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Sample Information

Patient Name: 羅雪香 Gender: Female ID No.: G201055697 History No.: 46942976

Age: 66

Ordering Doctor: DOC8518J 黃昱凱 Ordering REQ.: OBGCVRY

Signing in Date: 2021/05/20

Path No.: S110-98843 **MP No.:** F21042

Assay: Oncomine Focus Assay

Sample Type: FFPE Block No.: S110-66363E Percentage of tumor cells: 80%

Note:

Sample Cancer Type: Gastrointestinal Stromal Tumor

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Relevant Gastrointestinal Stromal Tumor Variants

Gene	Finding
KIT	KIT exon 11 deletion
NTRK1	Not detected
NTRK2	Not detected
NTRK3	Not detected
PDGFRA	Not detected

Relevant Biomarkers

Tier	Genomic Alteration	Relevant Therapies (In this cancer type)	Relevant Therapies (In other cancer type)	Clinical Trials
IA	KIT exon 11 deletion	imatinib	imatinib	11
	KIT proto-oncogene, receptor tyrosine kinase Allele Frequency: 34.63%			

Tier Reference: Li et al. Standards and Guidelines for the Interpretation and Reporting of Sequence Variants in Cancer: A Joint Consensus Recommendation of the Association for Molecular Pathology, American Society of Clinical Oncology, and College of American Pathologists. J Mol Diagn. 2017 Jan;19(1):4-23.

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Variant Details

DNA Sequence Variants							
Amino Acid Change	Coding	Variant ID	Locus	Allele Frequency	Transcript	Variant Effect	Coverage
p.(D579del)	c.1735_1737delGAT	COSM1294	chr4:55593666	34.63%	NM_000222.2	nonframeshift Deletion	1975
p.(=)	c.2199A>G		chr1:65310489	76.29%	NM_002227.3	synonymous	1881
p.(D1529E)	c.4587C>G		chr2:29416366	49.17%	NM_004304.4	missense	1997
p.(=)	c.3375C>A		chr2:29445458	46.07%	NM_004304.4	synonymous	1995
p.(P136L)	c.407C>T		chr5:176517797	99.05%	NM_213647.2	missense	2000
p.(N375S)	c.1124A>G		chr7:116340262	50.58%	NM_001127500.2	missense	1997
	Amino Acid Change p.(D579del) p.(=) p.(D1529E) p.(=) p.(P136L)	Amino Acid Change Coding p.(D579del) c.1735_1737delGAT p.(=) c.2199A>G p.(D1529E) c.4587C>G p.(=) c.3375C>A p.(P136L) c.407C>T	Amino Acid Change Coding Variant ID p.(D579del) c.1735_1737delGAT COSM1294 p.(=) c.2199A>G . p.(D1529E) c.4587C>G . p.(=) c.3375C>A . p.(P136L) c.407C>T .	Amino Acid Change Coding Variant ID Locus p.(D579del) c.1735_1737delGAT COSM1294 chr4:55593666 p.(=) c.2199A>G . chr1:65310489 p.(D1529E) c.4587C>G . chr2:29416366 p.(=) c.3375C>A . chr2:29445458 p.(P136L) c.407C>T . chr5:176517797	Amino Acid ChangeCodingVariant IDLocusAllele Frequencyp.(D579del)c.1735_1737delGATCOSM1294chr4:5559366634.63%p.(=)c.2199A>G.chr1:6531048976.29%p.(D1529E)c.4587C>G.chr2:2941636649.17%p.(=)c.3375C>A.chr2:2944545846.07%p.(P136L)c.407C>T.chr5:17651779799.05%	Amino Acid ChangeCodingVariant IDLocusAllele FrequencyTranscriptp.(D579del)c.1735_1737delGATCOSM1294chr4:5559366634.63%NM_000222.2p.(=)c.2199A>G.chr1:6531048976.29%NM_002227.3p.(D1529E)c.4587C>G.chr2:2941636649.17%NM_004304.4p.(=)c.3375C>A.chr2:2944545846.07%NM_004304.4p.(P136L)c.407C>T.chr5:17651779799.05%NM_213647.2	Amino Acid Change Coding Variant ID Locus Allele Frequency Transcript Variant Effect p.(D579del) c.1735_1737delGAT COSM1294 chr4:55593666 34.63% NM_000222.2 nonframeshift Deletion p.(=) c.2199A>G . chr1:65310489 76.29% NM_002227.3 synonymous p.(D1529E) c.4587C>G . chr2:29416366 49.17% NM_004304.4 missense p.(=) c.3375C>A . chr2:29445458 46.07% NM_004304.4 synonymous p.(P136L) c.407C>T . chr5:176517797 99.05% NM_213647.2 missense

