Lot-specific Information

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# Olerup SSP<sup>®</sup> DRB1\*09

Product number:	101.120-06 – including <i>Taq</i> polymerase
Lot number:	07G
Expiry date:	2011-July-01
Number of tests:	6
Number of wells per test:	8
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. 07G.

## CHANGES COMPARED TO THE PREVIOUS OLERUP SSP® DRB1\*09 LOT

The DRB1\*09 specificity and interpretation tables have been updated for the DRB alleles described since the previous *Olerup* SSP<sup>®</sup> DRB1\*09 lot was made **(Lot No. 18F)**.

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

Well	5'-primer	3'-primer	rationale
3	Exchanged	Exchanged	New primer pair for the DRB1*090201 and *090202 alleles.
5	-	Added	New primer added for the DRB1*0908 allele.

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# PRODUCT DESCRIPTION

# DRB1\*09 SSP subtyping

## CONTENT

The primer set contains 5'- and 3'-primers for identifying the DRB1\*0901 to DRB1\*0908 alleles.

## **PLATE LAYOUT**

Each test consists of 8 PCR reactions in an 8 well PCR plate.

1	2	3	4	5	6	7	8
							/

The 8 well PCR plate is marked with 'DR9' in silver/gray ink.

Well No. 1 is marked with the Lot No. '07G'.

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 the DRB1\*09 alleles will be amplified by the DRB1\*09 subtyping kit. Thus, the interpretation of DRB1\*09 subtypings is not influenced by other groups of DRB1 alleles or other DRB genes.

### UNIQUELY IDENTIFIED ALLELES

All the DRB1\*09 alleles, i.e. **DRB1\*0901 to DRB1\*0908**, recognized by the HLA Nomenclature Committee in May 2009<sup>1</sup> will give rise to unique amplification patterns by the primers in the DRB1\*09 subtyping kit.

The DRB1\*09 SSP subtyping kit cannot separate the DRB1\*090102 to DRB1\*090104 or the DRB1\*090201 and DRB1\*090202 alleles.

<sup>1</sup>DRB alleles listed on the IMGT/HLA web page 2009-May-11, release 2.25.2, <u>www.ebi.ac.uk/imgt/hla</u>.

### **RESOLUTION IN HOMO- AND HETEROZYGOTES**

The 8 DRB1\*09 alleles can be combined in 36 homozygous and heterozygous combinations. Eighteen of these genotypes do not give rise to unique amplification patterns.

++-++--- 0903,0904 = 0903,0908 ++--+-- 0901,0903 = 0903,0903 ++--++- 0904,0905 = 0905,0908 ++--+-+ 0904,0907 = 0906,0908 ++--+-- 0901,0904 = 0901,0908 = 0904,0904 = 0904,0908 ++---+- 0901,0906 = 0906,0906 ++---++ 0901,0907 = 0907,0907

0901 = 090102 - 090104, 0902 = 090201 - 090202



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# SPECIFICITY TABLE

# DRB1\*09 SSP subtyping

Specificities and sizes of the PCR products of the 8 primer mixes used for DRB1\*09 SSP subtyping

Primer Mix	Size of spec. PCR product <sup>1</sup>	Size of control band <sup>2</sup>	Amplified DRB1*09 alleles
1	195 bp	515 bp	090102-0904, 0906-0908
<b>2</b> <sup>3</sup>	130 bp	430 bp	090102-090104, 0903-0907
3	185 bp	430 bp	090201, 090202
4	215 bp	430 bp	0903
5 <sup>4</sup>	185, 215 bp	430 bp	0904, 0908
6	250 bp	430 bp	0905
7	220 bp	430 bp	0906
8	175 bp	430 bp	0907

<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 DRB1\*09 SSP typings.

When the primers in a primer mix can give rise to specific PCR products of more than one length this is indicated if the size difference is 20 base pairs or more. Size differences shorter than 20 base pairs are not given. For high resolution SSP kits the respective lengths of the specific PCR product(s) of the alleles amplified by these primer mixes are given.

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.

<sup>2</sup>The 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.

Well number 1 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\*09 subtyping.

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

<sup>3</sup>Specific PCR fragments shorter than 150 base pairs have a lower intensity than longer PCR bands.

<sup>4</sup>Primer mix 5: Specific PCR fragment of 185 bp in the DRB1\*0908 allele. Specific PCR fragment of 215 bp in the DRB1\*0904 allele.

