Lot-specific Information

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Olerup SSP® DQB1*04

| Product number: | 101.215-12 – including <i>Taq</i> polymerase |
|----------------------------------|--|
| Lot number: | 87M |
| Expiry date: | 2014-May-01 |
| Number of tests: | 12 |
| Number of wells per test: | 8 |
| 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. 87M.

CHANGES COMPARED TO THE PREVIOUS OLERUP SSP[®] DQB1*04 LOT

The DQB1*04 specificity and interpretation tables have been updated with the DQB1 alleles described since the previous *Olerup* SSP[®] DQB1*04 lot (Lot No. 54K) was made.

Three wells have been added to the HLA-DQB1*04 kit, wells 6 to 8.

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 | Added | - | Primer added for the DQB1*04:02:02 allele. |
| 6 | New | New | New primer pair for the DQB1*04:06 allele. |
| 7 | New | New | New primer pair for the DQB1*04:07 allele. |
| 8 | New | New | New primer pair for the DQB1*04:08 allele. |

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PRODUCT DESCRIPTION

DQB1*04 SSP subtyping

CONTENT

The primer set contains 5'- and 3'-primers for identifying the DQB1*04:01 to DQB1*04:08 alleles.

Please note that DQB1 amplifications usually are somewhat less pronounced than e.g. DRB and DQA1 amplifications even when using the same DNA preparation and exactly the same experimental procedures.

PLATE LAYOUT

Each test consists of 8 PCR reactions in an 8 well PCR plate.

| 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 |
|---|---|---|---|---|---|---|---|
| | | | - | | | | |

The 8 well PCR plate is marked with 'DQ4' in silver/gray ink.

Well No. 1 is marked with the Lot No. '87M'.

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 8 well PCR plate, make sure that the remaining plates stay covered. Use a scalpel or a similar instrument to carefully cut the foil between the plates.

INTERPRETATION

Only the DQB1*04 alleles will be amplified by the DQB1*04 subtyping kit, except DQB1*03 alleles will be amplified by primer mixes 4, 5 and 8. Thus, the interpretation of DQB1*04 subtypings is not influenced by other groups of the DQB1 alleles or the DQB2 and DQB3 genes.

UNIQUELY IDENTIFIED ALLELES

All the DQB1*04 alleles, i.e. **DQB1*04:01 to DQB1*04:08**, recognized by the HLA Nomenclature Committee in July 2011¹ will give rise to unique amplification patterns by the primers in the DQB1*04 subtyping kit.

The DQB1*04 subtyping kit cannot distinguish the DQB1*04:01:01 and DQB1*04:01:02 alleles or the DQB1*04:02:01 and 04:02:02 alleles.

¹HLA-DQB1 alleles listed on the IMGT/HLA web page 2011-July-14, release 3.5.0, <u>www.ebi.ac.uk/imgt/hla</u>.

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RESOLUTION IN HOMO- AND HETEROZYGOTES

A total of 11 alleles generate 9 amplification patterns that can be combined in 45 homozygous and heterozygous combinations. 19 of these genotypes do not give rise to unique amplification patterns. The different lengths of the specific PCR products were not considered in these calculations.

| +++ | *04:01:01, *04:04 = *04:02:01, *04:05 = *04:04, *04:05 |
|------|--|
| +++ | *04:01:01, *04:05 = *04:05, *04:05 |
| +++ | *04:01:01, *04:06 = *04:06, *04:06 |
| +++- | *04:01:01, *04:07 = *04:07, *04:07 |
| +++ | *04:01:01, *04:08 = *04:08, *04:08 |
| +-++ | *04:03:01, *04:04 = *04:03:02, *04:04 |
| +-++ | *04:02:01, *04:03:01 = *04:02:01, *04:03:02 = *04:03:01, *04:03:01 |
| | = *04:03:01, *04:03:02 |
| +-+ | *04:02:01, *04:04 = *04:04, *04:04 |
| | |

