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Lot No.: 48V Lot-specific information www.olerup-ssp.com

Olerup SSP® HLA-A*24:09N

Product number: 101.841-12 – including *Taq* polymerase

101.841-12u – without *Taq* polymerase

Lot number: 48V

Expiry date: 2016-August-01

Number of tests: 12 Number of wells per test: 2+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. 48V.

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-A*24:09N Lot (98R)

A well containing Negative Control primer pairs has been added.

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

One well has been added to HLA-A*24:09N, well 3.

The HLA-A*24:09N specificity and interpretation tables have been updated compared the previous *Olerup* SSP® HLA-A*24:09N lot **(Lot No. 98R)**.

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
3	New	New	Negative Control.

Change in revision R01 compared to R00:

1. Primer mix 1 does not amplify the A*24:09N allele. This has been corrected in the Specificity and Interpretation Tables. Thus, this lot of the HLA-A*24:09N kit cannot distinguish the A*24:02:01:01-24:02:83 alleles and the A*24:09N allele.

Change in revision R02 compared to R01:

1. Primer mix 1 weakly amplifies the A*24:09N allele. This has been corrected in the Specificity and Interpretation Tables.



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Well **3** 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-A*24:09N SSP subtyping

CONTENT

The primer set contains 5'- and 3'-primers for identifying the HLA-A*24:09N allele.

PLATE LAYOUT

Each test consists of 3 PCR reactions in an 8 well cut PCR plate. Wells 4 to 8 are empty.

1 2 NC empty empty empty empty empty

The 8 well cut PCR plate is marked with the Lot No. '48V' in silver/gray ink.

Well No. 1 is marked with the Lot No. '48V'.

Wells 1 to 2 – HLA-A*24:09N high resolution primers.

Well 3 – 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

The interpretation of HLA-A*24:09N SSP subtypings will be influenced by the $A*02:17:01^w-02:17:03^w$, A*23:14:01-23:14:02, the A*11:139, most A*24, the A*26:16, A*33:19 and the A*68:45 alleles.

UNIQUELY IDENTIFIED ALLELES

The HLA-A*24:09N allele will give rise to a unique amplification pattern by the primers in the HLA-A*24:09N kit^{1,2}.

¹HLA-A alleles listed on the IMGT/HLA web page 2014-January-17, release 3.15.0, 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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SPECIFICITY TABLE

HLA-A*24:09N SSP subtyping

Specificities and sizes of the PCR products of the 2+1 primer mixes used for HLA-A*24:09N SSP subtyping

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	Amplified HLA-A alleles ³
14	105 bp	800 bp	*24:09N ^w
2	175 bp, 205 bp	1070 bp	*02:17:01**-02:17:03**, 11:139, 23:14:01-23:14:02, 24:02:01:01-24:11N, 24:13:01-24:13:02, 24:17-24:50, 24:54-24:56, 24:58-24:63, 24:66-24:91, 24:93, 24:95-24:113, 24:115-24:137, 24:139-24:187, 24:189-24:210, 24:212-24:221, 24:223-24:227, 24:229-24:270, 26:16, 33:19, 68:45
3 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-A*24:09N SSP typings.

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 preheated.

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 mix 3 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', may be weakly amplified.

Product Insert

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101.841-12 – including *Taq* polymerase, IFU-01 **101.841-12u – without** *Taq* polymerase, IFU-02

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PRIMER SPECIFICATION

Well No.	1	2
Length of spec.	105	175
PCR product		205
Length of int.	800	1070
pos. control ¹		
5'-primer(s) ²	678	98
	^{5'} -AgA ^{3'}	^{5'} -CTC ^{3'}
		368
		^{5'} -gTT ^{3'}
3'-primer(s) ³	742	259
	^{5'} -CTA ^{3'}	^{5'} -gTT ^{3'}
		502
		^{5'} -CTT ^{3'}
		539
		^{5'} -TCT ^{3'}
Well No.	1	2

