**●LERUPSSP®**HLA-C\*15 Add-on Product Insert Page 1 of 12

**101.865-12 – including** *Taq* **polymerase**, IFU-01 **101.865-12u – without** *Taq* **polymerase**, IFU-02

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Lot No.: 91X Lot-specific information

# Olerup SSP® HLA-C\*15 Add-on

Product number: 101.865-12 – including *Taq* polymerase

101.865-12u – without *Taq* polymerase

Lot number: 91X

Expiry date: 2017-July-01

Number of tests: 12 Number of wells per test: 3+1

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. 91X.

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

# Changes compared to the previous *OLERUP* SSP® HLA-C\*15 Add-on Lot (47S)

A well containing Negative Control primer pairs has been added.

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

Two wells have been added to HLA-C\*15 Add-on, wells **3 to 4**.

The HLA-C\*15 Add-on primer set, specificity and interpretation tables have been updated for the HLA-C alleles described since the previous *Olerup* SSP<sup>®</sup> HLA-C\*15 Add-on lot was made (Lot No. 47S). The kit design is based on IMGT/HLA database 3.18.0.

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.

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**101.865-12 – including** *Taq* **polymerase**, IFU-01 **101.865-12u – without** *Taq* **polymerase**, IFU-02

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

| Well | 5'-primer | 3'-primer | rationale   |
|------|-----------|-----------|---|
| 3    | New       | New       | New primer pair for resolution of the C*15:87 allele. |
| 4    | New       | New       | Negative Control.                                     |

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**101.865-12 – including** *Taq* **polymerase**, IFU-01 **101.865-12u – without** *Taq* **polymerase**, IFU-02

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Well **4** contains <u>Negative Control primer pairs</u>, that will amplify more than 95% of the *Olerup* SSP<sup>®</sup> 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          | 105                              | 200                              | 105                              | 80                                | 75                                | 80                               | 85                               |
|------------------------|----------------------------------|----------------------------------|----------------------------------|-----------------------------------|-----------------------------------|----------------------------------|----------------------------------|
| product                |                                  |                                  |                                  |                                   |                                   |                                  |                                  |
| 5'-primer <sup>1</sup> | 164                              | 340                              | 440                              | 45                                | 45                                | 43                               | 36                               |
| -                      | 5'-CAC3'                         | 5'-Agg <sup>3'</sup>             | <sup>5'</sup> -TTA3'             | <sup>5</sup> '-Tgg <sup>3</sup> ' | <sup>5</sup> '-Tgg <sup>3</sup> ' | <sup>5'</sup> -Tgg <sup>3'</sup> | 5'-TAC3'                         |
|                        |                                  |                                  |                                  |                                   |                                   |                                  | 36                               |
|                        |                                  |                                  |                                  |                                   |                                   |                                  | <sup>5'</sup> -TAT <sup>3'</sup> |
| 3'-primer <sup>2</sup> | 231                              | 2 <sup>nd</sup> I                | 507                              | 59                                | 58                                | 57                               | 47                               |
| •                      | <sup>5'</sup> -TgC <sup>3'</sup> | <sup>5'</sup> -AAA <sup>3'</sup> | <sup>5'</sup> -TTg <sup>3'</sup> | 5'-CTC3'                          | <sup>5'</sup> -ggC <sup>3'</sup>  | 5'-CTC3'                         | 5'-ACA3'                         |
|                        |                                  |                                  |                                  |                                   |                                   |                                  | 48                               |
|                        |                                  |                                  |                                  |                                   |                                   |                                  | <sup>5'</sup> -gCA <sup>3'</sup> |
|                        |                                  |                                  |                                  |                                   |                                   |                                  | 48                               |
|                        |                                  |                                  |                                  |                                   |                                   |                                  | <sup>5'</sup> -gCC <sup>3'</sup> |
|                        |                                  |                                  |                                  |                                   |                                   |                                  | 52                               |
|                        |                                  |                                  |                                  |                                   |                                   |                                  | <sup>5'</sup> -TgT <sup>3'</sup> |
| A*                     | +                                | +                                | +                                |                                   |                                   |                                  |                                  |
| B*                     | +                                | +                                | +                                |                                   |                                   |                                  |                                  |
| C*                     | +                                | +                                | +                                |                                   |                                   |                                  |                                  |
| DRB1                   |                                  |                                  |                                  | +                                 | +                                 |                                  |                                  |
| DRB3                   |                                  |                                  |                                  | +                                 | +                                 |                                  |                                  |
| DRB5                   |                                  |                                  |                                  | +                                 |                                   |                                  |                                  |
| DQB1                   |                                  |                                  |                                  |                                   | +                                 |                                  |                                  |
| DPB1                   |                                  |                                  |                                  |                                   |                                   | +                                |                                  |
| DQA1                   |                                  |                                  |                                  |                                   |                                   |                                  | +                                |

