

101.628-06 – including *Taq* polymerase, IFU-01  
101.628-06u – without *Taq* polymerase, IFU-02

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

Lot No.: 4H9

Lot-specific information

## Olerup SSP<sup>®</sup> HLA-C\*17

Product number:	101.628-06 – including <i>Taq</i> polymerase 101.628-06u – without <i>Taq</i> polymerase
Lot number:	4H9
Expiry date:	2021-09-01
Number of tests:	6
Number of wells per test:	15+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. 4H9.

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

### CHANGES COMPARED TO THE PREVIOUS OLERUP SSP<sup>®</sup> HLA-C\*17 LOT (2G9)

The HLA-C\*17 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 3.35.0

The format of the Worksheet has been changed.

The HLA-C\*17 specificity and interpretation tables have been updated for the HLA-C alleles described since the previous *Olerup SSP<sup>®</sup>* HLA-C\*17 lot was made (**Lot No. 2G9**).

The HLA-C\*17 primer set is unchanged compared to the previous lot (**Lot No. 2G9**).

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Well **16** contains Negative Control primer pairs, that will amplify more than 95% of the *Olerup SSP*<sup>®</sup> HLA Class I, DRB, DQB1, DPB1 and DQA1 amplicons as well as all the amplicons generated by the control primer pairs matching the human growth hormone gene.

HLA-specific PCR product sizes range from 75 to 200 base pairs.

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

Length of PCR product	105	200	105	80	75	80	85
<b>5'-primer<sup>1</sup></b>	<b>164</b>	<b>340</b>	<b>440</b>	<b>45</b>	<b>45</b>	<b>43</b>	<b>36</b>
	5'-CAC <sup>3'</sup>	5'-Agg <sup>3'</sup>	5'-TTA <sup>3'</sup>	5'-Tgg <sup>3'</sup>	5'-Tgg <sup>3'</sup>	5'-Tgg <sup>3'</sup>	5'-TAC <sup>3'</sup>
							36
							5'-TAT <sup>3'</sup>
<b>3'-primer<sup>2</sup></b>	<b>231</b>	<b>2<sup>nd</sup> I</b>	<b>507</b>	<b>59</b>	<b>58</b>	<b>57</b>	<b>47</b>
	5'-TgC <sup>3'</sup>	5'-AAA <sup>3'</sup>	5'-TTg <sup>3'</sup>	5'-CTC <sup>3'</sup>	5'-ggC <sup>3'</sup>	5'-CTC <sup>3'</sup>	5'-ACA <sup>3'</sup>
							48
							5'-gCA <sup>3'</sup>
							48
							5'-gCC <sup>3'</sup>
							52
							5'-TgT <sup>3'</sup>
<b>A*</b>	<b>+</b>	<b>+</b>	<b>+</b>				
<b>B*</b>	<b>+</b>	<b>+</b>	<b>+</b>				
<b>C*</b>	<b>+</b>	<b>+</b>	<b>+</b>				
<b>DRB1</b>				<b>+</b>	<b>+</b>		
<b>DRB3</b>				<b>+</b>	<b>+</b>		
<b>DRB5</b>				<b>+</b>			
<b>DQB1</b>					<b>+</b>		
<b>DPB1</b>						<b>+</b>	
<b>DQA1</b>							<b>+</b>

<sup>1</sup>The nucleotide position for HLA class I genes and the codon for HLA class II genes, in the 2<sup>nd</sup> or 3<sup>rd</sup> 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](http://www.ebi.ac.uk/imgt/hla) web site. The sequence of the 3 terminal nucleotides of the primer is given.

<sup>2</sup>The nucleotide position for HLA class I genes and the codon for HLA class II genes, in the 2<sup>nd</sup> or 3<sup>rd</sup> exon or the 2<sup>nd</sup> 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](http://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-C\*17 SSP typing

#### CONTENT

The primer set contains 5'- and 3'-primers for identifying the C\*17:01 to C\*17:41 alleles.

#### PLATE LAYOUT

Each HLA-C\*17 test consists of 16 PCR reactions in a 16 well cut PCR plate.

