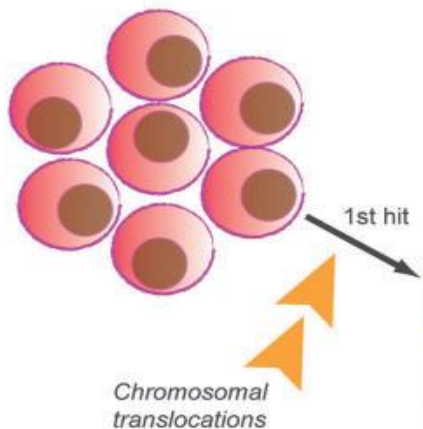


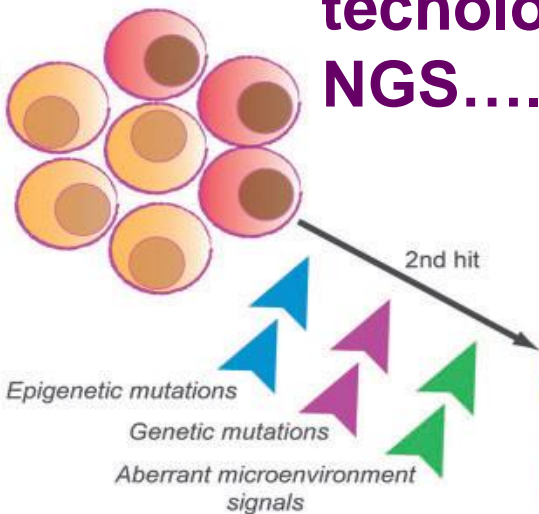
Dal 2008 al...

Caratterizzazione Biologica con la utilizzazione delle tecnologie PCR, GEP, NGS.....

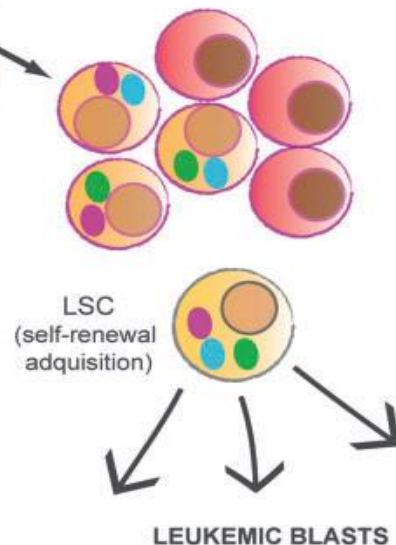
NORMAL BLOOD CELLS



PRELEUKEMIC STATE



ACUTE MYELOID LEUKEMIA



✓ Classificazione WHO 2008.....2016

✓ Score Prognostici ELN...

✓ New and Old Drugs

2008

10<sup>th</sup> EDITION

Highlights from EHA

**WILMS' TUMOR GENE MUTATIONS IN CHILDHOOD AML:  
CHARACTERISTICS, PROGNOSTIC VALUE AND CONSEQUENCES  
FOR MRD DETECTION (Hollink et al, #457)**

**AMONAFIDE: A TOPO II INHIBITOR WITH NOVEL  
PHARMACOLOGICAL PROPERTIES AND UNIQUE ACTIVITY FOR THE  
TREATMENT OF SEC. AML (Capizzi et al, #890)**

**PHASE II STUDY OF SINGLE AGENT CLOFARABINE IN UNTREATED  
ELDERLY PATIENTS WITH AML UNLIKELY TO BENEFIT FROM  
STANDARD INDUCTION CHEMOTHERAPY (Erba et al, #892)**

## **The molecular basis of AML**

**K.L. Rice**

## **Genetic markers in relations to the therapeutic management**

**B. Lowenberg,**

# *The* NEW ENGLAND JOURNAL *of* MEDICINE

ESTABLISHED IN 1812

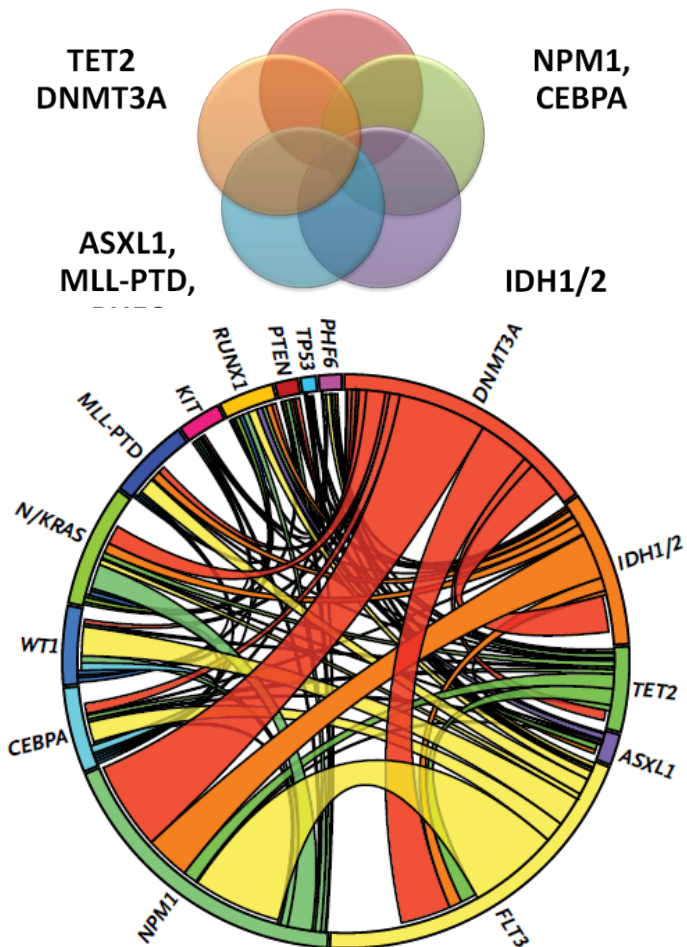
MARCH 22, 2012

VOL. 366 NO. 12

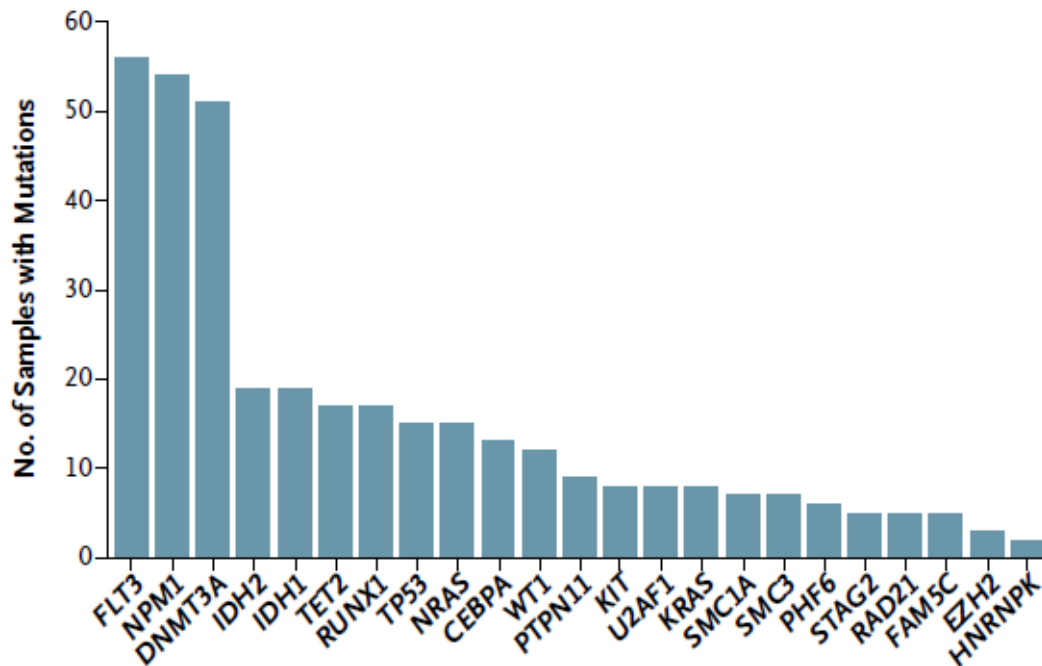
## Prognostic Relevance of Integrated Genetic Profiling in Acute Myeloid Leukemia

Jay P. Patel, Mithat Gönen, Ph.D., Maria E. Figueroa, M.D., Hugo Fernandez, M.D., Zhuoxin Sun, Ph.D., Janis Racevskis, Ph.D., Pieter Van Vlierberghe, Ph.D., Igor Dolgalev, B.S., Sabrena Thomas, B.S., Olga Aminova, B.S., Kety Huberman, B.S., Janice Cheng, B.S., Agnes Viale, Ph.D., Nicholas D. Socci, Ph.D., Adriana Heguy, Ph.D., Athena Cherry, Ph.D., Gail Vance, M.D., Rodney R. Higgins, Ph.D., Rhett P. Ketterling, M.D., Robert E. Gallagher, M.D., Mark Litzow, M.D., Marcel R.M. van den Brink, M.D., Ph.D., Hillard M. Lazarus, M.D., Jacob M. Rowe, M.D., Selina Luger, M.D., Adolfo Ferrando, M.D., Ph.D., Elisabeth Paietta, Ph.D., Martin S. Tallman, M.D., Ari Melnick, M.D., Omar Abdel-Wahab, M.D., and Ross L. Levine, M.D.

