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Lot No.: 22V Lot-specific information

Olerup SSP® HLA-C*17

Product number: 101.628-06 – including *Taq* polymerase

101.628-06u - without *Taq* polymerase

Lot number: 22V

Expiry date: 2016-July-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
 Product Insert
 RT

This Product Description is only valid for Lot No. 22V.

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*17 Lot (50R)

The HLA-C*17 kit is updated for new alleles to enable separation of:

- Confirmed¹ alleles as listed in the IMGT/HLA database
- Polymorphisms in exons outside of the region encoding the peptide binding domain
- Null and Alternatively expressed alleles

A well containing Negative Control primer pairs has been added.

The format of the Product Insert and Worksheet have been changed.

Two wells have been added to the HLA-C*17 kit, wells **15 and 16**.

The HLA-C*17 specificity and interpretation tables have been updated for the HLA-C alleles described since the previous *Olerup* SSP® HLA-C*17 lot was made (Lot No. 50R).

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¹As described in section Uniquely Identified Alleles.

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

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As of lot series V, the Specificity Table is included in the lot-specific Product Insert, and the Interpretation Table is included in the Worksheet.

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

Well	5'-primer	3'-primer	rationale
15	New	New	New primer pair for the C*17:22 allele.
16			Negative Control

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Well **16** contains <u>Negative Control primer pairs</u>, that will amplify more than 95% of the *Olerup* SSP[®] HLA Class I, DRB, DQB1 and DPB1 amplicons as well as amplicons generated by a control primer pair.

PCR product sizes range from 75 to 430 base pairs. The PCR product generated by the control primer pair is 430 base pairs.

Length of PCR	105	200	105	80	75	80
product						
5'-primer ¹	164	340	440	45	45	43
	5'-CAC3'	⁵ '-Agg ³ '	^{5'} -TTA ^{3'}	^{5'} -Tgg ^{3'}	⁵ '-Tgg ³ '	⁵ '-Tgg ³ '
3'-primer ²	231	2 nd I	507	59	58	57
_	^{5'} -TgC ^{3'}	^{5'} -AAA ^{3'}	^{5'} -TTg ^{3'}	^{5'} -CTC ^{3'}	^{5'} -ggC ^{3'}	^{5'} -CTC ^{3'}
A*	+	+	+			
B*	+	+	+			
C*	+	+	+			
DRB1				+	+	
DRB3				+	+	
DRB5				+		
DQB1					+	
DPB1						+

¹The nucleotide position for HLA class I genes and the codon for HLA class II genes, in the 2nd or 3rd exon, matching the specificity-determining 3'-end of the primer is given. Nucleotide and 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-C*17 SSP typing

CONTENT

The primer set contains 5'- and 3'-primers for identifying the C*17:01 to C*17:22 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. '22V'.

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 primer mixes 6, 8 and 15. In addition, a few HLA-B alleles will be amplified by primer mixes 8 and 15.

For further details see Specificity Table.

UNIQUELY IDENTIFIED ALLELES

All the HLA-C*17 alleles, i.e. **C*17:01 to C*17:22**, recognized by the HLA Nomenclature Committee in October 2013¹ 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. 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 polymorphisms in exons outside of the region encoding the peptide binding domain and of null and alternatively expressed alleles

The HLA-C*17 subtyping kit cannot distinguish the silent mutations in the C*17:01:01-17:01:09 alleles.

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¹HLA-C alleles listed on the IMGT/HLA web page 2013-October-11, release 3.14.0, www.ebi.ac.uk/imgt/hla.

ALLELE CONFIRMATION STATUS

Allele	Status ¹	Allele	Status ¹
C*17:01:01:01	Confirmed	C*17:11	Unconfirmed
C*17:01:01:02	Unconfirmed	C*17:12	Unconfirmed
C*17:01:01:03	Unconfirmed	C*17:13	Unconfirmed
C*17:01:02	Unconfirmed	C*17:14	Unconfirmed
C*17:01:03	Confirmed	C*17:15	Unconfirmed
C*17:01:04	Confirmed	C*17:16	Unconfirmed
C*17:01:05	Unconfirmed	C*17:17	Unconfirmed
C*17:01:06	Unconfirmed	C*17:18	Unconfirmed
C*17:01:07	Unconfirmed	C*17:19	Unconfirmed
C*17:01:08	Unconfirmed	C*17:20	Unconfirmed
C*17:01:09	Confirmed	C*17:21	Unconfirmed
C*17:02	Unconfirmed	C*17:22	Confirmed
C*17:03	Unconfirmed		
C*17:04	Unconfirmed		
C*17:05	Unconfirmed		
C*17:06	Unconfirmed		
C*17:07	Confirmed		
C*17:08	Unconfirmed		
C*17:09	Unconfirmed		
C*17:10	Unconfirmed		