Biomarker Descriptions

KIT (KIT proto-oncogene, receptor tyrosine kinase)

Background: The KIT gene, also known as CD117, encodes the KIT proto-oncogene receptor tyrosine kinase (c-KIT), a member of the PDGF receptor type III receptor tyrosine kinase family, which includes PDGFRA, PDGFRB, CSF1R, FLT1, FLT3, FLT4 and KDR¹.². KIT is a receptor for stem cell factor, important in regulating growth and development of hematopoietic cells³. The KIT gene is flanked by the PDGFRA and KDR genes on chromosome 4q12. Ligand binding to KIT results in kinase activation and stimulation of downstream pathways including the RAS/RAF/MEK/ERK and PI3K/AKT/MTOR pathways promoting cell proliferation and survival⁴.

Alterations and prevalence: Recurrent somatic KIT alterations are observed in both solid and hematological cancers and include activating mutations such as single nucleotide variants, small duplications, and complex in-frame insertions or deletions (indels). Mutations in KIT exons 8, 9, 11, and 17 disrupt auto-inhibitory mechanisms and lead to constitutive activity⁵. Gain of function mutations are found in up to 70% of mast cell tumors, 17% of nasal T-cell lymphomas, and 9% of dysgerminoma⁶. Somatic mutations in exon 11 occur in 60-70% of all gastrointestinal stromal tumor (GIST), whereas alterations in exons 8 and 17 are more common in myeloid cancers^{5,6,7}. A common kinase domain mutation that causes ligand-independent constitutive activation, D816V, occurs in 80-93% of aggressive forms of mastocytosis^{8,9}.

Potential relevance: Imatinib¹⁰ (2001) is approved for KIT positive malignant GIST and adult patients with aggressive systemic mastocytosis (SM) harboring D816V mutations. Imatinib is also recommended for KIT activating mutations in melanoma and exon 9 and 11 mutations in GIST^{11,12,13}. Mutations in exon 17 have been identified to confer resistance to imatinib and sunitinib¹⁴. Patients with acute myeloid leukemia (AML) that harbor KIT activating mutations with t(8;21) and inv(16) have an increased risk of relapse¹⁵. KIT D816V mutation is associated with the diagnosis of SM and aggressiveness of the disease^{16,17}.

Relevant Therapy Summary

in this cancer type	In other cancer type	in this cancer	type and other car	icer types	No eviden	ce
KIT exon 11 del	etion					
Relevant Therapy		FDA	NCCN	EMA	ESMO	Clinical Trials*
imatinib		×	•	×		(III)
avelumab, axitinib		×	×	×	×	(II)
cabozantinib		×	×	×	×	(II)

^{*} Most advanced phase (IV, III, II/III, II, I/II, I) is shown and multiple clinical trials may be available.

Relevant Therapy Summary (continued)

In this cancer type

In other cancer type

In this cancer type and other cancer types

X No evidence

KIT exon 11 deletion (continued)

Relevant Therapy	FDA	NCCN	EMA	ESMO	Clinical Trials*
dasatinib, sunitinib	×	×	×	×	(II)
nilotinib, pazopanib	×	×	×	×	(II)
ponatinib	×	×	×	×	(II)
sunitinib, regorafenib	×	×	×	×	(II)
spartalizumab, imatinib	×	×	×	×	(1/11)

^{*} Most advanced phase (IV, III, II/II, II, I/II, I) is shown and multiple clinical trials may be available.

