Recent Advances in Acute Leukemias and Myelodysplastic Syndromes



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Learning Objectives

- Review new classification systems and recently approved drugs for lower risk myelodysplastic syndromes
- Discuss treatment of NPM1-mutated AML
- Highlight shifting paradigms in the upfront treatment of B-ALL

Myelodysplastic Syndromes

(Myelodysplastic Neoplasms?)

MDS in the WHO5 Classification System

Table 3. Classification and defining features of myelodysplastic neoplasms (MDS).

| | Blasts | Cytogenetics | Mutations |
|---|---------------------------------------|---|---|
| MDS with defining genetic abnormalities | | | |
| MDS with low blasts and isolated 5q deletion (MDS-5q) | <5% BM and <2% PB | 5q deletion alone, or with 1 other abnormality other than monosomy 7 or 7q deletion | |
| MDS with low blasts and SF3B1 mutation ^a (MDS-SF3B1) | | Absence of 5q deletion, monosomy 7, or complex karyotype | SF3B1 |
| MDS with biallelic <i>TP53</i> inactivation (MDS-bi <i>TP53</i>) | <20% BM and PB | Usually complex | Two or more <i>TP53</i> mutations, or 1 mutation with evidence of <i>TP53</i> copy number loss or cnLOH |
| MDS, morphologically defined | | | |
| MDS with low blasts (MDS-LB) | <5% BM and <2% PB | | |
| MDS, hypoplastic ^b (MDS-h) | | | |
| MDS with increased blasts (MDS-IB) | | | |
| MDS-IB1 | 5-9% BM or 2-4% PB | | |
| MDS-IB2 | 10-19% BM or 5–19% PB or Auer rods | | |
| MDS with fibrosis (MDS-f) | 5–19% BM; 2–19% PB | | |
| | | | |

^aDetection of \geq 15% ring sideroblasts may substitute for *SF3B1* mutation. Acceptable related terminology: MDS with low blasts and ring sideroblasts.

^bBy definition, ≤25% bone marrow cellularity, age adjusted.

BM bone marrow, PB peripheral blood, cnLOH copy neutral loss of heterozygosity.

MDS in the ICC Classification System

Table 20. Myelodysplastic syndromes (MDS) and myelodysplastic syndrome/acute myeloid leukemia (MDS/AML)

| | Dysplastic lineages | Cytopenias | Cytoses* | BM and PB Blasts | Cytogenetics ^b *** | Mutations |
|--|------------------------|------------|------------------------|-------------------------------|--|---|
| MDS with mutated SF3B1 (MDS- SF3B1) | Typically >1° | ≥1 | 0 | <5% BM <2% PB | Any, except isolated del(5q), - 7/del(7q), abn3q26.2, or complex | SF3B1 (≥10% VAF), without multi-hit TP53, or RUNX1 |
| MDS with del(5q) [MDS- del(5q)] | Typically >1° | ≥1 | Thrombocytosis allowed | <5% BM <2% PB ^d | del(5q), with up to 1 additional, except -7/del(7q) | Any, except multi- hit <i>TP53</i> |
| MDS, NOS - without dysplasia | 0 | ≥1 | 0 | <5% BM <2% PB ^d | -7/del(7q) or complex | Any, except multi- hit <i>TP53</i> or <i>SF3B1</i> (≥10% VAF) |
| MDS, NOS - with single lineage dysplasia | 1 | ≥1 | 0 | <5% BM <2% PB ^d | Any, except not meeting criteria for MDS-del(5q) | Any, except multi- hit <i>TP53</i> ;not meeting criteria for MDS- <i>SF3B1</i> |
| MDS, NOS - with multilineage dysplasia | ≥2 | ≥1 | 0 | <5% BM <2% PB ^d | Any, except not meeting criteria for MDS-del(5q) | Any, except multi- hit <i>TP53</i> ,; not meeting criteria for MDS- <i>SF3B1</i> |

| MDS with excess blasts (MDS-EB) | Typically ≥1 ^c | ≥1 | 0 | 5-9% BM, 2-9% PB ^d | Any | Any, except multi- hit <i>TP53</i> |
|---------------------------------------|------------------------------|----|---|----------------------------------|---|--|
| MDS/AML | Typically ≥1 ^c | ≥1 | 0 | 10-19% BM or PB ^e | Any, except AML- defining ^f | Any, except NPM1, bZIP CEBPA or TP53 |

 a Cytoses: Sustained white blood count ≥13 x 10 9 /L, monocytosis (≥0.5 x 10 9 /L and ≥10% of leukocytes), or platelets ≥450 x 10 9 /L; thrombocytosis is allowed in MDS-del(5q) or in any MDS case with inv(3) or t(3;3) cytogenetic abnormality.

^bBCR::ABL1 rearrangement or any of the rearrangements associated with myeloid/lymphoid neoplasms with eosinophilia and tyrosine kinase gene fusions exclude a diagnosis of MDS, even in the context of cytopenia.

^cAlthough dysplasia is typically present in these entities, it is not required.

^dAlthough 2% PB blasts mandates classification of an MDS case as MDS-EB, the presence of 1% PB blasts confimed on two separate occasions also qualifies for MDS-EB.

For pediatric patients (<18 years), the blast thresholds for MDS-EB are 5-19% in BM and 2-19% in PB, and the entity MDS/AML does not apply.

^fAML-defining cytogenetics are listed in the AML section.

| Туре | Cytopenia | Blasts | Genetics |
|----------------------------------|-----------------|---|---|
| MDS with mutated TP53 | Any | 0-9% bone marrow and blood blasts | Multi-hit TP53 mutation ^a , or <i>TP53</i> mutation (VAF >10%) and complex karyotype often with loss of 17p ^b |
| MDS/AML with mutated <i>TP53</i> | Any | 10-19% bone marrow or blood blasts | Any somatic <i>TP53</i> mutation (VAF >10%) |
| AML with mutated <i>TP53</i> | Not required | ≥20% bone marrow or blood blasts or meets criteria for pure erythroid leukemia | Any somatic <i>TP53</i> mutation (VAF >10%) |

^aDefined as two distinct *TP53* mutations (each VAF >10%) OR a single *TP53* mutation with either 1) 17p deletion on cytogenetics; 2) VAF of >50%; or 3) Copy-neutral loss of heterozygosity (LOH) at the 17p *TP53* locus.

^bIf TP53 locus LOH information is not available

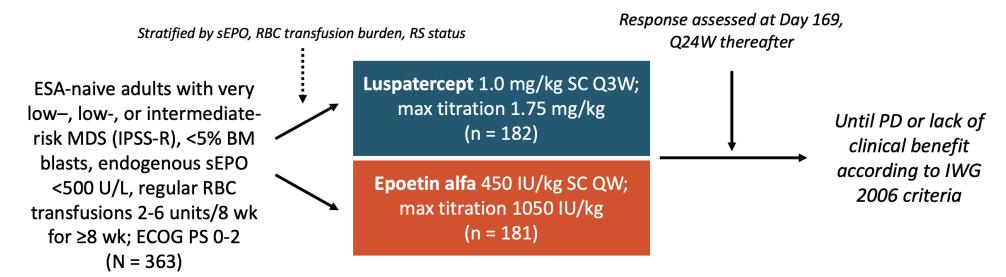
IPSS-M



COMMANDS: Study Design

<u>Luspatercept</u> is a first-in-class erythroid maturation agent

Global, open-label, randomized phase III trial; data cutoff March 31, 2023



- Primary endpoint: RBC-TI ≥12 wk with concurrent mean Hb increase ≥1.5 g/dL, Wk 1-24
- Key secondary endpoints:
 - HI-E ≥8 wk per IWG criteria; RBC-TI ≥12 wk and at 24 wk; RBC-TI ≥24 wk, Wk 1-48 (exploratory); safety