<sup>&</sup>lt;sup>1</sup>The nucleotide position for HLA class I genes and the codon for HLA class II genes, in the 2<sup>nd</sup> or 3<sup>rd</sup> exon, matching the specificity-determining 3'-end of the primer is given. Nucleotide and codonnumbering as on the <a href="https://www.ebi.ac.uk/imgt/hla">www.ebi.ac.uk/imgt/hla</a> web site. The sequence of the 3 terminal nucleotides of the primer is given.

<sup>&</sup>lt;sup>2</sup>The nucleotide position for HLA class I genes and the codon for HLA class II genes, in the 2<sup>nd</sup> or 3<sup>rd</sup> exon or the 2<sup>nd</sup> intron, matching the specificity-determining 3'-end of the primer is given in the anti-sense direction. Nucleotide and codon numbering as on the <a href="www.ebi.ac.uk/imgt/hla">www.ebi.ac.uk/imgt/hla</a> web site. The sequence of the 3 terminal nucleotides of the primer is given.

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# PRODUCT DESCRIPTION

# **HLA-C\*15 Add-on SSP subtyping**

#### CONTENT

The primer set contains 5'- and 3'-primers for distinguishing the HLA-C\*15:29 from the C\*15:05 alleles and the C\*15:87 from the C\*15:02 alleles.

#### PLATE LAYOUT

Each test consists of 4 PCR reactions in an 8 well cut PCR plate. Wells 5 to 8 are empty.

#### 1 2 3 NC empty empty empty empty

The 8 well cut PCR plate is marked with the Lot No. '91X' in silver/gray ink.

Well No. 1 is marked with the Lot No. '91X'.

Wells 1 to 3 – HLA-C\*15 Add-on high resolution primers.

Well 4 – 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 heat-sealed with a PCR-compatible foil.

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

#### INTERPRETATION

The interpretation of HLA-C\*15 Add-on SSP subtypings will be influenced by a few other C\*15 and HLA-C alleles. In addition, two HLA-B alleles will be amplified by primer mix 1.

For further details see Specificity Table.

#### **UNIQUELY IDENTIFIED ALLELES**

The HLA-C\* C\*15:02, 15:05, 15:29, and C\*15:87 alleles give different patterns in the HLA-C\*15 Add-on kit<sup>1,2</sup>.

The HLA-C\*15 Add-on kit cannot distinguish the silent mutations in the C\*15:02:01-15:02:20 or the C\*15:05:01-15:05:11 alleles.

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<sup>&</sup>lt;sup>1</sup>Based on HLA-C alleles listed on the IMGT/HLA web page 2014-October-10, release 3.18.0, <a href="https://www.ebi.ac.uk/imgt/hla">www.ebi.ac.uk/imgt/hla</a>.

Alleles 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 <a href="http://hla.alleles.org/alleles/deleted.html">http://hla.alleles.org/alleles/deleted.html</a>.

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# SPECIFICITY TABLE

# **HLA-C\*15 Add-on SSP subtyping**

Specificities and sizes of the PCR products of the 3+1 primer mixes used for HLA-C\*15 Add-on SSP subtyping

| Primer<br>Mix           | Size of spec.<br>PCR<br>product <sup>1</sup> | Size of<br>control<br>band <sup>2</sup> | Amplified HLA-<br>C*15:02/15:05/15:29/15:87<br>alleles <sup>3</sup> | Other amplified HLA-C alleles <sup>4</sup>  |
|-------------------------|--|---|---|---|
| 15                      | 100 bp                                       | 800 bp                                  | 15:05:01-15:05:11, 15:29  | 15:22-15:23, 15:36, 15:46, 15:54, 15:59, 15:69-15:70, 15:90, 04:108, <b>B*35:03:11, B*35:205</b>  |
| 2                       | 545 bp                                       | 1070 bp                                 | 15:29, 15:87  | *01:85, 08:22, 08:56,   |
| 3                       | 315 bp                                       | 800 bp                                  | 15:02:01-15:02:20   | *15:03, 15:07-15:08,<br>15:10:01-15:13, 15:15-<br>15:18, 15:21, 15:26, 15:28,<br>15:31-15:35, 15:37-15:39,<br>15:41-15:45, 15:47-15:53,<br>15:56-15:58, 15:60, 15:62-<br>15:63, 15:67-15:68, 15:71-<br>15:75, 15:78-15:89, 15:91-<br>15:101, 15:103, 02:06,<br>02:47, 03:19, 03:102,<br>04:178, 07:289, 12:15 |
| <b>4</b> <sup>6,7</sup> | -  | -                                       | Negative Control  |   |

<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 HLA-C\*15 Add-on SSP typings. 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 internal positive control bands are 1070 or 800 base pairs respectively, well distribution as outlined in the table. Well number 1 contains the shorter, 800 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.

