

101.549-06 – including *Taq* polymerase
101.549-06u – without *Taq* polymerase

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

Lot No.: **3R1**

Lot-specific information

Olerup SSP[®] HLA-B*53

Product number:	101.549-06 – including <i>Taq</i> polymerase 101.549-06u – without <i>Taq</i> polymerase
Lot number:	3R1
Expiry date:	2026-09-01
Number of tests:	6
Number of wells per test:	23+1
Storage - pre-aliquoted primers:	dark, between -15°C and -25°C
- PCR Master Mix:	between -15°C and -25°C
- Adhesive PCR seals	RT

This Product Description is only valid for Lot No. 3R1.

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*53 LOT (0L7).

- The product documentation has been updated for new alleles of IMGT 3.49.0
- The kit resolution focuses on common and well documented (CWD) alleles¹.

The HLA-B*53 specificity and interpretation tables have been updated for the HLA-B alleles described since the previous *Olerup SSP[®] HLA-B*53* lot was made (**Lot No. 0L7**).

The HLA-B*53 primer set is unchanged compared to the previous *Olerup SSP[®] HLA-B*53* (**Lot No. 0L7**).

¹S. J. Mack, P. Cano, J. A. Hollenbach et al.
Common and well-documented HLA alleles: 2012 update to
the CWD catalogue. Tissue Antigens, 2013, 81, 194–203



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Well **24** contains Negative Control primer pairs, that will amplify the majority of the Olerup SSP[®] HLA Class I, DRB, DQB1, DPB1 and DQA1 amplicons as well as all the amplicons generated by the control primer pairs matching the human growth hormone gene.

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

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

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

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



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PRODUCT DESCRIPTION

HLA-B*53 SSP typing

CONTENT

The primer set contains 5'- and 3'-primers for identifying the B*53:01 to B*53:70 alleles.

PLATE LAYOUT

Each HLA-B*53 test consists of 24 PCR reactions in a 24 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	NC

The 24 well PCR plate is marked with 'HLA-B*53' in silver/gray ink.

Well No. 1 is marked with the Lot No. '3R1'.

Wells 1 to 23 – HLA-B*53 high resolution primers.

Well 24 – 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 24 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*53 alleles will be amplified by some primer mixes. For further details see Specificity Table.

UNIQUELY IDENTIFIED ALLELES

All the HLA-B*53, i.e. **B*53:01 to B*53:70**, recognized by the HLA Nomenclature Committee in July 2022^{1,2} will be amplified by the primers in the HLA-B*53 SSP kit³.

The HLA-B*53 kit enables separation of the confirmed HLA-B*53 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*53 alleles is listed below.

The HLA-B*53 kit also enables identification of many null and alternatively expressed alleles.



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¹HLA-B alleles listed on the IMGT/HLA web page 2022-July-12, release 3.49.0, 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>.

³The HLA-B*53 primer set cannot separate the following alleles. These alleles can be distinguished by the HLA-B low resolution kit.

Alleles

B*53:01:01:01-53:01:29, 53:25, 53:27, 53:32, 53:34-53:35, 53:41, 53:43:01-53:47, 53:49-53:54, 53:57-53:61, 53:63-53:64, 53:66, B*58:100
B*53:62, B*35:471, B*51:10, 51:42, 51:81, 51:115, 51:181, 51:337, 51:359

