

101.568-06 – including *Taq* polymerase, IFU-01
101.568-06u – without *Taq* polymerase, IFU-02

Visit <https://labproducts.caredx.com> for
“Instructions for Use” (IFU)

Lot No.: **7F4**

Lot-specific information

Olerup SSP[®] HLA-B*58

Product number:	101.568-06 – including <i>Taq</i> polymerase 101.568-06u – without <i>Taq</i> polymerase
Lot number:	7F4
Expiry date:	2022-10-01
Number of tests:	6
Number of wells per test:	31+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. 7F4.

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-B*58 Lot (7E4)

The HLA-B*58 kit is updated for new alleles to enable separation of:

- Null and Alternatively expressed alleles
- The product documentation has been updated for new alleles of IMGT 3.32.0

The format of the Worksheet has been changed.

One well has been added to HLA-B*58, well **32**.

The HLA-B*58 primer set, specificity and interpretation tables have been updated for the HLA-B alleles described since the previous *Olerup SSP[®]* HLA-B*58 lot was made (**Lot No. 7E4**).

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

Well	5'-primer	3'-primer	rationale
1	-	Added	3'-primer added for the B*58:01:21 allele.
9	Added	-	5'-primer added for the B*58:74 allele.
11	Modified	Modified	Primer pair modified for improved HLA-specific amplification.
13	Added	-	5'-primer added for the B*58:01:22 allele.
15	Added	-	5'-primer added for the B*58:74 allele.
16	-	Added	3'-primer added for the B*58:84 allele.
17	-	Added	3'-primer added for the B*58:84 allele.
23	-	-	Positive control primer pair exchanged for improved HLA-specific amplification.
24	Moved	Moved	Primer pair moved to well 31 for decreased tendency of primer oligomer formation.
25	Added	Added	Primer pair added for the B*58:35 allele.
26	Added	Added	Primer pair added for the B*58:35 allele.
28	-	-	Positive control primer pair exchanged for decreased tendency of primer oligomer formation.
31	Added	Added	Negative control moved to well 32. Primer pair added from well 24. 3'-primer added for the B*58:93N allele.
32	-	-	Negative control added from well 31.

Changes in R01 compared to R00:

1. Primer mix 30 does not amplify the B*58:62 allele. Thus, this lot cannot discriminate between the B*58:62 and the 58:01:01:01-58:01:19, 58:01:21-58:01:23, 58:32-58:34, 58:37, 58:41-58:42, 58:45:01-58:45:02, 58:47-58:50, 58:53-58:56, 58:63-58:64, 58:67-58:71, 58:73, 58:75, 58:77, 58:79-58:83, 58:85-58:90, 58:92 alleles. This has been corrected in the specificity and interpretation tables.

Changes in revision R02 compared to R01:

1. The expiration date has been altered due to extension of shelf-life.

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Well **32** 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-	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								+

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.

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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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Lot-specific information

PRODUCT DESCRIPTION

HLA-B*58 SSP subtyping

CONTENT

The primer set contains 5'- and 3'-primers for identifying the B*58:01 to B*58:93N alleles.

PLATE LAYOUT

Each test consists of 32 PCR reactions in a 32 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	NC

The 32 well cut PCR plate is marked with 'HLA-B*58' in silver/gray ink.

Well No. 1 is marked with the Lot No. '7F4'.

Wells 1 to 31 – HLA-B*58 high resolution primers.

Well 32 – Negative Control (NC).

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 heat-sealed with a PCR-compatible foil.

Please note: When removing each 32 well PCR plate, make sure that the remaining plates stay sealed. Use a scalpel or a similar instrument to carefully cut the foil between the plates.

INTERPRETATION

Due to the sharing of sequence motifs between HLA-B alleles, non-HLA-B*58 alleles will be amplified by some primer mixes. For further details see Specificity Table.

UNIQUELY IDENTIFIED ALLELES

All the HLA-B*58 alleles, i.e. **B*58:01 to B*58:93N** recognized by the HLA Nomenclature Committee in April 2018^{1,2,3} will be amplified by the primers in the HLA-B*58 subtyping kit.

The HLA-B*58 kit enables separation of the confirmed HLA-B*58 alleles as listed in the IMGT/HLA database 3.25.0. An HLA allele is listed as confirmed by IMGT/HLA if it has been sequenced by more than a single laboratory or from multiple sources. Current allele confirmation status for HLA-B*58 alleles is listed below.

The HLA-B*58 kit also enables identification of null and alternatively expressed alleles.

The following HLA-B*58 alleles can be distinguished by the different sizes of the HLA-specific PCR product:

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Lot-specific information

Alleles	Primer mix
B*58:10N, 58:13	12
B*58:21, 58:29	20
B*58:58, 58:66	26

¹HLA-B alleles listed on the IMGT/HLA web page 2018-April-16, release 3.32.0, 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>.