¹Allele status "confirmed" or "unconfirmed" as listed on the IMGT/HLA web page 2013-October-11, release 3.14.0, 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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Lot-specific information 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 ¹	Size of control band ²	Amplified HLA-C*17 alleles ³	Other amplified HLA-C alleles ⁴
1	350 bp	800 bp	*17:01:01:01-17:01:09, 17:03, 17:04 [?] , 17:05, 17:06 [?] -17:22 [?]	
2 ⁵	70 bp	1070 bp	*17:02, 17:04 [?] , 17:06 [?] - 17:22 [?]	
3	300 bp	1070 bp	*17:03, 17:04 [?] , 17:06 [?] - 17:22 [?]	
4 ⁵	90 bp	1070 bp	*17:01:01:01-17:01:09, 17:04 [?] , 17:05, 17:06 [?] - 17:22 [?]	
5	155 bp	1070 bp	*17:04	
6 ⁵	65 bp	1070 bp	*17:05	*01:43, 05:01:12, 06:67, 07:101, 07:148, 07:161
7 ⁵	125 bp	800 bp	*17:06	
8 ⁵	100 bp	1070 bp	*17:07	*03:129, B*15:116, B*40:63, B*40:92, B*44:169, B*44:182, B*46:43, B*57:59
9	135 bp	1070 bp	*17:08	
10 ⁵	95 bp	1070 bp	*17:09	
11	160 bp	1070 bp	*17:10	
12	450 bp	1070 bp	*17:11	
13 ⁵	125 bp	1070 bp	*17:12	
14	180 bp	1070 bp	*17:13	
15 ⁵	55 bp	1070 bp	*17:22	*01:17, 01:21, 01:23, 02:12*, 03:27, 03:38:01-03:38:02, 03:69, 03:136, 04:29, 06:11, 07:01:01:01-07:02:40, 07:02:42-07:06, 07:08, 07:10-07:33N, 07:35-07:38:02, 07:41-07:48, 07:50-07:60, 07:62-07:75, 07:77-07:114, 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, 07:248-07:290, 07:292-07:294, 07:296-07:314,

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

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07:316-07:327, 07:329N-07:334, 07:336-07:337, 08:05, 08:21, 08:25, 12:02:01-12:03:28, 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^w, 12:73-12:97, 12:99-12:107, 12:109-12:111, 15:25, 16:15:01-16:15:02. B*07:13, B*67:02 16⁶ **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*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 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.

⁴Due to the sharing of sequence motifs between HLA-C alleles, non-HLA-C*17 alleles will be amplified by primer mixes 6, 8 and 15. In addition, a few HLA-B alleles will be amplified by primer mixes 8 and 15.

⁵HLA-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.

⁶Primer mix 16 contains a negative control, which will amplify more than 95% of HLA amplicons as well as the amplicons generated by control primer pairs. PCR product sizes range from 75 to 200 base pairs. The PCR product generated by the control primer pair is 430 base pairs. 'w', might be weakly amplified.

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

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

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PRIMER SPECIFICATION

Well No.	1	2	3	4	5	6	7	8	9	10	11	12
Length of spec.	350	70	300	90	155	65	125	100	135	95	160	450
PCR product												
Length of int.	800	1070	1070	1070	1070	1070	800	1070	1070	1070	1070	1070
pos. control ¹												
5'-primer(s) ²	20	28	70	20	126	176	412	499	406	445	80	341
	5' -CCA 3'	^{5'} -TCA ^{3'}	^{5'} -ggA ^{3'}	5' -CCA 3'	^{5'} -ggA ^{3'}	^{5'} -gCA ^{3'}	^{5'} -ATA ^{3'}	5' -TCT 3'	^{5'} -gCA ^{3'}	5' -TCA 3'	^{5'} -CCg ^{3'}	^{5'} -ggA ^{3'}
3'-primer(s) ³	201	59	201	70	239	201	495	559	499	499	201	499
. ,	5' -CTC 3'	5' -CgA 3'	5' -CTC 3'	^{5'} -ggC ^{3'}	5' -gCg 3'	5' -CTC 3'	^{5'} -ATA ^{3'}	5' -CAg 3'	^{5'} -ggA ^{3'}	5' -ggA 3'	5' -CTC 3'	^{5'} -ggA ^{3'}
	201											
	5' -CTT 3'											
Well No.	1	2	3	4	5	6	7	8	9	10	11	12

