

101.118-24 – including *Taq* polymerase, IFU-01
101.118-24u – without *Taq* polymerase, IFU-02

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

Lot No.: **58X**

Lot-specific information

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Olerup SSP® DRB1*07

| | |
|---|---|
| Product numbers: | 101.118-24 – including <i>Taq</i> pol. 101.118-24u – without <i>Taq</i> pol. |
| Lot number: | 58X |
| Expiry date: | 2017-September-01 |
| Number of tests: | 24 |
| Number of wells per test: | 15+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. 58X.

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*07 LOT (02V)

The DRB1*07 kit is updated to enable separation of:

- Confirmed DRB1*07 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

A well containing Negative Control primer pairs has been added.

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

¹As described in section Uniquely Identified Alleles.

The DRB1*07 primer set, specificity and interpretation tables have been updated for the DRB1 alleles described since the previous *Olerup SSP®* DRB1*07 lot was made (**Lot No. 02V**). The kit design is based on IMGT/HLA database 3.19.0.

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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 |
|------|-----------|-----------|--|
| 10 | Added | Added | Primer pair added for the DRB1*07:34 allele. |
| 11 | Added | Added | Primer pair added for the DRB1*07:34 allele. |
| 16 | - | - | Updated negative control. |

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Well **16** 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*07 SSP subtyping

CONTENT

The primer set contains 5'- and 3'-primers for identifying the DRB1*07:01 to DRB1*07:35 alleles.

PLATE LAYOUT

Each test consists of 16 PCR reactions in a 16 well cut PCR plate.

| | | | | | | | |
|----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|
| 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 |
| 9 | 10 | 11 | 12 | 13 | 14 | 15 | NC |

The 16 well cut PCR plate is marked with 'DRB1*07' in silver/gray ink.

Well No. 1 is marked with the Lot No. '58X'.

Wells 1 to 15 – DRB1*07 primers.

Well 16 – 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 16 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

The interpretation of DRB1*07 PCR-SSP subtypings will only be influenced the DRB1*04:90 and 09:08 alleles, when present on the other haplotype. Thus, the interpretation of DRB1*07 subtypings is only marginally influenced by other groups of DRB1 alleles.

For further details see Specificity Table.

UNIQUELY IDENTIFIED ALLELES

All the DRB1*07 alleles, i.e. **DRB1*07:01 to DRB1*07:35**, recognized by the HLA Nomenclature Committee in January 2015^{1,2} will be amplified by the primers in the DRB1*07 subtyping kit.

The DRB1*07 kit enables separation of the confirmed DRB1*07 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*07 alleles is listed below.

The DRB1*07 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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Lot No.: **58X**

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The following DRB1*07 alleles can be distinguished by the different sizes of the specific PCR product:

| Alleles | Primer mix |
|-------------------|------------|
| DRB1*07:14, 07:24 | 11 |
| DRB1*07:17, 07:27 | 14 |

The DRB1*07 subtyping kit cannot distinguish the silent mutations in the DRB1*07:01:01:01 to DRB1*07:01:08 alleles.

¹DRB1 alleles listed on the IMGT/HLA web page 2015-January-19, release 3.19.0.

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 |
|-------------------------|---------------------|--------------------|---------------------|------------|---------------------|
| DRB1*07:01:01:01 | Confirmed | DRB1*07:14 | Confirmed | DRB1*07:34 | Unconfirmed |
| DRB1*07:01:01:02 | Unconfirmed | DRB1*07:15 | Unconfirmed | DRB1*07:35 | Unconfirmed |
| DRB1*07:01:02 | Unconfirmed | DRB1*07:16 | Unconfirmed | | |
| DRB1*07:01:03 | Unconfirmed | DRB1*07:17 | Unconfirmed | | |
| DRB1*07:01:04 | Unconfirmed | DRB1*07:18 | Unconfirmed | | |
| DRB1*07:01:05 | Unconfirmed | DRB1*07:19 | Unconfirmed | | |
| DRB1*07:01:06 | Unconfirmed | DRB1*07:20 | Confirmed | | |
| DRB1*07:01:07 | Unconfirmed | DRB1*07:21 | Unconfirmed | | |
| DRB1*07:01:08 | Unconfirmed | DRB1*07:22 | Unconfirmed | | |
| DRB1*07:03 | Confirmed | DRB1*07:23 | Unconfirmed | | |
| DRB1*07:04 | Confirmed | DRB1*07:24 | Unconfirmed | | |
| DRB1*07:05 | Confirmed | DRB1*07:25 | Unconfirmed | | |
| DRB1*07:06 | Unconfirmed | DRB1*07:26N | Confirmed | | |
| DRB1*07:07 | Confirmed | DRB1*07:27 | Unconfirmed | | |
| DRB1*07:08 | Unconfirmed | DRB1*07:28 | Unconfirmed | | |
| DRB1*07:09 | Unconfirmed | DRB1*07:29 | Unconfirmed | | |
| DRB1*07:10N | Unconfirmed | DRB1*07:30 | Unconfirmed | | |
| DRB1*07:11 | Confirmed | DRB1*07:31 | Unconfirmed | | |
| DRB1*07:12 | Confirmed | DRB1*07:32 | Unconfirmed | | |
| DRB1*07:13 | Unconfirmed | DRB1*07:33 | Unconfirmed | | |

