

Olerup SSP® KIR Genotyping

| | |
|---|---|
| Product number: | 104.101-12 – including <i>Taq</i> polymerase 104.101-12u – without <i>Taq</i> polymerase |
| Lot number: | 97X |
| Expiry date: | 2017-July-01 |
| Number of tests: | 12 |
| Number of wells per test: | 23 + 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. 97X.

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

CHANGES COMPARED TO THE PREVIOUS OLERUP SSP® KIR GENOTYPING LOT (98V)

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

The KIR Genotyping kit design, specificity and interpretation tables are based on IPD-KIR database 2.4.0.

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.

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 | - | 5'-primer added for the 2DL3*00102 allele. |

Change in revision R01 compared to R00:

1. Primer mix 3, but not primer mix 2, has been changed compared to the previous lot of KIR Genotyping. This information has been corrected.

Changes in revision R02 compared to R01:

1. Primer mix 1 contains the 800 base pair positive control primer pair. This has been corrected in the Specificity and Primer Specification Tables.

Change in revision R03 compared to R02:

1. The primer set is not changed compared to previous lot. This information has been corrected in the front page.

Change in revision R04 compared to R03:

1. The footnotes in the Primer Specification have been corrected, reflecting that the primer positions refer to codon numbering.

Change in revision R05 compared to R04:

1. Primer mix 2 may give rise to a PCR product of 225 base pairs, in addition to product sizes of 65 and 150 base pairs.



Well **24** contains Negative Control primer pairs, that will produce exon 4 and/or exon 5 amplicons for more than 97% of applicable KIR alleles as well as amplicons generated by positive control primer pairs constituent of all primer mixes in the Olerup SSP® product range.

PCR product sizes: 280bp KIR specific amplicons
 430bp Positive control

| Length of PCR product | 280 | 280 | 280 | 280 |
|------------------------------|----------------------|----------------------|----------------------|----------------------|
| 5'-primer¹ | 110 | 109 | 208 | 208 |
| | 5'-CAg ^{3'} | 5'-CCT ^{3'} | 5'-CCA ^{3'} | 5'-CCg ^{3'} |
| 3'-primer | 187 | 187 | 288 | 288 |
| | 5'-ggT ^{3'} | 5'-ggT ^{3'} | 5'-gTC ^{3'} | 5'-gTC ^{3'} |
| | 187 | 187 | 288 | 288 |
| | 5'-ggT ^{3'} | 5'-ggT ^{3'} | 5'-ggT ^{3'} | 5'-ggT ^{3'} |
| | | | 288 | 288 |
| | | | 5'-gAT ^{3'} | 5'-gAT ^{3'} |
| 2DL1* | + | | + | |
| 2DL2* | + | | + | |
| 2DL3* | + | | + | |
| 2DL4* | N/A | N/A | | + |
| 2DL5A* | N/A | N/A | + | |
| 2DL5B* | N/A | N/A | + | |
| 2DS1* | + | | + | |
| 2DS2* | + | | + | |
| 2DS3* | + | | + | |
| 2DS4* | | + | + | |
| 2DS5* | + | | + | |
| 3DL1* | + | | + | |
| 3DL2* | + | | + | |
| 3DL3* | + | | + | |
| 3DS1* | + | | + | |
| 2DP1* | + | | + | |
| 3DP1* | + | | + | |

¹The codon position for KIR genes, in the 4th or 5th exon, matching the specificity-determining 3'-end of the primer is given. Codon numbering as on the www.ebi.ac.uk/ipd/kir web site. The sequence of the 3 terminal nucleotides of the primer is given.

PRODUCT DESCRIPTION

KIR Genotyping SSP typing

CONTENT

The primer set contains 5'- and 3'-primers for KIR Genotyping.

PLATE LAYOUT

Each test consists of 24 PCR reactions in a 24 well cut PCR plate.

| | | | | | | | |
|----|----|----|----|----|----|----|----|
| 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 |
| 9 | 10 | 11 | 12 | 13 | 14 | 15 | 16 |
| 17 | 18 | 19 | 20 | 21 | 22 | 23 | NC |

The 24 well cut PCR plate is marked with 'KIR GENOTYP' in silver/gray ink.

