

Olerup SSP® HLA-B low resolution

Product number:	101.501-48/12 – including <i>Taq</i> pol.
Lot number:	98M
Expiry date:	2014-July-01
Number of tests:	48 tests – Product No. 101.501-48 12 tests – Product No. 101.501-12
Number of wells per test:	47 + 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. 98M.

CHANGES COMPARED TO THE PREVIOUS OLERUP SSP® HLA-B LOW RESOLUTION LOT

The HLA-B low resolution specificity and interpretation tables have been updated for the HLA-B alleles reported since the previous *Olerup SSP® HLA-B low resolution lot (Lot No. 63M)* was made.

The HLA-B low primer set is unchanged compared to the previous lot.

Changes in revision R01 compared to R00:

1. Primer mix 19 does not amplify the B*44:10 allele. This has been corrected in the Specificity and Interpretation Tables.

Well 48 contains Negative Control primer pairs, that will amplify more than 95% of the *Olerup SSP®* HLA Class I, DRB, DQB1 and DPB1 amplicons as well as the amplicons generated by control primer pairs.

PCR product sizes range from 75 to 430 base pairs.

The PCR product generated by the control primer pair is 430 base pairs.

Length of PCR product	105	200	105	80	75	80
5'-primer¹	164	340	440	45	45	43
	5'-CAC ^{3'}	5'-Agg ^{3'}	5'-TTA ^{3'}	5'-Tgg ^{3'}	5'-Tgg ^{3'}	5'-Tgg ^{3'}
3'-primer²	231	2nd 	507	59	58	57
	5'-TgC ^{3'}	5'-AAA ^{3'}	5'-TTg ^{3'}	5'-CTC ^{3'}	5'-ggC ^{3'}	5'-CTC ^{3'}
A*	+	+	+			
B*	+	+	+			
C*	+	+	+			
DRB1					+	+
DRB3					+	+
DRB5					+	
DQB1						+
DPB1						+

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 codonnumbering 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.

PRODUCT DESCRIPTION

HLA-B low resolution SSP typing

CONTENT

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.

PLATE LAYOUT

Each test consists of 48 PCR reactions in a 48 well cut 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

Wells 1 to 47 – HLA-B low resolution primers.

Well 48 – Negative Control.

The 48 well cut PCR plate is marked with 'HLA-B low' in silver/gray ink.

Well No. 1 is marked with the Lot No. '98M'.

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.

Please note: When removing each 48 well PCR plate, make sure that the remaining plates stay covered. Use a scalpel or a similar instrument to carefully cut the foil between the plates.

INTERPRETATION

Only HLA-B alleles will be amplified by the HLA-B low resolution typing kit, except that primer mix 1 will amplify the C*02:23 and C*04:77 alleles, primer mix 3 will amplify the C*03:05, C*03:25 and C*03:27 alleles, primer mix 4 will amplify the C*01:30 allele, primer mix 6 will amplify the A*23:31, A*24:106 and C*16:10 alleles, the C*07:46 allele will be amplified by primer mix 8, primer mix 13 will amplify the A*24:174 allele, primer mix 17 will amplify the C*03:102 allele, primer mix 21 will amplify the C*15:51 allele, primer mix 25 will amplify the C*03:129 allele, primer mix 29 will amplify the C*15:25 allele, primer mix 31 will amplify the C*15:39 allele, primer mix 40 will amplify the C*15:02:04 allele, primer mix 41 will amplify the C*03:12, C*03:19 and C*03:102 alleles, primer mix 43 will weakly amplify the C*02:06 and C*02:47 alleles and primer mix 45 will amplify the A*26:68, A*68:56, C*06:20 and C*12:50 alleles.

Thus, the interpretation of HLA-B low resolution typings is only influenced by the A*23:31, A*24:106, A*24:174, A*26:68, A*68:56, C*01:30, C*02:06, C*02:23, C*02:47, C*03:05, C*03:12, C*03:19, C*03:25, C*03:27, C*03:102, C*03:129,

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C*04:77, C*06:20, C*07:46, C*12:50, C*15:02:04, C*15:25, C*15:39, C*15:51 and C*16:10 alleles and not by other alleles of other HLA class I genes.

UNIQUELY IDENTIFIED ALLELES

All the HLA-B alleles recognized by the HLA Nomenclature Committee in October 2011¹ will be amplified by the primers in the HLA-B low resolution SSP kit. The HLA-B alleles will be grouped into their corresponding serological specificities^{2,3}.

¹HLA-B alleles listed on the IMGT/HLA web page 2011-October-10, release 3.6.0, www.ebi.ac.uk/imgt/hla.

²It is not possible to identify all serological splits and combinations of the B15 group of alleles, as the serological reactivity of several of the B*15 alleles is not known.

The resolution has been updated to fulfill the NMDP medium resolution confirmatory typing requirements effective from February 1st 2001.

DNA-based HLA-B typing resolution required:

B*07:xx, B*08:xx, B*13:xx, B*14:xx, B*18:xx, B*27:xx, B*15:22, B*35:xx, B*37:xx, B*38:xx, B*39:xx, B*40:05, B*41:xx, B*42:xx, B*44:xx, B*45:xx, B*46:xx, B*47:xx, B*48:xx, B*49:xx, B*50:xx, B*51:xx, B*52:xx, B*53:xx, B*54:xx, B*55:xx, B*56:xx, B*57:xx, B*58:xx, B*59:xx, B*67:xx, B*73:xx, B*78:xx, B*81:xx, B*82:xx,	
B*40EJD ^a	40:01, 40:07, 40:10, 40:12, 40:14, 40:15, 40:16
B*40EJB ^b	40:02, 40:03, 40:04, 40:06, 40:08, 40:09, 40:11, 40:13, 40:18, 40:19, 40:20
B*15EJA ^c	15:01, 15:04, 15:05, 15:06, 15:07, 15:20, 15:24, 15:25, 15:26N, 15:27, 15:28, 15:30, 15:32, 15:33, 15:34, 15:35, 15:36, 15:38, 15:39, 15:40, 15:43, 15:45
B*15AHM ^d	15:16, 15:17
B*15EHZ ^e	15:03, 15:09, 15:10, 15:18, 15:23, 15:29, 15:37, 15:47
B*15CYP ^f	15:02, 15:08, 15:11, 15:15, 15:21, 15:31, 15:44
B*15AND ^g	15:12, 15:14, 15:19
B*15I3 ^h	

^aB60 serologic equivalent cluster.

^bB61 serologic equivalent cluster.

^cB62 serologic equivalent cluster.

^dB63 serologic equivalent cluster.

^eB70 serologic equivalent cluster.

^fB75 serologic equivalent cluster.

^gB76 serologic equivalent cluster.

^hB77 serologic equivalent cluster.

³The B*08:26, 08:50 and 08:62 and the B*42:07 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*13:21 and 13:35 and the B*44:135 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*14:08 and the B*39:25N, 39:30, 39:32-39:34, 39:47 and 39:50 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*18:29 and the B*35:32:01-35:32:02, 35:37, 35:53N, 35:64, 35:66, 35:68:01-35:68:02, 35:99, 35:118-35:119 and 35:174 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*41:09 and the B*45:02 and B*45:03 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

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The B*54:01:02 and the B*55:01:07, 55:02:01-55:02:06, 55:07, 55:10, 55:12, 55:16, 55:19, 55:26, 55:30, 55:35, 55:37, 55:39, 55:41-55:43, 55:47-55:48 and 55:50 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*55:04, 55:08, 55:13, 55:27, 55:46 and 55:49 and the B*56:15, 56:19N and 56:22 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*55:23 and 55:32 and the B*56:18, 56:31-56:32 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*57:01:01-57:01:04, 57:01:06-57:01:12, 57:06, 57:08, 57:10, 57:15-57:16, 57:18-57:20, 57:22-57:23, 57:25-57:27, 57:29-57:30, 57:33-57:38, 57:40-57:41, 57:43-57:44, 57:48-57:50 and 57:52 and the B*58:36 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

SPECIFICITY TABLE**HLA-B low resolution SSP typing**

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}
1 ^{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, 07:140-07:143, 15:138, 15:230, 37:07, 40:15-40:16, 40:23, 40:32, 40:98, 40:136, 40:158, 41:08, 42:05:01-42:05:02, 48:05, 48:08, 48:15, C*02:23, C*04:77
2	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, 08:66-08:78, 15:142, 15:180, 44:49, 51:68
3	140 bp, 235 bp	1070 bp	7, 8, 13, 15, 35, 4005, 44, 49, 61, 62, 77	*07:20, 07:24, 07:60, 07:100, 07:131, 08:21, 08:25, 13:01:01-13:04, 13:06-13:08Q, 13:10-13:23, 13:25-13:38, 13:40-13:52, 15:07:01-15:07:02, 15:36 ^w , 15:45, 15:55, 15:68, 15:89 ^w , 15:126, 15:205, 15:207, 35:05:01-35:05:02, 35:16-35:17, 35:22, 35:30, 35:51, 35:58, 35:72, 35:89, 35:97, 35:113-35:114, 35:125, 35:164, 40:05, 40:71, 40:174, 44:08 ^w , 44:54, 44:57 ^w , 44:60, 44:106, 44:110, 44:135, 46:12, 46:20, 51:64, 53:14, 55:51, 58:18, C*03:05, C*03:25, C*03:27
4 ^{7,8}	130 bp, 265 bp	800 bp	12, 13, 14, 17, 21, 35, 40, 41, 44, 45, 47, 49, 50, 60, 61, 64, 65	*13:01:01-13:04, 13:06-13:13, 13:15-13:23, 13:25-13:52, 14:01:01-14:04, 14:07N, 14:09, 14:11-14:12, 14:14-14:28, 15:46, 15:53, 15:106, 15:212, 18:44, 35:46-35:47, 35:63, 35:154, 40:01:01-40:01:23, 40:07, 40:10:01-40:10:02, 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, 40:116-40:118N, 40:121, 40:123-40:126, 40:128-40:130, 40:132, 40:134-40:136, 40:139-40:141, 40:146-40:147, 40:150-40:156, 40:158, 40:160, 40:163-40:164, 40:166, 40:168, 40:171-40:172, 40:175, 41:01-41:19, 44:02:01:01-44:03:06, 44:03:08-44:05:03, 44:09-44:39, 44:41:01-44:43:02, 44:45-44:56N, 44:58N-44:80, 44:82, 44:84-44:102, 44:104-44:110, 44:112-44:129, 44:132-44:133, 44:135-44:137, 45:01-45:13, 47:01:01:01-47:08, 49:01:01-49:02, 49:04-49:20, 50:01:01-50:02, 50:04-50:06, 50:08-50:15, 57:45, 57:51, C*01:30

