DRA Product Insert Page 1 of 8

101.131-24u - without *Taq* polymerase

Lot No.: **52F** Lot-specific information

## Olerup SSP® DRA

Product number: 101.131-24u – without *Taq* polymerase

Lot number: 52F

Expiry date: 2011-February-01

Number of tests: 24 Number of wells per test: 2

Storage - pre-aliquoted primers: dark at -20°C

- PCR Master Mix: -20°C

This Product Description is only valid for Lot No. 52F.

# CHANGES COMPARED TO THE PREVIOUS *OLERUP* SSP® DRA LOT

The DRA primer set as well as the specificity and interpretation tables are unchanged compared to the previous *Olerup* SSP® DRA lot (Lot No. Y51).

Lot No.: **52F** Lot-specific information

## PRODUCT DESCRIPTION

## **DRA SSP typing**

#### CONTENT

The primer set contains 5'- and 3'-primers for identifying the DRA\*0101 to DRA\*0102 alleles.

#### STRIP LAYOUT

Each test consists of 2 PCR reactions in an 8 well PCR plate. Wells 3 to 8 are empty.

1 2 empty empty empty empty empty empty

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

Well No. 1 is marked with the Lot No. '52F'.

The PCR plates are covered with a PCR-compatible foil.

**Please note:** When removing each two-well typing, make sure that the remaining plates/wells stay covered. Use a scalpel or a similar instrument to carefully cut the foil between the plates/wells.

#### INTERPRETATION

Only the DRA alleles will be amplified by the DRA typing kit. Thus, the interpretation of DRA typings is not influenced by the other HLA class II genes.

#### **UNIQUELY IDENTIFIED ALLELES**

All the DRA alleles, i.e. **DRA\*0101 to DRA\*0102**, recognized by the HLA Nomenclature Committee in February 2009<sup>1</sup> will give rise to unique amplification patterns by the primers in the DRA typing kit.

The DRA kit cannot distinguish the DRA\*010201 and DRA\*010202 alleles.

<sup>1</sup>DRA alleles listed on the IMGT/HLA web page 2009-January-16, release 2.24.0, www.ebi.ac.uk/imgt/hla.

#### RESOLUTION IN HOMO- AND HETEROZYGOTES

The 2 DRA alleles can be combined in 3 homozygous and heterozygous combinations. These 3 genotypes give rise to unique amplification patterns.

Lot No.: **52F** Lot-specific information

## SPECIFICITY TABLE

## **DRA SSP typing**

Specificities and sizes of the PCR products of the 2 primer mixes used for DRA SSP typing

| Primer<br>Mix  | Size of<br>spec. PCR<br>product <sup>1</sup> | Size of control band <sup>2</sup> | Amplified DRA alleles |
|----------------|--|-----------------------------------|-----------------------|
| 1 <sup>3</sup> | 65   | 515 bp                            | 0101                  |
| $2^3$          | 100  | 430 bp                            | 010201-010202         |

<sup>&</sup>lt;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 DRA SSP subtypings.

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

Nonspecific amplifications, i.e. a ladder or a smear of bands, may sometimes be seen. GC-rich primers have a higher tendency of giving rise to nonspecific amplifications than other primers.

PCR fragments longer than the control bands may sometimes be observed. Such bands should be disregarded and do not influence the interpretation of the SSP typings.

PCR fragments migrating faster than the control bands, but slower than a 400 bp fragment may be seen in some gel read-outs. Such bands can be disregarded and do not influence the interpretation of the SSP typings.

Some primers may give rise to primer oligomer artifacts. Sometimes this phenomenon is an inherit feature of the primer pair(s) of a primer mix. More often it is due to other factors such as too low amount of DNA in the PCR reactions, taking too long time in setting up the PCR reactions, working at elevated room temperature or using thermal cyclers that are not pre-heated.

<sup>2</sup>The internal positive control primer pairs amplify segments of the human growth hormone gene. The two different control primer pairs give rise to either an internal positive control band of 430 base pairs, for most wells, or a band of 515 base pairs, for some wells.

Well number 1 contains the primer pair giving rise to the longer, 515 bp, internal positive control band in order to help in the correct orientation of the DRA typing.

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

<sup>3</sup>Specific PCR fragments shorter than 125 base pairs are less intense and not as sharp as longer specific bands.