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INTERPRETATION TABLE										
DRB1*09 SSP subtyping										
Amplification patterns of the DRB1*09 alleles										
		Well <sup>5</sup>								
	1									
Length of spec.	195	130	185	215	185	250	220	175		
PCR product					215					
Length of int.	515	430	430	430	430	430	430	430		
pos. control <sup>1</sup>										
5'-primer <sup>2</sup>	26(164)	28(170)	9(112)	9(112)	9(112)	9(112)	26(165)	13(125)		
	<sup>5'</sup> -gTA <sup>3'</sup>	<sup>5'</sup> -gCA <sup>3'</sup>	<sup>5'</sup> -TgA <sup>3'</sup>	<sup>5'</sup> -TgA <sup>3'</sup>	<sup>5'</sup> -TgA <sup>3'</sup>	<sup>5'</sup> -TgA <sup>3'</sup>	<sup>5'</sup> -TAT <sup>3'</sup>	<sup>5'</sup> -gTA <sup>3'</sup>		
3'-primer(s) <sup>3</sup>	78(319)	57(257)	57(257)	67(286)	57(256)	78(319)	86(344)	57(257)		
	<sup>5'</sup> -CAC <sup>3'</sup>	<sup>5'</sup> -CgA <sup>3'</sup>	<sup>5'</sup> -CAT <sup>3'</sup>	<sup>5'</sup> -gAA <sup>3'</sup>	<sup>5'</sup> -gCT <sup>3'</sup>	<sup>5'</sup> -gTA <sup>3'</sup>	<sup>5'</sup> -CCA <sup>3'</sup>	<sup>5'</sup> -CgA <sup>3'</sup>		
			57(258)		67(286)					
			<sup>5'</sup> -gCg <sup>3'</sup>		<sup>5'</sup> -gAg <sup>3'</sup>					
Well No.	1	2	3	4	5	6	7	8		
DRB1 allele <sup>4</sup>										
*090102-090104	1	2								
*090201, 090202	1		3							
*0903	1	2		4						
*0904	1	2			5					
*0905		2				6				
*0906	1	2					7			
*0907	1	2						8		
*0908	1				5					
DRB1 allele <sup>4</sup>										
Well No.	1	2	3	4	5	6	7	8		

<sup>1</sup>The 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.

Well number 1 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\*09 subtyping.

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

<sup>2</sup>The codon, and in parenthesis the nucleotide, matching the specificity-determining 3'-end of the primer is given. Codon and nucleotide numbering as on the <u>www.ebi.ac.uk/imgt/hla</u> web site. The sequence of the 3 terminal nucleotides of the primer is shown.

<sup>3</sup>The codon, and in parenthesis the nucleotide, in the 2<sup>nd</sup> and 3<sup>rd</sup> exon, matching the specificitydetermining 3'-end of the primer is given in the anti-sense direction. Codon and nucleotide numbering as on the <u>www.ebi.ac.uk/imgt/hla</u> web site. The sequence of the 3 terminal nucleotides of the primer is shown.

<sup>4</sup>The nucleotide sequence of the DRB1\*090101 allele has been shown to contain errors and be identical to DRB1\*090102.

<sup>5</sup>Primer mix 5: Specific PCR fragment of 185 bp in the DRB1\*0908 allele. Specific PCR fragment of 215 bp in the DRB1\*0904 allele.