ALLELE CONFIRMATION STATUS

Allele	Status ¹	Allele	Status ¹	Allele	Status ¹
B*53:01:01	Confirmed	B*53:10	Unconfirmed	B*53:29	Confirmed
B*53:01:02	Unconfirmed	B*53:11	Unconfirmed	B*53:30	Unconfirmed
B*53:01:03	Unconfirmed	B*53:12	Unconfirmed	B*53:31	Unconfirmed
B*53:01:04	Unconfirmed	B*53:13	Unconfirmed	B*53:32	Unconfirmed
B*53:01:05	Confirmed	B*53:14	Unconfirmed	B*53:33	Confirmed
B*53:01:06	Confirmed	B*53:15	Confirmed	B*53:34	Unconfirmed
B*53:01:07	Confirmed	B*53:16	Unconfirmed	B*53:35	Unconfirmed
B*53:01:08	Unconfirmed	B*53:17:01	Unconfirmed	B*53:36	Unconfirmed
B*53:01:09	Unconfirmed	B*53:17:02	Confirmed	B*53:37	Unconfirmed
B*53:01:10	Unconfirmed	B*53:18	Confirmed	B*53:38	Unconfirmed
B*53:01:11	Confirmed	B*53:19	Confirmed	B*53:39	Unconfirmed
B*53:02	Unconfirmed	B*53:20	Confirmed	B*53:40	Unconfirmed
B*53:03	Unconfirmed	B*53:21	Unconfirmed	B*53:41	Unconfirmed
B*53:04	Unconfirmed	B*53:22	Confirmed	B*53:42	Unconfirmed
B*53:05	Confirmed	B*53:23	Unconfirmed	B*53:43	Unconfirmed
B*53:06	Unconfirmed	B*53:24	Confirmed		
B*53:07	Unconfirmed	B*53:25	Unconfirmed		
B*53:08:01	Unconfirmed	B*53:26	Confirmed		
B*53:08:02	Unconfirmed	B*53:27	Unconfirmed		
B*53:09	Confirmed	B*53:28	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*53 homo- and heterozygotes is available upon request.



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Lot-specific information
SPECIFICITY TABLE

HLA-B*53 SSP subtyping

Specificities and sizes of the PCR products of the 23+1 primer mixes used for
HLA-B*53 SSP subtyping

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	Amplified HLA-B*53 alleles ³	Other amplified HLA Class I alleles
1	145 bp	800 bp	*53:01:01:01-53:02, 53:04-53:08:02, 53:10, 53:14-53:29, 53:32-53:35, 53:37, 53:40-53:54, 53:56-53:66, 53:68-53:70	*18:67, 18:136, 18:194:01:01-18:194:01:02, 27:236, 35:471, 37:34, 38:06-38:07, 44:06, 49:03, 51:01:01:01-51:24:05, 51:26-51:46, 51:48-51:53, 51:55-51:77, 51:79-51:103, 51:105-51:111, 51:113-51:146, 51:148-51:266, 51:268-51:283, 51:285-51:340, 51:342-51:370, 51:372-51:374, 52:01:01:01-52:01:28, 52:01:30-52:01:51, 52:03-52:15, 52:17-52:19, 52:21:01-52:31:02, 52:33-52:109, 57:120, 58:01:01:01-58:02:02, 58:04-58:16:02, 58:18-58:29, 58:31N-58:35, 58:37-58:43, 58:45:01-58:63, 58:65-58:110, 58:112-58:136
2⁵	270 bp	1070 bp	*53:02, 53:06, 53:28	*14:10, 14:102, 18:22, 18:69, 18:105, 18:149-18:150, 18:204, 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, 44:203:01-44:203:02, 44:275, 44:290, 51:04:01-51:04:02, 51:46, 51:56:01-51:56:03, 51:139, 51:227, 51:281, 51:310, 52:88, 57:14:01-57:14:02, 58:09, 58:76, C*03:384, C*15:39, C*15:130, C*15:195
3^{4,7}	95bp 135 bp 290 bp 490 bp	800 bp	*53:42 *53:03, 53:05:01-53:05:02, 53:16, 53:33, 53:55 *53:15 *53:42	*07:12, 07:18:01-07:18:02, 07:163, 07:199, 07:227, 07:248, 07:350, 35:34, 35:39, 35:407, 40:120, 44:107, 44:123, C*03:81, C*03:388, C*04:422, C*15:06:01:01-15:06:03, C*15:158, C*15:177N, C*15:180, C*15:214 *18:54, 18:67, 18:108, 18:132, 18:195, 27:52, 35:203, 35:355, 37:01:01:01-37:01:05, 37:01:07-37:05, 37:07-37:09, 37:12-37:13, 37:15-37:27, 37:30N-37:33N, 37:35-37:36, 37:38-37:60, 37:62-37:66, 37:68-37:81, 37:83-37:99, 37:101-37:105, 51:185, 52:09:01-52:09:02 *08:49, 08:60, 08:76, 08:129, 08:181, 35:530, 41:48, 42:13, 44:342 *07:65, 07:134, 07:400, 18:205, 18:217, 35:34, 35:39, 35:199, 35:407, 37:07, 37:09, 51:90, 51:135, 52:12
4⁵	190 bp	800 bp	*53:04, 53:07, 53:18, 53:31, 53:33	*13:03, 13:48, 13:120, 13:146, 13:164, 14:19, 14:27, 15:58, 15:73:01:01-15:73:01:02, 15:133, 15:137, 15:229, 15:253, 15:303, 15:460, 15:486, 15:488, 15:552, 35:03:01:01-35:03:05, 35:03:07-35:03:35, 35:06, 35:13, 35:36, 35:38, 35:55-35:56, 35:59:01-35:59:02, 35:70, 35:74-35:75, 35:84-35:85, 35:98, 35:106, 35:109, 35:127-35:128, 35:136, 35:150:01-35:153, 35:155-35:156, 35:160, 35:163, 35:167, 35:169, 35:179, 35:181, 35:193, 35:195, 35:198, 35:204-35:205:03, 35:221, 35:223, 35:231,



