■LERUP SSP* DRB1*14:01/14:54

Product Insert Page 1 of 12

101.813-12 – including *Taq* **polymerase**, IFU-01 **101.813-12u – without** *Taq* **polymerase**, IFU-02

Visit https://labproducts.caredx.com for "Instructions for Use" (IFU)

Lot No.: 6H7 Lot-specific information

Olerup SSP® DRB1*14:01/14:54

Product number: 101.813-12 – including *Tag* pol.

101.813-12u - without *Tag* pol.

Lot number: 6H7

Expiry date: 2023-04-01

Number of tests: 12 Number of wells per test: 2+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. 6H7.

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*14:01/14:54 Lot (0G7)

The DRB1*14:01/14:54 kit is updated for new alleles to enable separation of:

- Null and Alternatively expressed alleles
- The product documentation has been updated for new alleles of IMGT 3.35.0

The format of the Worksheet has been changed.

The DRB1*14:01/14:54 specificity and interpretation tables have been updated for the DRB1 alleles since the previous *Olerup SSP*® DRB1*14:01/14:54 lot was made (**Lot No. 0G7**).

The DRB1*14:01/14:54 primer set is unchanged compared to the previous lot (Lot No. 0G7).

Changes in revision R01 compared to R00:

1. The expiration date has been altered due to extension of shelf-life.



101.813-12 – including *Taq* **polymerase**, IFU-01 **101.813-12u – without** *Taq* **polymerase**, IFU-02

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Lot No.: 6H7 Lot-specific information

Well **3** contains <u>Negative Control primer pairs</u>, 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 | 105 | 200 | 105 | 80 | 75 | 80 | 85 |
|------------------------|----------|----------------------------------|----------------------------------|-----------------------------------|-----------------------------------|-----------------------------------|----------------------------------|
| product | | | | | | | |
| 5'-primer ¹ | 164 | 340 | 440 | 45 | 45 | 43 | 36 |
| | 5'-CAC3' | ^{5'} -Agg ^{3'} | ^{5'} -TTA3' | ⁵ '-Tgg ³ ' | ⁵ '-Tgg ³ ' | ⁵ '-Tgg ³ ' | 5'-TAC3' |
| | | | | | | | 36 |
| | | | | | | | ^{5'} -TAT ^{3'} |
| 3'-primer ² | 231 | 2 nd I | 507 | 59 | 58 | 57 | 47 |
| • | 5'-TgC3' | ^{5'} -AAA ^{3'} | ^{5'} -TTg ^{3'} | 5'-CTC3' | ^{5'} -ggC ^{3'} | 5'-CTC3' | 5'-ACA3' |
| | | | | | | | 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 codonnumbering 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.

Page 3 of 12

101.813-12 – including *Taq* polymerase, IFU-01 **101.813-12u –** without *Taq* polymerase, IFU-02

Visit https://labproducts.caredx.com for "Instructions for Use" (IFU)

Lot No.: 6H7

Lot-specific information

PRODUCT DESCRIPTION

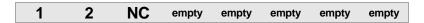
DRB1*14:01/14:54 SSP subtyping

CONTENT

The primer set contains 5'- and 3'-primers for separating the DRB1*14:01 and DRB1*14:54 alleles.

PLATE LAYOUT

Each test consists of 3 PCR reactions in an 8 well cut PCR plate. Wells 4 to 8 are empty.



The 8 well cut PCR plate is marked with '6H7' in silver/gray ink.

Well No. 1 is marked with the Lot No. '6H7'.

Wells 1 to 2 – DRB1*14:01/14:54 high resolution primers.

Well 3 – 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 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

Due to the sharing of sequence motifs some DRB1*11 alleles as well as many DRB1*14 alleles are amplified by the DRB1*14:01/14:54 primer mix 2. For further details see Specificity Table.

UNIQUELY IDENTIFIED ALLELES

The DRB1*14:01 and DRB1*14:54 alleles will give rise to unique amplification patterns in the DRB1*14:01/14:54 subtyping kit^{1,2}.

¹DRB alleles listed on the IMGT/HLA web page 2019-January-23, release 3.35.0, 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.

