■LERUP SSP*
HLA-C*04 Add-on Product Insert Page 1 of 12

101.862-12 – including *Taq* **polymerase**, IFU-01 **101.862-12u – without** *Taq* **polymerase**, IFU-02

Visit https://labproducts.caredx.com for "Instructions for Use" (IFU)

Lot No.: **0H8** Lot-specific information

Olerup SSP® HLA-C*04 Add-on

Product number: 101.862-12 – including *Taq* polymerase

101.862-12u – without *Tag* polymerase

Lot number: 0H8

Expiry date: 2022-10-01

Number of tests: 12 Number of wells per test: 4+1

Storage - pre-aliquoted primers: dark at -20°C

PCR Master Mix: -20°C
 Adhesive PCR seals
 Product Insert
 RT

This Product Description is only valid for Lot No. 0H8.

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-C*04 Add-on Lot (6E1)

The HLA-C*04 Add-on specificity and interpretation tables have been updated for the HLA-C alleles described since the previous *Olerup* SSP® HLA-C*04 Add-on lot was made (Lot No. 6E1). The kit design is based on IMGT/HLA database 3.33.0.

The HLA-C*04 Add-on primer set is unchanged compared to the previous *Olerup* SSP® HLA-C*04 Add-on (Lot No. 6E1).

Changes in revision R01 compared to R00:

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

June 2020 Rev. No.: 01



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Well **5** contains <u>Negative Control primer pairs</u>, that will amplify more than 95% of the *Olerup* SSP® HLA Class I, DRB, DQB1, DPB1 and DQA1 amplicons as well as all the amplicons generated by the control primer pairs matching the human growth hormone gene.

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

Length of PCR	105	200	105	80	75	80	85
product							
5'-primer ¹	164	340	440	45	45	43	36
•	5'-CAC3'	^{5'} -Agg ^{3'}	⁵'-TTA3'	^{5'} -Tgg ^{3'}	^{5'} -Tgg ^{3'}	^{5'} -Tgg ^{3'}	5'-TAC3'
							36
							^{5'} -TAT ^{3'}
3'-primer ²	231	2 nd I	507	59	58	57	47
	5'-TgC ^{3'}		⁵ '-TTg ³ '	^{5'} -CTC ^{3'}	5'-ggC ^{3'}	5'-CTC ^{3'}	5'-ACA3'
							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 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.

-C*04 Add-on Product Insert

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PRODUCT DESCRIPTION

HLA-C*04 Add-on SSP subtyping

CONTENT

The primer set contains 5'- and 3'-primers for distinguishing the HLA-C*04:09N, 04:30 and 04:82 alleles from the C*04:01 allele.

PLATE LAYOUT

Each test consists of 5 PCR reactions in an 8 well cut PCR plate. Wells 6 to 8 are empty.

1 2 3 4 NC empty empty empty

The 8 well cut PCR plate is marked with the Lot No. '0H8' in silver/gray ink.

Well No. 1 is marked with the Lot No. '0H8'.

Wells 1 to 4 – HLA-C*04 Add-on high resolution primers.

Well 5 – 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 8 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 most other C*04 alleles will be amplified by some primer mixes. For further details see Specificity Table.

UNIQUELY IDENTIFIED ALLELES

The HLA-C*04:01, 04:09N, 04:30 and 04:82 alleles give different amplification patterns in the HLA-C*04 Add-on kit^{1,2}.

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¹Based on HLA-C alleles listed on the IMGT/HLA web page 2018-July-11, release 3.33.0, 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.

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Lot-specific information

SPECIFICITY TABLE

HLA-C*04 Add-on SSP subtyping

Specificities and sizes of the PCR products of the 4+1 primer mixes used for HLA-C*04 Add-on SSP subtyping

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	Amplified HLA- C*04:01/04:09N/04:30/04:82 alleles ³	Other amplified HLA-C alleles
1	260 bp, 360 bp	800 bp	*04:01:01:01-04:01:94, 04:09N, 04:30, 04:82	*03:231, 03:248, 03:308, 04:03:01-04:08, 04:10-04:20, 04:23-04:29, 04:31-04:81, 04:83-04:212, 04:214-04:225N, 04:227-04:293, 05:78:01-05:78:02, 05:141, 06:120, 08:163, 12:194, 15:100, 16:34, 16:62
2	220 bp	1070 bp	*04:09N	
3	180 bp	1070 bp	*04:30	
44	105 bp	1070 bp	*04:82	*04:159
5 ⁵	-	-	Negative Control	

¹Alleles are assigned by the presence of specific PCR product(s). However, the sizes of the specific PCR products may be helpful in the interpretation of HLA-C*04 Add-on SSP subtypings.