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5	150 bp	800 bp	*53:09, 53:11-53:13, 53:22, 53:31, 53:36, 53:38, 53:67	*08:28, 08:35, 08:37, 08:89, 08:107, 08:291, 13:58, 15:51, 15:179:01-15:179:02, 15:199, 15:218Q, 15:518, 18:09, 18:56, 18:113, 35:115, 35:260, 37:10, 37:35, 37:56, 41:24, 41:30, 41:51, 42:09, 42:32, 44:17, 44:83, 44:134, 44:230, 44:308, 44:445, 45:09, 51:54, 51:78:01-51:78:02, 51:97, 51:341, 51:371, 52:20, 57:24, 57:119, 58:137, C*07:68, C*07:140:01-07:140:02, C*07:151, C*07:302, C*07:929
6	205 bp	800 bp	*53:02, 53:06, 53:28	*14:56, 15:38:01-15:38:02, 15:185, 15:335, 15:364, 15:368, 15:610, 18:01:01:01-18:01:49, 18:01:51-18:09, 18:12:01-18:15, 18:17N-18:20, 18:22-18:25, 18:27-18:34, 18:36-18:40, 18:42-18:55, 18:57:01-18:57:02, 18:59-18:60, 18:62-18:100, 18:102-18:109, 18:111-18:121, 18:123-18:147, 18:149-18:163, 18:167, 18:169-18:188, 18:190, 18:192-18:204, 18:206-18:216, 18:218-18:224, 35:21, 35:24:01-35:24:02, 35:188, 35:190, 35:287, 35:519, 39:43, 51:37, 51:45, 51:63:01-51:63:02, 51:97, 51:216, 51:251, 52:39, 52:97, 56:31,



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	240 bp		*53:03, 53:05:01-53:05:02, 53:12, 53:16, 53:33, 53:55 53:67	57:14:01-57:14:02, 58:09, 58:76, C*02:142, C*06:207, C*07:516, C*07:521:01-07:521:02, C*12:87, C*14:76 *08:52, 08:117, 08:126, 13:25, 13:156, 15:43, 15:87, 15:115, 15:594, 27:106, 27:224, 35:329, 37:55, 40:188, 40:394, 44:15:01:01-44:15:01:02, 44:109, 44:176, 44:257, 44:355, 44:537, 47:01:01:02-47:01:05, 47:05-47:10, 47:12, 49:67, 49:70, 49:75, 51:185, 52:09:01-52:09:02, 55:90, 55:121, 56:46, 57:60
7^{4,5}	125 bp	800 bp	*53:07, 53:10, 53:26, 53:65	*08:49, 08:59:01, 13:79, 15:32:01-15:32:02, 15:299, 15:583, 18:18:01:01-18:18:01:02, 18:204, 35:48, 35:108:01, 35:349, 35:433, 37:01:01:01-37:01:29, 37:03N-37:06:02, 37:08, 37:10-37:11, 37:13-37:18, 37:20-37:43, 37:45-37:55, 37:57-37:80, 37:82N-37:93, 37:95-37:105, 39:09:01:01-39:09:05, 39:88, 39:112, 39:158, 39:170-39:171, 39:176, 44:209, 44:248, 51:255, 58:121, C*07:168
8	170 bp	1070 bp	*53:06, 53:08:01-53:08:02, 53:28	*07:02:01:01-07:02:05, 07:02:07-07:24, 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, 07:262:01-07:300, 07:302-07:331, 07:333-07:345, 07:347-07:358, 07:360Q-07:362, 07:364-07:385, 07:387-07:406, 07:409-07:410, 07:412-07:419, 07:421-07:449, 07:451-07:465, 08:20:01-08:20:02, 08:53:01-08:53:02, 08:79:01-08:79:02, 08:156, 08:281, 08:296, 13:16, 13:20, 13:31, 13:48, 13:62, 13:128, 13:131, 13:145, 14:01:01:01-14:40, 14:42:01-14:42:02, 14:44-14:55, 14:57-14:58, 14:60-14:89, 14:91-14:111, 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, 15:23-15:30:01:03, 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, 15:89:01-15:90, 15:92-15:99, 15:101-15:104, 15:106, 15:108-15:110, 15:112-15:113, 15:115-15:122, 15:125:01-15:129, 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, 15:189:01-15:199, 15:201-15:217, 15:219-15:221, 15:223, 15:225-15:234, 15:236, 15:238-15:249, 15:251:01-15:274, 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, 15:422-15:448, 15:450-15:484, 15:486-15:498, 15:500-15:506, 15:508-15:517, 15:519-15:538, 15:540N-15:551, 15:553-15:576, 15:579-15:586, 15:588-15:598, 15:600-15:622, 15:624-15:627, 15:629-15:632, 15:634-15:635, 15:637, 18:15, 18:19, 18:21, 18:30, 18:57:01-18:57:02, 18:101, 18:152, 18:164, 27:04:01-27:04:06, 27:06:01:01-27:06:02, 27:10, 27:15, 27:18, 27:20-27:21:02, 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,