Well No. 1 is marked with the Lot No. '97X'.

Wells 1 to 23 – KIR Genotyping primers.

Well 24 – Negative Control.

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 24 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.

UNIQUELY IDENTIFIED ALLELES

All the KIR alleles available in the IPD KIR Sequence Database in April 2011¹ will be amplified by the primers in the KIR Genotyping SSP kit.

¹KIR alleles listed on the IPD KIR web page 2011-April-15, release 2.4.0, www.ebi.ac.uk/ipd/kir.

PROTOCOL

DNA EXTRACTION

Extracted, highly pure DNA is needed for SSP typings. DNA samples to be used for PCR-SSP HLA typing should be re-suspended in dH₂O. The A260/A280 ratio should be 1.6 – 2.0 by UV spectrophotometry for optimal band visualization during electrophoresis.

We recommend automated DNA extraction with the QIAGEN EZ1 DSP DNA Blood System. ACD blood should be used as starting material.

Alternatively, the DNA can be extracted by any preferred method yielding pure DNA. When using alternative methods, the DNA concentration should be adjusted to 30 ng/μl. **Do not use heparinised blood with these methods.**

Recommended DNA concentration using:

EZ1-extracted DNA, 15 ng/μl.

DNA extracted by other methods, 30 ng/μl.

Concentrations exceeding 50 ng/μl will increase the risk for nonspecific amplifications and weak extra bands. If necessary, dilute the extracted DNA in dH₂O.

DNA samples should not be re-suspended in solutions containing chelating agents such as EDTA, above 0.5 mM in concentration.

DNA samples may be used immediately after extraction or stored at +4°C for up to 2 weeks with no adverse effects on results. DNA samples can be stored at -20°C or colder for 9 months. The purity and concentration of extracted DNA samples that have been stored for a prolonged period should be tested for acceptability prior to HLA typing.

DNA samples should be shipped at +4°C or colder to preserve their integrity during transport.

PCR AMPLIFICATION

104.101-12 – including *Taq* polymerase

For one KIR Genotyping typing, begin by adding to well No. 24, i.e. the well with the negative control primer pairs:

7 μl dH₂O

3 μl PCR Master Mix complete with *Taq*,

then add at room temperature in a 0.5 ml tube:

27 x 2 μl = 54 μl DNA (30 ng/μl)

27 x 3 μl = 81 μl PCR Master Mix complete with *Taq* – mix well

before taking your aliquot

27 x 5 μl = 135 μl dH₂O

Mix well, dispense 10 µl of the DNA-PCR Master Mix-H₂O mixture into each of the 23 wells of an KIR Genotyping typing, i.e. wells 1 to 23. Cover the primer tray(s) with the provided adhesive seals. Check that all reaction wells are completely covered to prevent evaporative loss during PCR amplification.

104.101-12u – without *Taq* polymerase

For one KIR Genotyping typing, begin by adding at room temperature in a 0.5 ml tube:

28 x 3 µl = 84 µl PCR Master Mix without *Taq* – mix well before taking your aliquot

2.2 µl *Taq* polymerase (5 units/µl)

Mix well, dispense 3 µl of the PCR Master Mix-*Taq* mixture from the 0.5 ml tube into well No. 24, i.e. the well with the negative control primer pairs. Then add 7 µl dH₂O to well 24.

Then add at room temperature to the 0.5 ml tube containing 84 + 2.2 - 3 = 83,2 µl PCR Master Mix-*Taq* mixture:

27 x 2 µl = 54 µl DNA (30 ng/µl)

27 x 5 µl - 2,2 µl = 132.8 µl dH₂O

Mix well, dispense 10 µl of the DNA-PCR Master Mix-*Taq*-H₂O mixture into each of the 23 wells of an KIR Genotyping typing, i.e. wells 1 to 23. Cover the primer tray(s) with the provided adhesive seals. Check that all reaction wells are completely covered to prevent evaporative loss during PCR amplification.

Use a 96 well thermal cycler with a heated lid. The temperature gradient across the heating block should be < 1°C.

