19th Post-ASH Meeting

Acute Leukemia

Prof. Carlos Graux CHU UCL Namur (Godinne)

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Dinant • Godinne • Sainte-Elisabeth



The decision-making process in AML

Clinical evaluation

Genetic informations

o MRD data



- Do I treat my patient with an intensive (= possibly curative) versus a non intensive (= probably never curative) approach ?
- > Do I have therapeutic targets to improve the current treatment ?
- Is there an indication for SCT, maintenance therapy, ...





The headhache of prognostication

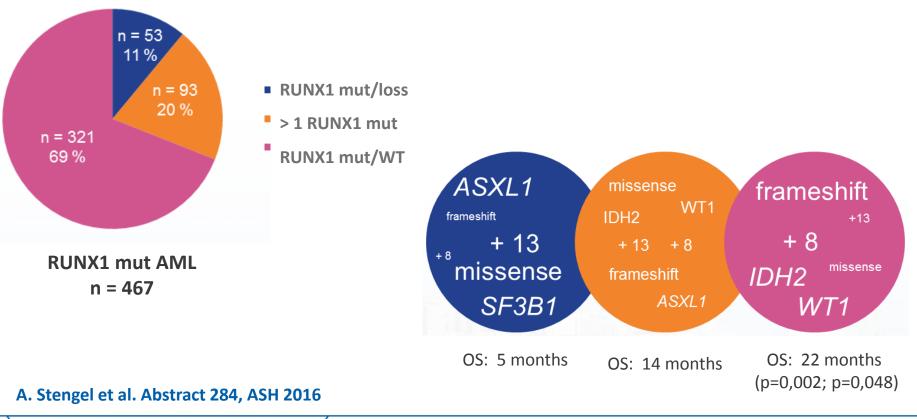
<u>RISK STATUS</u>	<u>CYTOGENETICS</u>	MOLECULAR ABNORMALITIES
Favorable-risk ⁵	Core binding factor: inv(16) ^{2,3} or t(16;16) ² or t(8;21) ² t(15;17)	Normal cytogenetics: NPM1 mutation in the absence of FLT3-ITD or isolated biallelic CEBPA mutation
Intermediate- risk	Normal cytogenetics +8 alone t(9;11) Other non-defined	
Poor-risk	Complex (\geq 3 clonal chromosomal abnormalities) Monosomal karyotype -5, 5q-, -7, 7q- 11q23 - non t(9;11) inv(3), t(3;3) t(6;9) t(9;22) ⁴	Normal cytogenetics: with FLT3-ITD mutation ⁶ TP53 mutation « AML with mutated RUNX1 »



« AML with mutated RUNX1 » ?

Critical transcription factor involved in HSC differentiation

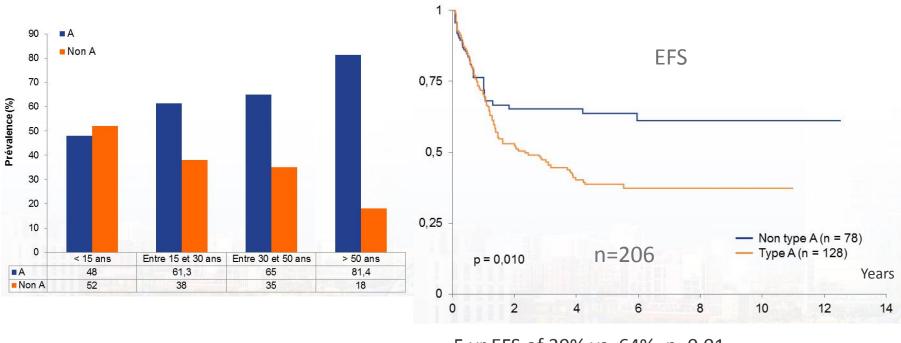
10-15% AML, associated with therapy-related disease, older age, increased cytopenias, shorter survival and chemoresistance





NPM1 insertion type do matter

Frameshift mutations are the result of 4 base pair (bp) insertions in exon 12 There are several genomic subsets of insertions, type A variant (insertion of TCTG) being the most common



