

101.111-24/06 – including *Taq* pol., IFU-01
101.111-24u/06u – without *Taq* pol., IFU-02

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“Instructions for Use” (IFU)

Lot No.: **9E8**

Lot-specific information

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Olerup SSP[®] DRB1*01

Product number:	101.111-24/06 – including <i>Taq</i> pol. 101.111-24u/06u – without <i>Taq</i> pol.
Lot number:	9E8
Expiry date:	2019-09-01
Number of tests:	24 test – Product No. 101.111-24/24u 6 tests – Product No. 101.111-06/06u
Number of wells per test:	23+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. 9E8.

Complete product documentation consists of generic Instructions for Use (IFU), lot specific Product Insert, Worksheet and Certificate.

**CHANGES COMPARED TO THE PREVIOUS OLERUP SSP[®]
DRB1*01 LOT (75Y)**

The DRB1*01 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.

¹As described in section Uniquely Identified Alleles.

The DRB1*01 specificity and interpretation tables have been updated for the DRB1 alleles described since the previous *Olerup SSP[®] DRB1*01* lot was made (**Lot No. 75Y**). The kit design is based on IMGT/HLA database 3.26.0.

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.

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Well	5'-primer	3'-primer	rationale
19	-	Added	3'-primer added for the DRB1*01:77 allele.
20	Added	Added	Primer pair added for the DRB1*01:77 allele.

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Well **24** contains Negative Control primer pairs, 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 product	105	200	105	80	75	80	85
5'-primer¹	164	340	440	45	45	43	36
	5'-CAC ^{3'}	5'-Agg ^{3'}	5'-TTA ^{3'}	5'-Tgg ^{3'}	5'-Tgg ^{3'}	5'-Tgg ^{3'}	5'-TAC ^{3'}
							36
							5'-TAT ^{3'}
3'-primer²	231	2nd I	507	59	58	57	47
	5'-TgC ^{3'}	5'-AAA ^{3'}	5'-TTg ^{3'}	5'-CTC ^{3'}	5'-ggC ^{3'}	5'-CTC ^{3'}	5'-ACA ^{3'}
							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 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

DRB1*01 SSP subtyping

CONTENT

The primer set contains 5'- and 3'-primers for identifying the DRB1*01:01 to DRB1*01:78 alleles.

PLATE LAYOUT

Each test consists of 24 PCR reactions in a 24 well PCR plate.

1	2	3	4	5	6	7	8
9	10	11	12	13	14	15	16
17	18	19	20	21	22	23	NC

The 24 well PCR plate is marked with 'DRB1*01' in silver/gray ink.

Well No. 1 is marked with the Lot No. '9E8'.

Wells 1 to 23 – DRB1*01 high resolution primers.

Well 24 – 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 covered with a PCR-compatible foil.

Please note: When removing each 24 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*01 alleles will be amplified by the DRB1*01 subtyping kit, except that the DRB1*13:165 will be amplified by primer mix 17 and DRB1*14:112 will be amplified by primer mix 8. Thus, the interpretation of DRB1*01 subtypings is only marginally influenced by other groups of DRB1 alleles.

UNIQUELY IDENTIFIED ALLELES

All the DRB1*01 alleles, i.e. **DRB1*01:01 to DRB1*01:78**, recognized by the HLA Nomenclature Committee in October 2016^{1,2} will be amplified by the primers in the DRB1*01 subtyping kit.

The DRB1*01 kit enables separation of the confirmed DRB1*01 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*01 alleles is listed below.

The DRB1*01 kit also enables identification of polymorphisms in exons outside of the region encoding the peptide binding domain and of null and alternatively expressed alleles.

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The following DRB1*01 alleles can be distinguished by the different sizes of the HLA-specific PCR product:

Alleles	Primer mix
DRB1*01:40N, 01:67	19

The DRB1*01 primer set cannot distinguish the silent mutations in DRB1*01:01:01-01:01:29 alleles, the DRB1*01:02:01-01:02:05 and 01:02:07-01:02:12 alleles, the DRB1*01:11:01-01:11:02, the DRB1*01:24:01-01:24:02 or the DRB1*01:44:01-01:44:02 alleles.