PCR cycling parameters:

| | | | |
|---------------|------|---------|-------------------------|
| 1. 1 cycle | 94°C | 2 min | denaturation |
| 2. 10 cycles | 94°C | 10 sec. | denaturation |
| | 65°C | 60 sec. | annealing and extension |
| 3. 20 cycles | 94°C | 10 sec. | denaturation |
| | 61°C | 50 sec. | annealing |
| | 72°C | 30 sec. | extension |
| 4. End - hold | RT | | if less than 8 hours |
| | 4°C | | if longer than 8 hours |

Total reaction volume in each well, 10 µl.

The same PCR cycling parameters are used for all the *Olerup* SSP kits.

AGAROSE GEL ELECTROPHORESIS

Prepare a 2% (w/v) agarose gel in 0.5 x TBE buffer. Dissolve the agarose by boiling in a microwave oven. Let the gel solution cool to 60°C. Stain the gel prior

to casting with ethidium bromide (10 mg/ml), 5 µl per 100 ml gel solution. For maximal ease of handling use our ethidium bromide dropper bottles (Product No. 103.301-10), 1 drop of ethidium bromide solution per 50-75 ml of gel. **Note: Ethidium bromide is a carcinogen. Handle with appropriate personal protective equipment.**

Load the PCR products, preferably using an 8-channel pipette. Load a DNA size marker (100 base pair ladder, Product No. 103.201-100) in one well per row.

Run the gel in 0.5 x TBE buffer, without re-circulation of the buffer, for 15-20 minutes at 8-10 V/cm.

DOCUMENTATION AND INTERPRETATION

Put the gel on a UV transilluminator and document by photography.

Record the presence and absence of specific PCR products. The relative lengths of the specific PCR products are helpful in the interpretation of the results.

Record the presence and relative lengths of the internal positive control bands. The differently sized control bands will help in the correct orientation of the typing as well as in kit identification.

Lanes without either control band or specific PCR products should be repeated.

Interpret the typings with the ***lot-specific Interpretation and Specificity Tables***.

PCR MASTER MIXES

The PCR Master Mix complete with *Taq* polymerase contains:

| | |
|-----------------------|--|
| <i>Taq</i> polymerase | 0.4 unit per 10 µl SSP reaction |
| nucleotides | final concentration of each dNTP is 200 µM |
| PCR buffer | final concentrations: 50 mM KCl, 1.5 mM MgCl ₂ , 10 mM Tris-HCl pH 8.3, 0.001% w/v gelatin |
| glycerol | final concentration of glycerol is 5% |
| cresol red | final concentration of cresol red is 100 µg/ml |

The same PCR Master Mix complete with *Taq* is used for all Olerup SSP kits.

The PCR Master Mix without *Taq* polymerase contains:

| | |
|-------------|--|
| nucleotides | final concentration of each dNTP is 200 µM |
| PCR buffer | final concentrations: 50 mM KCl, 1.5 mM MgCl ₂ , 10 mM Tris-HCl pH 8.3, 0.001% w/v gelatin |
| glycerol | final concentration of glycerol is 5% |
| cresol red | final concentration of cresol red is 100 µg/ml |

The same PCR Master Mix without *Taq* is used for all Olerup SSP kits.

SPECIFICITY TABLE

KIR Genotyping SSP typing

Specificities and sizes of the PCR products of the 23+1 primer mixes used for KIR SSP Genotyping SSP.

