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Lot No.: 72R Lot-specific information

Olerup SSP® DRB1*10

Product number: 101.129-06 – including *Taq* polymerase

101.129-06u - without *Taq* polymerase

Lot number: 72R

Expiry date: 2015-July-01

Number of tests: 6 Number of wells per test: 5

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. 72R.

Changes compared to the previous *OLERUP* SSP® DRB1*10 Lot (14N)

The DRB1*10 kit is updated to enable separation of:

- Confirmed DRB1*10 alleles as listed in the IMGT/HLA database¹
- Polymorphisms in exons outside of the region encoding the peptide binding domain
- Null and Alternatively expressed alleles

The DRB1*10 specificity and interpretation tables have been updated for the DRB1 alleles described since the previous *Olerup* SSP[®] DRB1*10 lot was made **(Lot No. 14N)**.

The DRB1*10 primer set is unchanged compared to the previous lot.

¹As described in section Uniquely Identified Alleles.

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PRODUCT DESCRIPTION

DRB1*10 SSP subtyping

CONTENT

The primer set contains 5'- and 3'-primers for identifying the DRB1*10:01 to DRB1*10:04 alleles.

PLATE LAYOUT

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

1 2 3 4 5 empty empty empty

The 8 well PCR plate is marked with 'DR10' in silver/gray ink.

Well No. 1 is marked with the Lot No. '72R'.

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 covered with a PCR-compatible foil.

The PCR plates are covered with a PCR-compatible foil.

Please note: When removing each 8 well PCR plate, make sure that the remaining plates stay covered. Use a scalpel or a similar instrument to carefully cut the foil between the plates.

INTERPRETATION

Only DRB1*10 alleles will be amplified by the DRB1*10 subtyping kit. Thus, the interpretation of DRB1*10 subtypings is not influenced by other groups of DRB1 alleles or other DRB genes.

UNIQUELY IDENTIFIED ALLELES

All the DRB1*10 alleles, i.e. **DRB1*10:01 to DRB1*10:04**, recognized by the HLA Nomenclature Committee in October 2012¹ will give rise to unique amplification patterns by the primers in the DRB1*10 subtyping kit.

The DRB1*10 kit enables separation of the confirmed DRB1*10 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 DRB1*10 alleles is listed below.

The DRB1*10 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 DRB1*10 SSP subtyping kit cannot distinguish the silent mutations in the DRB1*10:01:01 to DRB1*10:01:03 alleles.

¹DRB1 alleles listed on the IMGT/HLA web page 2012-October-17, release 3.10.0, www.ebi.ac.uk/imgt/hla.

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ALLELE CONFIRMATION STATUS

Allele	Status ¹
DRB1*10:01:01	Confirmed
DRB1*10:01:02	Unconfirmed
DRB1*10:01:03	Unconfirmed
DRB1*10:02	Unconfirmed
DRB1*10:03	Unconfirmed
DRB1*10:04	Unconfirmed

¹Allele status "confirmed" or "unconfirmed" as listed on the IMGT/HLA web page 2012-October-17, release 3.10.0, <u>www.ebi.ac.uk/imgt/hla</u>.

RESOLUTION IN HOMO- AND HETEROZYGOTES

A total of 6 alleles generate 4 amplification patterns that can be combined in 10 homozygous and heterozygous combinations. 4 of these genotypes do not give rise to unique amplification patterns.

```
++-+- *10:01:01, *10:03 = *10:03, *10:03
++--+ *10:01:01, *10:04 = *10:04, *10:04
```

^{*10:01:01 = *10:01:01-10:01:03}

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Lot No.: 72R Lot-specific information

SPECIFICITY TABLE

DRB1*10 SSP subtyping

Specificities and sizes of the PCR products of the 5 primer mixes used for DRB1*10 SSP subtyping

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	Amplified DRB1*10 alleles
1	210 bp	515 bp	*10:01:01-10:04
2	205 bp	515 bp	*10:01:01-10:01:03, 10:03-10:04
3	205 bp	430 bp	*10:02
4 ³	70 bp	430 bp	*10:03
5	160 bp	430 bp	*10:04

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 DRB1*10 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 respective lengths of the HLA-specific PCR product(s) are given for the alleles amplified by these primer mixes.

