

Department of Pathology and Laboratory Medicine No.201, Sec. 2, Shipai Rd., Beitou District, Taipei City, Taiwan 11217, R.O.C. Tel: 02-2875-7449

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Sample Information

Patient Name: 留麗絹 Gender: Female ID No.: P220769096 History No.: 18478267

Age: 61

Ordering Doctor: DOC6372L 張彥安 Ordering REQ.: OCHBPDY Signing in Date: 2023/03/16

Path No.: M112-00041 **MP No.**: MY23013

Assay: Oncomine Myeloid Assay **Sample Type:** Bone Marrow

Bone Marrow Aspirating Date: 2023/03/08

Reporting Doctor: DOC5444B 楊靜芬 (Phone: 8#5444)

Note:

Sample Cancer Type: Acute Myeloid Leukemia

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0 Clinical Trials

Relevant Acute Myeloid Leukemia Variants

Gene	Finding	Gene	Finding
ABL1	None detected	MECOM	None detected
ASXL1	None detected	MLLT3	None detected
CEBPA	None detected	MYH11	None detected
CREBBP	None detected	NPM1	NPM1 p.(W288Cfs*12) c.863_864insTCTG
FLT3	FLT3 ITD mutation	NUP214	None detected
IDH1	None detected	RARA	None detected
IDH2	None detected	RUNX1	None detected
KMT2A	None detected	TP53	None detected

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Relevant Biomarkers

Tier	Genomic Alteration	Relevant Therapies (In this cancer type)	Relevant Therapies (In other cancer type)	Clinical Trials
IA	FLT3 ITD mutation fms related receptor tyrosine kinase 3 Allele Frequency: 8.60%, 22.30% (2 variants)	gilteritinib 1,2 midostaurin + chemotherapy 1,2 cytarabine + daunorubicin cytarabine + daunorubicin + etoposide cytarabine + etoposide + idarubicin cytarabine + fludarabine + idarubicin + filgrastim cytarabine + idarubicin cytarabine + mitoxantrone gemtuzumab ozogamicin + chemotherapy sorafenib sorafenib + chemotherapy venetoclax + chemotherapy	None	0
	Prognostic significance: ELN 2017:	Favorable to Intermediate		
IA	NPM1 p.(W288Cfs*12) c.863_864insTCTG nucleophosmin 1 Allele Frequency: 44.61%	cytarabine + daunorubicin cytarabine + daunorubicin + etoposide cytarabine + etoposide + idarubicin cytarabine + fludarabine + idarubicin + filgrastim cytarabine + idarubicin cytarabine + mitoxantrone gemtuzumab ozogamicin + chemotherapy midostaurin + chemotherapy	None	0
	Prognostic significance: ELN 2017: Diagnostic significance: Acute Mye	Favorable to Intermediate		

Public data sources included in relevant therapies: FDA1, NCCN, EMA2, ESMO Public data sources included in prognostic and diagnostic significance: NCCN, ESMO

Prevalent cancer biomarkers without relevant evidence based on included data sources

TET2 p.(I1873N) c.5618T>A, TET2 p.(L11111Ffs*19) c.3332_3333insT

Variants (Exclude variant in Taiwan BioBank with >1% allele frequency)

DNA	Sequence Varia	ants						
Gene	Amino Acid Change	Coding	Variant ID	Locus	Allele Frequency	Transcript	Variant Effect	Coverage
TET2	p.(L1111Ffs*19)	c.3332_3333insT		chr4:106158429	50.25%	NM_001127208.2	frameshift Insertion	1986
TET2	p.(I1873N)	c.5618T>A		chr4:106197285	5.80%	NM_001127208.2	missense	2000
NPM1	p.(W288Cfs*12)	c.863_864insTCTG	COSM17559	chr5:170837547	44.61%	NM_002520.6	frameshift Insertion	408
FLT3	p.(E598_F612dup)	c1_0insAATATGAT CTCAAATGGGAGTT TCCAAGAGAAAATT TAGAGTTTG		chr13:28608218	8.60%	NM_004119.3	nonframeshift Insertion	1965