COMMANDS: Baseline Characteristics

| Characteristic | Luspatercept (n = 182) | Epoetin alfa (n = 181) |
|---|------------------------------------|------------------------------------|
| Median age, yr (range) | 74 (46 to 93) | 74 (31 to 91) |
| Female sex, n (%) | 73 (40.1) | 89 (49.2) |
| Median time since diagnosis, mo (range) | 7.97 (-0.4 to 243.1) | 5.13 (-0.3 to 171.6) |
| ECOG PS, n (%) ■ 0 ■ 1 ■ 2 | 74 (40.7) 104 (57.1) 4 (2.2) | 69 (38.1) 94 (51.9) 18 (9.9) |
| Median RBC TB, U/8 wk (range) | 3 (1-10) | 3 (0-14) |
| RBC TB, n (%) ■ <4 U/8 wk ■ ≥4 U/8 wk | 118 (64.8) 64 (35.2) | 111 (61.3) 70 (38.7) |
| Median Hb, g/dL (range) | 7.8 (4.7 to 9.2) | 7.8 (4.5 to 10.2) |

| Characteristic, n (%) | Luspatercept (n = 182) | Epoetin alfa (n = 181) |
|--|------------------------------------|------------------------------------|
| sEPO ■ ≤200 U/L ■ >200 to < 500 U/L | 145 (79.7) 37 (20.3) | 144 (79.6) 37 (20.4) |
| SF3B1 mutationMutatedWild typeMissing | 114 (62.6) 65 (35.7) 3 (1.6) | 101 (55.8) 72 (39.8) 8 (4.4) |
| RS RS+ RS- Missing | 133 (73.1) 49 (26.9) 0 | 130 (71.8) 50 (27.6) 1 (0.6) |

COMMANDS: Efficacy

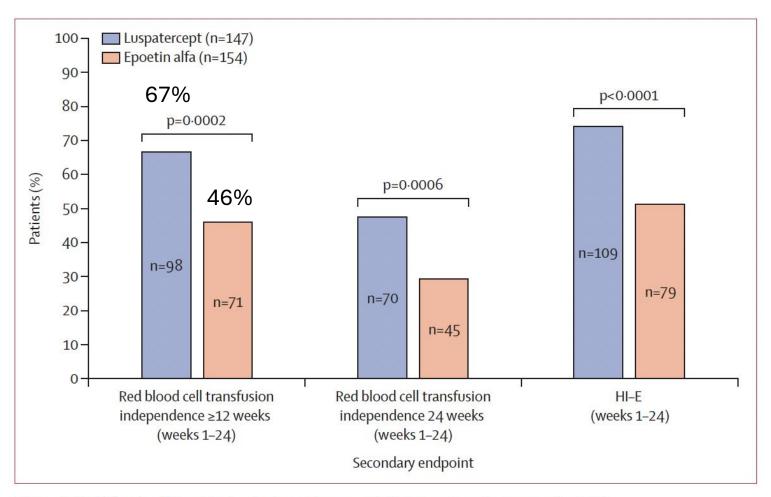


Figure 2: Red blood cell transfusion independence and HI–E response during weeks 1–24

Only patients who received their first dose of treatment at least 24 weeks (169 days) before the data cutoff (Aug 31, 2022), including those who discontinued treatment, were included in the analysis. HI-E=haematological improvement-erythroid.

Primary Endpoint:

RBC TI for at least 12 weeks and increase in Hgb at least 1.5g/dL 59% vs 31%, p<0.0001

Among 12-Wk Tl responders: Median DoR 127 wks

COMMANDS: Adverse Events

| | Luspaterce | pt (n=178) | Epoetin alf | a (n=176) | | | | |
|---|---------------|------------|-------------|-----------|--|--|--|--|
| | Any grade | Grade 3-4 | Any grade | Grade 3-4 | | | | |
| General disorder or administration site conditions | | | | | | | | |
| Fatigue | 26 (15%) | 1 (1%) | 12 (7%) | 1 (1%) | | | | |
| Peripheral oedema | 23 (13%) | 0 | 12 (7%) | 0 | | | | |
| Asthenia | 22 (12%) | 0 | 25 (14%) | 1 (1%) | | | | |
| Infections and infes | tations | | | | | | | |
| COVID-19 | 19 (11%) | 6 (3%) | 17 (10%) | 2 (1%) | | | | |
| Gastrointestinal disorders | | | | | | | | |
| Diarrhoea | 26 (15%) | 2 (1%) | 20 (11%) | 1 (1%) | | | | |
| Nausea | 21 (12%) | 0 | 13 (7%) | 0 | | | | |
| Respiratory, thoracic, or mediastinal disorders | | | | | | | | |
| Dyspnoea | 21 (12%) | 7 (4%) | 13 (7%) | 2 (1%) | | | | |
| Vascular disorders | | | | | | | | |
| Hypertension | 23 (13%) | 15 (8%) | 12 (7%) | 8 (5%) | | | | |
| Blood and lymphati | c system diso | orders | | | | | | |
| Anaemia | 17 (10%) | 13 (7%) | 17 (10%) | 12 (7%) | | | | |
| Data are n (%), where n=number of patients. Events of grade 1–4 severity (Common Terminology Criteria for Adverse Events version 4.03) occurring in at east 10% of patients in either group are shown. System organ classes and oreferred terms were coded with the Medical Dictionary for Regulatory Activities (version 25.0). Treatment-emergent adverse events were defined as adverse events that started on or after the first treatment of study medication until 42 days after the last dose of any study drug. A patient was counted only once for the maximum severity for multiple events under the same preferred term within system organ class. | | | | | | | | |

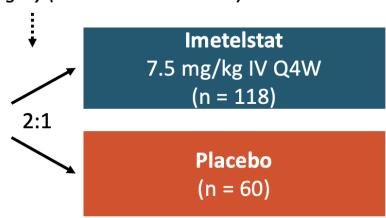
IMerge: Study Design

International, double-blind, randomized phase III trial

Imetelstat is a first-inclass competitive telomerase inhibitor

Stratified by transfusion burden (4-6 vs >6 U) and IPSS-R category (low vs intermediate 1)

Patients with low-risk or intermediate 1–risk MDS (IPSS-R); R/R to ESA or EPO >500 mU/mL (ESA ineligible); RBC transfusion dependent (≥4 U/8 wk over 16 wk prestudy); non-del(5q); no prior lenalidomide or HMAs (N = 178)



- Primary endpoint: 8-wk RBC-TI
- **Key secondary endpoints:** 24-wk RBC-TI, TI duration, HI-E, safety
- Key exploratory endpoints: changes in VAF, PRO (FACIT-Fatigue)



IMerge: Baseline Characteristics

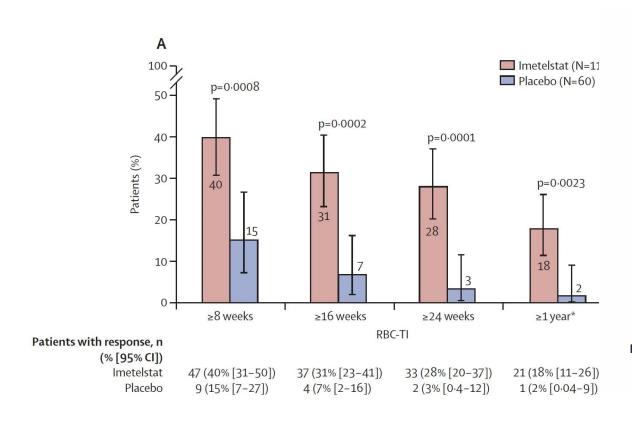
| | Imetelstat (n = 118) | Placebo (n = 60) |
|--|-------------------------|---------------------|
| Median age, yr (range) | 72 (44-87) | 73 (39-85) |
| Male, n (%) | 71 (60) | 40 (67) |
| Median time since MDS diagnosis, yr (range) | 3.5 (0.1-26.7) | 2.8 (0.2-25.7) |
| WHO classification, n (%) RS- RS+ | 44 (37) 73 (62) | 23 (38) 37 (62) |
| Baseline IPSS-R risk, n (%) ■ Low ■ Intermediate-1 | 80 (68) 38 (32) | 39 (65) 21 (35) |

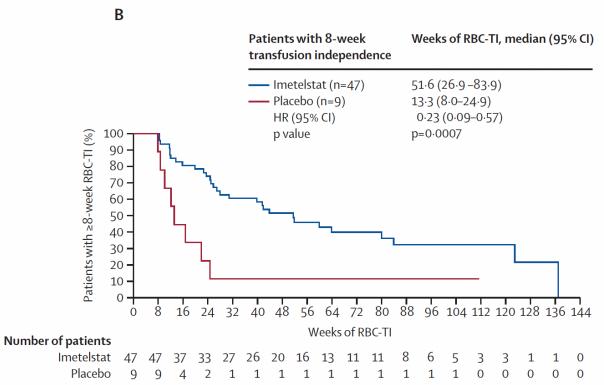
| Imetelstat (n = 118) | Placebo (n = 60) |
|-------------------------|---|
| 7.9 (5.3-10.1) | 7.8 (6.1-9.2) |
| 6 (4-33) | 6 (4-13) |
| 62 (53) 56 (48) | 33 (55) 27 (45) |
| 174.9 (6.0-4460.0) | 277.0 (16.9-5514.0) |
| 87 (74) 26 (22) | 36 (60) 22 (37) |
| 108 (92) | 52 (87) |
| 7 (6) | 4 (7) |
| | (n = 118) 7.9 (5.3-10.1) 6 (4-33) 62 (53) 56 (48) 174.9 (6.0-4460.0) 87 (74) 26 (22) 108 (92) |

^{*}Data missing for 5 patients in imetelstat group and 2 in placebo group.



