

Revised Technical Product Report

For Research Use Only; Not for use in Diagnostic Procedures

Product Description:	Seraseq® FFPE TMB RM Score 13		
Material Number:	0710-1618	Batch Number:	10594031
Material Description:	Reference standard made from a mutagenized reference cell line (GM24385) and its matched "normal" cell line. Cells were mixed at 30% mutagenized, fixed with 10% formalin, and embedded in paraffin.		
Fill Size:	2x10 µm FFPE curls		
Date of Manufacture:	24 NOV 2021	Expiration Date:	14 MAY 2024
Extraction and Concentration Test Method:	QIAamp DNA FFPE Tissue Kit for extraction and Qubit dsDNA HS Assay for concentration.		
Measured Yield per Curl:	Seraseq FFPE TMB RM Score 13: 190 ± 35 ng		
Whole Exome Sequencing (WES) Method & Analysis:	<ul style="list-style-type: none"> • Library preparation <ul style="list-style-type: none"> ○ Agilent SureSelectXT Human AllExon V6 + COSMIC (enrichment kit) ○ Capture Region = 32.4 Mb ○ DNA input = 1000 ng / run • Whole Exome Sequencing <ul style="list-style-type: none"> ○ ILMN NovaSeq 6000 ○ Average read depth ~300X • TMB Analysis Pipeline metrics and parameters (Based on FoCR TMB harmonization algorithm¹) <ul style="list-style-type: none"> ○ TMB variants = missense, nonsense, frameshift, in-frame insertion, in-frame deletion ○ FILTER=PASS; AF≥0.05; tDP≥25; AO≥3 ○ Aligner = bwa mem ○ Mutation Detection = Mutect2 (tumor-normal mode) ○ Mutation Annotation = SnpEff ○ Mutation Filtration = SnpSift 		
TMB Score by WES:	12.12 ± 0.25 (average and standard deviation of three replicates)		

¹<https://meetinglibrary.asco.org/record/172797/abstract>

Approval:



22 JUN 2023

Prepared By

Date