*04:01:01 = *04:01:01 and 04:01:02 *04:02:01 = *04:02:01 and 04:02:02

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SPECIFICITY TABLE

DQB1*04 SSP subtyping

Specificities and sizes of the PCR products of the 8 primer mixes used for DQB1*04 SSP subtyping

| Primer Mix | Size of spec. PCR product ¹ | Size of control band ² | Amplified DQB1*04 alleles ³ | Amplified non- DQB1*04 alleles ⁴ |
|-----------------------|--|---|--|--|
| 1 ^{6,7} | 210 bp, 245 bp | 515 bp | *04:01:01-04:08 | |
| 2 ⁶ | 205 bp | 515 bp | *04:01:01-04:01:02, 04:05-04:08 | |
| 3 | 205 bp | 430 bp | *04:02:01-04:03:01, 04:04 | |
| 4 | 195 bp | 430 bp | *04:03:01-04:03:02 | *03:06, 03:25 |
| 5 | 245 bp | 430 bp | *04:04-04:05 | *03:06, 03:25 |
| 6 ⁵ | 110 bp | 430 bp | *04:06 | |
| 7 | 160 bp | 430 bp | *04:07 | |
| 8 ⁵ | 95 bp | 430 bp | *04:08 | *03:06 [?] -03:08 [?] , 03:11 [?] - 03:18 [?] , 03:19, 03:20 [?] , 03:23 [?] , 03:26 [?] , 03:37 [?] |

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 DQB1*04 SSP typings.

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 lengths 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.

²The 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 DQB1*04 subtyping.

In addition, well number 2 contains the primer pair giving rise to the longer, 515 bp, internal positive control band in order to allow kit identification.

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

³For several DQB1 alleles only partial 2nd exon nucleotide sequences are available. In these instances it is not known whether some of the primers of the SSP sets are completely matched with

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the target sequences or not. We assume that unknown sequences of the 2nd exon of DQB1 alleles are conserved within allelic groups.

⁴Due to the sharing of sequence motifs between DQB1 alleles non-DQB1*04 alleles will be amplified by primer mixes 4, 5 and 8.

⁵ Short specific PCR fragments are less intense and not as sharp as longer specific bands.

⁶Primer mixes 1 and 2 may give a lower yield of HLA-specific PCR product than the other DQB1*04

primer mixes. ⁷Primer mix 1: Specific PCR fragment of 210 bp in the DQB1*04:04 and 04:05 alleles. Specific PCR fragment of 245 bp in the DQB1*04:02:02 and 04:03:02 alleles. Specific PCR fragments of 210 and 245 bp in the DQB1*04:01:01-04:02:01, 04:03:01 and 04:06-04:08 alleles.

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

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| INTERPRETATION TABLE | | | | | | | | |
|---|----------------------------------|----------------------------------|----------------------------------|----------------------------------|----------------------------------|----------------------------------|----------------------------------|----------------------------------|
| DQB1*04 SSP subtyping | | | | | | | | |
| Amplification patterns of the DQB1*04 alleles | | | | | | | | |
| | | | | We | ell ⁴ | | | |
| | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 |
| Length of spec. | 210 | 205 | 205 | 195 | 245 | 110 | 160 | 95 |
| PCR product(s) | 245 | | | | | | | |
| Length of int. | 515 | 515 | 430 | 430 | 430 | 430 | 430 | 430 |
| pos. control ¹ | | | | | | | | |
| 5'-primer ² | 9(122) | 23(164) | 23(164) | 26(173) | 9(122) | 55(259) | 145(529) | 167(596) |
| | ^{5'} -gTT ^{3'} | ^{5'} -gCT ^{3'} | ^{5′} -gCg ^{3′} | ^{5'} -TCT ^{3'} | ^{5'} -gTA ^{3'} | ^{5'} -ggT ^{3'} | ^{5'} -CCg ^{3'} | ^{5'} -gCA ^{3'} |
| | 21(159) | | 23(164) | | | | | |
| | ^{5'} -ACC ^{3'} | | ^{5'} -gCg ^{3'} | | | | | |
| | | | | | | | | |
| 3'-primer ³ | 77(327) | 77(327) | 77(327) | 77(327) | 77(327) | 77(327) | 185(650) | 185(650) |
| | ^{5'} -ACg ^{3'} | ^{5'} -CgA ^{3'} | ^{5'} -CgA ^{3'} |
| Well No. | 1 | 2 | 3 | 4 | 5 | 5 | 5 | 5 |
| DQB1 allele | | | | | | | | |
| *04:01:01-04:01:02 | 1 | 2 | | | | | | |
| *04:02:01-04:02:02 | 1 | | 3 | | | | | |
| *04:03:01 | 1 | | 3 | 4 | | | | |
| *04:03:02 | 1 | | | 4 | | | | |
| *04:04 | 1 | | 3 | | 5 | | | |
| *04:05 | 1 | 2 | | | 5 | | | |
| *04:06 | 1 | 2 | | | | 6 | | |
| *04:07 | 1 | 2 | | | | | 7 | |
| *04:08 | 1 | 2 | | | | | | 8 |
| *03:06 | | | | 4 | 5 | | | ? |
| *03:07-03:08, 03:11- | | | | | | | | |
| 03:18, 03:20, 03:23, | | | | | | | | ? |
| 03:26, 03:37 | | | | | | | | |
| *03:19 | | | | | | | | 8 |
| *03:25 | | | | 4 | 5 | | | |
| DQB1 allele | | | | | | | | |
| Well No. | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 |

¹The 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 DQB1*04subtyping.