| Primer Mix | Size of spec. PCR product ¹ | Size of control band ² | KIR Gene | Amplified KIR ³ alleles |
|--------------------|--|-----------------------------------|-----------------|--|
| 1 | 145 bp | 800 bp | 2DL1 | 001-025 |
| 2 ^{4,7,9} | 65 bp, 225 bp 150 bp | 1070 bp | 2DL2 | 004 0010101-010 |
| 3 ^{4,7} | 100 bp, 520 bp | 1070 bp | 2DL3 | 0010101-017 |
| 4 | 200 bp | 1070 bp | 2DL4 | 00101-022 |
| 5 ⁶ | 155 bp | 1070 bp | 2DL5A, 2DL5B | 0010101-00105, 0050101- 005010104, 01201-01202, 014-015 0020101-004, 00601-011, 01301- 01303, 016 |
| 6 ⁵ | 1650 bp | 430 bp | 2DL5A | 0010101-00105, 0050101- 005010104, 01201-01202, 014-015 |
| 7 ^{5,8} | 1650 bp | 515 bp | 2DL5B | 0020101-004, 00601-011, 01301- 01303, 016 |
| 8 ⁴ | 100 bp | 1070 bp | 2DS1 | 001-008 |
| 9 | 205 bp | 1070 bp | 2DS2 | 0010101-008 |
| 10 | 130 bp | 1070 bp | 2DS3 | 00101-005 |
| 11 | 215 bp | 1070 bp | 2DS4 | 0010101-00104, 01101-01102, 014, 015 |
| 12 | 200 bp | 1070 bp | 2DS4 | 0030101-0030104, 0040101- 0040102, 0060101-0060102, 007- 010, 012, 013 |
| 13 ^{4,7} | 110 bp | 1070 bp | 2DS5 | 001-011 |
| 14 | 135 bp | 1070 bp | 3DL1 | 0010101-002, 00401-00403, 0050101-009, 01501-044, 051-054, 056, 057, 059-068, 072-073 |
| 15 | 200 bp | 1070 bp | 3DL2 | 0010101-062 |
| 16 ⁴ | 115 bp | 1070 bp | 3DL3 | 00101-036, 041-055 |
| 17 | 130 bp | 1070 bp | 3DS1 | 010-014, 045-049N, 050, 055, 058 |
| 18 | 165 bp | 1070 bp | 2DP1 | 00101-010 |
| 19 ⁴ | 125 bp | 1070 bp | 3DP1 | 001-010 |
| 20 | 235 bp | 1070 bp | 3DP1 | 0030101-0030402, 005, 006, 008, 010 |

104.101-12 – including *Taq* polymerase
104.101-12u – without *Taq* polymerase

Lot No.: **97X**

Lot-specific information

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| | | | | |
|--------------------------|--------|---------|------|--|
| 21 | 145 bp | 1070 bp | 2DS1 | 001 |
| 22^{4,10} | 95 bp | 1070 bp | 2DS1 | 0020101-008 |
| | 235 bp | | 3DP1 | 001-002, 004, 007, 0090101-00902 |
| 23 | 210 bp | 1070 bp | 3DL1 | 00401-00403, 019, 021, 036, 037, 039, 056, 072 |
| 24^{6,11} | - | - | - | 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 KIR 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 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 1070, 800, 430 or 515 base pairs respectively, well distributed as outlined in the table. 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.

³KIR alleles listed on the IPD KIR web page 2011-April-15, release 2.4.0, www.ebi.ac.uk/ipd/kir.

⁴Specific PCR products shorter than 125 base pairs have a lower intensity and are less sharp than longer PCR products.

⁵The specific PCR product generated by primer mixes 6 and 7 are longer than the internal positive control band and the positive control band may be weaker than for other KIR primer mixes.

⁶Primer mixes 5 and 24 have a tendency to give rise to primer oligomer formation.

⁷Primer mixes 2, 3 and 13 may have tendencies of unspecific amplifications.

⁸Primer mix 7 may give rise to a lower yield of specific PCR product than the other KIR primer mixes.

⁹The 2DL2*004 and the 2DL2*0010101-010 alleles may be distinguished by the different sizes of the specific PCR product in primer mix 2; three specific PCR fragments of 65, 150 and 225 bp in the 2DL2*004 allele and one specific PCR fragment of 150 bp in the 2DL2*0010101-00304 and 005-010 alleles.

¹⁰The 2DS1 and the 3DP1 amplicons in primer mix 22 are differentiated by amplicon size; a specific PCR fragment of 95 bp for the 2DS1*0020101-008 alleles and a specific PCR fragment of 235 bp for the 3DP1*001-002, 004, 007 and 0090101-00902 alleles.

¹¹Well 24 contains negative control primer pairs, that will produce exon 4 and/or exon 5 amplicons for more than 97% of applicable KIR alleles as well as amplicons generated by positive control primer pairs.

'?', the 2nd intron sequence of the primer matching region is not known.

