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PATIENT	DOB	DISEASE	MRN	REPORT DATE	REPORT STATUS
[REDACTED]	[REDACTED]	[REDACTED]	TSO500		Final

REPORT SUMMARY

Executive Summary

[Please add your summary for this case.]

Other Biomarkers

BIOMARKER	LEVEL
TMB	High
MSI	Unstable

Genomic Findings

IA	IB	IIC	IID
No variants reported.	No variants reported.	BRCA1 p.K339Rfs*2 c.1016delA	TET1 p.K22Rfs*23 c.65delA
		RNF43 p.G659Vfs*4 1 c.1976delG	WT1 p.R369* c.1105C>T
		CSF3R p.P468Qfs* 5 c. 1403_1404d elCC	HNF1A p.P291Qfs* 51 c.864delG
		MLH1 p.R497Gfs* 11 c.1489delC	GLI1 p.G274Afs*6 c.821delG
		KDR p.Q472H c.1416A>T	NOTCH3 p.G2035Vfs* 50 c.6102delC
		SDHA p.T308M c.923C>T	NOTCH3 p.A1802Gfs* 8 c.5404dupG
		BCOR p.K839Sfs*1 7 c.2514delC	CIC p.P1529Lfs c.4586delC
		DNMT3A p.R729W c.2185C>T	PTPRS p.V1025Sfs* 18 c.3072delC
		TET2 p.Q1546* c.4636C>T	ASXL1 p.L815P c.2444T>C

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FLT4	p.G1154R c.3460G>A	BCORL1	p.P1681Qfs *20 c.5042delC
FGFR4	p.G388R c.1162G>A	MED12	p.Q2115del c. 6276_6278d elGCA
CARD11	p.R555Gfs* 45 c.1663delC	BRCA1	p.K1183R c.3548A>G
TP53	p.G244C c.730G>T	MUTYH	p.R245H c.734G>A
ERBB2	p.P1170A c.3508C>G	BCOR	p.P1621Qfs *53 c.4862delC

27 Clinical Trials

0 Clinical Trials

CLINICALLY RELEVANT RESULTS

Tier I - Strong Clinical Significance

No variants were reported for this classification tier.

Tier II - Potential Clinical Significance

VARIANT	CLINICAL IMPACT
<p>BRCA1</p> <p>p.K339Rfs*2 c.1016delA</p> <p>C</p> <p>NM_007294.3 VAF % 25.1 DEPTH 275</p>	<p>May benefit from</p> <ul style="list-style-type: none"> — Rucaparib or Olaparib <i>in Hormone refractory prostate cancer or Adenocarcinoma of prostate</i> — Talazoparib <i>in Human epidermal growth factor 2 negative carcinoma of breast, Malignant tumor of breast, Infiltrating duct carcinoma of breast, Invasive micropapillary carcinoma of breast, Mixed ductal and lobular carcinoma of breast, Mucinous carcinoma of breast, Inflammatory carcinoma of breast, Infiltrating</i>



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PATIENT

DOB

DISEASE

MRN
ONC4

REPORT DATE

REPORT STATUS
Final

REPORT SUMMARY

Executive Summary

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Other Biomarkers

BIOMARKER	LEVEL
TMB	Medium
MSI	Stable

Genomic Findings

IA	IB	IIC	IID
No variants reported.	<p>TP53</p> <p>p.R213* c.637C>T</p> <p>4 Clinical Trials</p>	<p>CDK6</p> <p>Copy number gain in CDK6 (3 copies)</p> <p>EGFR</p> <p>Copy number gain in EGFR (3 copies)</p> <p>MET</p> <p>Copy number gain in MET (3 copies)</p> <p>FGFR3</p> <p>p.F384L c.1150T>C</p> <p>6 Clinical Trials</p>	No variants reported.

CLINICALLY RELEVANT RESULTS

Tier I - Strong Clinical Significance

VARIANT	CLINICAL IMPACT
<p>TP53</p> <p>p.R213*</p>	<p>Unfavorable Prognosis in</p> <p>— Carcinoma of colon</p>



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PATIENT	DOB	DISEASE	MRN	REPORT DATE	REPORT STATUS
[REDACTED]	[REDACTED]	[REDACTED]	ONC3		Final

REPORT SUMMARY

Executive Summary

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Other Biomarkers

BIOMARKER	LEVEL
TMB	Low
MSI	Stable

Genomic Findings

IA	IB	IIC	IID
No variants reported.	No variants reported.	DNMT3A p.N552Kfs* 99 c.1656delC	No variants reported.

