	-
12	9052	DBB	*06:02			-	-	-	-
13	9004	JESTHOM	*01:02			-	-	-	+
14	9071	OLGA	*01:02	*03:04		-	-	-	+
15	9075	DKB	*03:04			-	-	-	+
16	9037	SWEIG007	*02:02			-	-	-	-
17	9282	CTM3953540	*03:03	*07:01		-	-	-	+
18	9257	32367	*01:02	*07:05		-	-	-	+
19	9038	BM16	*07:01			-	-	-	-
20	9059	SLE005	*03:04			-	-	-	+
21	9064	AMALA	*03:03			-	-	-	+
22	9056	KOSE	*12:03			-	-	-	-
23	9124	IHL	*01:02	*15:02		+	-	-	+
24	9035	JBUSH	*12:03			-	-	-	-
25	9049	IBW9	*08:02			-	-	-	+
26	9285	WT49	*07:01			-	-	-	-
27	9191	CH1007	*07:04	*15:05		+	-	-	+
28	9320	BEL5GB	*05:01	*16:01		-	+	-	+
29	9050	MOU	*16:01			-	+	-	-
30	9021	RSH	*17:01			-	-	+	-
31	9019	DUCAF	*05:01			-	-	-	+
32	9297	HAG	*17:01	*17:03		-	-	+	-
33	9098	MT14B	*03:04			-	-	-	+
34	9104	DHIF	*12:03			-	-	-	-
35	9302	SSTO	*05:01			-	-	-	+
36	9024	KT17	*03:03	*04:01		-	+	-	+
37	9065	HHKB	*07:02			-	-	-	-
38	9099	LZL	*03:03			-	-	-	+
39	9315	CML	*02:02	*07:01		-	-	-	-
40	9134	WHONP199	*01:02	*06:02		-	-	-	+
41	9055	H0301	*08:02			-	-	-	+
42	9066	TAB089	*01:02			-	-	-	-
43	9076	T7526	*01:02	*08:01		-	+	-	+
44	9057	TEM	*12:03			-	-	-	-
45	9239	SHJO	*06:02	*17:01		-	-	+	-
46	9013	SCHU	*07:02			-	-	-	+
47	9045	TUBO	*07:04	*15:02		+	-	-	+
48	9303	TER-ND	*04:01	*16:01		-	+	-	+

101.702-24/06 – including *Taq* pol., IFU-01
101.702-24u/06u – without *Taq* pol., IFU-02

Visit www.olerup.com for
“Instructions for Use” (IFU)

Lot No.: 3D7

Lot-specific information

¹The provided cell line HLA specificities are retrieved from the <http://www.ihwg.org/hla> web site. The specificity of an individual cell line may thus be subject to change.

²The specificity of each primer solution in the kit has been tested against 48 well characterized cell line DNAs and where applicable, additional cell line DNAs.

Additional 5'-primers and 3'-primers in primer solutions 73, 74, 79, 86 to 88 and 94 were tested by separately adding one additional 3'-primer, respectively one additional 5'-primer. Additional 3'-primers in primer solution 80, 90 and 91 were tested by separately adding one additional 5'-primer. Additional 5'-primers in primer solutions 78, 85 and 95 were tested by separately adding one additional 3'-primer.

In primer solutions 73, 88 and 95 one 5'-primer was not possible to test, and in primer solutions 75, 83 to 86, 88 and 90 one or two 3'-primers were not possible to test.

101.702-24/06 – including *Taq* pol., IFU-01
101.702-24u/06u – without *Taq* pol., IFU-02

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Lot-specific information

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