



## Sample Information

**Patient Name:** 陳正吉  
**Gender:** Male  
**ID No.:** Q100399873  
**History No.:** 43155153  
**Age:** 77

**Ordering Doctor:** DOC1654E 林庭安  
**Ordering REQ.:** H43591B  
**Signing in Date:** 2022/05/20

**Path No.:** S111-99367  
**MP No.:** MY22013  
**Assay:** Oncomine Myeloid Assay  
**Sample Type:** Bone Marrow  
**Bone Marrow Aspirating Date:** 2022/05/12

**Reporting Doctor:** DOC5466K 葉奕成 (Phone: 8#5466)

**Note:**

## Sample Cancer Type: Myelodysplastic/Myeloproliferative Neoplasm

Table of Contents	Page	Report Highlights
Variants (Exclude variant in Taiwan BioBank with >1% allele frequency)	2	2 Relevant Biomarkers
Biomarker Descriptions	2	18 Therapies Available
Relevant Therapy Summary	4	0 Clinical Trials
Relevant Therapy Details	5	
Alert Details	17	

## Relevant Myelodysplastic/Myeloproliferative Neoplasm Variants

Gene	Finding
PDGFRA	None detected
PDGFRB	None detected

## Relevant Biomarkers

Genomic Alteration	Relevant Therapies (In this cancer type)	Relevant Therapies (In other cancer type)	Clinical Trials
<i>TP53</i> p.(C277Y) c.830G>A tumor protein p53 Allele Frequency: 41.70%	None	<b>idelalisib + rituximab</b> <sup>2</sup> acalabrutinib allogeneic stem cells azacitidine cytarabine cytarabine + daunorubicin	0

Public data sources included in relevant therapies: FDA1, NCCN, EMA2, ESMO

## Relevant Biomarkers (continued)

Genomic Alteration	Relevant Therapies (In this cancer type)	Relevant Therapies (In other cancer type)	Clinical Trials
		cytarabine + daunorubicin + etoposide cytarabine + etoposide + idarubicin cytarabine + fludarabine + idarubicin + filgrastim cytarabine + idarubicin cytarabine + mitoxantrone decitabine gemtuzumab ozogamicin + chemotherapy ibrutinib obinutuzumab + venetoclax rituximab + venetoclax venetoclax venetoclax + chemotherapy	
<b><i>RUNX1</i> p.(R201Pfs*11) c.601_602insCC</b> None RUNX family transcription factor 1 Allele Frequency: 44.72%		allogeneic stem cells azacitidine cytarabine cytarabine + daunorubicin cytarabine + daunorubicin + etoposide cytarabine + etoposide + idarubicin cytarabine + fludarabine + idarubicin + filgrastim cytarabine + idarubicin cytarabine + mitoxantrone decitabine gemtuzumab ozogamicin + chemotherapy venetoclax + chemotherapy	0

Public data sources included in relevant therapies: FDA1, NCCN, EMA2, ESMO

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

### DNA Sequence Variants

Gene	Amino Acid Change	Coding	Variant ID	Locus	Allele Frequency	Transcript	Variant Effect	Coverage
TP53	p.(C277Y)	c.830G>A	.	chr17:7577108	41.70%	NM_000546.5	missense	2000
RUNX1	p.(R201Pfs*11)	c.601_602insCC	.	chr21:36231782	44.72%	NM_001754.4	frameshift Insertion	161
EZH2	p.(Y663C)	c.1988A>G	.	chr7:148507466	60.87%	NM_004456.5	missense	23
PTPN11	p.(T468P)	c.1402A>C	.	chr12:112926269	36.92%	NM_002834.5	missense	65

## Biomarker Descriptions

### RUNX1 (RUNX family transcription factor 1)

**Background:** The RUNX1 gene encodes the runt-related transcription factor (RUNX) 1, part of the RUNX family of transcription factors which also includes RUNX2 and RUNX3<sup>1</sup>. All RUNX proteins share several conserved regions with similar functionality including a highly conserved N-terminal 'runt' domain responsible for binding DNA, a C-terminal region composed of an activation domain, inhibitory domain, protein interacting motifs, and a nuclear matrix targeting signal<sup>2</sup>. Each of these proteins are capable of interacting