Page 4 of 12

101.813-12 – including *Taq* **polymerase**, IFU-01 **101.813-12u – without** *Taq* **polymerase**, IFU-02

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Lot No.: 6H7 Lot-specific information

SPECIFICITY TABLE

DRB1*14:01/14:54 SSP subtyping

Specificities and sizes of the PCR products of the 2+1 primer mixes used for DRB1*14:01/14:54 SSP subtyping

| Primer Mix | Size of spec. PCR product ¹ | Size of control band ² | Amplified DRB1*14:01/14:54 alleles | Other amplified DRB alleles ^{3,4} |
|-----------------------|----------------------------------------------|-----------------------------------|---------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 1 | 250 bp | 515 bp | *14:01:01-14:01:04 | |
| 2 | 210 bp | 430 bp | *14:01:01-14:01:04, 14:54:01:01-14:54:07 | *11:13:01-11:13:02, 11:17, 11:52, 11:89, 11:159, 11:202, 14:05:01:01-14:05:04, 14:07:01-14:08, 14:14, 14:18, 14:23:01-14:23:04, 14:26, 14:32:01-14:32:02, 14:34-14:36, 14:38:01-14:39, 14:42-14:45, 14:55 ^w , 14:56, 14:58-14:60, 14:62, 14:64-14:65, 14:70, 14:72, 14:75, 14:81-14:82, 14:86-14:88, 14:90-14:92N, 14:95-14:97, 14:99-14:101, 14:103, 14:110, 14:112-14:114, 14:117-14:118, 14:122-14:125, 14:127:01-14:127:02, 14:129-14:134, 14:139-14:140, 14:142, 14:147, 14:149-14:151, 14:153-14:155, 14:157-14:158, 14:160-14:163, 14:166N-14:169, 14:172, 14:178, 14:184-14:193, 14:197N, 14:202, 14:204, 14:207 |
| 3 ⁵ | - | - | 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 DRB1*14:01/14:54 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.

³For several DRB alleles 1st and/or 3rd exon(s) and beyond, as well as intron nucleotide





Product Insert Page 5 of 12

101.813-12 – including *Taq* **polymerase**, IFU-01 **101.813-12u – without** *Taq* **polymerase**, IFU-02

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Lot No.: 6H7 Lot-specific information

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 the sharing of sequence motifs some DRB1*11 alleles as well as many DRB1*14 alleles are amplified by the DRB1*14:01/14:54 primer mix 2.

⁵Primer mix 3 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. 'w', might be weakly amplified.

Page 6 of 12

101.813-12 – including *Taq* **polymerase**, IFU-01 **101.813-12u – without** *Taq* **polymerase**, IFU-02

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Lot No.: 6H7 Lot-specific information

PRIMER SPECIFICATION

| Well No. | 1 | 2 | |
|---------------------------|----------------------------------|----------------------------------|--|
| Length of spec. | 250 | 210 | |
| PCR product | | | |
| | | | |
| Length of int. | 515 | 430 | |
| pos. control ¹ | | | |
| 5'-primer(s) ² | 112(421) | 13(125) | |
| | 5' -ACT 3' | ^{5'} -gTC ^{3'} | |
| | | | |
| 3'-primer(s) ³ | 181(630) | | |
| | ^{5'} -CTT ^{3'} | ^{5'} -TCC ^{3'} | |
| Well No. | 1 | 2 | |

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.



101.813-12 – including *Taq* **polymerase**, IFU-01 **101.813-12u – without** *Taq* **polymerase**, IFU-02