[1 Clinical Trial](#)

CLINICALLY RELEVANT RESULTS

Tier I - Strong Clinical Significance

No variants were reported for this classification tier.

Tier II - Potential Clinical Significance

VARIANT	CLINICAL IMPACT
DNMT3A p.N552Kfs*99 c.1656delC C	Unfavorable Prognosis in — Myeloproliferative neoplasm or Myelosclerosis with myeloid metaplasia
NM_022552.4 VAF % 2.1 DEPTH 292	



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[REDACTED]	[REDACTED]	[REDACTED]	ONC1		Final

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Other Biomarkers

BIOMARKER	LEVEL
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TMB	Low
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MSI	Stable
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Genomic Findings

IA	IB	IIC	IID
KRAS p.G12D c.35G>A 14 Clinical Trials	FBXW7 p.R347C c.1039C>T TP53 p.Y220C c.659A>G 3 Clinical Trials	FGFR4 p.G388R c.1162G>A SMAD4 p.R361H c.1082G>A 2 Clinical Trials	No variants reported.

CLINICALLY RELEVANT RESULTS

Tier I - Strong Clinical Significance

VARIANT	CLINICAL IMPACT
KRAS p.G12D c.35G>A A NM_033360.2 VAF % 4 DEPTH 1.058	<p>Not likely to benefit from</p> <ul style="list-style-type: none"> — Cetuximab or Panitumumab <i>in Malignant tumor of colon</i> <p>Contraindicated</p> <ul style="list-style-type: none"> — Oxaliplatin + Panitumumab or Cetuximab + Oxaliplatin <i>in Malignant tumor of colon</i> <p>INTERPRETATION</p> <p>The KRAS protein has intrinsic GTPase activity and is an important mediator of growth factor receptor signaling resulting in the activation of the PI3K-mTOR and RAS-RAF-MEK pathways (RefSeq, Jul 2008). Cetuximab in combination with oxaliplatin is contraindicated per the EMA (Cetuximab, Revision 28) in metastatic colorectal cancer harboring a RAS mutation. Panitumumab in combination with</p>



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Other Biomarkers

BIOMARKER	LEVEL
TMB	Medium
MSI	Stable

Genomic Findings

	IA	IB	IIC	IID
KDR	p.Q472H c.1416A>T	TP53 p.R273C c.817C>T	ATM p.S707P c.2119T>C	BRCA2 Copy number gain in BRCA2 (4 copies)
KRAS	p.G12V c.35G>T	1 Clinical Trial	BRCA2 p.D1420Y c.4258G>T	CCND3 p.S259A c.775T>G
22 Clinical Trials			ETV6 p.L201P c.602T>C	FANCA p.S858R c.2574C>G
			FGF9 Copy number gain in FGF9 (5 copies)	FGF14 Copy number gain in FGF14 (4 copies)
			FGFR2 p.S57L c.170C>T	LAMP1 Copy number gain in LAMP1 (5 copies)
			MET p.N375S c.1124A>G	TCF7L2 p.? c.1162-2A>G
				1 Clinical Trial



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Other Biomarkers

BIOMARKER	LEVEL
TMB	High
MSI	Stable

Genomic Findings

IA		IB		IIC		IID	
KDR	p.Q472H c.1416A>T	TP53	p.R175H c.524G>A	RICTOR	Copy number gain in RICTOR (3 copies)	BRCA2	Copy number gain in BRCA2 (3 copies)
0 Clinical Trials		2 Clinical Trials		LRP1B	p.P4016L c.12047C>T	CCND3	p.S259A c.775T>G
				APC	p.E1397* c.4189G>T	FGF9	Copy number gain in FGF9 (3 copies)
				FGFR2	p.S57L c.170C>T	LAMP1	Copy number gain in LAMP1 (3 copies)
				1 Clinical Trial		0 Clinical Trials	

