

Batch Certificate For Research Use Only

PRODUCT INFORMATION AND QUALITY CONTROL

NAME OF PRODUCT	PIK3CA-H1047R 50%AF FFPE Reference Standard
DESCRIPTION	Human FFPE Reference Standard (curl)
CATALOG NUMBER	SID-000102
BATCH NUMBER	00062
MANUFACTURING CONDITIONS	<ul style="list-style-type: none"> • Manufactured and sealed according to internal quality standards related to EN ISO 13485 • At room temperature
PACKAGE SIZE AND TYPE	<ul style="list-style-type: none"> • 2D barcoded tube with screw cap • Material: Polypropylen (PP)
DATE OF MANUFACTURE	14.04.2020
EXPIRY DATE	13.04.2021
FORMAT	10 µm section / 1 curl
MUTATION	PIK3CA p.H1047R (COSM775*, COSV55873195*, substitution, c.3140A>G, Exon 20) <small>* GRCh38 COSMIC v90</small>
ALLELIC FREQUENCY	50.0 %
QUALITY	DNA quantity metrologically traceable to internationally certified reference material ¹
STORAGE CONDITIONS	+ 2-8 °C
MANUFACTURING AND QUALITY CONTROL SITES	SensID GmbH Schillingallee 68, 18057 Rostock, Germany

¹ ERM_AD442K
Phone: +49 (0) 381 377 182 01

TEST METHOD AND ACCEPTANCE CRITERIA	Quality Control	Test Method	Acceptance criteria
	Cell Density	Visual	> 60 %
	Quality	Agarose gel electrophoresis 1% Gel with fluorescent DNA stain in 1 % TAE buffer	Bright band of high-molecular-weight gDNA ≥ 20 kb
	Quantification	dsDNA measurement ² : Qubit dsDNA BR Assay Kit (Invitrogen) RNA measurement ² : Qubit RNA BR Assay Kit (Invitrogen)	dsDNA: > 400 ng/μl RNA: > 400 ng/μl
	Allelic Frequency	ddPCR Analysis ³ Fehler! Textmarke nicht definiert. using BioRad QX200™ System	AF 50.0 % (45.0-55.0 %)

RESULTS OF ANALYSIS	Result		PASS/FAIL				
	Cell Density	Visual: > 60 %		PASS			
Quality	Bright band of high-molecular-weight gDNA ≥ 20 kb		PASS				
Quantity	707.3 ng/μl (dsDNA) 427.8 ng/μl (RNA)		PASS				
Allelic Frequency	<table border="1"> <thead> <tr> <th>Mutation</th> <th>AF in %</th> </tr> </thead> <tbody> <tr> <td>PIK3CA H1047R</td> <td>51.1%</td> </tr> </tbody> </table>	Mutation	AF in %	PIK3CA H1047R	51.1%		PASS
Mutation	AF in %						
PIK3CA H1047R	51.1%						

COMMENTS/REMARKS						
<p>Additional information:</p> <p>Theoretical DNA yield from 1 curl under the assumption of a diploid chromosome set: 3122 ng/μl (dsDNA)</p> <p>Copy numbers (CN) of the respective measurements</p> <p><i>Table 1 indicates the values of the QC assays performed by SensID GmbH with a DNA input of ~10 ng. The value for the respective mutation results from the mean value of five measured replicates (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></p> <table border="1"> <thead> <tr> <th>Mutation</th> <th>CN wt³/ ng extracted DNA</th> <th>CN mut⁴/ ng extracted DNA</th> </tr> </thead> <tbody> <tr> <td>PIK3CA H1047R</td> <td>214</td> <td>224</td> </tr> </tbody> </table>	Mutation	CN wt ³ / ng extracted DNA	CN mut ⁴ / ng extracted DNA	PIK3CA H1047R	214	224
Mutation	CN wt ³ / ng extracted DNA	CN mut ⁴ / ng extracted DNA				
PIK3CA H1047R	214	224				

² Measured after extraction with Qiagen AllPrep DNA/RNA FFPE Kit

³ Wild Type

⁴ Mutation



Bringing Precision to MD_x

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

Mr. Björn Nowack

Date of batch release: 09.06.2020

Signature batch release: Björn Nowack

This document was created electronically and is valid without a signature.

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