Well No.	13	14	15
Length of spec.	125	180	55
PCR product			
Length of int.	1070	1070	1070
pos. control ¹			
5'-primer(s) ²	228	359	289
	^{5'} -ATA ^{3'}	^{5'} -CCg ^{3'}	5' -Agg 3'
3'-primer(s) ³	312	499	302
	^{5'} -AgT ^{3'}	^{5'} -ggA ^{3'}	^{5'} -ggC ^{3'}
Well No.	13	14	15

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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Lot No.: ZZV Lot-specific information																			
CELL LINE VALIDATION SHEET																			
			H	ILA-C*	17	SS	Pρ	rim	ner	set	t ²								
							•					Wel	ı						
					1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
				Prod. No.:	201328901	201328902	201328903	201328904	201328905	201328906	201328907	201328908	201328909	201328910	201328911	201328912	201328913	201328914	201328915
				rod	013	013	013	013	013	013	013	013	013	013	013	013	013	013	013
	IHV	/C cell line ¹		С*	2	- 21	-21	N	- 21	N	-21	N		N	N	N	-21	N	-21
1	9001		*07:02	<u> </u>	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
2		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		LKT3	*01:02			-	-	-	-	-	-	-	-	-	-	-	-	-	-
11	9051	PITOUT	*16:01		-	-	-	-	-	-	-	-	•	-	-	-	-	-	-
12	9052		*06:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
13		JESTHOM	*01:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
14		OLGA	*01:02	*03:04	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
15	9075		*03:04		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
16		SWEIG007	*02:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
17		CTM3953540	*03:03	*07:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
18		32367	*01:02	*07:05	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
19		BM16	*07:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
20		SLE005	*03:04		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
21		AMALA	*03:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
22		KOSE	*12:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
23	9124		*01:02	*15:02	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
24		JBUSH	*12:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
25	9049		*08:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
26		WT49	*07:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
27		CH1007	*07:04	*15:05	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
28		BEL5GB	*05:01	*16:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
29	9050		*16:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
30	9021		*17:01		+	-	•	+	-	-	-	-	-	-	•	-	-	-	-
31		DUCAF	*05:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
32	9297		*17:01	*17:03	+	-	+	W	-	-	-	-	<u> </u>	-	-	-	-	-	-
33		MT14B	*03:04		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
34	9104		*12:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
35		SSTO	*05:01	*04.04	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
36		KT17	*03:03	*04:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
37		HHKB	*07:02		-	-	-	-	-	-	-	-	_	-	-	-	-	-	-
38	9099		*03:03	+07.04	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
39	9315		*02:02	*07:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
40		WHONP199	*01:02	*06:02	_	-	-	-	-	-	-	-	<u> </u>	-	-	-	-	-	-
41		H0301	*08:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
42		TAB089	*01:02	*00 0 °	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
43		T7526	*01:02	*08:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
44	9057		*12:03	447.01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
45		SHJO	*06:02	*17:01	+	-	-	+	-	-	-	-	-	-	-	-	-	-	-
46		SCHU	*07:02	*45.55	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+
47		TUBO	*07:04	*15:02	-	-	-	-	-	-	-	-	_	-	-	-	-	-	+
48	9303	TER-ND	*04:01	*16:01	-	-	-	-	-	-	-	-	_	-	-	-	-	-	-

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.

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

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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, 12 and 13 were 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'-primer, the 3'-primer was not possible to test. In primer solutions 9 to 11 and 14 it was only possible to test the 3'-primer, the 5'-primer was not possible to test. One additional 3'-primer in primer solution 1 was tested by separately adding one 5'-primer.

OLERUPSSP® HLA-C*17

Product Insert

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

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

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Lot No.: 22V Lot-specific information

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