CLINICALLY RELEVANT RESULTS

Tier I - Strong Clinical Significance



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Other Biomarkers

BIOMARKER	LEVEL
TMB	Medium
MSI	Stable

Genomic Findings

IA	IB	IIC	IID
KDR p.Q472H c.1416A>T 2 Clinical Trials	No variants reported.	FGFR4 p.G388R c.1162G>A 1 Clinical Trial	CCND3 p.S259A c.775T>G 0 Clinical Trials

CLINICALLY RELEVANT RESULTS

Tier I - Strong Clinical Significance

VARIANT	CLINICAL IMPACT
KDR p.Q472H c.1416A>T A NM_002253.2 VAF% 43.5 DEPTH 347	INTERPRETATION KDR encodes vascular endothelial growth factor receptor (VEGFR-2) which functions as the main mediator of VEGF-induced endothelial proliferation and angiogenesis (RefSeq, May 2009). A missense substitution in KDR, Q472H is identified. Q472H (rs1870377) is a known polymorphism (population frequencies: ExAC = 22%; NHLBI = 19.4%). In literature, KDR Q472H (rs1870377) has been classified as a functional polymorphism and as an activating mutation. KDR Q472H has been demonstrated to be an activating mutation in vitro, resulting in increased phosphorylation of KDR (PMID: 21712447; http://cancerres.aacrjournals.org/content/75/15_Supplement/73). KDR Q472H was associated with proliferation and invasion in vitro; and with the increased microvessel density in non-small cell lung cancer (NSCLC) and melanoma patients (PMID: 21712447; 26631613). KDR Q472H has been reported in colorectal cancer, haemangioblastoma, rhabdomyosarcoma, hematopoietic and lymphoid neoplasms (COSMIC, accessed Oct 2017). A study reported an inverse relationship of



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Other Biomarkers

BIOMARKER	LEVEL
TMB	Low
MSI	Stable

Genomic Findings

IA	IB	IIC		IID	
No variants reported.	No variants reported.	APC	p.R283* c.847C>T	CCND3	p.S259A c.775T>G
		APC	p.Q1367* c.4099C>T	CHEK1	Copy number gain in CHEK1 (3 copies)
		ATM	p.S49C c.146C>G	RET	p.R982C c.2944C>T
		ERBB2	p.A386D c.1157C>A	ZNF217	p.E914_P915delinsDS c.2742_2743delinsTT
		SMARCB1	p.R374Q c.1121G>A		
		TP53	p.P278R c.833C>G		
		1 Clinical Trial		0 Clinical Trials	

CLINICALLY RELEVANT RESULTS

Tier I - Strong Clinical Significance

No variants were reported for this classification tier.



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BIOMARKER	LEVEL
TMB	Medium
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Genomic Findings

IA	IB	IIC	IID
No variants reported.	<p>PIK3CA p.G1049R c.3145G>C</p> <p>2 Clinical Trials</p>	<p>APC p.R876* c.2626C>T</p> <p>ATM p.F858L c.2572T>C</p> <p>MUTYH p.Y179C c.536A>G</p> <p>NOTCH1 p.R621H c.1862G>A</p> <p>ZRSR2 p.S447_R44 8dup c. 1338_1343d up6</p> <p>ABL1 p.(=) c.3324A>G</p>	<p>BRCA2 Copy number gain in BRCA2 (3 copies)</p> <p>CCND3 p.S259A c.775T>G</p> <p>LAMP1 Copy number gain in LAMP1 (3 copies)</p> <p>0 Clinical Trials</p>



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Other Biomarkers

BIOMARKER	LEVEL
TMB	Medium
MSI	Stable

Genomic Findings

IA		IB		IIC		IID	
KDR	p.Q472H c.1416A>T	TP53	p.? c.559+1G>T	APC	p.E536* c.1606G>T	CCND3	p.S259A c.775T>G
KRAS	p.G12V c.35G>T	0 Clinical Trials		APC	p.L1489Yfs* 18 c.4466delT	FANCA	p.S858R c.2574C>G
3 Clinical Trials				ATM	p.R1618* c.4852C>T	MDC1	p.P1177S c. 3528_3529d elinsAT
				FGFR4	p.G388R c.1162G>A	0 Clinical Trials	
				MYC	p.N26S c.77A>G		
				PIK3R1	p.W583del c. 1748_1750d elGGT		
				ABL1	p.(-) c.3324A>G		