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 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 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.



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⁵Primer mix 5 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.

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101.862-12 – including *Taq* **polymerase**, IFU-01 **101.862-12u – without** *Taq* **polymerase**, IFU-02

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Lot No.: 0H8 Lot-specific information

PRIMER SPECIFICATION

Well No.	1	2	3	4
Length of spec.	260	220	180	105
PCR product	360			
Length of int.	800	1070	1070	1070
pos. control ¹				
5'-primer(s) ²	28	1018	1018	900
	5' -TCA 3'	^{5'} -gTg ^{3'}	^{5'} -gTg ^{3'}	^{5'} -CCg ^{3'}
3'-primer(s) ³	112	1092	1052	963
	5' -CCA 3'	^{5'} -TTA ^{3'}	^{5'} -Tgg ^{3'}	^{5'} -gCT ^{3'}
	118			
	^{5'} -gCT ^{3'}			
	218			
	^{5'} -gCT ^{3'}			
Well No.	1	2	3	4

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.



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9001 9280 9011 9275 9009 9353 9020 9025 9026 9107 9051 9052	QBL DEU YAR LKT3		*07:02	1 1 102424101 - · · · · · · · · · · · · · · · · · ·	201547702 5	201547703 8	201547704
9001 9280 9011 9275 9009 9353 9020 9025 9026 9107 9051 9052	SA LK707 E4181324 GU373 KAS011 SM QBL DEU YAR LKT3	*07:02 *07:01 *12:02 *03:04 *06:02 *03:04 *05:01 *04:01	*15:05 *04:01	201547701	201547702	3	201547704
9001 9280 9011 9275 9009 9353 9020 9025 9026 9107 9051 9052	SA LK707 E4181324 GU373 KAS011 SM QBL DEU YAR LKT3	*07:02 *07:01 *12:02 *03:04 *06:02 *03:04 *05:01 *04:01	*15:05 *04:01	201547701	201547702		201547704
9001 9280 9011 9275 9009 9353 9020 9025 9026 9107 9051 9052	SA LK707 E4181324 GU373 KAS011 SM QBL DEU YAR LKT3	*07:02 *07:01 *12:02 *03:04 *06:02 *03:04 *05:01 *04:01	*15:05 *04:01	-		201547703	-
9001 9280 9011 9275 9009 9353 9020 9025 9026 9107 9051 9052	SA LK707 E4181324 GU373 KAS011 SM QBL DEU YAR LKT3	*07:02 *07:01 *12:02 *03:04 *06:02 *03:04 *05:01 *04:01	*15:05	- - + -	- - - -	- - - -	-
9001 9280 9011 9275 9009 9353 9020 9025 9026 9107 9051 9052	SA LK707 E4181324 GU373 KAS011 SM QBL DEU YAR LKT3	*07:02 *07:01 *12:02 *03:04 *06:02 *03:04 *05:01 *04:01	*15:05	- - + -	- - - -	- - -	-
9280 9011 9275 9009 9353 9020 9025 9026 9107 9051 9052	LK707 E4181324 GU373 KAS011 SM QBL DEU YAR LKT3	*07:01 *12:02 *03:04 *06:02 *03:04 *05:01 *04:01	*04:01	- + -	-	-	-
9011 9275 9009 9353 9020 9025 9026 9107 9051 9052	E4181324 GU373 KAS011 SM QBL DEU YAR LKT3	*12:02 *03:04 *06:02 *03:04 *05:01 *04:01	*04:01	+	-	-	-
9275 9009 9353 9020 9025 9026 9107 9051 9052	GU373 KAS011 SM QBL DEU YAR LKT3	*06:02 *03:04 *05:01 *04:01		+	-	-	
9353 9020 9025 9026 9107 9051 9052	SM QBL DEU YAR LKT3	*03:04 *05:01 *04:01		-	-		-