PRIMER SPECIFICATION

| Well No. | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
|--|-----------------------------|-----------------------------|-----------------------------|-----------------------------|-----------------------------|-----------------------------|-----------------------------|-----------------------------|-----------------------------|-----------------------------|-----------------------------|-----------------------------|
| Length of spec. PCR product | 145 | 65 150 225 | 100 520 | 200 | 155 | 1650 | 1650 | 100 | 205 | 130 | 215 | 200 |
| Length of int. pos. control ¹ | 800 | 1070 | 1070 | 1070 | 1070 | 430 | 515 | 1070 | 1070 | 1070 | 1070 | 1070 |
| 5'-primer(s) ² | 130 5'-gAA ^{3'} | 208 5'-CCA ^{3'} | 226 5'-CCA ^{3'} | 208 5'-CCg ^{3'} | 226 5'-CCA ^{3'} | -16 5'-TCA ^{3'} | -16 5'-TCg ^{3'} | 165 5'-gAg ^{3'} | 140 5'-gTA ^{3'} | 236 5'-CAC ^{3'} | 229 5'-CTA ^{3'} | 234 5'-TCT ^{3'} |
| | 130 5'-TAA ^{3'} | 156 5'-AAA ^{3'} | 332 5'-TCg ^{3'} | | | | -16 5'-Tgg ^{3'} | 165 5'-gAA ^{3'} | | | | |
| | | 262 5'-ggA ^{3'} | 344 5'-CTg ^{3'} | | | | | | | | | |
| | | | 344 5'-CTg ^{3'} | | | | | | | | | |
| 3'-primer(s) ³ | 165 5'-gCg ^{3'} | 243 5'-ACA ^{3'} | 246 5'-AgA ^{3'} | 262 5'-ggA ^{3'} | 276 5'-gAg ^{3'} | 27 5'-ACA ^{3'} | 27 5'-ACA ^{3'} | 185 5'-gAC ^{3'} | 195 5'-ATg ^{3'} | 266 5'-CCT ^{3'} | 288 5'-ggA ^{3'} | 288 5'-ggA ^{3'} |
| | | 195 5'-ATg ^{3'} | 350 5'-CAA ^{3'} | | | | | | | | | |
| | | 269 5'-TAC ^{3'} | 351 5'-ACC ^{3'} | | | | | | | | | |
| Well No. | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |

| Well No. | 13 | 14 | 15 | 16 | 17 | 18 | 19 | 20 | 21 | 22 | 23 |
|--|-----------------------------|-----------------------------|-----------------------------|-----------------------------|-----------------------------|----------------------------|----------------------------|---|-----------------------------|---|----------------------------|
| Length of spec. PCR product | 110 | 135 | 200 | 115 | 130 | 165 | 125 | 235 | 145 | 95 235 | 210 |
| Length of int. pos. control ¹ | 1070 | 1070 | 1070 | 1070 | 1070 | 1070 | 1070 | 1070 | 1070 | 1070 | 1070 |
| 5'-primer(s) ² | 142 5'-ACC ^{3'} | 136 5'-CAA ^{3'} | 110 5'-ggg ^{3'} | 156 5'-CCC ^{3'} | 133 5'-TCT ^{3'} | 29 5'-CAT ^{3'} | 25 5'-Tgg ^{3'} | 2 nd I 5'-gCC ^{3'} | 130 5'-gAA ^{3'} | 130 5'-gAA ^{3'} | 31 5'-TCA ^{3'} |
| | | 208 5'-CCA ^{3'} | | | | | | | | 2 nd I 5'-TCC ^{3'} | 31 5'-TCA ^{3'} |
| 3'-primer(s) ³ | 165 5'-gTg ^{3'} | 166 5'-CAA ^{3'} | 164 5'-CAA ^{3'} | 181 5'-gTA ^{3'} | 163 5'-ggA ^{3'} | 71 5'-TAC ^{3'} | 54 5'-TAC ^{3'} | 54 5'-TAC ^{3'} | 165 5'-gCC ^{3'} | 54 5'-TAC ^{3'} | 86 5'-CCA ^{3'} |
| | | 238 5'-CCg ^{3'} | | | | | | | | 165 5'-gCT ^{3'} | |
| Well No. | 13 | 14 | 15 | 16 | 17 | 18 | 19 | 20 | 21 | 22 | 23 |