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	CELL LINE VALIDATION SHEET											
	DRB1*09 SSP subtyping kit											
					Well							
					1	2	3	4	5	6	7	8
					Ξ	N	e	4	ß	9	5	8
				Production Vo.	200616901	200616902	200961603	200616904	200961605	200961606	200616907	200852208
				quo	61	6	96	61	96	96	61	85.
				Pro.	50	8	20	8	8	8	8	200
	інм	C cell line		<u> </u>								
1	9001		*0101		-	-	-	-	-	-	-	-
2		LK707	*1502	*0405	-	-	-	-	-	-	-	-
3		E4181324	*1502	0.00	-	-	-	-	-	-	-	-
4		GU373	*0301		-	-	-	-	-	-	-	-
5		KAS011	*1601		-	-	-	-	-	-	-	-
6	9353		*0407	*0803	-	-	-	-	-	-	-	-
7	9020	QBL	*0301		-	-	-	-	-	-	-	-
8	9025	DEU	*0401		-	-	-	-	-	-	-	-
9	9026	YAR	*0402		-	-	-	-	-	-	-	-
10	9107	LKT3	*0405		-	-	-	-	-	-	-	-
11	9051	PITOUT	*0701		-	-	-	-	-	-	-	-
12	9052	DBB	*0701		-	-	-	-	-	-	-	-
13	9004	JESTHOM	*0101		-	-	-	-	-	-	-	-
14		OLGA	*0802		-	-	-	-	-	-	-	-
15	9075	DKB	*0901		+	+	-	-	-	-	-	-
16	9037	SWEIG007	*1101		-	-	-	-	-	-	-	-
17		CTM3953540	*0301	*1301	-	-	-	-	-	-	-	-
18		32367	*0901	*1101	+	+	-	-	-	-	-	-
19		BM16	*1201		-	-	-	-	-	-	-	-
20		SLE005	*1302		-	-	-	-	-	-	-	-
21		AMALA	*1402		-	-	-	-	-	-	-	-
22		KOSE	*1302	*1401	-	-	-	-	-	-	-	-
23	9124		*0803	*1414	-	-	-	-	-	-	-	-
24		JBUSH	*1101		-	-	-	-	-	-	-	-
25		IBW9	*0701		-	-	-	-	-	-	-	-
26		WT49	*0301	*4004	-	-	-	-	-	-	-	-
27		CH1007	*0405	*1001	-	-	-	-	-	-	-	-
28		BEL5GB	*0416	*0701	-	-	-	-	-	-	-	-
29	9050	MOU	*0701		-	-	-	-	-	-	-	-
30 31		-	*0302		-	-	-	-	-	-	-	-
32		DUCAF HAG	*0301		-	-	-	-	-	-	-	-
33		MT14B	*0404		-	-	-	-	-	-	-	-
34	9104		*1101		-	-	-	-	-	-	-	-
35		SSTO	*0403		-	-	-	-	-	-	-	-
36		KT17	*0403	*0406	-	-	-	-	-	-	-	-
37		ННКВ	*1301	0400	-	-	-	-	-	-	-	-
38	9099		*1402		-	-	-	-	-	-	-	-
39	9315		*0301	*0401	-	-	-	-	-	-	-	-
40		WHONP199	*0701	*0901	+	+	-	-	-	-	-	-
41		H0301	*1302		-	-	-	-	-	-	-	-
42		TAB089	*0803		-	-	-	-	-	-	-	-
43		T7526	*0901		+	+	-	-	-	-	-	-
44	9057		*1401		-	-	-	-	-	-	-	-
45		SHJO	*0701		-	-	-	-	-	-	-	-
46		SCHU	*1501		-	-	-	-	-	-	-	-
47		TUBO	*1104	*1201	-	-	-	-	-	-	-	-
48		TER-ND	*0103		-	-	-	-	-	-	-	-



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# CERTIFICATE OF ANALYSIS

# **Olerup SSP<sup>®</sup> DRB1\*09 SSP**

Product number: 101.120-06 – including *Taq* polymerase Lot number: 07G 2011-July-01 Expiry date: Number of tests: 6 Number of wells per test: 8

### Well specifications:

Well No.	Production No.					
1	2006-169-01					
2	2006-169-02					
3	2009-616-03					
4	2006-169-04					
5	2009-616-05					
6	2009-616-06					
7	2006-169-07					
8	2008-522-08					

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

No DNAs carrying the allele to be amplified by primer solutions 3 to 8 were available. The specificities of the primers in primer solutions 3 to 8 were tested by separately adding one additional 5'-primer, respectively one additional 3'primer.

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

Date of approval: 2009-July-20

Approved by:

**Quality Control, Supervisor** 



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# **Declaration of Conformity**

Product name: Product number: Lot number:	<i>Olerup</i> SSP <sup>®</sup> DRB1*09 101.120-06 07G
Intended use:	DRB1*09 high resolution histocompatibility testing
Manufacturer:	<i>Olerup</i> SSP AB Hasselstigen 1 SE-133 33 Saltsjöbaden, Sweden <i>Phone:</i> +46-8-717 88 27 <i>Fax:</i> +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 9001:2000 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, Hasselstigen 1, SE-133 33 Saltsjöbaden, Sweden.

The Authorized Representative located within the Community is: *Olerup* SSP AB.

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

Saltsjöbaden, Sweden 2009-July-20

Olle Olerup Managing Director



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