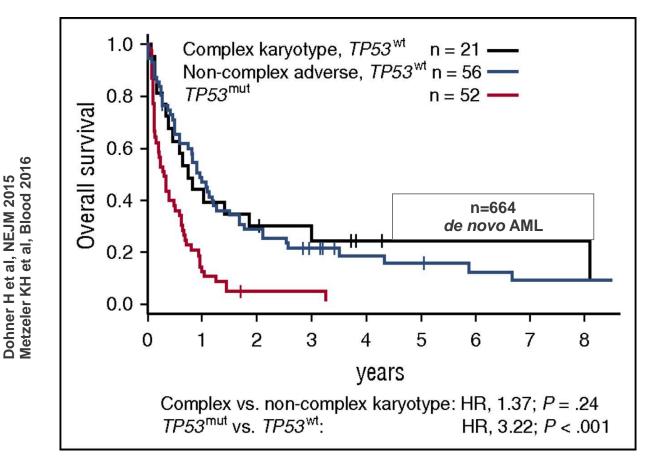
5 yr EFS of 39% vs. 64%, p=0.01 5 yr RFS of 41% vs. 70%, p=0.002 ... regardless of age or FLT3/ITD status

D. Selim et al. Abstract 285, ASH 2016



TP53 mutations and prediction of response

Key TS involved in genomic stability, cell cycle regulation, metabolism and DNA repair Associated with advanced age, complex karyotype, chemo-resistance, and dismal survival





TP53 mutations and prediction of response

Prognostic value of TP53 mutation in the AZA-AML-001 study comparing azacytidine (AZA) vs conventional care regimen (CCR) in older patients with AML (Dombret et al, Blood 2015) The biomarker cohort comprised 156 of all 488 pts (32%; AZA n=83, CCR n=73)

Table. Median OS associated with mutated genes: AZA vs CCR					
Mutated Gene	Median OS AZA (mos)	Median OS CCR (mos)	P value		
TP53	7.2 (3.9, 18.6)	2.4 (1.5, 7.1)	0.069		
NRAS	11.8 (7.7, NR)	4.3 (2.3, NR)	0.047		
FLT3 (-ITD or -TKD)	5.4 (4.5, NR)	6.4 (3.8, NR)	0.17		
TET2	9.6 (4.5, 13.5)	11.1 (2.8, NR)	0.082		
IDH1	11.1 (1.3, NR)	14.6 (3.7, NR)	0.773		
IDH2	12.6 (4.4, NR)	12.5 (5.6, NR)	0.914		
DNMT3A	12.6 (7.0, 20.8)	10.3 (3.8, NR)	0.963		
Any DNA methylation gene (IDH1, IDH2, DNMT3A, TET2)	11.1 (5.8, 15.4)	12.5 (4.3,17.6)	0.497		

Similar observations have been reported for decitabine in TP53 mutated and unfavorable karyotype AML

Welch JS et al. NEJM 2016

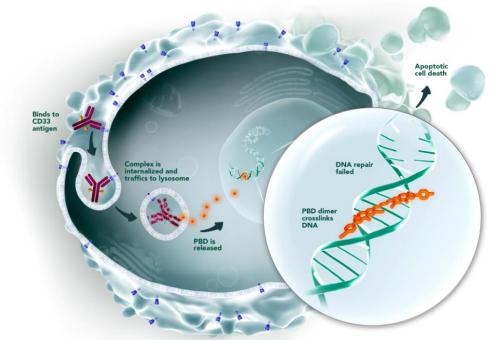
L. Tang et al. Abstract 2859, ASH 2016



The long parade of HMA combined therapies

HMA + vadastuximab talirine (33A)

New CD33-directed antibody-drug conjugate that links to 2 molecules of a pyrrolobenzodiazepine (PBD) dimer, a new agent that crosslinks DNA



Vadastuximab talirine (SGN-CD33A; 33A) is an investigational agent, and its safety and efficacy have not yet been established. ©2015 Seattle Genetics, Inc. All rights reserved.