In addition, well number 2 contains the primer pair giving rise to the longer, 515 bp, internal positive control band in order to allow kit identification.

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²The codon, and in parenthesis the nucleotide, in the 2nd or 3rd exon, matching the specificitydetermining 3'-end of the primer is given. Codon and nucleotide numbering as on the <u>www.ebi.ac.uk/imgt/hla</u> web site. The sequence of the 3 terminal nucleotides of the primer is given.

given. ³The codon, and in parenthesis the nucleotide, in the 2nd or 3rd exon, matching the specificitydetermining 3'-end of the primer is given in the anti-sense direction. Codon and nucleotide numbering as on the <u>www.ebi.ac.uk/imgt/hla</u> web site. The sequence of the 3 terminal nucleotides of the primer is given.

⁴Primer mix 1: Specific PCR fragment of 210 bp in the DQB1*04:04 and 04:05 alleles. Specific PCR fragment of 245 bp in the DQB1*04:02:02 and 04:03:02 alleles. Specific PCR fragments of 210 and 245 bp in the DQB1*04:01:01-04:02:01, 04:03:01 and 04:06-04:08 alleles.

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

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| | CELL LINE VALIDATION SHEET | | | | | | | | | | | |
|----------|----------------------------|-------------------|--------|-------------------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|
| | DQB1*04 SSP subtyping kit | | | | | | | | | | | |
| | | | | | | | | W | ell | | | |
| | | | | | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 |
| | | | - | Production No. | 201077501 | 201077502 | 201192603 | 200846904 | 201077505 | 201192606 | 201192607 | 201192608 |
| | IHW | C cell line | | B1 | | | | | | | | |
| 1 | 9001 | | *05:01 | | - | - | - | - | - | - | - | - |
| 2 | | LK707 | *06:01 | *02:02 | - | - | - | - | - | - | - | - |
| 3 | | E4181324 | *06:01 | | - | - | - | - | - | - | - | - |
| 4 | | GU373 | *02:01 | | - | - | - | - | - | - | - | - |
| 5 | | KAS011 | *05:02 | | - | - | - | - | - | - | - | - |
| 6 | 9353 | | *03:02 | *06:01 | - | - | - | - | - | - | - | - |
| 7 | 9020 | | *02:01 | | - | - | - | - | - | - | - | - |
| 8 | 9025 | - | *03:01 | | - | - | - | - | - | - | - | - |
| 9 | | YAR | *03:02 | | - | - | - | - | - | - | - | - |
| 10 | | LKT3 | *04:01 | | + | + | - | - | - | - | - | - |
| 11 | | PITOUT | *02:02 | | - | - | - | - | - | - | - | - |
| 12 | 9052 | | *03:03 | | - | - | - | - | - | - | - | - |
| 13 | | JESTHOM | *05:01 | | - | - | - | - | - | - | - | - |
| 14 | 9071 | OLGA | *04:02 | | + | - | + | - | - | - | - | - |
| 15 | 9075 | | *03:03 | | - | - | - | - | - | - | - | - |
| 16 | 9037 | SWEIG007 | *03:01 | | - | - | - | - | - | - | - | - |
| 17 | 9282 | CTM3953540 | *02:01 | *06:03 | - | - | - | - | - | - | - | - |
| 18 | 9257 | 32367 | *06:02 | *02:02 | - | - | - | - | - | - | - | - |
| 19 | 9038 | BM16 | *03:01 | | - | - | - | - | - | - | - | - |
| 20 | 9059 | SLE005 | *06:04 | | - | - | - | - | - | - | - | - |
| 21 | 9064 | AMALA | *03:01 | | - | - | - | - | - | - | - | - |
| 22 | 9056 | KOSE | *05:03 | *06:04 | - | - | - | - | - | - | - | - |
| 23 | 9124 | | *05:03 | *06:01 | - | - | - | - | - | - | - | - |
| 24 | 9035 | JBUSH | *03:01 | | - | - | - | - | - | - | - | - |
| 25 | 9049 | IBW9 | *02:02 | | - | - | - | - | - | - | - | - |
| 26 | | WT49 | *02:01 | | - | - | - | - | - | - | - | - |
| 27 | | CH1007 | *04:01 | *05:01 | + | + | - | - | - | - | - | - |
| 28 | | BEL5GB | *02:02 | *03:01 | - | - | - | - | - | - | - | - |
| 29 | | MOU | *02:02 | | - | - | - | - | - | - | - | - |
| 30 | 9021 | | *04:02 | | + | - | + | - | - | - | - | - |
| 31 | | DUCAF | *02:01 | | <u> </u> | - | - | - | - | - | - | - |
| 32 | 9297 | | *03:01 | | - | - | - | - | - | - | - | - |
| 33 | | MT14B | *03:02 | | - | - | - | - | - | - | - | - |
| 34 | 9104 | | *03:02 | | - | - | - | - | - | - | - | - |
| 35 | | SSTO | *03:05 | | - | - | - | - | - | - | - | _ |
| 36 | | KT17 | *03:03 | | - | - | - | - | - | - | - | - |
| 30 | | ННКВ | *06:03 | | - | - | - | - | - | - | - | |
| 38 | 9005 | | *03:01 | | | - | - | - | - | - | - | |
| 30 | 9099 | | *02:01 | *03:01 | | - | - | - | - | - | - | _ |
| 39 40 | | WHONP199 | | *03:01 | | - | | | - | - | | |
| | | WHONP199 H0301 | *02:02 | 03.03 | - | - | - | - | - | - | - | - |
| 41 | | | *06:09 | | - | - | - | - | - | - | - | - |
| 42 | | TAB089 | *06:01 | | - | - | - | - | - | - | - | - |
| 43 | | T7526 | *03:03 | | - | - | - | - | - | - | - | - |
| 44 | 9057 | | *05:03 | | - | - | - | - | - | - | - | - |
| 45 | | SHJO | *02:02 | | - | - | - | - | - | - | - | - |
| 46 | | SCHU | *06:02 | | - | - | - | - | - | - | - | - |
| 47 | | TUBO | *03:01 | | - | - | - | - | - | - | - | - |
| 48 | 9303 | TER-ND | *05:01 | | - | - | - | - | - | - | - | - |

