■LERUP SSP*
KIR Genotyping

Product Insert

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

Lot No.: 8D0 Lot-specific information www.olerup.com

Olerup SSP® KIR Genotyping

Product number: 104.101-12 – including *Taq* polymerase

104.101-12u – without *Tag* polymerase

Lot number: 8D0

Expiry date: 2019-01-01

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
 Product Insert
 RT

This Product Description is only valid for Lot No. 8D0.

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 (97X)

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 2.6.0.

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
5	Modified	-	5'-primer modified for increased yield.
6	-	-	Strength of control primers optimized.
16	Added	Added	5'-primer added for increased yield of the 3DL3*00103 and 3DL3*00904 alleles, 3'-primer added for increased yield of the 3DL3*00601-00602, 01501, 0601, 02102 and 0250101 alleles.

Change in revision R01 compared to R00:

1. Typing errors in the Interpretation and Specificity Tables and the Cell Line Validation sheet have been corrected.

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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'-primer ¹	110	109	208	208
o primer			^{5'} -CCA ^{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'}
ODI 4*			_	
2DL1*	+		+	
2DL2*	+		+	
2DL3*	+	N1/A	+	
2DL4*	N/A	N/A		+
2DL5A*	N/A	N/A	+	
2DL5B*	N/A	N/A	+	
2DS1*	+		+	
2DS2*	+		+	
2D\$3*	+		+	
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.

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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. '8D0'.

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 2015¹ will be amplified by the primers in the KIR Genotyping SSP kit.

¹KIR alleles listed on the IPD KIR web page 2015-January-19, release 2.6.0, www.ebi.ac.uk/ipd/kir.

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KIR Genotyping

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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 dH $_2$ O. 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 $ng/\mu l$ will increase the risk for nonspecific amplifications and weak extra bands. If necessary, dilute the extracted DNA in dH_2O .

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:

 $7 \mu l dH₂O$

3 μl PCR Master Mix complete with Taq,

then add at room temperature in a 0.5 ml tube:

 $27 \times 2 \mu I = 54 \mu I DNA (30 ng/\mu I)$

27 x 3 μ l = 81 μ l PCR Master Mix complete with Taq – mix well

before taking your aliquot

 $27 \times 5 \mu l = 135 \mu l dH_2O$

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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:

 $28 \times 3 \mu I = 84 \mu I$ PCR Master Mix without Taq – mix well before taking your aliquot

2.2 μl *Taq* polymerase (5 units/μl)

Mix well, dispense 3 μ l 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 μ l dH₂O to well 24.

Then add at room temperature to the 0.5 ml tube containing 84 + 2.2 - 3 = 83.2 μ l PCR Master Mix-Tag mixture:

$$27 \times 2 \mu l = 54 \mu l DNA (30 ng/\mu l)$$

$$27 \times 5 \mu I - 2.2 \mu I = 132.8 \mu I dH2O$$

Mix well, dispense 10 μ l of the DNA-PCR Master Mix-Taq- H_2O 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 65°C	10 sec. 60 sec.	denaturation annealing and extension
3. 20 cycles	94°C 61°C 72°C	10 sec. 50 sec. 30 sec.	denaturation annealing extension
4. End - hold	RT 4°C		if less than 8 hours 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

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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:

Tag polymerase 0.4 unit per 10 μl SSP reaction

 $\begin{array}{ll} \text{nucleotides} & \text{final concentration of each dNTP is 200 } \mu\text{M} \\ \text{PCR buffer} & \text{final concentrations: 50 mM KCI, 1.5 mM MgCI}_2, \\ \end{array}$

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 Tag is used for all Olerup SSP kits.

The PCR Master Mix without *Taq* polymerase contains:

nucleotides final concentration of each dNTP is 200 μ M PCR buffer final concentrations: 50 mM KCl, 1.5 mM MgCl₂,

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.