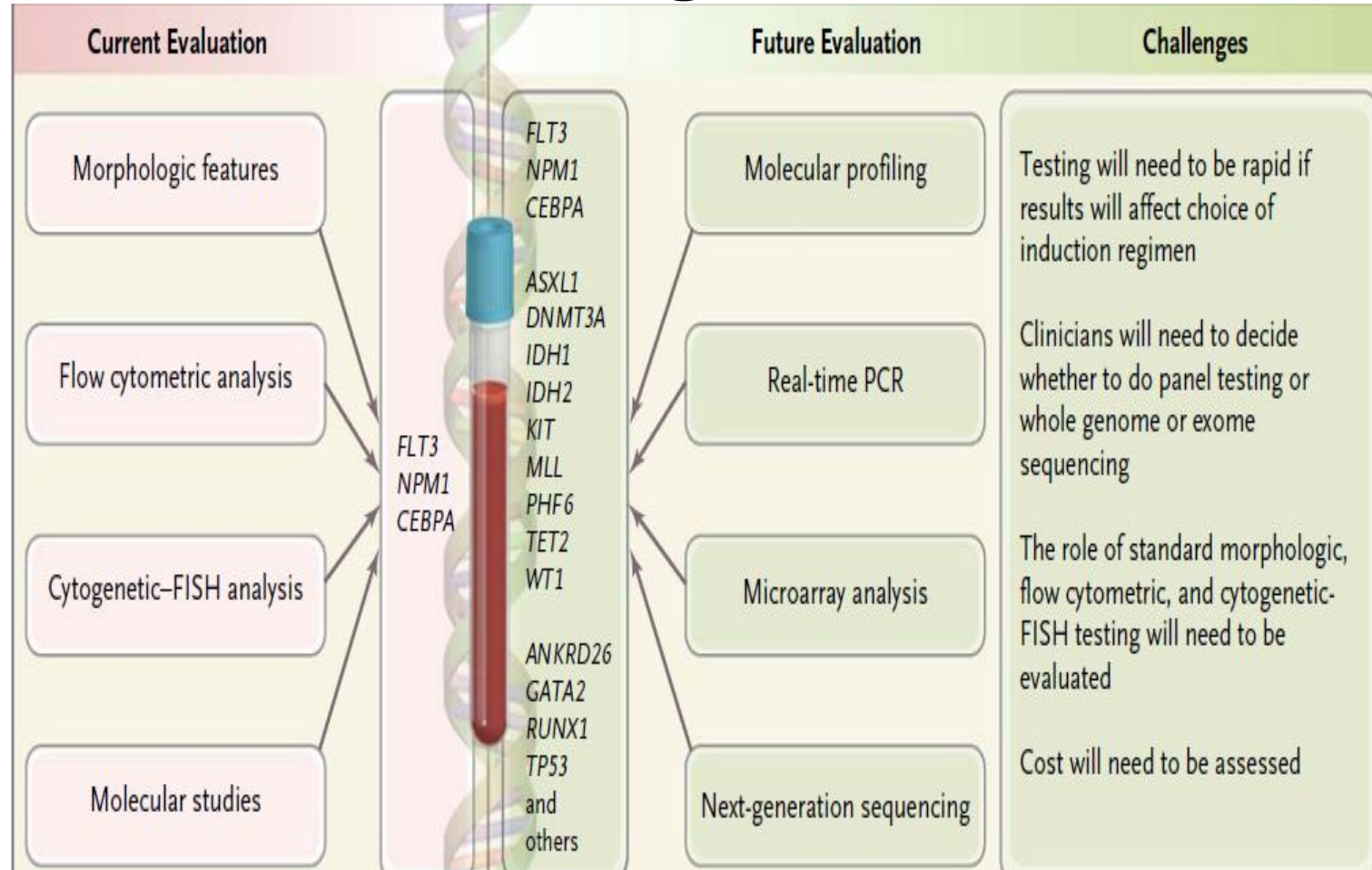
**FLT3-ITD**



**B Significantly Mutated Genes**



# AML Diagnosis



## Do they have prognostic value?

### Poor Survival:

FLT3<sup>+</sup> or MLL and in those with point mutations of ASXL1 or PHF6.

### Favorable Survival:

CEBPA or IDH2 mutations; NPM1 mutations with concurrent IDH1 or IDH2 mutations.

## Lestaurtinib / Sorafenib

Inhibition of constitutively activated FLT3, lestaurtinib in relapsed AML and sorafenib in newly diagnosed older AML, have failed to demonstrated significant benefit when combined to intensive chemotherapy.



## Midostaurin / Quizartinib

- Phase III randomized study of midostaurin restricted to FLT3 mutated pts younger than 60 yrs is ongoing.
- Phase II study of quizartinib or AC220, the most selective FLT3 inhibitor available, in relapsed AML have confirmed that clonal responses could be observed with monotherapy.

## **Best of EHA in acute myeloid leukemia (AML)**

### **Biological studies:**

*Murine models of NPM1-mutated AML*

*Mutations in AML secondary to congenital neutropenia  
(Plenary Session)*

*Mutations in AML secondary to Down syndrome  
(Plenary session)*

### **Clinical Studies:**

*Value of MRD monitoring in NPM1-mutated AML*

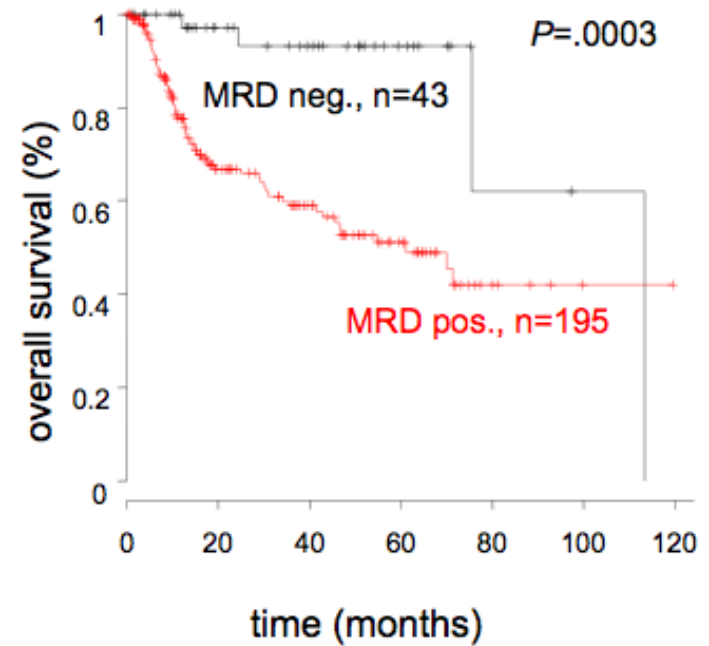
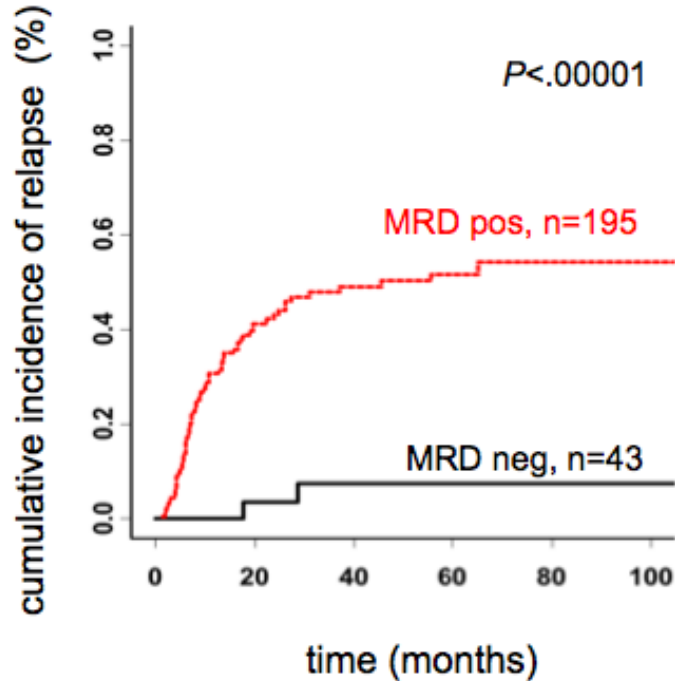
*First trial with PI3K inhibitor Volasertib in relapsed/refractory AML*

*Development of bispecific CLL-1 x CD3 antibody for therapy of AML*

MRD Monitoring in *NPM1* mutated AML:

A Study of the German-Austrian  
AML Study Group (AMLSG)

After double induction in patients in CR (n=238)



**2014**

- **Conventional & novel hypomethylating agents**
- **Novel targeted therapies**
- **AML founding mutations and HSC**

RESULTS OF A PHASE III, MULTICENTER, RANDOMIZED, OPEN-LABEL STUDY OF AZACITIDINE (AZA) VS CONVENTIONAL CARE REGIMENS (CCR) IN OLDER PATIENTS WITH NEWLY DIAGNOSED AML

PHASE I/II STUDY OF VOLASERTIB, A POLO-LIKE KINASE INHIBITOR (PLK), IN PATIENTS WITH RELAPSED/REFRACTORY AML: UPDATED PHASE I RESULTS FOR VOLASERTIB MONOTHERAPY

A PHASE I STUDY OF AG-221, A FIRST IN CLASS, POTENT INHIBITOR OF THE IDH2-MUTANT PROTEIN, IN PATIENTS WITH IDH2 MUTANT POSITIVE ADVANCED HEMATOLOGIC MALIGNANCIES

2015

10<sup>th</sup> EDITION

Highlights from EHA

***Dnmt3a R882*** Mutations Promote  
Chemoresistance and Therapeutic Relapse  
Through Impaired DNA Damage Sensing

***DNMT3A<sup>mut</sup>*** AML patients are less sensitive to  
anthracyclines and benefit from dose-intensification

**What's new in the WHO classification ?**

*Clara Bloomfield*

*Wthe Ohio State University, Columbus, Ohio, USA)*

## WHO 2016: proposed changes in the category of “AML with recurrent genetic abnormalities”

- Switching from provisional to distinct entities:

*AML with mutated NPM1 (distinct entity)*

*AML with double mutated CEBPA (distinct entity)\**

2016

## Clinical trials

- Benefit of HD-DAU in FLT3-ITD<sup>mut</sup> AML (NCRI-AML17)
- Benefit of CPX-351 in FLT3<sup>mut</sup> AML (update on phase 3/HR-AML)
- Volasertib+LDAC (phase 3/elderly AML)
- Vosaroxin+Decitabine (phase 1-2/elderly AML+HR-MDS)

## Novel targeted agents to watch....

- SGN-CD33A in combination with HMA (phase 1/CD33+ AML)



## New approaches starting to bear fruit...



Amadori S. 2016

## **MRD: nuovo endpoint surrogato (OS, EFS) nella AML?**

Quali tecnologie?; quali time-points?