¹Allele status “confirmed” or “unconfirmed” as listed on the IMGT/HLA web page 2015-January-19, release 3.19.0, www.ebi.ac.uk/imgt/hla.

RESOLUTION IN HOMO- AND HETEROZYGOTES

Results file with resolution in DRB1*07 homo- and heterozygotes is available upon request.

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Lot No.: **58X**

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SPECIFICITY TABLE

DRB1*07 SSP subtyping

Specificities and sizes of the PCR products of the 15+1 primer mixes used for DRB1*07 SSP subtyping

| Primer Mix | Size of spec. PCR product ¹ | Size of control band ² | Amplified DRB1*07 alleles ³ | Amplified non-DRB1*07 alleles ⁴ |
|------------------|--|-----------------------------------|--|--|
| 1 ⁶ | 235 bp | 515 bp | *07:01:01:01-07:01:08, 07:03-07:21, 07:23-07:24, 07:26N-07:35 | |
| 2 | 185 bp | 430 bp | *07:01:01:01-07:01:08, 07:05-07:10N, 07:12-07:24, 07:26N-07:35 | |
| 3 | 185 bp | 430 bp | *07:03, 07:09, 07:26N | |
| 4 ⁵ | 90 bp 230 bp | 515 bp | *07:10N *07:04, 07:25 | |
| 5 | 190 bp 235 bp | 430 bp | *07:05 *07:11, 07:25 | |
| 6 | 170 bp | 430 bp | *07:06 | |
| 7 | 160 bp | 430 bp | *07:07, 07:28 | |
| 8 | 175 bp | 430 bp | *07:08, 07:26N | |
| 9 ^{5,6} | 105 bp 240 bp | 430 bp | *07:12 *07:22-07:23 | *04:90, 09:08 |
| 10 ⁵ | 120 bp 165 bp 220 bp | 515 bp | *07:34 *07:13 *07:18 | |
| 11 ⁵ | 110 bp 260 bp | 515 bp | *07:14, 07:34 *07:24 | |
| 12 | 210 bp 245 bp | 430 bp | *07:15, 07:21 07:23 | |
| 13 | 210 bp | 430 bp | *07:16, 07:21 | |
| 14 ⁵ | 125 bp 215 bp 260 bp | 430 bp | *07:27 *07:17 *07:20 | |
| 15 | 155 bp 220 bp 260 bp | 430 bp | *07:19, 07:28 *07:18 *07:20 | |
| 16 ⁷ | - | - | Negative Control | |

¹ 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*07 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.

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Lot No.: **58X**

Lot-specific information

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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. 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 sharing of sequence motifs, the DRB1*04:90 and 09:08 alleles are amplified by primer mix 9.

⁵HLA-Specific PCR fragments shorter than 125 base pairs have a lower intensity and are less sharp than longer PCR bands.

⁶Primer mixes 1 and 9 have a tendency to giving rise to primer oligomer formation.

⁷Primer mix 16 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.