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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	001-026N
2 ^{4,7,9}	65 bp, 225 bp 150 bp	1070 bp	2DL2 2DL2	004 0010101-013
3 ^{4,7}	100 bp, 520 bp	1070 bp	2DL3	0010101-032
4	200 bp	1070 bp	2DL4	00101-027
5 ⁶	155 bp	1070 bp	2DL5A, 2DL5B	0010101-00105, 0050101-0050104, 01201-01202, 014-015 0020101-004, 00601-011, 01301-01303, 016-0018
6 ⁵	1650 bp	430 bp	2DL5A	0010101-00105, 0050101- 005010104, 01201-01202, 014-015
7 ^{5,6,7,8}	1650 bp	515 bp	2DL5B	0020101-004, 00601-011, 01301- 01303, 016-018
84	100 bp	1070 bp	2DS1	001-008
9	205 bp	1070 bp	2DS2	0010101-008
10	130 bp	1070 bp	2DS3	00101-007
11	215 bp	1070 bp	2DS4	0010101-00104, 01101-01102, 014- 016
12	200 bp	1070 bp	2DS4	0030101-0030104, 0040101- 0040102, 0060101-0060102, 007- 010, 012, 013
13 ^{4,7}	110 bp	1070 bp	2DS5	001-012
14	135 bp	1070 bp	3DL1	0010101-002, 00401-00403, 0050101-009, 01501-044, 051-054, 056, 057, 059-077, 079
15	200 bp	1070 bp	3DL2	0010101-063
16 ⁴	115 bp	1070 bp	3DL3	00101-057
17	130 bp	1070 bp	3DS1	010-014, 045-049N, 050, 055, 058, 078, 082-085
18	165 bp	1070 bp	2DP1	00101-014
19 ⁴	125 bp	1070 bp	3DP1	001-014

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20	235 bp	1070 bp	3DP1	0030101-0030402, 005, 006, 008, 010, 013-014
21	145 bp	1070 bp	2DS1	001
22 ^{4,10}	95 bp 235 bp	1070 bp	2DS1 3DP1	0020101-008 001-002, 004, 007, 0090101-00902, 011-012
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 KIR-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 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.

³KIR alleles listed on the IPD KIR web page 2015-January-19, release 2.6.0, <u>www.ebi.ac.uk/ipd/kir</u>. 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, 7 and 24 have a tendency to giving rise to primer oligomer formation.

⁷Primer mixes 2, 3, 7 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*0010101-013 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 and 2DL2*011 alleles and one specific PCR fragment of 150 bp in the 2DL2*0010101-00304, 005-010 and 012-013 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, 0090101-00902 and 011-012 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.

