

101.122-24/06 – including *Taq* pol., IFU-01
101.122-24u/06u – without *Taq* pol., IFU-02

Visit www.olerup.com for
“Instructions for Use” (IFU)

Lot No.: **8E8**

Lot-specific information
Olerup SSP® DRB4

Product number:	101.122-24/06 – including <i>Taq</i> pol. 101.122-24/06u – without <i>Taq</i> pol.
Lot number:	8E8
Expiry date:	2019-10-01
Number of tests:	24 test – Product No. 101.122-24 6 tests – Product No. 101.122-06
Number of wells per test:	30+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. 8E8.

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

CHANGES COMPARED TO THE PREVIOUS OLERUP SSP® DRB4 LOT (3D3)

Sixteen wells have been added to DRB4, well **16 to 31**.

The DRB4 primer set, specificity and interpretation tables have been updated for the DRB alleles described since the previous *Olerup SSP®* DRB4 lot was made (**Lot No. 3D3**). The kit design is based on IMGT/HLA database 3.26.0.

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

Well	5'-primer	3'-primer	rationale
6	-	Added	3'-primers added for the DRB4*01:12 and DRB4*01:14 alleles.
12	-	Added	3'-primers added for the DRB4*01:07:02 and DRB4*01:14 alleles.
13	-	Added	3'-primers added for the DRB4*01:12 and DRB4*01:39 alleles.
15	Added	Added	Updated negative control moved to well 31, primer pairs added for the DRB4*01:10, DRB4*01:21 and DRB4*01:30 alleles.
16	New	New	New primer pairs added for the DRB4*01:13 and DRB4*01:17 alleles.

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17	New	New	New primer pairs added for the DRB4*01:17, DRB4*01:21, DRB4*01:22 and DRB4*01:26 alleles.
18	New	New	New primer pairs added for the DRB4*01:29, DRB4*01:30 and DRB4*01:31 alleles.
19	New	New	New primer pairs added for the DRB4*01:27, DRB4*01:28 and DRB4*01:29 alleles.
20	New	New	New primer pairs added for the DRB4*01:22, DRB4*01:23 and DRB4*01:27 alleles.
21	New	New	New primer pairs added for the DRB4*01:20, DRB4*01:36 and DRB4*01:42 alleles.
22	New	New	New primer pairs added for the DRB4*01:33, DRB4*01:38N and DRB4*01:40 alleles.
23	New	New	New primer pairs added for the DRB4*01:24 and DRB4*01:37 alleles.
24	New	New	New primer pairs added for the DRB4*01:32, DRB4*01:35 and DRB4*01:43 alleles.
25	New	New	New primer pairs added for the DRB4*01:15, DRB4*01:35 and DRB4*01:44 alleles.
26	New	New	New primer pairs added for the DRB4*01:18, DRB4*01:25 and DRB4*01:34 alleles.
27	New	New	New primer pairs added for the DRB4*01:19, DRB4*01:34 and DRB4*01:42 alleles.
28	New	New	New primer pairs added for the DRB4*01:16N and DRB4*01:25 alleles.
29	New	New	New primer pair added for the DRB4*01:11 allele.
30	New	New	New primer pair added for the DRB4*01:41 allele.
31	-	-	Updated negative control added from well 15.

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Well **31** 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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Lot-specific information

PRODUCT DESCRIPTION

DRB4 SSP subtyping

CONTENT

The primer set contains 5'- and 3'-primers for identifying the DRB4*01:01:01:01 to DRB4*03:01N alleles.

PLATE LAYOUT

Each test consists of 31 PCR reactions in a 32 well cut PCR plate. Well 32 is empty.

1	2	3	4	5	6	7	8
9	10	11	12	13	14	15	16
17	18	19	20	21	22	23	24
25	26	27	28	29	30	NC	empty

The 16 well cut PCR plate is marked with 'DRB4' in silver/gray ink.

Well No. 1 is marked with the Lot No. '8E8'.

Wells 1 to 30 – DRB4 high resolution primers.

Well 31 – 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 32 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

Due to the sharing of sequence motifs between DRB4 and DRB1 alleles, primer mix 20 will amplify DRB1 alleles¹.

For further details see Specificity Table.