¹The internal positive control primer pairs amplify segments of the human growth hormone gene. The internal positive control bands are 1070, 800, 430 or 515 base pairs respectively, well distribution as outlined in the table. 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 codon position, in the 1st, 3rd, 4th, 5th or 7th exon or the 2nd intron matching the specificity-determining 3'-end of the primer is given. Codon numbering as on the KIR web page 2011-April-25, release 2.4.0., www.ebi.ac.uk/ipd/kir. The sequence of the 3 terminal nucleotides of the primer is given.

³The codon position, in the 3rd, 4th, 5th or 8th exon, matching the specificity-determining 3'-end of the primer. Codon numbering as on the KIR web page 2011-April-15, release 2.4.0., www.ebi.ac.uk/ipd/kir. The sequence of the 3 terminal nucleotides of the primer is given in the anti-sense direction.

| | | CELL LINE VALIDATION SHEET | | | | | | | | | | | | | | | |
|-----------------------------|-----------------|--|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|
| | | KIR Genotyping primer set ² | | | | | | | | | | | | | | | |
| | | Well | | | | | | | | | | | | | | | |
| | | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | 13 | 14 | 15 | 16 |
| | | 201324201 | 201437602 | 201448203 | 201324204 | 201324205 | 201324206 | 201448207 | 201324208 | 201324209 | 201324210 | 201324211 | 201324212 | 201324213 | 201324214 | 201324215 | 201324216 |
| IHCW cell line ¹ | | | | | | | | | | | | | | | | | |
| 1 | 9001 SA | + | - | + | + | - | - | - | - | - | - | + | - | - | + | + | + |
| 2 | 9280 LK707 | - | + | - | + | + | - | + | + | + | - | - | - | + | + | + | + |
| 3 | 9011 E4181324 | + | - | + | + | - | - | - | - | - | - | + | + | - | + | + | + |
| 4 | 9275 GU373 | + | - | + | + | - | - | - | - | - | - | + | + | - | + | + | + |
| 5 | 9009 KAS011 | + | - | + | + | + | + | - | + | - | - | - | + | + | + | + | + |
| 6 | 9353 SM | + | - | + | + | + | + | - | + | - | + | + | - | - | + | + | + |
| 7 | 9020 QBL | + | + | + | + | + | + | + | + | + | + | - | + | + | + | + | + |
| 8 | 9025 DEU | + | + | + | + | - | - | - | - | + | - | - | + | - | + | + | + |
| 9 | 9026 YAR | + | - | + | + | - | - | - | - | - | - | + | + | - | + | + | + |
| 10 | 9107 LKT3 | + | - | + | + | - | - | - | - | - | - | + | - | - | + | + | + |
| 11 | 9051 PITOUT | + | + | + | + | - | - | - | - | + | - | - | + | - | + | + | + |
| 12 | 9052 DBB | + | + | + | + | + | - | + | - | + | + | - | + | - | + | + | + |
| 13 | 9025 JESTHOM | + | + | - | + | + | - | + | - | + | + | + | + | - | + | + | + |
| 14 | 9071 OLGA | + | - | + | + | + | + | - | + | - | - | - | + | + | + | + | + |
| 15 | 9075 DKB | + | - | + | + | - | - | - | - | - | - | + | - | - | + | + | + |