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PRIMER SPECIFICATION

Well No.	1	2	3	4	5	6	7	8	9	10	11	12
Length of spec.	145	65	100	200	155	1650	1650	100	205	130	215	200
PCR product		150	520									
		225										
Length of int.	800	1070	1070	1070	1070	430	515	1070	1070	1070	1070	1070
pos. control ¹												
5'-primer(s) ²	130	208	226	208	226	-16	-16	165	140	236	229	234
. , ,	^{5'} -gAA ^{3'}	5'-CCA3'	5'-CCA3'	5'-CCg3'	5'-CCA3'	5'-TCA3'	5'-TCg ^{3'}	5'-gAg ^{3'}	^{5'} -gTA ^{3'}	5'-CAC3'	5'-CTA3'	5'-TCT3'
	130	156	332				-16	165				
	^{5'} -TAA ^{3'}	^{5'} -AAA ^{3'}	5'-TCg3'				5'-Tgg ^{3'}	5'-gAA3'				
		262	344									
		5'-ggA ^{3'}	5'-CTg3'									
			344									
			5'-CTg ^{3'}									
3'-primer(s) ³	165	243	246	262	276	27	27	185	195	266	288	288
. ,	^{5'} -gCg ^{3'}	5'-ACA3'	5'-AgA ^{3'}	^{5'} -ggA ^{3'}	^{5'} -gAg ^{3'}	5'-ACA3'	5'-ACA3'	5'-gAC3'	⁵ '-ATg ³ '	5'-CCT3'	^{5'} -ggA ^{3'}	5'-ggA ^{3'}
		195	350									
		5'-ATg ^{3'}	5'-CAA3'									
		269	351									
		^{5'} -TAC ^{3'}	5'-ACC3'									
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.	110	135	200	115	130	165	125	235	145	95	210
PCR product										235	
Length of int.	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070
pos. control ¹											
5'-primer(s) ²	142	136	110	156	133	29	25	2 nd I	130	130	31
. ,	5'-ACC3'	5'-CAA3'	⁵ '-ggg ^{3'}	5'-CCC3'	5'-TCT3'	5'-CAT3'	⁵ '-Tgg ³ '	^{5'} -gCC ^{3'}	^{5'} -gAA ^{3'}	⁵ '-gAA ³ '	5'-TCA3'
		208								2 nd I	31
		5'-CCA3'								5'-TCC3'	5'-TCA3'
3'-primer(s) ³	165	166	164	181	163	71	54	54	165	54	86
	⁵ '-gTg ³ '	5'-CAA3'	5'-CAA3'	^{5'} -gTA ^{3'}	^{5'} -ggA ^{3'}	5'-TAC3'	5'-TAC3'	5'-TAC3'	^{5'} -gCC ^{3'}	5'-TAC3'	5'-CCA3'
		238								165	
		5'-CCg3'								^{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 2015-January-19, release 2.6.0,, 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 2015-January-19, release 2.6.0, www.ebi.ac.uk/ipd/kir. The sequence of the 3 terminal nucleotides of the primer is given in the antisense direction.