Lot No.: **52F** Lot-specific information

| INTERPRETATION TABLE                      |                                   |                                  |                        |  |  |  |  |  |
|---|-----------------------------------|----------------------------------|------------------------|--|--|--|--|--|
| DRA SSP typing                            |                                   |                                  |                        |  |  |  |  |  |
| Amplification patterns of the DRA alleles |                                   |                                  |                        |  |  |  |  |  |
|   | well                              |                                  |                        |  |  |  |  |  |
|   | 1                                 | 2                                |                        |  |  |  |  |  |
| Length of spec.                           | 65                                | 100                              | Length of spec.        |  |  |  |  |  |
| PCR product                               |                                   |                                  | PCR product            |  |  |  |  |  |
| Length of int.                            | 515                               | 430                              | Length of int.         |  |  |  |  |  |
| pos. control                              |                                   |                                  | pos. control           |  |  |  |  |  |
| 5'-primer <sup>2</sup>                    | 217                               | 197                              | 5'-primer <sup>2</sup> |  |  |  |  |  |
|   | <sup>5'</sup> -gA g <sup>3'</sup> | 5'-CC C3'                        |                        |  |  |  |  |  |
|   |                                   |                                  |                        |  |  |  |  |  |
| 3'-primer <sup>3</sup>                    | 224                               | 217                              | 3'-primer <sup>3</sup> |  |  |  |  |  |
|   | <sup>5'</sup> -g TT <sup>3'</sup> | <sup>5'</sup> -CAA <sup>3'</sup> |                        |  |  |  |  |  |
| Well No.                                  | 1                                 | 2                                | Well No.               |  |  |  |  |  |
| DRA allele                                |                                   |                                  | DRA allele             |  |  |  |  |  |
| *0101                                     | 1                                 |                                  | *0101                  |  |  |  |  |  |
| *010201-010202                            |                                   | 2                                | *010201-010202         |  |  |  |  |  |
| DRA allele                                |                                   |                                  | DRA allele             |  |  |  |  |  |
| Well No.                                  | 1                                 | 2                                | Well No.               |  |  |  |  |  |

<sup>&</sup>lt;sup>1</sup>The internal positive control primer pairs amplify segments of the human growth hormone gene. The two different control primer pairs give rise to either an internal positive control band of 430 base pairs, for most wells, or a band of 515 base pairs, for some wells.

Well number 1 contains the primer pair giving rise to the longer, 515 bp, internal positive control band in order to help in the correct orientation of the DRA typing.

<sup>&</sup>lt;sup>2</sup>The codon, in the 4<sup>th</sup> exon, matching the specificity-determining 3'-end of the primer is given. Codon numbering 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. Empty spaces indicate codon boundaries.

<sup>&</sup>lt;sup>3</sup>The codon, in the 4<sup>th</sup> exon, matching the specificity-determining 3'-end of the primer is given in the anti-sense direction. 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. Empty spaces indicate codon boundaries.