9353 9020 9025 9026 9107 9051 9052	SM QBL DEU YAR LKT3	*05:01 *04:01	*07:02	-		-	-
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9025 9026 9107 9051 9052	DEU YAR LKT3	*04:01		-	-	-	-
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9052	PITOUT	*16:01		Ι-	-	-	_
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9071	OLGA	*01:02	*03:04	-	-	-	-
9075		*03:04	00.04	-	-	-	-
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		*05:01		-	-	-	-
				-	-	-	-
		*03:04		-	-	-	-
				-	-	-	-
		*05:01		-	-	-	-
		*03:03	*04:01	+	-	-	-
		*07:02		-	-	-	-
		*03:03		-	-	-	-
		*02:02	*07:01	-	-	-	-
9134	WHONP199	*01:02	*06:02	Ŀ	-	-	
9055	H0301	*08:02		-	-	-	-
9066	TAB089	*01:02		-	-	-	-
9076	T7526	*01:02	*08:01	-	-	-	-
9057	TEM	*12:03		-	-	-	-
9239	SHJO	*06:02	*17:01	-	-	-	-
9013	SCHU	*07:02		-	-	-	-
		*07:04	*15:02	-	-	-	-
		*04:01	*16:01	+	-	_	
	9038 9059 9064 9056 9124 9035 9049 9285 9191 9320 9050 9021 9019 9297 9098 9104 9302 9024 9065 9099 9315 9059 9076 9057 9057 9239 9013 9045	9257 32367 9038 BM16 9059 SLE005 9064 AMALA 9056 KOSE 9124 IHL 9035 JBUSH 9049 IBW9 9285 WT49 9191 CH1007 9320 BEL5GB 9050 MOU 9021 RSH 9019 DUCAF 9297 HAG 9098 MT14B 9104 DHIF 9302 SSTO 9024 KT17 9065 HHKB 9099 LZL 9315 CML 9134 WHONP199 9055 H0301 9066 TAB089 9076 T7526 9057 TEM 9239 SHJO 9013 SCHU 9045 TUBO 9303 TER-ND	9038 BM16 *07:01 9059 SLE005 *03:04 9064 AMALA *03:03 9056 KOSE *12:03 9124 IHL *01:02 9035 JBUSH *12:03 9049 IBW9 *08:02 9285 WT49 *07:18 9191 CH1007 *07:04 9320 BEL5GB *05:01 9050 MOU *16:01 9021 RSH *17:01 9019 DUCAF *05:01 9297 HAG *17:03 9098 MT14B *03:04 9104 DHIF *12:03 9302 SSTO *05:01 9024 KT17 *03:03 9055 HHKB *07:02 9099 LZL *03:03 9315 CML *02:02 9134 WHONP199 *01:02 9055 H0301 *08:02 9066	9038 BM16 *07:01 9059 SLE005 *03:04 9064 AMALA *03:03 9056 KOSE *12:03 9124 IHL *01:02 *15:02 9035 JBUSH *12:03 9049 IBW9 *08:02 9285 WT49 *07:18 9191 CH1007 *07:04 *15:29 9320 BEL5GB *05:01 *16:01 9021 RSH *17:01 9019 DUCAF *05:01 9297 HAG *17:03 9098 MT14B *03:04 9104 DHIF *12:03 9302 SSTO *05:01 9024 KT17 *03:03 *04:01 9025 HHKB *07:02 9099 LZL *03:03 9315 CML *02:02 *07:01 9134 WHONP199 *01:02 *06:02 9055 H0301 *08:02 9066 TAB089 *01:02 9076 T7526 *01:02 *08:01 9013 SCHU *07:02 9091 SCHU *07:02 9093 SHJO *06:02 *17:01 9013 SCHU *07:02 9045 TUBO *07:04 *15:02	9038 BM16	9038 BM16	9038 BM16



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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 3 and 4 were available. In primer solutions 3 and 4 it was only possible to test the 5'-primer, the 3'-primer was not possible to test.



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Lot No.: 0H8 Lot-specific information

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