■LERUP SSP*
KIR Genotyping

Product Insert

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

Lot No.: **9E6** Lot-specific information

www.olerup.com

Olerup SSP® KIR Genotyping

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

104.101-12u – without *Taq* polymerase

Lot number: 9E6

Expiry date: 2019-11-01

Number of tests: 12 Number of wells per test: 25 + 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. 9E6.

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 (8D0)

Two wells have been added to KIR Genotyping, wells **25 to 26**.

The KIR Genotyping kit design, specificity and interpretation tables are based on IPD-KIR database 2.6.1.

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	Exchanged	Exchanged	Exchanged primer pair for the 2DL3*012 alleles.
24	Added	Added	Negative control moved to well 26, primer pair added for improved resolution of 2DL4 alleles.
25	New	New	Primer pair added for improved resolution of 2DL5B alleles.
26	-	-	Negative control added from well 24.

Change in revision R01 compared to R00:

- 1. The Protocol has been corrected, to reflect that two wells have been added to the kit. Change in revision R02 compared to R01:
- 1. A footnote has been added to the Specificity and Interpretation Tables, explaining that an "?" reflects that nucleotide sequence information is not available for the primer matching sequence. Change in revision R03 compared to R02:
- 1. Typing errors in the Interpretation and Specificity Tables and the Cell Line Validation sheet have been corrected.

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Well **26** contains <u>Negative Control primer pairs</u>, 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
	5'-CAg3'	^{5'} -CCT ^{3'}	5'-CCA3'	^{5'} -CCg ^{3'}
3'-primer	187	187	288	288
	^{5'} -ggT ^{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.

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

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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 26 PCR reactions in a 32 well cut PCR plate. Wells 27 to 32 are empty.

1	2	3	4	5	6	7	8
9	10	11	12	13	14	15	16
17	18	19	20	21	22	23	24
25	NC	empty	empty	empty	empty	empty	empty

The 32 well cut PCR plate is marked with 'KIR GENOTYP' in silver/gray ink.

Well No. 1 is marked with the Lot No. '9E6'.

Wells 1 to 25 – KIR Genotyping primers.

Well 26 – 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 32 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 February 2015¹ will be amplified by the primers in the KIR Genotyping SSP kit.

¹KIR alleles listed on the IPD KIR web page 2015-Febuary-17, release 2.6.1, www.ebi.ac.uk/ipd/kir.

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

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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_2O . 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/ μ l. **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

July 2018

Rev. No.: 03

104.101-12 – including Tag polymerase

For one KIR Genotyping typing, begin by adding to well No. 26, i.e. the well with the negative control primer pairs:

7 ul dH2O

3 μl PCR Master Mix complete with Taq,

then add at room temperature in a 0.5 ml tube:

 $30 \times 2 \mu I = 60 \mu I DNA (30 ng/\mu I)$

30 x 3 μ l = 90 μ l PCR Master Mix complete with Taq – mix well before taking your aliquot

Mix well, dispense 10 μ l of the DNA-PCR Master Mix-H₂O mixture into each of the 25 wells of an KIR Genotyping typing, i.e. wells 1 to 25. Cover the primer tray(s) with the provided

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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:

30 x 3 μ l = 90 μ l PCR Master Mix without Taq – mix well before taking your aliquot

2.4 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. 26, i.e. the well with the negative control primer pairs. Then add 7 μ l dH₂O to well 26.

Then add at room temperature to the 0.5 ml tube containing 90 + 2.4 - 3 = 89.4 μ l PCR Master Mix-*Tag* mixture:

30 x 2
$$\mu$$
l = 60 μ l DNA (30 ng/ μ l)
30 x 5 μ l – 2.4 μ l = 147.6 μ l dH₂O

Mix well, dispense 10 μ l of the DNA-PCR Master Mix-Taq-H₂O mixture into each of the 25 wells of an KIR Genotyping typing, i.e. wells 1 to 25. 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 to casting with ethidium bromide (10 mg/ml), 5 μ l 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.