Lot No.: **52F** Lot-specific information

| <b>CELL LINE VALIDATION SHEET</b> |                |                 |                |                   |           |           |  |  |
|-----------------------------------|----------------|-----------------|----------------|-------------------|-----------|-----------|--|--|
| DRA SSP typing kit                |                |                 |                |                   |           |           |  |  |
|                                   |                |                 |                |                   | W         | ell       |  |  |
|                                   |                |                 |                |                   | 1         | 2         |  |  |
|                                   |                |                 |                | ction             | 200855701 | 200855702 |  |  |
|                                   |                |                 |                | Production<br>No. | 2008      | 2008      |  |  |
|                                   | IHWC cell line |                 | _              | DRA               |           |           |  |  |
| 1                                 | 9001           |                 | *0101          |                   | +         | -         |  |  |
| 2                                 |                | LK707           | *0101          |                   | +         | -         |  |  |
| 3                                 |                | E4181324        | *0101          |                   | +         | -         |  |  |
| 5                                 |                | GU373<br>KAS011 | *0101          |                   | +         | -         |  |  |
| 6                                 | 9359           |                 | *0101<br>*0102 |                   | +         | -         |  |  |
| 7                                 | 9020           |                 | *0102          |                   | -         | +         |  |  |
| 8                                 | 9020           |                 | *0101          |                   | +         | +         |  |  |
| 9                                 | 9007           |                 | *0102          |                   | +         | +         |  |  |
| 10                                |                | LKT3            | *0101          |                   | +         | H         |  |  |
| 11                                |                | PITOUT          | *0101          |                   | +         | -         |  |  |
| 12                                | 9052           |                 | *0101          |                   | +         | -         |  |  |
| 13                                | 9067           | BTB             | *0101          |                   | _         | +         |  |  |
| 14                                |                | OLGA            | *0102          |                   | _         | +         |  |  |
| 15                                | 9075           |                 | *0101          | *0102             | +         | +         |  |  |
| 16                                |                | SWEIG007        | *0101          | 0102              | +         | Ė         |  |  |
| 17                                |                | WILJON          | *0101          |                   | _         | +         |  |  |
| 18                                | 9257           | 32367           | *0102          |                   | _         | +         |  |  |
| 19                                |                | BM16            | *0101          |                   | +         | -         |  |  |
| 20                                |                | SLE005          | *0101          |                   | +         | -         |  |  |
| 21                                |                | AMALA           | *0101          |                   | +         | -         |  |  |
| 22                                |                | KOSE            | *0102          |                   | Ė         | +         |  |  |
| 23                                | 9124           |                 | *0101          | *0102             | +         | +         |  |  |
| 24                                |                | JBUSH           | *0101          | 0.02              | +         | -         |  |  |
| 25                                |                | IBW9            | *0101          |                   | +         | -         |  |  |
| 26                                |                | WT49            | *0101          | *0102             | +         | +         |  |  |
| 27                                |                | CH1007          | *0102          |                   | -         | +         |  |  |
| 28                                | 9320           | BEL5GB          | *0101          |                   | +         | -         |  |  |
| 29                                |                | MOU             | *0101          |                   | +         | -         |  |  |
| 30                                | 9021           | RSH             | *0101          | *0102             | +         | +         |  |  |
| 31                                | 9019           | DUCAF           | *0101          |                   | +         | -         |  |  |
| 32                                | 9297           | HAG             | *0101          |                   | +         | -         |  |  |
| 33                                | 9098           | MT14B           | *0101          |                   | +         | -         |  |  |
| 34                                | 9104           | DHIF            | *0101          |                   | +         | -         |  |  |
| 35                                | 9302           | SSTO            | *0101          |                   | +         | -         |  |  |
| 36                                | 9024           | KT17            | *0101          |                   | +         | -         |  |  |
| 37                                | 9065           | HHKB            | *0102          |                   | -         | +         |  |  |
| 38                                | 9099           | LZL             | *0102          |                   | -         | +         |  |  |
| 39                                | 9315           | CML             | *0101          |                   | +         | -         |  |  |
| 40                                | 9062           | WDV             | *0101          |                   | +         | -         |  |  |
| 41                                | 9055           | H0301           | *0102          |                   | -         | +         |  |  |
| 42                                | 9066           | TAB089          | *0101          | *0102             | +         | +         |  |  |
| 43                                | 9076           | T7526           | *0102          |                   | -         | +         |  |  |
| 44                                | 9057           | TEM             | *0102          |                   | -         | +         |  |  |
| 45                                | 9239           | SHJO            | *0102          |                   | -         | +         |  |  |
| 46                                |                | SCHU            | *0101          | *0102             | +         | +         |  |  |
| 47                                |                | TUBO            | *0102          |                   | -         | +         |  |  |
| 48                                | 9303           |                 | *0101          |                   | +         | -         |  |  |

Lot No.: **52F** Lot-specific information

## **CERTIFICATE OF ANALYSIS**

Olerup SSP® DRA SSP

Product number: 101.131-24u – without Taq polymerase

Lot number: 52F

Expiry date: 2011-February-01

Number of tests: 24 Number of wells per test: 2

### Well specifications:

| Well No. | Production No. |  |  |
|----------|----------------|--|--|
| 1        | 2008-557-01    |  |  |
| 2        | 2008-557-02    |  |  |

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

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

Date of approval: 2009-May-25

Approved by:

**Quality Control, Supervisor** 

Lot No.: **52F** Lot-specific information

## **Declaration of Conformity**

**Product name:** Olerup SSP® DRA

Product number: 101.131-24u

Lot number: 52F

**Intended use:** DRA high resolution histocompatibility testing

Manufacturer: Olerup SSP AB

Hasselstigen 1

SE-133 33 Saltsjöbaden, Sweden

**Phone:** +46-8-717 88 27 **Fax:** +46-8-717 88 18

We, *Olerup* SSP AB, hereby declare that this product, to which this Declaration of Conformity relates is in conformity with the following Standard(s) and other normative document(s) ISO 9001:2000 and ISO 13485:2003, following the provisions of the 98/79/EC Directive on *in vitro* diagnostic medical devices, Annex II List B, conformity assessed using Annex IV, as transposed into the national laws of the Member States of the European Union.

The Technical Documentation File is maintained at *Olerup* SSP AB, Hasselstigen 1, SE-133 33 Saltsjöbaden, Sweden.

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

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

Saltsjöbaden, Sweden 2009-May-25

Olle Olerup Managing Director

Lot No.: **52F** Lot-specific information

ADDRESSES:

Manufacturer:

Olerup SSP AB, Hasselstigen 1, SE-133 33 Saltsjöbaden, Sweden.

**Tel:** +46-8-717 88 27 **Fax:** +46-8-717 88 18

*E-mail:* info-ssp@olerup.com *Web page:* http://www.olerup.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: <a href="http://www.olerup.com">http://www.olerup.com</a>

For information on Olerup SSP distributors worldwide, contact Olerup GmbH.