## Biomarker Descriptions (continued)

with core binding factor beta (CBFβ) to form the core binding factor (CBF) complex. Consequently, RUNX1, RUNX2, and RUNX3 are collectively known as core binding factor alpha (CBFα) since they can each function as the alpha subunit of CBF. Specifically, CBFβ binds to the 'runt' domain of RUNX1 leading to RUNX1 stabilization and increased affinity of the CBF complex for promoters involved in hematopoietic differentiation and cell cycle regulation<sup>3,4</sup>. RUNX1 is frequently mutated in various hematological malignancies<sup>4</sup>. Germline mutations in RUNX1 result in a rare autosomal dominant condition known as familial platelet disorder, with predisposition to acute myeloid leukemia (FPD/AML)<sup>5,6</sup>. Somatic mutations and chromosomal translocations in RUNX1 are often observed in myelodysplastic syndrome (MDS), acute myeloid leukemia (AML), acute lymphoblastic leukemia (ALL), and chronic myelomonocytic leukemia (CMML)<sup>4</sup>.

**Alterations and prevalence:** RUNX1 is frequently rearranged in hematological malignancies with over 50 different observed translocations<sup>7</sup>. The most recurrent translocation, t(12;21)(q34;q11), results in ETV6-RUNX1 fusion and is observed in 20-25% of childhood ALL<sup>8,9,10</sup>. This translocation is also observed in adult ALL at a lower frequency (2%)<sup>9,10</sup>. Another recurrent translocation, t(8;21)(q22;q22), results in RUNX1-RUNX1T1 fusion and is observed in 5-10% of AML<sup>11</sup>. The RUNX1-RUNX1T1 fusion, consists of the RHD domain of RUNX1 and the majority of RUNX1T1, which promotes oncogenesis by altering transcriptional regulation of RUNX1 target genes<sup>4,11</sup>. Somatic mutations in RUNX1 include missense, nonsense, and frameshift mutations resulting in loss of function or dominant negative effects<sup>4</sup>. RUNX1 mutations are reported in approximately 10% of de novo AML as well as 10-15% of MDS<sup>4,12,13,14</sup>.

**Potential relevance:** The t(8;21)(q22;q22.1)/RUNX1-RUNX1T1 translocation is recognized as a distinct AML disease category by the World Health Organization (WHO)<sup>15</sup>. Additionally, AML with RUNX1 mutations is a provisional entity in the WHO<sup>15</sup>. Translocations involving RUNX1, specifically t(8;21)(q22;q22)/RUNX1-RUNX1T1 in AML and t(12;21)(q34;q11)/ETV6-RUNX1 in ALL, are associated with favorable risk<sup>12,16</sup>. On the other hand, mutations in RUNX1 confer poor prognosis in AML, MDS, and systemic mastocytosis (SM)<sup>12,13,17</sup>.

### TP53 (tumor protein p53)

**Background:** The TP53 gene encodes the p53 tumor suppressor protein that binds to DNA and activates transcription in response to diverse cellular stresses to induce cell cycle arrest, apoptosis, or DNA repair. In unstressed cells, TP53 is kept inactive by targeted degradation via MDM2, a substrate recognition factor for ubiquitin-dependent proteolysis. Alterations in TP53 is required for oncogenesis as they result in loss of protein function and gain of transforming potential<sup>18</sup>. Germline mutations in TP53 are the underlying cause of Li-Fraumeni syndrome, a complex hereditary cancer predisposition disorder associated with early-onset cancers<sup>19,20</sup>.

**Alterations and prevalence:** TP53 is the most frequently mutated gene in the cancer genome with approximately half of all cancers experiencing TP53 mutations. Ovarian, head and neck, esophageal, and lung squamous cancers have particularly high TP53 mutation rates (60-90%)<sup>14,21,22,23,24,25</sup>. Approximately two-thirds of TP53 mutations are missense mutations and several recurrent missense mutations are common including substitutions at codons R158, R175, Y220, R248, R273, and R282<sup>14,21</sup>. Invariably, recurrent missense mutations in TP53 inactivate its ability to bind DNA and activate transcription of target genes<sup>26,27,28,29</sup>.