IMerge: Efficacy and Duration of Response





Among 8-Wk TI responders: Hgb rise median 3.6 vs 0.8 Hgb peak median 11.3 vs 8.9

IMerge: Adverse Events

| | Imetelstat (| N=118) | Placebo (N=59) | |
|--|--------------|-----------|----------------|-----------|
| | Any grade | Grade 3–4 | Any grade | Grade 3-4 |
| Haematological | | | | |
| Thrombocytopenia | 89 (75%) | 73 (62%) | 6 (10%) | 5 (8%) |
| Neutropenia | 87 (74%) | 80 (68%) | 4 (7%) | 2 (3%) |
| Anaemia | 24 (20%) | 23 (19%) | 6 (10%) | 4 (7%) |
| Leukopenia | 12 (10%) | 9 (8%) | 1 (2%) | 0 |
| General disorders and administration site conditions | | | | |
| Asthenia | 22 (19%) | 0 | 8 (14%) | 0 |
| Oedema peripheral | 13 (11%) | 0 | 8 (14%) | 0 |
| Pyrexia | 9 (8%) | 2 (2%) | 7 (12%) | 0 |
| COVID-19 | 22 (19%)† | 3 (3%)‡ | 8 (14%)† | 3 (5%)‡ |
| Gastrointestinal disorders | | | | |
| Diarrhoea | 14 (12%) | 1 (1%) | 7 (12%) | 1 (2%) |
| Constipation | 9 (8%) | 0 | 7 (12%) | 0 |
| Headache | 15 (13%) | 1 (1%) | 3 (5%) | 0 |
| Alanine aminotransferase increased | 14 (12%) | 3 (3%) | 4 (7%) | 2 (3%) |
| Hyperbilirubinaemia | 11 (9%) | 1 (1%) | 6 (10%) | 1 (2%) |

^{*}Includes all patients who received at least one dose of study drug. †Includes COVID-19, asymptomatic COVID-19, and COVID-19 pneumonia. ‡Only COVID-19 pneumonia events were classified as grade 3–4 events for COVID-19.

Table 3: Number of patients with treatment-emergent adverse events occurring in at least 10% of patients in the safety population*

- Median duration of G3/4 thrombocytopenia and neutropenia was <2wks and >80% resolved to G2 or better within 4wks
- Occurred most often during C1-3
- Infection and bleeding similar between arms
- 49% required dose reductions due to AEs and 8% discontinued treatment due to AEs of neutropenia or thrombocytopenia

Recent FDA approvals in MDS:

- *Luspatercept* is approved for:
 - Anemia without previous ESA in adults with very low to intermediate risk MDS who may require RBC transfusions
 - Anemia failing an ESA and requiring 2 or more RBC units over 8 weeks in adults with very low to intermediate risk MDS-RS or MDS/MPN-RS-T
- *Imetelstat* is approved for treatment of adult patients with low-to intermediate-1 risk MDS with transfusion-dependent anemia requiring 4 or more RBC units over 8 weeks who have not responded to or have lost response to or are ineligible for ESA.

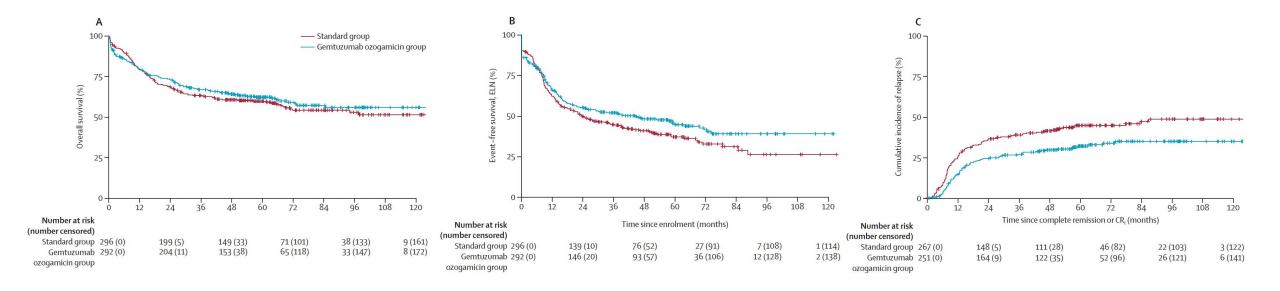
Acute Myeloid Leukemia

AML with NPM1Mutation

- WHO5 Acute myeloid leukemia with NPM1 mutation
 - Can be diagnosed irrespective of blast count
- ICC 2022 Acute myeloid leukemia with mutated NPM1
 - Can be diagnosed with 10% or more blasts
- Favorable risk by ELN 2022, with caveats
 - Favorable risk in the context of intensively treated patients and in younger patients
 - Adverse risk if it has concurrent adverse risk cytogenetic abnormalities
 - Intermediate risk if is has concurrent FLT3-ITD mutation

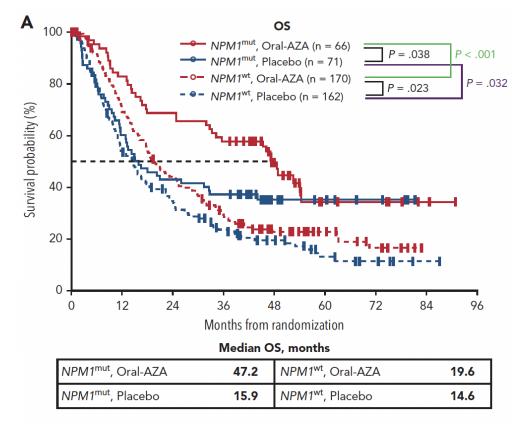
GO for AML with NPM1 mutation

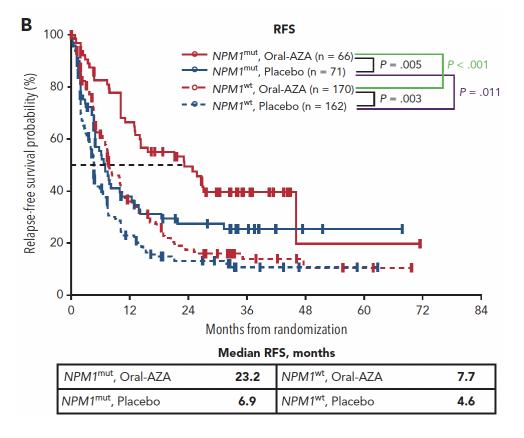
AMLSG 09-09 Study – Added a dose of GO 3mg/m2 to 2 cycles of 7+3 based induction and the first of 3 cycles of HiDAC No OS or EFS advantage but addition of GO significantly decreased cumulative incidence of relapse



Oral Azacitidine for AML with NPM1 mutation

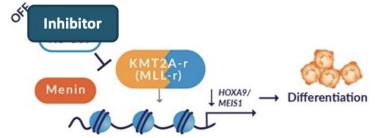
- QUAZAR phase 3 trial led to the approval of oral Aza for maintenance for patients with AML in remission
- Post hoc analysis of NPM1-mutated patients on the QUAZAR trial showed significant improvements in OS and RFS vs placebo





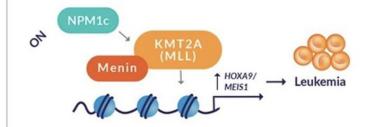
Menin Inh for AML with NPM1 mutation

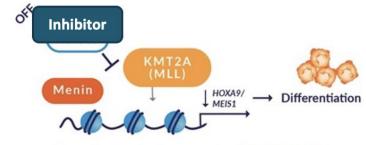
KMT2A-r (MLL-r) Menin HOXA9/ MEIS1 Leukemia



Targeting the menin-KMT2A(MLL) interaction to reverse epigenetic dysregulation in MLL-rearranged AML

NPM1 Mutant AML





A central role for menin-KMT2A(MLL) interaction in epigenetic dysregulation in NPM1-mutant AML

Kühn. Cancer Discov. 2016;6:1166. Thorsteinsdottir. Mol Cell Biol. 2001;21:224. Patel. Curr Hematol Malig Rep. 2020;15:350. Brunetti. Cancer Cell. 2018;34:499.