Variants (Exclude variant in Taiwan BioBank with >1% allele frequency) (continued)

DNA Sequence Variants (continued)

Gene	Amino Acid Change	Coding	Variant ID	Locus	Allele Frequency	Transcript	Variant Effect	Coverage
FLT3	p.(Y597_L610dup)	c.1787_1787delAinsA ATATGAATATGATC TCAAATGGGAGTTT CCAAGAGAAAATTT	۸.	chr13:28608269	22.30%	NM_004119.3	nonframeshift Block Substitution	
ASXL1	p.(N986S)	c.2957A>G		chr20:31023472	50.10%	NM_015338.6	missense	2000

Biomarker Descriptions

FLT3 (fms related receptor tyrosine kinase 3)

<u>Background</u>: The FLT3 gene encodes the fms related tyrosine kinase 3, a tyrosine kinase receptor that is a member of the class III receptor tyrosine kinase family that also includes PDGFR, FMS, and KIT¹. FLT3 is highly expressed in hematopoietic progenitor cells². Genomic alterations in FLT3 activate downstream oncogenic pathways including PI3K/AKT/mTOR and RAS/RAF/MEK/ERK pathways which promote cellular proliferation, survival, and inhibition of differentiation¹.

Alterations and prevalence: Somatic mutations occur in approximately 30% of acute myeloid leukemia (AML), 7-10% of melanoma, and up to 8% of uterine cancer^{3,4,5,6}. The most common activating FLT3 mutations are internal tandem duplications (ITD) that range from 3 to 400 base pairs in length within exons 14 and 15 in the juxtamembrane (JM) domain⁷. The second most frequent mutations are point mutations in exon 20 within the tyrosine kinase domain (TKD)⁸. FLT3 is amplified in up to 8% of colorectal cancer, 3% of stomach cancer, and is commonly overexpressed in AML^{5,6,9}.

Potential relevance: The presence of FLT3-ITD confers poor prognosis in myelodysplastic syndrome (MDS) and AML¹0,¹¹. Similarly, the FLT3 TKD mutation D835 confers poor prognosis in MDS¹0. Midostaurin¹² (2017) and gilteritinib¹³ (2018) are kinase inhibitors approved for AML patients with FLT3-ITD and TKD mutations including D835 and I836 mutations. The FDA granted fast track designations in 2017 to crenolanib¹⁴ and in 2022 to tuspetinib (HM43239)¹⁵ for FLT3 mutation-positive relapsed or refractory AML. In 2018 the FDA granted breakthrough therapy designation to quizartinib¹⁶ for AML with FLT3-ITD. A phase II trial testing crenolanib in 34 patients with FLT3-ITD and TKD mutated relapsed/refractory AML, reported that FLT3 inhibitor naïve patients demonstrated a longer overall survival (OS) and event free survival (EFS) in comparison to previously treated patients (median OS: 55 weeks vs 13 weeks; median EFS: 13 weeks vs 7 weeks)¹७. Another phase II trial of crenolanib with chemotherapy in newly diagnosed FLT3 mutated AML reported complete remission in 24/29 (83%) patients¹в. Several multi-targeted tyrosine kinase inhibitors such as sorafenib (2005), sunitinib (2006), cabozantinib (2012), and ponatinib (2012) are FDA approved and include FLT3 as a target. Sorafenib is recommended in combination with chemotherapy in FLT3-ITD mutated AML¹¹1.