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 two different control primer pairs give rise to either an internal positive control band of 430 base pairs, for most wells, or a band of 515 base pairs, for some wells.

Wells number 1 and 2 contain the primer pair giving rise to the longer, 515 bp, internal positive control band in order to help in the correct orientation of the DRB1*10 subtyping and in order to allow kit identification.

In the presence of a specific amplification the intensity of the control band often decreases.

³HLA-specific PCR products shorter than 125 base pairs have a lower intensity and are less sharp than longer PCR products.

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Lot No.: 72R Lot-specific information

INTERPRETATION TABLE						
DRB1*10 SSP subtyping						
Amplification patterns of the DRB1*10 alleles						
	Well					
	1	2	3	4	5	
Length of spec.	210	205	205	70	160	
PCR product						
Length of int.	515	515	430	430	430	
pos. control ¹						
5'-primer ²	30(178)	30(178)	30(178)	30(178)	30(178)	
	^{5'} -gCg ^{3'}					
3'-primer ³	86(346)	86(344)	86(344)	40(206)	70(296)	
	^{5'} -CTC ^{3'}	^{5'} -CAC ^{3'}	^{5'} -CCA ^{3'}	^{5'} -CgA ^{3'}	^{5'} -TCT ^{3'}	
Well No.	1	2	3	4	5	
DRB1 allele ⁴						
*10:01:01 -10:01:03	1	2				
*10:02	1		3			
*10:03	1	2		4		
*10:04	1	2			5	
DRB1 allele						
Well No.	1	2	3	4	5	

¹The internal positive control primer pairs amplify segments of the human growth hormone gene. The two different control primer pairs give rise to either an internal positive control band of 430 base pairs, for most wells, or a band of 515 base pairs, for some wells.

Wells number 1 and 2 contains the primer pair giving rise to the longer, 515 bp, internal positive control band in order to help in the correct orientation of the DRB1*10 subtyping and in order to allow kit identification.

²The codon, and in parenthesis the nucleotide, in the 2nd exon, matching the specificity-determining 3'-end of the primer is given. Codon and 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.

given.

The codon, and in parenthesis the nucleotide, in the 2nd exon, matching the specificity-determining 3'-end of the primer is given in the anti-sense direction. Codon and 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.

⁴DRB1*10 alleles in bold lettering are listed as confirmed alleles on the on the IMGT/HLA web page www.ebi.ac.uk/imgt/hla, release 3.10.0, October 2012.