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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	No.: 4			Lot-		
CE	LL L	INE VAL	IDATIO	ON SH	ΙΕΙ	ΕT
	H	ILA-A24:0	9N SSP	kit ²		
					W	ell
					1	2
				ot No.:	201214701	201214702
	IHW	/C cell line ¹	A*	A*	<u>``</u>	.,
1	9001		*24:02		-	+
2		LK707	*02:01		-	Ė
3		E4181324	*01:01		-	-
4		GU373	*30:01		-	-
5		KAS011	*01:01		-	-
6	9353		*02:01	*26:03	-	-
7	9020		*26:01		-	-
8	9025		*31:01		-	-
9		YAR	*26:01		-	-
10		LKT3	*24:02		-	+
11		PITOUT	*29:02		-	-
12	9052		*02:01		-	-
13		JESTHOM	*02:01		-	-
14		OLGA	*31:01		-	-
15	9075		*24:02		-	+
16		SWEIG007	*29:02		-	Ė
17		CTM3953540	*03:01	*80:01	-	-
18		32367	*33:03	*74:01	-	-
19		BM16	*02:01	74.01	-	-
20		SLE005	*02:01		-	-
21		AMALA	*02:17		-	w
22		KOSE	*02:01		-	-
23	9124		*02:01	*34:01	-	-
24		JBUSH	*32:01	01.01	-	-
25		IBW9	*33:01		-	-
26		WT49	*02:05		-	-
27		CH1007	*24:10	*29:01	-	+
28		BEL5GB	*02:01	*29:02	_	Ė
29		MOU	*29:02	23.02		-
30		RSH	*30:01	*68:02	-	-
31		DUCAF	*30:02	00.02	-	-
32		HAG	*02:01			-
33		MT14B	*31:01		-	-
34		DHIF	*31:01			Ē
35		SSTO	*32:01		-	-
36		KT17	*02:06	*11:01	-	-
37		HHKB	*03:01	11.01	-	-
38	9099		*02:17		-	w
39	9315		*01:01	*03:01		-
40		WHONP199	*02:07	*30:01	-	-
41		H0301	*03:01	30.01	-	-
41		TAB089				-
42			*02:07	*02:07	ļ-	Ė
_		T7526	*02:06	*02:07	Ē	Ė
44	9057		*66:01	*24.02	-	-
45		SHJO	*23:01	*24:02	-	+
46		SCHU	*03:01	*02.04	-	-
47		TUBO	*02:16	*03:01	-	-
48	9303	TER-ND	*02:01	*11:01		

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





Product Insert

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101.841-12 – including *Taq* **polymerase**, IFU-01 **101.841-12u – without** *Taq* **polymerase**, IFU-02

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DNAs and where applicable, additional cell line DNAs

No DNAs carrying the alleles to be amplified by primer solution 1 were available. In primer solution 1 it was only possible to test the 5'-primer by separately adding one additional 3'-primer, the 3'-primer was not possible to test. Additional primers in primer mix 2 were tested by separately adding one additional 5'-primer, respectively one additional 3'-primer.

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Addresses:

Manufacturer:

Olerup SSP AB, Franzengatan 5, SE-112 51 Stockholm, Sweden.

Tel: +46-8-717 88 27 **Fax:** +46-8-717 88 18

E-mail: info-ssp@olerup.com

Web page: http://www.olerup-ssp.com

Distributed by:

Olerup GmbH, Löwengasse 47 / 6, AT-1030 Vienna, Austria.

Tel: +43-1-710 15 00 **Fax:** +43-1-710 15 00 10

E-mail: support-at@olerup.com **Web page:** http://www.olerup.com

Olerup Inc., 901 S. Bolmar St., Suite R, West Chester, PA 19382

Tel: 1-877-OLERUP1 **Fax:** 610-344-7989

E-mail: info.us@olerup.com

Web page: http://www.olerup.com

For information on Olerup SSP distributors worldwide, contact Olerup GmbH.