**Potential relevance:** The small molecule p53 reactivator, PC14586, received a fast track designation (2020) by the FDA for advanced tumors harboring a TP53 Y220C mutation<sup>30</sup>. The FDA has granted fast track designation (2019) to the p53 reactivator, eprentapopt<sup>31</sup> and breakthrough designation<sup>32</sup> (2020) in combination with azacitidine or azacitidine and venetoclax for acute myeloid leukemia patients (AML) and myelodysplastic syndrome (MDS) harboring a TP53 mutation, respectively. In addition to investigational therapies aimed at restoring wild-type TP53 activity, compounds that induce synthetic lethality are also under clinical evaluation<sup>33,34</sup>. TP53 mutations confer poor prognosis in multiple blood cancers including AML, MDS, myeloproliferative neoplasms (MPN), and chronic lymphocytic leukemia (CLL)<sup>12,13,35,36</sup>. In mantle cell lymphoma, TP53 mutations are associated with poor prognosis when treated with conventional therapy including hematopoietic cell transplant<sup>37</sup>. Mono- and bi-allelic mutations in TP53 confer unique characteristics in MDS, with multi-hit patients also experiencing associations with complex karyotype, few co-occurring mutations, and high-risk disease presentation as well as predicted death and leukemic transformation independent of the IPSS-R staging system<sup>38</sup>.

## Relevant Therapy Summary

☒ In this cancer type
 ☐ In other cancer type
 ☒ In this cancer type and other cancer types
 ☒ No evidence

### TP53 p.(C277Y) c.830G>A

Relevant Therapy	FDA	NCCN	EMA	ESMO	Clinical Trials*
Allogeneic hematopoietic stem cell transplantation	×	○	×	×	×
azacitidine	×	○	×	×	×
cytarabine	×	○	×	×	×
cytarabine + daunorubicin	×	○	×	×	×
cytarabine + daunorubicin + etoposide	×	○	×	×	×
cytarabine + etoposide + idarubicin	×	○	×	×	×
cytarabine + fludarabine + idarubicin + filgrastim	×	○	×	×	×
cytarabine + idarubicin	×	○	×	×	×
cytarabine + mitoxantrone	×	○	×	×	×
decitabine	×	○	×	×	×
gemtuzumab ozogamicin + cytarabine + daunorubicin	×	○	×	×	×
venetoclax + azacitidine	×	○	×	×	×
venetoclax + cytarabine	×	○	×	×	×
venetoclax + decitabine	×	○	×	×	×
idelalisib + rituximab	×	×	○	○	×
acalabrutinib	×	×	×	○	×
ibrutinib	×	×	×	○	×
obinutuzumab + venetoclax	×	×	×	○	×
rituximab + venetoclax	×	×	×	○	×
venetoclax	×	×	×	○	×

### RUNX1 p.(R201Pfs\*11) c.601\_602insCC

Relevant Therapy	FDA	NCCN	EMA	ESMO	Clinical Trials*
Allogeneic hematopoietic stem cell transplantation	×	○	×	×	×
azacitidine	×	○	×	×	×
cytarabine	×	○	×	×	×
cytarabine + daunorubicin	×	○	×	×	×
cytarabine + daunorubicin + etoposide	×	○	×	×	×
cytarabine + etoposide + idarubicin	×	○	×	×	×

## Relevant Therapy Summary (continued)

☒ In this cancer type
 ☐ In other cancer type
 ☒ In this cancer type and other cancer types
 ☒ No evidence

### RUNX1 p.(R201Pfs\*11) c.601\_602insCC (continued)

Relevant Therapy	FDA	NCCN	EMA	ESMO	Clinical Trials*
cytarabine + fludarabine + idarubicin + filgrastim	×	○	×	×	×
cytarabine + idarubicin	×	○	×	×	×
cytarabine + mitoxantrone	×	○	×	×	×
decitabine	×	○	×	×	×
gemtuzumab ozogamicin + cytarabine + daunorubicin	×	○	×	×	×
venetoclax + azacitidine	×	○	×	×	×
venetoclax + cytarabine	×	○	×	×	×
venetoclax + decitabine	×	○	×	×	×