## **L'era della chemio intensiva di prima linea ("AML Dogma") è ormai prossima alla fine?**

Eccellenti risultati (CR/CRi, tossicità) con nuovi farmaci mirati (Venetoclax in primis) in combinazione con HMAs o chemio a bassa intensità (LDAC) in pazienti anziani poor-risk (età/fitness, biologia)

**Tutti gli anziani?**

**Anche nei giovani?**

**Chi dovrebbe continuare ad essere trattato con chemio intensiva?**



## **Acute myeloid leukemia**

Gert Ossenkoppele (Coordinating Author)

### ***Molecular diagnostics in acute myeloid leukemia***

Lars Bullinger

*Department of Internal Medicine III, Ulm University, Germany*

### ***Targeting mutated FLT3 in acute myeloid leukemia***

Mark Levis

*Johns Hopkins University, Baltimore, USA*

### ***3+7 and beyond***

Norbert Vey

*Institut Paoli Calmettes and Aix-Marseille Université, Marseille, France*

# Risk Stratification and Cytogenetic and Molecular Abnormalities in AML

## Favorable Genetic Risk

**Frequency:** 15%

**Survival:** 65%

- t(8;21)(q22;q22); *RUNX1-RUNX1T1*
- inv(16)(p13.1q22) or t(16;16)(p13.1;q22); *CBFB-MYH11*
- Mutated *NPM1* without *FLT3* ITD or with *FLT3-ITD*<sup>low</sup>
- Biallelic mutated *CEBPA* (normal karyotype)

## Intermediate Genetic Risk

**Frequency:** 55%

**Survival:** 50%

- Mutated *NPM1* and *FLT3-ITD*<sup>high</sup> (normal karyotype)
- Wild-type *NPM1* without *FLT3-ITD* or with *FLT3-ITD*<sup>low</sup>
- t(9;11)(p21.3;q23.3); *MLL3-KMT2A*

## Adverse Genetic Risk Group

**Frequency:** 30%

**Survival:** 20%

- t(6;9)(p23;q34.1); *DEK-NUP214*
- t(v;11;q23.3); *KMT2A* rearranged
- t(9;22)(q34.1;q11.2) *BCR-ABL1*
- inv(3)(q21.3q26.2) or t(3;3)(q21.3;q26.2); *GATA2 MECOM (EVI1)*
- -5 or del(5q)-7; -17/abn(17p)
- Complex and/or monosomal karyotype
- Wild-type *NPM1* and *FLT3-ITD*<sup>high</sup>
- Mutations in *RUNX1*, *ASXL1*, *TP53*

- ✓ **Genomic knowledge** does now also facilitate monitoring of **MRD** ( DPCR, NGS, qRT-PCR, MFC ).
- ✓ Comprehensive and individualized **MRD assessment** is useful to identify pts at high relapse risk at early time points.
- ✓ **Genomic knowledge** will allow us to better guide the use of novel drugs

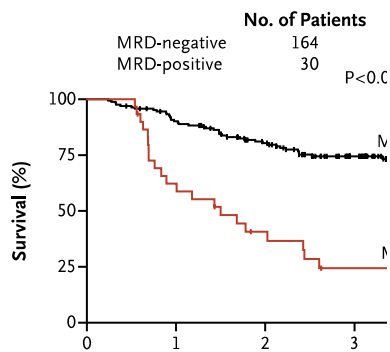
The NEW ENGLAND JOURNAL of MEDICINE

# Assessment of Minimal Residual Disease in Standard-Risk AML

A. Ivey, R.K. Hills, M.A. Simpson, J.V. Jovanovic, A. Gilkes, A. Grech, Y. Patel,  
N. Bhudia, H. Farah, J. Mason, K. Wall, S. Akiki, M. Griffiths, E. Solomon,  
F. McCaughan, D.C. Linch, R.E. Gale, P. Vyas, S.D. Freeman, N. Russell,  
A.K. Burnett, and D. Grimwade, for the UK National Cancer Research Institute

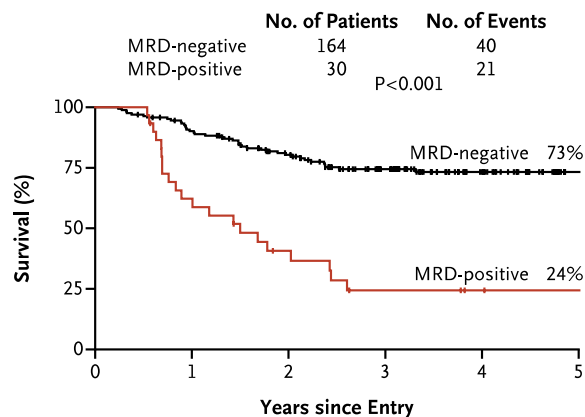
AML

**A Overall Survival**



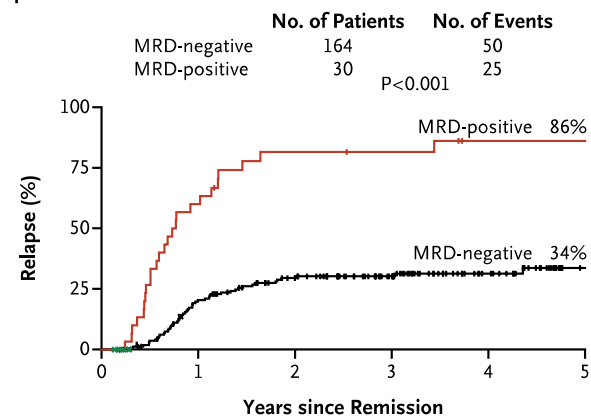
No. at Risk	Years since Entry			
	0	1	2	3
MRD-negative	164	144	116	77
MRD-positive	30	18	10	5

**A Overall Survival**



No. at Risk	Years since Entry					
	0	1	2	3	4	5
MRD-negative	164	144	116	77	39	8
MRD-positive	30	18	10	5	3	2

**B Relapse in All Patients**



No. at Risk	Years since Remission					
	0	1	2	3	4	5
MRD-negative	164	120	93	64	33	6
MRD-positive	30	12	5	4	1	1

## Problematiche nella real life per la valutazione della MRD

**Quali BioMarkers** PML/RAR $\alpha$ , NPM1,

**MFC** ma in tutti i Laboratori è standardizzata ? no

Sono necessari altri trials?

Potremmo utilizzare la piattaforma LabNet AML anche per standardizzare ?

**L'era della chemio intensiva di prima linea (“AML Dogma”) è ormai prossima alla fine?**

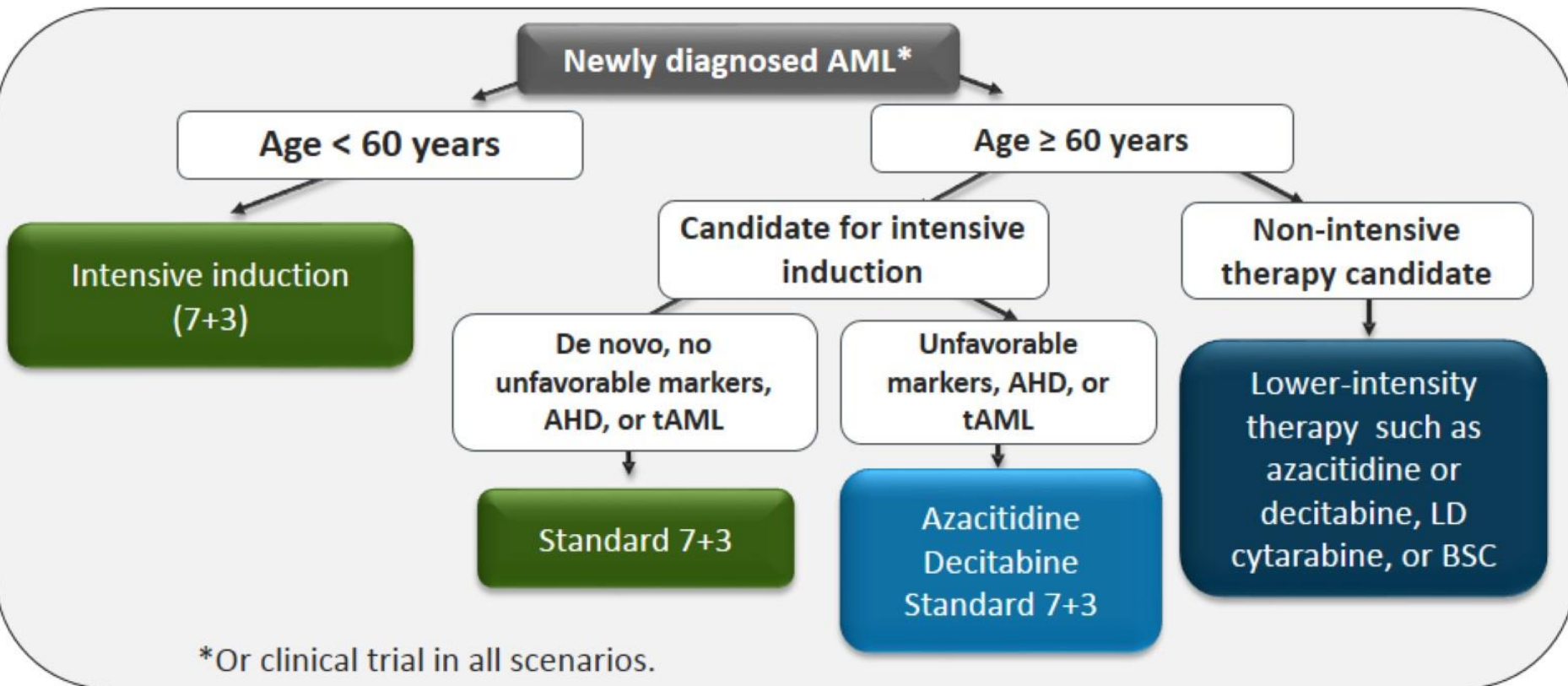
**Chi dovrebbe continuare ad essere trattato con CHT Intensiva?**

**Tutti i pazienti elegibili per la CHT standard/intensiva !!  
DNR/IDA (> 60mg/m, 12 mg/m) ; ARA-C 100-200-1000mg)  
FLAG-Ida ?**

