

Lot No.: **40E**

Lot-specific information

## **Olerup SSP<sup>®</sup> DR low resolution**

<b>Product number:</b>	101.101-48/12 – including <i>Taq</i> pol. 101.101-48u/12u – without <i>Taq</i> pol.
<b>Lot number:</b>	40E
<b>Expiry date:</b>	2010-May-01
<b>Number of tests:</b>	48 tests – Product No. 101.101-48 12 tests – Product No. 101.101-12
<b>Number of wells per test:</b>	23 + 1
<b>Storage - pre-aliquoted primers:</b>	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. 40E.**

### **CHANGES COMPARED TO THE PREVIOUS *OLERUP SSP*<sup>®</sup> DR LOW RESOLUTION LOT**

The DR low resolution specificity and interpretation tables have been updated for the HLA-DRB1 alleles described since the previous *Olerup SSP*<sup>®</sup> DR low resolution lot was made (**Lot No. Y39**).

The DR low resolution primer set is unchanged compared to the previous lot.

Changes in revision R01 compared to R00:

1. The length of the specific PCR product in primer mix 12 has been corrected to 205 bp in the Interpretation Table.

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Well **24** contains Negative Control primer pairs, that will amplify more than 95% of the *Olerup* SSP<sup>®</sup> HLA Class I, DRB, DQB1 and DPB1 amplicons as well as the amplicons generated by control primer pairs.

PCR product sizes range from 75 to 430 base pairs.  
 The PCR product generated by the control primer pair is 430 base pairs.

Length of PCR product	105	200	105	80	75	80
<b>5'-primer<sup>1</sup></b>	<b>164</b>	<b>340</b>	<b>440</b>	<b>45</b>	<b>45</b>	<b>43</b>
	5'-CAC <sup>3'</sup>	5'-Agg <sup>3'</sup>	5'-TTA <sup>3'</sup>	5'-Tg g <sup>3'</sup>	5'-Tg g <sup>3'</sup>	5'-Tg g <sup>3'</sup>
<b>3'-primer<sup>2</sup></b>	<b>231</b>	<b>2<sup>nd</sup> I</b>	<b>507</b>	<b>59</b>	<b>58</b>	<b>57</b>
	5'-TgC <sup>3'</sup>	5'-AAA <sup>3'</sup>	5'-TTg <sup>3'</sup>	5'-CTC <sup>3'</sup>	5'-ggC <sup>3'</sup>	5'-CTC <sup>3'</sup>
<b>A*</b>	<b>+</b>	<b>+</b>	<b>+</b>			
<b>B*</b>	<b>+</b>	<b>+</b>	<b>+</b>			
<b>Cw*</b>	<b>+</b>	<b>+</b>	<b>+</b>			
<b>DRB1</b>				<b>+</b>	<b>+</b>	
<b>DRB3</b>				<b>+</b>	<b>+</b>	
<b>DRB5</b>				<b>+</b>		
<b>DQB1</b>					<b>+</b>	
<b>DPB1</b>						<b>+</b>

<sup>1</sup>The nucleotide position for HLA class I genes and the codon for HLA class II genes, in the 2<sup>nd</sup> or 3<sup>rd</sup> exon, matching the specificity-determining 3'-end of the primer is given. Nucleotide and codon numbering as on the [www.ebi.ac.uk/imgt/hla](http://www.ebi.ac.uk/imgt/hla) web site. The sequence of the 3 terminal nucleotides of the primer is given.

<sup>2</sup>The nucleotide position for HLA class I genes and the codon for HLA class II genes, in the 2<sup>nd</sup> or 3<sup>rd</sup> exon or the 2<sup>nd</sup> 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](http://www.ebi.ac.uk/imgt/hla) web site. The sequence of the 3 terminal nucleotides of the primer is given.

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## PRODUCT DESCRIPTION

### DR low resolution

#### CONTENT

The primer set contains 5'- and 3'-primers for grouping the DRB1\*0101 to DRB1\*1001 alleles into the corresponding serological groups DR1 to DR18 as well as primer pairs for recognizing the DRB3, DRB4 and DRB5 groups of alleles.

#### PLATE LAYOUT

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

<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>	<b>7</b>	<b>8</b>
<b>9</b>	<b>10</b>	<b>11</b>	<b>12</b>	<b>13</b>	<b>14</b>	<b>15</b>	<b>16</b>
<b>17</b>	<b>18</b>	<b>19</b>	<b>20</b>	<b>21</b>	<b>22</b>	<b>23</b>	<b>24</b>

Wells 1 to 23 – DR low resolution primers.

Well 24 – Negative Control.

The 24 well cut PCR plate is marked with 'DR low'.

Well No. 1 is marked with the Lot No. '40E'.

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.

#### INTERPRETATION

Only HLA-DRB alleles will be amplified by the 23 wells of the DR low resolution primer set, **wells 1 to 23**. Thus, the interpretation of DR low resolution typings is not influenced by other HLA class II genes.

#### UNIQUELY IDENTIFIED ALLELES

All the HLA-DRB1, -DRB3, -DRB4<sup>1</sup> and -DRB5 alleles, i.e. **DRB1\*010101 to 1002, DRB3\*010101 to DRB3\*0303, DRB4\*010101 to DRB4\*0107, DRB5\*010101 to DRB5\*0205**, recognized by the HLA Nomenclature Committee in April 2008<sup>2</sup> will be amplified by the primers in the DR low resolution SSP kit. The HLA-DRB alleles will be grouped into their corresponding serological specificities<sup>3</sup>.

<sup>1</sup>The DRB4\*0201N and DRB4\*0301N null alleles will not be amplified by the DR low resolution primer set.

<sup>2</sup>DRB alleles listed on the IMGT/HLA web page 2008-April-08, release 2.21.0, [www.ebi.ac.uk/imgt/hla](http://www.ebi.ac.uk/imgt/hla).

<sup>3</sup>The DRB1\*0809, DRB1\*0821 and DRB1\*1415 alleles yield identical amplification patterns except for the specific PCR product yielded by the DRB3 gene in linkage disequilibrium with the DRB1\*1415 allele.

The DRB1\*0820, DRB1\*1318, DRB1\*1347 and DRB1\*1355 alleles yield identical amplification patterns except for the specific PCR product yielded by the DRB3 gene in linkage disequilibrium with the DRB1\*1318, DRB1\*1347 and DRB1\*1355 alleles.