## Relevant Therapy Details

### Current NCCN Information

☒ In this cancer type
 ☐ In other cancer type
 ☒ In this cancer type and other cancer types

NCCN information is current as of 2022-02-28. For the most up-to-date information, search [www.nccn.org](http://www.nccn.org).  
For NCCN International Adaptations & Translations, search [www.nccn.org/global/international\\_adaptations.aspx](http://www.nccn.org/global/international_adaptations.aspx).

### TP53 p.(C277Y) c.830G>A + RUNX1 p.(R201Pfs\*11) c.601\_602insCC

#### ○ decitabine

Cancer type: Acute Myeloid Leukemia

Variant classes: RUNX1 & TP53 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

#### ○ venetoclax + azacitidine

Cancer type: Acute Myeloid Leukemia

Variant classes: RUNX1 & TP53 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**TP53 p.(C277Y) c.830G>A**☐ **cytarabine + daunorubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

☐ **cytarabine + daunorubicin + etoposide**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

☐ **cytarabine + etoposide + idarubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

☐ **cytarabine + idarubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

☐ **Allogeneic hematopoietic stem cell transplantation**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Consolidation therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**TP53 p.(C277Y) c.830G>A (continued)****○ cytarabine**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Consolidation therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + daunorubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Consolidation therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + daunorubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + idarubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + mitoxantrone**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**TP53 p.(C277Y) c.830G>A (continued)****○ gemtuzumab ozogamicin + cytarabine + daunorubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ venetoclax + cytarabine**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ venetoclax + decitabine**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ azacitidine**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 2B

Population segment (Line of therapy):

- (Induction therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + daunorubicin + etoposide**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 2B

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]



**TP53 p.(C277Y) c.830G>A (continued)****○ cytarabine + etoposide + idarubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 2B

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + fludarabine + idarubicin + filgrastim**

Cancer type: Acute Myeloid Leukemia

Variant class: TP53 mutation

NCCN Recommendation category: 2B

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**RUNX1 p.(R201Pfs\*11) c.601\_602insCC****○ cytarabine + daunorubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + daunorubicin + etoposide**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**RUNX1 p.(R201Pfs\*11) c.601\_602insCC (continued)****○ cytarabine + etoposide + idarubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + idarubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 1

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ Allogeneic hematopoietic stem cell transplantation**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Consolidation therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Consolidation therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + daunorubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Consolidation therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**RUNX1 p.(R201Pfs\*11) c.601\_602insCC (continued)****○ cytarabine + daunorubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + idarubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + mitoxantrone**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ gemtuzumab ozogamicin + cytarabine + daunorubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy); Preferred intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ venetoclax + cytarabine**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**RUNX1 p.(R201Pfs\*11) c.601\_602insCC (continued)****○ venetoclax + decitabine**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 2A

Population segment (Line of therapy):

- (Induction therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ azacitidine**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 2B

Population segment (Line of therapy):

- (Induction therapy)

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + daunorubicin + etoposide**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 2B

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + etoposide + idarubicin**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 2B

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

**○ cytarabine + fludarabine + idarubicin + filgrastim**

Cancer type: Acute Myeloid Leukemia

Variant class: RUNX1 mutation

NCCN Recommendation category: 2B

Population segment (Line of therapy):

- (Induction therapy); Other recommended intervention

Reference: NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]

## Current EMA Information

☒ In this cancer type    ☐ In other cancer type    ☐ In this cancer type and other cancer types

EMA information is current as of 2022-03-16. For the most up-to-date information, search [www.ema.europa.eu/ema](http://www.ema.europa.eu/ema).