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IHW C cell line	CELL LINE VALIDATION SHEET									
			DRB1*10 S	SP sul	otyping	j ki				
IHWC cell line							\	_	<u> </u>	
IHW C cell line						1	2	3	4	5
1 9001 SA					Production No.	201183001	201183002	201183003	201183004	201211805
1 9001 SA		IHW	C cell line	DI	RB1					
3 9011 E4181324 *15:02 - - - - - - - - -	1					-	-	-	-	-
3 9011 E4181324 *15:02 - - - - - - - - -	2	9280	LK707	*15:02	*04:05	-	-	-	-	-
5 9009 KAS011 *16:01 -	3	9011	E4181324			-	-	-	-	-
6 9353 SM *04:07 *08:03 -	4	9275	GU373	*03:01		-	-	-	-	-
7 9020 QBL *03:01 - <td< th=""><th>5</th><th>9009</th><th>KAS011</th><th>*16:01</th><th></th><th>-</th><th>-</th><th>-</th><th>-</th><th>-</th></td<>	5	9009	KAS011	*16:01		-	-	-	-	-
8 9025 DEU *04:01 - <td< th=""><th>6</th><th>9353</th><th>SM</th><th></th><th>*08:03</th><th>-</th><th>-</th><th>-</th><th>-</th><th>-</th></td<>	6	9353	SM		*08:03	-	-	-	-	-
9 9026 YAR	7	9020	QBL			-	-	-	-	-
10	8	9025	DEU			-	-	-	-	-
11 9051 PTTOUT *07:01	9	9026	YAR	*04:02		-	-	-	-	-
12 9052 DBB *07:01 - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -	10	9107	LKT3	*04:05		L-	-	_	-	-
13 9004 JESTHOM *01:01 -	11	9051	PITOUT	*07:01		-	-	-	-	-
14 9071 OLGA *08:02 - <	12	9052	DBB	*07:01		-	-	-	-	-
15 9075 DKB *09:01 - <t< th=""><th>13</th><th>9004</th><th>JESTHOM</th><th>*01:01</th><th></th><th>-</th><th>-</th><th>-</th><th>-</th><th>-</th></t<>	13	9004	JESTHOM	*01:01		-	-	-	-	-
16 9037 SWEIGO07 *11:01 -	14	9071	OLGA	*08:02		-	-	-	-	-
17 9282 CTM3953540 *03:01 *13:01 - - - - - - - - - - - - - - - - -	15					-	-	-	-	-
17 9282 CTM3953540 *03:01 *13:01 - - - - - - - - - - - - - - - - -	16	9037	SWEIG007	*11:01		-	-	-	-	-
18 9257 32367 *09:01 *11:01 -	17				*13:01	-	-	-	-	-
20 9059 SLE005 *13:02 -	18	9257	32367	*09:01		-	-	-	-	-
21 9064 AMALA *14:02 22 9056 KOSE *13:02 *14:54 23 9124 IHL *08:03 *14:14 24 9035 JBUSH *11:01 - 25 9049 IBW9 *07:01 - 26 9285 WT49 *03:01 - 27 9191 CH1007 *04:05 *10:01 + + - 28 9320 BEL5GB *04:16 *07:01 - - 29 9050 MOU *07:01 -	19	9038	BM16	*12:01		-	-	-	-	-
21 9064 AMALA *14:02 22 9056 KOSE *13:02 *14:54 23 9124 IHL *08:03 *14:14 24 9035 JBUSH *11:01 - 25 9049 IBW9 *07:01 - 26 9285 WT49 *03:01 - 27 9191 CH1007 *04:05 *10:01 + + - 28 9320 BEL5GB *04:16 *07:01 - - 29 9050 MOU *07:01 -	20	9059	SLE005	*13:02		-	-	-	-	-
23 9124 IHL *08:03 *14:14 -	21	9064	AMALA			-	-	-	-	-
24 9035 JBUSH *11:01 -	22	9056	KOSE			-	-	-	-	-
25 9049 IBW9 *07:01 - <	23	9124	IHL	*08:03	*14:14	-	-	-	-	-
26 9285 WT49 *03:01	24	9035	JBUSH	*11:01		-	-	-	-	-
27 9191 CH1007 *04:05 *10:01 + + -	25	9049	IBW9	*07:01		-	-	-	-	-
28 9320 BEL5GB *04:16 *07:01 - - - - - - - - - - - - - - - - - <th>26</th> <th>9285</th> <th>WT49</th> <th>*03:01</th> <th></th> <th>-</th> <th>-</th> <th>-</th> <th>-</th> <th>-</th>	26	9285	WT49	*03:01		-	-	-	-	-
29 9050 MOU *07:01 - <t< th=""><th>27</th><th>9191</th><th>CH1007</th><th>*04:05</th><th>*10:01</th><th>+</th><th>+</th><th>-</th><th>-</th><th>-</th></t<>	27	9191	CH1007	*04:05	*10:01	+	+	-	-	-
30 9021 RSH *03:02 - <t< th=""><th>28</th><th>9320</th><th>BEL5GB</th><th>*04:16</th><th></th><th>-</th><th>-</th><th>-</th><th>-</th><th>-</th></t<>	28	9320	BEL5GB	*04:16		-	-	-	-	-
31 9019 DUCAF *03:01 -	29	9050	MOU			-	-	-	-	-
31 9019 DUCAF *03:01 -	30	9021	RSH	*03:02		-	-	-	-	-
32 9297 HAG *13:03	31	9019	DUCAF			-	-	-	-	-
33 9098 MT14B *04:04 -						-	-	-	-	-
34 9104 DHIF *11:01 - <		9098	MT14B			-	-	-	-	-
35 9302 SSTO *04:03 - <		9104	DHIF	*11:01		-	-	-	-	-
36 9024 KT17 *04:03 *04:06 -	35	9302	SSTO			-	-	-	-	-
37 9065 HHKB *13:01		9024	KT17		*04:06	-	-	-	-	-
38 9099 LZL *14:02 - <t< th=""><th></th><th></th><th></th><th></th><th></th><th>-</th><th>-</th><th>-</th><th>-</th><th>-</th></t<>						-	-	-	-	-
39 9315 CML *03:01 *04:01 -						-	-	-	-	-
40 9134 WHONP199 *07:01 *09:01 - - - - 41 9055 H0301 *13:02 -	39	9315	CML		*04:01	-	-	-	-	-
41 9055 H0301 *13:02 - 42 9066 TAB089 *08:03 - 43 9076 T7526 *09:01 - 44 9057 TEM *14:01 - 45 9239 SHJO *07:01 -		9134	WHONP199	*07:01		-	-	-	-	-
43 9076 T7526 *09:01 - - 44 9057 TEM *14:01 - - - - - - - 45 9239 SHJO *07:01 - - - -	41	9055	H0301	_		-	-	-	-	-
43 9076 T7526 *09:01 - - 44 9057 TEM *14:01 - - - - - - - 45 9239 SHJO *07:01 - - - -						-	-	-	-	-
44 9057 TEM *14:01 -						-	-	-	-	-
45 9239 SHJO *07:01						l -	-	-	-	-
							-	-	-	-
	46			*15:01		-	-	-	-	-
47 9045 TUBO *11:04 *12:01	_				*12:01		-	-	-	-
48 9303 TER-ND *01:03					12.01		-		-	-