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5⁸	185 bp, 235 bp	800 bp	7, 16, 17, 27, 2708, 37, 38, 3901, 3902, 39, 64, 67	*14:01:01-14:01:02, 14:07N-14:08, 14:10, 14:12, 14:14, 14:19, 14:26, 27:01-27:05:15, 27:05:17-27:06, 27:08-27:10, 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, 27:82-27:86, 37:02, 38:01:01-38:02:02, 38:03, 38:07-38:24, 38:26-38:32, 38:34N-38:37, 39:01:01:01- 39:01:01:02L, 39:01:03-39:01:08, 39:01:10-39:02:01, 39:03, 39:05:01-39:09, 39:11, 39:14-39:15, 39:18, 39:19:02, 39:22, 39:24:01-39:44, 39:46-39:48, 39:50-39:62, 39:64-39:68, 47:04-47:05, 48:21, 48:26, 67:01:01, 67:03
6	190 bp	800 bp	14, 35, 38, 39, 65	*07:28, 14:02:01-14:02:02, 14:02:04-14:02:05, 14:03-14:06:02, 14:09, 14:11, 14:13, 14:15-14:18, 14:20, 14:22-14:23, 14:25, 14:27, 15:77, 15:189, 15:233, 35:26, 38:05, 38:33, 39:04, 44:16, 44:37, 44:64:01-44:64:02, 44:91, 44:132, 52:26, 57:04, A*23:31, A*24:106, C*16:10
7	290 bp	1070 bp	15, 22, 62, 63, 71, 72, 75, 76, 77, 46	*15:01:01:01-15:01:03, 15:01:06-15:02:05, 15:03:03- 15:08, 15:10:02-15:11:03, 15:11:05-15:15, 15:17:01:01-15:17:02, 15:19-15:21, 15:24-15:28, 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, 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, 46:01:01-46:30
8	165 bp, 220 bp, 330 bp	1070 bp	5, 8, 12, 21, 22, 37, 41, 42, 44, 45, 48, 51, 56, 57, 60, 62, 70, 71, 72, 82	*08:01:01-08:05, 08:08N-08:12:03, 08:15-08:19N, 08:21-08:24, 08:26-08:27, 08:29-08:36, 08:38-08:39, 08:41-08:48, 08:50-08:54, 08:56-08:69, 08:71-08:73, 08:75-08:76, 08:78, 13:46, 15:03:01-15:03:03, 15:37, 15:46-15:47, 15:49, 15:51-15:54, 15:61- 15:62, 15:64, 15:69, 15:72, 15:74, 15:80, 15:83, 15:90-15:91, 15:93, 15:98-15:99, 15:103, 15:106, 15:108, 15:114-15:115, 15:119, 15:123-15:124, 15:127, 15:131-15:134, 15:143, 15:151, 15:153, 15:156, 15:158, 15:161, 15:173, 15:176, 15:186, 15:197-15:198, 15:200, 15:210, 15:212, 15:220- 15:221, 15:226N, 15:229, 15:235, 35:87, 37:12, 38:30, 40:12, 40:136, 41:01-41:03:02, 41:05-41:09,

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9^{8,9,11}	165 bp, 190 bp, 390 bp	1070 bp	5, 17, 21, 35, 51, 5102, 5103, 52, 56, 62, 63, 70, 78	*15:01:02, 15:09, 15:16:01-15:17:02, 15:67, 15:95, 15:162, 15:168, 15:177, 15:196, 15:208, 15:216, 15:222, 15:230, 35:01:10, 35:04:02, 40:26, 40:28, 44:62, 49:18, 50:14, 51:01:01-51:01:03, 51:01:05- 51:11N-51:12, 51:13:02-51:14, 51:16-51:23, 51:24:03-51:24:04, 51:26-51:33, 51:37-51:41N, 51:43-51:44N, 51:46, 51:48-51:61, 51:63-51:80, 51:82-51:91, 51:94- 51:98N, 51:100-51:126, 52:01:01:01-52:01:04, 52:01:06-52:13, 52:15-52:24, 52:26-52:27, 56:05:01- 56:06, 56:21, 58:08:01-58:08:02, 78:01:01-78:07
10	180 bp	800 bp	13, 22, 55, 60	*07:78, 13:01:01-13:02:12, 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, 40:48, 45:10, 49:07, 55:09, 55:22, 55:24
11⁶	105 bp, 195 bp	1070 bp	8, 12, 27, 38, 39, 3902, 40, 44, 45, 48, 60, 70, 71, 72, 82	*07:27, 07:50, 08:04, 08:17, 08:54, 15:03:01- 15:03:03, 15:47, 15:49, 15:54, 15:61-15:62, 15:64, 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, 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, 39:49, 40:12, 40:46, 40:93, 42:11, 44:10, 44:15, 44:18, 44:40, 44:44, 44:130, 45:01, 45:05-45:07, 45:11-45:13, 48:01:01-48:05, 48:07- 48:27, 49:20, 50:02, 52:16, 55:18, 82:01-82:03
12	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, 72, 75, 76, 77, 78	*07:09, 07:11, 07:17, 08:28, 08:35, 08:37, 13:04, 13:10, 13:26, 15:01:01-15:01:04, 15:01:06-15:08, 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, 15:49-15:57, 15:60-15:62, 15:64-15:72, 15:74-15:76, 15:78:01-15:82, 15:84-15:85, 15:87- 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, 15:197-15:207, 15:209N-15:215, 15:217-15:223, 15:225-15:228, 15:231-15:232, 15:234-15:236, 18:01:01-18:15, 18:17N-18:25, 18:27-18:40, 18:42-18:60, 18:62-18:68, 27:41, 35:01:01-35:01:29, 35:05:01-35:05:02, 35:07- 35:08:05, 35:10-35:11:02, 35:14:01-35:17, 35:19- 35:21, 35:23-35:30, 35:32:01-35:32:02, 35:35, 35:37,

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38	300 bp	1070 bp	7, 12, 13, 15, 17, 18, 21, 27, 2708, 35, 37, 38, 39, 3902, 40, 4005, 41, 44, 45, 47, 48, 49, 50, 52, 60, 61, 62, 70, 72, 76	44:129, 44:131, 45:01-45:13, 47:02, 47:03 ^w , 50:01:01-50:02, 50:04-50:05, 50:07-50:15
				*07:54, 07:123, 08:17, 08:38, 08:54, 13:01:01-13:04, 13:06, 13:08Q-13:23, 13:25-13:35, 13:37-13:52, 15:01:01:01-15:01:04, 15:01:06-15:01:18, 15:01:20- 2708, 15:01:24, 15:03:01-15:07:02, 15:12, 15:14, 15:19- 37, 38, 39, 15:20, 15:24-15:27:03, 15:30, 15:32-15:36, 3902, 15:38:01-15:40, 15:42-15:43, 15:45-15:50, 15:53- 4005, 15:54, 15:56-15:58, 15:60-15:63, 15:65-15:66, 44, 45, 47, 15:68-15:71, 15:73-15:75, 15:77-15:79N, 15:81- 48, 49, 50, 15:87, 15:91-15:92, 15:94N, 15:96-15:98, 15:101- 52, 60, 61, 15:107, 15:109-15:111N, 15:113, 15:116-15:118, 62, 70, 72, 15:122-15:123, 15:125-15:129, 15:131-15:132, 76 15:135-15:138, 15:140-15:142, 15:145-15:147, 15:150-15:152, 15:154-15:160, 15:163-15:167, 15:169, 15:171-15:175, 15:178-15:179, 15:181N- 15:185, 15:187-15:188, 15:190N, 15:192-15:193, 15:195, 15:199, 15:201-15:207, 15:210-15:212, 15:217-15:220, 15:224-15:225, 15:227-15:228, 15:231-15:236, 18:12, 27:01-27:21, 27:24-27:86, 35:10, 35:13, 35:16, 35:28, 35:63, 35:69, 35:80, 37:01:01-37:07, 37:09-37:31, 38:03-38:04, 38:25, 39:02:01-39:02:02, 39:08, 39:13:01-39:13:02, 39:22- 39:23, 39:49, 40:01:01-40:07, 40:09-40:16, 40:18- 40:24, 40:26-40:40, 40:42-40:67, 40:70-40:105, 40:107-40:165, 40:167-40:177, 41:01-41:19, 42:11, 44:02:01:01-44:05:03, 44:07-44:11, 44:13-44:137, 45:01-45:13, 47:01:01-47:08, 48:01:01-48:05, 48:07-48:15, 48:17-48:27, 49:01:01-49:20, 50:01:01- 50:02, 50:04-50:14, 52:01:01-52:27, 53:17:01- 53:17:02, 55:18, 55:34, 55:40, 78:05-78:06
39 ^{8,9}	160 bp, 330 bp	1070 bp	22, 44, 45, 47, 54	*44:02:01:01-44:02:21, 44:06, 44:08-44:09, 44:11- 44:12, 44:16, 44:19N-44:24, 44:27:01-44:27:02, 44:33-44:34, 44:41:01-44:41:02, 44:44, 44:48-44:49, 44:52N-44:53, 44:55, 44:59, 44:63, 44:66-44:68, 44:71-44:74, 44:80, 44:83-44:84, 44:86-44:87, 44:89-44:91, 44:93, 44:95, 44:97, 44:99-44:102, 44:104, 44:106, 44:112-44:113, 44:116, 44:118- 44:119, 44:121, 44:126-44:127, 44:131-44:132, 44:137, 54:01:01, 54:02-54:24, 83:01
40	180 bp, 210 bp	1070 bp	13, 15, 18, 35, 39, 40, 54, 55, 56, 59, 78	*07:65, 13:06, 15:42, 15:86, 15:224, 18:04, 35:08:05, 35:42:02, 35:60, 35:185, 39:17, 39:63, 40:01:11, 40:58, 45:08, 46:18, 54:01:01-54:02, 54:05N, 54:07- 54:08N, 54:10, 54:12-54:13, 54:16-54:24, 55:01:01- 55:03, 55:05, 55:07, 55:10-55:12, 55:15-55:16, 55:18-55:19, 55:21, 55:25-55:26, 55:29-55:31, 55:33-55:45, 55:47-55:48, 55:50, 55:52-55:54, 56:05:01, 56:06, 56:10, 56:23,