| 16 | 9037 SWEIG007 | + | - | + | + | - | - | - | - | - | - | - | + | - | + | + | + |
| 17 | 9282 CTM3953540 | + | + | + | + | + | + | + | + | + | + | + | + | + | + | + | + |
| 18 | 9257 32367 | + | - | + | + | - | - | - | - | - | - | - | + | - | + | + | + |
| 19 | 9038 BM16 | + | - | + | + | - | - | - | - | - | - | - | + | - | + | + | + |
| 20 | 9059 SLE005 | + | - | + | + | - | - | - | - | - | - | + | + | - | + | + | + |
| 21 | 9064 AMALA | + | + | + | + | + | + | - | + | + | - | + | - | + | + | + | + |
| 22 | 9056 KOSE | + | + | + | + | - | - | - | - | + | - | - | + | - | + | + | + |
| 23 | 9124 IHL | + | + | + | + | - | - | - | - | + | - | + | + | - | + | + | + |
| 24 | 9035 JBUSH | + | - | + | + | - | - | - | - | - | - | + | + | - | + | + | + |
| 25 | 9049 IBW9 | + | - | + | + | - | - | - | - | - | - | + | + | - | + | + | + |
| 26 | 9285 WT49 | + | + | + | + | + | - | + | + | + | - | - | + | + | + | + | + |
| 27 | 9191 CH1007 | + | + | + | + | + | - | + | - | + | + | + | + | - | + | + | + |
| 28 | 9320 BEL5GB | + | + | - | + | + | - | + | - | + | + | - | + | - | + | + | + |
| 29 | 9050 MOU | + | - | + | + | - | - | - | - | - | - | - | + | - | + | + | + |
| 30 | 9021 RSH | + | + | + | + | + | - | + | - | + | - | + | + | + | + | + | + |
| 31 | 9019 DUCAF | + | - | + | + | - | - | - | - | - | - | - | + | - | + | + | + |
| 32 | 9297 HAG | + | - | + | + | - | - | - | - | - | - | + | + | - | + | + | + |
| 33 | 9098 MT14B | + | - | + | + | - | - | - | - | - | - | + | - | - | + | + | + |
| 34 | 9104 DHIF | + | + | + | + | - | - | - | - | + | - | + | + | - | + | + | + |
| 35 | 9302 SSTO | + | + | + | + | - | - | - | - | + | - | + | + | - | + | + | + |
| 36 | 9024 KT17 | + | - | + | + | + | + | - | + | - | + | - | + | - | + | + | + |
| 37 | 9065 HHKB | + | + | + | + | + | + | - | + | + | - | - | + | + | + | + | + |
| 38 | 9099 LZL | + | + | + | + | + | - | + | + | + | + | - | - | + | + | + | + |
| 39 | 9315 CML | + | + | - | + | + | + | + | + | + | + | - | + | - | + | + | + |
| 40 | 9134 WHONP199 | + | - | + | + | - | - | - | - | - | - | + | - | - | + | + | + |
| 41 | 9055 H0301 | + | + | - | + | + | - | + | - | + | + | + | - | - | + | + | + |
| 42 | 9066 TAB089 | + | - | + | + | - | - | - | - | - | - | - | + | - | + | + | + |
| 43 | 9076 T7526 | + | - | + | + | + | + | - | + | - | - | + | - | + | + | + | + |
| 44 | 9057 TEM | + | + | + | + | + | - | + | - | + | + | - | + | - | + | + | + |
| 45 | 9239 SHJO | + | + | + | + | + | - | + | - | + | - | + | + | + | + | + | + |
| 46 | 9013 SCHU | + | - | + | + | - | - | - | - | - | - | + | + | - | + | + | + |
| 47 | 9045 TUBO | + | - | + | + | - | - | - | - | - | - | - | + | - | + | + | + |
| 48 | 9303 TER-ND | + | - | + | + | - | - | - | - | - | - | - | + | - | + | + | + |