³This lot of the B*58 lot cannot discriminate between the B*58:62 and the B*58:01:01:01-58:01:19, 58:01:21-58:01:23, 58:32-58:34, 58:37, 58:41-58:42, 58:45:01-58:45:02, 58:47-58:50, 58:53-58:56, 58:63-58:64, 58:67-58:71, 58:73, 58:75, 58:77, 58:79-58:83, 58:85-58:90 and 58:92 alleles.

ALLELE CONFIRMATION STATUS

Allele	Status ¹	Allele	Status ¹	Allele	Status ¹	Allele	Status ¹
B*58:01:01:01	Confirmed	B*58:09	Unconfirmed	B*58:38	Unconfirmed	B*58:66	Unconfirmed
B*58:01:01:02	Unconfirmed	B*58:10N	Unconfirmed	B*58:39N	Unconfirmed	B*58:67	Unconfirmed
B*58:01:01:03	Unconfirmed	B*58:11	Confirmed	B*58:40	Confirmed	B*58:68	Unconfirmed
B*58:01:02	Unconfirmed	B*58:12	Confirmed	B*58:41	Unconfirmed	B*58:69	Unconfirmed
B*58:01:03	Unconfirmed	B*58:13	Unconfirmed	B*58:42	Unconfirmed	B*58:70	Unconfirmed
B*58:01:04	Unconfirmed	B*58:14	Unconfirmed	B*58:43	Unconfirmed	B*58:71	Unconfirmed
B*58:01:05	Unconfirmed	B*58:15	Unconfirmed	B*58:44	Unconfirmed	B*58:72N	Unconfirmed
B*58:01:06	Unconfirmed	B*58:16:01	Unconfirmed	B*58:45:01	Unconfirmed	B*58:73	Unconfirmed
B*58:01:07	Unconfirmed	B*58:16:02	Unconfirmed	B*58:45:02	Unconfirmed	B*58:74	Unconfirmed
B*58:01:08	Confirmed	B*58:17N	Unconfirmed	B*58:46	Unconfirmed	B*58:75	Unconfirmed
B*58:01:09	Unconfirmed	B*58:18	Confirmed	B*58:47	Unconfirmed	B*58:76	Unconfirmed
B*58:01:10	Unconfirmed	B*58:19	Unconfirmed	B*58:48	Unconfirmed	B*58:77	Unconfirmed
B*58:01:11	Unconfirmed	B*58:20	Unconfirmed	B*58:49	Unconfirmed	B*58:78	Unconfirmed
B*58:01:12	Unconfirmed	B*58:21	Confirmed	B*58:50	Unconfirmed	B*58:79	Unconfirmed
B*58:01:13	Unconfirmed	B*58:22	Unconfirmed	B*58:51	Unconfirmed	B*58:80	Unconfirmed
B*58:01:14	Unconfirmed	B*58:23	Confirmed	B*58:52	Unconfirmed	B*58:81	Unconfirmed
B*58:01:15	Unconfirmed	B*58:24	Confirmed	B*58:53	Unconfirmed		
B*58:01:16	Unconfirmed	B*58:25	Unconfirmed	B*58:54	Unconfirmed		
B*58:01:17	Unconfirmed	B*58:26	Confirmed	B*58:55	Unconfirmed		
B*58:01:18	Unconfirmed	B*58:27	Unconfirmed	B*58:56	Unconfirmed		
B*58:01:19	Unconfirmed	B*58:28:01	Unconfirmed	B*58:57	Unconfirmed		
B*58:01:20	Unconfirmed	B*58:28:02	Confirmed	B*58:58	Unconfirmed		
B*58:02:01	Confirmed	B*58:29	Unconfirmed	B*58:59:01	Confirmed		
B*58:02:02	Unconfirmed	B*58:31N	Unconfirmed	B*58:59:02	Unconfirmed		
B*58:04	Confirmed	B*58:32	Unconfirmed	B*58:60	Unconfirmed		
B*58:05	Unconfirmed	B*58:33	Unconfirmed	B*58:61	Unconfirmed		
B*58:06	Confirmed	B*58:34	Unconfirmed	B*58:62	Confirmed		
B*58:07	Confirmed	B*58:35	Unconfirmed	B*58:63	Unconfirmed		
B*58:08:01	Unconfirmed	B*58:36	Unconfirmed	B*58:64	Unconfirmed		
B*58:08:02	Confirmed	B*58:37	Unconfirmed	B*58:65	Unconfirmed		

¹Allele status “confirmed” or “unconfirmed” as listed on the IMGT/HLA web page 2016-July-14, release 3.25.0, www.ebi.ac.uk/imgt/hla.