¹The DRB1*15:01:01:01-15:01:30, 15:02:01:01-15:02:17, 15:03:01:01-15:137N and the DRB1*16:01:01-16:05:02, 16:07-16:41N alleles might be faintly amplified by primer mix 7.

UNIQUELY IDENTIFIED ALLELES

All the DRB4 alleles, i.e. **DRB4*01:01:01:01 to DRB4*03:01N**, recognized by the HLA Nomenclature Committee in October 2016^{1,2} will give rise to unique amplification patterns by the primers in the DRB4 subtyping kit.

The following DRB4 alleles can be distinguished by the different sizes of the HLA-specific PCR product:

Alleles	Primer mix
DRB4*01:20, 01:36	21
DRB4*01:38N, 01:40	22

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The DRB4 subtyping kit cannot distinguish the silent mutation in the DRB4*01:03:01:01 and the 01:03:01:03-01:03:04 alleles.

¹DRB4 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>.

RESOLUTION IN HOMO- AND HETEROZYGOTES

Results file with resolution in DRB4 homo- and heterozygotes is available upon request.

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SPECIFICITY TABLE

DRB4 SSP subtyping

Specificities and sizes of the PCR products of the 30+1 primer mixes used for DRB4 SSP subtyping

Primer Mix	Size of spec. PCR product ¹	Size of control band ²	Amplified DRB4 ³ alleles	Other amplified DRB alleles ⁴
1	185 bp	515 bp	*01:01:01:01-01:01:02, 01:03:01:01-01:03:04, 01:05-01:26, 01:28-01:44	
2⁸	140 bp	430 bp	*01:02	
3	130 bp	430 bp	*01:01:01:01, 01:01:02 [?] , 01:04 [?] -01:05 [?] , 01:06, 01:07:01 [?] -01:08 [?] , 01:11 [?] -01:44 [?] , 02:01N, 03:01N	
4	245 bp	515 bp	*01:01:01:01-01:03:01:01, 01:03:01:03-01:04, 01:05 [?] , 01:06-01:44, 02:01N	
5⁷	155 bp	430 bp	*01:03:01:02N	
6⁵	115 bp 200 bp	430 bp	*01:12 *01:04, 01:14	
7^{6,7}	155 bp	430 bp	*01:02-01:03:04, 01:04 [?] -01:05 [?] , 01:07:01 [?] -01:08 [?] , 01:09-01:10, 01:11 [?] -01:36 [?] , 01:37-01:44	
8⁹	290 bp	515 bp	*01:01:01:01, 01:01:02 [?] , 01:04 [?] -01:44 [?] , 02:01N [?] , 03:01N	
9⁸	155 bp	515 bp	*01:05	
10⁵	85 bp	515 bp	*02:01N	
11⁵	110 bp	430 bp	*01:06	
12	210 bp	430 bp	*01:07:01-01:07:02, 01:14	
13⁵	110 bp 215 bp	430 bp	*01:12, 01:39 *01:08	
14	220 bp	430 bp	*01:09	
15	190 bp 245 bp	430 bp	*01:10, 01:30 *01:02 [?] , 01:21	
16⁵	125 bp 265 bp	430 bp	*01:13 *01:02 [?] , 01:17	
17	255 bp	430 bp	*01:02 [?] , 01:17, 01:21-01:22, 01:26	
18	200 bp	430 bp	*01:29-01:31	
19	210 bp	430 bp	*01:27-01:29	
20	225 bp 250 bp	430 bp	*01:23, 01:27 *01:02 [?] , 01:22	DRB1*04:109, DRB1*15:42
21⁵	105 bp 225 bp	430 bp	*01:20 *01:05 [?] , 01:07:01 [?] , 01:36, 01:42	
22⁵	100 bp 230 bp	430 bp	*01:33, 01:40 *01:38N	
23⁵	105 bp 240 bp	430 bp	*01:24 *01:37	
24⁵	75 bp 175 bp	430 bp	*01:32 *01:35, 01:43	

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Well	Size (bp)	Size (bp)	Allele
25 ⁵	75 bp 170 bp	430 bp	*01:44 *01:15, 01:35
26	135 bp 165 bp	430 bp	*01:25 *01:18, 01:34
27	160 bp 220 bp	430 bp	*01:19, 01:34 *01:42
28	140 bp	430 bp	*01:16N, 01:25
29	140 bp	430 bp	*01:11
30	235 bp	430 bp	*01:41
31 ¹⁰	-	-	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 DRB4 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 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. 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. The complete 2nd exon nucleotide sequence of the DRB4*01:05 allele is not known. Thus, it is not known whether the DRB4*01:05 allele will be amplified by primer mix 4 or not.