1	2	3	4	5	6	7	8
9	10	11	12	13	14	15	NC

The 16 well PCR plate is marked with ‘HLA-C17’ in silver/gray ink.

Well No. 1 is marked with the Lot No. ‘4H9’.

Wells 1 to 15 – HLA-C\*17 high resolution primers.

Well 16 – 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 16 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-C alleles, non-HLA-C\*17 alleles will be amplified by some primer mixes. For further details see Specificity Table.

#### UNIQUELY IDENTIFIED ALLELES

All the HLA-C\*17 alleles, i.e. **C\*17:01 to C\*17:41**, recognized by the HLA Nomenclature Committee in January 2019<sup>1,2</sup> will be amplified by the primers in the HLA-C\*17 SSP kit.

The HLA-C\*17 kit enables separation of the confirmed HLA-C\*17 alleles as listed in the IMGT/HLA database 3.24.0. 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-C\*17 alleles is listed below.

The HLA-C\*17 kit also enables identification of null and alternatively expressed alleles.

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<sup>1</sup>HLA-C alleles listed on the IMGT/HLA web page 2019-January-23, release 3.35.0, [www.ebi.ac.uk/imgt/hla](http://www.ebi.ac.uk/imgt/hla).

<sup>2</sup>Alleles 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>.

## ALLELE CONFIRMATION STATUS

Allele	Status <sup>1</sup>	Allele	Status <sup>1</sup>	Allele	Status <sup>1</sup>
C*17:01:01:01	Confirmed	C*17:07	Confirmed	C*17:27N	Unconfirmed
C*17:01:01:02	Confirmed	C*17:08	Confirmed	C*17:28	Unconfirmed
C*17:01:01:03	Unconfirmed	C*17:09	Confirmed	C*17:29	Unconfirmed
C*17:01:01:04	Unconfirmed	C*17:10	Unconfirmed	C*17:30	Unconfirmed
C*17:01:01:05	Confirmed	C*17:11	Confirmed	C*17:31	Unconfirmed
C*17:01:02	Unconfirmed	C*17:12	Unconfirmed		
C*17:01:03	Confirmed	C*17:13	Unconfirmed		
C*17:01:04	Confirmed	C*17:14	Unconfirmed		
C*17:01:05	Unconfirmed	C*17:15	Confirmed		
C*17:01:06	Unconfirmed	C*17:16	Unconfirmed		
C*17:01:07	Unconfirmed	C*17:17	Confirmed		
C*17:01:08	Unconfirmed	C*17:18	Unconfirmed		
C*17:01:09	Confirmed	C*17:19	Unconfirmed		
C*17:01:10	Unconfirmed	C*17:20	Unconfirmed		
C*17:01:11	Unconfirmed	C*17:21	Unconfirmed		
C*17:02	Unconfirmed	C*17:22	Confirmed		
C*17:03	Confirmed	C*17:23	Unconfirmed		
C*17:04	Unconfirmed	C*17:24	Unconfirmed		
C*17:05	Unconfirmed	C*17:25	Unconfirmed		
C*17:06	Unconfirmed	C*17:26	Unconfirmed		

<sup>1</sup>Allele status “confirmed” or “unconfirmed” as listed on the IMGT/HLA web page 2016-April-15, release 3.24.0, [www.ebi.ac.uk/imgt/hla](http://www.ebi.ac.uk/imgt/hla).

## RESOLUTION IN HOMO- AND HETEROZYGOTES

Results file with resolution in HLA-C\*17 homo- and heterozygotes is available upon request.

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**SPECIFICITY TABLE**

**HLA-C\*17 SSP subtyping**

Specificities and sizes of the PCR products of the 15+1 primer mixes used for HLA-C\*17 SSP subtyping