59:01:01:01-59:01:01:02, 59:05, 78:01:01-78:01:02,
78:02:02-78:03, 78:07, **C*15:02:04**

41	180 bp	1070 bp	12, 15, 21, 22, 44, 45, 49, 50, 51, 56, 61, 62, 82	*13:03, 13:48, 15:73, 40:71, 44:10, 44:15, 44:18, 45:01, 45:04-45:07, 45:11-45:13, 46:11, 49:01:01-49:03, 49:06, 49:08-49:17, 49:19N-49:20, 50:01:01-50:02, 50:04-50:08, 50:10-50:13, 50:15, 51:15, 51:62, 51:106, 52:25, 54:03, 56:01:01-56:02, 56:04, 56:07-56:08, 56:13-56:14, 56:16-56:17, 56:20:01-56:20:02, 56:24-56:30, 59:04, 82:01-82:03, C*03:12, C*03:19, C*03:102
42^{6,10}	90 bp, 240 bp	800 bp	57	*55:14, 57:01:01-57:44, 57:46-57:50, 57:52, 58:36
43^{6,13}	90 bp	1070 bp	5, 13, 15, 17, 39, 55, 56, 58, 60, 61, 62, 71	*13:11, 15:18:03, 15:73, 15:224, 39:17, 40:48, 40:71, 51:62, 52:25, 56:02, 56:04, 56:10, 58:01:01-58:02, 58:04-58:35, C*02:06^w, C*02:47^w
44⁶	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, 18:01:01-18:11, 18:13-18:15, 18:17N-18:36, 18:38-18:47, 18:49-18:65, 18:67-18:68, 35:01:01-01-35:08:04, 35:09:01-35:09:03, 35:11:01-35:12:03, 35:14:01-35:15, 35:17-35:18, 35:20:01-35:24:02, 35:27, 35:29-35:45, 35:48, 35:50-35:62, 35:64-35:68:02, 35:70-35:72, 35:74-35:75, 35:76 ^w , 35:77-35:79, 35:81-35:153, 35:155-35:185, 37:08, 38:06-38:07, 39:19:01-39:19:02, 44:06, 51:01:01-51:24:04, 51:26-51:46, 51:48-51:103, 51:105-51:111, 51:113-51:117, 51:119-51:126, 53:01:01-53:16, 53:18-53:27, 56:06 ^w , 78:01:01-78:04, 78:07
45^{6,8}	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, 15:01:01:01-15:01:04, 15:01:06-15:01:16, 15:01:18-15:01:24, 15:04, 15:07:01-15:08, 15:11:01-15:12, 15:14-15:15, 15:19, 15:24, 15:26N-15:28, 15:30, 15:32, 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, 18:19, 27:25, 27:75, 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, 39:18, 39:36, 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, 46:27- 46:30, 51:61, 52:21, 54:06, 55:21, 56:03, 73:01- 73:02, A*26:68, A*68:56, C*06:20, C*12:50
46¹⁴	360 bp	1070 bp	Bw4	
47	350 bp	1070 bp	Bw6	
48¹⁵	-	-		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-B low resolution SSP typings.

When the primers in a primer mix can give rise to specific PCR products of more than one length this is indicated if the size difference is 20 base pairs or more. Size differences shorter than 20 base pairs are not given. For high resolution SSP kits the respective lengths of the specific PCR product(s) of the alleles amplified by these primer mixes are given.

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 inherit 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 two different control primer pairs give rise to either an internal positive control band of 1070 base pairs, for most wells, or a band of 800 base pairs, for some wells.

Well number 1 contains the primer pair giving rise to the shorter, 800 bp, internal positive control band in order to help in the correct orientation of the HLA-B low resolution typing.

In addition, wells number 4 to 6, 10, 17, 20, 21, 30, 35 and 42 contain the primer pair giving rise to the shorter, 800 bp, internal positive control band in order to allow 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.

⁴Nucleotide sequence information is available for only exons 2 and 3 of many HLA Class I alleles and for only exon 2 of many HLA Class II alleles and not for other exons or for the introns of these alleles. We assume that unknown sequences in these exons and in the introns are conserved within loci and within allelic groups.

The B*08:26, 08:50 and 08:62 and the B*42:07 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*13:21 and 13:35 and the B*44:135 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*14:08 and the B*39:25N, 39:30, 39:32-39:34, 39:47 and 39:50 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*18:29 and the B*35:32:01-35:32:02, 35:37, 35:53N, 35:64, 35:66, 35:68:01-35:68:02, 35:99, 35:118-35:119 and 35:174 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*41:09 and the B*45:02 and B*45:03 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

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The B*54:01:02 and the B*55:01:07, 55:02:01-55:02:06, 55:07, 55:10, 55:12, 55:16, 55:19, 55:26, 55:30, 55:35, 55:37, 55:39, 55:41-55:43, 55:47-55:48 and 55:50 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*55:04, 55:08, 55:13, 55:27, 55:46 and 55:49 and the B*56:15, 56:19N and 56:22 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*55:23 and 55:32 and the B*56:18, 56:31-56:32 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

The B*57:01:01-57:01:04, 57:01:06-57:01:12, 57:06, 57:08, 57:10, 57:15-57:16, 57:18-57:20, 57:22-57:23, 57:25-57:27, 57:29-57:30, 57:33-57:38, 57:40-57:41, 57:43-57:44, 57:48-57:50 and 57:52 and the B*58:36 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

⁵Due to sharing of sequence motifs, primer mix 1 will amplify the C*02:23 and C*04:77 alleles, primer mix 3 will amplify the C*03:05, C*03:25 and C*03:27 alleles, primer mix 4 will amplify the C*01:30 allele, primer mix 6 will amplify the A*23:31, A*24:106 and C*16:10 alleles, the C*07:46 allele will be amplified by primer mix 8, primer mix 13 will amplify the A*24:174 allele, primer mix 17 will amplify the C*03:102 allele, primer mix 21 will amplify the C*15:51 allele, primer mix 25 will amplify the C*03:129 allele, primer mix 29 will amplify the C*15:25 allele, primer mix 31 will amplify the C*15:39 allele, primer mix 40 will amplify the C*15:02:04 allele, primer mix 41 will amplify the C*03:12, C*03:19 and C*03:102 alleles, primer mix 43 will weakly amplify the C*02:06 and C*02:47 alleles and primer mix 45 will amplify the A*26:68, A*68:56, C*06:20 and C*12:50 alleles.

⁶Specific PCR fragments shorter than 125 base pairs have a lower intensity and are less sharp than longer PCR bands.

⁷Primer mix 4 may yield less 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 4, 5, 9, 39 and 45 give a lower yield of specific PCR product than the other HLA-B low resolution primer mixes.

⁹Primer mixes 9, 20 and 39 may give rise to nonspecific amplifications, most pronounced in primer mix 9.

¹⁰Primer mixes 1, 15, 19, 20 and 42 may give rise to a primer oligomer artifact.

¹¹The B*57 and B*58 alleles might be faintly amplified by primer mix 9.

¹²The C*17:01 to C*17:04 alleles might be faintly amplified by primer mix 25.

¹³Primer mixes 32 and 43 may generate a false positive band of about 800 base pairs. This band should be disregarded when interpreting HLA-B low resolution typings.

¹⁴The Bw4-associated HLA-A specificities A9, A23, A24, A2403, A25 and A32 are not amplified by the primer pair in primer mix 46.

¹⁵Primer mix 48 contains a negative control, which will amplify more than 95% of HLA amplicons as well as the amplicons generated by control primer pairs. PCR product sizes range from 75 to 200 base pairs. The PCR product generated by the control primer pair is 430 base pairs.

'w', might be weakly amplified.