Menin Inhibitors in Clinical Development

| Trial Name (NCT #) | Agent (Route) | Phase I/II Expansion Cohorts for R/R Disease | Phase/# Patients | Current Trial Status |
|-----------------------------------|--|--|-------------------------|---|
| AUGMENT-101 (NCT04065399) | Revumenib (SNDX-5613) PO BID | (a)ALL or MPAL with <i>KMT2Ar</i> (b)AML with <i>KMT2Ar; (c) NPM1</i> | Phase I/II (n = 413) | In Ph II expansion FDA breakthrough NDA pending |
| KOMET-001 (NCT04067336) | Ziftomenib (KO-539) PO QD | (a)AML with KMT2Ar (b)AML with NPM1c | Phase I/II (n = 199) | In Ph II expansion (52 sites) Actively recruiting |
| CR108998 (NCT04811560) | JNJ-75276617 PO QD | (a)AML/ALL with <i>KMT2Ar</i> (b)AML with <i>NPM1c</i> | Phase I/II (n = 150) | Recruiting (24 sites) |
| DSP-5336-101 (NCT04988555) | DSP-5336 PO QD | RR-AML/R/R-ALL Ph II : <i>NPM1/KMT2Ar</i> | Phase I/II (n = 70) | Recruiting (11 sites) |
| COVALENT 101 (NCT05153330) | BMF-219 PO | (a) AML/ ALL (KMT2Ar, <i>NPM1</i>) (b)DLBCL; (c) MM; (d) CLL/SLL | Phase I (n = 177) | Multiple cohorts Actively enrolling |

Open at UCD:

COVALENT 101

and

ClinicalTrials.gov.

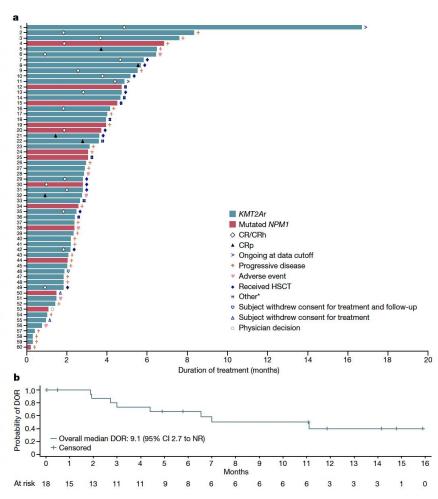
CTEP P1 trial of revumenib plus 7+3 for NPM1-mutated AML (OSU: Dr. Alice Mims)

Slide credit: clinicaloptions.com

AUGMENT-101 - Revumenib

Table 2 | Responses to treatment

| Response | Efficacy population (n=60) | KMT2Ar (n=46) | Mutated NPM1 (n=14) |
|---|----------------------------|----------------|------------------------|
| Overall response* | 32 (53%) | 27 (59%) | 5 (36%) |
| Median time to first morphologic response (range), months | 0.95 (0.9–3.7) | 0.95 (0.9–3.7) | 0.99 (1.0–1.9) |
| Best response* | | | |
| CR/CRh | 18 (30%) | 15 (33%) | 3 (21%) |
| CR | 12 (20%) | 9 (20%) | 3 (21%) |
| CRh | 6 (10%) | 6 (13%) | 0 |
| Median time to CR or CRh (range), months | 1.9 (0.9–4.9) | 2.0 (0.9-4.9) | 1.9 (1.0-1.9) |
| CRi | 0 | 0 | 0 |
| CRp | 5 (8%) | 5 (11%) | 0 |
| MLFS | 9 (15%) | 7 (15%) | 2 (14%) |
| Partial remission | 0 | 0 | 0 |
| No response | 19 (32%) | 12 (26%) | 7 (50) |
| Progressive disease | 7 (12%) | 6 (13%) | 1 (7%) |
| Missing | 2 (3%) | 1(2%) | 1 (7%) |
| MRD [†] neg. rate within CR/CRh | 14/18 (78%) | 11/15 (73%) | 3/3 (100%) |
| Median time to MRD [†] neg. among patients with CR/CRh (range), months | 1.9 (0.9–4.9) | 1.9 (0.9–4.9) | 1.9 (1.0-2.8) |



$\label{thm:continuous} \textbf{Table 1} | \textbf{Any-grade treatment-related and TEAEs, regardless} \\ \textbf{of causality}$

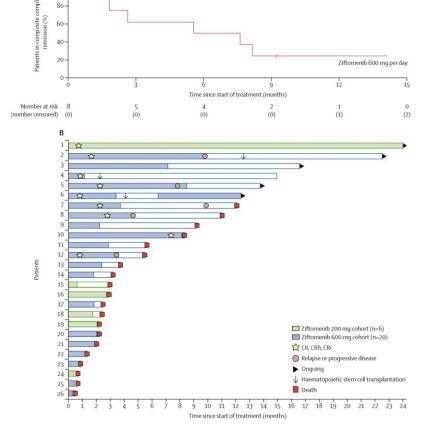
| Event | Overall population $(n=68)$ |
|------------------------------|-----------------------------|
| Any-grade TRAE (5% or over) | 53 (77.9%) |
| ECG QT prolonged | 36 (52.9%) |
| Nausea | 18 (26.5%) |
| Differentiation syndrome | 11 (16.2%) |
| Vomiting | 11 (16.2%) |
| Diarrhoea | 7 (10.3%) |
| Decreased appetite | 5 (7.4%) |
| Dysgeusia | 5 (7.4%) |
| Any-grade TEAE (20% or over) | 67 (98.5%) |
| ECG QT prolonged | 38 (55.9%) |
| Nausea | 34 (50.0%) |
| Vomiting | 27 (39.7%) |
| Febrile neutropenia | 21 (30.9%) |
| Diarrhoea | 20 (29.4%) |
| Fatigue | 18 (26.5%) |
| ALT increased | 17 (25.0%) |
| Headache | 16 (23.5%) |
| Hyperphosphataemia | 16 (23.5%) |
| Hypokalaemia | 15 (22.1%) |
| Hyponatraemia | 15 (22.1%) |
| Thrombocytopenia | 15 (22.1%) |
| Epistaxis | 14 (20.6%) |
| | 14 (20.6%) |