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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 *Tag* polymerase contains:

Tag polymerase 0.4 unit per 10 μl SSP reaction

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

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 Taq 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 Tag 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 25+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 150 bp 225 bp	1070 bp	2DL2 2DL2 2DL2	004, 011 0010101-013 004, 011
3 ^{4,7}	90 bp 520 bp	1070 bp	2DL3 2DL3	01201-01202 0010101-011, 013-032
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, 0050101-0050104, 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 135 bp	1070 bp 1070 bp	2DS5 3DL1	001-012 0010101-002, 00401-00403, 0050101-009, 01501-044, 051-054,
15	200 bp	1070 bp	3DL2	056, 057, 059-077, 079 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 19 ⁴	165 bp 125 bp	1070 bp 1070 bp	2DP1 3DP1	00101-014 001-014



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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
244	100 bp	1070 bp	2DL4	00101-00602, 010, 01201-01202, 014-016, 018, 021-026
25	195 bp	1070 bp	2DL5B 3DP1	0020101-0020105, 0020106?, 0020107, 00202?, 004,00601, 00603-0070101, 0070102?, 0080101-00802, 00803?, 009- 01302, 01303?, 016-018? 001, 002, 004, 007, 0090101- 00902, 011-014?
26 ^{6,11}				,
∠0°,	-	-	-	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, 515 or 430 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-February-17, release 2.6.1, <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 26 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.

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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 26 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.

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

104.101-12 - including Taq polymerase

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PRIMER SPECIFICATION

Well No.	1	2	3	4	5	6	7	8	9	10	11	12	13
Length of spec.	145	65	90	200	155	1650	1650	100	205	130	215	200	110
PCR product		150	520										
		225											
Length of int.	800	1070	1070	1070	1070	430	515	1070	1070	1070	1070	1070	1070
pos. control ¹													
5'-primer(s) ²	130	208	332	208	226	-16	-16	165	140	236	229	234	142
	^{5'} -gAA ^{3'}	5'-CCA3'	5'-TCg3'	5'-CCg3'	5'-CCA3'	5'-TCA3'	5'-TCg3'	5'-gAg ^{3'}	^{5'} -gTA ^{3'}	5'-CAC3'	5'-CTA3'	5'-TCT3'	5'-ACC
	130	156	344				-16	165					
	5'-TAA3'	^{5'} -AAA ^{3'}	5'-CTg3'				5'-Tgg ^{3'}	^{5'} -gAA ^{3'}					
		262	344										
		5'-ggA ^{3'}	5'-CTg3'										
			378										
			5'-TAT3'										
3'-primer(s) ³	165	243	350	262	276	27	27	185	195	266	288	288	165
	5'-gCg3'	5'-ACA3'	5'-CAA3'	^{5'} -ggA ^{3'}	5'-gAg ^{3'}	5'-ACA3'	5'-ACA3'	5'-gAC3'	5'-ATg ^{3'}	5'-CCT3'	^{5'} -ggA ^{3'}	5'-ggA ^{3'}	⁵ '-gTg ³
		195	351										
		5'-ATg ^{3'}	5'-ACC3'										
		269	394										
		5'-TAC3'	5'-gAA3'										
Well No.	1	2	3	4	5	6	7	8	9	10	11	12	13
Well No.	14	15	16	17	18	19	20	21	22	23	24	25	7
Length of spec.	135	200	115	130	165	125	235	145	95	210	100	195	
PCR product									235				
Length of int.	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	1070	
pos. control ¹													
5'-primer(s) ²	136	110	156	133	29	25	2 nd I	130	130	31	324	up ⁴	
- p(-)	5'-CAA3'	^{5'} -ggg ^{3'}	5'-CCC3'	5'-TCT3'	5'-CAT3'	5'-Tgg ^{3'}	5'-gCC3'	^{5'} -gAA ^{3'}	5'-gAA3'	5'-TCA3'	5'-gTA3'	5'-AAg ³	•
	208								2 nd I	31			
	5'-CCA3'								5'-TCC3'	5'-TCA3'			
3'-primer(s) ³	166	164	181	163	71	54	54	165	54	86	344	1 st I	
3 -primer(s)	5'-CAA3'			5'-ggA ^{3'}	5'-TAC3'		5'-TAC3'		5'-TAC3'				-
	238	UAA	giA	997	140	140	IAG	900	165	JUA	. 99	IOA	
	230								100				

¹The internal positive control primer pairs amplify segments of the human growth hormone gene. The internal positive control bands are 1070, 800, 515 or 430 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.