NPM1 (nucleophosmin 1)

Background: The NPM1 gene encodes the nucleophosmin protein, a histone chaperone of the nucleophosmin/nucleoplasmin family, which also includes NPM2 and NPM3¹⁹. NPM1 functions as an oncogene and tumor suppressor, and is important in maintaining genomic stability, DNA repair, and apoptosis^{19,20}. NPM1 has a highly conserved N-terminal region which constitutes the core domain responsible for oligomerization, an acidic domain, a nuclear localization signal, and a disorganized C-terminal region which is required for nucleolar localization¹⁹. Oligomerization of NPM1 localizes the protein in the nucleus of proliferating cells where it binds to Akt in response to growth factor stimulation and escapes proteolytic degradation by caspase activity, thereby promoting cell survival^{19,20}. NPM1 is one of the most frequently altered genes in hematological cancers²¹. Most NPM1 mutations occur in the C-terminus, impacting protein folding or the nucleolar localization signal, and result in the localization of NPM1 to the cytoplasm (NPMc) instead of to the nucleus¹⁹.

Alterations and prevalence: NPM1 mutations are observed in 45-60% of AML with a normal karyotype (NK-AML), 28-35% of de novo acute myeloid leukemia (AML) and are frequently co-mutated with DNMT3A and/or FLT3-ITD^{11,22,23}. NPM1 fusions are associated with distinct partner genes in acute promyelocytic leukemia (APL), anaplastic large-cell lymphoma (ALCL), AML, and myelodysplasia²¹. Specifically, NPM1-ALK fusion is found in 30% of all ALCL and this specific fusion is observed in 85% of ALK-positive ALCL¹⁹. The t(5;17)(q35;q21) translocation that results in NPM1-RARA fusion is observed in APL²⁴.

Potential relevance: NPM1 mutated AML is recognized as a distinct diagnostic disease entity by the World Health Organization (WHO)²⁵. NPM1 mutations are associated with better outcomes, increased complete remission, and improved overall survival in AML^{11,23}. NPM1 without FLT3-ITD mutations or with <0.5 allelic ratio FLT3-ITD mutations are associated with favorable risk in AML¹¹. Concurrent NPM1 and with >0.5 allelic ratio FLT3-ITD mutations confer intermediate risk in AML, whereas wild-type NPM1 confers

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X No evidence

Biomarker Descriptions (continued)

poor/adverse risk¹¹. The NPM1 frameshift mutation W288fs*12 is associated with poor prognosis in myelodysplastic syndrome (MDS)¹⁰. The ALK-NPM1 fusion, and translocation t(2;5)(p23;q35) which leads to an ALK-NPM1 fusion, is diagnostic of ALK-positive anaplastic large cell lymphoma^{26,27}.

TET2 (tet methylcytosine dioxygenase 2)

Background: TET2 encodes the tet methylcytosine dioxygenase 2 protein and belongs to a family of ten-eleven translocation (TET) proteins that also includes TET1 and TET3²⁸. TET2 is involved in DNA methylation, specifically in the conversion of 5-methylcytosine to 5-hydroxymethylcytosine^{29,30}. The TET proteins contain a C-terminal core catalytic domain that contains a cysteine-rich domain and a double stranded ß-helix domain (DSBH)³¹. TET2 is a tumor suppressor gene. Loss of function mutations in TET2 are associated with loss of catalytic activity and transformation to hematological malignancies^{28,29,30}

Alterations and prevalence: Somatic TET2 mutations, including nonsense, frameshift, splice site, and missense, are observed in 20-25% of myelodysplastic syndrome (MDS) associated diseases, including 40%-60% chronic myelomonocytic leukemia (CMML)¹⁰. TET2 mutations at H1881 and R1896 are frequently observed in myeloid malignancies^{29,32}. TET2 mutations are also observed in 9% of uterine, 8% of melanoma and acute myeloid leukemia (AML), as well as 6% of diffuse large B-cell lymphoma (DLBCL).

