

Guide to the MOCHA Solid Tumor Biomarker Report



Molecular Characterization Laboratory

Frederick National Laboratory for Cancer Research Leidos Biomedical Research, Inc. 459 Miller Drive, Frederick, MD 21701 CLIA Laboratory ID 21 D2097127 301.846.7689

OCAv3 Next-Generation Sequencing Assay Results

NCI Protocol #10323 Report Date: 8/28/2019

INVESTIGATIONAL USE ONLY

The OCAv3 next-generation sequencing (NGS) assay identifies more than 3000 annotated mutations of interest (MOIs) broadly categorized into 5 mutation types: single nucleotide variants (SNV), small insertions/deletions (Indels), large (>3 bases) insertions/deletions (Large Indels), copy number variants (CNV), and gene fusions. This report summarizes annotated mutations identified in the tumor specimen identified below.

Patient Name: James Johnson Referring Physician: Rober			Patient ID: MD123-0001 pert Smith		Specimen ID: 10323-SK78WG02-1		MoCha Sample ID: OCA-15002	
					Telephone:	202-555-1234	Fax:	202-555-4321
Biopsy Site:	Skin		Date Collected:	Aug 07 2019	Primary Diagnosis:	Melanoma	Tumor Content (%) ^{1,2} :	75

Single Nucleotide Variants (SNVs) & Small/Large Indels ^{3,4,5} :										
Gene	ID Code	VAF ³	Variant Class ⁴	Function	HGV\$⁵	Transcript ID	Protein Change			
NRAS	COSM566	37.63%	Hotspot	missense	c.35G>T	NM_002524.4	p.(G12V)			
BRAF	COSM476	68.14%	Hotspot	missense	c.1799T>A	NM_004333.4	p.(V600E)			
TP53		6.07%	Hotspot	missense	c.557A>G	NM_000546.5	p.(D186G)			

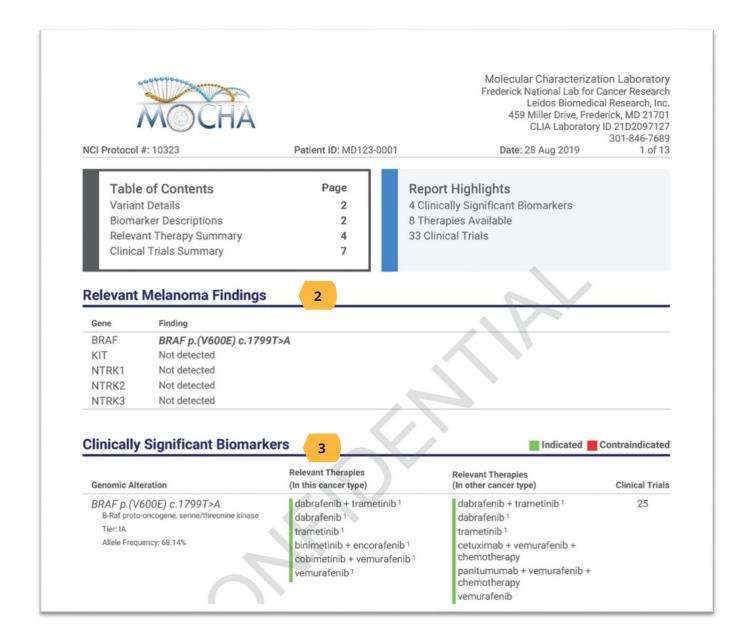
1. Mutations of Interest (MOIs) Found in the Tumor

This table lists <u>DNA</u> and/or <u>RNA</u> biomarkers found in the tumor that may have important information about a person's cancer and possible treatments.

The biomarker test looks for three types of biomarkers:

- Single Nucleotide Variants which are variations of a gene
- Copy Number Variants which is the number of times a gene is repeated
- Gene fusions when two separate genes combine together. Get more information on the genetics of cancer.