<sup>3</sup>For several HLA Class I alleles 1<sup>st</sup> and/or 4<sup>th</sup> 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.

<sup>4</sup>The interpretation of HLA-C\*15 Add-on SSP subtypings will be influenced by a few other C\*15 and HLA-C alleles. In addition, two HLA-B alleles will be amplified by primer mix 1.

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

<sup>6</sup>Primer mix <sup>4</sup> has a tendency to giving rise to primer oligomer formation.

<sup>7</sup>Primer mix 4 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.

**Product Insert** 

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**101.865-12 – including** *Taq* **polymerase**, IFU-01 **101.865-12u – without** *Taq* **polymerase**, IFU-02

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### PRIMER SPECIFICATION

| Well No.                  | 1                                | 2                                | 3                                |
|---------------------------|----------------------------------|----------------------------------|----------------------------------|
| Length of spec.           | 100                              | 545                              | 315                              |
| PCR product               |                                  |                                  |                                  |
|                           |                                  |                                  |                                  |
| Length of int.            | 800                              | 1070                             | 800                              |
| pos. control <sup>1</sup> |                                  |                                  |                                  |
| 5'-primer(s) <sup>2</sup> | 420                              | 972                              | 420                              |
|                           | 5' -TTC 3'                       | <sup>5'</sup> -CTA <sup>3'</sup> | <sup>5'</sup> -TTA <sup>3'</sup> |
|                           |                                  |                                  | 420                              |
|                           |                                  |                                  | <sup>5'</sup> -TTA <sup>3'</sup> |
|                           |                                  |                                  |                                  |
| 3'-primer(s) <sup>3</sup> | 477                              | 1034                             | 3 <sup>rd</sup> I                |
|                           | <sup>5'</sup> -gCg <sup>3'</sup> | <sup>5'</sup> -AgT <sup>3'</sup> | <sup>5'</sup> -CTC <sup>3'</sup> |
| Well No.                  | 1                                | 2                                | 3                                |

<sup>1</sup>The internal positive control primer pairs amplify segments of the human growth hormone gene. The internal positive control bands are 1070 or 800 base pairs respectively, well distribution as outlined in the table. Well number 1 contains the shorter, 800 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.

<sup>2</sup>The nucleotide position matching the specificity-determining 3'-end of the primer is given. Nucleotide numbering as on the <a href="www.ebi.ac.uk/imgt/hla">www.ebi.ac.uk/imgt/hla</a> web site. The sequence of the 3 terminal nucleotides of the primer is given.

<sup>3</sup>The nucleotide position matching the specificity-determining 3'-end of the primer is given in the anti-sense direction. Nucleotide numbering as on the <a href="www.ebi.ac.uk/imgt/hla">www.ebi.ac.uk/imgt/hla</a> web site. The sequence of the 3 terminal nucleotides of the primer is given.