KOMET-001 - Ziftomenib

100

| | Patients with NPM1 mutation or KMT2A rearrangement with 600 mg ziftomenib (n=38) | Patients with NPM1 mutation with 200 mg ziftomenib (n=6) | Patients with NPM1 mutation with 600 mg ziftomenib (n=20) | Patients with NPM1 mutation with 200 mg of 600 mg ziftomenib (n=26) |
|--|--|--|---|--|
| CR | 7 (18%; 7·7-34·3) | 1 (17%; 0·4–64·1) | 7 (35%; 15·4–59·2) | 8 (31%; 14·3–51·8) |
| CR/CRh | 9 (24%; 11·4-40·2) | 1 (17%; 0·4-64·1) | 7 (35%; 15·4–59·2) | 8 (31%; 14·3-51·8) |
| CRc* | 11 (29%; 15·4-45·9) | 1 (17%; 0·4-64·1) | 8 (40%; 19·1–63·9) | 9 (35%; 17·2–55·7) |
| Overall response† | 12 (32%; 17·5-48·7) | 2 (33%; 4·3-77·7) | 9 (45%; 23·1-68·5) | 11 (42%; 23·4-63·1) |
| MRD negative CR/CRh‡ | 6/9 (67%; 29·9–92·5) | 1/1 (100%; 2·5–100) | 4/7 (57%;§ 18·4–90·1) | 5/8 (63%; 24·5–91·5) |
| CR/CRh in patients with previous venetoclax therapy | 4/26 (15%; 4·4-34·9) | | 2/13 (15%; 1·9–45·4) | |
| CR/CRh in patients with no previous venetoclax therapy | 5/12 (42%; 15·2–72·3) | | 5/7 (71%; 29·0–96·3) | |
| Median duration of CRc (95% CI) | 5.6 (1.2-8.2) | 32·6¶ | 6.6 (1.0-NE) | 7·7 (1·0-NE) |
| Median duration of CRc censored at HSCT (95% CI) | 3.1 (1.2-7.7) | | 5·6 (1·0-NE) | 5.6 (1.0-8.2) |
| Median duration of CR/CRh (95% CI) | 3.1 (0.6–8.2) | 32·6¶ | 5.6 (0.6–8.2) | 6-6 (0-6-NE) |
| Median duration of CR/CRh censored at HSCT (95% CI) | 3.1 (0.6–7.7) | | 5·6 (0·6–NE) | 5.6 (0.6–8.2) |
| Median overall survival, months (95% CI) | 5.7 (3.2-8.4) | 2·7 (0·7-NE) | 5.6 (2.1–12.1) | 3.5 (2.3-9.3) |
| Transfusion independence rate for patients who we | ere independent at b | aseline | | |
| Platelet | 2/8 (25%) | 1/3 (33%) | 2/6 (33%) | 3/9 (33%) |
| RBC | 1/3 (33%) | 0/1 | 1/3 (33%) | 1/4 (25%) |
| Platelet or RBC | 1/3 (33%) | 0/1 | 1/3 (33%) | 1/4 (25%) |
| Transfusion independence rate for patients who we | ere dependent at bas | seline | | |
| Platelet | 7/30 (23%) | 0/3 | 4/14 (29%) | 4/17 (24%) |
| RBC | 10/35 (29%) | 1/5 (20%) | 6/17 (35%) | 7/22 (32%) |
| Platelet or RBC | 7/35 (20%) | 1/5 (20%) | 5/17 (29%) | 6/22 (27%) |





HIT vs LIT?

High intensity therapy vs low intensity therapy in AML – which is better and which groups? PARADIGM trial – randomized phase 2 trial comparing Aza-Ven to IC for newly diagnosed fit adults with AML

UCD Retrospective Study

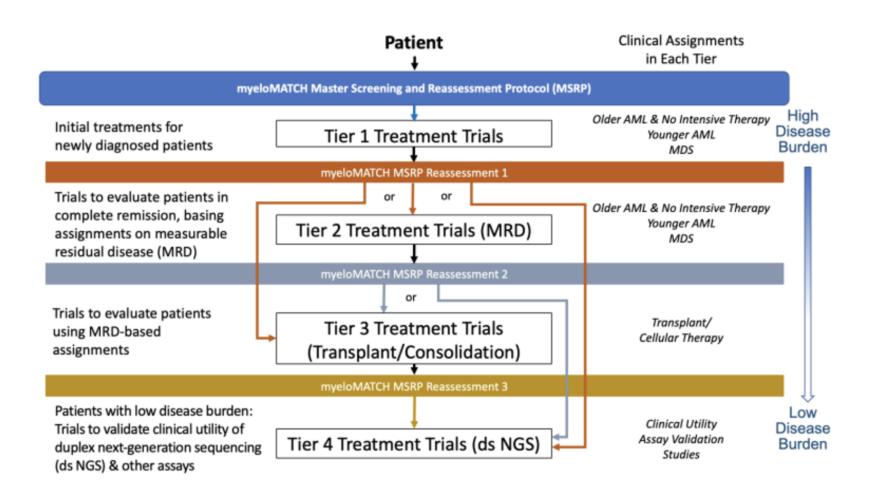
IM Resident: Margaret Krackeler

| Table 1. Patient demographics | HMA-Ven | 7+3 | CPX-351 |
|-------------------------------------|-------------------|-------------------|---------------|
| | (n = 49) | (n = 60) | (n = 7) |
| Age | | | |
| Median (range) – yr | 72 (64-90) | 56 (20-79) | 60 (48-68) |
| ≥75 yr - no. (%) | 16 (33) | 1 (2) | 0 (0) |
| AML type - no. (%) | | | |
| De novo | 37 (76) | 56 (93) | 2 (29) |
| Secondary | 12 (24) | 4 (7) | 5 (71) |
| Secondary AML - no./total no. (%) | | | • |
| Hx of MDS or CMML | 8/12 (67) | 4/4 (100) | 3/5 (60) |
| Therapy-related AML | 4/12 (33) | 0/4 (0) | 2/5 (40) |
| ECOG performance-status - no. (%) | | | |
| 0-1 | 42 (86) | 58 (97) | 7 (100) |
| 2-3 | 7 (14) | 2 (3) | 0 (0) |
| Bone marrow blast count - no. (%) | | | |
| <30% | 14 (29) | 7 (12) | 2 (29) |
| ≥30-50% | 14 (29) | 14 (29) | 2 (29) |
| ≥50% | 21 (43) | 39 (65) | 3 (43) |
| Cytogenetic risk category - no. (%) | | 740.50 | 1000 |
| Intermediate | 31 (63) | 33 (55) | 3 (43) |
| Normal karyotype - no. | 30 | 27 | 3 |
| Trisomy 8; +8 alone - no. | 1 | 6 | 0 |
| Poor | 10 (20) | 10 (17) | 4 (57) |
| 7 or 7g deletion - no. | 6 | 4 | 1 |
| 5 or 5g deletion - no. | 3 | 0 | 0 |
| Complex, >3 clonal abnormal - no. | 2 | 6 | 3 |
| Somatic mutations - no. (%) | | | |
| IDH1 or IDH2 | 17 (35) | 15 (25) | 1 (14) |
| FLT3-ITD or TKD | 4 (8) | 19 (32) | 1 (14) |
| NPM1 | 4 (8) | 17 (28) | 2 (29) |
| TP53 | 4 (8) | 1 (2) | 0 (0) |
| Baseline cytopenia grade ≥ 3 | | | |
| Anemia - no. (%) | 29 (59) | 24 (40) | 2 (29) |
| Neutropenia - no. (%) | 32 (65) | 19 (32) | 4 (57) |
| Thrombocytopenia - no. (%) | 23 (47) | 30 (50) | 3 (43) |
| Only reached CRi | 13 (27) | 0 (0) | 0 (0) |

| Table 2. Primary and Secondary HRU Outcomes | | HMA-Ven (n = 49) | 7+3/CPX-351 (n = 67) | p-value |
|---|--|--|--|----------------------------|
| HRU to CR/CRi median (range) | pRBC units PIt units Hospital Days | 4.0 (0-20) 2.0 (0-19) 9.0 (0-45) | 9.0 (4-24) 9.0 (3-30) 33 (23-57) | <0.001 <0.001 <0.001 |
| HRU 60 days post-CR/CRi median (range) | pRBC units Plt units Hospital Days | 0.0 (0-8) 0.0 (0-12) 0.0 (0-34) | 2.0 (0-11) 2.0 (0-23) 10 (0-44) | 0.01 <0.001 <0.001 |
| HRU 90 days post-CR/CRi median (range) | pRBC units PIt units Hospital Days | 0.0 (0-15) 0.0 (0-12) 0.0 (0-41) | 3.0 (0-17) 4.0 (0-44) 16 (0-64) | <0.001 <0.001 <0.001 |
| Time to count recovery days (range) | ANC > 1000 Plt > 50 | 41 (3-284) 24 (1-168) | 27 (19-67) 23 (17-92) | <0.001 0.94 |
| Time to CR/CRi days (range) | | 31 (21-284) | 34 (23-190) | 0.026 |
| Duration of remission months (range) | | 11.9 (1.1-47) | 14.6 (1.1-117) | 0.46 |
| Non-responders | | | | |
| HRU 30 days post-tx start median (range) | pRBC units Plt units Hospital Days | n = 19 2.0 (0-9) 4.0 (0-12) 9.0 (4-20) | n = 60 9.0 (1-25) 11 (0-47) 30 (15-30) | <0.001 0.003 <0.001 |
| HRU 60 days post-tx start median (range) | pRBC units Plt units Hospital Days | n = 11 8.0 (0-24) 8.0 (0-81) 14 (4-60) | n = 49 12 (6-36) 17 (0-88) 49 (24-60) | 0.037 0.008 <0.001 |