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Lot No.: **58X**

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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 | 235 | 185 | 185 | 90 | 190 | 170 | 160 | 175 | 105 | 120 | 110 | 210 |
| | | | | 235 | 235 | | | | 240 | 165 | 260 | 245 |
| | | | | | | | | | | 220 | | |
| Length of int. pos. control ¹ | 515 | 430 | 430 | 515 | 430 | 430 | 430 | 430 | 430 | 515 | 515 | 430 |
| 5'-primer(s) ² | 13(127) 5'-ATA 3' | 29(174) 5'-AgA 3' | 29(173) 5'-AAg 3' | 13(127) 5'-ATA 3' | 13(127) 5'-ATA 3' | 13(127) 5'-ATA 3' | 13(127) 5'-ATA 3' | 29(173) 5'-AAg 3' | 10(116) 5'-gCg 3' | 13(127) 5'-ATA 3' | 13(127) 5'-ATA 3' | 10(116) 5'-gCg 3' |
| | 13(127) 5'-ATA 3' | | 29(174) 5'-AgT 3' | | | | | 36(196) 5'-Agg 3' | 13(127) 5'-gTA 3' | 106(405) 5'-ACT 3' | 106(405) 5'-ACT 3' | 19(145) 5'-ACC 3' |
| | | | 29(175) 5'-gAT 3' | | | | | | 56(256) 5'-CTA 3' | | | 21(149) 5'-gAA 3' |
| 3'-primer(s) ³ | 77(319) 5'-CAC 3' | 77(319) 5'-CAC 3' | 77(319) 5'-CAC 3' | 29(174) 5'-..T 3' | 64(278) 5'-TCC 3' | 57(257) 5'-CAg 3' | 54(248) 5'-gCT 3' | 77(319) 5'-CAC 3' | 77(319) 5'-CAC 3' | 55(251) 5'-gCA 3' | 35(191) 5'-CCC 3' | 77(319) 5'-CAC 3' |
| | 77(319) 5'-gTA 3' | | | 77(317) 5'-AAT 3' | 77(317) 5'-Agg 3' | | 54(250) 5'-CCA 3' | | | 72(304) 5'-gCT 3' | 86(344) 5'-CAA 3' | |
| | | | | 77(319) 5'-CAA 3' | 77(319) 5'-CAA 3' | | | | | 133(485) 5'-TCT 3' | 133(485) 5'-TCT 3' | |
| Well No. | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |

| Well No. | 13 | 14 | 15 |
|---|----------------------|----------------------|----------------------|
| Length of spec. PCR product | 210 | 125 | 155 |
| | | 215 | 220 |
| | | 260 | 260 |
| Length of int. pos. control ¹ | 430 | 430 | 430 |
| 5'-primer(s) ² | 19(145) 5'-ACC 3' | 13(127) 5'-ATA 3' | 13(127) 5'-ATA 3' |
| | 22(154) 5'-Agg 3' | | |
| 3'-primer(s) ³ | 77(319) 5'-CAC 3' | 42(213) 5'-TCC 3' | 49(235) 5'-CAA 3' |
| | | 71(301) 5'-CCA 3' | 54(248) 5'-gCT 3' |
| | | 86(344) 5'-CAT 3' | 72(304) 5'-gCT 3' |
| | | | 86(344) 5'-CAT 3' |
| Well No. | 13 | 14 | 15 |

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.

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Lot No.: **58X**

Lot-specific information

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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*07 SSP subtyping kit ² | | | | | | | | | | | | | | | | | | |
| | | | | Well | | | | | | | | | | | | | | |
| | | | | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | 13 | 14 | 15 |
| | | | | 201444101 | 201444102 | 201444103 | 201444104 | 201444105 | 201444106 | 201444107 | 201444108 | 201444109 | 201444110 | 201444111 | 201444112 | 201444113 | 201444114 | 201444115 |
| | | | Production No. | | | | | | | | | | | | | | | |
| | IHWC 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 PTOUT | *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 | | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - |

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.

101.118-24 – including *Taq* polymerase, IFU-01
101.118-24u – without *Taq* polymerase, IFU-02

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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 3 to 15 were available. The specificities of the primers in primer solutions 4, 5, 6 and 9 were tested by separately adding one additional 5'-primer, respectively one additional 3'-primer. In primer solutions 3, 8, 12 and 13 it was only possible to test the 3'-primers, the 5'-primers were not possible to test. In primer solutions 7, 10, 11 and 14 to 15 it was only possible to test the 5'-primers, the 3'-primers were not possible to test. In primer solutions 4 and 5, two 3'-primers were not possible to test, and in primer solution 9, two 5'-primers were not possible to test. In addition, one 3'-primer in primer solution 1 was tested by separately adding one additional 5'-primer.

101.118-24 – including *Taq* polymerase, IFU-01
101.118-24u – without *Taq* polymerase, IFU-02

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Lot No.: **58X**

Lot-specific information

www.olerup-ssp.com

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**.