The DRB1\*0831 and DRB1\*1167 alleles yield identical amplification patterns except for the specific PCR product yielded by the DRB3 gene in linkage disequilibrium with the DRB1\*1167 allele.

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## SPECIFICITY TABLE

### DR low resolution primer set

Specificities and sizes of the PCR products of the 23 primer mixes of the DR low resolution primer set

Primer Mix	Size of spec. PCR product <sup>1</sup>	Size of control band <sup>2</sup>	DR serology <sup>3</sup>	Amplified HLA-DRB <sup>4</sup> alleles
<b>1</b>	205 bp	<b>515 bp</b>	1	010101-010204, 0104-0119
<b>2</b>	195 bp	430 bp	103	0103
<b>3</b>	215, 260 bp	430 bp	15	150101-1527
<b>4</b>	210 bp	430 bp	16	160101-160502, 1607-1613N
<b>5</b>	220 bp	430 bp	3, 17, 18, 11	030101-0339, 1107, 1153
<b>6<sup>5,6</sup></b>	80, 210 bp	430 bp	3, 17, 11, 13, 14	030101-030106, 0304-0306, 0308-0316, 0318-0320, 0322, 0323, 0325, 0326, 0328, 0330, 0331, 0333, 0334, 0336, 0337, 110201-1103, 111101-111102, 111401-111402, 1116, 1120, 1121, 1136, 1140, 1141, 1148, 1159, 1163, 116501-116502, 1168, 130101-1304, 1308, 1310, 1315-1317, 1319, 1320, 1322-1324, 1327-1329, 1331-1341, 1343, 1345, 1348, 1351-1354, 1357, 1359, 1361, 1363-1366, 1368-1376, 1378-1381, 1383, 1416, 1419, 1421
<b>7<sup>5,6</sup></b>	80, 210 bp	430 bp	3, 18, 11, 13, 14	030201-0303, 0327, 0329, 0338, 1126, 1134, 1315, 1319, 1326, 1344, 1353, 1357, 1402-140302, 1406, 1409, 1412, 1413, 1417-1421, 1424, 1427, 1429, 1430, 1433, 1440, 1441, 1447-1449, 1451, 1463, 1467, 1477
<b>8<sup>5,6</sup></b>	100, 175 bp	430 bp	4	040101-0473
<b>9</b>	205, 230 bp	430 bp	7, 13, 14	070101-070102, 0703-0715, 1317, 1450
<b>10</b>	165, 215, 250	<b>515 bp</b>	8,	080101-080203, 080302-0819,

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	bp		11, 12, 14	0821-0834, 1167, 1204, 1216, 1411, 1415, 1468
<b>11<sup>5</sup></b>	85, 130, 180 bp	430 bp	3, 9, 11	0308, 090102-0906, 1107, 1153
<b>12</b>	205 bp	430 bp	10	100101-1002
<b>13<sup>5,8</sup></b>	100, 170 bp	430 bp	11, 3, 8	0308, 0831, 110101-1168
<b>14<sup>5</sup></b>	85, 105 bp	430 bp	12, 8	0832, 120101-120202, 120302-1216
<b>15</b>	210 bp	430 bp	13, 8, 11, 14	0820, 110101-110404, 110601-110602, 110801-111202, 111401-1116, 1118-1121, 1123-1125, 112701- 1133, 1135-1151, 115401- 115402, 1156-1166, 1168, 130101-1308, 1310-1316, 1318- 1343, 1345-1383, 140301-140302, 1412, 1416, 1419, 1421, 1422, 1425, 1427, 1440, 1453, 1463, 1467, 1469, 1474, 1477
<b>16<sup>6</sup></b>	195, 210 bp	430 bp	13, 8, 11, 12, 14	080101-080203, 080401-0809, 0811, 0816, 0817, 0820-0822, 0824, 0826, 0828, 0831, 110101-110602, 1109-111202, 111401-1116, 1120, 1121, 1123- 1125, 112701-1130, 1132, 1133, 1135-1141, 1143, 1144, 1146- 1151, 115401-1156, 1158-1163, 116501-1168, 120201-120202, 1213, 1215, 1216, 130101-130201, 130203, 1304- 130502, 130701-1309, 1311, 131401-1324, 1326-1329, 1331, 1332, 1334-1336, 1338-1343, 1345-1355, 1357, 1359, 1361- 1365, 1367-1376, 1378-1380, 1383, 1415, 1416, 1422, 1424, 1425, 1427, 1437, 1453, 1473
<b>17</b>	175 bp	430 bp	3, 8, 13, 14	030101-0307, 0309, 0311-0339, 0820, 130101-1316, 1318-1342, 1344,

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				1346-1366, 1368-1383, 1402-140302, 140501-1406, 1409, 1412-1414, 1417-1421, 142301, 1424, 1427, 1429, 1430, 1433, 1436, 1437, 1440-1445, 1447, 1448, 1451, 1456, 1459, 1463-1467, 1477
<b>18<sup>5-7</sup></b>	100, 140 bp	430 bp	14, 4, 8, 13	0462, 0469, 0473, 0808, 1345, 140101-140102, 1404, 140701- 140702, 1410, 1416, 1422, 1425, 1426, 1428, 1431-143202, 1435, 1437-1439, 1449, 1450, 1453- 1455, 1457, 1458, 1460-1462, 1468-1471, 1473-1476
<b>19<sup>5-9</sup></b>	110, 135, 170 bp	<b>515 bp</b>	14, 3, 9, 11, 12, 13, 15	0308, 0310, 090102-0902, 0904-0906, 110404, 1109, 1110, 111202, 1113, 1116, 1117, 1120, 1122, 1140, 1146, 1152, 1158, 1159, 1204, 1343, 140101-1402, 1404-1411, 1413, 1414, 1416-142302, 1426, 1428- 1436, 1438, 1439, 1441, 1443- 1452, 1454-1457, 1459-1462, 1464, 1465, 1468, 1470-1476, 1527
<b>20<sup>5-7</sup></b>	110, 175, 225 bp	430 bp	14, 3, 8, 11, 13, 15, 16	0310, 0809, 0820, 0821, 0832, 1113, 1117, 1123, 1125, 1131, 1145, 1152, 1155, 1164, 1313, 1318, 1343, 1345, 1347, 1355, 140101-140103, 140301-140503, 140701-1408, 1410-1412, 1414- 1416, 1418, 1422-142302, 1425- 1428, 1431-143202, 1434-1436, 1438-1440, 1442-1445, 1449, 1450, 1453-1477, 1521 <sup>weakly</sup> , 1604 <sup>weakly</sup>
<b>21</b>	230 bp	430 bp	52	DRB3*01010201-0111, 0201- 0223, 030101-0303
<b>22<sup>10</sup></b>	215 bp	430 bp	53	DRB4*01010101-0107
<b>23</b>	260 bp	430 bp	51	DRB5*010101-0113, 0202-0205

101.101-48/12 – including *Taq* polymerase101.101-48u/12u – without *Taq* polymeraseLot No.: **40E**

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<sup>1</sup>Alleles 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 DR low resolution SSP subtypings.