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21

²The codon position matching the specificity-determining 3'-end of the primer is given. Codon numbering as on the KIR web page 2015-February-17, release 2.6.1, www.ebi.ac.uk/ipd/kir. The sequence of the 3 terminal nucleotides of the primer is given.

³The codon position matching the specificity-determining 3'-end of the primer. Codon numbering as on the KIR web page 2015-February-17, release 2.6.1, www.ebi.ac.uk/ipd/kir. The sequence of the 3 terminal nucleotides of the primer is given in the anti-sense direction.

⁴Primer located upstream of the 1st exon, 84 nucleotides upstream of codon -21.



5'-gCT3'

22

23

24

25

Well No.

5'-CCg³

14

15

16

17

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		CELL	LI	N	ΕV	VA	LI	D	ΑT	'IO	N	SI	HE	E	T				
		K	IR	Ge	ene	otv	nia	na	pr	ime	er :	set	2						
							<u> </u>	-9	Ρ.		W		-						
				1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
				1	2	6	4	2	9	7	8	6	0	_	7	က	4	5	9
			Prod. No.:	120	209	80	120	24	40	940	120	120	121	121	121	121	121	121	7,7
			ط. ا	324	437		32	999	999	999	35	32	35	35	35	35	324	324	999
			입	201324201	201437602	201777803	201324204	201666405	201666406	201666407	201324208	201324209	201324210	201324211	201324212	201324213	201324214	201324215	201666416
	IHV	/C cell line ¹		- ,		.,	- ,	- ,	- ,	- 1		.,	- ,		.,		- 1 1		
1	9001			+	-	+	+	-	-	-	-	-	-	+	-	-	+	+	+
2	9280	LK707		-	+	-	+	+	-	+	+	+	-	-	+	+	+	+	+
3		E4181324		+	-	+	+	-	-	-	-	-	-	+	+	-	+	+	+
4		GU373		+	-	+	+	-	-	-	-	-	-	+	+	-	+	+	+
5		KAS011		+	-	+	+	+	+	-	+	-	-	-	+	+	+	+	+
6	9353			+	-	+	+	+	+	-	+	-	+	+	-	-	+	+	+
7 8	9020 9025			+	+	+	+	+	+	+	+	+	+	-	+	+	+	+	+
9	9025			+	-	+	+			-		+		÷	+		+	+	+
10		LKT3		+	-	+	+	-	-	-	-	-	-	+	-	-	+	+	+
11		PITOUT	\dashv	+	+	+	+	-	-	-	-	+	-	÷	+	-	+	+	+
12	9052			+	+	+	+	+	-	+	-	+	+	-	+	-	+	+	+
13		JESTHOM		+	+	-	+	+	-	+	-	+	+	+	+	-	+	+	+
14		OLGA		+	-	+	+	+	+	-	+	•	-	-	+	+	+	+	+
15	9075			+	-	+	+	-	-	-	Ŀ	-	-	+	-	-	+	+	+
16		SWEIG007		+	-	+	+	Ŀ	÷	-	÷	-	Ŀ	-	+	÷	+	+	+
17 18		CTM3953540 32367		+	+	+	+	+	+	+	+	+	+	÷	+	+	+	+	+
19		BM16		+	-	+	+	H	÷	-	H		H	÷	+	÷	+	+	+
20		SLE005		+	-	+	+	-	-	-	-	-	-	+	+	-	+	+	+
21		AMALA		+	+	+	+	+	+	-	+	+	-	+	-	+	+	+	+
22	9056	KOSE		+	+	+	+	-	-	-	-	+	-	-	+	-	+	+	+
23	9124	IHL		+	+	+	+	-	-	-	-	+	-	+	+	-	+	+	+
24		JBUSH		+	-	+	+	-	-	-	-	•	-	+	+	-	+	+	+
25		IBW9		+	-	+	+	-	-	-	Ŀ	-	-	+	+	-	+	+	+
26		WT49		+	+	+	+	+	-	+	+	+	÷	÷	+	+	+	+	+
27 28		CH1007 BEL5GB		+	+	+	+	+	-	+	Ŀ	+	+	+	+		+	+	+
29	9050			+	+	+	+	+	-	+	-	+	+	÷	+		+	++	+
30	9021			+	+	+	+	+	-	+	-	+		+	+	+	+	+	+
31		DUCAF		+	÷	+	+	÷	-	-	-	÷	-	÷	+	÷	+	÷	+
32	9297			+	-	+	+	-	-	-	-	-	-	+	+	-	+	+	+
33	9098	MT14B		+	-	+	+	-	-	-	-	-	-	+	-	-	+	+	+
34	9104			+	+	+	+	-	-	-	-	+	-	+	+	-	+	+	+
35		SSTO		+	+	+	+	-	-	-	-	+	-	+	+	-	+	+	+
36		KT17		+	-	+	+	+	+	-	+	Ŀ	+	-	+	-	+	+	+
37 38	9065	HHKB		+	+	+	+	+	+	-	+	+	-	÷	+	+	+	+	+
39	9099			+	+	+	+	+	+	+	+	+	+	+	+		+	+	+
40		WHONP199		+	-	+	+	-	-	-	-	-	-	+	-		+	+	+
41		H0301		+	+	÷	+	+	-	+	-	+	+	+	-	-	+	+	+
42		TAB089		+	-	+	+	-	-	-	-	-	-	-	+	-	+	+	+
43		T7526		+	-	+	+	+	+	-	+	-	-	+	-	+	+	+	+
44	9057			+	+	+	+	+	-	+	-	+	+	-	+	-	+	+	+
45		SHJO		+	+	+	+	+	-	+	-	+	-	+	-	+	+	+	+
46		SCHU		+	-	+	+	-	-	-	-	-	-	+	+	-	+	+	+
47		TUBO		+	-	+	+	-	-	-	-	-	-	-	+	-	+	+	+
48	9303	TER-ND		+	-	+	+					-			+		+	+	+