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Lot No.: 72R Lot-specific information

CERTIFICATE OF ANALYSIS

Olerup SSP® DRB1*10 SSP

Product number: 101.129-06 – including Taq polymerase

101.129-06u - without *Taq* polymerase

Lot number: 72R

Expiry date: 2015-July-01

Number of tests: 6 Number of wells per test: 5

Well specifications:

Well No.	Production No.
1	2011-830-01
2	2011-830-02
3	2011-830-03
4	2011-830-04
5	2012-118-05

The specificity of each primer solution of the kit has been tested against 48 well characterized IHWC cell line DNAs.

No DNAs carrying the allele to be amplified by primer solution 3 to 5 were available. The specificities of the primers in primer solutions 3 to 5 were tested by separately adding additional 5'-primers, respectively additional 3'-primers.

Results: No false positive or false negative amplifications were obtained.

Date of approval: 2013-January-31

Approved by:

Production Quality Control

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Lot No.: **72R** Lot-specific information

Declaration of Conformity

Product name: Olerup SSP® DRB1*10

Product number: 101.129-06/06u

Lot number: 72R

Intended use: DRB1*10 high resolution histocompatibility testing

Manufacturer: Olerup SSP AB

Franzengatan 5

SE-112 51 Stockholm, Sweden

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

We, *Olerup* SSP AB, hereby declare that this product, to which this Declaration of Conformity relates is in conformity with the following Standard(s) and other normative document(s) ISO 9001:2008 and ISO 13485:2003, following the provisions of the 98/79/EC Directive on *in vitro* diagnostic medical devices, Annex II List B, conformity assessed using Annex IV, as transposed into the national laws of the Member States of the European Union.

The Technical Documentation File is maintained at *Olerup* SSP AB, Franzengatan 5, SE-112 51 Stockholm, Sweden.

Notified Body: Lloyd's Register Quality Assurance Limited, Hiramford, Middlemarch Office Village, Siskin Drive, Coventry CV3 4FJ, United Kingdom. (Notified Body number: 0088.)

Stockholm, Sweden 2013-January-31

Ann-Cathrin Jareman Head of QA and Regulatory Affairs

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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.