¹DRB1 alleles listed on the IMGT/HLA web page 2016-October-14, release 3.26.0, 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>.

ALLELE CONFIRMATION STATUS

Allele	Status ¹	Allele	Status ¹	Allele	Status ¹	Allele	Status ¹
DRB1*01:01:01	Confirmed	DRB1*01:02:12	Unconfirmed	DRB1*01:38	Unconfirmed	DRB1*01:76	Unconfirmed
DRB1*01:01:02	Unconfirmed	DRB1*01:03	Confirmed	DRB1*01:39N	Unconfirmed	DRB1*01:77	Unconfirmed
DRB1*01:01:03	Unconfirmed	DRB1*01:04	Confirmed	DRB1*01:40N	Unconfirmed	DRB1*01:78	Unconfirmed
DRB1*01:01:04	Unconfirmed	DRB1*01:05	Unconfirmed	DRB1*01:41	Unconfirmed		
DRB1*01:01:05	Unconfirmed	DRB1*01:06	Unconfirmed	DRB1*01:42	Unconfirmed		
DRB1*01:01:06	Confirmed	DRB1*01:07	Confirmed	DRB1*01:43	Unconfirmed		
DRB1*01:01:07	Confirmed	DRB1*01:08	Unconfirmed	DRB1*01:44:01	Unconfirmed		
DRB1*01:01:08	Confirmed	DRB1*01:09	Unconfirmed	DRB1*01:44:02	Unconfirmed		
DRB1*01:01:09	Unconfirmed	DRB1*01:10	Unconfirmed	DRB1*01:45	Unconfirmed		
DRB1*01:01:10	Confirmed	DRB1*01:11:01	Unconfirmed	DRB1*01:46	Unconfirmed		
DRB1*01:01:11	Unconfirmed	DRB1*01:11:02	Confirmed	DRB1*01:47	Unconfirmed		
DRB1*01:01:12	Confirmed	DRB1*01:12	Unconfirmed	DRB1*01:48	Unconfirmed		
DRB1*01:01:13	Unconfirmed	DRB1*01:13	Unconfirmed	DRB1*01:49	Unconfirmed		
DRB1*01:01:14	Confirmed	DRB1*01:14	Unconfirmed	DRB1*01:50	Unconfirmed		
DRB1*01:01:15	Unconfirmed	DRB1*01:15	Confirmed	DRB1*01:51	Unconfirmed		
DRB1*01:01:16	Confirmed	DRB1*01:16	Confirmed	DRB1*01:52N	Unconfirmed		
DRB1*01:01:17	Confirmed	DRB1*01:17	Unconfirmed	DRB1*01:53	Unconfirmed		
DRB1*01:01:18	Confirmed	DRB1*01:18	Unconfirmed	DRB1*01:54	Unconfirmed		
DRB1*01:01:19	Confirmed	DRB1*01:19	Confirmed	DRB1*01:55	Unconfirmed		
DRB1*01:01:20	Unconfirmed	DRB1*01:20:01	Confirmed	DRB1*01:56	Unconfirmed		
DRB1*01:01:21	Unconfirmed	DRB1*01:20:02	Unconfirmed	DRB1*01:57	Unconfirmed		
DRB1*01:01:22	Unconfirmed	DRB1*01:21	Unconfirmed	DRB1*01:58	Unconfirmed		
DRB1*01:01:23	Unconfirmed	DRB1*01:22	Unconfirmed	DRB1*01:59	Unconfirmed		
DRB1*01:01:24	Unconfirmed	DRB1*01:23	Confirmed	DRB1*01:60	Unconfirmed		
DRB1*01:01:25	Unconfirmed	DRB1*01:24:01	Confirmed	DRB1*01:61	Unconfirmed		
DRB1*01:01:26	Unconfirmed	DRB1*01:24:02	Unconfirmed	DRB1*01:62N	Unconfirmed		
DRB1*01:01:27	Confirmed	DRB1*01:25	Unconfirmed	DRB1*01:63	Unconfirmed		
DRB1*01:01:28	Unconfirmed	DRB1*01:26	Confirmed	DRB1*01:64	Unconfirmed		
DRB1*01:01:29	Confirmed	DRB1*01:27	Confirmed	DRB1*01:65:01	Unconfirmed		
DRB1*01:02:01	Confirmed	DRB1*01:28	Unconfirmed	DRB1*01:65:02	Unconfirmed		
DRB1*01:02:02	Unconfirmed	DRB1*01:29:01	Confirmed	DRB1*01:66	Unconfirmed		
DRB1*01:02:03	Unconfirmed	DRB1*01:29:02	Unconfirmed	DRB1*01:67	Unconfirmed		
DRB1*01:02:04	Unconfirmed	DRB1*01:30	Unconfirmed	DRB1*01:68N	Unconfirmed		
DRB1*01:02:05	Unconfirmed	DRB1*01:31	Unconfirmed	DRB1*01:69	Unconfirmed		
DRB1*01:02:06	Confirmed	DRB1*01:32	Confirmed	DRB1*01:70	Unconfirmed		
DRB1*01:02:07	Unconfirmed	DRB1*01:33N	Unconfirmed	DRB1*01:71	Unconfirmed		
DRB1*01:02:08	Unconfirmed	DRB1*01:34	Unconfirmed	DRB1*01:72	Unconfirmed		
DRB1*01:02:09	Confirmed	DRB1*01:35	Confirmed	DRB1*01:73	Unconfirmed		
DRB1*01:02:10	Unconfirmed	DRB1*01:36	Unconfirmed	DRB1*01:74	Confirmed		
DRB1*01:02:11	Confirmed	DRB1*01:37	Unconfirmed	DRB1*01:75	Unconfirmed		