Vadastuximab talirine + HMA

Open-label, phase I combination cohort study

Previously untreated CD33+AML pts $aged \ge 60 yrs$ ineligible for intensive induction/consolidation therapy(n=53)

Azacitadine 75 mg/m² SQ/IV x 7 days or Decitabine 20 mg/m² IV x 5 days + Vadastuximab talirine 10 µg/kg IV on last HMA day 4 x 4-wk cycles If clinical benefit, continue until recurrence

Follow-up every 3 months

AT. Fathi et al. Abstract 591, ASH 2016



Vadastuximab talirine + HMA

Outcome, %	Evaluable Pts (n = 49)	Secondary AML Pts [‡] (n = 22)	Pts with <i>FLT3/ITD</i> + AML (n = 5)	Pts Aged ≥ 75 Yrs (n = 26)
Remission rate (CR + CRi)	73	77	100	65
CR	47	50	80	38
CRi (p)*	20	18	20	19
CRi (n)†	6	9	0	8
mLFS	2	5	0	4
ORR (CR+CRi+mLFS)	76	82	100	69

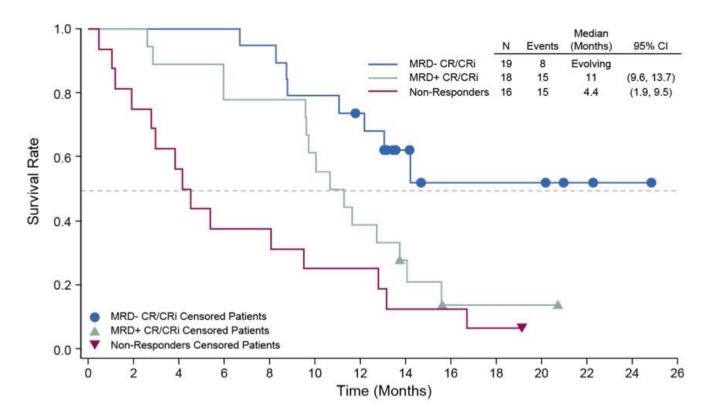
- > 50% of pts with response achieved MRD negativity by FCT
- Thrombocytopenia was the main toxicity. No VOD signals

AT. Fathi et al. Abstract 591, ASH 2016



Vadastuximab talirine + HMA

Overall Survival by MRD Status



AT. Fathi et al. Abstract 591, ASH 2016

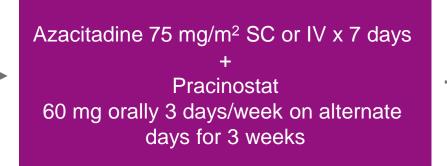


Pracinostat + AZA

Potent oral HDAC inhibitor expected to induce a reexpression of silenced genes in a synergistic fashion

Phase 2 study

Untreated *de novo* or secondary AML pts \geq 65 yrs ____ not eligible for induction (n=50)



If clinical benefit, continue until recurrence

G. Garcia Manero. Abstract 100, ASH 2016



Pracinostat + AZA

	CR Rate	cCR Rate	Survival (months) (Median, 95% CI)
Overall population (N = 50)	42%	52%	19.4 (10.0-NR)
Cytogenetics			
Intermediate (N = 27)	48.1%	59.3%	NR (10.7, NR)
High risk (N = 21)	38.1%	47.6%	13.5 (2.4, NR)
Age			
≥75 (N = 26)	42.3%	57.7%	13.5 (9.0, 21.5)
66-74 (N = 24)	41.7%	45.8%	26.5 (8.0, 26.5)
Type AML			
De novo (N = 33)	42.4%	51.5%	13.02 (5.7, 26.5)
Secondary (N = 17)	41.2%	52.9%	NR (>16.4, NR)
ECOG Performance Status			
0-1 (N = 42)	40.5%	50.0%	19.08 (10.0, 19.1)
2 (N = 8)	50.0%	62.5%	13.0 (8.0, 26.5)

