

Revised Technical Product Report

For Research Use Only; Not for use in Diagnostic Procedures

Product Description:

Seraseg® ctDNA Mutation Mix v2 WT

Material Number:

0710-0144

Batch Number:

10678693

Material Description:

ctDNA-like human genomic DNA from the reference cell line, GM24385.

AF%1

0.00

0.00

Volume:

25 µL

Date of Manufacture:

10 JUL 2023

Expiration Date:

10 JUL 2027

Storage:

-20°C

Concentration

Nominal Concentration: 10 ng/µL

(Qubit dsDNA BR Assay):

Average Measured Concentration: 14.9 ng/µL

COSMIC Identifier

COSM33765

Average fragment size (Agilent Bioanalyzer DNA

150 bp

ΔKT1

1000 Analysis):

Acceptance criteria for

average fragment size:

140-200 bp

Gene ID

AKIT	COSM33765	0.00
APC	COSM13127	0.00
APC	COSM18561	0.00
ATM	COSM21924	0.00
BRAF	COSM476	0.00
CTNNB1	COSM5664	0.00
EGFR	COSM6224	0.00
EGFR	COSM12378	0.00
EGFR	COSM6225	0.00
EGFR	COSM6240	0.00
ERBB2	COSM20959	0.00
FGFR3	COSM715	0.00
FLT3	COSM783	0.00
FOXL2	COSM33661	0.00
GNA11	COSM52969	0.00
GNAQ	COSM28758	0.00
GNAS	COSM27887	0.00
IDH1	COSM28747	0.00
JAK2	COSM12600	0.00
KIT	COSM1314	0.00
KRAS	COSM521	0.00
MPL	COSM18918	0.00
NCOA4-RET	N/A	0.00
NPM1	COSM17559	0.00

Digital PCR testing using BioRad QX200™ Droplet Digital™ PCR System:

COSM584

NRAS



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PDGFRA	COSM736	0.00
PDGFRA	COSM28053	0.00
PIK3CA	COSM763	0.00
PIK3CA	COSM12464 ³	0.00
PIK3CA	COSM775	0.00
PTEN	COSM4986	0.00
PTEN	COSM5809	0.00
RET	COSM965	0.00
SMAD4	COSM14105	0.00
TP53	COSM10648	0.00
TP53	COSM10660	0.00
TP53	COSM10662	0.00
TP53	COSM6530	0.00
TP53	COSM18610	Not Tested ²
TPR-ALK	N/A	0.00

Variant allele frequencies > 0.00% for this wildtype negative control are within the expected range for stochastic positive dPCR reactions.

NGS was performed as an orthogonal verification step. Results confirm no variants were detected above 0.1%, except for the TP53 COSM10648 variant detected at 0.1% in the WT sample; the corresponding dPCR data was 0.05%. Thus, we associate the allele frequency for the TP53 COSM10648 variant with the background from the assay using the conditions listed below.

Next Generation
Sequencing testing using
Archer® Reveal ctDNA™
28 Kit run on an Illumina®
MiSeq™ using v2 (2x150
bp) PE chemistry
reagents:

NGS Parameters:

DNA input = 50 ng # of samples / flow cell = 2 # of total reads / sample = 21.1 M Average read depth = 6151X On-target reads = ~96% Q30 score = ~89% Analysis = Archer Analysis Suite v6.2.7

Default analysis settings except for the following:

- NORMALIZATION_LEVEL is set to 10000000 (default is 3500000; cfDNA pipeline)
- VARIANT_DOWNSTREAM_ROI_SIZE is set to 150 (default=400; cfDNA pipeline)

Please see the poster from NIST for more information about assay sensitivity:

LGC Clinical Diagnostics, Inc • 910 Clopper Road, Gaithersburg, MD 20878 • 508.244.6400 • 800.676.1881 Toll Free

² NGS testing alone is used for QC of variant COSM18610.

³ As of June 2019 this mutation is no longer listed in the COSMIC database.



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Product Description:

Seraseq® ctDNA Mutation Mix v2 WT

https://digital.seracare.com/multilab-assessment-reference-materials-

ctdna-poster2018

Approval:

Prepared By

19-NOV-2024

Date