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CERTIFICATE OF ANALYSIS

| <i>Olerup</i> SSP [®] DQB1*04 SSP | |
|--|-----------|
| Product number: | 101.215-1 |
| Lot number: | 87M |
| Expiry date: | 2014-May |
| Number of tests: | 12 |
| Number of wells per test: | 8 |

12 – including *Taq* polymerase v-01

Well specifications:

| Well No. | Production No. | | |
|----------|----------------|--|--|
| 1 | 2010-775-01 | | |
| 2 | 2010-775-02 | | |
| 3 | 2011-926-03 | | |
| 4 | 2008-469-04 | | |
| 5 | 2010-775-05 | | |
| 6 | 2011-926-06 | | |
| 7 | 2011-926-07 | | |
| 8 | 2011-926-08 | | |

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

No DNAs carrying the alleles to be amplified by primer solutions 4 to 7 were available. The specificities of the primers in primer solutions 4 and 5 were tested by separately adding one additional 5'-primer, respectively one additional 3'-primer. In primer solutions 6 and 7 it was only possible to test the 3'-primer, the 5'-primer was not possible to test. In primer solution 3 one 5'-primer was not possible to test.

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

Date of approval: 2011-December-02

Approved by:

Production Quality Control

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Declaration of Conformity

| Product name: Product number: Lot number: | <i>Olerup</i> SSP [®] DQB1*04 101.215-12 87M |
|---|---|
| Intended use: | DQB1*04 resolution histocompatibility testing |
| Manufacturer: | <i>Olerup</i> SSP AB Franzengatan 5 SE-112 51 Stockholm, Sweden <i>Phone:</i> +46-8-717 88 27 <i>Fax:</i> +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 9001:2008 and ISO 13485:2003, following the provisions of the 98/79/EC Directive on *in vitro* diagnostic medical devices, Annex III, as transposed into the national laws of the Member States of the European Union.

The Technical Documentation File is maintained at *Olerup* SSP AB, Franzengatan 5, SE-112 51 Stockholm, Sweden.

Stockholm, Sweden 2012-January-09

Ann-Cathrin Jareman Head of QA and Regulatory Affairs

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| | IFU-01 Rev. | No. 03 can be downloaded from |
| Lot No.: 87M | Lot-specific Information | www.olerup-ssp.com |

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