When the primers in a primer mix can give rise to specific PCR products of more than one length this is indicated if the size difference is 20 base pairs or more. Size differences shorter than 20 base pairs are not given. For high resolution SSP kits the respective length of the specific PCR product(s) of the alleles amplified by these primer mixes are given.

Nonspecific amplifications, i.e. a ladder or a smear of bands, may sometimes be seen. GC-rich primers have a higher tendency of giving rise to nonspecific amplifications than other primers, e.g. the primers in wells 3, 18, 19 and 20.

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.

<sup>2</sup>The internal positive control primer pairs amplify segments of the human growth hormone gene. The two different control primer pairs give rise to either an internal positive control band of 430 base pairs, for most wells, or a band of 515 base pairs, for some wells.

Well number 1 contains the primer pair giving rise to the longer, 515 bp, internal positive control band in order to help in the correct orientation of the DR low resolution typing.

In addition, wells number 10 and 19 contain the primer pair giving rise to the longer, 515 bp, internal positive control band in order to allow kit identification.

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

<sup>3</sup>The serological reactivity of all DRB alleles is not known. In this table we use the information in the HLA Dictionary 2004 on the [www.ebi.ac.uk/imgt/hla](http://www.ebi.ac.uk/imgt/hla) web site and the information available at the [www.worldmarrow.org](http://www.worldmarrow.org) web site and have also inferred the serological grouping from the naming of the sequence-defined allele.

<sup>4</sup>For several DRB alleles only partial second exon nucleotide sequences are available. In these instances it is not known whether some of the primers of the SSP set are completely matched with the target sequences or not. We assume that unknown sequences in the first hyperpolymorphic region of the second exon of DRB alleles are conserved within allelic groups and that unknown sequences of codons 87 to 92 are identical with the DRB1\*0101 consensus sequence.

The DRB1\*0809, DRB1\*0821 and DRB1\*1415 alleles yield identical amplification patterns except for the specific PCR product yielded by the DRB3 gene in linkage disequilibrium with the DRB1\*1415 allele.

The DRB1\*0820, DRB1\*1318, DRB1\*1347 and DRB1\*1355 alleles yield identical amplification patterns except for the specific PCR product yielded by the DRB3 gene in linkage disequilibrium with the DRB1\*1318, DRB1\*1347 and DRB1\*1355 alleles.

The DRB1\*0831 and DRB1\*1167 alleles yield identical amplification patterns except for the specific PCR product yielded by the DRB3 gene in linkage disequilibrium with the DRB1\*1167 allele.

<sup>5</sup>Specific PCR fragments shorter than 125 base pairs have a lower intensity and are less sharp than longer PCR bands.

<sup>6</sup>Individual alleles can give rise to two differently sized specific PCR fragments in these primer mixes.

<sup>7</sup>These primer mixes may give rise to nonspecific amplifications.

<sup>8</sup>Primer mix 13 may give rise to a primer oligomer formation.

<sup>9</sup>Primer mix 19 has a tendency of primer oligomer formation and also has an intense primer cloud due to the high number of primers present in the primer mix.

<sup>10</sup>The DRB4\*01030102N allele is amplified by the primer pair in well No. 22, whereas the DRB4\*0201N and DRB4\*0301N null alleles are not amplified by this primer pair.