⁴Due to the sharing of sequence motifs between DRB4 and DRB1 alleles, primer mix 20 will amplify DRB1 alleles.

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

⁶The DRB1*15:01:01:01-15:01:30, 15:02:01:01-15:02:17, 15:03:01:01-15:137N and the DRB1*16:01:01-16:05:02, 16:07-16:41N alleles might be faintly amplified by primer mix 7.

⁷Primer mixes 5 and 7 may have tendencies of unspecific amplifications.

⁸Primer mixes 2 and 9 have a tendency to giving rise to primer oligomer formation.

⁹Primer mix 8 may give rise to a lower yield of HLA-specific PCR product than the other DRB4 primer mixes.

¹⁰Primer mix 31 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.

‘?’, nucleotide sequence information not available for the primer matching sequence.

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PRIMER SPECIFICATION

Well No.	1	2	3	4	5	6	7	8	9	10	11	12
Length of spec. PCR product	185	140	130	245	155	115 200	155	290	155	85	110	210
Length of int. pos. control ¹	515	430	430	515	430	430	430	515	515	515	430	430
5'-primer(s) ²	28(170) 5'-gAT 3'	42(213) 5'-AgT 3'	105(401) 5'-AAA 3'	1 st I 5'-ggg 3'	5(101) 5'-CAA 3'	28(170) 5'-gAT 3'	96(375) 5'-CAA 3'	2 nd I 5'-TgA 3'	42(213) 5'-AgT 3'	28(170) 5'-gAT 3'	111(421) 5'-ACT 3'	28(170) 5'-gAT 3'
3'-primer(s) ³	76(314) 5'-TgT 3'	76(314) 5'-TgC 3'	134(490) 5'-gCT 3'	1 st I 5'-TgC 3'	42(213) 5'-TCA 3'	52(244) 5'-CAT 3'	134(490) 5'-gCC 3'	2 nd I 5'-TTC 3'	80(328) 5'-gTg 3'	42(213) 5'-TCA 3'	134(490) 5'-gCT 3'	83(335) 5'-CgC 3'
						77(317) 5'-AgT 3'						83(337) 5'-CCg 3'
						83(335) 5'-CgC 3'						83(337) 5'-CCT 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	24
Length of spec. PCR product	110 215	220	190 245	125 265	255	200	210	225 250	105 225	100 230	105 240	75 175
Length of int. pos. control ¹	430	430	430	430	430	430	430	430	430	430	430	430
5'-primer(s) ²	28(170) 5'-gAT 3'	96(375) 5'-CAA 3'	17(138) 5'-TTg 3'	11(121) 5'-CTg 3'	11(121) 5'-CTg 3'	29(175) 5'-gAg 3'	28(170) 5'-gAg 3'	17(137) 5'-TTC 3'	28(170) 5'-gAT 3'	28(170) 5'-gAT 3'	28(170) 5'-gAT 3'	28(170) 5'-gAT 3'
			36(195) 5'-gAC 3'	57(259) 5'-ACC 3'	15(131) 5'-gTA 3'	36(195) 5'-gAC 3'	29(175) 5'-gAg 3'	23(155) 5'-gCA 3'				
			37(199) 5'-ACA 3'		17(137) 5'-TTC 3'	36(196) 5'-AgC 3'	29(175) 5'-gAC 3'	28(170) 5'-gAg 3'				
					17(138) 5'-TTg 3'							
3'-primer(s) ³	51(239) 5'-CCA 3'	156(555) 5'-gTA 3'	86(344) 5'-CCA 3'	86(344) 5'-CCA 3'	86(344) 5'-CCA 3'	86(344) 5'-CCA 3'	86(344) 5'-CCA 3'	86(344) 5'-CCA 3'	49(233) 5'-CCA 3'	45(223) 5'-CTT 3'	48(232) 5'-CgT 3'	39(203) 5'-AgT 3'
	52(244) 5'-CAT 3'								88(350) 5'-Agg 3'	48(230) 5'-CCC 3'	94(368) 5'-CTT 3'	71(299) 5'-gCT 3'
	86(346) 5'-CTT 3'								93(365) 5'-gCT 3'	91(361) 5'-CTA 3'		74(309) 5'-ACA 3'
Well No.	13	14	15	16	17	18	19	20	21	22	23	24