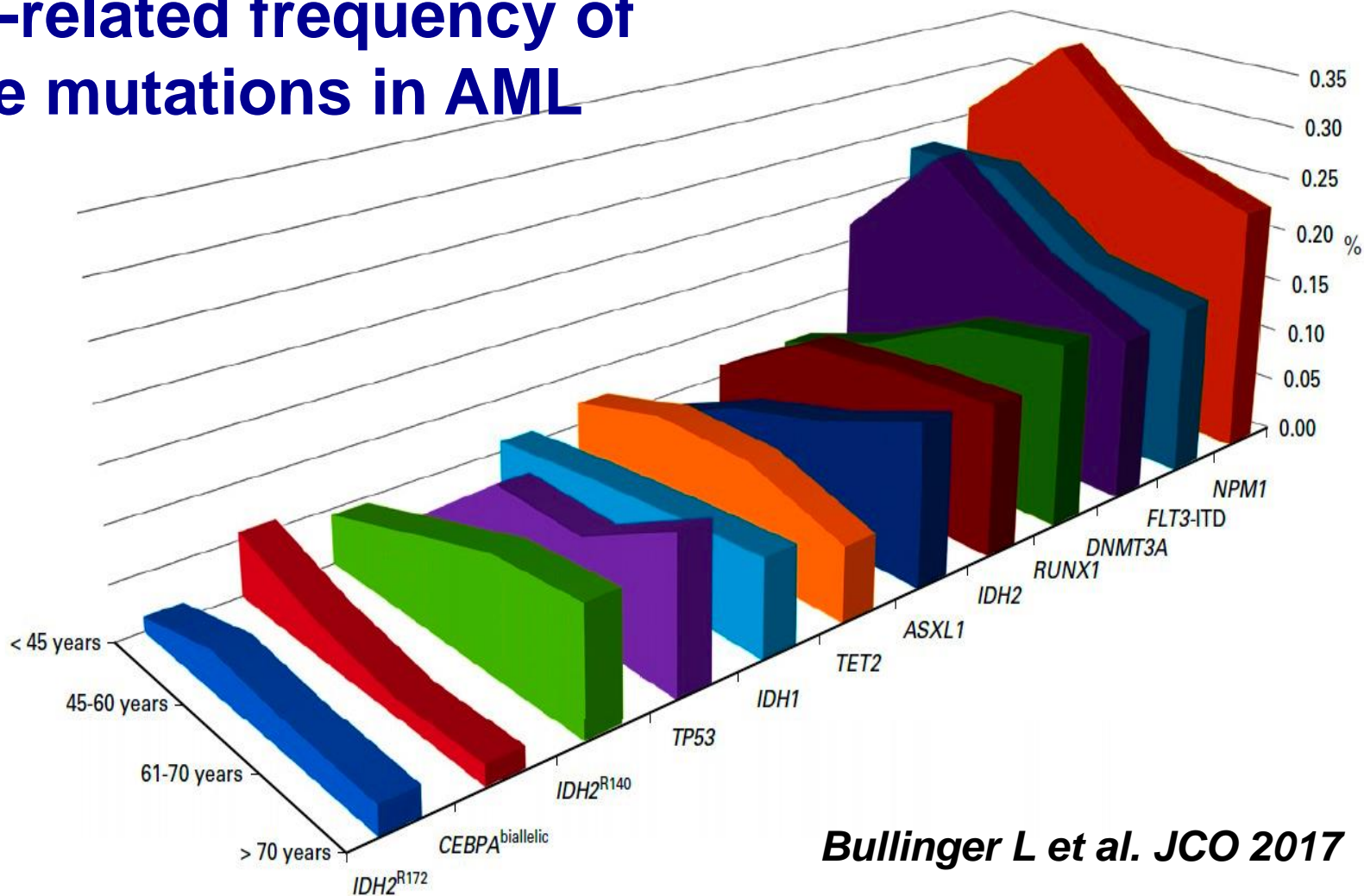


# Treating Newly Diagnosed AML

## Current Paradigms



# Age-related frequency of gene mutations in AML



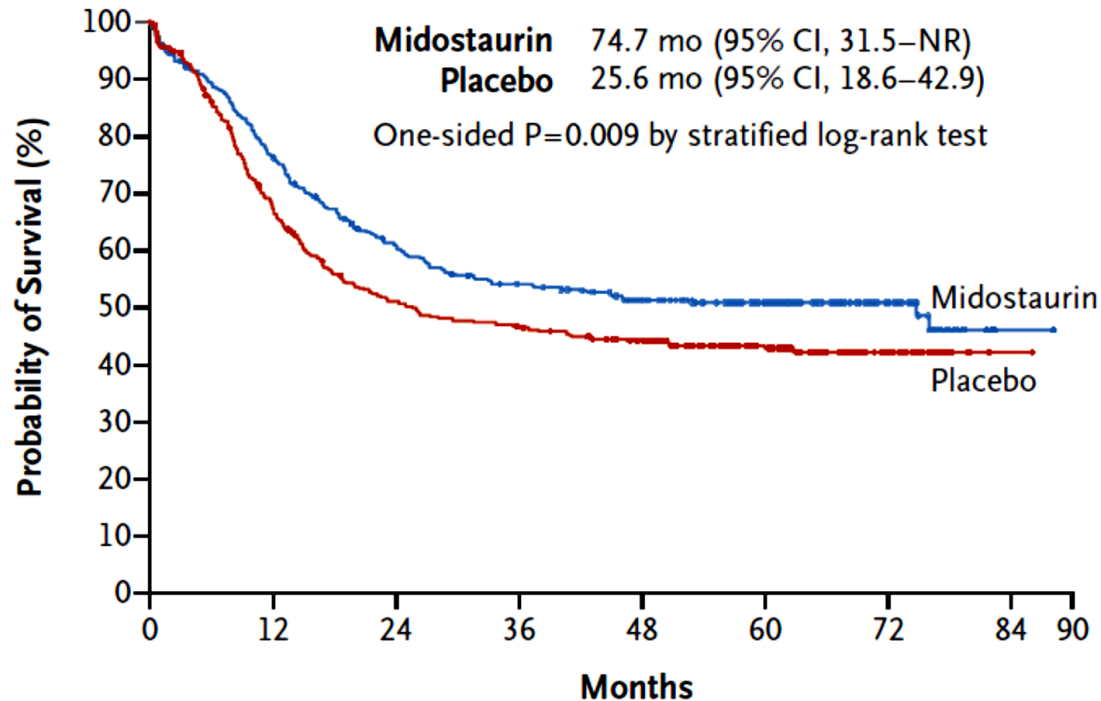
**Bullinger L et al. JCO 2017**

## Examples of Novel Targeted Therapies in AML

- FLT3 inhibitors
- IDH inhibitors
- Venetoclax
- Monoclonal antibody-drug conjugates, such as gemtuzumab-ozogamicin and SGN-CD33A
- BiTE antibodies
- Immune checkpoint inhibitors
- Novel formulations of cytotoxic agents
- CPX-351 (combination of daunorubicin and cytarabine)
- Vosaroxin, a TP53-independent drug that may be particularly useful in patients with relapsed disease and those older than 60 years
- Hedgehog pathway/MEK pathway inhibitors
- MDM2 inhibitors

# Midostaurin plus Chemotherapy for AML with a FLT3 Mutation

## Median Overall Survival

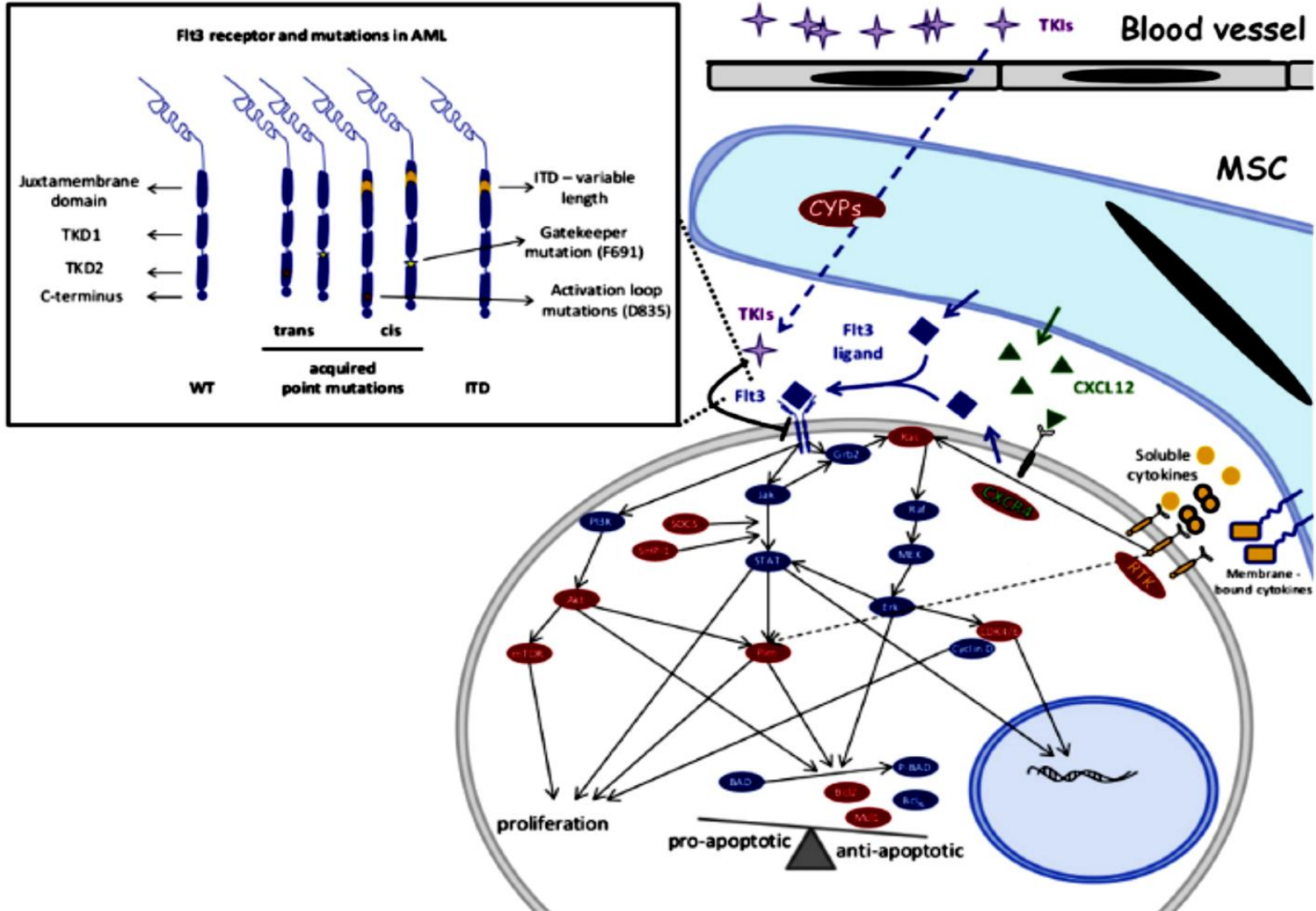


## No. at Risk

Midostaurin	360	269	208	181	151	97	37	1
Placebo	357	221	163	147	129	80	30	1

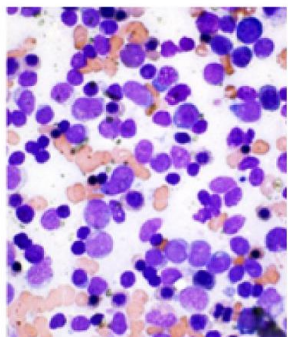
The addition of the multitargeted kinase inhibitor midostaurin to standard chemotherapy significantly prolonged overall and event-free survival among patients with AML and a FLT3 mutation