RESOLUTION IN HOMO- AND HETEROZYGOTES

Results file with resolution in HLA-B*58 homo- and heterozygotes is available upon request.

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SPECIFICITY TABLE

HLA-B*58 SSP subtyping

Specificities and sizes of the PCR products of the 31+1 primer mixes used for HLA-B*58 SSP subtyping

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	Amplified HLA-B*58 alleles ³	Other amplified HLA-B alleles
1⁴	90 bp	800 bp	*58:01:01:01-58:02:02, 58:05-58:29, 58:31N-58:35, 58:37-58:38, 58:40-58:43, 58:45:01-58:93N	
2	150 bp	800 bp	*58:01:01:01-58:01:23, 58:04-58:05, 58:09-58:15, 58:17N, 58:19, 58:21-58:24, 58:28:01-58:29, 58:31N-58:37, 58:39N-58:42, 58:44-58:45:02, 58:47-58:59:02, 58:62-58:93N	*07:235, 08:49, 13:01:01:01-13:01:13, 13:06-13:07N, 13:12:01-13:13:02, 13:17, 13:20, 13:22:01-13:23, 13:25-13:26:02, 13:28-13:29, 13:36, 13:39, 13:43, 13:50-13:52, 13:57, 13:60-13:61, 13:63N, 13:73, 13:76N-13:80, 13:83, 13:86-13:87, 13:90, 13:92, 13:95, 13:106-13:107, 13:109-13:110, 14:10, 15:02:01-15:02:11, 15:13:01-15:13:02, 15:20-15:21:01:02, 15:25:01-15:25:03, 15:36, 15:44, 15:62, 15:77, 15:80, 15:85, 15:88-15:89, 15:106, 15:112, 15:121, 15:139, 15:144, 15:154, 15:165, 15:170, 15:194, 15:204, 15:213-15:214, 15:223, 15:240, 15:250, 15:265, 15:271, 15:283, 15:289, 15:291, 15:297, 15:301-15:302N, 15:308, 15:319, 15:325, 15:328, 15:330, 15:341, 15:345, 15:357-15:358, 15:366, 15:374, 15:378, 15:393-15:394, 15:399, 15:402, 15:404, 15:407, 15:418, 15:420, 15:425, 15:437-15:438, 18:22, 18:69, 18:105, 27:19, 27:30, 27:127, 27:150, 35:01:01:01-35:04:03, 35:06-35:08:10, 35:10-35:17:02, 35:19-35:21, 35:23-35:30, 35:33-35:36, 35:38-35:42:02, 35:45-35:50, 35:52, 35:54-35:57, 35:59:01-35:59:02, 35:61:01-35:63, 35:65Q, 35:69-35:71, 35:74, 35:76-35:78, 35:80-35:85, 35:90-35:96, 35:98, 35:100-35:101:02, 35:103-35:113, 35:115-35:116, 35:120-35:126, 35:128-35:134N, 35:136-35:150:02, 35:152-35:173:02N, 35:175-35:184, 35:186-35:198, 35:200-35:204, 35:206-35:229, 35:231, 35:233, 35:236-35:250, 35:252-35:259, 35:261-35:264, 35:266-35:272, 35:274-35:275, 35:277-35:291, 35:293-35:320, 35:322-35:330, 35:332-35:366, 35:370, 37:01:01:01-37:01:07, 37:01:09, 37:01:11-37:01:12, 37:03N-37:06:02, 37:08, 37:10-37:11, 37:13-37:43, 37:45-37:50, 37:52-37:71, 38:20, 39:42, 40:28, 44:02:01:01-44:02:33, 44:02:35-44:14, 44:16-44:17, 44:19N, 44:21-44:30, 44:32-44:40, 44:42-44:46, 44:48-44:52N, 44:55-44:64:02, 44:66-44:98, 44:101-44:105, 44:107-44:134, 44:136-44:137,