¹Allele status “confirmed” or “unconfirmed” as listed on the IMGT/HLA web page 2016-October-14, release 3.26.0, www.ebi.ac.uk/imgt/hla.

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RESOLUTION IN HOMO- AND HETEROZYGOTES

Results file with resolution in DRB1*01 homo- and heterozygotes is available upon request.

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SPECIFICITY TABLE

DRB1*01 SSP subtyping

Specificities and sizes of the PCR products of the 23+1 primer mixes used for DRB1*01 SSP subtyping

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	Amplified DRB1*01 alleles ³	Other amplified DRB1 alleles ⁴
1⁶	255 bp	515 bp	*01:01:01-01:01:29, 01:03, 01:05, 01:07-01:19, 01:21, 01:22 ^w , 01:24:01-01:25, 01:27-01:32, 01:36-01:41, 01:45, 01:47-01:53, 01:55-01:56, 01:59-01:60, 01:62N, 01:64-01:73, 01:75, 01:77-01:78	
2	235 bp	430 bp	*01:02:01-01:02:05, 01:02:07-01:02:12, 01:20:02, 01:23, 01:26, 01:34, 01:43-01:44:02, 01:46, 01:54, 01:57, 01:74, 01:76	
3	200 bp	515 bp	*01:01:01-01:02:12, 01:04-01:14, 01:16-01:21, 01:22 ^w , 01:25-01:29:01, 01:30-01:32, 01:34-01:38, 01:40N-01:41, 01:43-01:50, 01:52N-01:53, 01:56-01:60, 01:62N-01:78	
4	180 bp 210 bp	430 bp	*01:62N *01:03, 01:10, 01:39N, 01:42, 01:61	
5	210 bp 230 bp	430 bp	*01:29:01, 01:74 *01:04, 01:11:01-01:11:02, 01:16, 01:35, 01:55	
6	135 bp 215 bp	430 bp	*01:05, 01:31 *01:29:01-01:29:02, 01:74	
7	210 bp	430 bp	*01:03, 01:15, 01:17, 01:39N, 01:42, 01:61	
8⁵	115 bp 175 bp 210 bp	430 bp	*01:50 *01:27 *01:07	*14:112
9⁵	110 bp 255 bp	430 bp	*01:08 *01:04, 01:06, 01:20:01-01:20:02, 01:61	
10	210 bp 250 bp	430 bp	*01:06, 01:09, 01:15 *01:52N	
11	140 bp 210 bp	430 bp	*01:18, 01:44:01-01:44:02 *01:10, 01:68N	
12	140 bp 180 bp 215 bp	430 bp	*01:31 *01:26, 01:37 *01:12	
13⁵	85 bp 150 bp 215 bp	430 bp	*01:13 *01:28 *01:23, 01:51	
14	170 bp 205 bp	430 bp	*01:14 *01:23-01:24:02, 01:51, 01:54, 01:68N	
15	150 bp	430 bp	*01:28	