Potential relevance: The presence of TET2 mutations may be used as one of the major diagnostic criteria in pre-primary myelofibrosis (pre-PMF) and overt PMF in the absence of JAK2/CALR/MPL mutations^{25,33}. TET2 mutations are associated with poor prognosis in PMF and increased rate of transformation to leukemia^{33,34}

In this cancer type and other cancer types

Relevant Therapy Summary

O In other cancer type

In this cancer type

FLT3 ITD mutation					
Relevant Therapy	FDA	NCCN	EMA	ESMO	Clinical Trials
gilteritinib					×
midostaurin + cytarabine + daunorubicin		•	•		×
cytarabine + daunorubicin	×	•	×	×	×
cytarabine + daunorubicin + etoposide	×	•	×	×	×
cytarabine + etoposide + idarubicin	×	•	×	×	×
cytarabine + fludarabine + idarubicin + filgrastim	×		×	×	×
cytarabine + idarubicin	×	•	×	×	×
cytarabine + mitoxantrone	×	•	×	×	×
gemtuzumab ozogamicin + cytarabine + daunorubicin	×	•	×	×	×
midostaurin + cytarabine	×	•	×	×	×
sorafenib	×	•	×	×	×
sorafenib + azacitidine	×	•	×	×	×
sorafenib + decitabine	×	•	×	×	×
venetoclax + azacitidine	×	•	×	×	×
venetoclax + cytarabine	×		×	×	×

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Relevant Therapy Summary (continued)

In this cancer type

O In other cancer type

In this cancer type and other cancer types

X No evidence

FLT3 ITD mutation (continued)

Relevant Therapy	FDA	NCCN	EMA	ESMO	Clinical Trials*
venetoclax + decitabine	×		×	×	×

NPM1 p.(W288Cfs*12) c.863_864insTCTG

Relevant Therapy	FDA	NCCN	EMA	ESMO	Clinical Trials*
cytarabine + daunorubicin	×		×	×	×
cytarabine + daunorubicin + etoposide	×		×	×	×
cytarabine + etoposide + idarubicin	×	•	×	×	×
cytarabine + fludarabine + idarubicin + filgrastim	×		×	×	×
cytarabine + idarubicin	×		×	×	×
cytarabine + mitoxantrone	×	•	×	×	×
gemtuzumab ozogamicin + cytarabine + daunorubicin	×	•	×	×	×
midostaurin + cytarabine + daunorubicin	×	•	×	×	×

Relevant Therapy Details

Current FDA Information

In this cancer type In other cancer type

In this cancer type and other cancer types

FDA information is current as of 2023-01-18. For the most up-to-date information, search www.fda.gov.

FLT3 ITD mutation

gilteritinib

Cancer type: Acute Myeloid Leukemia Label as of: 2022-01-12 Variant class: FLT3 ITD mutation

Indications and usage:

XOSPATA® is a kinase inhibitor indicated for the treatment of adult patients who have relapsed or refractory acute myeloid leukemia (AML) with a FLT3 mutation as detected by an FDA-approved test.

Reference:

https://www.accessdata.fda.gov/drugsatfda_docs/label/2022/211349s003lbl.pdf

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FLT3 ITD mutation (continued)

midostaurin + cytarabine + daunorubicin

Cancer type: Acute Myeloid Leukemia Label as of: 2021-11-15 Variant class: FLT3 ITD mutation

Indications and usage:

RYDAPT® is a kinase inhibitor indicated for the treatment of adult patients with:

Newly diagnosed acute myeloid leukemia (AML) that is FLT3 mutation-positive as detected by an FDA-approved test, in combination with standard cytarabine and daunorubicin induction and cytarabine consolidation.

Limitations of Use: RYDAPT® is not indicated as a single-agent induction therapy for the treatment of patients with AML.

 Aggressive systemic mastocytosis (ASM), systemic mastocytosis with associated hematological neoplasm (SM-AHN), or mast cell leukemia (MCL).

Reference:

https://www.accessdata.fda.gov/drugsatfda_docs/label/2021/207997s008lbledt.pdf

Current NCCN Information

In this cancer type

O In other cancer type

In this cancer type and other cancer types

NCCN information is current as of 2023-01-03. For the most up-to-date information, search www.nccn.org. For NCCN International Adaptations & Translations, search www.nccn.org/global/international_adaptations.aspx.

