

Multiplex cfDNA Reference Standard

IMPORTANT: Centrifuge tube prior to opening to prevent loss of product

PRODUCT INFORMATION

Catalog ID	HD786
Batch ID	37408
Product Description	Structural Multiplex cfDNA Reference Standard
Date of Manufacture	17-May-2019
Manufacturing QMS Status	ISO 9001

PROPERTIES

Quantity	350 ng
Concentration	20 ng/ μ L
Average Fragment Length	160 bp
Buffer	Tris-EDTA (10 mM Tris-HCl, 1 mM EDTA), pH 8.0
Storage Conditions	+4°C
Use by	For the latest information on product shelf life, please visit our website: https://www.horizondiscovery.com/reference-standards/coa-information

CUSTOMER SUPPORT

Technical Support	technical@horizondiscovery.com
Customer Service	orders@horizondiscovery.com

ADDITIONAL INFORMATION

N/A

Horizon Discovery

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QUALITY CONTROL

 Allele Frequency (AF) / Copy Number and Genotype: **Droplet Digital™ PCR**

Gene	Variant	Expected AF (%) / Copy Number	Actual AF (%) / Copy Number	Acceptance Criteria (%) / Copy Number		Pass/Fail
				Low	High	
GNA11	Q209L	5.6	5.54	3.90	7.30	PASS
AKT1	E17K	5.0	4.51	3.50	6.50	PASS
PIK3CA	E545K	5.6	4.81	3.90	7.30	PASS
EGFR	V769_D770insASV	5.6	4.48	3.90	7.30	PASS
EGFR	ΔE746-A750	5.3	5.51	3.70	6.90	PASS
ROS1	SLC34A2/ROS1 Fusion	5.6	5.86	3.90	7.30	PASS
RET	CCDC6/RET Fusion	5.0	4.16	3.50	6.50	PASS
MET	Amplification	4.5 copies	4.90 copies	3.20 copies	5.90 copies	PASS
MYC-N	Amplification	9.5 copies	10.60 copies	7.60 copies	11.40 copies	PASS

 Fragmentation Size: **D1000 DNA ScreenTape assay (Agilent Technologies) TapeStation software version A.02.02**

Acceptance Criteria	Reading	Pass/Fail
160 bp ± 10% (144 bp – 176 bp)	166 bp	PASS

 Quantification: **Qubit® dsDNA BR assay (Invitrogen)**

Acceptance Criteria	Reading	Pass/Fail
20 ng/μL ± 15% (17.0 – 23.0 ng/μL)	18.7 ng/μL	PASS

FOR RESEARCH USE ONLY

Manufactured in the UK

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