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51:04:02, 51:06:01:01-51:07:02, 51:09:01-51:19,
51:21-51:24:05, 51:26-51:28, 51:30-51:35, 51:37-
51:43, 51:45-51:46, 51:48-51:52, 51:53^w, 51:56:01-
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51:87^w, 51:88-51:90, 51:91^w, 51:92:01-51:96, 51:98N-
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52:01:24-52:01:25, 52:01:27-52:01:32, 52:01:34-52:18,
52:20, 52:22-52:23, 52:25:01-52:27, 52:28^w-52:29^w,
52:30-52:43, 52:45-52:60, 52:62-52:68, 52:70-52:96N,
52:98, 52:100-52:109, 54:01:01:01-54:05N, 54:07-
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101.549-06 – including *Taq* polymerase
 101.549-06u – without *Taq* polymerase

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17	145 bp	1070 bp	*53:30, 53:39	*40:13, 40:19, 40:109, 40:117, 40:292, 40:340, 40:394, 41:46, 41:60, 44:18:01:01-44:18:01:02, 44:25, 44:50:01, 44:50:03, 44:95, 44:355, 49:01:01:01-49:01:23, 49:04:01-49:14, 49:16-49:25, 49:27-49:69, 49:71-49:72, 49:74-49:78, 51:112, 57:45, 57:51, 57:69, 58:111
	275 bp		*53:20	*51:187, A*25:37, A*32:138
18⁴	110bp 140 bp	1070 bp	*53:48N *53:21	*35:381N, C*15:177N *07:134, 07:400, 18:01:01:01-18:01:50, 18:01:52-18:03:02, 18:05:01:01-18:15, 18:17N-18:24, 18:26-18:36, 18:38-18:47, 18:49-18:65, 18:67, 18:69-18:106Q, 18:108-18:124, 18:126-18:142, 18:144-18:177, 18:179-18:224, 27:52, 27:236, 35:50, 35:84, 35:162, 35:197, 35:231, 35:267, 35:423, 35:494, 37:01:01:01-37:01:14, 37:01:16-37:05, 37:07-37:27, 37:29-37:30N, 37:32-37:41, 37:43-37:54, 37:56-37:60, 37:62-37:73, 37:75-37:84, 37:86N-37:98, 37:100-37:105, 38:06-38:07, 39:19:01-39:19:02, 39:186, 40:209, 40:350, 51:57, 51:103, 51:116, 51:218, 51:354
19^{4,7}	115 bp	1070 bp	*53:23, 53:48N	*07:65, 07:134, 07:400, 08:32, 08:180, 08:239, 18:03:01:01-18:03:02, 18:132, 18:142, 18:152, 18:171, 18:198, 18:203, 27:236, 35:29:01-35:29:03, 35:69, 35:76, 35:136, 35:186 ^w , 35:229, 35:294, 35:316, 35:381N, 35:472, 35:497, 37:37, 39:19:01-39:19:02, 46:40:01 ^w -46:40:02 ^w , 51:20, 51:208, 56:06, 56:45, 78:01:01:02-78:01:02, 78:03, 78:07-78:08, C*07:1011^w, C*15:177N
20⁴	60 bp	1070 bp	*53:11, 53:16, 53:24, 53:55	*07:20, 07:24, 07:60, 07:220, 07:358, 15:65, 15:84, 15:99, 15:221, 15:237, 15:379, 15:569, 35:08:01:01-35:08:15, 35:18, 35:45, 35:61:01-35:61:02, 35:80, 35:99-35:100, 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, 35:232, 35:300-35:301, 35:328, 35:337:01-35:338, 35:342, 35:346, 35:356, 35:386, 35:388, 35:411, 35:437, 35:444-35:445, 35:466, 35:481, 35:488, 35:493-35:494, 35:556Q, 37:39, 39:33, 40:496, 41:62, 44:28:01-44:28:02, 44:76, 46:62, 49:41, 51:05:01:01-51:05:01:02, 51:29, 51:54, 51:217, 51:279, 52:24, 56:51, 56:63, 57:02:01:01-57:02:02, 57:04:01-57:05, 57:12, 57:19, 57:28N, 57:30, 57:42, 57:63, 58:22, C*01:107, C*03:28, C*03:49, C*03:549, C*04:08, C*04:147, C*05:39, C*08:115, C*14:20, C*18:08
21^{4,8}	125 bp	800 bp	*53:30	*08:155, 18:205, 35:01:10, 35:03:32, 35:04:02, 35:46-35:47, 35:154, 35:207, 35:217, 35:251:01, 35:257,