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3	240 bp	800 bp	*58:02:01-58:02:02, 58:06-58:07, 58:25, 58:38, 58:43, 58:46, 58:60	*14:19, 14:27, 50:51, 57:11, C*01:35, C*01:107, C*03:17:01-03:17:02, C*03:71, C*03:249, C*03:271, C*03:302, C*06:96, C*06:197, C*12:122, C*14:20
4	190 bp 230 bp	1070 bp	*58:04 *58:15	
5⁴	90 bp	1070 bp	*58:05	*15:124, 18:88, 38:43, 40:77, 40:87:01-40:87:02, 40:222, 40:237, 40:368, 44:247, 46:43, 48:14, 48:26, 48:29, 49:29, 51:181, 52:74, 57:59, C*07:81, C*07:168, C*07:450
6	170 bp	1070 bp	*58:06, 58:08:01-58:08:02, 58:19, 58:46, 58:78, 58:91	*07:02:01:01-07:24, 07:26-07:47, 07:49N-07:50, 07:52-07:154, 07:156-07:163, 07:165-07:201N, 07:203-07:208, 07:210-07:219, 07:221-07:222, 07:224-07:225, 07:227-07:234, 07:236-07:260, 07:262-07:300, 07:302-07:311, 08:20, 08:53:01-08:53:02, 08:79, 08:156, 13:16, 13:20, 13:31, 13:48, 13:62, 14:01:01:01-14:40, 14:42, 14:44-14:55, 14:57-14:58, 14:60-14:61, 15:01:01:01-15:01:04, 15:01:06-15:04:04, 15:06-15:17:04, 15:17:06-15:19, 15:21:01:01-15:21:01:02, 15:23-15:30:01:02, 15:32:01-15:40:02, 15:42-15:47:02, 15:49-15:50, 15:53-15:54, 15:56-15:58, 15:60-15:74, 15:76-15:82, 15:85, 15:87, 15:89-15:90, 15:92-15:99, 15:101-15:104, 15:106, 15:108-15:110, 15:112-15:113, 15:115-15:122, 15:125-15:129, 15:131-15:135, 15:137-15:144, 15:146-15:150, 15:152-15:154, 15:156-15:161, 15:163-15:175, 15:177-15:178, 15:180-15:184, 15:187, 15:189-15:199, 15:201-15:217, 15:219-15:221, 15:223, 15:225-15:234, 15:236, 15:238-15:249, 15:251-15:274, 15:276-15:282, 15:284-15:311, 15:313-15:331, 15:333-15:347, 15:349:01-15:382, 15:384-15:392, 15:394-15:395, 15:397-15:406, 15:408-15:410, 15:412-15:414, 15:416-15:420, 15:422-15:445, 18:15, 18:19, 18:21, 18:30, 18:57, 18:101, 27:04:01-27:04:06, 27:06, 27:10, 27:15, 27:18, 27:20-27:21, 27:24-27:25, 27:40, 27:54, 27:63, 27:66N, 27:68-27:69, 27:79, 27:86, 27:91-27:92, 27:100, 27:103, 27:105-27:109, 27:112-27:115, 27:120, 27:138, 27:149, 27:153, 27:165, 27:168, 27:173, 35:11:01-35:11:03, 35:14:01-35:14:02, 35:21, 35:43:01-35:44, 35:58, 35:67, 35:79, 35:86, 35:96, 35:99, 35:102, 35:117-35:118, 35:135, 35:167, 35:185, 35:188, 35:213,

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7 ⁴	80 bp	800 bp	*58:06, 58:08:01-58:08:02, 58:19, 58:46, 58:91	*07:02:35 ^w , 07:68:02 ^w , 07:151:02 ^w , 07:262 ^w , 13:16, 13:20, 13:48, 13:62, 14:01:01:01-14:02:06, 14:02:08-14:11, 14:12 ^w , 14:13-14:42, 14:44-14:46, 14:47 ^w , 14:48-14:55, 14:57-14:58, 14:60-14:61, 15:02:09, 15:03:02, 15:03:04, 15:13:02, 15:39:02, 15:42, 15:435, 18:15, 27:153, 35:11:01, 35:11:03 ^w , 35:21, 35:58, 35:96, 35:167, 35:188, 35:277, 35:337:02 ^w , 38:10, 38:32, 39:35, 39:107, 40:28, 40:95, 42:19, 44:76 ^w , 44:79:01, 48:25, 49:01:01:01-49:01:01:02, 49:01:03-49:10, 49:12-49:14, 49:16-49:21, 49:23-49:40, 49:41 ^w , 49:42-49:44, 49:46-49:50, 50:01:01:01-50:02, 50:04-50:20, 50:31, 50:32 ^w , 50:33-50:46, 50:48, 50:50-50:55, 51:01:01:01-51:01:24, 51:01:26-51:01:36, 51:01:38-51:01:46, 51:01:47 ^w , 51:01:48-51:01:51, 51:01:53-51:02:04, 51:03-51:04, 51:06:01-51:07:02, 51:11N-51:14, 51:16-51:18, 51:21-51:24:01, 51:24:04-51:24:05, 51:26-51:28, 51:29 ^w , 51:30, 51:32-51:35, 51:37-51:39, 51:41N, 51:43, 51:45-51:46, 51:48-51:53, 51:55 ^w , 51:56:01-51:60, 51:62-51:70, 51:74-51:80, 51:82 ^w , 51:83-51:86, 51:88-51:90, 51:92:01-51:92:02, 51:94-51:96, 51:98N-51:107, 51:109-51:114, 51:116-51:121, 51:123-51:125, 51:127-51:138, 51:140,