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Well No.	25	26	27	28	29	30
Length of spec. PCR product	75	135	160	140	140	235
Length of int. pos. control ¹	430	430	430	430	430	430
5'-primer(s) ²	28(170) 5'-gAT 3'	28(170) 5'-gAT 3'	28(170) 5'-gAT 3'	28(170) 5'-gAT 3'	14(130) 5'-AgC 3'	21(149) 5'-gAT 3'
3'-primer(s) ³	38(202) 5'-gCA 3'	60(266) 5'-AgC 3'	68(290) 5'-CCg 3'	60(266) 5'-AgC 3'	48(230) 5'-CCT 3'	86(344) 5'-CCA 3'
	71(299) 5'-gCT 3'	68(290) 5'-CCg 3'	69(295) 5'-CCC 3'	60(267) 5'-CAC 3'		
	71(301) 5'-CCA 3'	69(295) 5'-CTg 3'	88(350) 5'-Agg 3'			
Well No.	25	26	27	28	29	30

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																
DRB4 SSP kit ²																
					Well											
					1	2	3	4	5	6	7	8	9	10	11	12
				Prod. No.:	201777101	201777102	201777103	201777104	201777105	201777106	201777107	201777108	201777109	201777110	201777111	201777112
					201777113	201777114	201777115	201777116								
	IHWC cell line ¹	DRB4														
1	9001 SA				-	-	-	-	-	-	-	-	-	-	-	-
2	9280 LK707	*01:03			+	-	-	+	-	-	+	-	-	-	-	-
3	9011 E4181324				-	-	-	-	-	-	f	-	-	-	-	-
4	9275 GU373				-	-	-	-	-	-	-	-	-	-	-	-
5	9009 KAS011				-	-	-	-	-	-	f	-	-	-	-	-
6	9353 SM	*01:03			+	-	-	+	-	-	+	-	-	-	-	-
7	9020 QBL				-	-	-	-	-	-	-	-	-	-	-	-
8	9025 DEU	*01:03			+	-	-	+	-	-	+	-	-	-	-	-
9	9026 YAR	*01:03			+	-	-	+	-	-	+	-	-	-	-	-
10	9107 LKT3	*01:03			+	-	-	+	-	-	+	-	-	-	-	-
11	9051 PITOUT	*01:01			+	-	+	+	-	-	-	+	-	-	-	-
12	9052 DBB	*01:03N			+	-	-	-	+	-	+	-	-	-	-	-
13	9004 JESTHOM				-	-	-	-	-	-	-	-	-	-	-	-
14	9071 OLGA				-	-	-	-	-	-	-	-	-	-	-	-
15	9075 DKB	*01:03			+	-	-	+	-	-	+	-	-	-	-	-
16	9037 SWEIG007				-	-	-	-	-	-	-	-	-	-	-	-
17	9282 CTM3953540				-	-	-	-	-	-	-	-	-	-	-	-
18	9257 32367	*01:01			+	-	+	+	-	-	-	+	-	-	-	-
19	9038 BM16				-	-	-	-	-	-	-	-	-	-	-	-
20	9059 SLE005				-	-	-	-	-	-	-	-	-	-	-	-
21	9064 AMALA				-	-	-	-	-	-	-	-	-	-	-	-
22	9056 KOSE				-	-	-	-	-	-	-	-	-	-	-	-
23	9124 IHL				-	-	-	-	-	-	-	-	-	-	-	-
24	9035 JBUSH				-	-	-	-	-	-	-	-	-	-	-	-
25	9049 IBW9	*01:01			+	-	+	+	-	-	-	+	-	-	-	-
26	9285 WT49				-	-	-	-	-	-	-	-	-	-	-	-
27	9191 CH1007	*01:03			+	-	-	+	-	-	+	-	-	-	-	-
28	9320 BEL5GB	*01:01			+	-	+	+	-	-	+	+	-	-	-	-
29	9050 MOU	*01:01			+	-	+	+	-	-	-	+	-	-	-	-
30	9021 RSH				-	-	-	-	-	-	-	-	-	-	-	-
31	9019 DUCAF				-	-	-	-	-	-	-	-	-	-	-	-
32	9297 HAG				-	-	-	-	-	-	-	-	-	-	-	-
33	9098 MT14B	*01:03			+	-	-	+	-	-	+	-	-	-	-	-
34	9104 DHIF				-	-	-	-	-	-	-	-	-	-	-	-
35	9302 SСТО	*01:03			+	-	-	+	-	-	+	-	-	-	-	-
36	9024 KT17	*01:03			+	-	-	+	-	-	+	-	-	-	-	-
37	9065 HHKB				-	-	-	-	-	-	-	-	-	-	-	-
38	9099 LZL				-	-	-	-	-	-	-	-	-	-	-	-
39	9315 CML	*01:02			-	+	-	+	-	-	+	-	-	-	-	-
40	9134 WHONP199	*01:03			+	-	-	+	-	-	+	-	-	-	-	-
41	9055 H0301				-	-	-	-	-	-	-	-	-	-	-	-
42	9066 TAB089				-	-	-	-	-	-	-	-	-	-	-	-
43	9076 T7526	*01:03			+	-	-	+	-	-	+	-	-	-	-	-
44	9057 TEM				-	-	-	-	-	-	-	-	-	-	-	-
45	9239 SHJO	*01:01	*01:03		+	-	+	+	-	-	+	+	-	-	-	-
46	9013 SCHU				-	-	-	-	-	-	f	-	-	-	-	-
47	9045 TUBO				-	-	-	-	-	-	-	-	-	-	-	-
48	9303 TER-ND				-	-	-	-	-	-	-	-	-	-	-	-

