



<b>Well No.</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>
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<b>INTERPRETATION TABLE</b>					
<b>DRB1*16 SSP subtyping</b>					
<b>Amplification patterns of the DRB1*1601 to 1613N alleles</b>					
<b>Well</b>					
<b>9</b>	<b>10</b>	<b>11</b>	<b>12</b>	<b>13</b>	
<b>140</b>	<b>115</b>	<b>215</b>	<b>220</b>	<b>155</b>	<b>Length of spec. PCR product</b>
<b>430</b>	<b>430</b>	<b>430</b>	<b>515</b>	<b>430</b>	<b>Length of int. pos. control<sup>1</sup></b>
<b>13</b>	<b>47</b>	<b>14</b>	<b>13</b>	<b>13</b>	<b>5'-primer<sup>2</sup></b>
5' -Agg 3'	5' -g TT 3'	5' -gg A 3'	5' -AAg 3'	5' -Agg 3'	
<b>47</b>	<b>72</b>	<b>72</b>	<b>72</b>	<b>52</b>	<b>3'-primer(s)<sup>3</sup></b>
5' -g gA 3'	5' -gC g 3'	5' -gC g 3'	5' -gC g 3'	5' -CTA 3'	
<b>9</b>	<b>10</b>	<b>11</b>	<b>12</b>	<b>13</b>	<b>Well No.</b>
					<b>DRB1 allele<sup>4</sup></b>
					<b>*160101-160102</b>
					<b>*160201-160202</b>
					<b>*1603</b>
					<b>*1604</b>
					<b>*160501-160502</b>
					<b>*1607</b>
					<b>*1608</b>
<b>9</b>	<b>10</b>				<b>*1609</b>
<b>9</b>	<b>10</b>				<b>*1610</b>
		<b>11</b>			<b>*1611</b>
			<b>12</b>		<b>*1612</b>
				<b>13</b>	<b>*1613N</b>
	<b>10</b>				<b>*110103, 111902, 120101, 120201-120203, 120302-1210, 1212-1215, 1217-1219, 130202, 1377</b>
<b>9</b>					<b>*15010101-150106, 15030101-1506, 1512, 1513, 1516-1518, 1520, 1522-1525, 1532</b>
<b>9</b>					<b>*150201-150206, 1508, 1514, 1515, 1519, 1526, 1527, 1529-1531</b>
<b>9</b>					<b>*1510</b>
					<b>*1511</b>
<b>9</b>					<b>*1521</b>
					<b>DRB1 allele<sup>4</sup></b>

9	10	11	12	13	Well No.
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