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8	250 bp	1070 bp	*58:07	C*06:165, C*14:61
9^{4,5}	90 bp 160 bp 190 bp	800 bp	*58:44 *58:08:01, 58:23 *58:72N, 58:74	*07:219 *57:01:20, A*02:01:33
10⁴	115 bp 270 bp	800 bp	*58:24 *58:09, 58:76	C*03:141, C*03:233, C*15:18 *14:10, 18:22, 18:69, 18:105, 35:21, 35:24:01-35:24:02, 35:81, 35:96, 35:109, 35:157, 35:188, 35:190, 35:233, 35:287, 37:04:01-37:04:02, 40:28, 44:190, 44:203, 44:275, 51:04, 51:46, 51:56:01-51:56:03, 51:139, 51:227, 53:02, 53:06, 53:28, 57:14:01-57:14:02, C*03:384, C*15:39, C*15:130
11	210 bp	1070 bp	*58:02:01-58:02:02, 58:07, 58:25, 58:38, 58:43, 58:60	*14:43, 14:59, 18:44:01-18:44:02, 40:243, 54:38, 57:11, A*24:64^w, C*01:06^w, C*06:79N, C*06:153, C*12:175, C*14:53, C*14:71^w, C*16:43^w
12	145 bp 235 bp	1070 bp	*58:13 *58:10N	*49:06
13	165 bp	1070 bp	*58:01:01:01-58:02:02, 58:04-58:10N, 58:12-58:19, 58:21-58:29, 58:31N-58:93N	*15:20, 15:228, 15:305, 15:435, 35:01:01:01-35:01:27, 35:01:29-35:30, 35:32:01-35:34, 35:36-35:42:02, 35:45, 35:47-35:56, 35:58-35:59:02, 35:61:01-35:72, 35:74-35:78, 35:80-35:86, 35:88-35:134N, 35:136-35:184, 35:186-35:278, 35:280-35:358, 35:360-35:370, 48:02:01-48:02:03, 51:01:01:01-51:24:05, 51:26-51:41N, 51:43-51:46, 51:48-51:156, 51:158:01-51:163, 51:165-51:219, 51:221-51:234, 52:01:01:01-52:06:03, 52:08-52:75, 53:01:01-53:50, 56:05:01-56:05:02, 56:21, 56:36, 67:05, 78:01:01:01, 78:01:02-78:09, 81:03
14⁶	165 bp	1070 bp	*58:11, 58:20	*07:02:01:01-07:57, 07:59-07:311, 08:01:01:01-08:05, 08:07-08:190, 08:192-08:195, 13:01:01:01-13:01:11, 13:01:13-13:04, 13:06-13:23, 13:25-13:60, 13:62-13:110, 14:01:01:01-14:61, 15:01:01:01-15:01:04, 15:01:06-15:19, 15:21:01:01-15:21:01:02, 15:23-15:40:02, 15:42-15:58, 15:60-15:99, 15:101, 15:103-15:129, 15:131-15:227, 15:229-15:283, 15:285-15:304N, 15:306-15:434, 15:436-15:445, 18:01:01:01-18:15, 18:17N-18:145, 27:01-27:05:30,