### TP53 p.(C277Y) c.830G>A

#### ☐ idelalisib + rituximab

Cancer type: Chronic Lymphocytic Leukemia    Label as of: 2021-10-06

Variant class: TP53 mutation

Reference:

[https://www.ema.europa.eu/en/documents/product-information/zydelig-epar-product-information\\_en.pdf](https://www.ema.europa.eu/en/documents/product-information/zydelig-epar-product-information_en.pdf)

## 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 2022-02-28. For the most up-to-date information, search [www.esmo.org](http://www.esmo.org).

### TP53 p.(C277Y) c.830G>A

#### ☐ acalabrutinib

**Cancer type:** Chronic Lymphocytic Leukemia, Small Lymphocytic Lymphoma **Variant class:** TP53 mutation

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

**Population segment (Line of therapy):**

- Relapsed (Subsequent therapy)

**Reference:** ESMO Clinical Practice Guidelines - ESMO-Chronic Lymphocytic Leukaemia [Annals of Oncology (2020), <https://doi.org/10.1016/j.annonc.2020.09.019>.]

#### ☐ ibrutinib

**Cancer type:** Chronic Lymphocytic Leukemia, Small Lymphocytic Lymphoma **Variant class:** TP53 mutation

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

**Population segment (Line of therapy):**

- Relapsed (Subsequent therapy)

**Reference:** ESMO Clinical Practice Guidelines - ESMO-Chronic Lymphocytic Leukaemia [Annals of Oncology (2020), <https://doi.org/10.1016/j.annonc.2020.09.019>.]

#### ☐ rituximab + venetoclax

**Cancer type:** Chronic Lymphocytic Leukemia, Small Lymphocytic Lymphoma **Variant class:** TP53 mutation

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

**Population segment (Line of therapy):**

- Relapsed (Subsequent therapy)

**Reference:** ESMO Clinical Practice Guidelines - ESMO-Chronic Lymphocytic Leukaemia [Annals of Oncology (2020), <https://doi.org/10.1016/j.annonc.2020.09.019>.]

#### ☐ ibrutinib

**Cancer type:** Chronic Lymphocytic Leukemia, Small Lymphocytic Lymphoma **Variant class:** TP53 mutation

**ESMO Level of Evidence/Grade of Recommendation:** II / B

**Population segment (Line of therapy):**

- Relapsed (Subsequent therapy)

**Reference:** ESMO Clinical Practice Guidelines - ESMO-Chronic Lymphocytic Leukaemia [Annals of Oncology (2020), <https://doi.org/10.1016/j.annonc.2020.09.019>.]

## TP53 p.(C277Y) c.830G>A (continued)

### ○ idelalisib + rituximab

**Cancer type:** Chronic Lymphocytic Leukemia, Small **Variant class:** TP53 mutation  
Lymphocytic Lymphoma

**ESMO Level of Evidence/Grade of Recommendation:** II / B

**Population segment (Line of therapy):**

- Relapsed (Subsequent therapy)

**Reference:** ESMO Clinical Practice Guidelines - ESMO-Chronic Lymphocytic Leukaemia [Annals of Oncology (2020), <https://doi.org/10.1016/j.annonc.2020.09.019>.]

### ○ acalabrutinib

**Cancer type:** Chronic Lymphocytic Leukemia, Small **Variant class:** TP53 mutation  
Lymphocytic Lymphoma

**ESMO Level of Evidence/Grade of Recommendation:** III / A

**Population segment (Line of therapy):**

- (First-line therapy)

**Reference:** ESMO Clinical Practice Guidelines - ESMO-Chronic Lymphocytic Leukaemia [Annals of Oncology (2020), <https://doi.org/10.1016/j.annonc.2020.09.019>.]

### ○ ibrutinib

**Cancer type:** Chronic Lymphocytic Leukemia, Small **Variant class:** TP53 mutation  
Lymphocytic Lymphoma

**ESMO Level of Evidence/Grade of Recommendation:** III / A

**Population segment (Line of therapy):**

- (First-line therapy)

**Reference:** ESMO Clinical Practice Guidelines - ESMO-Chronic Lymphocytic Leukaemia [Annals of Oncology (2020), <https://doi.org/10.1016/j.annonc.2020.09.019>.]