# Resistance to FLT3 Inhibitors

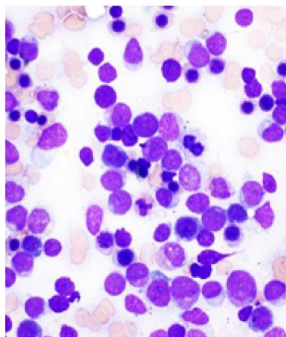


## Enasidenib induces AML cell differentiation to promote clinical response

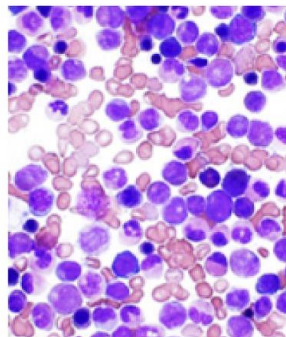
Screening  
37% BM blasts



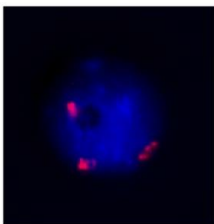
Cycle 1 Day 15  
Evidence of cellular differentiation



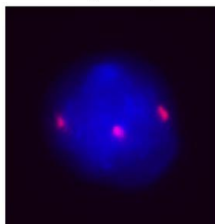
Cycle 3 Day 1  
4% BM blasts



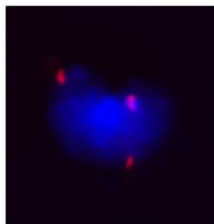
Blasts



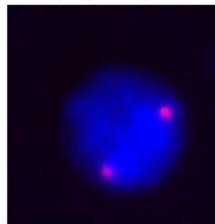
Promyelocytes



Mature Granulocytes



Lymphocytes

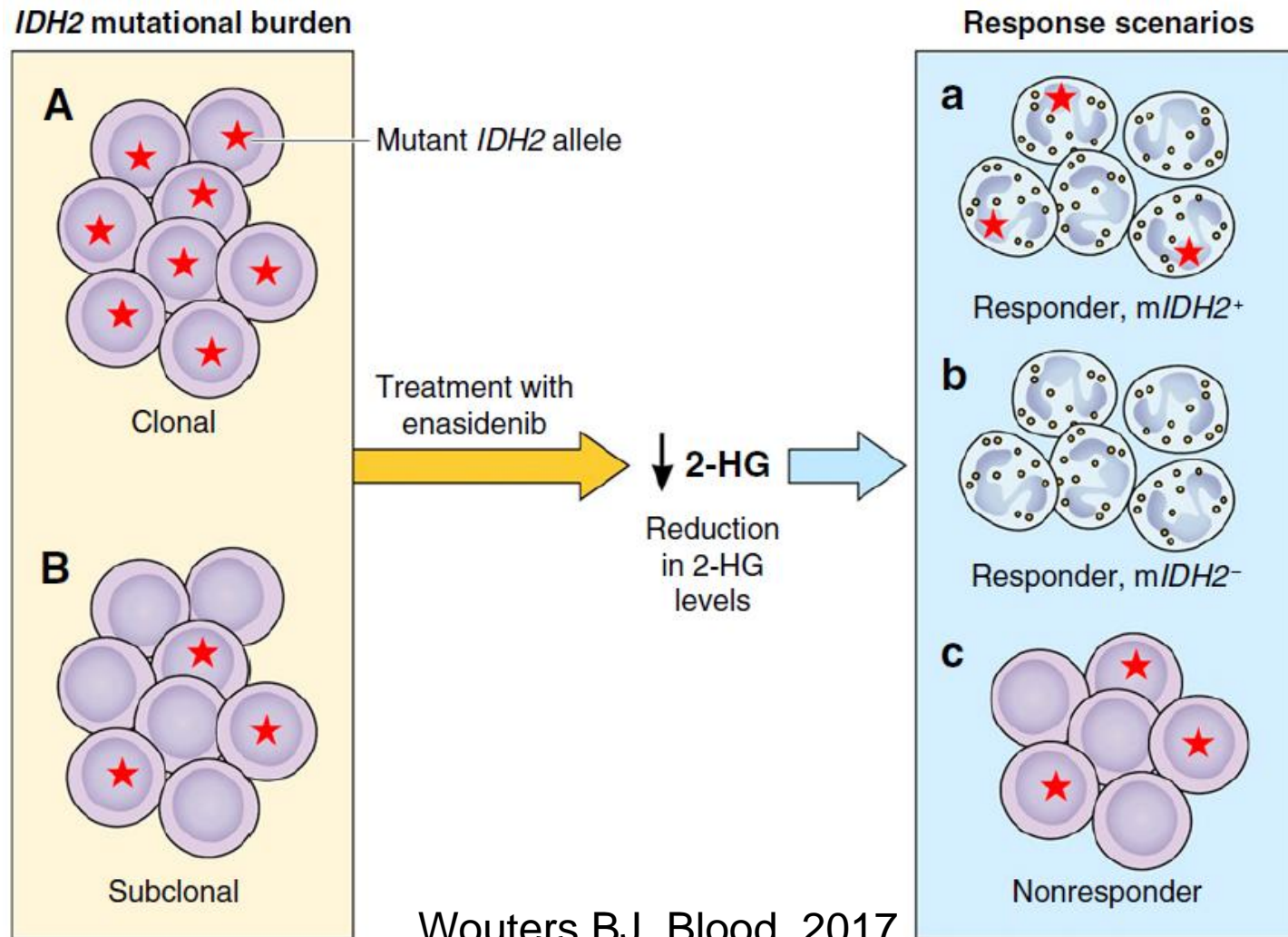


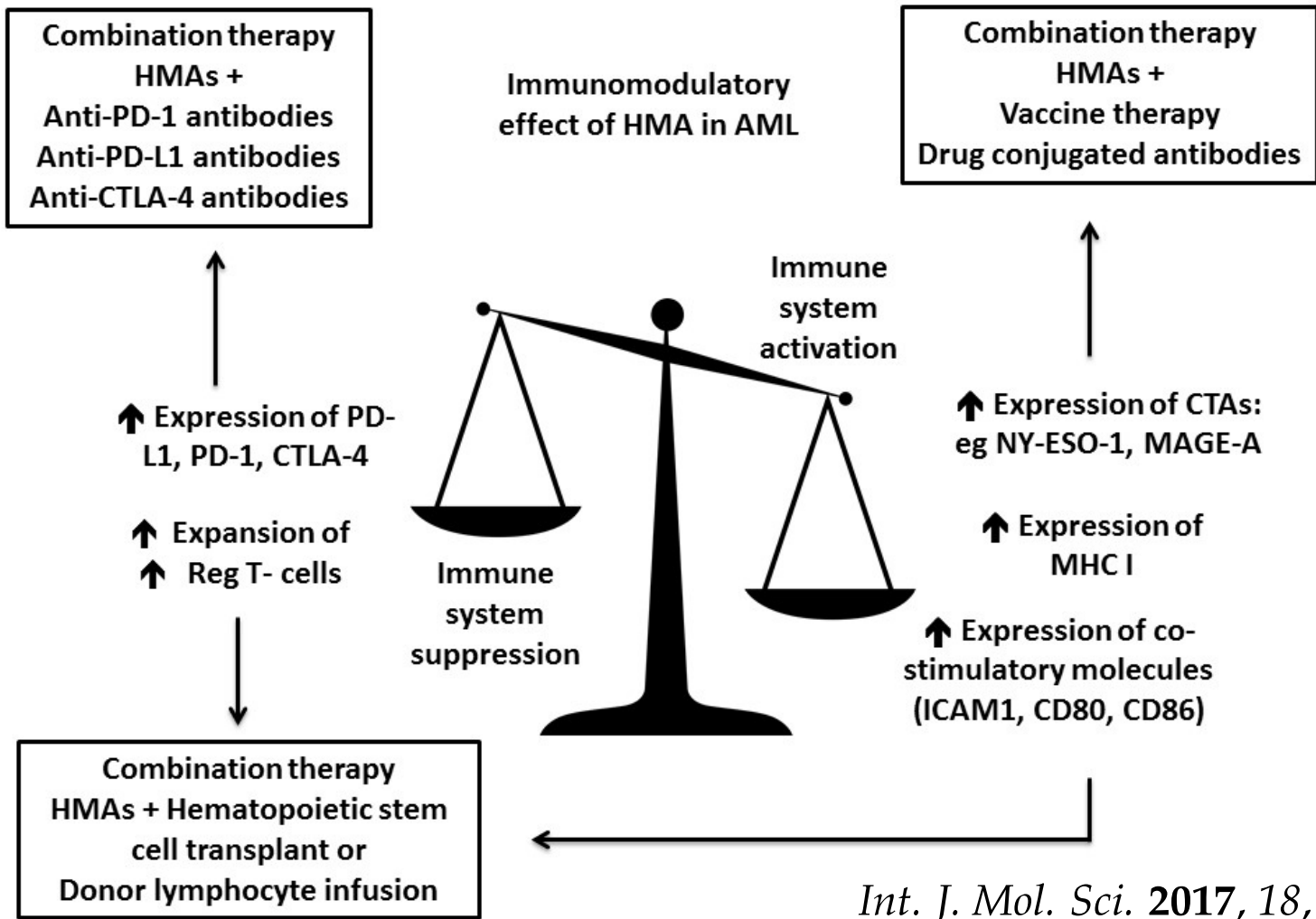
CR with persistence of mIDH2 and normalization of hematopoietic stem and progenitor compartments with emergence of functional mIDH2 neutrophils were observed. In a subset of CR patients, mIDH2 allele burden was reduced and remained undetectable with response.

Co-occurring mutations in NRAS and other MAPK pathway effectors were enriched in nonresponding patients, consistent with RAS signaling contributing to primary therapeutic resistance.