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

INTERPRETATION TABLE

HLA-B low resolution SSP

Amplification patterns of the B*07:02 to B*83:01 alleles¹⁵

		Well																							
		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24
Length of spec.		110																							
PCR product(s)		215																							
Length of int. pos. control ¹		235																							
5'-primer(s) ²		235	140																						
3'-primer(s) ³		265	130																						
Well No.	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	
	5' -gCT 3' 605	5' -gAC 3' 540	5' -TgA 3' 527	800																					
	5' -TgA 3' 272	5' -Tgg 3' 272	5' -TCg 3' 97	1070																					
	5' -CAG 3' 559	5' -gTg 3' 309	5' -AgC 3' 363	363	5' -AgC 3' 209	209																			
	5' -gTg 3' 583	5' -CgT 3' 193	5' -CgT 3' 193	361	5' -CCg 3' 103	103	5' -CCg 3' 103	800																	
	5' -CTC 3' 559	5' -TAT 3' 246	5' -TAT 3' 246	363	5' -CCT 3' 103	103	5' -CCT 3' 103	800																	
	5' -gCC 3' 572	5' -TAT 3' 246	5' -Agg 3' 418	418	5' -CCg 3' 103	103	5' -CCg 3' 103	800																	
	5' -gTC 3' 538	5' -CCT 3' 206	5' -ATA 3' 412	357	5' -gGA 3' 45	45	5' -gGA 3' 45	1070																	
	5' -gTg 3' 583	5' -CAG 3' 559	5' -TCC 3' 266	435	5' -gGA 3' 45	45	5' -gGA 3' 45	1070	390	190	165														
	5' -gCg 3' 572	5' -Tgg 3' 272	5' -TTA 3' 420	420	5' -AgA 3' 206	206	5' -AgA 3' 206	1070	195	105															
	5' -gCg 3' 572	5' -TAT 3' 3 rd l	5' -gTC 3' 419	419	5' -TCT 3' 142	142	5' -TCT 3' 142	1070																	
	5' -gCg 3' 572	5' -Tgg 3' 272	5' -TCT 3' 3 rd l	420	5' -gTC 3' 419	419	5' -gTC 3' 419	1070																	
	5' -gCg 3' 572	5' -Tgg 3' 272	5' -Tgg 3' 272	301	5' -gTC 3' 301	301	5' -gTC 3' 301	301	5' -gTC 3' 301	301	5' -gTC 3' 301	301	5' -gTC 3' 301	301	5' -gTC 3' 301	301	5' -gTC 3' 301	301	5' -gTC 3' 301	301	5' -gTC 3' 301	301	5' -gTC 3' 301	301	
	5' -gCT 3' 605	5' -ATC 3' 309	5' -Tgc 3' 272	540	5' -gAC 3' 540	540	5' -gAC 3' 540	540	5' -gAC 3' 540	540	5' -gAC 3' 540	540	5' -gAC 3' 540	540	5' -gAC 3' 540	540	5' -gAC 3' 540	540	5' -gAC 3' 540	540	5' -gAC 3' 540	540	5' -gAC 3' 540	540	
	5' -Tgg 3' 272	5' -Tgg 3' 272	5' -TCT 3' 234	363	5' -Cgg 3' 161	161	5' -Cgg 3' 161	1070	150	115															
	5' -gCT 3' 204	5' -TCT 3' 204	5' -gCT 3' 167	167	5' -gCT 3' 167	167	5' -gCT 3' 167	1070																	
	5' -gCT 3' 463	5' -TCA 3' 355	5' -TCA 3' 355	355	5' -TCA 3' 355	355	5' -TCA 3' 355	800																	
	5' -ggC 3' 302	5' -gAC 3' 206	5' -gAC 3' 206	206	5' -gAC 3' 206	206	5' -gAC 3' 206	1070																	
	5' -gTC 3' 538	5' -ggg 3' 311	5' -gTT 3' 557	557	5' -gTT 3' 557	557	5' -gTT 3' 557	368	5' -gGA 3' 45	45															
	5' -gCg 3' 572	5' -gTA 3' 292	5' -Tgg 3' 272	266	5' -gTT 3' 259	259	5' -gTT 3' 259	368	5' -ggC 3' 409	409															
	5' -gCg 3' 572	5' -gTA 3' 292	5' -Tgg 3' 272	317	5' -ggC 3' 44	44	5' -ggC 3' 44	1070	110	110															
	5' -gCg 3' 477	5' -ggT 3' 302	5' -ggC 3' 302	357	5' -Tgg 3' 357	357	5' -Tgg 3' 357	800	170	170															

INTERPRETATION TABLE

HLA-B low resolution SSP

Amplification patterns of the B*07:02 to B*83:01 alleles¹⁵

		Well																										Length of spec. PCR product(s)												
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	Length of int. pos. control ¹				5'-primer(s) ²												
5' -TgT 3'	193	5' -Cgt 3'	193	5' -Agg 3'	499	5' -TCT 3'	499	5' -Agg 3'	418	5' -TCA 3'	355	5' -TCA 3'	355	5' -gCT 3'	72	1070	260	225	195	115	180	290	105	105	105	105	105	105	105	105	105	105	105							
5' -gCT 3'	605	5' -gTg 3'	603	5' -GCC 3'	282	5' -ATC 3'	309	5' -TCA 3'	355	5' -ggg 3'	209	5' -ggC 3'	209	5' -ATT 3'	141	800	215	215	215	215	215	215	215	215	215	215	215	215	215	215	215	215	215	215	215					
5' -gCg 3'	572	5' -gTC 3'	538	5' -gCC 3'	282	5' -ggA 3'	499	5' -gTT 3'	259	5' -AgC 3'	363	5' -ggC 3'	209	5' -CAG 3'	1st I	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070						
5' -gTC 3'	538	5' -ATC 3'	309	5' -gTC 3'	299	5' -CTC 3'	299	5' -TCA 3'	355	5' -ggg 3'	209	5' -ggC 3'	209	5' -Tgg 3'	357	141	800	270	130	410	90	210	120	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	
5' -gTg 3'	583	5' -TCA 3'	299	5' -gTC 3'	418	5' -gCC 3'	282	5' -ggA 3'	499	5' -gTT 3'	259	5' -ggC 3'	209	5' -ggA 3'	206	5' -gCC 3'	48	1070	430	145	210	90	210	120	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48
5' -CCT 3'	527	5' -TCT 3'	435	5' -ATC 3'	309	5' -ggA 3'	499	5' -gTT 3'	259	5' -AgC 3'	363	5' -ggC 3'	209	5' -Tgg 3'	357	800	1070	175	90	210	120	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48			
5' -Tgc 3'	262	5' -CTC 3'	259	5' -Ctg 3'	259	5' -ggT 3'	311	5' -ggA 3'	362	5' -ggT 3'	362	5' -ggA 3'	209	5' -ggC 3'	209	5' -gAA 3'	206	1070	145	145	145	145	145	145	145	145	145	145	145	145	145	145	145	145	145	145	145			
5' -Cgt 3'	559	5' -CTT 3'	259	5' -ggt 3'	256	5' -CAG 3'	559	5' -ggT 3'	311	5' -ggA 3'	362	5' -ggA 3'	209	5' -ggC 3'	209	5' -CAG 3'	1st I	1070	300	300	300	300	300	300	300	300	300	300	300	300	300	300	300	300	300	300	300			
5' -gCT 3'	420	5' -CTC 3'	256	5' -ggT 3'	256	5' -ggT 3'	259	5' -ggT 3'	272	5' -ggA 3'	317	5' -ggT 3'	311	5' -TAC 3'	369	5' -ggA 3'	209	1070	210	180	210	180	210	180	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48
Negative Control																																			Well No.					
Neg. Control																																			Well No.					