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15⁴	90 bp	1070 bp	*58:36	*57:01:01:01-57:01:22, 57:01:24-57:04:01, 57:04:03-57:15, 57:17-57:19, 57:21-57:35, 57:37-57:44, 57:46-57:50, 57:52-57:61, 57:63-57:68, 57:70-57:88, 57:90-57:96
	155 bp		*58:12	*07:219, 57:58, 57:80, A*02:42, A*02:310
	190 bp		*58:74	
16⁴	100 bp	1070 bp	*58:27-58:28:02, 58:65	*07:05:01:01-07:06:03, 07:32, 07:34, 07:40, 07:53, 07:69, 07:80, 07:90, 07:97, 07:105, 07:112, 07:123, 07:137-07:138, 07:140, 07:176, 07:182N, 07:201N, 07:206-07:207, 07:209-07:210, 07:213, 07:222, 07:249, 07:258, 07:262, 07:264, 07:269-07:270, 07:278, 07:283-07:284, 07:293, 07:304, 08:01:01:01-08:05, 08:08N, 08:10-08:11, 08:13, 08:15, 08:17-08:27, 08:29, 08:31-08:34, 08:36, 08:38-08:48, 08:50-08:68, 08:70-08:83, 08:85-08:86N, 08:88, 08:90-08:106, 08:108-08:110, 08:112-08:155, 08:157-08:183, 08:185-08:194, 13:13:01-13:13:02, 13:21, 13:71, 13:86, 13:108, 14:13, 14:53, 15:324, 15:329, 27:07:01-27:07:06, 27:11, 27:24, 27:33, 27:43, 27:164, 35:02:01:01-35:02:08, 35:04:01-35:04:03, 35:09:01-35:09:03, 35:12:01-35:12:04, 35:18, 35:22, 35:31, 35:81, 35:83, 35:87-35:88, 35:95, 35:129N, 35:149, 35:154, 35:157, 35:162, 35:172, 35:182-35:184, 35:201, 35:211, 35:220, 35:230, 35:233, 35:258, 35:266, 35:270, 35:285, 35:309, 35:311, 35:316, 35:321, 35:323, 35:335, 35:339, 35:357, 35:361, 35:366, 37:12, 39:14:01:01-39:14:01:02, 40:02:01:01-40:02:15, 40:02:17-40:02:25, 40:04-40:05:01:02, 40:08, 40:13, 40:15-40:16, 40:19, 40:28-40:30, 40:34-40:35:02, 40:37, 40:39-40:40, 40:45, 40:50, 40:56-40:58, 40:64:01:01-40:64:01:02, 40:68, 40:78, 40:80, 40:82, 40:85, 40:89-40:91, 40:94, 40:97, 40:99, 40:104, 40:107, 40:111, 40:115, 40:119, 40:122, 40:129, 40:133Q, 40:137, 40:142N-40:145, 40:157,

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	225 bp		*58:84	40:160:01-40:160:02, 40:164, 40:169, 40:173-40:174, 40:176, 40:181, 40:189, 40:200-40:203, 40:205-40:206, 40:209, 40:211, 40:214, 40:219-40:220, 40:224, 40:226, 40:229, 40:232, 40:246, 40:248, 40:254-40:255, 40:266, 40:271, 40:274, 40:276, 40:283, 40:287, 40:289-40:293, 40:296-40:297, 40:302-40:305, 40:309, 40:320, 40:322, 40:331, 40:334-40:335, 40:337N, 40:343, 40:345N, 40:347, 40:356, 40:359, 40:369, 41:02:01-41:02:07, 41:10-41:11, 41:13, 41:15, 41:19, 41:23, 41:27, 41:31, 41:36, 41:38-41:43, 41:45N-41:49, 41:52, 41:55, 42:01:01-42:02:01:02, 42:05:01-42:05:02, 42:07-42:08, 42:10-42:15, 42:17-42:19, 42:21-42:25, 44:62, 44:77, 44:82, 44:166, 44:184, 44:213, 48:01:01:01-48:01:06, 48:04:01-48:06, 48:09-48:13, 48:15-48:16, 48:18-48:20, 48:22, 48:24, 48:27-48:39, 48:41-48:45, 51:04, 51:46, 51:56:01-51:56:03, 51:64, 51:81, 51:139, 51:148, 53:19, 53:36, 55:04, 55:49, 55:51, 55:86, 56:12, 57:02:01-57:03:03, 57:05, 57:07, 57:09, 57:12, 57:17, 57:39, 57:42, 57:46, 57:57, 57:63, 57:66, 57:70, 57:80, 57:84, 57:94, 57:96, 81:01-81:08, C*03:317, C*03:382, C*15:24 *08:55, 15:265, 35:35, 37:40, 38:19-38:20, 39:03:01:01-39:03:01:02, 39:14:01:01-39:14:01:02, 39:24:01-39:24:02, 39:29, 39:37, 39:42, 39:76, 39:120, C*15:51
17⁵	225 bp	1070 bp	*58:22, 58:84	*07:12, 07:14, 07:18:01-07:18:02, 07:137, 07:162-07:163, 07:199, 07:225, 07:227, 07:248, 13:73, 15:265, 35:08:01:01-35:08:10, 35:18, 35:35, 35:45, 35:61:01-35:61:02, 35:80, 35:100, 35:105, 35:142, 35:156, 35:158, 35:164, 35:172, 35:174, 35:176, 35:187, 35:192, 35:206, 35:225, 35:229, 35:262, 35:300-35:301, 35:328, 35:333Q, 35:337:01-35:338, 35:341-35:342, 35:346, 35:356, 37:39-37:40, 38:20, 39:42, 44:04, 44:28:01-44:28:02, 44:56N, 44:76, 44:150, 53:11, 53:16, 53:24, 57:02:01-57:02:02, 57:04:01-57:05, 57:12-57:13, 57:19, 57:28N, 57:30, 57:42, 57:63, A*01:41, C*01:49:01-01:50, C*03:28, C*03:49, C*03:117, C*03:251, C*03:293, C*04:80, C*04:100, C*04:110, C*05:10, C*05:148, C*08:126, C*14:54, C*15:08, C*15:51, C*15:143, C*15:148
18⁴	70 bp 140 bp	1070 bp	*58:25 *58:17N	
19⁴	95 bp	800 bp	*58:18	*07:09, 07:11, 07:17, 07:237, 07:287, 08:28, 08:35, 08:37, 15:07:01:01-15:07:03, 15:55, 15:68, 15:126, 15:197:02, 15:207, 15:331, 15:360, 15:383, 15:405, 15:421, 15:431, 18:14, 35:05:01:01-35:05:03, 35:51, 35:58, 35:66, 35:72, 35:89, 35:97, 35:114, 35:232, 35:368, 40:03:01:01-40:03:01:02, 40:105, 40:267, 40:284,