Together, these data support differentiation as the main mechanism of enasidenib efficacy in relapsed/refractory AML patients and provide insight into resistance mechanisms to inform future mechanism-based combination treatment studies

## Enasidenib induces AML cell differentiation to promote clinical response







# New approaches starting to bear fruit...



Amadori S. 2016



# 10<sup>th</sup> EDITION

# Highlights from EHA

## Cytotoxic agents

Liposomal D+A Topo-II inhibitor	CPX 351 Vosaroxin	HR elderly AML frontline R/R AML	Rando Phase 2 Phase 3 R/R
------------------------------------	----------------------	-------------------------------------	------------------------------

## Monoclonal antibodies

AntiCD33 mAb	lintuzumab	Misc.	Phase 3
AntiCD33 ADC	GO	frontline	Phase 3
	SGN-33A	R/R AML +frontline	Phase 3 combo
AntiCD33/CD3	AMG330	R/R AML	Phase 1 single agent
Anti-CD123 mAb	Talacotuzumab	R/R AML	Phase 2 combo
Anti-CD123 ADC	SGN-CD123A	R/R AML	Phase 1 single agent
Anti-CD3/CD123	MGD006	R/R AML	Phase 1 single agent
	JNJ-63709178	R/R AML	Phase 1 single agent

## Apoptosis targeting agents

BCL2-i	Venetoclax	R/R AML	Phase 2 combos
	S55746	R/R AML	Phase 1
MCL1-i	S64315	R/R AML	Phase 1
MDM2-i	Idasanutlin	R/R AML	Phase 3 combo

## Kinase/Cell cycle-i

PIM kinase-i	CLGH447	R/R AML	Phase 1 combo
MEK-i	Cobimetinib	R/R AML	Phase 1 combo
PI3K/RAS-i	Rigosertib	R/R AML	Phase 1
CDK-i	Palbociclib	R/R AML	Phase 1

## Epigenetic drugs

Oral azacitidine	CC486	Frontline	Phase 3 combo
Decitabine prodrug	SGI-110	Frontline elderly	Phase 3
Bromodomaine-i	OTX015	R/R AML	Phase 1
DOTL1-i	EPZ-5676	R/R MLL AML	Phase 1

# 10<sup>th</sup> EDITION

# Highlights from EHA

<b>Immunotherapy</b>			
<b>ICB</b>			
Anti-CTLA4	Ipilimumab	R/R AML	Phase 1-2
Anti-PD1	Nivolumab	R/R + frontline AML	Phase 1-2 combo
Anti-KIR	IPH2101	R/R AML	Phase 1
Anti-NKG2A	Lirilumab	frontline elderly AML	Phase 2-3
CAR-T cells	Monalizumab	Maintenance post allo	Phase 1
Anti-CD33	CART33	R/R AML	Phase 1
Anti-CD123	CART123	R/R AML	Phase 1
Anti-CD133	CART133	R/R AML	Phase 1

**Table 2**  
**Selected upfront acute myeloid leukemia clinical studies with FLT3 inhibitors**

<b>Drug, Reference</b>	<b>Patients</b>	<b>FLT3 Status</b>	<b>Phase and Treatment Regimen</b>	<b>Treatment Response</b>
Sorafenib <sup>49</sup>	18 yo+	+/- <i>FLT3</i> mutation	II: Sor 400 mg po bid plus AraC + Ida for induction, Sor plus cytarabine for consolidation, and Sor alone as maintenance × 1 y	<ul style="list-style-type: none"> <li>• CR in 79% (n = 49 of 61) and CRp in 8% (n = 5 of 61), including CR/CRp 95% (n = 18 of 19) and 84% (n = 36 of 43) with and without <i>FLT3</i>-ITD, respectively</li> <li>• Median OS: 29 mo</li> <li>• Median DFS: 13.8 mo</li> </ul>
Sorafenib <sup>50</sup>	60 yo+	+/- <i>FLT3</i> mutation	II: randomized to Sor 400 mg PO BID vs placebo after DNR + AraC, after cytarabine for consolidation and as maintenance × 1 y	Placebo vs sorafenib: <ul style="list-style-type: none"> <li>• ORR (CR + CRi) 64 of 95 vs 57 of 102 (<i>P</i> = .34), respectively</li> <li>• EFS 7 m vs 5 m (HR 1.26; 95% CI, 0.94–1.70)</li> <li>• OS 15 m vs 13 m (HR 1.03; 95% CI, 0.73–1.44)</li> <li>• Sorafenib arm had higher 60-d mortality (<i>P</i> = .035) attributable to infections (<i>P</i> = .015)</li> </ul>
Sorafenib <sup>51</sup>	60 yo+	+ <i>FLT3</i> -ITD or + <i>FLT3</i> TKD	II: sorafenib 400 mg po bid days 1–7 plus 7 + 3, followed by Sor plus intermediate dose AraC for consolidation and Sor alone as maintenance × 1 y	<ul style="list-style-type: none"> <li>• CR or CRi in 69% (n = 37 of 54)</li> <li>• 1 y observed OS: 62% for <i>FLT3</i>-ITD and 71% for <i>FLT3</i>-TKD</li> <li>• Favorable outcome (1-y OS) compared with historical controls for <i>FLT3</i>-ITD (62% vs 30%; <i>P</i> &lt; .0001)</li> </ul>
Sorafenib <sup>52</sup>	18–60 yo	+/- <i>FLT3</i> mutation	II: randomized to Sor 400 mg po bid vs placebo after 7 + 3 for induction, after cytarabine for consolidation and as maintenance × 1 y	Placebo (n = 133) vs sorafenib (n = 134): <ul style="list-style-type: none"> <li>• CR: 59 vs 60%</li> <li>• 3-y EFS 22% (95% CI, 13–32) vs 40% (HR 0.64; 95% CI, 0.45–0.91; <i>P</i> = .013)</li> <li>• No OS difference</li> </ul>

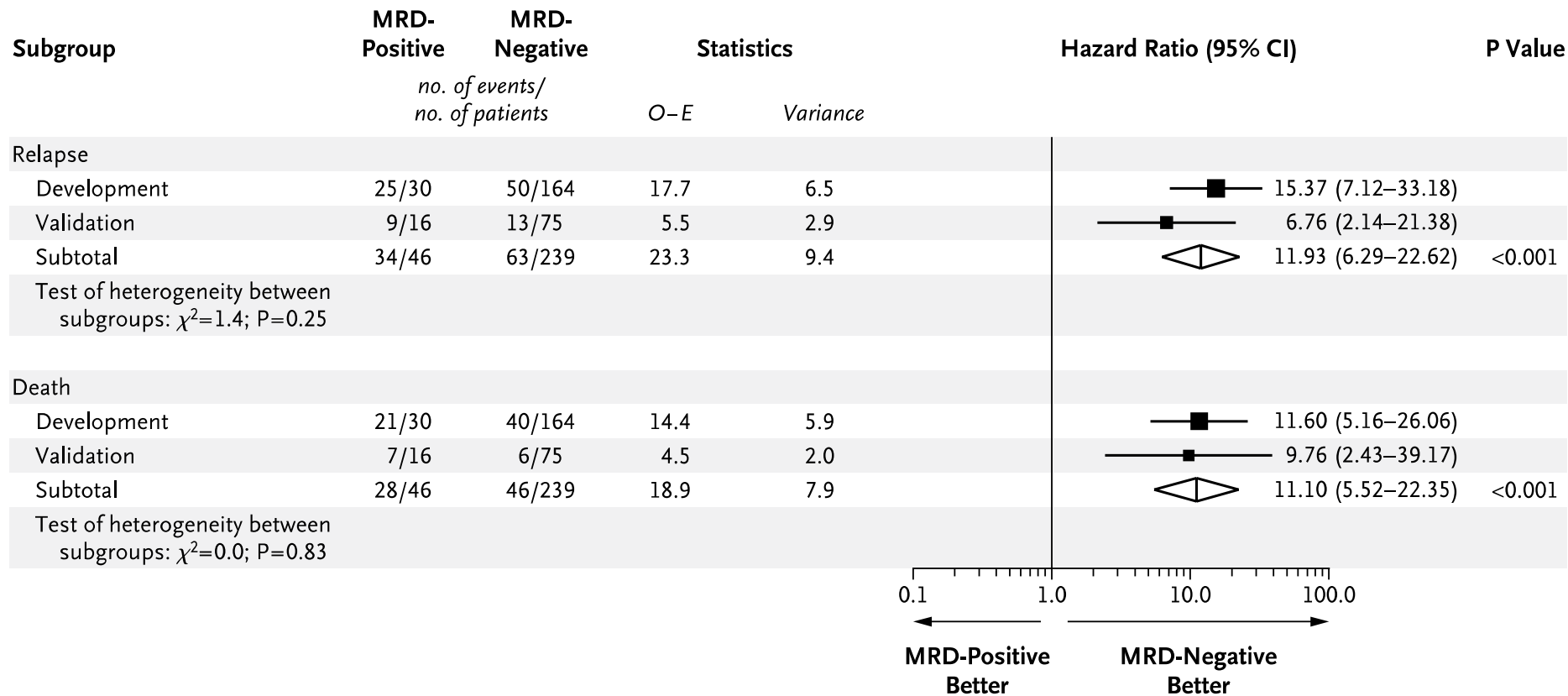
# 10<sup>th</sup> EDITION Highlights from EHA

Midostaurin <sup>55</sup>	18–60 yo	+/- <i>FLT3</i> mutation	Ib: M 50–100 mg, po bid, either concomitantly or sequentially with 7 + 3, M with HiDAC consolidation and M alone as maintenance	<ul style="list-style-type: none"> <li>• 100-mg cohort: CR 45% (n = 13 of 29, including 8 of 23 with <i>FLT3</i> WT and 5 of 6 with <i>FLT3</i>-mutant)</li> <li>• 50-mg cohort: CR 80% (n = 32 of 40, including 20 of 27 with <i>FLT3</i> WT and 12 of 13 with <i>FLT3</i>-mutant)</li> <li>• <i>FLT3</i>-mutant cohort: 1-y OS of 0.85 (95% CI, 0.65–1.0); 2-y OS of 0.62 (95% CI, 0.35–0.88); 1-y DFS of 0.50 (95% CI, 0.22–0.78)</li> <li>• <i>FLT3</i> WT cohort: 1-y OS of 0.78 (95% CI, 0.62–0.93); 2-y OS of 0.52 (95% CI, 0.33–0.71) in <i>FLT3</i> WT; 1-y DFS of 0.60 (95% CI, 0.39–0.81)</li> </ul>
Midostaurin <sup>56</sup>	18–60 yo	+ any activating <i>FLT3</i> mutation	III: M 50 mg po bid vs placebo after 7 + 3 for induction, after HiDAC for consolidation, and as maintenance	<p>M vs placebo:</p> <ul style="list-style-type: none"> <li>• CR: 59% vs 54%; <i>P</i> = .18</li> <li>• 5-y OS: median 74.7 mo vs 26.0 mo, HR 0.77 (1-sided; <i>P</i> = .007)</li> <li>• 5-y EFS: median 8.0 mo vs 3.0 mo, HR 0.80 (1-sided; <i>P</i> = .004)</li> </ul>
Midostaurin <sup>57</sup>	18–70 yo	+ <i>FLT3</i> -ITD mutation	II: M 50 mg po bid after 7 + 3 for induction, after HiDAC for consolidation, and as maintenance after chemo or allo-HCT	<ul style="list-style-type: none"> <li>• Overall CR 75% after induction</li> </ul>