101.122-24/06 – including *Taq* pol., IFU-01
101.122-24u/06u – without *Taq* pol., IFU-02

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Lot No.: **8E8**

Lot-specific information

CELL LINE VALIDATION SHEET																		
DRB4 SSP kit ²																		
					Well													
					17	18	19	20	21	22	23	24	25	26	27	28	29	30
				Prod. No.:	201777117	201777118	201777119	201777120	201777121	201777122	201777123	201777124	201777125	201777126	201777127	201777128	201777129	201777130
	IHWC cell line ¹	DRB4																
1	9001 SA				-	-	-	-	-	-	-	-	-	-	-	-	-	-
2	9280 LK707	*01:03			-	-	-	-	-	-	-	-	-	-	-	-	-	-
3	9011 E4181324				-	-	-	-	-	-	-	-	-	-	-	-	-	-
4	9275 GU373				-	-	-	-	-	-	-	-	-	-	-	-	-	-
5	9009 KAS011				-	-	-	-	-	-	-	-	-	-	-	-	-	-
6	9353 SM	*01:03			-	-	-	-	-	-	-	-	-	-	-	-	-	-
7	9020 QBL				-	-	-	-	-	-	-	-	-	-	-	-	-	-
8	9025 DEU	*01:03			-	-	-	-	-	-	-	-	-	-	-	-	-	-
9	9026 YAR	*01:03			-	-	-	-	-	-	-	-	-	-	-	-	-	-
10	9107 LKT3	*01:03			-	-	-	-	-	-	-	-	-	-	-	-	-	-
11	9051 PITOUT	*01:01			-	-	-	-	-	-	-	-	-	-	-	-	-	-
12	9052 DBB	*01:03N			-	-	-	-	-	-	-	-	-	-	-	-	-	-
13	9004 JESTHOM				-	-	-	-	-	-	-	-	-	-	-	-	-	-
14	9071 OLGA				-	-	-	-	-	-	-	-	-	-	-	-	-	-
15	9075 DKB	*01:03			-	-	-	-	-	-	-	-	-	-	-	-	-	-
16	9037 SWEIG007				-	-	-	-	-	-	-	-	-	-	-	-	-	-
17	9282 CTM3953540				-	-	-	-	-	-	-	-	-	-	-	-	-	-
18	9257 32367	*01:01			-	-	-	-	-	-	-	-	-	-	-	-	-	-
19	9038 BM16				-	-	-	-	-	-	-	-	-	-	-	-	-	-
20	9059 SLE005				-	-	-	-	-	-	-	-	-	-	-	-	-	-
21	9064 AMALA				-	-	-	-	-	-	-	-	-	-	-	-	-	-
22	9056 KOSE				-	-	-	-	-	-	-	-	-	-	-	-	-	-
23	9124 IHL				-	-	-	-	-	-	-	-	-	-	-	-	-	-
24	9035 JBUSH				-	-	-	-	-	-	-	-	-	-	-	-	-	-
25	9049 IBW9	*01:01			-	-	-	-	-	-	-	-	-	-	-	-	-	-
26	9285 WT49				-	-	-	-	-	-	-	-	-	-	-	-	-	-
27	9191 CH1007	*01:03			-	-	-	-	-	-	-	-	-	-	-	-	-	-
28	9320 BEL5GB	*01:01			-	-	-	-	-	-	-	-	-	-	-	-	-	-
29	9050 MOU	*01:01			-	-	-	-	-	-	-	-	-	-	-	-	-	-
30	9021 RSH				-	-	-	-	-	-	-	-	-	-	-	-	-	-
31	9019 DUCAF				-	-	-	-	-	-	-	-	-	-	-	-	-	-
32	9297 HAG				-	-	-	-	-	-	-	-	-	-	-	-	-	-
33	9098 MT14B	*01:03			-	-	-	-	-	-	-	-	-	-	-	-	-	-
34	9104 DHIF				-	-	-	-	-	-	-	-	-	-	-	-	-	-
35	9302 SСТО	*01:03			-	-	-	-	-	-	-	-	-	-	-	-	-	-