<b>INTERPRETATION TABLE</b>													
<b>DR low resolution SSP typing</b>													
<b>Amplification patterns of the DRB1*0101 to DRB1*1002 alleles</b>													
		<b>Well</b>											
		<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>	<b>7</b>	<b>8</b>	<b>9</b>	<b>10</b>	<b>11</b>	<b>12</b>
<b>Length of spec.</b>		<b>205</b>	<b>125</b>	<b>215</b>	<b>210</b>	<b>220</b>	<b>80</b>	<b>80</b>	<b>100</b>	<b>205</b>	<b>165</b>	<b>85</b>	<b>205</b>
<b>PCR product(s)</b>				<b>260</b>			<b>210</b>	<b>210</b>	<b>175</b>	<b>230</b>	<b>215</b>	<b>130</b>	
											<b>250</b>	<b>180</b>	
<b>Length of int.</b>		<b>515</b>	<b>430</b>	<b>430</b>	<b>430</b>	<b>430</b>	<b>430</b>	<b>430</b>	<b>430</b>	<b>430</b>	<b>515</b>	<b>430</b>	<b>430</b>
<b>pos. control<sup>1</sup></b>													
<b>5'-primer(s)<sup>2</sup></b>		<b>14</b>	<b>14</b>	<b>13</b>	<b>13</b>	<b>5</b>	<b>13</b>	<b>13</b>	<b>13</b>	<b>14</b>	<b>16</b>	<b>26</b>	<b>31</b>
		5'-gAA <sup>3'</sup>	5'-gAA <sup>3'</sup>	5'-Agg <sup>3'</sup>	5'-Agg <sup>3'</sup>	5'-g TC <sup>3'</sup>	5'-g TC <sup>3'</sup>	5'-g TC <sup>3'</sup>	5'-A CA <sup>3'</sup>	5'-AT A <sup>3'</sup>	5'-gT T <sup>3'</sup>	5'-TAT <sup>3'</sup>	5'-gC g <sup>3'</sup>
					<b>13</b>		<b>16</b>		<b>13</b>	<b>14</b>	<b>16</b>	<b>58</b>	
					5'-AAg <sup>3'</sup>		5'-gT T <sup>3'</sup>		5'-A CC <sup>3'</sup>	5'-AT A <sup>3'</sup>	5'-gT T <sup>3'</sup>	5'-gAg <sup>3'</sup>	
									<b>13</b>	<b>16</b>			
									5'-A TA <sup>3'</sup>	5'-gT T <sup>3'</sup>			
<b>3'-primer(s)<sup>3</sup></b>		<b>67</b>	<b>67</b>	<b>70</b>	<b>67</b>	<b>73</b>	<b>26</b>	<b>28</b>	<b>33</b>	<b>71</b>	<b>58</b>	<b>57</b>	<b>86</b>
		5'-gAg <sup>3'</sup>	5'-gAT <sup>3'</sup>	5'-CTg <sup>3'</sup>	5'-gAA <sup>3'</sup>	5'-g gC <sup>3'</sup>	5'-g gT <sup>3'</sup>	5'-CT C <sup>3'</sup>	5'-gTg <sup>3'</sup>	5'-CTC <sup>3'</sup>	5'-C CT <sup>3'</sup>	5'-C gA <sup>3'</sup>	5'-C AC <sup>3'</sup>
		<b>71</b>		<b>71</b>	<b>67</b>	<b>73</b>	<b>71</b>	<b>70</b>	<b>58</b>	<b>73</b>	<b>74</b>	<b>73</b>	<b>86</b>
		5'-gC g <sup>3'</sup>		5'-CgC <sup>3'</sup>	5'-gAg <sup>3'</sup>	5'-g gC <sup>3'</sup>	5'-g CT <sup>3'</sup>	5'-CTg <sup>3'</sup>	5'-C gg <sup>3'</sup>	5'-g gC <sup>3'</sup>	5'-CAg <sup>3'</sup>	5'-g gC <sup>3'</sup>	5'-C CA <sup>3'</sup>
				<b>86</b>	<b>72</b>					<b>77</b>	<b>86</b>	<b>78</b>	
				5'-C CA <sup>3'</sup>	5'-gC g <sup>3'</sup>					5'-A AT <sup>3'</sup>	5'-C AC <sup>3'</sup>	5'-CAC <sup>3'</sup>	
										<b>78</b>			
										5'-CAC <sup>3'</sup>			
<b>Well No.</b>	<b>DR</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>	<b>7</b>	<b>8</b>	<b>9</b>	<b>10</b>	<b>11</b>	<b>12</b>
<b>DRB1 allele<sup>4</sup></b>	<b>ser<sup>5</sup></b>												
<b>*010101-010204, 0104-0119</b>	<b>1</b>	<b>1</b>											
<b>*0103</b>	<b>103</b>		<b>2</b>										
<b>*030101-030106, 0304-0306, 0309, 0311-0316, 0318-0320, 0322, 0323, 0325, 0326, 0328, 0330, 0331, 0333, 0334, 0336, 0337</b>	<b>17</b>					<b>5</b>	<b>6</b>						
<b>*030201-0303, 0327, 0329, 0338</b>	<b>18</b>					<b>5</b>		<b>7</b>					
<b>*0307, 0317, 0321, 0324, 0332, 0335, 0339</b>	<b>-</b>					<b>5</b>							
<b>Well No.</b>	<b>DR</b>	<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>	<b>7</b>	<b>8</b>	<b>9</b>	<b>10</b>	<b>11</b>	<b>12</b>



<b>INTERPRETATION TABLE</b>												
<b>DR low resolution SSP typing</b>												
<b>Amplification patterns of the DRB1*0101 to DRB1*1002 alleles</b>												
<b>Well</b>												
13	14	15	16	17	18	19	20	21	22	23		
100	85	210	195	175	100	110	110	230	215	260		Length of spec.
170	105		210		140	135	175					PCR product(s)
						170	225					
430	430	430	430	430	430	515	430	430	422	430		Length of int.
												pos. control <sup>1</sup>
13	16	10	10	13	37	26	13	10	28	13		5'-primer(s) <sup>2</sup>
5'-g TC <sup>3'</sup>	5'-gT T <sup>3'</sup>	5'-g CT <sup>3'</sup>	5'-g CT <sup>3'</sup>	5'-g TC <sup>3'</sup>	5'-g TA <sup>3'</sup>	5'-g TA <sup>3'</sup>	5'-g TC <sup>3'</sup>	5'-g CT <sup>3'</sup>	5'-g AT <sup>3'</sup>	5'-g TA <sup>3'</sup>		
16		13	13		37	34	34	10		13		
5'-gT C <sup>3'</sup>		5'-g TC <sup>3'</sup>	5'-g TC <sup>3'</sup>		5'-g TT <sup>3'</sup>	5'-CAG <sup>3'</sup>	5'-CAG <sup>3'</sup>	5'-g CT <sup>3'</sup>		5'-g TA <sup>3'</sup>		
38			16									
5'-C gT <sup>3'</sup>			5'-gT T <sup>3'</sup>									
			16									
			5'-gT C <sup>3'</sup>									
58	30	70	67	58	57	57	57	73	87	87		3'-primer(s) <sup>3</sup>
5'-C CT <sup>3'</sup>	5'-gTg <sup>3'</sup>	5'-gTC <sup>3'</sup>	5'-gAA <sup>3'</sup>	5'-C gg <sup>3'</sup>	5'-C Ag <sup>3'</sup>	5'-C Ag <sup>3'</sup>	5'-C Ag <sup>3'</sup>	5'-g gC <sup>3'</sup>	5'-CTC <sup>3'</sup>	5'-CTC <sup>3'</sup>		
58	38	71	67	58	71	58	60	73				
5'-C CT <sup>3'</sup>	5'-CAG <sup>3'</sup>	5'-g CT <sup>3'</sup>	5'-gAA <sup>3'</sup>	5'-C Ag <sup>3'</sup>	5'-CgC <sup>3'</sup>	5'-C CT <sup>3'</sup>	5'-gTg <sup>3'</sup>	5'-g gC <sup>3'</sup>				
			71			70	70					
			5'-CTC <sup>3'</sup>			5'-CTg <sup>3'</sup>	5'-T CC <sup>3'</sup>					
			71			70	74					
			5'-CgC <sup>3'</sup>			5'-T CC <sup>3'</sup>	5'-CAG <sup>3'</sup>					
13	14	15	16	17	18	19	20	21	22	23	DR	Well No.
											ser <sup>5</sup>	DRB1 allele <sup>4</sup>
											1	*010101-010204, 0104-0119
											103	*0103
				17							17	*030101-030106, 0304-0306, 0309, 0311-0316, 0318-0320, 0322, 0323, 0325, 0326, 0328, 0330, 0331, 0333, 0334, 0336, 0337
				17							18	*030201-0303, 0327, 0329, 0338
				17							-	*0307, 0317, 0321, 0324, 0332, 0335, 0339
13	14	15	16	17	18	19	20	21	22	23	DR	Well No.