Primer Mix	Size of spec. PCR product <sup>1</sup>	Size of control band <sup>2</sup>	Amplified HLA-C*17 alleles <sup>3</sup>	Other amplified HLA-C alleles
1	350 bp	800 bp	*17:01:01:02-17:01:14, 17:03:01:01-17:05, 17:06 <sup>2</sup> -17:16:01 <sup>2</sup> , 17:16:02-17:17, 17:18 <sup>2</sup> - 17:20 <sup>2</sup> , 17:21, 17:22 <sup>2</sup> - 17:29 <sup>2</sup> , 17:30, 17:31 <sup>2</sup> - 17:36 <sup>2</sup> , 17:37-17:41	*06:239, 07:665
2 <sup>4</sup>	70 bp	1070 bp	*17:02, 17:06 <sup>2</sup> -17:16:01 <sup>2</sup> , 17:18 <sup>2</sup> -17:20 <sup>2</sup> , 17:22 <sup>2</sup> - 17:29 <sup>2</sup> , 17:31 <sup>2</sup> -17:36 <sup>2</sup>	
3	300 bp	1070 bp	*17:03:01:01-17:03:02, 17:06 <sup>2</sup> -17:16:01 <sup>2</sup> , 17:16:02-17:17, 17:18 <sup>2</sup> - 17:20 <sup>2</sup> , 17:22 <sup>2</sup> -17:29 <sup>2</sup> , 17:31 <sup>2</sup> -17:36 <sup>2</sup> , 17:39- 17:41	
4 <sup>4</sup>	90 bp	1070 bp	*17:01:01:02-17:01:14, 17:04-17:05, 17:06 <sup>2</sup> - 17:16:01 <sup>2</sup> , 17:18 <sup>2</sup> -17:20 <sup>2</sup> , 17:21, 17:22 <sup>2</sup> -17:29 <sup>2</sup> , 17:30, 17:31 <sup>2</sup> -17:36 <sup>2</sup> , 17:37-17:38	<b>B*51:250</b>
5	155 bp 200 bp	1070 bp	*17:04 *17:15	*15:135
6 <sup>4</sup>	65 bp	1070 bp	*17:05	*01:43, 03:280, 05:01:12, 06:67, 07:101, 07:148, 07:161, 08:02:11
7 <sup>4</sup>	125 bp 200 bp	800 bp	*17:06 *17:15	
8 <sup>4</sup>	100 bp	1070 bp	*17:07	*03:129, 03:234, <b>B*15:116</b> , <b>B*15:352</b> , <b>B*15:412</b> , <b>B*35:331</b> , <b>B*40:63</b> , <b>B*40:92</b> , <b>B*40:324</b> , <b>B*44:169</b> , <b>B*44:182</b> , <b>B*46:43</b> , <b>B*57:59</b>
9 <sup>4</sup>	85 bp 135 bp 495 bp	1070 bp	*17:27N *17:08 *17:27N	
10 <sup>4</sup>	90 bp 495 bp	1070 bp	*17:09, 17:27N *17:27N	<b>B*40:260</b>
11	160 bp	1070 bp	*17:10	*03:319
12	450 bp	1070 bp	*17:11	
13 <sup>4</sup>	125 bp	1070 bp	*17:12	*15:98, <b>B*42:22</b>
14 <sup>4</sup>	80 bp 260 bp	1070 bp	*17:17 *17:13	
15 <sup>4,5</sup>	55 bp	1070 bp	*17:22	*01:17, 01:21, 01:23, 01:128, 01:152, 01:157, 02:12 <sup>w</sup> , 02:115, 03:27, 03:38:01-03:38:02, 03:69, 03:136, 03:246, 03:274, 03:431, 04:29, 04:172, 06:11, 06:122 <sup>w</sup> ,