MyeloMatch

Precision Medicine initiative in AML and MDS from the NCI and NCTN



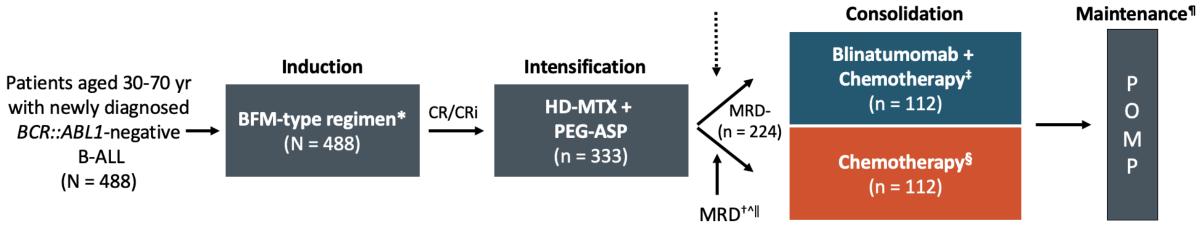
Acute Lymphoblastic Leukemia

ECOG-ACRIN E1910: Study Design

Blinatumomab is a CD3-CD19 BiTE

Multicenter, randomized, open-label phase III trial

Stratified by age (< or >55 yr), CD20 status, rituximab use, HSCT intent, MRD at randomization



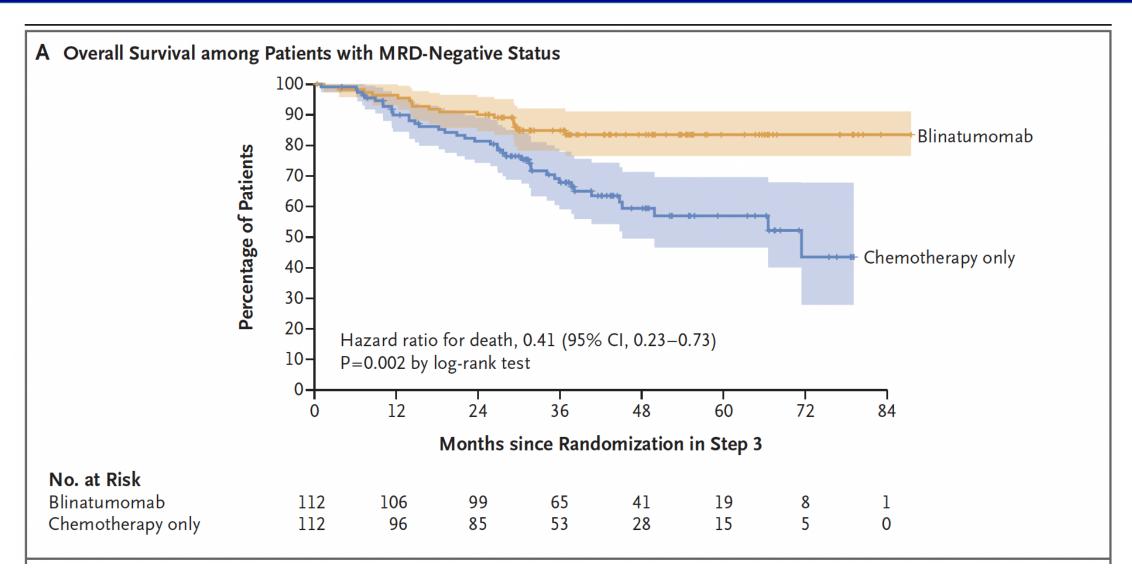
^{*}Regimen adapted from E2993/UKALLXII trial, including extended remission induction, addition of PEG-ASP for patients <55 yr of age, and addition of rituximab for CD20+ disease. †MRD assessed centrally by 6-color flow cytometry, with cutoff of ≤0.01% for MRD negativity. ^286 patients underwent MRD assessment, with 224 being negative and 62 being positive. ^{||}After blinatumomab regulatory approval in March 2018 for MRD^{pos} BCR::ABL1^{neg} B-ALL, MRD^{pos} patients were assigned to blina arm and no longer randomized. [‡]Two 28-day cycles of blina → 3 cycles of CT → 1 cycle of blina → 1 cycle of blina; 72% received 72-hr or 96-hr infusions. [§]Four cycles of consolidation CT. ^{||}2.5 yr of maintenance POMP timed from start of intensification. Patients could undergo alloHSCT at discretion of treating physician, ideally after first 2 cycles of blina in experimental arm or at any time following intensification in control arm.

Primary endpoint: OS in MRD^{neg} patients

Key secondary endpoints: MRD status, RFS



E1910: Primary Endpoint

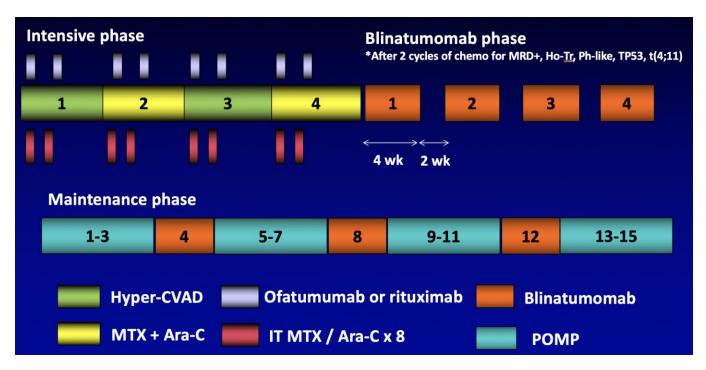


E1910: Adverse Events

| Event | Blinatumomab + Chemotherapy (N = 112) | | | Chemotherapy Only (N=112) | | |
|-----------------------------------|--|---------|---------------|---------------------------|---------|---------|
| | Grade 3 | Grade 4 | Grade 5 | Grade 3 | Grade 4 | Grade 5 |
| | | | percentage of | patients | | |
| Anemia | 20 | 1 | 0 | 35 | 2 | 0 |
| Leukopenia | 4 | 27 | 0 | 2 | 52 | 0 |
| Neutropenia | 3 | 55 | 0 | 1 | 86 | 0 |
| Lymphopenia | 3 | 8 | 0 | 6 | 17 | 0 |
| Thrombocytopenia | 9 | 40 | 0 | 10 | 59 | 0 |
| Febrile neutropenia | 16 | 1 | 0 | 21 | 2 | 0 |
| Sepsis | 0 | 4 | 1 | 0 | 6 | 1 |
| Hyperglycemia | 3 | 1 | 0 | 6 | 2 | 0 |
| Fatigue | 3 | 0 | 0 | 4 | 0 | 0 |
| ALT increased | 3 | 0 | 0 | 5 | 1 | 0 |
| AST increased | 1 | 0 | 0 | 1 | 2 | 0 |
| Hypertriglyceridemia | 0 | 3 | 0 | 1 | 3 | 0 |
| Nausea | 3 | 0 | 0 | 1 | 0 | 0 |
| Vomiting | 2 | 0 | 0 | 3 | 0 | 0 |
| Headache | 3 | 0 | 0 | 5 | 0 | 0 |
| Syncope | 3 | 0 | 0 | 3 | 0 | 0 |
| Other infection | 2 | 1 | 0 | 2 | 1 | 0 |
| Catheter-related infection | 1 | 0 | 0 | 3 | 1 | 0 |
| Upper respiratory tract infection | 1 | 0 | 0 | 3 | 0 | 0 |

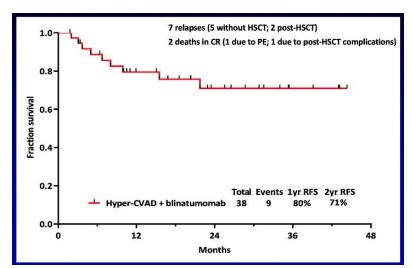
Treatment-related neurologic or psychiatric adverse event of grade 3 or higher in 23% of patients receiving blinatumomab vs 5% in the chemotherapy-only group (p < 0.001).