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	220 bp		*01:16, 01:21, 01:55
16	200 bp 230 bp	430 bp	*01:24:01-01:24:02, 01:54 *01:19, 01:21
17⁵	125 bp 230 bp 260 bp	430 bp	*01:30 *01:27 *01:22
18^{5,7}	115 bp 230 bp	430 bp	*01:50 *01:25
19⁵	125 bp 235 bp 270 bp	430 bp	*01:67 *01:77 *01:39N-01:40N
20	205 bp 235 bp	430 bp	*01:33N *01:77
21⁵	90 bp 180 bp 250 bp	430 bp	*01:34 *01:62N *01:32
22	205 bp 250 bp	430 bp	*01:36 *01:52N
23	255 bp	430 bp	*01:02:01-01:02:12, 01:23, 01:26, 01:34-01:35, 01:42-01:44:02, 01:46, 01:54, 01:57-01:58, 01:74, 01:76
24^{6,8}	-	-	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 DRB1*01 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 inherent 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 430 or 515 base pairs respectively, well distribution as outlined in the table. Well number 1 contains the longer, 515 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 DRB1 alleles 1st and/or 3rd 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 DRB1 alleles non-DRB1*01 alleles are amplified by primer mixes 8 and 17.

⁵HLA-specific PCR products shorter than 125 base pairs have a lower intensity and are less sharp than longer PCR products.

⁶Primer mixes 1 and 24 has a tendency of giving rise to primer oligomer formation.

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⁷Primer mix 18 may have tendency of unspecific amplification and may give rise to a lower yield of HLA-specific PCR product than the other DRB1*01 primer mixes.

⁸Primer mix 24 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.

'w', might be weakly amplified.

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PRIMER SPECIFICATION

Well No.	1	2	3	4	5	6	7	8	9	10	11	12
Length of spec.	255	235	200	180	210	135	210	115	110	210	140	140
PCR product				210	230	215		175	255	250	210	180
								210				215
Length of int. pos. control ¹	515	430	515	430	430	430	430	430	430	430	430	430
5'-primer(s) ²	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'	9(115) 5'-ggg 3'	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'
								22(154) 5'-Agg 3'				
								110(417) 5'-CAC 3'				
3'-primer(s) ³	86(344) 5'-CAC 3'	78(321) 5'-CAA 3'	66(286) 5'-gAg 3'	61(270) 5'-TTT 3'	70(296) 5'-TCC 3'	44(220) 5'-CCT 3'	66(286) 5'-gAT 3'	66(286) 5'-gAg 3'	37(197) 5'-CgT 3'	70(298) 5'-CgC 3'	47(227) 5'-ggA 3'	48(230) 5'-CCT 3'
			66(286) 5'-gAg 3'	71(299) 5'-gCT 3'	77(317) 5'-AAT 3'	48(230) 5'-CCT 3'	66(286) 5'-gAT 3'	134(490) 5'-gCC 3'	85(341) 5'-CAA 3'	83(336) 5'-CCC 3'	69(295) 5'-CTA 3'	60(266) 5'-Agg 3'
					77(317) 5'-AgT 3'	70(296) 5'-TCC 3'	74(308) 5'-CCT 3'				70(298) 5'-CTT 3'	72(302) 5'-CCA 3'
						72(303) 5'-gCA 3'						
Well No.	1	2	3	4	5	6	7	8	9	10	11	12