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				41:24, 41:51, 42:09, 46:12, 48:14, 53:14, C*02:60, C*07:294, C*07:526, C*07:530
20⁴	85 bp 145 bp 175 bp 275 bp	1070 bp	*58:29 *58:39N *58:21, 58:29 *58:21	*18:51, 53:34 *53:34
21	440 bp	1070 bp	*58:14	*35:208
22	160 bp	1070 bp	*58:51	
23	150 bp 260 bp	800 bp	*58:26 *58:31N, 58:52	*35:75, 39:106, 44:138Q *51:32
24	190 bp 280 bp	1070 bp	*58:72N *58:40	
25	130 bp 260 bp 375 bp	1070 bp	*58:57 *58:52 *58:35	*51:32
26⁴	75 bp 170 bp 375 bp	1070 bp	*58:66 *58:58 *58:35	
27	590 bp	1070 bp	*58:08:01-58:09, 58:76	*57:14:01-57:14:02
28	365 bp	800 bp	*58:01:20, 58:16:01- 58:16:02, 58:18	*57:82
29	385 bp	1070 bp	*58:08:01- 58:08:02, 58:20, 58:27, 58:59:01- 58:59:02	*07:219, 15:17:01:01-15:17:06, 15:162, 15:168, 15:177, 15:196, 15:208, 15:216, 15:230, 15:268, 15:273, 15:356, 15:361, 15:396, 15:403, 15:411, 15:423-15:424, 57:01:01:01-57:01:09, 57:01:11- 57:04:03, 57:06-57:66, 57:68-57:96
30	170 bp	1070 bp		
31	145 bp 550 bp	1070 bp	*58:39N *58:93N	
32⁷	-	-	-	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*58 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.

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

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matched with the target sequences or not. Assumption is made that unknown sequences in these regions are conserved within allelic groups.

⁴HLA-specific PCR products shorter than 125 base pairs have a lower intensity and are less sharp than longer PCR products.

⁵Primer mixes 9 and 17 may have tendencies of primer oligomer formation.

⁶Primer mix 14 may have tendencies of unspecific amplifications.

⁷Primer mix 32 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.

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

'w', might be weakly amplified.

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PRIMER SPECIFICATION

Well No.	1	2	3	4	5	6	7	8	9	10	11	12
Length of spec. PCR product	90	150	240	190	90	170	80	250	90	115	210	145
				230					160	270		235
									190			
Length of int. pos. control ¹	800	800	800	1070	1070	1070	800	1070	800	800	1070	1070
5'-primer(s) ²	207	354	361	106	463	527	486	361	105	355	361	367
	5'-ACC 3'	5'-ATA 3'	5'-AgT 3'	5'-CCA 3'	5'-TgA 3'	5'-TgA 3'	5'-ACC 3'	5'-AgT 3'	5'-gCT 3'	5'-TCA 3'	5'-AgT 3'	5'-T.T 3'
	209	355							141			454
	5'-ggA 3'	5'-TCA 3'							5'-ATT 3'			5'-ACA 3'
									206			
									5'-AgA 3'			
									355			
									5'-TCA 3'			
3'-primer(s) ³	256	463	559	256	512	3 rd I	527	572	256	430	527	559
	5'-CCC 3'	5'-gCT 3'	5'-CAg 3'	5'-CTC 3'	5'-CCA 3'	5'-TAT 3'	5'-CCT 3'	5'-gCg 3'	5'-CCC 3'	5'-gCg 3'	5'-CCA 3'	5'-CAg 3'
	256			296					479	583	540	
	5'-CCC 3'			5'-CTg 3'					5'-CCA 3'	5'-gTg 3'	5'-CTT 3'	
									506			
									5'-TgC 3'			
Well No.	1	2	3	4	5	6	7	8	9	10	11	12