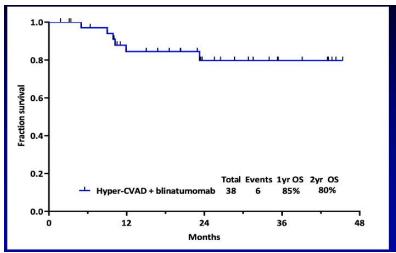
Hyper-CVAD plus Blinatumomab



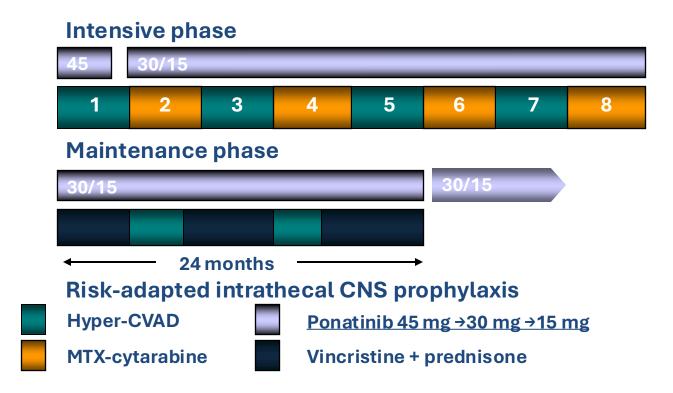
N=38, Median age 37 (17-59) CRS 13%, 3% G3 Blin-related Neuro events 45%, 13% G3

| Response* | n/N (%) |
|--------------------------------|-------------|
| CR after induction | 26/32 (81) |
| CR at any time | 32/32 (100) |
| MRD negativity after induction | 24/34 (71) |
| MRD negativity at anytime | 33/34 (97) |
| 30-day mortality | 0 |





Hyper-CVAD plus Ponatinib



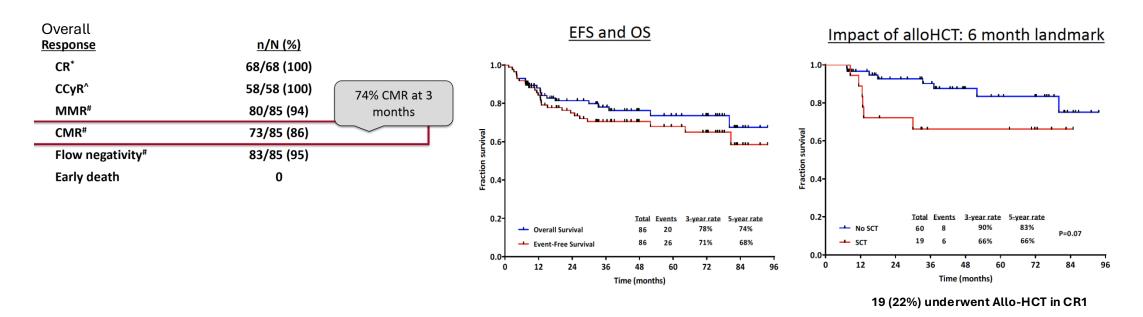
| <u>Characteristic</u> | <u>Category</u> | N (%) / median [range] |
|-----------------------------|------------------------|------------------------|
| Age (years) | | 46 (21-80) |
| | ≥60 | 20 (23) |
| Performance status | 0-1 | 78 (91) |
| | 2 | 8 (9) |
| WBC (x10 ⁹ /L) | | 13.6 [0.9-629.4] |
| CNS involvement | | 5 (6) |
| CD20 positivity | | 30 (35) |
| BCR-ABL transcript | p190 | 63 (73) |
| | p210 | 21 (24) |
| | Unknown | 2 (2) |
| Karyotype | Ph+ | 58 (67) |
| | Diploid/IM (FISH/PCR+) | 28 (33) |
| CNS disease at diagnosis | | 6 (7) |
| ≥ 1 baseline CV risk factor | | 56 (65) |

N=86

After the emergence of vascular toxicity, protocol was amended: Beyond induction, ponatinib 30 mg daily, then 15 mg daily once in CMR

12 doses of IT chemo (d2 and d7 cycles 1-6) 8 doses of R (cycles 1-4) for CD20+

Hyper-CVAD plus Ponatinib: Efficacy and Safety



3 relapses on ponatinib and no CNS relapses (12 IT ppx)

Toxicities – VTE (13%), Arterial CV events (7%); Grade 3+ infections (88%), AST/ALT elevation (29%), pancreatitis (15%), hyperbilirubinemia (15%), bleeding (14%), HTN (14%), rash (10%)

73% of VTE events at 45mg Pon; 67% of arterial CV events at 30-45mg Pon

No treatment related deaths after amendment of Pon dosing (2 died from MI prior)

Hyper-CVAD plus Ponatinib at UCDCCC

0.25

0.00

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Clinical experience with frontline Hyper-CVAD-based regimens, including Hyper-CVAD plus ponatinib, in patients with acute lymphoblastic leukemia treated at a comprehensive cancer center

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- * University of California Davis School of Medicine, Sacramento, CA, USA

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A. Overall survival

1.00

0.75

Aliging and 0.50

1.00

1.00

1.00

1.00

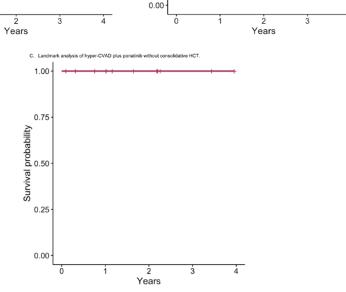
1.00

1.00

1.00

1.00

1.00

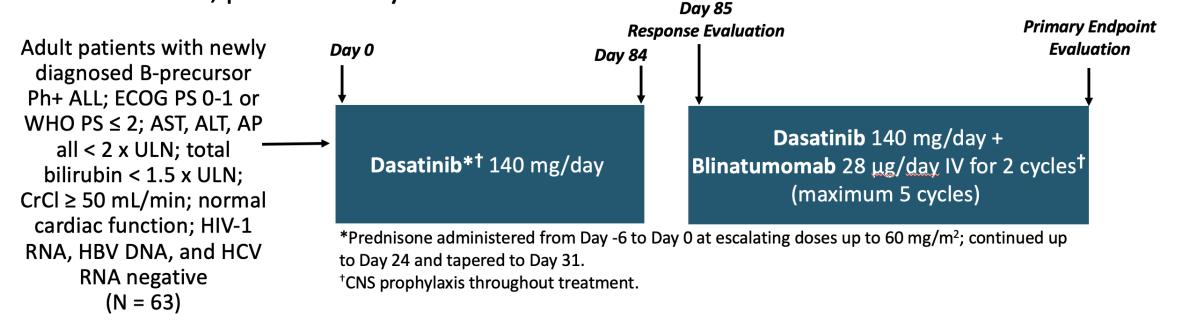


0.25

Fig. 2.: Impact of hyper-CVAD plus ponatinib on Ph+ ALL A. Overall survival. B. Event-free survival. C Landmark analysis of hyper-CVAD plus ponatinib without consolidative HCT.