Well No.	13	14	15	16	17	18	19	20	21	22	23
Length of spec.	85	170	150	200	125	115	125	205	90	205	255
PCR product	150	205	220	230	230	230	235	235	180	250	
					260		270		250		
Length of int. pos. control ¹	430	430	430	430	430	430	430	430	430	430	430
5'-primer(s) ²	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'	12(123) 5'-AAC 3'	22(154) 5'-AgT 3'	9(114) 5'-TgA 3'	12(124) 5'-AT 3'	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'	14(129) 5'-gAA 3'
					22(154) 5'-Agg 3'	110(417) 5'-CAC 3'	105(401) 5'-AAA 3'	105(401) 5'-AAA 3'			
					58(262) 5'-CCA 3'						
3'-primer(s) ³	29(175) 5'-gAg 3'	57(257) 5'-CAg 3'	51(239) 5'-CCA 3'	66(286) 5'-gAT 3'	86(344) 5'-CAC 3'	86(344) 5'-CAC 3'	86(344) 5'-CAC 3'	66(286) 5'-gAg 3'	30(178) 5'-gAC 3'	69(293) 5'-gCC 3'	85(341) 5'-CAg 3'
	51(239) 5'-CCA 3'	66(286) 5'-gAT 3'	73(305) 5'-ggC 3'	75(313) 5'-gTT 3'		134(490) 5'-gCC 3'	133(485) 5'-TCT 3'	169(595) 5'-AAA 3'	61(270) 5'-TTT 3'	83(336) 5'-CCC 3'	
	72(303) 5'-gCg 3'	69(295) 5'-CTA 3'	77(319) 5'-CAC 3'	77(319) 5'-CAC 3'			169(595) 5'-AAA 3'		83(337) 5'-CCT 3'		
		72(303) 5'-gCg 3'									
Well No.	13	14	15	16	17	18	19	20	21	22	23

¹The internal positive control primer pairs amplify segments of the human growth hormone gene. The internal positive control bands are 430 or 515 base pairs respectively, well distribution as outlined in the table. Well number 1 contains the longer, 515 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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CELL LINE VALIDATION SHEET																				
DRB1*01 SSP subtyping kit ²																				
				Well																
				1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	
				Production No.	201314001	201314002	201314003	201556004	201436505	201436506	201314007	201314008	201314009	201314010	201556011	201314012	201314013	201778014	201556015	201314016
	IHCW cell line ¹	DRB1																		
1	9001 SA	*01:01		+	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
2	9280 LK707	*15:02	*04:05	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
3	9011 E4181324	*15:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
4	9275 GU373	*03:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
5	9009 KAS011	*16:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
6	9353 SM	*04:07	*08:03	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
7	9020 QBL	*03:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
8	9025 DEU	*04:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
9	9026 YAR	*04:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
10	9107 LKT3	*04:05		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
11	9051 PITOUT	*07:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
12	9052 DBB	*07:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
13	9004 JESTHOM	*01:01		+	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
14	9071 OLGA	*08:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
15	9075 DKB	*09:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
16	9037 SWEIG007	*11:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
17	9282 CTM3953540	*03:01	*13:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
18	9257 32367	*09:01	*11:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
19	9038 BM16	*12:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
20	9059 SLE005	*13:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
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		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
30	9021 RSH	*03:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
31	9019 DUCAF	*03:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
32	9297 HAG	*13:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
33	9098 MT14B	*04:04		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
34	9104 DHIF	*11:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
35	9302 SSTO	*04:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
36	9024 KT17	*04:03	*04:06	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
37	9065 HHKB	*13:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
38	9099 LZL	*14:02		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
39	9315 CML	*03:01	*04:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
40	9134 WHONP199	*07:01	*09:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
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		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
46	9013 SCHU	*15:01		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
47	9045 TUBO	*11:04	*12:01	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
48	9303 TER-ND	*01:03		+	-	-	+	-	-	+	-	-	-	-	-	-	-	-	-	-