Lot No.: **40E**

Lot-specific information

Length of spec.		205	125	215	210	220	80	80	100	205	165	85	205
PCR product(s)				260			210	210	175	230	215	130	
Well No.	DR	1	2	3	4	5	6	7	8	9	10	11	12
*0308	-					5	6					11	
*0310	-					5	6						
*040101-0461, 0463-0468, 0470-0472	4								8				
*0462, 0469, 0473	-								8				
*070101-070102, 0703-0715	7									9			
*080101-080203, 080401-0807, 0811, 0816, 0817, 0822, 0824, 0826, 0828	8										10		
*080302, 0810, 0812-0815, 0818, 0819, 0823, 0825, 0827, 0829, 0830, 0833, 0834	8										10		
*0808	-										10		
*0809, 0821, 1415	8										10		
*0820, 1318, 1347, 1355	8,13												
*0831, 1167	-										10		
*0832	-										10		
*090102-0902, 0904-0906	9											11	
*0903	9											11	
*100101-1002	10												12
*110101-110107, 110401-110403, 110601-110602, 111201, 1115, 1124, 112701-1130, 1132, 1133, 1135, 1137-1139, 1143, 1144, 1147, 1149-1151, 115401-115402, 1156, 1160-1162, 1166	11												
*110201-1103, 111101-111102, 111401-111402, 1121, 1136, 1141, 1148, 1163, 116501-116502, 1168	11						6						
Well No.	DR	1	2	3	4	5	6	7	8	9	10	11	12

Lot No.: **40E**

Lot-specific information

100	85	210	195	175	100	110	110	230	215	260		Length of spec.
170	105		210		140	135	175					PCR product(s)
						170	225					
13	14	15	16	17	18	19	20	21	22	23	DR	Well No.
13						19					-	*0308
						19	20				-	*0310
											4	*040101-0461, 0463-0468, 0470-0472
					18						-	*0462, 0469, 0473
											7	*070101-070102, 0703-0715
			16								8	*080101-080203, 080401-0807, 0811, 0816, 0817, 0822, 0824, 0826, 0828
											8	*080302, 0810, 0812-0815, 0818, 0819, 0823, 0825, 0827, 0829, 0830, 0833, 0834
			16		18						-	*0808
			16				20				8	*0809, 0821, 1415
		15	16	17			20				8,13	*0820, 1318, 1347, 1355
13			16								-	*0831, 1167
	14						20				-	*0832
						19					9	*090102-0902, 0904-0906
											9	*0903
											10	*100101-1002
13		15	16								11	*110101-110107, 110401-110403, 110601-110602, 111201, 1115, 1124, 112701-1130, 1132, 1133, 1135, 1137-1139, 1143, 1144, 1147, 1149-1151, 115401-115402, 1156, 1160-1162, 1166
13		15	16								11	*110201-1103, 111101-111102, 111401-111402, 1121, 1136, 1141, 1148, 1163, 116501-116502, 1168
13	14	15	16	17	18	19	20	21	22	23	DR	Well No.

Lot No.: **40E**

Lot-specific information

Length of spec.		205	125	215	210	220	80	80	100	205	165	85	205
PCR product(s)				260			210	210	175	230	215	130	
											250	180	
Well No.	DR	1	2	3	4	5	6	7	8	9	10	11	12
*110404, 1109, 1110, 111202, 1146, 1158	11												
*1105	11												
*1107, 1153	-					5						11	
*110801-110802, 1118- 111902, 1142, 1157	11												
*1113, 1117, 1152	11												
*1116, 1120, 1140, 1159	-						6						
*1122	-												
*1123, 1125	11												
*1126, 1134	11							7					
*1131, 1145, 1164	-												
*1155	-												
*120101-120102, 120302, 1205-1212, 1214	12												
*120201-120202, 1213, 1215	12												
*1204	-										10		
*1216	-										10		
*130101-130201, 130203, 1304, 1308, 1316, 1320, 1322-1324, 1327-1329, 1331, 1332, 1334-1336, 1338-1341, 1348, 1351, 1352, 1354, 1359, 1361, 1363-1365, 1368-1376, 1378-1380, 1383	13						6						
*130202, 130301- 130302, 1310, 1333, 1337, 1366, 1381	13						6						
*130501-130502, 130701-130702, 1311, 131401-131403, 1321, 1342, 1346, 1349- 135002, 1362	13												
Well No.	DR	1	2	3	4	5	6	7	8	9	10	11	12

Lot No.: **40E**

Lot-specific information

100	85	210	195	175	100	110	110	230	215	260		Length of spec.
170	105		210		140	135	175					PCR product(s)
						170	225					
13	14	15	16	17	18	19	20	21	22	23	DR	Well No.
13		15	16			19					11	*110404, 1109, 1110, 111202, 1146, 1158
13			16								11	*1105
13											-	*1107, 1153
13		15									11	*110801-110802, 1118-111902, 1142, 1157
13						19	20				11	*1113, 1117, 1152
13		15	16			19					-	*1116, 1120, 1140, 1159
13						19					-	*1122
13		15	16				20				11	*1123, 1125
13											11	*1126, 1134
13		15					20				-	*1131, 1145, 1164
13			16				20				-	*1155
	14										12	*120101-120102, 120302, 1205-1212, 1214
	14		16								12	*120201-120202, 1213, 1215
	14					19					-	*1204
	14		16								-	*1216
		15	16	17							13	*130101-130201, 130203, 1304, 1308, 1316, 1320, 1322-1324, 1327-1329, 1331, 1332, 1334-1336, 1338-1341, 1348, 1351, 1352, 1354, 1359, 1361, 1363-1365, 1368-1376, 1378-1380, 1383
		15		17							13	*130202, 130301-130302, 1310, 1333, 1337, 1366, 1381
		15	16	17							13	*130501-130502, 130701-130702, 1311, 131401-131403, 1321, 1342, 1346, 1349-135002, 1362
13	14	15	16	17	18	19	20	21	22	23	DR	Well No.