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22^{4,5}	115 bp 205 bp	1070 bp	*53:37 *53:19, 53:36	*35:512, 51:336 *13:86, 35:02:01:01-35:02:07, 35:02:08 ^w , 35:02:09-35:02:19, 35:81, 35:83, 35:95, 35:129N, 35:146, 35:154, 35:162, 35:172, 35:182-35:184, 35:211, 35:220, 35:258, 35:270, 35:273, 35:285, 35:311, 35:323:01-35:323:02, 35:335, 35:339, 35:357, 35:361, 35:366-35:367, 35:372, 35:374, 35:377-35:379, 35:387, 35:391, 35:394, 35:396, 35:403, 35:407, 35:410, 35:419, 35:443, 35:483, 35:487, 35:492, 35:501, 35:505, 35:508N, 35:522, 35:546, 35:550, 35:553, 35:557, 35:562, 50:36, 58:65
23	165 bp	1070 bp	*53:29	*08:181, 35:208:01, 44:96, 44:153, 55:14, 57:01:19, 57:14:01, 58:14
24⁹	-	-		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*53 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 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 2, 4, 7 and 22 may have tendencies of unspecific amplifications.

⁶Primer mixes 11 and 14 may give rise to a lower yield of HLA-specific PCR product than the other HLA-B*53 primer mixes.

⁷Primer mixes 3 and 19 have a tendency to giving rise to primer oligomer formation.



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⁸Primer mix 21 may give rise to a long fragment of approx. 600 bp in some HLA-B alleles. This band should not be considered in the interpretation of HLA-A*53 typings.

⁹Primer mix 24 contains a negative control, which will amplify the majority 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 200 base pairs.

Abbreviations

w: may be weakly amplified.



