

## Batch Certificate For Research Use Only

### PRODUCT INFORMATION AND QUALITY CONTROL

NAME OF PRODUCT	EGFR-Multiplex 5% AF cfDNA
DESCRIPTION	SensID cfDNA (human) is highly characterized human DNA from cell lines.
CATALOG NUMBER	SID-000019
BATCH NUMBER	00037
MANUFACTURING CONDITIONS	<ul style="list-style-type: none"> <li>• Manufactured and sealed in class 2 safety cabinet</li> <li>• At room temperature</li> </ul>
PACKAGE SIZE AND TYPE	<ul style="list-style-type: none"> <li>• 2D barcoded tube with screw cap</li> <li>• Material: Polypropylen (PP)</li> </ul>
DATE OF MANUFACTURE	08.01.2020
EXPIRY DATE	07.01.2022
CONCENTRATION	20 ng/μl (ds DNA)
QUANTITY	400 ng (ds DNA)
NOMINAL VOLUME	20 μl
MUTATION	<p>EGFR</p> <p>p.G719S (COSM6252*, COSV51767289*, substitution, c.2155G&gt;A, Exon 18)</p> <p>p.E746_A750delELREA (COSM6225*, COSV51765066*, deletion, c.2236_2250del15, Exon 19)</p> <p>p.S752_I759delSPKANKEI (COSM6256*, COSV51774879*, deletion, c.2254_2277del24, Exon 19)</p> <p>p.S768I (COSM6241*, COSV51768106* substitution, c.2303G&gt;T, Exon 20)</p> <p>p.V769_D770insASV (COSM20884*, COSV51850427* Insertion, c.2303_2304insTGTGGCCAG, Exon 20)</p> <p>p.T790M (COSM6240*, COSV51765492*, substitution, c.2369C&gt;T, Exon 20)</p> <p>p.L858R (COSM6224*, COSV51765161*, substitution, c.2573T&gt;G, Exon 21)</p> <p>p.L861Q (COSM6213*, COSV51766344*, substitution, c.2582T&gt;A, Exon 21)</p> <p><small>* GRCh38 COSMIC v90</small></p>
ALLELE FREQUENCY	5%
QUALITY	<p>DNA quantity metrological traceable to internationally certified reference material<sup>1</sup></p> <p>The copy number values are metrologically traceable to the natural units count 1 and ratio 1 and International System of Units (SI) derived units of volume.</p>
STORAGE CONDITIONS	+ 2-8 °C

<sup>1</sup> ERM\_AD442K  
Phone: +49 (0) 381 377 182 01

MANUFACTURING AND QUALITY CONTROL SITES	SensID GmbH Schillingallee 68, 18057 Rostock, Germany																																	
TEST METHOD AND ACCEPTANCE CRITERIA	Quality Control	Test Method	Acceptance criteria																															
	Fragmentation	Fragment Length Analysis Agilent High Sensitivity DNA Kit (Agilent Technologies)	peak size 167 bp ± 10% (151 bp – 181 bp)																															
	Quantification	Total DNA measurement: Spectrophotometry ssDNA [ng/μl] = (A260-A320)*38 <sup>2,3</sup> dsDNA measurement: Qubit dsDNA BR Assay Kit (Invitrogen)	Total DNA: n.a. <sup>4</sup> dsDNA: 17.5 – 22.5 ng/μl																															
	Allele Frequency	dPCR Analysis using BioRad QX200™ System	AF 5% ±30% (3.5-6.5%)																															
RESULTS OF ANALYSIS	<table border="1"> <thead> <tr> <th></th> <th>Result</th> <th>PASS/FAIL</th> </tr> </thead> <tbody> <tr> <td>Fragmentation</td> <td>179 bp</td> <td>PASS</td> </tr> <tr> <td>Quantity</td> <td>29.6 ng/μl (total DNA) 20.3 ng/μl (dsDNA)</td> <td>PASS</td> </tr> <tr> <td rowspan="8">Allele Frequency</td> <td><b>Mutation</b></td> <td><b>AF in %</b></td> </tr> <tr> <td>L858R</td> <td>4.4</td> </tr> <tr> <td>L861Q</td> <td>5.3</td> </tr> <tr> <td>S768I</td> <td>5.3</td> </tr> <tr> <td>E746_A750delELREA</td> <td>3.9</td> </tr> <tr> <td>T790M</td> <td>4.0</td> </tr> <tr> <td>G719S</td> <td>4.6</td> </tr> <tr> <td>V769_D770insASV</td> <td>4.6</td> </tr> <tr> <td>S752_I759delSPANKEI</td> <td>5.3</td> </tr> <tr> <td></td> <td></td> <td>PASS</td> </tr> </tbody> </table>				Result	PASS/FAIL	Fragmentation	179 bp	PASS	Quantity	29.6 ng/μl (total DNA) 20.3 ng/μl (dsDNA)	PASS	Allele Frequency	<b>Mutation</b>	<b>AF in %</b>	L858R	4.4	L861Q	5.3	S768I	5.3	E746_A750delELREA	3.9	T790M	4.0	G719S	4.6	V769_D770insASV	4.6	S752_I759delSPANKEI	5.3			PASS
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COMMENTS/REMARKS	Additional information:  <b>Copy numbers (CN) of the respective measurements</b>  <i>Table 1 indicates the values of the QC assays performed by SensID GmbH with an DNA input of ~20 ng. The value for the respective mutation results from the mean value of five measured QC samples (CN values are rounded). CN concentration values per microliter (μl), are based on droplet digital (ddPCR) assay counts dilution factors, and droplet volume measurements. The detection of the amount of CNs may vary depending on the assay used. Therefore, due to assay properties, there may be deviations in the observed number of copies and allele frequencies compared to the values given here.</i> <table border="1" data-bbox="587 1787 1406 1854"> <thead> <tr> <th>Mutation</th> <th>CN wt<sup>5</sup>/μl</th> <th>CN mut<sup>6</sup>/μl</th> </tr> </thead> <tbody> <tr> <td>L858R</td> <td>4631</td> <td>213</td> </tr> </tbody> </table>			Mutation	CN wt <sup>5</sup> /μl	CN mut <sup>6</sup> /μl	L858R	4631	213																									
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<sup>2</sup> Measured before spiking in

<sup>3</sup> Protocol NK603 – Community Reference Laboratory for GM Food and Feed

<sup>4</sup> not applicable

<sup>5</sup> Wild Type

<sup>6</sup> Mutation

L861Q	5968	332
S768I	4555	253
E746_A750delELREA	4444	180
T790M	4671	195
G719S	4634	225
V769_D770insASV	3683	181
S752_I759delSPANKEI	2509	137

Name and position/title of Person authorising the batch release:

Mr. Björn Nowack

Date of batch release: 13.01.2020

Signature batch release: Björn Nowack

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