### ○ idelalisib + rituximab

**Cancer type:** Chronic Lymphocytic Leukemia, Small **Variant class:** TP53 mutation  
Lymphocytic Lymphoma

**ESMO Level of Evidence/Grade of Recommendation:** III / A

**Population segment (Line of therapy):**

- (First-line therapy)

**Reference:** ESMO Clinical Practice Guidelines - ESMO-Chronic Lymphocytic Leukaemia [Annals of Oncology (2020), <https://doi.org/10.1016/j.annonc.2020.09.019>.]

## TP53 p.(C277Y) c.830G>A (continued)

### ○ obinutuzumab + venetoclax

**Cancer type:** Chronic Lymphocytic Leukemia, Small Lymphocytic Lymphoma  
**Variant class:** TP53 mutation

**ESMO Level of Evidence/Grade of Recommendation:** III / A

**Population segment (Line of therapy):**

- (First-line therapy)

**Reference:** ESMO Clinical Practice Guidelines - ESMO-Chronic Lymphocytic Leukaemia [Annals of Oncology (2020), <https://doi.org/10.1016/j.annonc.2020.09.019>.]

### ○ venetoclax

**Cancer type:** Chronic Lymphocytic Leukemia, Small Lymphocytic Lymphoma  
**Variant class:** TP53 mutation

**ESMO Level of Evidence/Grade of Recommendation:** III / A

**Population segment (Line of therapy):**

- (First-line therapy)

**Reference:** ESMO Clinical Practice Guidelines - ESMO-Chronic Lymphocytic Leukaemia [Annals of Oncology (2020), <https://doi.org/10.1016/j.annonc.2020.09.019>.]

### ○ venetoclax

**Cancer type:** Chronic Lymphocytic Leukemia, Small Lymphocytic Lymphoma  
**Variant class:** TP53 mutation

**ESMO Level of Evidence/Grade of Recommendation:** III / B

**Population segment (Line of therapy):**


- Relapsed (Subsequent therapy)

**Reference:** ESMO Clinical Practice Guidelines - ESMO-Chronic Lymphocytic Leukaemia [Annals of Oncology (2020), <https://doi.org/10.1016/j.annonc.2020.09.019>.]



## Alerts Informed By Public Data Sources

### Current FDA Information

 Contraindicated
  Not recommended
  Resistance
  Breakthrough
  Fast Track

FDA information is current as of 2022-03-16. For the most up-to-date information, search [www.fda.gov](http://www.fda.gov).

#### TP53 p.(C277Y) c.830G>A

##### eprenetapopt + azacitidine, eprenetapopt + venetoclax + azacitidine

**Cancer type:** Myelodysplastic Syndrome

**Variant class:** TP53 mutation

##### Supporting Statement:

The FDA has granted Breakthrough Designation to the small molecule inhibitor, eprenetapopt, in combination with azacitidine or azacitidine and venetoclax for TP53 mutant myelodysplastic syndrome (MDS).

##### Reference:

<http://vp280.alertir.com/en/pressreleases/karolinska-development%27s-portfolio-company-aprea-therapeutics-receives-fda-breakthrough-therapy-designation-1769167>

##### eprenetapopt + azacitidine, eprenetapopt + venetoclax + azacitidine

**Cancer type:** Acute Myeloid Leukemia

**Variant class:** TP53 mutation

##### Supporting Statement:

The FDA has granted Fast Track Designation to the small molecule inhibitor, eprenetapopt, in combination with azacitidine or azacitidine and venetoclax for TP53 mutant acute myeloid leukemia (AML).

##### Reference:

<https://ir.aprea.com//news-releases/news-release-details/aprea-therapeutics-receives-fda-fast-track-designation>

### Current NCCN Information

 Contraindicated
  Not recommended
  Resistance
  Breakthrough
  Fast Track

NCCN information is current as of 2022-02-28. For the most up-to-date information, search [www.nccn.org](http://www.nccn.org). For NCCN International Adaptations & Translations, search [www.nccn.org/global/international\\_adaptations.aspx](http://www.nccn.org/global/international_adaptations.aspx).