36	9024 KT17	*01:03			-	-	-	-	-	-	-	-	-	-	-	-	-	-
37	9065 HHKB				-	-	-	-	-	-	-	-	-	-	-	-	-	-
38	9099 LZL				-	-	-	-	-	-	-	-	-	-	-	-	-	-
39	9315 CML	*01:02			-	-	-	-	-	-	-	-	-	-	-	-	-	-
40	9134 WHONP199	*01:03			-	-	-	-	-	-	-	-	-	-	-	-	-	-
41	9055 H0301				-	-	-	-	-	-	-	-	-	-	-	-	-	-
42	9066 TAB089				-	-	-	-	-	-	-	-	-	-	-	-	-	-
43	9076 T7526	*01:03			-	-	-	-	-	-	-	-	-	-	-	-	-	-
44	9057 TEM				-	-	-	-	-	-	-	-	-	-	-	-	-	-
45	9239 SHJO	*01:01	*01:03		-	-	-	-	-	-	-	-	-	-	-	-	-	-
46	9013 SCHU				-	-	-	-	-	-	-	-	-	-	-	-	-	-
47	9045 TUBO				-	-	-	-	-	-	-	-	-	-	-	-	-	-
48	9303 TER-ND				-	-	-	-	-	-	-	-	-	-	-	-	-	-

101.122-24/06 – including *Taq* pol., IFU-01
101.122-24u/06u – without *Taq* pol., IFU-02

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Lot No.: **8E8**

Lot-specific information

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.

† The DRB1*15:01:01:01-15:01:30, 15:02:01:01-15:02:17, 15:03:01:01-15:137N, DRB1*16:01:01-16:05:02, 16:07-16:41N alleles might be weakly amplified by primer mix 7.

No DNAs carrying the alleles to be amplified by primer solutions 6 and 9 to 30 were available. The specificities of the primers in primer solutions 6, 9, 22 and 24 to 26 were tested by separately adding one additional 5'-primer, respectively, one additional 3'-primer. In primer solutions 10, 11, 15 to 20, 29 and 30, it was only possible to test the 3'-primers, the 5'-primers were not possible to test. In primer solutions 12 to 14, 21, 23, 27 and 28 it was only possible to test the 5'-primer, the 3'-primers were not possible to test. In primer solutions 6, 22, 24 to 26 one or two 3'-primers were not possible to test.

One additional 3'-primer in primer solution 1 was tested by adding one additional 5'-primer.

101.122-24/06 – including *Taq* pol., IFU-01
101.122-24u/06u – without *Taq* pol., IFU-02

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Lot No.: 8E8

Lot-specific information

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101.122-24u/06u – without *Taq* pol., IFU-02

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Lot No.: **8E8**

Lot-specific information

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