101.111-24/06 – including *Taq* pol., IFU-01
101.111-24u/06u – without *Taq* pol., IFU-02

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CELL LINE VALIDATION SHEET											
DRB1*01 SSP subtyping kit²											
				Well							
				17	18	19	20	21	22	23	
				Production No.	201778017	201314018	201778019	201778020	201556021	201556022	201314023
IHWC cell line¹			DRB1								
1	9001	SA	*01:01		-	-	-	-	-	-	-
2	9280	LK707	*15:02	*04:05	-	-	-	-	-	-	-
3	9011	E4181324	*15:02		-	-	-	-	-	-	-
4	9275	GU373	*03:01		-	-	-	-	-	-	-
5	9009	KAS011	*16:01		-	-	-	-	-	-	-
6	9353	SM	*04:07	*08:03	-	-	-	-	-	-	-
7	9020	QBL	*03:01		-	-	-	-	-	-	-
8	9025	DEU	*04:01		-	-	-	-	-	-	-
9	9026	YAR	*04:02		-	-	-	-	-	-	-
10	9107	LKT3	*04:05		-	-	-	-	-	-	-
11	9051	PITOUT	*07:01		-	-	-	-	-	-	-
12	9052	DBB	*07:01		-	-	-	-	-	-	-
13	9004	JESTHOM	*01:01		-	-	-	-	-	-	-
14	9071	OLGA	*08:02		-	-	-	-	-	-	-
15	9075	DKB	*09:01		-	-	-	-	-	-	-
16	9037	SWEIG007	*11:01		-	-	-	-	-	-	-
17	9282	CTM3953540	*03:01	*13:01	-	-	-	-	-	-	-
18	9257	32367	*09:01	*11:01	-	-	-	-	-	-	-
19	9038	BM16	*12:01		-	-	-	-	-	-	-
20	9059	SLE005	*13:02		-	-	-	-	-	-	-
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		-	-	-	-	-	-	-
30	9021	RSH	*03:02		-	-	-	-	-	-	-
31	9019	DUCAF	*03:01		-	-	-	-	-	-	-
32	9297	HAG	*13:03		-	-	-	-	-	-	-
33	9098	MT14B	*04:04		-	-	-	-	-	-	-
34	9104	DHIF	*11:01		-	-	-	-	-	-	-
35	9302	SSTO	*04:03		-	-	-	-	-	-	-
36	9024	KT17	*04:03	*04:06	-	-	-	-	-	-	-
37	9065	HHKB	*13:01		-	-	-	-	-	-	-
38	9099	LZL	*14:02		-	-	-	-	-	-	-
39	9315	CML	*03:01	*04:01	-	-	-	-	-	-	-
40	9134	WHONP199	*07:01	*09:01	-	-	-	-	-	-	-
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		-	-	-	-	-	-	-
46	9013	SCHU	*15:01		-	-	-	-	-	-	-
47	9045	TUBO	*11:04	*12:01	-	-	-	-	-	-	-
48	9303	TER-ND	*01:03		-	-	-	-	-	-	-

101.111-24/06 – including *Taq* pol., IFU-01
101.111-24u/06u – without *Taq* pol., IFU-02

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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 5, 6 and 8 to 22 were available. The specificities of the primers in primer solutions 5, 6, 9 to 16, 19 and 20 were tested by separately adding one or more additional 5'-primers, respectively one additional 3'-primer. In primer solution 8, 17 and 18 it was only possible to test the 3'-primers, the 5'-primers were not possible to test. In primer solution 21 and 22 it was only possible to test the 5'-primer, the 3'-primers were not possible to test. In primer solutions 4, 6, 10 to 16, 19 and 20 one or two 3'-primers were not possible to test. In primer solution 19 and 20, one 5'-primer was not possible to test. Additional 3'-primers in primer solution 7 were tested by separately adding additional 5'-primers.

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Lot No.: **9E8**

Lot-specific information

www.olerup.com

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