FLT3 ITD mutation + NPM1 p.(W288Cfs*12) c.863_864insTCTG

cytarabine + daunorubicin

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

(Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

cytarabine + daunorubicin + etoposide

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

(Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

cytarabine + etoposide + idarubicin

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

(Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

cytarabine + idarubicin

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

(Induction therapy); Other recommended intervention

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FLT3 ITD mutation + NPM1 p.(W288Cfs*12) c.863_864insTCTG (continued)

cytarabine + daunorubicin

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

(Induction therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

cytarabine + idarubicin

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

(Induction therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

cytarabine + mitoxantrone

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

■ (Induction therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

gemtuzumab ozogamicin + cytarabine + daunorubicin

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

(Induction therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

midostaurin + cytarabine + daunorubicin

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

■ (Induction therapy)

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FLT3 ITD mutation + NPM1 p.(W288Cfs*12) c.863_864insTCTG (continued)

cytarabine + daunorubicin + etoposide

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

NCCN Recommendation category: 2B

Population segment (Line of therapy):

(Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

cytarabine + etoposide + idarubicin

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

NCCN Recommendation category: 2B

Population segment (Line of therapy):

(Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

cytarabine + fludarabine + idarubicin + filgrastim

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

NCCN Recommendation category: 2B

Population segment (Line of therapy):

■ (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

FLT3 ITD mutation

gilteritinib

Cancer type: Acute Myeloid Leukemia Variant class: FLT3 ITD mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

Relapsed, Refractory (Line of therapy not specified)

FLT3 ITD mutation (continued)

midostaurin + cytarabine

Cancer type: Acute Myeloid Leukemia Variant class: FLT3 ITD mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

(Consolidation therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

midostaurin + cytarabine + daunorubicin

Cancer type: Acute Myeloid Leukemia Variant class: FLT3 ITD mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

(Subsequent therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

sorafenib

Cancer type: Acute Myeloid Leukemia Variant class: FLT3 ITD mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

■ (Maintenance therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

sorafenib + azacitidine

Cancer type: Acute Myeloid Leukemia Variant class: FLT3 ITD mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention
- Relapsed, Refractory (Line of therapy not specified)

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FLT3 ITD mutation (continued)

sorafenib + decitabine

Cancer type: Acute Myeloid Leukemia Variant class: FLT3 ITD mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

(Induction therapy); Other recommended intervention

Relapsed, Refractory (Line of therapy not specified)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

venetoclax + azacitidine

Cancer type: Acute Myeloid Leukemia Variant class: FLT3 mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

■ (Induction therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

venetoclax + cytarabine

Cancer type: Acute Myeloid Leukemia Variant class: FLT3 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

(Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

venetoclax + decitabine

Cancer type: Acute Myeloid Leukemia Variant class: FLT3 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

■ (Induction therapy); Preferred intervention

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Current EMA Information

In this cancer type

O In other cancer type

In this cancer type and other cancer types

EMA information is current as of 2023-01-18. For the most up-to-date information, search www.ema.europa.eu/ema.

FLT3 ITD mutation

gilteritinib

Cancer type: Acute Myeloid Leukemia Label as of: 2021-09-08 Variant class: FLT3 ITD mutation

Reference:

https://www.ema.europa.eu/en/documents/product-information/xospata-epar-product-information_en.pdf

midostaurin + cytarabine + daunorubicin

Cancer type: Acute Myeloid Leukemia Label as of: 2022-09-23 Variant class: FLT3 ITD mutation

Reference:

https://www.ema.europa.eu/en/documents/product-information/rydapt-epar-product-information_en.pdf

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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 2023-01-03. For the most up-to-date information, search www.esmo.org.