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	07:01:01:01-07:01:40, 07:01:42-07:01:69, 07:01:71-07:02:40, 07:02:42-07:02:99, 07:02:101-07:04:13, 07:04:15-07:06:02, 07:08, 07:10-07:33N, 07:35-07:38:02, 07:41-07:48, 07:50-07:60, 07:62-07:75, 07:77-07:78:01, 07:79-07:114, 07:116-07:162, 07:164N-07:176, 07:178-07:209, 07:211-07:222, 07:224, 07:226-07:237, 07:239-07:246:02, 07:248-07:290, 07:292-07:294, 07:296-07:314:03, 07:316-07:327, 07:329N-07:334, 07:336-07:356, 07:358-07:360, 07:362-07:376, 07:378-07:402, 07:404-07:405, 07:407-07:414, 07:416-07:477, 07:478 <sup>w</sup> , 07:479-07:517, 07:519-07:527, 07:529-07:554, 07:556-07:558:01:02, 07:559 <sup>w</sup> , 07:560-07:562, 07:564-07:597, 07:599-07:653, 07:655, 07:656 <sup>w</sup> , 07:657-07:724, 08:05, 08:21, 08:25, 08:137, 12:02:01-12:02:22, 12:02:23 <sup>w</sup> , 12:02:24-12:03:55, 12:06-12:08, 12:10:01-12:20, 12:22-12:26, 12:28-12:32, 12:34-12:40, 12:42Q-12:48, 12:50-12:53, 12:55-12:59, 12:61-12:62, 12:64-12:71, 12:72 <sup>w</sup> , 12:73-12:97, 12:99:01-12:107, 12:109-12:134, 12:135 <sup>w</sup> , 12:136-12:145, 12:147-12:153, 12:154 <sup>w</sup> , 12:155Q-12:187, 12:189-12:267, 14:73, 15:25, 16:15:01-16:15:02, 16:64, <b>B*07:13, B*67:02</b>
16 <sup>6</sup>	<b>Negative Control</b>

<sup>1</sup>Alleles are assigned by the presence of specific PCR product(s). However, the sizes of the specific PCR products may be helpful in the interpretation of HLA-C\*17 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.

<sup>2</sup>The 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.

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<sup>3</sup>For several HLA Class I alleles 1<sup>st</sup> and/or 4<sup>th</sup> 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.

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

In order to have as little co-amplification as possible of non-HLA-C\*17 alleles by the primer pairs of the HLA-C\*17 subtyping kit, many of the specific PCR products are shorter than 100 base pairs.

<sup>5</sup>Primer mix 15 may give rise to a lower yield of HLA-specific PCR product than the other HLA-C\*17 primer mixes in the C\*07:02 alleles.

<sup>6</sup>Primer mix 16 contains a negative control, which will amplify more than 95% of HLA amplicons as well as the amplicons generated by the control primer pairs matching the human growth hormone gene. HLA-specific PCR product sizes range from 75 to 200 base pairs and the PCR product generated by the HGH positive control primer pair is 430 base pairs.

'w', might be weakly amplified.

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

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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	350	70	300	90	155	65	125	100	85	90	160	450
					200		200		135	495		
									495			
Length of int. pos. control <sup>1</sup>	800	1070	1070	1070	1070	1070	800	1070	1070	1070	1070	1070
5'-primer(s) <sup>2</sup>	20	28	70	20	126	176	126	499	295	295	80	341
	5'-CCA 3'	5'-TCA 3'	5'-ggA 3'	5'-CCA 3'	5'-ggA 3'	5'-gCA 3'	5'-ggA 3'	5'-TCT 3'	5'-ACT 3'	5'-ACT 3'	5'-CCg 3'	5'-ggA 3'
							412		406	445		
							5'-ATA 3'		5'-gCA 3'	5'-TCA 3'		
3'-primer(s) <sup>3</sup>	201	59	201	70	239	201	287	559	341	341	201	499
	5'-CTC 3'	5'-CgA 3'	5'-CTC 3'	5'-ggC 3'	5'-gCg 3'	5'-CTC 3'	5'-CCC 3'	5'-CAg 3'	5'-Cgg 3'	5'-Cgg 3'	5'-CTC 3'	5'-ggA 3'
	201				287		495		499	499		
	5'-CTT 3'				5'-CCC 3'		5'-ATA 3'		5'-ggA 3'	5'-ggA 3'		
Well No.	1	2	3	4	5	6	7	8	9	10	11	12

Well No.	13	14	15
Length of spec. PCR product	125	80	55
		260	
Length of int. pos. control <sup>1</sup>	1070	1070	1070
5'-primer(s) <sup>2</sup>	228	359	289
	5'-ATA 3'	5'-CCg 3'	5'-Agg 3'
		539	
		5'-gCg 3'	
3'-primer(s) <sup>3</sup>	312	580	302
	5'-AgT 3'	5'-TCC 3'	5'-ggC 3'
Well No.	13	14	15

<sup>1</sup>The 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.