# 10<sup>th</sup> EDITION Highlights from EHA

Midostaurin <sup>55</sup>	18–60 yo	+/- <i>FLT3</i> mutation	Ib: M 50–100 mg, po bid, either concomitantly or sequentially with 7 + 3, M with HiDAC consolidation and M alone as maintenance	<ul style="list-style-type: none"> <li>• 100-mg cohort: CR 45% (n = 13 of 29, including 8 of 23 with <i>FLT3</i> WT and 5 of 6 with <i>FLT3</i>-mutant)</li> <li>• 50-mg cohort: CR 80% (n = 32 of 40, including 20 of 27 with <i>FLT3</i> WT and 12 of 13 with <i>FLT3</i>-mutant)</li> <li>• <i>FLT3</i>-mutant cohort: 1-y OS of 0.85 (95% CI, 0.65–1.0); 2-y OS of 0.62 (95% CI, 0.35–0.88); 1-y DFS of 0.50 (95% CI, 0.22–0.78)</li> <li>• <i>FLT3</i> WT cohort: 1-y OS of 0.78 (95% CI, 0.62–0.93); 2-y OS of 0.52 (95% CI, 0.33–0.71) in <i>FLT3</i> WT; 1-y DFS of 0.60 (95% CI, 0.39–0.81)</li> </ul>
Midostaurin <sup>56</sup>	18–60 yo	+ any activating <i>FLT3</i> mutation	III: M 50 mg po bid vs placebo after 7 + 3 for induction, after HiDAC for consolidation, and as maintenance	<p>M vs placebo:</p> <ul style="list-style-type: none"> <li>• CR: 59% vs 54%; <i>P</i> = .18</li> <li>• 5-y OS: median 74.7 mo vs 26.0 mo, HR 0.77 (1-sided; <i>P</i> = .007)</li> <li>• 5-y EFS: median 8.0 mo vs 3.0 mo, HR 0.80 (1-sided; <i>P</i> = .004)</li> </ul>
Midostaurin <sup>57</sup>	18–70 yo	+ <i>FLT3</i> -ITD mutation	II: M 50 mg po bid after 7 + 3 for induction, after HiDAC for consolidation, and as maintenance after chemo or allo-HCT	<ul style="list-style-type: none"> <li>• Overall CR 75% after induction</li> </ul>

# 10<sup>th</sup> EDITION Highlights from EHA



Pts 346 ( 2569 samples)



Clinical trial (preferred)

or

Standard-dose cytarabine 100–200 mg/m<sup>2</sup> continuous infusion x 7 days with idarubicin 12 mg/m<sup>2</sup> or daunorubicin 60–90 mg/m<sup>2</sup> x 3 days<sup>rr,ss</sup> (category 1)

or

Standard-dose cytarabine 200 mg/m<sup>2</sup> continuous infusion x 7 days with daunorubicin 60 mg/m<sup>2</sup> x 3 days and cladribine 5 mg/m<sup>2</sup> x 5 days (category 2A)<sup>tt</sup>

or

Age<sup>nn,oo</sup> <60 y ▶ High-dose cytarabine (HiDAC)<sup>ss,uu</sup> 2 g/m<sup>2</sup> every 12 hours x 6 days<sup>vv</sup> or 3 g/m<sup>2</sup> every 12 h x 4 days<sup>ww</sup> with idarubicin 12 mg/m<sup>2</sup> or daunorubicin 60 mg/m<sup>2</sup> x 3 days (1 cycle) (category 1 for patients ≤45 y, category 2B for other age groups)

or

AML ≥60 y  
[See AML-11](#)  
Standard dose cytarabine 200 mg/m<sup>2</sup> continuous infusion x 7 days with daunorubicin 60 mg/m<sup>2</sup> x 3 days and oral midostaurin 50 mg every 12 hours, days 8-21<sup>xx</sup> (FLT3-mutated AML)

or

Fludarabine 30 mg/m<sup>2</sup> IV days 2–6, HiDAC 2 g/m<sup>2</sup> over 4 hours starting 4 hours after fludarabine on days 2–6, idarubicin 8 mg/m<sup>2</sup> IV days 4–6, and G-CSF SC daily days 1–7 (category 2B)<sup>yy</sup>

# Risk Stratification and Treatment Selection

---

- Genetic risk informs likelihood of responding to intensive chemotherapy
- Performance status and comorbidities (and possibly age) inform likelihood of treatment benefit outweighing risk

## Induction

**All risk groups** (patients considered eligible for intensive chemotherapy)

**Treatment:** 7+3 chemotherapy

## Consolidation

**Favorable risk:** IDAC chemotherapy (regimen adjusted for patient age)

**Intermediate risk, age 18-60/65:**  
IDAC or allo-HSCT

**Intermediate risk, age >60/65:**  
Allo-HSCT

**Adverse risk:** Allo-HSCT

## Risk Stratification and Treatment Selection

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Allo-HSCT

**Adverse risk:** Allo-HSCT

# CPX-351 vs 7+3

## Older ND High-Risk (Secondary) AML, Phase 3

**CPX-351: liposome-encapsulated 5:1 fixed molar ratio of cytarabine:daunorubicin**

### Key Eligibility

- Secondary AML
- Newly diagnosed secondary AML\*
- Ages 60-75 years
- Able to tolerate intensive therapy
- PS 0-2

**CPX-351**  
n = 153

**7+3**  
n = 156

Induction  
(1-2 cycles)

Patients in CR or CRi:  
Consolidation  
(1-2 cycles)