FLT3 ITD mutation

gilteritinib

Cancer type: Acute Myeloid Leukemia Variant class: FLT3 ITD mutation

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

Population segment (Line of therapy):

■ Relapsed, Refractory (Line of therapy not specified)

Reference: ESMO Clinical Practice Guidelines - ESMO-Acute Myeloblastic Leukaemia in Adult Patients [Ann Oncol (2020); 31(6): 697-712.]

midostaurin + cytarabine + daunorubicin

Cancer type: Acute Myeloid Leukemia Variant class: FLT3 ITD mutation

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

Population segment (Line of therapy):

(Induction therapy)

Reference: ESMO Clinical Practice Guidelines - ESMO-Acute Myeloblastic Leukaemia in Adult Patients [Ann Oncol (2020); 31(6): 697-712.]

Prognostic Details

Current NCCN Information

NCCN information is current as of 2023-01-03. For the most up-to-date information, search www.nccn.org. For NCCN International Adaptations & Translations, search www.nccn.org/global/international_adaptations.aspx.

FLT3 ITD mutation + NPM1 p.(W288Cfs*12) c.863_864insTCTG

Prognostic significance: ELN 2017: Intermediate

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

NCCN Recommendation category: 2A

Summary:

FLT3-ITD^{high}; High defined as allelic ratio (≥0.5).

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 2.2022]

Current ESMO Information

ESMO information is current as of 2023-01-03. For the most up-to-date information, search www.esmo.org.

FLT3 ITD mutation + NPM1 p.(W288Cfs*12) c.863_864insTCTG

Prognostic significance: ELN 2017: Intermediate

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

Summary:

FLT3-ITD^{High}; High defined as allelic ratio (≥0.5)

Reference: ESMO Clinical Practice Guidelines - ESMO-Acute Myeloblastic Leukaemia in Adult Patients [Ann Oncol (2020); 31(6): 697-712.]

Prognostic significance: ELN 2017: Favorable

Cancer type: Acute Myeloid Leukemia Variant classes: FLT3 ITD mutation & NPM1 mutation

Summary:

■ FLT3-ITD^{Low}; Low defined as allelic ratio (<0.5)

Reference: ESMO Clinical Practice Guidelines - ESMO-Acute Myeloblastic Leukaemia in Adult Patients [Ann Oncol (2020); 31(6): 697-712.]

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Diagnostic Details

Current ESMO Information

ESMO information is current as of 2023-01-03. For the most up-to-date information, search www.esmo.org.

NPM1 p.(W288Cfs*12) c.863_864insTCTG

Diagnostic significance: Acute Myeloid Leukemia

Variant class: NPM1 mutation

Diagnostic notes:

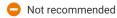
AML with recurrent genetic abnormalities; WHO classification of AML

Reference: ESMO Clinical Practice Guidelines - ESMO-Acute Myeloblastic Leukaemia in Adult Patients [Ann Oncol (2020); 31(6): 697-712.]

Alerts Informed By Public Data Sources

Current FDA Information











Variant class: FLT3 mutation

FDA information is current as of 2023-01-18. For the most up-to-date information, search www.fda.gov.

FLT3 ITD mutation

A crenolanib

Cancer type: Acute Myeloid Leukemia

Supporting Statement:

The FDA has granted Fast Track Designation to the benzimidazole type I kinase inhibitor, crenolanib, for:

- FLT3 mutation-positive relapsed or refractory acute myeloid leukemia (AML)
- PDGFRA D842V mutated unresectable or metastatic gastrointestinal stromal tumors (GIST)

Reference:

https://www.globenewswire.com/news-release/2017/12/01/1216122/0/en/Arog-Pharmaceuticals-Receives-FDA-Fast-Track-Designation-for-Crenolanib-in-Relapsed-or-Refractory-FLT3-Positive-AML.html

tuspetinib

Cancer type: Acute Myeloid Leukemia

Variant class: FLT3 mutation

Supporting Statement:

The FDA has granted Fast Track Designation to tuspetinib (HM43239), a myeloid kinome inhibitor, for relapsed or refractory (R/R) acute myeloid leukemia (AML) with FLT3 mutation.

Reference

https://www.aptose.com/news-media/press-releases/detail/230/aptose-receives-fast-track-designation-for-hm43239-in

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Signatures

Testing Personnel:

Laboratory Supervisor:

Pathologist:

References

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