Dasatinib + Blinatumomab for Ph+ ALL: Study Design

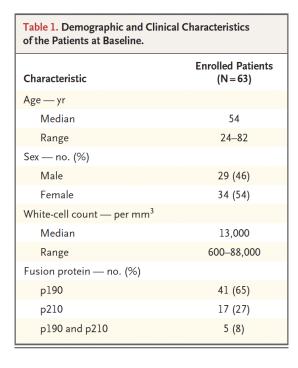
Multicenter, phase II study

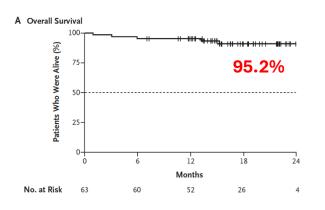


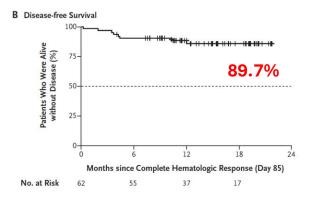
- Primary endpoint: CMR and MRD negativity after 2 cycles
- Secondary endpoints: CMR after dasatinib induction, CMR duration, OS, DFS, CIR, safety, MRD change after blinatumomab

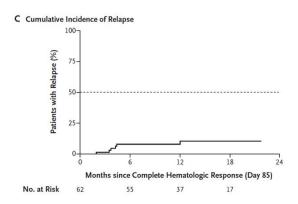
Slide credit: clinicaloptions.com

D-ALBA – Dasatinib plus Blinatumomab









Induction: 98% CHR, 29% molecular response

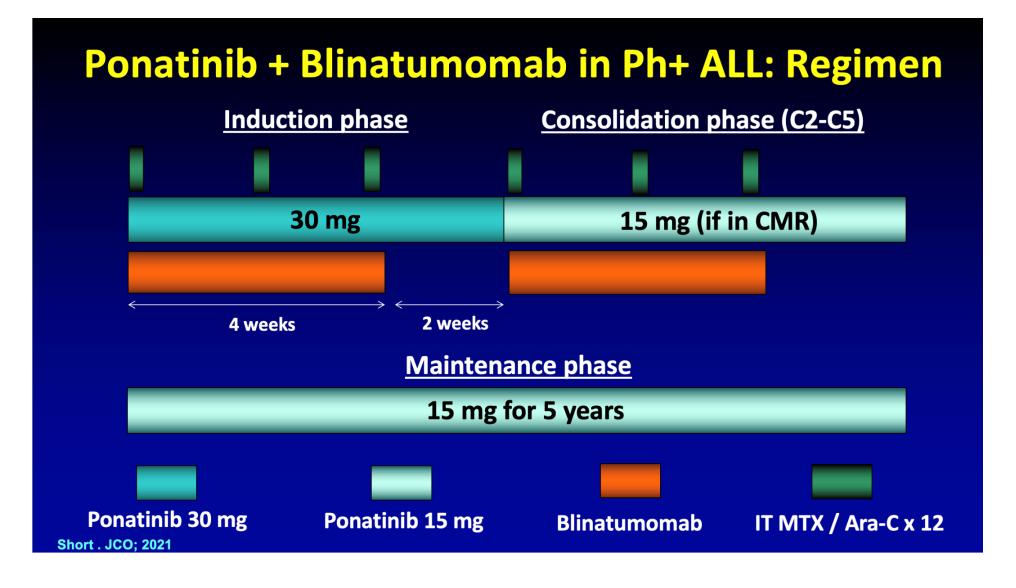
60.4% with CMR or non-quantifiable level after Cycle 2

Overall: 52% with molecular response

Median follow-up 14.3mo

Allo-HCT in 24pts

Patients aged ≥18 yr
with newly
diagnosed Ph+ ALL,
R/R Ph+ ALL, or
lymphoid AP/BP
CML; ECOG PS 0-2
(N = 55)



Primary outcome measures: CMR rate (newly diagnosed Ph+ and/or BCR-ABL+ ALL), ORR (R/R ALL), relapse-free survival, EFS, and OS

Ponatinib + Blinatumomab in Ph+ ALL: MRD Response Rates • 50 pts with ND Ph+ (n=30) median age 73 yrs (22-83), R/R Ph+ ALL (n=14), CML-BP (n=6) CMR ■ MMR ■ No MMR 100% 4% 10% 8% 8% 15% 18% 90% 8% 80% 40% 21% 70% 60% 80% 50% 20% 86% 85% 40% 77% 30% 61% 20% 40% 10% 20% 0% **FL ALL** R/R ALL **FL ALL** R/R ALL CML-LBC CML-LBC After 1st Cycle Overall Short. Blood 140: abst 2298; 2021

N=60 FL Ph+ B-ALL pts
39 untreated
21 in CR after 1-2 course of
chemotherapy within a median 49d
and a median duration of prior TKI
therapy 49d

TABLE 2. Hematologic and Molecular Responses

| Parameter | n/N (%) |
|------------------------------------|---------------|
| Overall response rate ^a | |
| CR | 37/39 (95) |
| CRi | 1/39 (2) |
| Early death | 1/39 (3) |
| CMR ^b | |
| After cycle 1 | 36/54 (67) |
| Overall | 45/54 (83) |
| MRD negativity by NGS/ClonoSEQ | |
| After cycle 1 | 10/22 (45) |
| Overall | 44/45 (98) |
| EFS | |
| 3-year rate, % (95% CI) | 77 (60 to 87) |
| No. of events | 10 (17) |
| OS | |
| 3-year rate, % (95% CI) | 91 (76 to 97) |
| No. of events | 4 (7) |
| | |

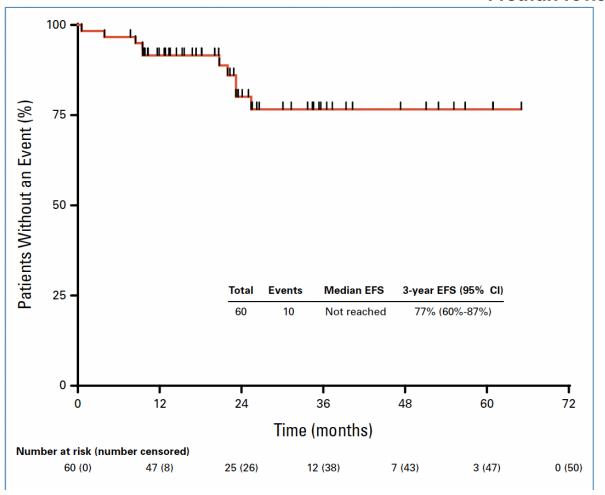
Abbreviations: CMR, complete molecular response; CR, complete remission; CRi, CR with incomplete hematologic recovery; EFS, event-free survival; MRD, measurable residual disease; NGS, next-generation sequencing; OS, overall survival.

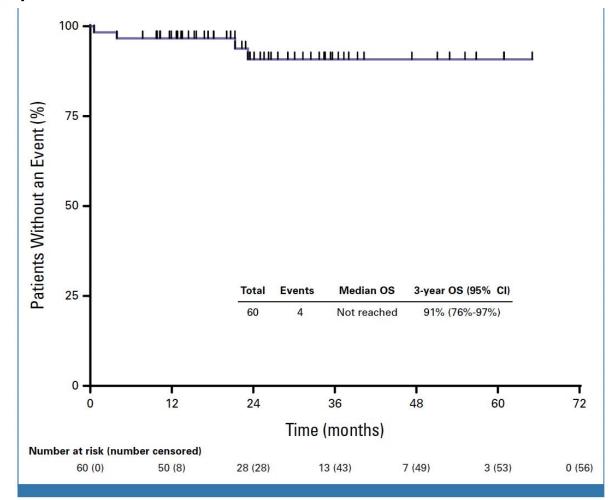
^aTwenty-one patients in CR at the start of therapy.

^bSix patients in CMR at the start of therapy.

Blinatumomab plus Ponatinib - Outcomes

Median follow-up 24 months





Recent FDA approvals in ALL and Ongoing Questions

- **Blinatumomab** is approved for CD19+ Ph- B-ALL in the consolidation phase of multiagent chemotherapy
 - Question how to give the Blina?
 - With E1910
 - With Hyper-CVAD
 - With other regimens, such as CALGB10403?
- **Ponatinib** is approved for newly diagnosed Ph+ ALL, in combination with chemotherapy
 - Question how to give the ponatinib?
 - With Hyper-CVAD or a Hyper-CVAD-like regimen
 - With Blinatumomab
 - Randomized EA9181 trial comparing TKI+Chemo vs TKI+Blina
 - How many IT chemo treatments?