<sup>2</sup>The nucleotide position matching the specificity-determining 3'-end of the primer is given. Nucleotide numbering as on the [www.ebi.ac.uk/imgt/hla](http://www.ebi.ac.uk/imgt/hla) web site. The sequence of the 3 terminal nucleotides of the primer is given.

<sup>3</sup>The 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](http://www.ebi.ac.uk/imgt/hla) web site. The sequence of the 3 terminal nucleotides of the primer is given.



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CELL LINE VALIDATION SHEET																			
HLA-C*17 SSP primer set <sup>2</sup>																			
				Well															
				1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	
				Prod. No.:	201891001	201891002	201891003	201891004	201891005	201891006	201891007	201891008	201891009	201891010	201891011	201891012	201891013	201891014	201891015
IHCW cell line <sup>1</sup>				C*															
1	9001	SA	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
2	9280	LK707	*07:01	*15:05	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
3	9011	E4181324	*12:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
4	9275	GU373	*03:04	*04:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
5	9009	KAS011	*06:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
6	9353	SM	*03:04	*07:02	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
7	9020	QBL	*05:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
8	9025	DEU	*04:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
9	9026	YAR	*12:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
10	9107	LKT3	*01:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
11	9051	PITOUT	*16:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
12	9052	DBB	*06:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
13	9004	JESTHOM	*01:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
14	9071	OLGA	*01:02	*03:04	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
15	9075	DKB	*03:04		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
16	9037	SWEIG007	*02:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
17	9282	CTM3953540	*03:03	*07:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
18	9257	32367	*01:02	*07:05	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
19	9038	BM16	*07:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
20	9059	SLE005	*03:04		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
21	9064	AMALA	*03:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
22	9056	KOSE	*12:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
23	9124	IHL	*01:02	*15:02	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
24	9035	JBUSH	*12:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
25	9049	IBW9	*08:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
26	9285	WT49	*07:18		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
27	9191	CH1007	*07:04	*15:29	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
28	9320	BEL5GB	*05:01	*16:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
29	9050	MOU	*16:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
30	9021	RSH	*17:01		+	-	-	+	-	-	-	-	-	-	-	-	-	-	-
31	9019	DUCAF	*05:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
32	9297	HAG	*17:03		+	-	+	-	-	-	-	-	-	-	-	-	-	-	-
33	9098	MT14B	*03:04		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
34	9104	DHIF	*12:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
35	9302	SSTO	*05:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
36	9024	KT17	*03:03	*04:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
37	9065	HHKB	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
38	9099	LZL	*03:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
39	9315	CML	*02:02	*07:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
40	9134	WHONP199	*01:02	*06:02	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
41	9055	H0301	*08:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
42	9066	TAB089	*01:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
43	9076	T7526	*01:02	*08:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
44	9057	TEM	*12:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
45	9239	SHJO	*06:02	*17:01	+	-	-	+	-	-	-	-	-	-	-	-	-	-	-
46	9013	SCHU	*07:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
47	9045	TUBO	*07:04	*15:02	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
48	9303	TER-ND	*04:01	*16:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-

101.628-06 – including *Taq* polymerase, IFU-01  
101.628-06u – without *Taq* polymerase, IFU-02

Visit <https://labproducts.caredx.com> for  
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Lot No.: 4H9

**Lot-specific information**

<sup>1</sup>The 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.

<sup>2</sup>The 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 and 5 to 14 were available.

The specificity of the primers in primer solutions 2, 6, 8, and 12 to 14 was tested by separately adding one additional 5'-primer, respectively one additional 3'-primer. In primer solutions 5 and 7 it was only possible to test the 5'-primers, the 3'-primers were not possible to test. In primer solutions 9 to 11 it was only possible to test the 3'-primers, the 5'-primers were not possible to test. In primer solution 14 one 5'-primer was not possible to test. One additional 3'-primer in primer solution 1 was tested by separately adding one 5'-primer.

101.628-06 – including *Taq* polymerase, IFU-01  
101.628-06u – without *Taq* polymerase, IFU-02

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Lot No.: 4H9

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