Lot No.: **40E**

Lot-specific information

Length of spec.		205	125	215	210	220	80	80	100	205	165	85	205
PCR product(s)				260			210	210	175	230	215	130	
											250	180	
Well No.	DR	1	2	3	4	5	6	7	8	9	10	11	12
*1306, 1312, 1325, 1330, 1356, 1358, 1360, 1377, 1382	13												
*1309	13												
*1313	-												
*1315, 1319, 1353, 1357	-						6	7					
*1317	13						6			9			
*1326	-							7					
*1343	-						6						
*1344	-							7					
*1345	-						6						
*1367	-												
*140101-140102, 1404, 140701-140702, 1410, 1426, 1428, 1431- 143202, 1435, 1438, 1439, 1454, 1455, 1457, 1460-1462, 1470, 1471, 1475, 1476	14												
*140103, 1408, 142302, 1434, 1472	-												
*1402, 1406, 1409, 1413, 1417, 1420, 1429, 1430, 1433, 1441, 1447, 1448, 1451	14							7					
*140301-140302, 1412, 1440, 1463, 1467, 1477	14							7					
*140501-140503, 1414, 142301, 1436, 1443- 1445, 1456, 1459, 1464, 1465	14												
*1411	-										10		
*1416	6						6						
*1418	-							7					
*1419, 1421	-						6	7					
*1422	-												
*1424	-							7					
*1425, 1453	-												
Well No.	DR	1	2	3	4	5	6	7	8	9	10	11	12

Lot No.: **40E**

Lot-specific information

100	85	210	195	175	100	110	110	230	215	260		Length of spec.
170	105		210		140	135	175					PCR product(s)
						170	225					
13	14	15	16	17	18	19	20	21	22	23	DR	Well No.
		15		17							13	*1306, 1312, 1325, 1330, 1356, 1358, 1360, 1377, 1382
			16	17							13	*1309
		15		17			20				-	*1313
		15	16	17							-	*1315, 1319, 1353, 1357
			16								13	*1317
		15	16	17							-	*1326
		15	16			19	20				-	*1343
				17							-	*1344
		15	16		18		20				-	*1345
		15	16								-	*1367
					18	19	20				14	*140101-140102, 1404, 140701-140702, 1410, 1426, 1428, 1431- 143202, 1435, 1438, 1439, 1454, 1455, 1457, 1460-1462, 1470, 1471, 1475, 1476
						19	20				-	*140103, 1408, 142302, 1434, 1472
				17		19					14	*1402, 1406, 1409, 1413, 1417, 1420, 1429, 1430, 1433, 1441, 1447, 1448, 1451
		15		17			20				14	*140301-140302, 1412, 1440, 1463, 1467, 1477
				17		19	20				14	*140501-140503, 1414, 142301, 1436, 1443- 1445, 1456, 1459, 1464, 1465
						19	20				-	*1411
		15	16		18	19	20				6	*1416
				17		19	20				-	*1418
		15		17		19					-	*1419, 1421
		15	16		18	19	20				-	*1422
			16	17							-	*1424
		15	16		18		20				-	*1425, 1453
13	14	15	16	17	18	19	20	21	22	23	DR	Well No.

Lot No.: **40E**

Lot-specific information

Length of spec.		205	125	215	210	220	80	80	100	205	165	85	205
PCR product(s)				260			210	210	175	230	215	130	
											250	180	
Well No.	DR	1	2	3	4	5	6	7	8	9	10	11	12
*1427	14							7					
*1437	-												
*1442													
*1446, 1452	-												
*1449	-							7					
*1450	-									9			
*1458	-												
*1468	-										10		
*1469	-												
*1473	-												
*1474	-												
*150101-1520, 1522-1526	15			3									
*1521	-			3									
*1527	-			3									
*160101-1603, 160501-160502, 1607-1613N	16				4								
*1604	16				4								
Well No.		1	2	3	4	5	6	7	8	9	10	11	12
DRB1 allele <sup>4</sup>	ser <sup>5</sup>												
DRB3*01010201-0111, 020101-0223, 030101-0303	52												
DRB4*01010101-0107	53												
DRB5*010101-0113, 0202-0205	51												
Well No.	ser <sup>5</sup>	1	2	3	4	5	6	7	8	9	10	11	12

<sup>1</sup>The internal positive control primer pairs amplify segments of the human growth hormone gene. The two different control primer pairs give rise to either an internal positive control band of 430 base pairs, for most wells, or a band of 515 base pairs, for some wells.

Well number 1 contains the primer pair giving rise to the longer, 515 bp, internal positive control band in order to help in the correct orientation of the DR low resolution typing.

In addition, wells number 10 and 19 contain the primer pair giving rise to the longer, 515 bp, internal positive control band in order to allow kit identification.

<sup>2</sup>The codon, in the 2<sup>nd</sup> exon, matching the specificity-determining 3'-end of the primer is given. Codon numbering as on the [www.ebi.ac.uk/imgt/hla](http://www.ebi.ac.uk/imgt/hla) web site. The sequence of the 3 terminal nucleotides of the primer is given. Empty spaces indicate codon boundaries.

<sup>3</sup>The codon, in the 2<sup>nd</sup> exon, matching the specificity-determining 3'-end of the primer is given in the anti-sense direction. Codon numbering as on the [www.ebi.ac.uk/imgt/hla](http://www.ebi.ac.uk/imgt/hla) web site. The sequence of the 3 terminal nucleotides of the primer is given. Empty spaces indicate codon boundaries.



Lot No.: **40E**

Lot-specific information

100	85	210	195	175	100	110	110	230	215	260		Length of spec.
170	105		210		140	135	175					PCR product(s)
						170	225					
13	14	15	16	17	18	19	20	21	22	23	DR	Well No.
		15	16	17			20				14	*1427
			16	17	18						-	*1437
				17			20					*1442
						19					-	*1446, 1452
					18	19	20				-	*1449
					18	19	20				-	*1450
					18		20				-	*1458
					18	19	20				-	*1468
		15			18		20				-	*1469
			16		18	19	20				-	*1473
		15			18	19	20				-	*1474
											15	*150101-1520, 1522-1526
							w				-	*1521
						19					-	*1527
											16	*160101-1603, 160501-160502, 1607-1613N
							w				16	*1604
13	14	15	16	17	18	19	20	21	22	23		Well No.
											ser <sup>5</sup>	DRB1 allele <sup>4</sup>
								21			52	DRB3*01010201-0111, 020101-0223, 030101-0303
									22		53	DRB4*01010101-0107
										23	51	DRB5*010101-0113, 0202-0205
13	14	15	16	17	18	19	20	21	22	23	ser <sup>5</sup>	Well No.