Relevant Therapy Details

Current NCCN Information

NCCN information is current as of 2021-04-01. For the most up-to-date information, search www.nccn.org. For NCCN International Adaptations & Translations, search www.nccn.org/global/international_adaptations.aspx.

KIT exon 11 deletion

imatinib

Cancer type: Gastrointestinal Stromal Tumor Variant class: KIT exon 11 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

Progression, Resectable (Neoadjuvant therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Gastrointestinal Stromal Tumor [Version 1.2021]

O imatinib

Cancer type: Melanoma Variant class: KIT exon 11 activating mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

 Cutaneous; Metastatic, Unresectable, Progression (Second-line therapy, Subsequent therapy); Useful in certain circumstances

Reference: NCCN Guidelines® - NCCN-Cutaneous Melanoma [Version 2.2021]

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Current ESMO Information

In this cancer type
In other cancer type
In this cancer type and other cancer types

ESMO information is current as of 2021-04-01. For the most up-to-date information, search www.esmo.org.

KIT exon 11 deletion

imatinib

Cancer type: Gastrointestinal Stromal Tumor Variant class: KIT exon 11 deletion

ESMO Level of Evidence/Grade of Recommendation: I / A

Population segment (Line of therapy):

Advanced, Metastatic (Line of therapy not specified)

Reference: ESMO Clinical Practice Guidelines - ESMO-EUROCAN-Gastrointestinal Stromal Tumours [Ann Oncol (2018) 29 (Suppl 4): iv68-iv78. (Corrigendum: 05 September 2018)]

Clinical Trials Summary

KIT exon 11 deletion

NCT ID	Title	Phase
NCT02413736	Three versus Five Years of Adjuvant Imatinib as Treatment of Patients with Operable GIST with a High Risk for Recurrence: A Randomised Phase III Study	III
NCT02712112	Randomized Phase 2 Study of Intermittent vs Continuous Dosing Schedule of Imatinib in Patients With Tyrosine Kinase Inhibitor Refractory Gastrointestinal Stromal Tumors (GISTs)	II
NCT03171389	Phase II Trial of Ponatinib in Patients With Metastatic and/or Unresectable Gastrointestinal Stromal Tumor (GIST) Following Failure or Intolerance of Prior Therapy With Imatinib (POETIG trial – POnatinib after rEsisTance to Imatinib in GIST)	II
NCT03609424	A Phase Ib/II Study Of PDR001 Plus Imatinib For Metastatic Or Unresectable GIST With Prior Failure Of Imatinib, Sunitinib And Regorafenib	1/11
NCT04258956	A Phase II, Single Arm Study of Avelumab In Combination With Axitinib in Patients With Unresectable/ Metastatic Gastrointestinal Stromal Tumor After Failure of Standard Therapy - AXAGIST	II
NCT04116541	MegaMOST - A Multicenter, Open-label, Biology Driven, Phase II Study Evaluating the Activity of Anti- cancer Treatments Targeting Tumor Molecular Alterations /Characteristics in Advanced / Metastatic Tumors.	II
NCT02461849	A Phase II, Open-label, Study in Patients With Refractory, Metastatic Cancer Harboring KIT Mutation or Amplification to Investigate the Clinical Efficacy and Safety of Imatinib Therapy.	II
NCT02029001	A Two-period, Multicenter, Randomized, Open-label, Phase II Study Evaluating the Clinical Benefit of a Maintenance Treatment Targeting Tumor Molecular Alterations in Patients With Progressive Locally-advanced or Metastatic Solid Tumors MOST: My own specific treatment	II
NCT02693535	Targeted Agent and Profiling Utilization Registry (TAPUR) Study	II
NCT03297606	Canadian Profiling and Targeted Agent Utilization Trial (CAPTUR): A Phase II Basket Trial	II
NCT02272998	Phase II Study Of Ponatinib For Advanced Cancers With Genomic Alterations In Fibroblastic Growth Factor Receptor (FGFR) And Other Genomic Targets (KIT, Pdgfra, RET FLT3, ABL1)	II

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Signatures

Pathologist:

Testing Personnel:

Laboratory Supervisor:

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