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Lot No.: **3R1**

Lot-specific information
PRIMER SPECIFICATION

Well No.	1	2	3	4	5	6	7	8	9	10	11	12
Length of spec. PCR product	145	270	95	190	150	205	125	170	75	150	85	105
			135			240						
			290									
			490									
Length of int. pos. control ¹	800	1070	800	800	800	800	800	1070	1070	1070	800	1070
5'-primer(s) ²	206 5'-gAC 3'	355 5'-TCA 3'	206 5'-gAC 3'	49 5'-CAg 3'	206 5'-gAC 3'	106 5'-CCA 3'	106 5'-CCA 3'	527 5'-TgA 3'	527 5'-TgT 3'	419 5'-gTC 3'	419 5'-gTC 3'	357 5'-Tgg 3'
			355 5'-TCA 3'	419 5'-gTT 3'	419 5'-gTC 3'	419 5'-gTC 3'	368 5'-gTC 3'					
				419 5'-gTT 3'			368 5'-gTC 3'					
				419 5'-gTT 3'								
3'-primer(s) ³	309 5'-ATC 3'	583 5'-gTg 3'	301 5'-gTC 3'	79 5'-gTA 3'	309 5'-gTg 3'	302 5'-ggC 3'	180 5'-TCC 3'	3 rd I 5'-TAT 3'	559 5'-CAg 3'	527 5'-CCA 3'	463 5'-gCT 3'	420 5'-gCT 3'
	312 5'-gCA 3'		302 5'-ggC 3'	559 5'-CAg 3'	538 5'-gTC 3'	311 5'-ggg 3'	197 5'-gAA 3'			527 5'-CCA 3'		
			409 5'-ATg 3'			583 5'-gTg 3'	463 5'-gCT 3'			531 5'-TCT 3'		
			603 5'-gTg 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
Length of spec. PCR product	90	95	105	160	145	110	115	60	125	115	165
		130			275	140				205	
		190									
Length of int. pos. control ¹	1070	1070	1070	1070	1070	1070	1070	1070	800	1070	1070
5'-primer(s) ²	207 5'-ACC 3'	106 5'-CCA 3'	206 5'-gAC 3'	539 5'-gCT 3'	77 5'-CTT 3'	106 5'-CCg 3'	209 5'-ggA 3'	539 5'-gCg 3'	193 5'-CCA 3'	397 5'-gCT 3'	361 5'-Agg 3'
	208 5'-CgT 3'	363 5'-AgC 3'			206 5'-gAA 3'	355 5'-TCA 3'	355 5'-TCA 3'		435 5'-AAA 3'	652 5'-CCg 3'	
	209 5'-ggA 3'										
3'-primer(s) ³	256 5'-CTC 3'	197 5'-gAA 3'	272 5'-Tgg 3'	3 rd I 5'-TAT 3'	309 5'-ATC 3'	206 5'-CCg 3'	292 5'-gTC 3'	559 5'-CAg 3'	272 5'-TgA 3'	559 5'-CAg 3'	486 5'-gCg 3'
	256 5'-CCC 3'	256 5'-CCC 3'				426 5'-TCT 3'	292 5'-gTC 3'		527 5'-CCA 3'	728 5'-CCT 3'	
		419 5'-CgA 3'					426 5'-TCT 3'				
		419 5'-Cgg 3'									
Well No.	13	14	15	16	17	18	19	20	21	22	23

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



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Lot-specific information

numbering as on the www.ebi.ac.uk/imgt/hla web site. The sequence of the 3 terminal nucleotides of the primer is given.

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



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101.549-06 – including *Taq* polymerase
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“Instructions for Use” (IFU)

Lot No.: **3R1**

Lot-specific information

CELL LINE VALIDATION SHEET			HLA-B*53 SSP subtyping kit ²																
			Well																
			1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	
			Prod. No.:	201906001	201906002	201906003	201906004	201906005	201906006	201906007	201906008	201906009	201906010	201906011	201906012	201906013	201906014	201906015	201906016
	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	-	-	-	-	-	-	-	-	-	-	+	+	+	-	-	-	+



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Lot-specific information

CELL LINE VALIDATION SHEET											
HLA-B*53 SSP subtyping kit²											
				Well							
				17	18	19	20	21	22	23	
				Prod. No.:	201906017	201906018	201906019	201906020	201906021	201906022	201906023
	IHWC cell line¹	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 PTOU	*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	-	-	-	-	-	-	-	



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101.549-06 – including *Taq* polymerase
101.549-06u – without *Taq* polymerase

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

Lot No.: 3R1

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 2, 5 and 23 were available. The specificities of the primers in primer solutions 2, 5 and 23 were tested by separately adding one or two additional 3'-primers, and one or two additional 5'-primers, respectively. In primer solutions 1, 4, 7, 10, 13, 18, 19 and 22 one of the 3'-primers was not possible to be tested, and in primer mixes 13 and 17 one or two 5'-primers were not possible to be tested. In addition, one or more primers in primer solutions 3, 4, 6, 14, 18, 19, 21 and 22 were tested by separately adding one additional 3'-primer and/or one additional 5'-primer.



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For *In Vitro* Diagnostic Use
MA123 v02 SSP PI Template
Date: October 2022, Rev. No: 00

101.549-06 – including *Taq* polymerase
101.549-06u – without *Taq* polymerase

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“Instructions for Use” (IFU)

Lot No.: **3R1**

Lot-specific information

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