

Date of Birth
1978-12-04

Sex
Male

Physician
Dr. Philip Marquez

Institution
Bishop-Thompson

Tumor specimen:
source Pancreatic
CollectedDate 2023-05-09
ReceivedDate 2023-05-09
TumorPercentage 5%

Normal specimen:
source Blood
CollectedDate 2023-05-11
ReceivedDate 2023-05-11

GENOMIC VARIANTS

Somatic - Potentially Actionable		variant allele fraction	
ARID2	c.798G>A p.W266* Stopgain-LOF	35.9%	<div><div></div></div>
IDH1	c.394_395delinsTC p.R132H Nonsense-GOF	32.59%	<div><div></div></div>
TOP2A	c.3113A>G p.A1515S Stopgain-LOF	25.8%	<div><div></div></div>
Somatic - Biologically Relevant			
FGFR3	c.1111A>T p.A1515S Spliceregionvariant-LOF	15.97%	<div><div></div></div>
IDH1	c.394_395delinsTC p.R132C Nonsense-LOF	2.0%	<div><div></div></div>

Germline - Pathogenic

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

Pertinent Negatives

PKLR BRCA1

IMMUNOTHERAPY MARKERS

Tumor Mutational Burden	Microsatellite Instability Status
48 m/Mb 17%	<div><div>Stable</div><div>Equivocal</div><div>High</div></div>

FDA-APPROVED THERAPIES, Current Diagnosis

KRAS G12C Inhibitors	Sotorasib	NCCN, Consensus, Non-Small Cell Lung Cancer MSK OncoKB, Level 1 KRASp.G12C G12C-GOF
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FDA-APPROVED THERAPIES, Other Indications

KRAS G12C Inhibitors	Sotorasib	NCCN, Consensus, Non-Small Cell Lung Cancer MSK OncoKB, Level 1 KRASp.G12C G12C-GOF
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ADDITIONAL INDICATORS

Unfavorable Prognosis	NCCN, Consensus, Non-Small Cell Lung Cancer KRASp.G12C Gain-of-function
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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	Mutation effect	Variant allele fraction
MYD88	c.794T>C p.L265P Spliceregionvariant-LOF NM_001011645	11.73% <div></div>
ERCC2	c.3113A>G p.F332V Missensevariant(exon2)-GOF NM_001011645	11.11% <div></div>
PAX5	c.547G>A p.P80R Spliceregionvariant-GOF NM_001011645	5.47% <div></div>
PAX5	c.964G>A p.G183S Frameshift-GOF NM_001011645	9.02% <div></div>
H3.3	c.3113A>G p.G35W Spliceregionvariant-LOF NM_001011645	8.69% <div></div>
HSP90B1	c.3113A>G p.I66T Missensevariant(exon2)-GOF NM_001011645	5.96% <div></div>
PKLR	c.1436G>A p.R479H Stopgain-LOF NM_001011645	4.81% <div></div>
EZH2	c.1936T>A p.Y646F Nonsense-LOF NM_001011645	3.14% <div></div>
Germline	Mutation effect	Condition
JAK3	c.3113A>G p.L857P Missensevariant(exon2)-GOF NM_001011645	anyone

LOW COVERAGE REGIONS

ZEB2 TOP2A

SOMATIC VARIANT DETAILS - POTENTIALLY ACTIONABLE

ARID2

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

VAF: 35.9%

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.

IDH1

c.394_395delinsTC p.R132H Nonsense-GOF

VAF: 32.59%

KRAS is a GDP/GTP binding protein that acts as an intracellular signal transducer. KRAS is involved in several pathways involved in cellular proliferation and survival, including the PI3K-AKT-mTOR pathway and the Ras-Raf-MEK-ERK pathway. Activating mutations, copy number gains, and overexpression of KRAS are associated with cancer progression.

TOP2A

c.3113A>G p.A1515S Stopgain-LOF

VAF: 25.8%

KRAS is a GDP/GTP binding protein that acts as an intracellular signal transducer. KRAS is involved in several pathways involved in cellular proliferation and survival, including the PI3K-AKT-mTOR pathway and the Ras-Raf-MEK-ERK pathway. Activating mutations, copy number gains, and overexpression of KRAS are associated with cancer progression.

SOMATIC VARIANT DETAILS - BIOLOGICALLY RELEVANT

FGFR3

c.1111A>T p.A1515S Spliceregionvariant-LOF

VAF: 15.97%

RBM10 encodes a protein that contains a RNA-binding motif and interacts with RNA homopolymers, and is thought to function in regulating alternative splicing. Loss of function mutations and copy number loss of RBM10 are associated with cancer progression.

IDH1

c.394_395delinsTC p.R132C Nonsense-LOF

VAF: 2.0%

NFE2L2 acts as a transcription factor for proteins that contain an antioxidant response element (ARE) within their promoter sequence. Genes that contain ARE are involved in injury and inflammation response. Activating mutations and overexpression of NFE2L2 are associated with cancer progression.

CLINICAL HISTORY

Diagnosed on
2023-05-06