Well No.	13	14	15	16	17	18	19	20	21	22	23	24
Length of spec. PCR product	165	165	90	100	225	70	95	85	440	160	150	190
			155	225		140		145			260	280
			190					175				
								275				
Length of int. pos. control ¹	1070	1070	1070	1070	1070	1070	800	1070	1070	1070	800	1070
5'-primer(s) ²	652	652	105	355	355	209	363	106	206	653	355	355
	5'-CCg 3'	5'-CCA 3'	5'-gCT 3'	5'-TCA 3'	5'-TCA 3'	5'-ggA 3'	5'-AgC 3'	5'-CCA 3'	5'-gAC 3'	5'-CgC 3'	5'-TCC 3'	5'-TCA 3'
	652		142	363				209			652	
	5'-CTg 3'		5'-TCT 3'	5'-AgC 3'				5'-ggA 3'			5'-CCg 3'	
			209									
			5'-ggC 3'									
3'-primer(s) ³	774	774	256	412	538	239	419	208	362	774	463	506
	5'-ggT 3'	5'-ggT 3'	5'-CCC 3'	5'-gTT 3'	5'-CCg 3'	5'-gCg 3'	5'-Cgg 3'	5'-CTA 3'	5'-TCA 3'	5'-ggT 3'	5'-gCT 3'	5'-TgC 3'
				544	544	309		252			870	596
				5'-ggT 3'	5'-ggT 3'	5'-A.C 3'		5'-TCg 3'			5'-gA 3'	5'-TCT 3'
								340			872	
								5'-ggT 3'			5'-TCA 3'	
Well No.	13	14	15	16	17	18	19	20	21	22	23	24

101.568-06 – including Taq polymerase, IFU-01
101.568-06u – without Taq polymerase, IFU-02

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Lot No.: 7F4

Lot-specific information

Well No.	25	26	27	28	29	30	31
Length of spec.	130	75	590	365	385	170	145
PCR product	260	170					550
	375	375					
Length of int.	1070	1070	1070	800	1070	1070	1070
pos. control ¹							
5'-primer(s) ²	2 nd I	2 nd I	281	281	281	209	106
	5'-CCg 3'	5'-CCg 3'	5'-CTC 3'	5'-CTC 3'	5'-CTC 3'	5'-ggA 3'	5'-CCA 3'
	652	652					
	5'-CCg 3'	5'-CCg 3'					
3'-primer(s) ³	559	559	583	353	379	339	208
	5'-CgT 3'	5'-CgT 3'	5'-gTg 3'	5'-ggg 3'	5'-CAC 3'	5'-gCg 3'	5'-CTA 3'
	739	685		369	379		369
	5'-gTA 3'	5'-ggA 3'		5'-CCg 3'	5'-CAC 3'		5'-CCT 3'
	872	782					
	5'-TCA 3'	5'-CTT 3'					
Well No.	25	26	27	28	29	30	31

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.

101.568-06 – including *Taq* polymerase, IFU-01
101.568-06u – without *Taq* polymerase, IFU-02

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Lot No.: 7F4

Lot-specific information

[illegible]

101.568-06 – including *Taq* polymerase, IFU-01
101.568-06u – without *Taq* polymerase, IFU-02

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Lot No.: **7F4**

Lot-specific information

CELL LINE VALIDATION SHEET																																
HLA-B*58 SSP subtyping kit ²																																
					Well																											
					Prod. No.:	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31												
						201786017	201786018	201786019	201786020	201786021	201786022	201786023	201786024	201786025	201786026	201786027	201786028	201786029	201786030	201786031												
	IHWC cell line ¹		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	9025	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	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-

101.568-06 – including *Taq* polymerase, IFU-01
101.568-06u – without *Taq* polymerase, IFU-02

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Lot No.: **7F4**

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.

No DNAs carrying the alleles to be amplified by primer solutions 4, 5, 8 to 10, 12, 18, 20 to 28, 30 and 31 were available. The specificities of the primers in primer solutions 5, 8 to 10, 23 and 25 to 28 were tested by separately adding one or two additional 5'-primers, respectively one, two or three additional 3'-primers. In primer solution 4, 18, 20, 21, 24, 30 and 31 it was only possible to test the 5'-primers, the 3'-primer were not possible to test. In primer solutions 12 and 22 it was only possible to test the 3'-primers, the 5'-primers were not possible to test. In primer solutions 9 to 11, 23, 25 and 26 one or two 3'-primers were not possible to test. In primer solution 1, 2, 13, 25 and 26 one 5'-primer was not possible to test. Additional primers in primer solutions 15 to 17 and 29 were tested by separately adding one additional 5'-primer or one additional 3'-primer.

101.568-06 – including *Taq* polymerase, IFU-01
101.568-06u – without *Taq* polymerase, IFU-02

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“Instructions for Use” (IFU)

Lot No.: **7F4**

Lot-specific information

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