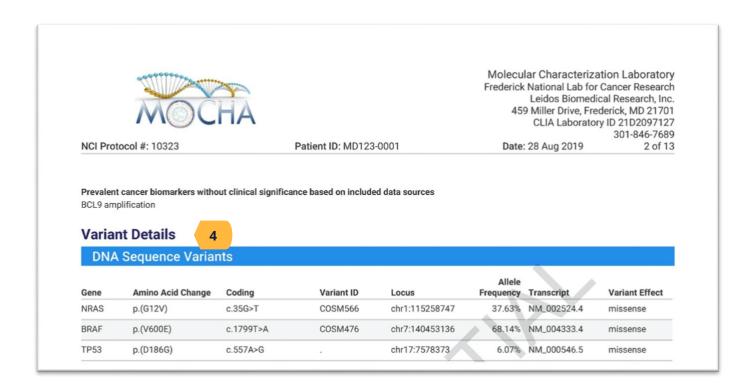
2. Relevant Findings

This table lists key cancer genes tested for. If a gene tested for was found, the description and location of any gene mutations are included.

3. Clinically Significant Biomarkers

This table lists therapies (treatments) based on the cancer type and specific to biomarkers found in the tumor. The list includes therapies for the patient's specific cancer type as well as for other types of cancers that have the same biomarkers. Different cancer types that share biomarkers may respond to the same biomarker-targeted therapies.

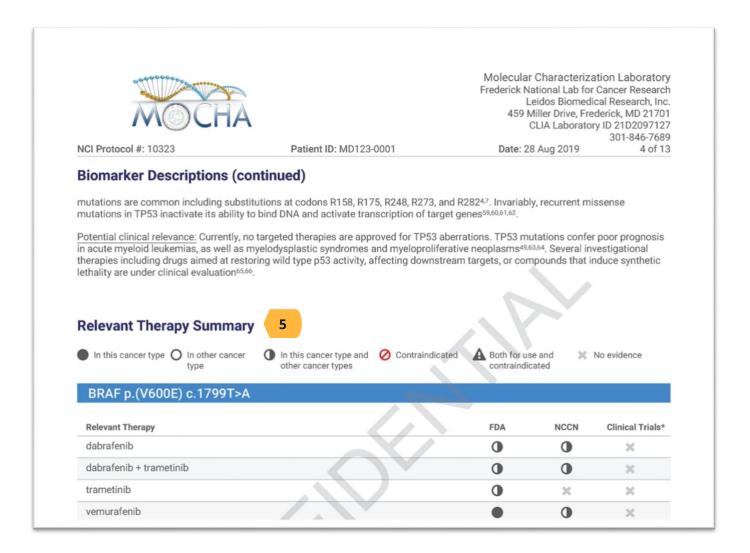




4. Variant Details

The biomarker report lists genes that are known to be related to cancer in some way. It also lists changes (variants) in those genes found in the tumor. Doctors may find this information useful when recommending treatment. There are still some changes in cancer genes that are not understood yet. They may be included in the report if they were detected, since they have been found in many cancers. This is another reason why research on these biomarkers is important.

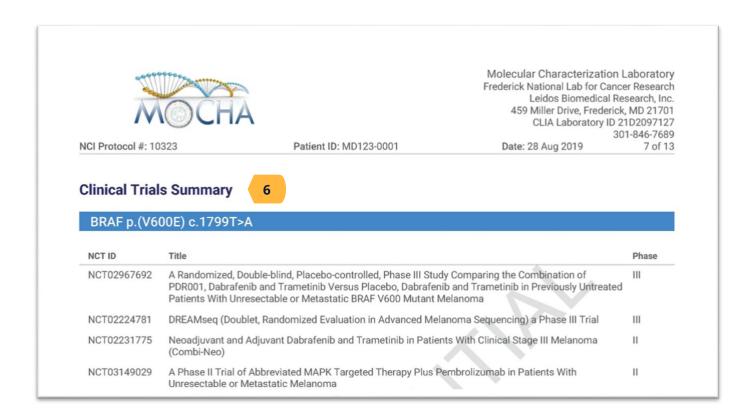




5. Relevant Therapy Summary

The Food and Drug and Administration (FDA) and National Comprehensive Cancer Network (NCCN) publish therapy guidelines for specific cancer types. This table lists their recommended therapies based on the <u>biomarkers</u> found in the tumor. The table also shows whether there may be clinical trials available. A clinical trial is a type of research study that tests potential new therapies.





6. Clinical Trials Summary

This table lists clinical trials that may be available for a specific type of cancer, based on gene <u>mutations</u> and or <u>biomarkers</u> found in the tumor. A clinical trial is a type of research study that tests potential new therapies.

More information on each clinical trial, such as where the clinical trial is being done, can be found at <u>ClinicalTrials.gov</u>.