| CELL LINE VAL. SHEET | | | | | | | | | |
|--|------|------------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|
| KIR Genotyping primer set ² | | | | | | | | | |
| | | | Well | | | | | | |
| | | | 17 | 18 | 19 | 20 | 21 | 22 | 23 |
| | | | 201324217 | 201324218 | 201324219 | 201324220 | 201324221 | 201437622 | 201324223 |
| IHC cell line ¹ | | | | | | | | | |
| 1 | 9001 | SA | - | + | + | + | - | - | - |
| 2 | 9280 | LK707 | - | - | + | - | - | + | - |
| 3 | 9011 | E4181324 | + | + | + | + | - | + | + |
| 4 | 9275 | GU373 | - | + | + | + | - | - | - |
| 5 | 9009 | KAS011 | + | + | + | + | - | + | - |
| 6 | 9353 | SM | + | + | + | + | - | + | - |
| 7 | 9020 | QBL | + | + | + | + | - | + | + |
| 8 | 9025 | DEU | - | + | + | + | - | + | - |
| 9 | 9026 | YAR | - | + | + | + | - | - | - |
| 10 | 9107 | LKT3 | - | + | + | + | - | - | - |
| 11 | 9051 | PITOUT | - | + | + | + | - | + | + |
| 12 | 9052 | DBB | - | + | + | + | - | - | + |
| 13 | 9025 | JESTHOM | - | + | + | + | - | + | - |
| 14 | 9071 | OLGA | + | + | + | + | - | + | - |
| 15 | 9075 | DKB | - | + | + | + | - | - | - |
| 16 | 9037 | SWEIG007 | - | + | + | + | - | - | + |
| 17 | 9282 | CTM3953540 | + | + | + | + | - | + | + |
| 18 | 9257 | 32367 | - | + | + | + | - | - | + |
| 19 | 9038 | BM16 | - | + | + | + | - | - | - |
| 20 | 9059 | SLE005 | - | + | + | + | - | - | - |
| 21 | 9064 | AMALA | + | + | + | + | - | + | - |
| 22 | 9056 | KOSE | - | + | + | + | - | + | + |
| 23 | 9124 | IHL | - | + | + | + | - | + | + |
| 24 | 9035 | JBUSH | - | + | + | + | - | - | - |
| 25 | 9049 | IBW9 | - | + | + | + | - | - | + |
| 26 | 9285 | WT49 | - | + | + | + | - | + | - |
| 27 | 9191 | CH1007 | - | + | + | + | - | - | - |
| 28 | 9320 | BEL5GB | - | + | + | + | - | + | + |
| 29 | 9050 | MOU | - | + | + | + | - | - | + |
| 30 | 9021 | RSH | - | + | + | + | - | - | - |
| 31 | 9019 | DUCAF | - | + | + | + | - | - | - |
| 32 | 9297 | HAG | - | + | + | + | - | - | - |
| 33 | 9098 | MT14B | - | + | + | + | - | - | - |
| 34 | 9104 | DHIF | - | + | + | + | - | + | - |
| 35 | 9302 | SSTO | - | + | + | + | - | + | - |
| 36 | 9024 | KT17 | + | + | + | + | - | + | - |
| 37 | 9065 | HHKB | + | + | + | + | - | + | - |
| 38 | 9099 | LZL | - | + | + | + | - | + | - |
| 39 | 9315 | CML | + | + | + | + | - | + | + |
| 40 | 9134 | WHONP199 | - | + | + | + | - | - | - |
| 41 | 9055 | H0301 | - | + | + | + | - | + | - |
| 42 | 9066 | TAB089 | - | + | + | + | - | - | - |
| 43 | 9076 | T7526 | + | + | + | + | - | + | - |
| 44 | 9057 | TEM | - | + | + | + | - | - | - |
| 45 | 9239 | SHJO | - | + | + | + | - | - | - |
| 46 | 9013 | SCHU | - | + | + | + | - | - | - |
| 47 | 9045 | TUBO | - | + | + | + | - | - | - |
| 48 | 9303 | TER-ND | - | + | + | + | - | - | - |

104.101-12 – including *Taq* polymerase104.101-12u – without *Taq* polymeraseLot No.: **97X**

Lot-specific information

www.olerup-ssp.com

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.

Additional primers in primer solution 3 were tested by separately adding another 5'-primer respective another 3'-primer.

In primer solution 21 it was only possible to test the 5'-primer, the 3'-primer was not possible to test.

In primer solutions 1, 3 and 7, one of the 5'-primers was not possible to test, and in primer solution 2 one 5'-primer and one 3'-primer was not possible to test.

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GUARANTEE

Olerup SSP AB guarantees that the primers in the *Olerup SSP*® typing trays have the specificities given in the lot-specific Specificity and Interpretation Tables of the product insert.

When stored at –20°C, the dried primers are stable for 30 months from the date of manufacture.

When stored at –20°C, the PCR Master Mix including *Taq* polymerase and the PCR Master Mix without *Taq* polymerase are stable for 33 months from the date of manufacture.

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