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Lot-specific information

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Well No.		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24
HLA-B allele ⁴	ser. ⁵																								
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*35:60	B35																			18						
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*35:163	-																		17	18						
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28					32														w					47	B35, 22	*	35:76	
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				29															44					47	-	*	35:115	
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	29																	44					46	B37	*	37:08		
	29													38										46	B37, -	*	37:10, 37:22, 37:29	
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*39:01:01:01- 39:01:01:02L, 39:01:03- 39:01:08, 39:01:10- 39:01:12, 39:03, 39:05:01- 39:06:02, 39:09, 39:11, 39:14-39:15, 39:24:01, 39:26-39:29, 39:31, 39:35, 39:37-39:38Q, 39:40N-39:41, 39:44, 39:46, 39:48, 39:51- 39:57, 39:59-39:62, 39:64- 39:68	B3901, B39, Low B39, Null, -					5															21	23	24		
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*39:02:01, 39:08	B3902, B39					5				11											21		24		
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*39:04	B39					6															21	23	24		
*39:07	B16					5				12											21	23	24		
*39:10:01-39:10:02, 39:16, 39:20	B39, -																				21		24		
*39:17	B39																				21		24		
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*39:19:01	B39																		18		21	23	24		
*39:19:02	B39					5													18		21	23	24		
*39:22	B39					5														21	23	24			
*39:24:02	-					5															21	23			
*39:36	-					5																23	24		
*39:39	B39					5				11											21		24		
*39:42	B39					5													17		21	23	24		
*39:43	-					5				12												23	24		
*39:58	-					5															21		24		
*39:63	-																				21		24		
*40:01:01-40:01:06, 40:01:08-40:01:10, 40:01:12-40:01:17, 40:01:19-40:01:23, 40:07, 40:51, 40:66, 40:73, 40:79, 40:124:01, 40:139- 40:140, 40:146-40:147, 40:150, 40:152-40:153, 40:155N	B60, Null, -					4																		24	
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*40:01:11	-				4																				24
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26												37	38									47		B61, B41, Null, -	*40:02:01-40:02:12, 40:04, 40:39, 40:89, 40:94, 40:97, 40:122, 40:142N-40:144N	
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*40:21, 40:137	B40, B15, -																								?	
*40:23	B40	1		4																					24	
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*40:40	B61																									
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*40:46	B60										11				16										?	
*40:47	B40			4																						
*40:48	B60			4						10															?	
*40:53, 40:72:02	B61, B60			4																						24
*40:58	B40			4																						?
*40:59-40:60	B40, -			4								12														24
*40:68	-																									?
*40:71	B61		3																							?
*40:74, 40:149	B60, -																									?
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*40:95	-																									?
*40:96	-																									24
*40:98	-	1																								?
*40:105	-												12													?
*40:109	-																									24
*40:110	-			4																						24
*40:117	-			4																						
*40:132	-			4																						?
*40:135	-			4																						
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*40:158	-	1	4												12											?
*40:159	-												12													24
*40:166	-			4																						?
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*41:01, 41:06-41:07, 41:12, 41:14, 41:16-41:17	B41, -			4				8																		24
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	?					31						37 38										47		-		*40:28		
	?							33				37 38										47		B60, -		*40:31, 40:45, 40:80		
25	?											37 38										47		B60		*40:38, 40:52		
												37 38										47		B61		*40:40		
													37 38															*40:44, 40:75, 40:86, 40:103, 40:127, 40:131, 40:162, 40:165, 40:167
25												38										47		B61, -		B60		*40:46
25	?											38										46		B40		*40:47		
25	?											37 38				43						47		B60		*40:48		
	?											37 38										47		B61,		*40:53, 40:72:02		
	?											37 38	40									47		B60		*40:58		
25												37 38										47		B40, -		*40:59-40:60		
	?											37										47		-		*40:68		
	?											37 38		41	43						47		B61		*40:71			
25	?											37 38										47		B60, -		*40:74, 40:149		
25	?												38										47		B60		*40:76	
	?											37 38										47		-		*40:83, 40:177		
	?												38										47		-		*40:93	
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	?												38										46		-		*40:96	
	?											37 38										47		-		*40:98		
	?											37 38										47		-		*40:105		
	?											34	36	38								46		-		*40:109		
	?												38										46		-		*40:110	
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	?	27										37	38									47		-		*40:136		
25												37	38									47		-		*40:138		
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	?												38										46		-		*40:157	
	?											37	38									47		-		*40:158		
	?											37	38									47		-		*40:159		
25	?			28								37										47		-		*40:166		
25												37	38									47		-		*40:170		
	?											37	38									47		-		*40:174		
	27			30								37	38									47		B41, -		*41:01, 41:06-41:07, 41:12, 41:14, 41:16-41:17		
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*41:02:01-41:03:02, 41:11, 41:13, 41:15, 41:19	B41, -				4				8																24
*41:04, 41:10, 41:18	B41, -				4																				24
*41:05	B41				4				8																24
*41:08	B41	1			4				8								14								24
*41:09, 45:02-45:03 ¹⁰	B45, -				4				8																24
*42:01:01-42:02, 42:06, 42:10, 42:12, 42:14-42:15	B42, -								8																24
*42:04	B42								8																24
*42:05:01-42:05:02	B42	1							8							14									24
*42:08	B42								8																24
*42:09	B42														12										24
*42:11	-								8			11													24
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*44:03:07, 44:07, 44:81, 44:103, 44:111, 44:134	B44, -																								17
*44:06	B44																								17
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Well No.		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24

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		27										37	38											47	B41, -	*41:02:01-41:03:02, 41:11, 41:13, 41:15, 41:19	
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		27	28																						47	B42, -	*42:01:01-42:02, 42:06, 42:10, 42:12, 42:14-42:15
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			28																						47	B42	*42:08
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												31		36						44		46				B53	*53:02	
														36						44		46				B53, B35	*53:03, 53:09, 53:11- 53:13	
												30	31		36					44		46				B53, –	*53:04, 53:07, 53:19	
														36						44		46				B53, B51	*53:06	
														36	38					46						–	*53:14	
												29		36						44		46				–	*53:17:01-53:17:02	
															36												–	*53:22
												28		32			39	40								B54, Null, –	*54:01:01, 54:02, 54:05N, 54:07-54:08N, 54:10, 54:13, 54:16-54:19, 54:21- 54:24	
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48		Well No.			

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Well No.		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24
*54:01:02, 55:01:07, 55:02:01-55:02:06, 55:07, 55:10, 55:12, 55:16, 55:19, 55:26, 55:30, 55:35, 55:37, 55:39, 55:41-55:43, 55:47-55:48, 55:50 ¹¹	B54, B55, –																								
*54:03	B22																								
*54:04, 54:15	B54, –																								
*54:06	–												12												
*54:09, 54:14	–												12												
*54:11	B54																							24	
*54:12	–																								
*54:20	–																								
*55:01:01-55:01:06, 55:01:08, 55:03, 55:05, 55:11, 55:15, 55:25, 55:29, 55:31, 55:33, 55:36, 55:38, 55:44- 55:45, 55:52-55:54	B55, –																								
*55:04, 55:08, 55:13, 55:27, 55:46, 55:49, 56:15, 56:19N, 56:22 ¹²	B55, B56, Null, –																								
*55:09, 55:24	B22, B55												10												
*55:14	–												12										17		
*55:17, 55:28	B55, –																								
*55:18	–												11												
*55:20	B22												8												
*55:21	B55																								
*55:22	B22												10												
*55:23, 55:32, 56:18, 56:31-56:32 ¹³	B55, –												12												
*55:34	–																								
*55:40	–																								
*55:51	–		3																						
*56:01:01-56:01:04, 56:08, 56:14, 56:16, 56:20:01, 56:24, 56:26- 56:30	B56, Null, –																								
*56:02, 56:04	B56																								
*56:03	B22, 15														12										
*56:05:01	B56												9												
*56:05:02	B56												9												
*56:06	B78												9									18			
*56:07	B56																								
Well No.		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24

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25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48		Well No.
			28		32								40								47				
			28		32								39		41							47		B22	*54:03
														39								47		B54, -	*54:04, 54:15
			28		32								39								45	47		-	*54:06
															39							47		-	*54:09, 54:14
			28		32									39								47		B54	*54:11
					32	34								39	40						46		-	*54:12	
			28		32	35								39	40							47		-	*54:20
			28		32	35							40									47		B55, -	*55:01:01-55:01:06, 55:01:08, 55:03, 55:05, 55:11, 55:15, 55:25, 55:29, 55:31, 55:33, 55:36, 55:38, 55:44- 55:45, 55:52-55:54
			28		32																	47		B55, B56, Null, -	*55:04, 55:08, 55:13, 55:27, 55:46, 55:49, 56:15, 56:19N, 56:22 ¹²
					32	35																47		B22, B55	*55:09, 55:24
			28		32																	47		-	*55:14
					32	35																47		B55, -	*55:17, 55:28
			28		32											38	40					47		-	*55:18
				27	28	30	32														47		B22	*55:20	
				28		32	w							40							45	47		B55	*55:21
			28		32																47		B22	*55:22	
				28		32																47		B55, -	*55:23, 55:32, 56:18, 56:31-56:32 ¹³
				28		32										38	40					47		-	*55:34
				28		32	35							38	40						47		-	*55:40	
				28		32																47		-	*55:51
			28		32																	47		B56, Null, -	*56:01:01-56:01:04, 56:08, 56:14, 56:16, 56:20:01, 56:24, 56:26- 56:30
				28		32																47		B56	*56:02, 56:04
			28		32																	47		B22, 15	*56:03
				28		32	35							40							45	47		B56	*56:05:01
			28		32	35															47		B56	*56:05:02	
				28		32	35							40			w				47		B78	*56:06	
				28		32												41				46		B56	*56:07
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48		Well No.

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Well No.		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	
*56:09	B56												12					17								
*56:10	B55																									
*56:11-56:12	B56, B22																						17			
*56:13	B56									8																
*56:17	B56																									
*56:20:02	-																									24
*56:21	-												9													
*56:23	-																									
*56:25	-																									
*57:01:01-57:01:04, 57:01:06-57:01:12, 57:06, 57:08, 57:10, 57:15- 57:16, 57:18-57:20, 57:22- 57:23, 57:25-57:27, 57:29- 57:30, 57:33-57:38, 57:40- 57:41, 57:43-57:44, 57:48- 57:50, 57:52, 58:36 ¹⁴	B57, -													12					17							
*57:01:05, 57:11, 57:13, 57:21, 57:31, 57:47	B57, -													12												
*57:02:01-57:03:02, 57:05, 57:17, 57:28N, 57:32, 57:39, 57:42, 57:46	B57, Null, -																			17						
*57:04	B57							6												17						
*57:07	B57																			17						
*57:09	B57								8											17						
*57:12	B57																			17						
*57:14	-													12						17						
*57:24	-													12						17						
*57:45, 57:51	-					4								12						17						
*58:01:01-58:01:02, 58:01:04-58:01:09, 58:04- 58:05, 58:10N-58:15, 58:19, 58:21-58:24, 58:29- 58:33, 58:35	B58, Null, -													12					17							
*58:01:03, 58:02, 58:06, 58:16, 58:25-58:26	B58, -													12												
*58:07	B58													12	13											
*58:08:01	B17, B5											9														
*58:08:02	-											9														
*58:09	B17													12					17							
*58:17N	Null													12					17							
*58:18	-				3									12												
*58:20	-																			21						
Well No.		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	