#### TP53 p.(C277Y) c.830G>A

##### chemoimmunotherapy

**Cancer type:** Chronic Lymphocytic Leukemia

**Variant class:** TP53 mutation

##### Summary:

NCCN Guidelines® include the following supporting statement(s):

- "Chemoimmunotherapy is not recommended since del(17p)/TP53 mutation is associated with low response rates."

**Reference:** NCCN Guidelines® - NCCN-Chronic Lymphocytic Leukemia/Small Lymphocytic Lymphoma [Version 2.2022]

## Current ESMO Information

 Contraindicated    Not recommended    Resistance    Breakthrough    Fast Track

ESMO information is current as of 2022-02-28. For the most up-to-date information, search [www.esmo.org](http://www.esmo.org).

### TP53 p.(C277Y) c.830G>A

#### lenalidomide

Cancer type: Myelodysplastic Syndrome

Variant class: TP53 mutation

##### Summary:

ESMO Clinical Practice Guidelines include the following supporting statement:

- "TP53 gene mutations, found in ~20% of lower-risk MDS with del(5q), confer resistance to LEN and a higher risk of AML progression."

**Reference:** ESMO Clinical Practice Guidelines - ESMO-Myelodysplastic Syndromes [Annals of Oncology (2020), <https://doi.org/10.1016/j.annonc.2020.11.002>]

## Signatures

Testing Personnel:

Laboratory Supervisor:

Pathologist:

## References

1. de et al. Runx transcription factors in the development and function of the definitive hematopoietic system. *Blood*. 2017 Apr 13;129(15):2061-2069. PMID: 28179276
2. Chuang et al. RUNX family: Regulation and diversification of roles through interacting proteins. *Int. J. Cancer*. 2013 Mar 15;132(6):1260-71. PMID: 23180629
3. Jung et al. Prognostic factor analysis in core-binding factor-positive acute myeloid leukemia. *Anticancer Res*. 2014 Feb;34(2):1037-45. PMID: 24511052
4. Sood et al. Role of RUNX1 in hematological malignancies. *Blood*. 2017 Apr 13;129(15):2070-2082. PMID: 28179279
5. Béri-Dexheimer et al. Clinical phenotype of germline RUNX1 haploinsufficiency: from point mutations to large genomic deletions. *Eur. J. Hum. Genet*. 2008 Aug;16(8):1014-8. PMID: 18478040
6. Hayashi et al. Myeloid neoplasms with germ line RUNX1 mutation. *Int. J. Hematol*. 2017 Aug;106(2):183-188. PMID: 28534116
7. De et al. RUNX1 translocations and fusion genes in malignant hemopathies. *Future Oncol*. 2011 Jan;7(1):77-91. PMID: 21174539
8. De et al. ETV6 fusion genes in hematological malignancies: a review. *Leuk. Res*. 2012 Aug;36(8):945-61. PMID: 22578774
9. Pui et al. Acute lymphoblastic leukemia. *N. Engl. J. Med*. 2004 Apr 8;350(15):1535-48. PMID: 15071128
10. NCCN Guidelines® - Acute Lymphoblastic Leukemia [Version 2.2019]. 2019 May 15
11. Huret et al. Atlas of genetics and cytogenetics in oncology and haematology in 2013. *Nucleic Acids Res*. 2013 Jan;41(Database issue):D920-4. PMID: 23161685
12. NCCN Guidelines® - NCCN-Acute Myeloid Leukemia [Version 1.2022]
13. NCCN Guidelines® - NCCN-Myelodysplastic Syndromes [Version 3.2022]
14. Weinstein et al. The Cancer Genome Atlas Pan-Cancer analysis project. *Nat. Genet*. 2013 Oct;45(10):1113-20. PMID: 24071849
15. Arber et al. The 2016 revision to the World Health Organization classification of myeloid neoplasms and acute leukemia. *Blood*. 2016 May 19;127(20):2391-405. PMID: 27069254
16. NCCN Guidelines® - NCCN-Acute Lymphoblastic Leukemia [Version 4.2021]
17. NCCN Guidelines® - NCCN-Systemic Mastocytosis [Version 1.2020]
18. Muller et al. Mutant p53 in cancer: new functions and therapeutic opportunities. *Cancer Cell*. 2014 Mar 17;25(3):304-17. PMID: 24651012
19. Olivier et al. TP53 mutations in human cancers: origins, consequences, and clinical use. *Cold Spring Harb Perspect Biol*. 2010 Jan;2(1):a001008. PMID: 20182602
20. Guha et al. Inherited TP53 Mutations and the Li-Fraumeni Syndrome. *Cold Spring Harb Perspect Med*. 2017 Apr 3;7(4). PMID: 28270529
21. Cerami et al. The cBio cancer genomics portal: an open platform for exploring multidimensional cancer genomics data. *Cancer Discov*. 2012 May;2(5):401-4. PMID: 22588877
22. Peter et al. Comprehensive genomic characterization of squamous cell lung cancers. *Nature*. 2012 Sep 27;489(7417):519-25. PMID: 22960745
23. Cancer Genome Atlas Network. Comprehensive genomic characterization of head and neck squamous cell carcinomas. *Nature*. 2015 Jan 29;517(7536):576-82. PMID: 25631445
24. Campbell et al. Distinct patterns of somatic genome alterations in lung adenocarcinomas and squamous cell carcinomas. *Nat. Genet*. 2016 Jun;48(6):607-16. PMID: 27158780
25. Cancer Genome Atlas Research Network. Integrated genomic characterization of oesophageal carcinoma. *Nature*. 2017 Jan 12;541(7636):169-175. doi: 10.1038/nature20805. Epub 2017 Jan 4. PMID: 28052061
26. Olivier et al. The IARC TP53 database: new online mutation analysis and recommendations to users. *Hum. Mutat*. 2002 Jun;19(6):607-14. PMID: 12007217
27. Rivlin et al. Mutations in the p53 Tumor Suppressor Gene: Important Milestones at the Various Steps of Tumorigenesis. *Genes Cancer*. 2011 Apr;2(4):466-74. PMID: 21779514
28. Petitjean et al. TP53 mutations in human cancers: functional selection and impact on cancer prognosis and outcomes. *Oncogene*. 2007 Apr 2;26(15):2157-65. PMID: 17401424
29. Soussi et al. Recommendations for analyzing and reporting TP53 gene variants in the high-throughput sequencing era. *Hum. Mutat*. 2014 Jun;35(6):766-78. PMID: 24729566
30. <https://www.globenewswire.com/news-release/2020/10/13/2107498/0/en/PMV-Pharma-Granted-FDA-Fast-Track-Designation-of-PC14586-for-the-Treatment-of-Advanced-Cancer-Patients-that-have-Tumors-with-a-p53-Y220C-Mutation.html>

## References (continued)

31. <https://ir.aprea.com//news-releases/news-release-details/aprea-therapeutics-receives-fda-fast-track-designation>
32. <http://vp280.alertir.com/en/pressreleases/karolinska-development%27s-portfolio-company-aprea-therapeutics-receives-fda-breakthrough-therapy-designation-1769167>
33. Parrales et al. Targeting Oncogenic Mutant p53 for Cancer Therapy. *Front Oncol.* 2015 Dec 21;5:288. doi: 10.3389/fonc.2015.00288. eCollection 2015. PMID: 26732534
34. Zhao et al. Molecularly targeted therapies for p53-mutant cancers. *Cell. Mol. Life Sci.* 2017 Nov;74(22):4171-4187. PMID: 28643165
35. NCCN Guidelines® - NCCN-Myeloproliferative Neoplasms [Version 2.2021]
36. NCCN Guidelines® - NCCN-Chronic Lymphocytic Leukemia/Small Lymphocytic Lymphoma [Version 2.2022]
37. NCCN Guidelines® - NCCN-B-Cell Lymphomas [Version 5.2021]
38. Bernard et al. Implications of TP53 allelic state for genome stability, clinical presentation and outcomes in myelodysplastic syndromes. *Nat. Med.* 2020 Aug 3. PMID: 32747829