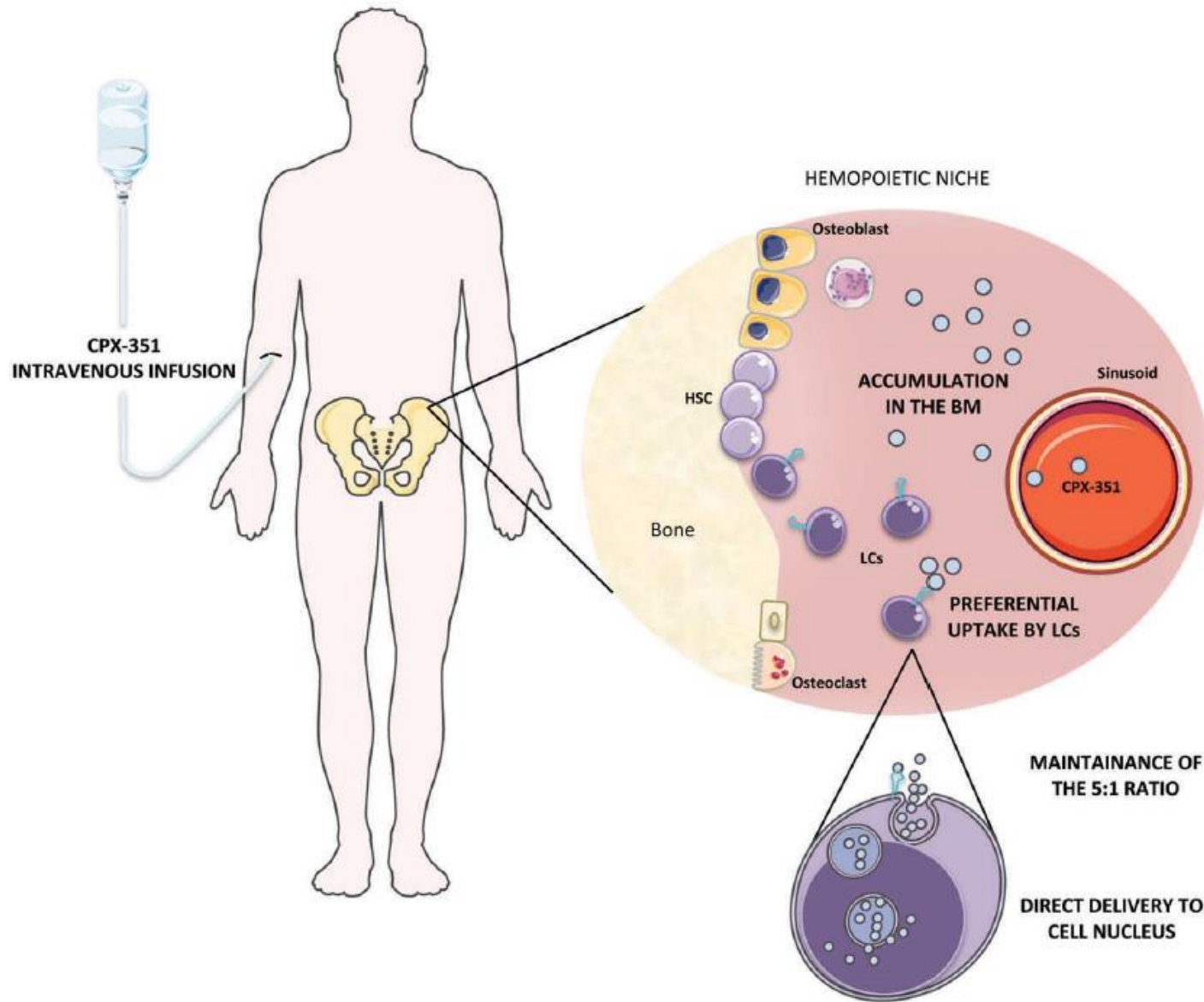
Follow-up:  
Death  
OR  
5 years

Primary Endpoint: OS

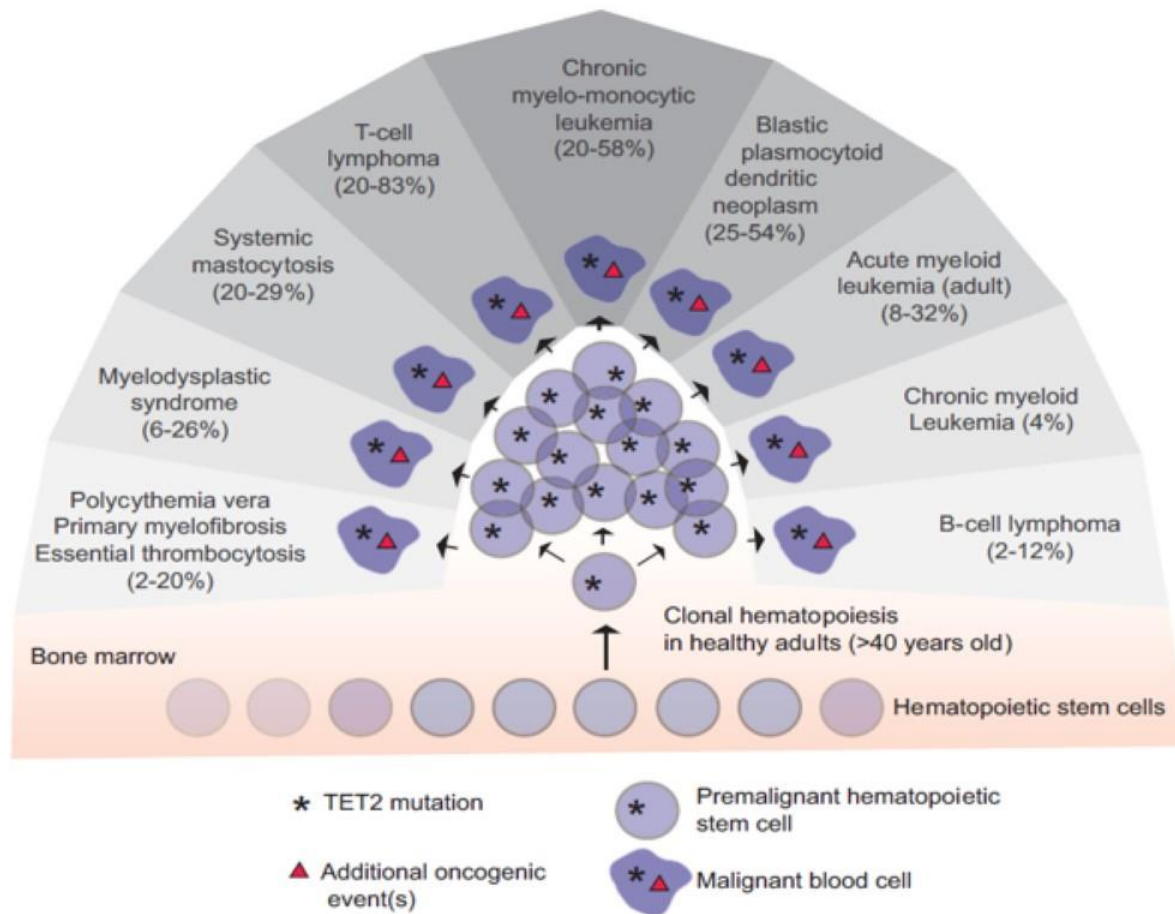
\*Secondary AML defined as having a history of prior cytotoxic treatment, antecedent MDS +/- a prior history of HMA, or AML with WHO-defined MDS-

	CPX-351 n = 153	7+3 n = 156	HR; P Value
Median OS, mo	9.56	5.95	.69; .005
Median EFS, mo	2.53	1.31	.74; .021
CR, %	37.3	25.6	.040
CR + CRi, %	47.7	33.3	.016

# CPX-351 in AML



# TET2 Mutation in Hematologic Malignancies

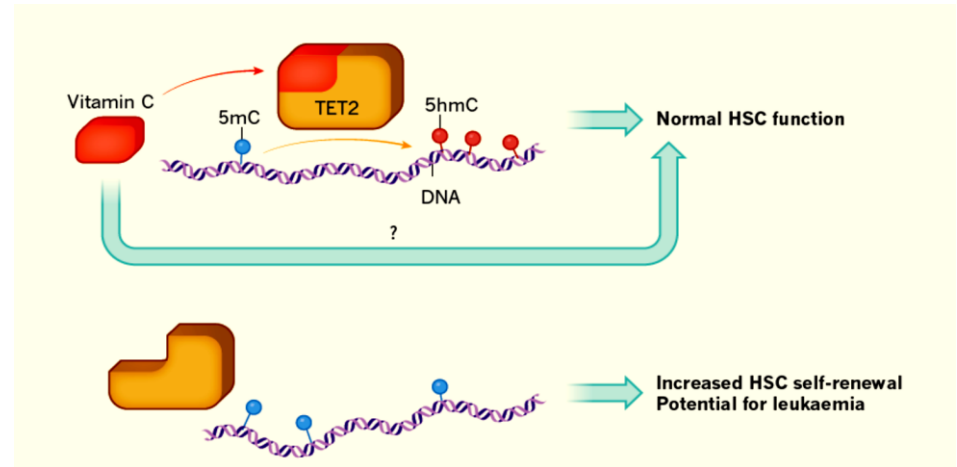
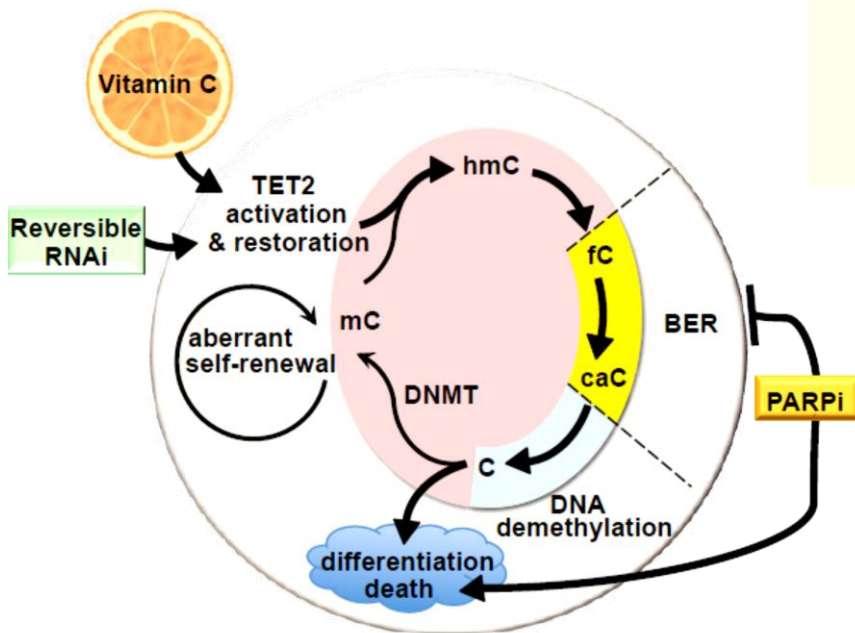


- A somatic mutation in *TET2* results in premalignant hematopoiesis and clonal expansion
- Additional oncogenic events cooperate with the initial *TET2* mutation to drive the onset of a wide variety of hematopoietic malignancies

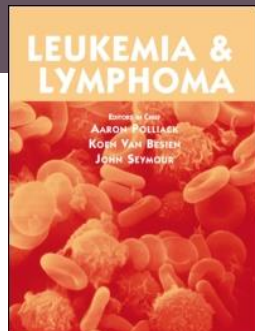
# Role of TET2 in DNA Methylation

- TET2 is one of a family of TET proteins that catalyzes the hydroxylation of 5-methyl cytosine, promoting hypomethylation of DNA
- Precise regulation of DNA methylation patterns is important for normal development
  - Methylated DNA provides protection against cellular transformation
- *TET2* mutation and altered gene expression is common in myeloid neoplasms
- *TET2* mutation is common in MDS and AML
  - Often "first hit" founder mutations in cancer development

## Restoration of TET2 Function Blocks Aberrant Self-Renewal and Leukemia Progression



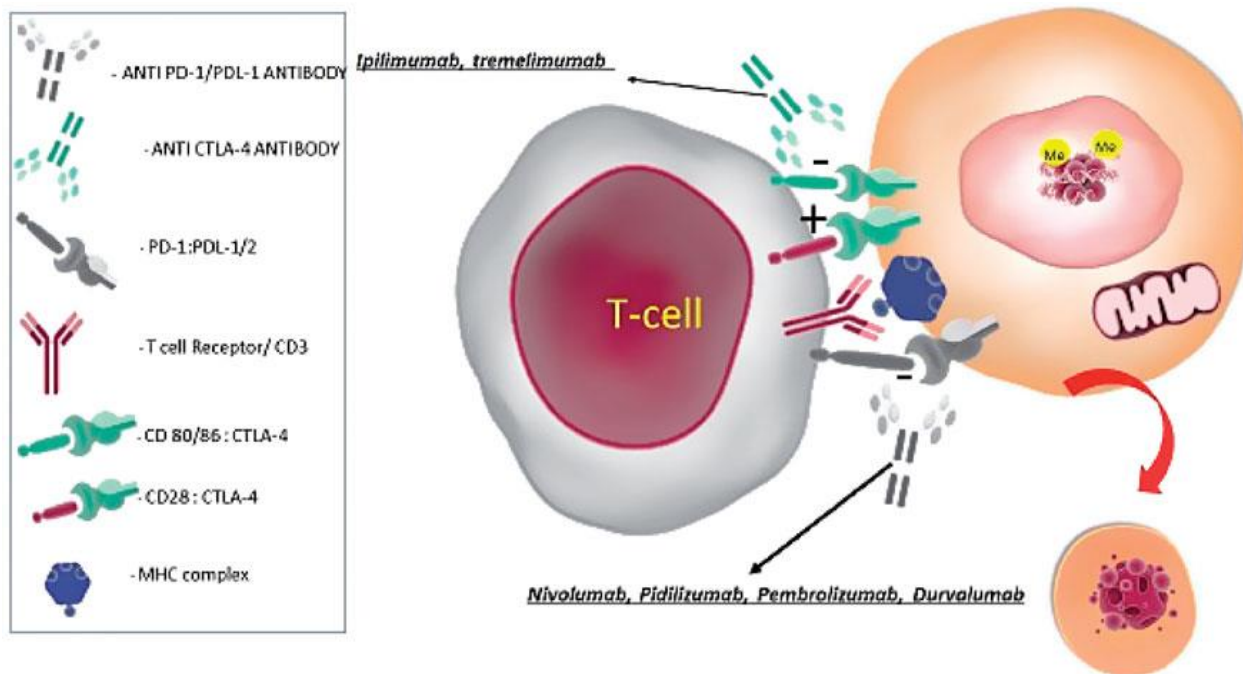




REVIEW

## The emerging role of immune checkpoint based approaches in AML and MDS

Prajwal Boddu<sup>a</sup> , Hagop Kantarjian<sup>a</sup>, Guillermo Garcia-Manero<sup>a</sup>, James Allison<sup>a</sup>, Padmanee Sharma<sup>b</sup> and Naval Daver<sup>a</sup>



10<sup>th</sup> EDITION  
**Highlights** from EHA

