

Batch Certificate For Research Use Only

PRODUCT INFORMATION AND QUALITY CONTROL

NAME OF PRODUCT EGFR-T790M 1% AF cfDNA in Plasma

DESCRIPTION Human proteins in common plasma concentrations, electrolytes,

EDTA, cfDNA / ctDNA in common plasma concentrations

CATALOG NUMBER SID-000012

BATCH NUMBER 00077

MANUFACTURING

• Manufactured and sealed in class 2 safety cabinet

CONDITIONS • Bottled with qualified Dispenser

At room temperature

PACKAGE SIZE AND

• 2D barcoded tube with screw cap

• Material: Polypropylen (PP)

DATE OF MANUFACTURE | 22.07.2020

EXPIRY DATE 21.07.2021

CONCENTRATION 80 ng/ml (dsDNA)

QUANTITY 400 ng (dsDNA)

NOMINAL VOLUME 23.0 µl in 5 ml plasma (518 ng)

MUTATION p.T790M (COSM6240*, COSV51765492*, substitution, c.2369C>T, Exon 20)

* GRCh38 COSMIC v91

ALLELE FREQUENCY 1.0%

QUALITY DNA quantity metrologically traceable to internationally certified

reference material¹

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.

STORAGE CONDITIONS + 2-8 °C

MANUFACTURING AND | SensID GmbH

QUALITY CONTROL Schillingallee 68, 18057 Rostock, Germany

SITES

¹ ERM_AD442K



TEST METHOD AND	Quality Control	Test Method	Acceptance	
ACCEPTANCE CRITERIA			criteria	
		Fragment Length Analysis ²	peak size 167 bp ±	
	Fragmentation	Agilent High Sensitivity DNA Kit	10%	
		(Agilent Technologies)	(151 bp - 181 bp)	
	Quantification	Total DNA measurement:	ssDNA:	
		Spectrophotometry	n.a. ⁴	
		ssDNA [ng/µl] = (A260-A320)*38 ^{2,3}		
		dsDNA measurement ² : Qubit	dsDNA:	
		dsDNA BR Assay Kit (Invitrogen)	n.a.⁴	
	Allele Frequency	ddPCR Analysis²	AF 1.0%	±40%
		using BioRad QX200™ System	(0.6-1.4%)	
RESULTS OF ANALYSIS		D t-		DACO /EAU
		Result		PASS/FAIL
	Fragmentation	171 bp		PASS
	Quantity	35.7 ng/µl (total DNA)		
		22.5 ng/μl (dsDNA)		n.a. ⁴
	Allele	Mutation AF i	n % PASS	
	Frequency	T790M 0.		
COMMENTS/REMARKS	Additional information:			
	Copy numbers (CN) of the respective measurements			
	Table 1 indicates the values of the QC assays performed by SensID GmbH with an DNA input			
	of ~10 ng. The value for the respective mutation results from the magnitude of three			

of ~40 ng. The value for the respective mutation results from the mean value of three measured replicates (CN values are rounded). CN concentration values per milliliter (ml), 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 conics and allele frequencies compared to the values given here.

 $copies\ and\ allele\ frequencies\ compared\ to\ the\ values\ given\ here.$

Mutation	CN wt⁵/ml	CN mut ⁶ /ml
EGFR T790M	18147	110

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

Mr. Björn Nowack

Date of batch release: 28.07.2020

Signature batch release: Björn Nowack

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

6 Mutation

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²Measured before spiking in

³ Protocol NK603 – Community Reference Laboratory for GM Food and Feed

⁴ not applicable

⁵ Wild Type