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

Lot No.: **9E6** Lot-specific information www.olerup.com

Strain S	+ - + + + - + + + + - + 201777824	25
IHWC cell line1	+ - + + + - + + + + - + 201777824	+ + + + 201777827
IHWC cell line ¹	+ - + + + - + + + + - + 201777824	+ + + + 201777827
IHWC 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 +	+ + + - +	+ +
13 9025 JESTHOM - + + + - + - + - 14 9071 OLGA + + + + + - + - + - 15 9075 DKB - + + + + 16 9037 SWEIG007 - + + + + + + 17 9282 CTM3953540 + + + + + + + 18 9257 32367 - + + + + + 19 9038 BM16 - + + + + 19 9059 SLE005 - + + + + 12 9064 AMALA + + + + + - + - + - 12 9056 KOSE - + + + + - + + + - + + + - + + 12	+ + + + - + +	+ - -
14 9071 OLGA + + + - + - + -<	+	-
15 9075 DKB - + + - </th <th>+</th> <th>- - +</th>	+	- - +
16 9037 SWEIG007 - + + + - - + 17 9282 CTM3953540 + - - + + + - - - + + - - - - + + - - - + + - - + + - - + + - - + + - - + + - - + + - - + + - - - + - - - + -	+	- +
17 9282 CTM3953540 + + + + - + + + - + + + - - + + - - + + - - + + -	+	+
18 9257 32367 - + + + + + - + + + + + + 19 9038 BM16 - + + + + 20 9059 SLE005 - + + + + 21 9064 AMALA + + + + + - + - + - - + + + + - + - 22 9056 KOSE - + + + + - + + - + + -	-	+
19 9038 BM16 - + + + - - 20 9059 SLE005 - + + + - - 21 9064 AMALA + + + + + - + 22 9056 KOSE - + + + + + +	-	_
20 9059 SLE005 - + + + 21 9064 AMALA + + + + - + - + - 22 9056 KOSE - + + + - + - + +		
21 9064 AMALA + + + + - + - 22 9056 KOSE - + + + - + +		-
22 9056 KOSE - + + + - + +	+	-
	+	+
00 0404	. -	+
23 9124 HL - + + + - + +	+	+
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 - + + +	-	-



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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 and 7 one of the 5'-primers was not possible to test, in primer mix 24 one 3'-primer was not possible to test, and in primer solution 2, 3 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.

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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 *Tag* polymerase are stable for 33 months from the date of manufacture.

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

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