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25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48				Well No.	
			28		32																		47			B56	*56:09	
			28		32										40		43						47			B55	*56:10	
			28		32																		47			B56, B22	*56:11-56:12	
			28	30	32										41								47			B56	*56:13	
															41								47			B56	*56:17	
			28		32										41								47			-	*56:20:02	
			28		32										41								47			-	*56:21	
					32	34	35								40							46			-	*56:23		
			28		32		35								41							47			-	*56:25		
															36													
																42								46			B57, -	*57:01:01-57:01:04, 57:01:06-57:01:12, 57:06, 57:08, 57:10, 57:15- 57:16, 57:18-57:20, 57:22 57:23, 57:25-57:27, 57:29 57:30, 57:33-57:38, 57:40 57:41, 57:43-57:44, 57:48 57:50, 57:52, 58:36 ¹⁴
															36		42						46			B57, -	*57:01:05, 57:11, 57:13, 57:21, 57:31, 57:47	
															36		42						46			B57, Null, -	*57:02:01-57:03:02, 57:05, 57:17, 57:28N, 57:32, 57:39, 57:42, 57:46	
															36		42						46			B57	*57:04	
			29												36		42						46			B57	*57:07	
			29												36		42						46			B57	*57:09	
																42								47			B57	*57:12
					31										36		42						46			-	*57:14	
					29										36		42						46			-	*57:24	
							34	36															46			-	*57:45, 57:51	
																36												
																36		43					46			B58, Null, -	*58:01:01-58:01:02, 58:01:04-58:01:09, 58:04- 58:05, 58:10N-58:15, 58:19, 58:21-58:24, 58:29 58:33, 58:35	
																36		43					46			B58, -	*58:01:03, 58:02, 58:06, 58:16, 58:25-58:26	
																36		43					46			B58	*58:07	
																35	36						46			B17, B5	*58:08:01	
																35	36						46			-	*58:08:02	
																36		43					46			B17	*58:09	
																36		43					46			Null	*58:17N	
																36		43					46			-	*58:18	
																36		43					46			-	*58:20	
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48				Well No.	

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Well No.		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	
*58:27	-																									
*58:28, 58:34	-																	17								
*59:01:01:01-59:01:01:02, 59:05	B59, -																									
*59:02-59:03	B59, -																									
*59:04	-																									
*67:01:01, 67:03	B67, -			5																21		24				
*67:01:02-67:02	B67, -																			21		24				
*73:01-73:02	B73, -																						24			
*78:01:01-78:01:02, 78:02:02-78:03, 78:07	B78, -								9									18								
*78:02:01, 78:04	B78, B35								9								18									
*78:05	-								9								18									
*78:06	-								9																	
*81:01	B81																									
*81:02	B81																						24			
*81:03-81:05	Null, -																									
*82:01-82:03	B82, -					8			11									19								
*83:01	-																17									
A*23:31, A*24:106, C*16:10						6																				
A*24:174																13										
A*26:68, A*68:56, C*06:20, C*12:50																										
C*01:30					4																					
C*02:06, C*02:47																										
C*02:23, C*04:77		1																								
C*03:05, C*03:25, C*03:27			3																							
C*03:12, C*03:19																										
C*03:102																		17								
C*03:129																										
C*07:46					8																					
C*15:02:04																										
C*15:25																										
C*15:39																										
C*15:51																				21						
Well No.		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	

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25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48		Well No.	
								36							43									-	*58:27	
								36							43									-	*58:28, 58:34	
								34				40												B59, -	*59:01:01:01-59:01:01:02, 59:05	
								34																	B59, -	*59:02-59:03
								34							41									-	*59:04	
	28																								B67, -	*67:01:01, 67:03
	28																								B67, -	*67:01:02-67:02
	28																								B73, -	*73:01-73:02
								35				40				44								B78, -	*78:01:01-78:01:02, 78:02:02-78:03, 78:07	
								35								44								B78, B35	*78:02:01, 78:04	
								35				38												-	*78:05	
								35				38												-	*78:06	
	28		32	33																				B81	*81:01	
	28			33																				B81	*81:02	
	28			33																				Null, -	*81:03-81:05	
	28		32												41									B82, -	*82:01-82:03	
	28	29	32									39												-	*83:01	
																										A *23:31, A *24:106, C *16:10 A *24:174
																										A *26:68, A *68:56, C *06:20, C *12:50 C *01:30
																										C *02:06, C *02:47 C *02:23, C *04:77
																										C *03:05, C *03:25, C *03:27
																										C *03:12, C *03:19
																										C *03:102
																										C *03:129
																										C *07:46
																										C *15:02:04
				29																						C *15:25
							31																			C *15:39
																										C *15:51
25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48		Well No.	

¹The internal positive control primer pairs amplify segments of the human growth hormone gene. The two different control primer pairs give rise to either an internal positive control band of 1070 base pairs, for most wells, or a band of 800 base pairs, for some wells.

Well number 1 contains the primer pair giving rise to the shorter, 800 bp, internal positive control band in order to help in the correct orientation of the HLA-B low resolution typing.

In addition, wells number 4 to 6, 10, 17, 20, 21, 30, 35 and 42 contain the primer pair giving rise to the shorter, 800 bp, internal positive control band in order to allow kit identification.

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²The nucleotide position, in the 1st, 2nd or 3rd exon or in the 1st intron, matching the specificity-determining 3'-end of the primer is given. Nucleotide numbering as on the www.ebi.ac.uk, [imgt](http://imgt.hc.cnr.it), [hla](http://hla.helsinki.fi) web site. The sequence of the 3 terminal nucleotides of the primer is given.

³The nucleotide position, in the 2nd, or 3rd exon or the 1st or 3rd introns, 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](http://imgt.hc.cnr.it), [hla](http://hla.helsinki.fi) web site. The sequence of the 3 terminal nucleotides of the primer is given.

⁴The sequence of the B*0701 allele has been shown to be in error.

The sequence of the B*08:06 allele has been shown to be identical to B*08:20.

The sequence of the B*1305 allele has been shown to be identical to B*13:04.

The B*1324 allele has been renamed B*13:22:02

The B*150105 allele has been corrected and renamed B*15:120

The B*1522 allele has been renamed B*35:43.

The sequence of the B*1541 allele has been shown to be identical to B*15:39.

The B*1559 allele has been renamed B*35:44.

The B*15:100 allele has never been assigned.

The sequence of the B*1816 allele has been shown to be identical to B*18:14.

The sequence of the B*27051 allele has been shown to be identical to B*27:05:02.

The sequence of the B*2722 allele has been shown to be identical to the corrected B*27:06 sequence.

The B*35:43:02 allele has been renamed B*35:185.

The B*3573 allele has been renamed B*35:08:03.

The sequence of the B*39012 allele has been shown to be identical to B*39:01:01:01.

The sequence of the B*3921 allele has been shown to be identical to B*39:24.

The sequence of the B*4017 allele has been shown to be identical to B*40:16.

The sequence of the B*4041 allele has been shown to be identical to B*40:40.

The sequence of the B*4203 allele has never been assigned.

The sequence of the B*4401 allele has been shown to be identical to B*44:02:01:01.

The sequence of the B*5003 allele has been shown to be identical to B*50:02.

The sequence of the B*5125 allele has been shown to be identical to B*51:22.

The B*5147 allele has been renamed B*51:09:02.

The sequence of the B*5506 allele has been shown to be identical to B*55:04.

The sequence of the B*5803 allele has never been assigned.

The B*7901 allele has been renamed B*15:18:01.

The B*9530 allele has been renamed B*15:27:02.

⁵The 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.

⁶The B*08:26, 08:50 and 08:62 and the B*42:07 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

⁷The B*13:21 and 13:35 and the B*44:135 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

⁸The B*14:08 and the B*39:25N, 39:30, 39:32-39:34, 39:47 and 39:50 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

⁹The B*18:29 and the B*35:32:01-35:32:02, 35:37, 35:53N, 35:64, 35:66, 35:68:01-35:68:02, 35:99, 35:118-35:119 and 35:174 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

¹⁰The B*41:09 and the B*45:02 and B*45:03 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

¹¹The B*54:01:02 and the B*55:01:07, 55:02:01-55:02:06, 55:07, 55:10, 55:12, 55:16, 55:19, 55:26, 55:30, 55:35, 55:37, 55:39, 55:41-55:43, 55:47-55:48 and 55:50 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

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¹²The B*55:04, 55:08, 55:13, 55:27, 55:46 and 55:49 and the B*56:15, 56:19N and 56:22 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

¹³The B*55:23 and 55:32 and the B*56:18, 56:31-56:32 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

¹⁴The B*57:01:01-57:01:04, 57:01:06-57:01:12, 57:06, 57:08, 57:10, 57:15-57:16, 57:18-57:20, 57:22-57:23, 57:25-57:27, 57:29-57:30, 57:33-57:38, 57:40-57:41, 57:43-57:44, 57:48-57:50 and 57:52 and the B*58:36 alleles give rise to identical amplification patterns with the HLA-B low resolution primer set. These alleles can be separated by the respective high resolution primer sets.

¹⁵Primer mix 48 contains a negative control, which will amplify more than 95% of HLA amplicons as well as the amplicons generated by control primer pairs. PCR product sizes range from 75 to 200 base pairs. The PCR product generated by the control primer pair is 430 base pairs.

'ser', serological HLA specificity.

'w', might be weakly amplified.