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Lot No.: **6H7**

Lot-specific information

| CELL LINE VALIDATION SHEET | | | | | | | | |
|-------------------------------------------------|------|--------------------------|------------------|------------|-----------|-----------|--|--|
| DRB1*14:01/14:54 SSP subtyping kit ² | | | | | | | | |
| | | | | | | Well | | |
| | | | | | 1 | 2 | | |
| | | | | | ŀ. | | | |
| | | | | .: | β | 201438802 | | |
| | | | | Prod. No.: | 201438801 | 388 | | |
| | | | | 9 | 14 | 4 | | |
| | | | | <u> </u> | 20 | 8 | | |
| | IHW | C cell line ¹ | DI | DRB1 | | | | |
| 1 | 9001 | SA | *01:01 | | - | - | | |
| 2 | 9280 | LK707 | *15:02 | *04:05 | - | - | | |
| 3 | 9011 | E4181324 | *15:02 | | - | - | | |
| 4 | | GU373 | *03:01 | | - | - | | |
| 5 | 9009 | KAS011 | *16:01 | | - | - | | |
| 6 | 9353 | | *04:07 | *08:03 | - | - | | |
| 7 | 9020 | | *03:01 | | - | - | | |
| 8 | 9025 | | *04:01 | | - | - | | |
| 9 | | YAR | *04:02 | | - | - | | |
| 10 | | LKT3 | *04:05 | | - | <u>-</u> | | |
| 11 | | PITOUT | *07:01 | | - | - | | |
| 12 | 9052 | | *07:01 | | - | - | | |
| 13 | | JESTHOM | *01:01 | | <u> </u> | - | | |
| 14 | | OLGA | *08:02 | | <u> </u> | - | | |
| 15 | 9075 | | *09:01 | | <u> </u> | - | | |
| 16 | | SWEIG007 | *11:01 | | ļ - | - | | |
| 17 | | CTM3953540 | *03:01 | *13:01 | <u> </u> | - | | |
| 18 | | 32367 | *09:01 | *11:01 | <u> </u> | - | | |
| 19 | | BM16 | *12:01 | | <u> </u> | - | | |
| 20 | | SLE005 | *13:02 | | <u> </u> | - | | |
| 21 | | AMALA | *14:02 | *4454 | <u> </u> | - | | |
| 22 | | KOSE | *13:02 | *14:54 | <u> </u> | + | | |
| 23 | 9124 | | *08:03 | *14:14 | <u> </u> | + | | |
| 24 25 | | JBUSH IBW9 | *11:01 *07:01 | | <u> </u> | - | | |
| 26 | | WT49 | *03:01 | | ⊢⁼ | - | | |
| 27 | | CH1007 | *04:05 | *10:01 | ⊢⁼ | - | | |
| 28 | | BEL5GB | | *07:01 | | - | | |
| 29 | | MOU | *04:16 *07:01 | 07.01 | Ŀ | - | | |
| 30 | 9050 | | *07:01 | | Ŀ | Ė | | |
| 31 | | DUCAF | *03:02 | | ۱ | Ė | | |
| 32 | | HAG | *13:03 | | Ė | - | | |
| 33 | | MT14B | *04:04 | | H | _ | | |
| 34 | 9104 | | *11:01 | | Ē | - | | |
| 35 | | SSTO | *04:03 | | Ė | - | | |
| 36 | | KT17 | *04:03 | *04:06 | Ι. | - | | |
| 37 | | HHKB | *13:01 | 07.00 | <u> </u> | - | | |
| 38 | 9099 | | *14:02 | | + | - | | |
| 39 | 9315 | | *03:01 | *04:01 | - | - | | |
| 40 | | WHONP199 | *07:01 | *09:01 | - | - | | |
| 41 | | H0301 | *13:02 | 55.01 | +- | - | | |
| 42 | | TAB089 | *08:03 | | - | - | | |
| 43 | | T7526 | *09:01 | | - | - | | |
| 44 | 9057 | | *14:01 | | + | + | | |
| 45 | | SHJO | *07:01 | | - | - | | |
| 46 | | SCHU | *15:01 | | - | - | | |
| 415 | | | 10.01 | | | | | |
| 46 | | TUBO | *11:04 | *12:01 | - | - | | |



Page 8 of 12

101.813-12 - including Taq polymerase, IFU-01 101.813-12u - without *Taq* polymerase, IFU-02

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Lot No.: 6H7 **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.



Page 9 of 12

101.813-12 – including *Taq* **polymerase**, IFU-01 **101.813-12u – without** *Taq* **polymerase**, IFU-02

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Lot No.: **6H7** Lot-specific information





ert Page 10 of 12

101.813-12 – including *Taq* **polymerase**, IFU-01 **101.813-12u – without** *Taq* **polymerase**, IFU-02

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Lot No.: 6H7 Lot-specific information





rt Page 11 of 12

101.813-12 – including *Taq* **polymerase**, IFU-01 **101.813-12u – without** *Taq* **polymerase**, IFU-02

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Lot No.: 6H7 Lot-specific information





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Page 12 of 12

101.813-12 – including *Taq* **polymerase**, IFU-01 **101.813-12u – without** *Taq* **polymerase**, IFU-02

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Lot No.: 6H7 Lot-specific information

ADDRESSES:

Manufacturer:

CareDx AB, Franzengatan 5, SE-112 51 Stockholm, Sweden.

Tel: +46-8-508 939 00 Fax: +46-8-717 88 18

E-mail: orders-se@caredx.com

Web page: https://labproducts.caredx.com/

CareDx Lab Solutions Inc., 901 S. Bolmar St., Suite R, West Chester, PA 19382

Tel: 1-877-653-78171 **Fax:** 610-344-7989

E-mail: orders-us@caredx.com

Web page: https://labproducts.caredx.com/

For information on CareDx distributors worldwide, contact CareDx AB.