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		CELL L	IN	Ε,	VД	۱LI	D	ΑT	'IC	N	SI	HE	Ε	Т				
		KIR	G	en	oty	pii	ng	pr	im	ers	set	2						
						•		•		W								
			1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
			201324201	201437602	201448203	201324204	201666405	201666406	201666407	201324208	201324209	201324210	201324211	201324212	201324213	201324214	201324215	201666416
	IHV	/C cell line ¹																
1	9001		+	-	+	+	-	-	-	-	-	-	+	-	-	+	+	+
2	9280	LK707	-	+	-	+	+	-	+	+	+	-	-	+	+	+	+	+
3		E4181324	+	-	+	+	-	-	-	-	•	-	+	+	-	+	+	+
4		GU373	+	-	+	+	-	-	-	-	-	-	+	+	-	+	+	+
5		KAS011	+	-	+	+	+	+	-	+	-	-	-	+	+	+	+	+
6	9353		+	-	+	+	+	+	-	+	-	+	+	-	-	+	+	+
8	9020 9025		+	+	+	+	+	+	+	+	+	+	-	+	+	+	+	+
9	9025		+	-	+	+		-	-	÷	-	÷	÷	+	÷	+	+	+
10		LKT3	+	-	+	+	-	-	-	Ė	-	E	+	-	Ė	+	+	+
11		PITOUT	+	+	+	+	-	-	-	-	+	-	÷	+	-	+	÷	+
12	9052		+	+	+	+	+	-	+	-	+	+	-	+	-	+	+	+
13	9025	JESTHOM	+	+	-	+	+	-	+	-	+	+	+	+	-	+	+	+
14		OLGA	+	-	+	+	+	+	-	+	-	-	-	+	+	+	+	+
15	9075	DKB	+	-	+	+	-	-	-	-	-	-	+	-	-	+	+	+
16		SWEIG007	+	-	+	+	-	-	-	-	-	-	-	+	-	+	+	+
17	9282	CTM3953540	+	+	+	+	+	+	+	+	+	+	-	+	+	+	+	+
18	9257	32367	+	-	+	+	-	-	-	-	-	-	-	+	-	+	+	+
19		BM16	+	-	+	+	-	-	-	-	•	-	-	+	-	+	+	+
20		SLE005	+	-	+	+	-	-	-	<u> </u>	•	-	+	+	<u> </u>	+	+	+
21		AMALA	+	+	+	+	+	+	-	+	+	-	+	-	+	+	+	+
22		KOSE	+	+	+	+	-	-	-	-	+	-	-	+	-	+	+	+
23	9124		+	+	+	+	-	-	-	-	+	-	+	+	-	+	+	+
24 25	9035	JBUSH	+	-	+	+	-	-	-	-	-	÷	+	+	-	+	+	+
26		WT49	+		+	+	+	-	+	-	÷	÷	+	+	-	+	+	+
27		CH1007	Ŧ	+	+	+	+	-	Ŧ	+	Ŧ	+	÷	Ŧ	+	+	+	+
28		BEL5GB	+	+	-	+	+	-	+		+	+	-	+		+	+	+
29	9050		+	-	+	+	-	-	-	-	Ė	÷	-	÷	-	+	+	+
30	9021		<u> </u>	+	+	+	+	-	+	-	+	-	+	+	+	+	÷	+
31		DUCAF	÷	Ė	+	+	Ė	-	Ė	-	Ė	-	÷	+	÷	+	+	+
32	9297		+	-	+	+	-	-	-	-	-	-	+	+	-	+	+	+
33	9098	MT14B	+	-	+	+	-	-	-	-	-	-	+	-	-	+	+	+
34	9104		+	+	+	+	-	-	-	-	+	-	+	+	-	+	+	+
35		SSTO	+	+	+	+	-	-	-	-	+	-	+	+	-	+	+	+
36		KT17	+	-	+	+	+	+	-	+	-	+	-	+	-	+	+	+
37		HHKB	+	+	+	+	+	+	-	+	+	-	-	+	+	+	+	+
38	9099		+	+	+	+	+	-	+	+	+	+	+	-	-	+	+	+
39	9315		+	+	-	+	+	+	+	+	+	+	÷	+	<u> </u>	+	+	+
40		WHONP199	+	-	+	+	-	-	-	-	-	-	+	-	-	+	+	+
41		H0301 TAB089	+	+	-	+	+	-	+	Ė	+	+	+	÷	-	+	+	+
42		T7526	+	-	+	+	+	+	-	+		÷	+	+	+	+	+	+
44	9057		Ŧ	+	+	+	+	-	+	Ξ.	÷	+	Ξ.	+	Ξ	+	+	+
45		SHJO	+	+	+	+	+	-	+	-	+	-	+	-	+	+	+	+
46		SCHU	÷	÷	+	+	-	-	÷	-	Ė	-	÷	+	÷	+	+	+
47		TUBO	+	-	+	+	-	-	-	-	-	-	÷	+	-	+	+	+
48		TER-ND	+	-	+	+	-	-	-	-	-	-	-	+	-	+	+	+



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KIR Genotyping Pr 104.101-12 – including *Taq* polymerase 104.101-12u – without *Taq* polymerase