<sup>4</sup>The sequence of the DRB1\*0702 allele has been shown to be identical to DRB1\*070101.  
 The sequence of the DRB1\*080301 allele has been shown to be identical to DRB1\*080302.  
 The sequence of the DRB1\*090101 allele has been shown to be identical to DRB1\*090102.  
 The sequence of the DRB1\*120301 allele has been shown to be identical to DRB1\*1201.  
 The sequence of the DRB1\*1606 allele has been shown to be identical to DRB1\*1605.  
 The DRB4\*0101102N allele has been renamed DRB4\*0103102N.  
 The sequence of the DRB5\*0201 allele has been shown to be identical to DRB5\*0202.  
 The DRB1\*0809, DRB1\*0821 and DRB1\*1415 alleles yield identical amplification patterns except for the specific PCR product yielded by the DRB3 gene in linkage disequilibrium with the DRB1\*1415 allele.  
 The DRB1\*0820, DRB1\*1318, DRB1\*1347 and DRB1\*1355 alleles yield identical amplification patterns except for the specific PCR product yielded by the DRB3 gene in linkage disequilibrium with the DRB1\*1318, DRB1\*1347 and DRB1\*1355 alleles.  
 The DRB1\*0831 and DRB1\*1167 alleles yield identical amplification patterns except for the specific PCR product yielded by the DRB3 gene in linkage disequilibrium with the DRB1\*1167 allele.

101.101-48/12 – including *Taq* polymerase101.101-48u/12u – without *Taq* polymeraseLot No.: **40E**

Lot-specific information

<sup>5</sup>The serological reactivity of all DRB alleles is not known. In this table we use the information in the HLA Dictionary 2004 on the [www.ebi.ac.uk/imgt/hla](http://www.ebi.ac.uk/imgt/hla) web site and the information available at the [www.worldmarrow.org](http://www.worldmarrow.org) web site and have also inferred the serological grouping from the naming of the sequence-defined allele.

'ser', serological HLA specificity.

'w', may be weakly amplified.

101.101-48/12 – including *Taq* polymerase

101.101-48u/12u – without *Taq* polymerase

Lot No.: **40E**

Lot-specific information

CELL LINE VALIDATION SHEET																					
DR low resolution primer set																					
				Prod. No.:	Well																
					1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	
					200848301	200848302	200848303	200848304	200848305	200848306	200848307	200848308	200848309	200848310	200848311	200848312	200848313	200848314	200848315	200848316	
	IHWC cell line		DRB1																		
1	9001 SA		*0101		+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
2	9280 LK707		*1502	*0405	+	-	+	-	-	-	-	+	-	-	-	-	-	-	-	-	-
3	9011 E4181324		*1502		-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
4	9275 GU373		*0301		-	-	-	-	+	+	-	-	-	-	-	-	-	-	-	-	-
5	9009 KAS011		*1601		-	-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-
6	9353 SM		*0407	*0803	-	-	-	-	-	-	-	+	-	+	-	-	-	-	-	-	-
7	9020 QBL		*0301		-	-	-	-	+	+	-	-	-	-	-	-	-	-	-	-	-
8	9007 DEM		*0401	*1602	-	-	-	+	-	-	-	+	-	-	-	-	-	-	-	-	-
9	9026 YAR		*0402		-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-
10	9107 LKT3		*0405		-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-
11	9051 PITOUT		*0701		-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-
12	9052 DBB		*0701		-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-
13	9067 BTB		*0801		-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	+
14	9071 OPGA		*0802		-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	+
15	9075 DKB		*0901		-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-
16	9037 SWEIG007		*1101		-	-	-	-	-	-	-	-	-	-	-	-	+	-	+	+	+
17	9008 WILJON		*1501		-	-	-	-	-	+	-	-	-	-	-	-	+	-	+	+	+
18	9257 32367		*0901	*1101	-	-	-	-	-	-	-	-	-	-	+	-	+	-	+	+	+
19	9038 BM16		*1201		-	-	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-
20	9059 SLE005		*1302		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	+	+
21	9064 AMALA		*1402		-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
22	9056 KOSE		*1302	*1401	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	+	+
23	9124 IHL		*0803	*1414	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-
24	9035 JBUSH		*1101		-	-	-	-	-	-	-	-	-	-	-	-	+	-	+	+	+
25	9049 IBW9		*0701		-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-
26	9285 WT49		*0301		-	-	-	-	+	+	-	-	-	-	-	-	-	-	-	-	-
27	9191 CH1007		*0405	*1001	-	-	-	-	-	-	-	+	-	-	-	+	-	-	-	-	-
28	9320 BEL5GB		*0416	*0701	-	-	-	-	-	-	-	+	+	-	-	-	-	-	-	-	-
29	9050 MOU		*0701		-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-
30	9021 RSH		*0302		-	-	-	-	+	-	+	-	-	-	-	-	-	-	-	-	-
31	9019 DUCAF		*0301		-	-	-	-	+	+	-	-	-	-	-	-	-	-	-	-	-
32	9297 HAG		*1303		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	+	-
33	9098 MT14B		*0404		-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-
34	9104 DHIF		*1101		-	-	-	-	-	-	-	-	-	-	-	-	+	-	+	+	+
35	9302 SSTO		*0403		-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-
36	9024 KT17		*0403	*0406	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-
37	9065 HHKB		*1301		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	+	+
38	9099 LZL		*1402		-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-
39	9315 CML		*0301	*0401	-	-	-	-	+	+	-	+	-	-	-	-	-	-	-	-	-
40	9134 WHONP199		*0701	*0901	-	-	-	-	-	-	-	-	+	-	+	-	-	-	-	-	-
41	9055 H0301		*1302		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	+	+
42	9066 TAB089		*0803		-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-
43	9076 T7526		*0901		-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-
44	9057 TEM		*1401		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
45	9239 SHJO		*0701		-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	-
46	9013 SCHU		*1501		-	-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-
47	9045 TUBO		*1104	*1201	-	-	-	-	-	-	-	-	-	-	-	-	+	+	+	+	+
48	9303 TER-ND		*0103		-	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-