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| CELL LINE VALIDATION SHEET           |      |                           |        |          |           |           |           |  |
|--------------------------------------|------|---------------------------|--------|----------|-----------|-----------|-----------|--|
| HLA-C*15 Add-on SSP kit <sup>2</sup> |      |                           |        |          |           |           |           |  |
|                                      |      |                           |        |          |           |           |           |  |
|                                      |      |                           |        |          | 1         | 2         | 3         |  |
|                                      |      |                           |        | Lot No.: | 201296601 | 201296602 | 201547603 |  |
|                                      | IHV  | /C cell line <sup>1</sup> |        | C*       |           | -         |           |  |
| 1                                    | 9001 |                           | *07:02 | 1        | -         | -         | -         |  |
| 2                                    | 9280 | LK707                     | *07:01 | *15:05   | +         | -         | -         |  |
| 3                                    |      | E4181324                  | *12:02 |          | -         | -         | -         |  |
| 4                                    | 9275 | GU373                     | *03:04 | *04:01   | -         | -         | -         |  |
| 5                                    | 9009 | KAS011                    | *06:02 |          | -         | -         | -         |  |
| 6                                    | 9353 | SM                        | *03:04 | *07:02   | -         | -         | -         |  |
| 7                                    | 9020 | QBL                       | *05:01 |          | -         | -         | -         |  |
| 8                                    | 9025 | DEU                       | *04:01 |          | -         | -         | -         |  |
| 9                                    | 9026 | YAR                       | *12:03 |          | -         | -         | -         |  |
| 10                                   | 9107 | LKT3                      | *01:02 |          | -         | -         | -         |  |
| 11                                   | 9051 | PITOUT                    | *16:01 |          | -         | -         | -         |  |
| 12                                   | 9052 | DBB                       | *06:02 |          | -         | -         | -         |  |
| 13                                   | 9004 | JESTHOM                   | *01:02 |          | -         | -         | -         |  |
| 14                                   | 9071 | OLGA                      | *01:02 | *03:04   | -         | -         | -         |  |
| 15                                   | 9075 | DKB                       | *03:04 |          | -         | -         | -         |  |
| 16                                   | 9037 | SWEIG007                  | *02:02 |          | -         | -         | -         |  |
| 17                                   | 9282 | CTM3953540                | *03:03 | *07:01   | -         | -         | -         |  |
| 18                                   | 9257 | 32367                     | *01:02 | *07:05   | -         | -         | -         |  |
| 19                                   |      | BM16                      | *07:01 |          | -         | -         | -         |  |
| 20                                   |      | SLE005                    | *03:04 |          | -         | -         | -         |  |
| 21                                   |      | AMALA                     | *03:03 |          | -         | -         | -         |  |
| 22                                   |      | KOSE                      | *12:03 |          | -         | -         | -         |  |
| 23                                   | 9124 |                           | *01:02 | *15:02   | -         | -         | +         |  |
| 24                                   |      | JBUSH                     | *12:03 |          | -         | -         | -         |  |
| 25                                   |      | IBW9                      | *08:02 |          | -         | -         | -         |  |
| 26                                   |      | WT49                      | *07:01 |          | -         | -         | -         |  |
| 27                                   |      | CH1007                    | *07:04 | *15:29   | +         | +         | -         |  |
| 28                                   |      | BEL5GB                    | *05:01 | *16:01   | -         | -         | -         |  |
| 29                                   | 9050 |                           | *16:01 |          | -         | -         | -         |  |
| 30                                   | 9021 |                           | *17:01 |          | -         | -         | -         |  |
| 31                                   |      | DUCAF                     | *05:01 |          | -         | -         | -         |  |
| 32                                   |      | HAG                       | *17:01 | *17:03   | -         | -         | -         |  |
| 33                                   |      | MT14B                     | *03:04 | 17.00    | -         | -         | -         |  |
| 34                                   | 9104 |                           | *12:03 |          | -         |           | -         |  |
| 35                                   |      | SSTO                      | *05:01 |          | -         |           | -         |  |
| 36                                   |      | KT17                      | *03:03 | *04:01   | -         |           | -         |  |
| 37                                   |      | HHKB                      | *07:02 | UT.U1    | -         |           | -         |  |
| 38                                   | 9099 |                           | *03:03 |          | -         | Ė         | Ē         |  |
| 39                                   | 9315 |                           | *02:02 | *07:01   |           | Ė         | Ė         |  |
| 40                                   |      | WHONP199                  | *01:02 | *06:02   |           | Ė         | Ė         |  |
| 41                                   |      | H0301                     | *08:02 | 00.02    | Ė         | Ť         | Ė         |  |
| 42                                   |      | TAB089                    | *01:02 |          |           | Ė         |           |  |
| 42                                   |      | T7526                     | *01:02 | *08.01   | Ē         | Ē         |           |  |
| 43                                   |      |                           |        | *08:01   | <u> </u>  | Ė         | -         |  |
|                                      | 9057 |                           | *12:03 | *17.04   | -         | Ē         | Ē         |  |
| 45                                   |      | SHJO                      | *06:02 | *17:01   | -         | Ė         | -         |  |
| 46                                   |      | SCHU                      | *07:02 | *45,00   | -         | <u> </u>  | -         |  |
| 47                                   | 9045 | TUBO                      | *07:04 | *15:02   | ı - I     |           | +         |  |

<sup>&</sup>lt;sup>1</sup>The provided cell line HLA specificities are retrieved from the <a href="http://www.ihwg.org/hla">http://www.ihwg.org/hla</a> web site. The specificity of an individual cell line may thus be subject to change.



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<sup>2</sup>The 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. In primer solution 3 one 5'-primers was not possible to test.

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Lot No.: 91X Lot-specific information

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.