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	CE	LL LINE V	ΆL	\$	SH	ΙΕΙ	ΕT	•	
	KIR	Genotypin	g p	rin	nei	S	et²		
			Ĺ			Vе			
			17	18	19	20	21	22	23
			201324217	201324218	201324219	201324220	201324221	201437622	201324223
	IHV	/C cell line ¹							
1	9001		1-	+	+	+	-	-	-
2	9280	LK707	-	-	+	-	-	+	-
3	9011	E4181324	+	+	+	+	-	+	+
4		GU373	-	+	+	+	-	-	-
5		KAS011	+	+	+	+	-	+	-
6	9353		+	+	+	+	-	+	-
7	9020		+	+	+	+	-	+	+
8	9025		-	+	+	+	-	+	-
9	9026		-	+	+	+	-	-	-
10		LKT3	ᆣ	+	+	+	-	-	-
11		PITOUT	-	+	+	+	-	+	+
12	9052		+-	+	+	+	-	-	+
13		JESTHOM OLGA	+-	+	+	+	-	+	-
14			+	+	+	+	-	+	-
15	9075	SWEIG007	-	+	+	+	-	-	-
16 17		CTM3953540	+-	+	+	+	-	-	+
18		32367	+	+	+	+	-	+	+
19		32367 BM16	+-	+	+	+	-	-	+
20		SLE005	╀	+	+	+	-		-
21		AMALA	+	+	+	+	÷	+	÷
22		KOSE	i i	+	+	+	-	+	+
23	9124		+-	+	+	+	-	+	+
24		JBUSH	-	+	+	+	-	-	÷
25		IBW9	-	+	+	+	-	-	+
26		WT49	-	+	+	+	-	+	-
27		CH1007	-	+	+	+	-	-	-
28		BEL5GB	-	+	+	+	-	+	+
29	9050		-	+	+	+	-	-	+
30	9021	RSH	-	+	+	+	-	-	-
31		DUCAF	1-	+	+	+	-	-	-
32	9297	HAG	1-	+	+	+	-	-	-
33	9098	MT14B	1-	+	+	+	-	-	-
34	9104		-	+	+	+	-	+	-
35	9302	SSTO	-	+	+	+	-	+	-
36	9024	KT17	+	+	+	+	-	+	-
37	9065	HHKB	+	+	+	+	-	+	-
38	9099	LZL	-	+	+	+	-	+	-
39	9315		+	+	+	+	-	+	+
40		WHONP199	-	+	+	+	-	-	-
41		H0301	F	+	+	+	-	+	-
42		TAB089	-	+	+	+	-	-	-
43		T7526	+	+	+	+	-	+	-
44	9057		-	+	+	+	-	-	-
45		SHJO	-	+	+	+	-	-	-
46		SCHU	-	+	+	+	-	-	-
47		TUBO	-	+	+	+	-	-	-
48	9303	TER-ND	<u> </u>	+	+	+	-	-	-



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

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¹The provided cell line KIR specificities are retrieved from the www.ebi.ac.uk/ipd/kir 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 and 16 one 5'-primer and one 3'-primer were not possible to test.

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

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TRADEMARKS USED IN THIS DOCUMENT/PRODUCT

Olerup SSP[®] is a registered trademark of *Olerup* SSP AB. Qiagen[™] is a trademark of QIAGEN.

WARRANTY

Olerup SSP AB warrants its products to the original purchaser against defects in materials and workmanship under normal use and application. Olerup SSP AB's sole obligation under this warranty shall be to replace, at no charge, any product that does not meet the performance standards stated on the product specification sheet.

This warranty applies only to products that have been handled and stored in accordance with *Olerup* SSP AB's recommendations, and does not apply to products that have been the subject of alternation, misuse, or abuse.

All claims under this warranty must be directed to *Olerup* SSP AB in writing and must be accompanied by a copy of the purchaser's invoice. This warranty is in lieu of all other warranties, expressed or implied, including the warranties of merchantability and fitness for a particular purpose. In no case shall *Olerup* SSP AB be liable for incidental or consequential damages.

This product may not be reformulated, repacked or resold in any form without the written consent of *Olerup* SSP AB, Franzengatan 5, SE-112 51 Stockholm, Sweden.

Handle all samples as if capable of transmitting disease. All work should be performed wearing gloves and appropriate protection.

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.



KIR Genotyping Product Insert 104.101-12 – including *Taq* polymerase 104.101-12u – without *Taq* polymerase

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Lot No.: **8D0**

KIR Genotyping Product Insert 104.101-12 – including *Taq* polymerase 104.101-12u – without *Taq* polymerase

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

Lot No.: **8D0** Lot-specific information www.olerup.com

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