9048

Date of Birth 1996-08-11

Leukemia Sample Jennifer Houston

Sex

**Female** 

Physician

Dr. Christopher Reynolds

Institution

White, Jones and Carlson

Tumor specimen: source Leukemia CollectedDate 2024-03-08 ReceivedDate 2024-03-14 TumorPercentage 50%

Normal specimen: source Blood CollectedDate 2024-03-12 ReceivedDate 2024-03-12

#### **GENOMIC VARIANTS**

**Somatic - Potentially Actionable** variant allele fraction ARID2 c.798G>A p.W266\* Stopgain-LOF 5.87% Somatic - Biologically Relevant c.1154\_1155insTTGTC p.L367fs\*46 **CALR** 10.76% Spliceregionvariant-LOF CSF3R c.3113A>G p.W266\* Missensevariant(exon2)-GOF 25.98% CDC73 c.1A>G p.W266\* Stopgain-LOF 17.65%

## Germline - Pathogenic

No Germline - Pathogenic variants were found in the limited set of genes on which we report.

## **Pertinent Negatives**

**PDGFRB** PTCH1 SF3B1 CALR

## **IMMUNOTHERAPY MARKERS**

**Tumor Mutational Burden** Microsatellite Instability Status

41 m/Mb 23% Stable Equivocal High

# FDA-APPROVED THERAPIES, Current Diagnosis

KRAS G12C Sotorasib NCCN, Consensus, Non-Small Cell Lung Cancer **Inhibitors** 

MSK OncoKB, Level 1 KRASp.G12C G12C-GOF

## FDA-APPROVED THERAPIES, Other Indications

KRAS G12C Sotorasib NCCN, Consensus, Non-Small Cell Lung Cancer

> MSK OncoKB, Level 1 KRASp.G12C G12C-GOF

**Inhibitors** 

## **ADDITIONAL INDICATORS**

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Intove	vrahla	Progn	veie.
Jillavi	JI abie	FIUGII	USIS.

NCCN, Consensus, Non-Small Cell Lung Cancer KRASp.G12C Gain-of-function

## **CLINICAL TRIALS**

A Study of VS-6766 v. VS-6766 + Defactinib in Recurrent G12V, Other KRAS and BRAF Non-Small Cell Lung Cancer

Phase 2 City, state - x mi KRAS mutation

A Phase 1/2 Study of MRTX849 in Patients With Cancer Having a KRAS G12C Mutation KRYSTAL-1

Phase 1/2 City, state - x mi KRAS mutation STK11 mutation

First-in-human Study of DRP-104 (Sirpiglenastat) as Single Agent and in Combination With Atezolizumab in Patients With Advanced Solid Tumors. (NCT04471415)

Phase 1/2 City, state - x mi NFE2L2 mutation STK11 mutation

## **VARIANTS OF UNKNOWN SIGNIFICANCE**

Somatic SRSF2	Mutation effect c.3113A>G p.P95X Spliceregionvariant-LOF NM_001011645	Variant allele fraction 21.92%
ARID2	c.798G>A p.W266* Spliceregionvariant-LOF NM_001011645	6.58% -
CDC73	c.1A>G p.M1V Missensevariant(exon2)-GOF NM_001011645	2.18%
PDGFRB	c.3113A>G p.T681I Frameshift-LOF NM_001011645	3.56%
IDH1	c.356G>A p.R132X Nonsense-LOF NM_001011645	7.26%
JAK3	c.3113A>G p.L857P Nonsense-LOF NM_001011645	2.48%
Germline	Mutation effect	Condition
GNAS	c.679C>G p.Q227R Spliceregionvariant-LOF NM_001011645	report

#### LOW COVERAGE REGIONS

ERCC2

IDH1

### **SOMATIC VARIANT DETAILS - POTENTIALLY ACTIONABLE**

ARID2

c.798G>A p.W266\* Stopgain-LOF

VAF: 5.87% -

ARID2 encodes a protein that is a subunit of the SWI/SNF chromatin remodeling complex SWI/SNF-B or PBAF. This complex functions in ligand-dependent transcriptional activation. Loss of function mutations and copy number loss of ARID2 are associated with cancer progression.

#### **SOMATIC VARIANT DETAILS - BIOLOGICALLY RELEVANT**

CALR

c.1154\_1155insTTGTC p.L367fs\*46 Spliceregionvariant-LOF

VAF: 10.76%

PTEN encodes a phosphatase that acts as a tumor suppressor by negatively regulating the PI3K-AKT-mTOR pathway. Loss of function mutations, copy number loss, and underexpression of PTEN are associated with cancer progression.

CSF3R

c.3113A>G p.W266\* Missensevariant(exon2)-GOF

VAF: 25.98%

PTEN encodes a phosphatase that acts as a tumor suppressor by negatively regulating the PI3K-AKT-mTOR pathway. Loss of function mutations, copy number loss, and underexpression of PTEN are associated with cancer progression.

CDC73

c.1A>G p.W266\* Stopgain-LOF

VAF: 17.65%

STK11 (LKB1) encodes an enzyme in the serine/threonine kinase family that is responsible for maintaining energy metabolism and cellular polarization through the activation of AMP-activated protein kinase and other members of the AMPK family. The enzyme also acts as a tumor suppressor by regulating cell growth. Loss of function mutations, copy number loss, epigenetic variation, and underexpression of STK11 are associated with cancer progression.

## **CLINICAL HISTORY**

Diagnosed on

2024-03-07