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

CELL LINE VALIDATION SHEET																	
HLA-B low resolution SSP kit																	
																Well	
																1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16	
																	Prod. No.:
																	201190001 201190002 201190003 201190004 201190005 201190006 201190007 201190008 201190009 201190010 201190011 201190012 201190013 201190014 201190015 201190016
IHWC cell line	B*																
1 9001	SA	*07:02		+	-	-	-	-	-	-	-	-	-	-	-	-	
2 9280	LK707	*52:01	*73:01	-	-	-	-	-	-	-	+	-	-	-	-	-	
3 9011	E4181324	*52:01		-	-	-	-	-	-	-	+	-	-	-	-	-	
4 9275	GU373	*15:10	*53:01	-	-	-	-	-	-	+	-	-	-	+	-	-	
5 9009	KAS011	*37:01		-	-	-	-	-	-	-	-	-	-	+	-	-	
6 9353	SM	*39:01	*51:01	-	-	-	-	+	-	-	+	-	-	-	-	-	
7 9020	QBL	*18:01		-	-	-	-	-	-	-	-	-	+	-	-	+	
8 9025	DEU	*35:01		-	-	-	-	-	-	-	-	-	+	-	-	-	
9 9026	YAR	*38:01		-	-	-	+	-	-	-	-	-	-	-	-	-	
10 9107	LKT3	*54:01		-	-	-	-	-	-	-	-	-	-	-	-	-	
11 9051	PITOUT	*44:03		-	-	-	+	-	-	-	-	-	-	-	-	-	
12 9052	DBB	*57:01		-	-	-	-	-	-	-	-	-	+	-	-	-	
13 9004	JESTHOM	*27:05		-	-	-	-	+	-	-	-	-	-	-	-	+	
14 9071	OLGA	*15:01	*15:20	-	-	-	-	-	-	+	-	-	-	+	-	-	
15 9075	DKB	*40:01		-	-	-	+	-	-	-	-	-	-	-	-	-	
16 9037	SWEIG007	*40:02		-	-	-	-	-	-	-	-	-	-	-	-	-	
17 9282	CTM3953540	*08:01	*55:01	-	+	-	-	-	-	+	-	-	-	-	-	-	
18 9257	32367	*14:01	*56:01	-	-	-	+	+	-	-	-	-	-	-	-	-	
19 9038	BM16	*18:01		-	-	-	-	-	-	-	-	-	+	-	-	+	
20 9059	SLE005	*40:01		-	-	-	+	-	-	-	-	-	-	-	-	-	
21 9064	AMALA	*15:01		-	-	-	-	-	-	+	-	-	-	+	-	-	
22 9056	KOSE	*35:03		-	-	-	-	-	-	-	-	-	-	-	-	-	
23 9124	IHL	*40:02	*56:02	-	-	-	-	-	-	-	-	-	-	-	-	-	
24 9035	JBUSH	*38:01		-	-	-	-	+	-	-	-	-	-	-	-	-	
25 9049	IBW9	*14:02		-	-	-	+	-	+	-	-	-	-	-	-	-	
26 9285	WT49	*58:01		-	-	-	-	-	-	-	-	-	+	-	-	-	
27 9191	CH1007	*07:05	*51:01	+	-	-	-	-	-	-	+	-	-	-	-	-	
28 9320	BEL5GB	*44:02	*44:03	-	-	-	+	-	-	-	-	-	-	-	-	-	
29 9050	MOU	*44:03		-	-	-	+	-	-	-	-	-	-	-	-	-	
30 9021	RSH	*42:01		-	-	-	-	-	-	+	-	-	-	-	-	-	
31 9019	DUCAF	*18:01		-	-	-	-	-	-	-	-	-	+	-	-	+	
32 9297	HAG	*41:02		-	-	-	+	-	-	+	-	-	-	-	-	-	
33 9098	MT14B	*40:01		-	-	-	+	-	-	-	-	-	-	-	-	-	
34 9104	DHIF	*38:01		-	-	-	-	+	-	-	-	-	-	-	-	-	
35 9302	SSTO	*44:02		-	-	-	+	-	-	-	-	-	-	-	-	-	
36 9024	KT17	*15:01	*35:01	-	-	-	-	-	-	+	-	-	-	+	-	-	
37 9065	HHKB	*07:02		+	-	-	-	-	-	-	-	-	-	-	-	-	
38 9099	LZL	*15:01		-	-	-	-	-	+	-	-	-	+	-	-	-	
39 9315	CML	*08:01	*27:05	-	+	-	-	+	-	-	+	-	-	-	-	+	
40 9134	WHONP199	*13:02	*46:01	-	-	+	-	-	+	-	-	+	-	+	-	-	
41 9055	H0301	*14:02		-	-	-	+	-	+	-	-	-	-	-	-	-	
42 9066	TAB089	*46:01		-	-	-	-	-	+	-	-	-	+	-	-	-	
43 9076	T7526	*46:01		-	-	-	-	-	-	+	-	-	+	-	-	-	
44 9057	TEM	*38:01		-	-	-	-	+	-	-	-	-	-	-	-	-	
45 9239	SHJO	*42:01	*50:01	-	-	-	+	-	-	+	-	-	-	-	-	-	
46 9013	SCHU	*07:02		+	-	-	-	-	-	-	-	-	-	-	-	-	
47 9045	TUBO	*51:01		-	-	-	-	-	-	-	+	-	-	-	-	-	
48 9303	TER-ND	*35:01	*44:03	-	-	-	+	-	-	-	-	-	+	-	-	-	

CELL LINE VALIDATION SHEET																				
HLA-B low resolution SSP kit																				
				Prod. No.:	Well															
					17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32
					201190017	201190018	201190019	201195520	201190021	201190022	201190023	201190024	201190025	201190026	201190027	201190028	201190029	201190030	201190031	201190032
	IHWC cell line	B*																		
1	9001	SA	*07:02		-	-	-	-	-	-	-	+	-	-	-	+	-	-	-	-
2	9280	LK707	*52:01	*73:01	-	-	-	-	-	-	-	+	-	-	-	+	-	-	-	-
3	9011	E4181324	*52:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
4	9275	GU373	*15:10	*53:01	+	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-
5	9009	KAS011	*37:01		+	-	-	+	-	-	-	-	-	-	-	-	+	-	-	-
6	9353	SM	*39:01	*51:01	-	-	-	-	+	-	+	+	-	-	-	-	-	+	-	-
7	9020	QBL	*18:01		-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
8	9025	DEU	*35:01		+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
9	9026	YAR	*38:01		-	-	-	-	+	+	+	-	-	-	-	-	-	-	-	-
10	9107	LKT3	*54:01		-	-	-	-	-	-	-	-	-	-	-	+	-	-	+	-
11	9051	PITOUT	*44:03		+	-	-	-	-	-	-	-	-	-	-	-	-	+	-	-
12	9052	DBB	*57:01		+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
13	9004	JESTHOM	*27:05		-	-	-	-	-	-	-	-	-	-	-	+	-	-	+	-
14	9071	OLGA	*15:01	*15:20	+	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-
15	9075	DKB	*40:01		-	-	-	-	-	-	-	+	+	-	-	-	-	-	-	-
16	9037	SWEIG007	*40:02		-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-
17	9282	CTM3953540	*08:01	*55:01	-	-	-	-	-	-	-	+	-	-	+	+	-	-	+	-
18	9257	32367	*14:01	*56:01	-	-	-	-	-	-	+	+	-	-	-	+	-	-	+	-
19	9038	BM16	*18:01		-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
20	9059	SLE005	*40:01		-	-	-	-	-	-	+	+	-	-	-	-	-	-	-	-
21	9064	AMALA	*15:01		-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-
22	9056	KOSE	*35:03		+	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
23	9124	IHL	*40:02	*56:02	-	-	-	-	-	-	-	-	-	+	-	+	-	-	+	-
24	9035	JBUSH	*38:01		-	-	-	-	+	+	+	-	-	-	-	-	-	-	-	-
25	9049	IBW9	*14:02		-	-	-	-	-	-	+	+	-	-	-	-	-	-	-	-
26	9285	WT49	*58:01		+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
27	9191	CH1007	*07:05	*51:01	-	-	-	-	-	-	+	-	-	-	+	-	+	-	-	-
28	9320	BEL5GB	*44:02	*44:03	+	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-
29	9050	MOU	*44:03		+	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-
30	9021	RSH	*42:01		-	-	-	-	-	-	+	-	-	+	-	+	-	-	-	-
31	9019	DUCAF	*18:01		-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
32	9297	HAG	*41:02		-	-	-	-	-	-	+	-	-	+	-	-	-	-	-	-
33	9098	MT14B	*40:01		-	-	-	-	-	-	-	+	-	+	-	-	-	-	-	-
34	9104	DHIF	*38:01		-	-	-	-	+	+	+	-	-	-	-	-	-	-	-	-
35	9302	SSTO	*44:02		+	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-
36	9024	KT17	*15:01	*35:01	+	+	+	-	-	-	-	-	-	-	-	-	-	-	-	-
37	9065	HHKB	*07:02		-	-	-	-	-	-	+	-	-	-	-	+	-	-	-	-
38	9099	LZL	*15:01		-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-
39	9315	CML	*08:01	*27:05	-	-	-	-	-	-	+	-	-	+	+	-	-	-	+	-
40	9134	WHONP199	*13:02	*46:01	-	-	-	-	-	-	-	-	-	+	-	-	-	-	+	-
41	9055	H0301	*14:02		-	-	-	-	-	-	+	+	-	-	-	-	-	-	-	-
42	9066	TAB089	*46:01		-	-	-	-	-	-	-	-	-	-	-	+	-	-	+	-
43	9076	T7526	*46:01		-	-	-	-	-	-	-	-	-	-	-	+	-	-	+	-
44	9057	TEM	*38:01		-	-	-	-	+	+	+	-	-	-	-	-	-	-	-	-
45	9239	SHJO	*42:01	*50:01	-	-	-	-	-	-	+	-	-	-	+	+	-	-	-	-
46	9013	SCHU	*07:02		-	-	-	-	-	-	+	-	-	-	-	+	-	-	-	-
47	9045	TUBO	*51:01		-	-	-	-	-	-	-	-	-	-	-	-	-	+	-	-
48	9303	TER-ND	*35:01	*44:03	+	+	-	-	-	-	-	-	-	-	-	-	+	-	-	-