Lot No.: **40E**

Lot-specific information

<b>CELL LINE VALIDATION SHEET</b>											
<b>DR low resolution primer set</b>											
				Prod. No.:	Well						
					17	18	19	20	21	22	23
					200848317	200848318	200848319	200848320	200848321	200848322	200848323
	<b>IHWC cell line</b>	<b>DRB1</b>									
1	9001 SA	*0101			-	-	-	-	-	-	-
2	9280 LK707	*1502	*0405		-	-	-	-	-	+	+
3	9011 E4181324	*1502			-	-	-	-	-	-	+
4	9275 GU373	*0301			+	-	-	-	+	-	-
5	9009 KAS011	*1601			-	-	-	-	-	-	+
6	9353 SM	*0407	*0803		-	-	-	-	-	+	-
7	9020 QBL	*0301			+	-	-	-	+	-	-
8	9007 DEM	*0401	*1602		-	-	-	-	-	+	+
9	9026 YAR	*0402			-	-	-	-	-	+	-
10	9107 LKT3	*0405			-	-	-	-	-	+	-
11	9051 PITOUT	*0701			-	-	-	-	-	+	-
12	9052 DBB	*0701			-	-	-	-	-	+	-
13	9067 BTB	*0801			-	-	-	-	-	-	-
14	9071 OLGA	*0802			-	-	-	-	-	-	-
15	9075 DKB	*0901			-	-	+	-	-	+	-
16	9037 SWEIG007	*1101			-	-	-	-	+	-	-
17	9008 WILJON	*1501			-	-	-	-	-	-	+
18	9257 32367	*0901	*1101		-	-	-	-	+	+	-
19	9038 BM16	*1201			-	-	-	-	+	-	-
20	9059 SLE005	*1302			+	-	-	-	+	-	-
21	9064 AMALA	*1402			+	-	+	-	+	-	-
22	9056 KOSE	*1302	*1401		+	+	+	+	+	-	-
23	9124 IHL	*0803	*1414		+	-	+	+	+	-	-
24	9035 JBUSH	*1101			-	-	-	-	+	-	-
25	9049 IBW9	*0701			-	-	-	-	-	+	-
26	9285 WT49	*0301			+	-	-	-	+	-	-
27	9191 CH1007	*0405	*1001		-	-	-	-	-	+	-
28	9320 BEL5GB	*0416	*0701		-	-	-	-	-	+	-
29	9050 MOU	*0701			-	-	-	-	-	+	-
30	9021 RSH	*0302			+	-	-	-	+	-	-
31	9019 DUCAF	*0301			+	-	-	-	+	-	-
32	9297 HAG	*1303			+	-	-	-	+	-	-
33	9098 MT14B	*0404			-	-	-	-	-	+	-
34	9104 DHIF	*1101			-	-	-	-	+	-	-
35	9302 SSTO	*0403			-	-	-	-	-	+	-
36	9024 KT17	*0403	*0406		-	-	-	-	-	+	-
37	9065 HHKB	*1301			+	-	-	-	+	-	-
38	9099 LZL	*1402			+	-	+	-	+	-	-
39	9315 CML	*0301	*0401		+	-	-	-	+	+	-
40	9134 WHONP199	*0701	*0901		-	-	-	-	-	+	-
41	9055 H0301	*1302			+	-	-	-	+	-	-
42	9066 TAB089	*0803			-	-	-	-	-	-	-
43	9076 T7526	*0901			-	-	+	-	-	+	-
44	9057 TEM	*1401			-	+	+	+	+	-	-
45	9239 SHJO	*0701			-	-	-	-	-	+	-
46	9013 SCHU	*1501			-	-	-	-	-	-	+
47	9045 TUBO	*1104	*1201		-	-	-	-	+	-	-
48	9303 TER-ND	*0103			-	-	-	-	-	-	-

## CERTIFICATE OF ANALYSIS

### **Olerup SSP<sup>®</sup> DR low resolution**

**Product number:** 101.101-48/12 – including *Taq* pol.  
101.101-48u/12u – without *Taq* pol.  
**Lot number:** 40E  
**Expiry date:** 2010-May-01  
**Number of tests:** 48 tests – Product No. 101.101-48  
12 tests – Product No. 101.101-12  
**Number of wells per test:** 23 + 1

### **Well specifications:**

Well No.	Production No.	Well No.	Production No.	Well No.	Production No.
1	2008-483-01	9	2008-483-09	17	2008-483-17
2	2008-483-02	10	2008-483-10	18	2008-483-18
3	2008-483-03	11	2008-483-11	19	2008-483-19
4	2008-483-04	12	2008-483-12	20	2008-483-20
5	2008-483-05	13	2008-483-13	21	2008-483-21
6	2008-483-06	14	2008-483-14	22	2008-483-22
7	2008-483-07	15	2008-483-15	23	2008-483-23
8	2008-483-08	16	2008-483-16		

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

The reactivities of one additional 3'-primer in primer solution 3, one additional 3'-primer in primer solution 4, one additional 5'-primer in primer solution 6, one additional 3'-primer in primer solution 9, one additional primer pair in primer solutions 8, 9 and 11, two additional 5'-primers in primer solution 13, one additional primer pair in primer solution 18, one additional 3'-primer in primer solution 19 and one additional 3'-primer in primer solution 20 were tested by separately adding another 5'-primer or 3'-primer. One of the 5'-primers in primer solution 10 and one of the 5'-primers in primer solution 13 were not possible to test.

The negative control primer pairs, **Production No. 2008-417-01**, can detect contamination with PCR products diluted  $10^{-7}$ .

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

**Date of approval:** 2009-August-05

**Approved by:**

**Quality Control Supervisor**

Lot No.: **40E**

Lot-specific information

## Declaration of Conformity

**Product name:** *Olerup* SSP<sup>®</sup> DR low resolution  
**Product number:** 101.101-48/12, 101.101-48u/12u  
**Lot number:** 40E

**Intended use:** DRB1 low resolution histocompatibility testing

**Manufacturer:** *Olerup* SSP AB  
Hasselstigen 1  
SE-133 33 Saltsjöbaden, Sweden  
**Phone:** +46-8-717 88 27  
**Fax:** +46-8-717 88 18

We, *Olerup* SSP AB, hereby declare that this product, to which this Declaration of Conformity relates is in conformity with the following Standard(s) and other normative document(s) ISO 9001:2000 and ISO 13485:2004, following the provisions of the 98/79/EC Directive on *in vitro* diagnostic medical devices, Annex II List B, as transposed into the national laws of the Member States of the European Union.

The Technical Documentation File is maintained at *Olerup* SSP AB, Hasselstigen 1, SE-133 33 Saltsjöbaden, Sweden.

The Authorized Representative located within the Community is: *Olerup* SSP AB.

Notified Body: Lloyd's Register Quality Assurance Limited, Hiramford, Middlemarch Office Village, Siskin Drive, Coventry CV3 4FJ, United Kingdom. (Notified Body number: 0088.)

Saltsjöbaden, Sweden  
2009-August-05

Olle Olerup  
Managing Director

101.101-48/12 – including *Taq* polymerase

101.101-48u/12u – without *Taq* polymerase

Lot No.: **40E**

Lot-specific information

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For information on *Olerup* SSP distributors worldwide, contact **QIAGEN Vertriebs GmbH**.