CELL LINE VALIDATION SHEET												
HLA-B low resolution SSP kit												
												Well
												33 34 35 36 37 38 39 40 41 42 43 44 45 46 47
												Prod. No.:
1	9001	SA	*07:02		-	-	-	-	-	-	-	201190033
2	9280	LK707	*52:01	*73:01	-	-	+	+	-	+	-	201190034
3	9011	E4181324	*52:01		-	-	+	+	-	+	-	201190035
4	9275	GU373	*15:10	*53:01	-	-	-	+	-	-	-	201190036
5	9009	KAS011	*37:01		-	-	-	-	+	-	-	201190037
6	9353	SM	*39:01	*51:01	-	-	+	+	-	-	-	201190038
7	9020	QBL	*18:01		-	-	-	-	-	-	-	201190039
8	9025	DEU	*35:01		-	-	-	-	-	-	-	201190040
9	9026	YAR	*38:01		-	-	-	-	-	-	-	201190041
10	9107	LKT3	*54:01		-	-	-	-	+	+	-	201190042
11	9051	PITOUT	*44:03		-	-	-	-	+	-	-	201190043
12	9052	DBB	*57:01		-	-	-	+	-	-	-	201190044
13	9004	JESTHOM	*27:05		-	-	-	-	+	-	-	201190045
14	9071	OLGA	*15:01	*15:20	-	-	-	-	+	-	-	201190046
15	9075	DKB	*40:01		-	-	-	+	+	-	-	201190047
16	9037	SWEIG007	*40:02		-	-	-	+	+	-	-	
17	9282	CTM3953540	*08:01	*55:01	-	-	+	-	-	+	-	
18	9257	32367	*14:01	*56:01	-	-	-	-	-	+	-	
19	9038	BM16	*18:01		-	-	-	-	-	-	+	
20	9059	SLE005	*40:01		-	-	-	+	+	-	-	
21	9064	AMALA	*15:01		-	-	-	-	+	-	-	
22	9056	KOSE	*35:03		-	-	-	-	-	-	+	
23	9124	IHL	*40:02	*56:02	-	-	-	+	+	-	-	
24	9035	JBUSH	*38:01		-	-	-	-	-	-	-	
25	9049	IBW9	*14:02		-	-	-	-	-	-	-	
26	9285	WT49	*58:01		-	-	+	-	-	-	+	
27	9191	CH1007	*07:05	*51:01	-	-	+	+	-	-	+	
28	9320	BEL5GB	*44:02	*44:03	-	-	-	-	+	+	-	
29	9050	MOU	*44:03		-	-	-	+	-	-	-	
30	9021	RSH	*42:01		-	-	-	-	-	-	-	
31	9019	DUCAF	*18:01		-	-	-	-	-	-	+	
32	9297	HAG	*41:02		-	-	-	+	+	-	-	
33	9098	MT14B	*40:01		-	-	-	+	+	-	-	
34	9104	DHIF	*38:01		-	-	-	-	-	-	+	
35	9302	SSTO	*44:02		-	-	-	-	+	+	-	
36	9024	KT17	*15:01	*35:01	-	-	-	-	+	-	-	
37	9065	HHKB	*07:02		-	-	-	-	-	-	-	
38	9099	LZL	*15:01		-	-	-	-	+	-	-	
39	9315	CML	*08:01	*27:05	-	-	-	-	+	-	-	
40	9134	WHONP199	*13:02	*46:01	-	-	-	-	+	-	-	
41	9055	H0301	*14:02		-	-	-	-	-	-	-	
42	9066	TAB089	*46:01		-	-	-	-	-	-	+	
43	9076	T7526	*46:01		-	-	-	-	-	-	+	
44	9057	TEM	*38:01		-	-	-	-	-	-	-	
45	9239	SHJO	*42:01	*50:01	-	-	+	-	+	-	-	
46	9013	SCHU	*07:02		-	-	-	-	-	-	-	
47	9045	TUBO	*51:01		-	-	+	+	-	-	+	
48	9303	TER-ND	*35:01	*44:03	-	-	-	-	+	-	+	

CERTIFICATE OF ANALYSIS

Olerup SSP® HLA-B low resolution SSP

Product number: 101.501-48/12 – including Taq pol.
Lot number: 98M
Expiry date: 2014-July-01
Number of tests: 48 tests – Product No. 101.501-48
12 tests – Product No. 101.501-12
Number of wells per test: 47 + 1

Well specifications:

Well No.	Production No.	Well No.	Production No.	Well No.	Production No.
1	2011-900-01	17	2011-900-17	33	2011-900-33
2	2011-900-02	18	2011-900-18	34	2011-900-34
3	2011-900-03	19	2011-900-19	35	2011-900-35
4	2011-900-04	20	2011-955-20	36	2011-900-36
5	2011-900-05	21	2011-900-21	37	2011-900-37
6	2011-900-06	22	2011-900-22	38	2011-900-38
7	2011-900-07	23	2011-900-23	39	2011-900-39
8	2011-900-08	24	2011-900-24	40	2011-900-40
9	2011-900-09	25	2011-900-25	41	2011-900-41
10	2011-900-10	26	2011-900-26	42	2011-900-42
11	2011-900-11	27	2011-900-27	43	2011-900-43
12	2011-900-12	28	2011-900-28	44	2011-900-44
13	2011-900-13	29	2011-900-29	45	2011-900-45
14	2011-900-14	30	2011-900-30	46	2011-900-46
15	2011-900-15	31	2011-900-31	47	2011-900-47
16	2011-900-16	32	2011-900-32		

The specificity of each primer solution of the kit has been tested against 48 well characterized IHWC cell line DNAs.

Additional 5'-primers and 3'-primers in primer solutions 3, 6, 14, 19, 29, 31 and 40 were tested by separately adding one additional 3'-primer, respectively one additional 5'-primer. Additional 3'-primers in primer solutions 2, 13, 15 and 35 were tested by separately adding one additional 5'-primer. Additional 5'-primers in primer solutions 1 21 and 25 were tested by separately adding one additional 3'-primer. In primer solution 31, one 5'-primer was not possible to test, in primer solution 38, one 3'-primer was not possible to test, and in primer solution 42, one 5'-primer and one 3'-primer was not possible to test.

The negative control primer pairs, **Production No. 2011-917-01**, can detect contamination with PCR products diluted 10⁻⁷.

Results: No false positive or false negative amplifications were obtained.

Date of approval: 2012-January-09

Approved by:

Production Quality Control

Declaration of Conformity

Product name: Olerup SSP® HLA-B low resolution

Product number: 101.501-48/12

Lot number: 98M

Intended use: HLA-B low resolution histocompatibility testing

Manufacturer: Olerup SSP AB
Franzengatan 5
SE-112 51 Stockholm, Sweden

Phone: +46-8-717 88 27

Fax: +46-8-717 88 18

We, Olerup SSP AB, hereby declare that this product, to which this Declaration of Conformity relates is in conformity with the following Standard(s) and other normative document(s) ISO 9001:2008 and ISO 13485:2003, following the provisions of the 98/79/EC Directive on *in vitro* diagnostic medical devices, Annex II List B, conformity assessed using Annex IV, as transposed into the national laws of the Member States of the European Union.

The Technical Documentation File is maintained at Olerup SSP AB, Franzengatan 5, SE-112 51 Stockholm, Sweden.

Notified Body: Lloyd's Register Quality Assurance Limited, Hiramford, Middlemarch Office Village, Siskin Drive, Coventry CV3 4FJ, United Kingdom.
(Notified Body number: 0088.)

Stockholm, Sweden
2012-January-09

Ann-Cathrin Jareman
Head of QA and Regulatory Affairs

**HLA-B low resolution
101.501-48/12 – including *Taq* polymerase**

Lot No.: 98M

Product Insert

Lot-specific information

General “**Instructions for Use**”

IFU-01 Rev. No. 03 can be downloaded from

www.olerup-ssp.com

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April 2014
Rev. No.: 01



0088

For *In Vitro* Diagnostic Use.

**HLA-B low resolution
101.501-48/12 – including *Taq* polymerase**

Lot No.: 98M

Product Insert

Lot-specific information

General “**Instructions for Use**”

IFU-01 Rev. No. 03 can be downloaded from

www.olerup-ssp.com

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April 2014
Rev. No.: 01



0088

For *In Vitro* Diagnostic Use.

**HLA-B low resolution
101.501-48/12 – including *Taq* polymerase**

Lot No.: 98M

Product Insert

Lot-specific information

General “**Instructions for Use**”

IFU-01 Rev. No. 03 can be downloaded from

www.olerup-ssp.com

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April 2014
Rev. No.: 01



0088

For *In Vitro* Diagnostic Use.

ADDRESSES:

Manufacturer:

Olerup SSP AB, Franzengatan 5, SE-112 51 Stockholm, Sweden.

Tel: +46-8-717 88 27

Fax: +46-8-717 88 18

E-mail: info-ssp@olerup.com

Web page: <http://www.olerup-ssp.com>

Distributed by:

Olerup GmbH, Löwengasse 47 / 6, AT-1030 Vienna, Austria.

Tel: +43-1-710 15 00

Fax: +43-1-710 15 00 10

E-mail: support-at@olerup.com

Web page: <http://www.olerup.com>

Olerup Inc., 901 S. Bolmar St., Suite R, West Chester, PA 19382

Tel: 1-877-OLERUP1

Fax: 610-344-7989

E-mail: info.us@olerup.com

Web page: <http://www.olerup.com>

For information on *Olerup SSP* distributors worldwide, contact **Olerup GmbH**.