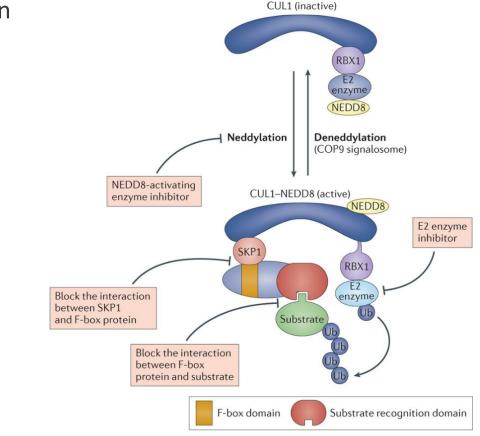
CI = Confidence Interval, NR = Not Reached

- cCR (CR/CRi/morphologic leukemia free state) was achieved in 52% (42%/4%/6%)
- Median duration of cCR was 13.2 months, CI [10.9-21.5]
- Median OS of 19.1 months CI [10.0-not reached, median follow-up of 21 months)



Pevonedistat + AZA

First-in-class NEDD8-activating enzyme inhibitor disrupting proteasome-mediated protein degradation



Nature Reviews | Drug Discovery



Pevonedistat + AZA

o Phase I

Older AML pts unlikely to benefit from standard induction therapy (n=61) Azacitadine 75 mg/m² SQ/IV x 7 days or Decitabine 20 mg/m² IV x 5 days + pevonedistat 20 or 30 mg/m IV on days 1, 3 and 5

- CR/CRi was obtain in 44% of response-evaluable pts (n=52).
- Median OS was 7.0 months for the MTD cohort
- There was limited additional toxicity

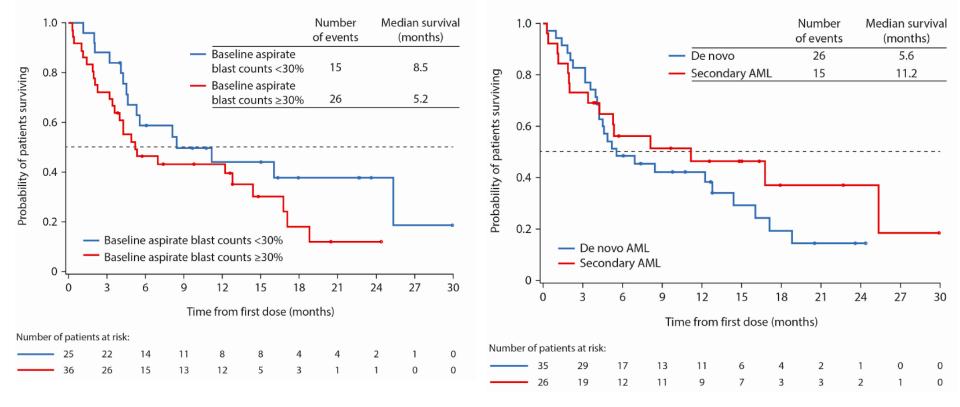
RT. Swordset al. Abstract 98, ASH 2016



Pevonedistat + AZA

Figure 2: Kaplan-Meier curve of overall survival for MTD cohort (n=61) by baseline aspirate blast counts subgroup (<30% vs \geq 30% marrow blasts)

Figure 3: Kaplan-Meier curve of overall survival for MTD cohort (n=61) by disease characteristic subgroup (de novo vs secondary AML)



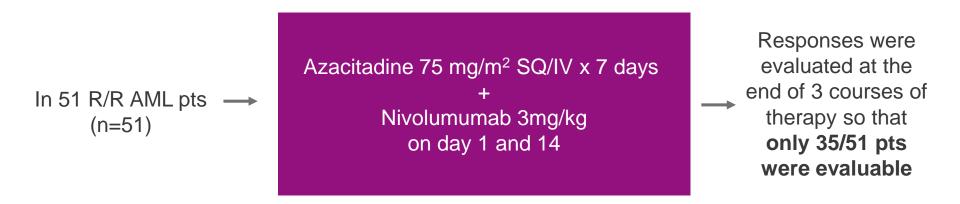
RT. Swordset al. Abstract 98, ASH 2016



Nivolumumab + AZA

Monoclonal antibody targeting programmed cell death protein 1 (PD-1) AZA upregulates PD-1 and PD-L1 in AML \rightarrow resistance to therapy

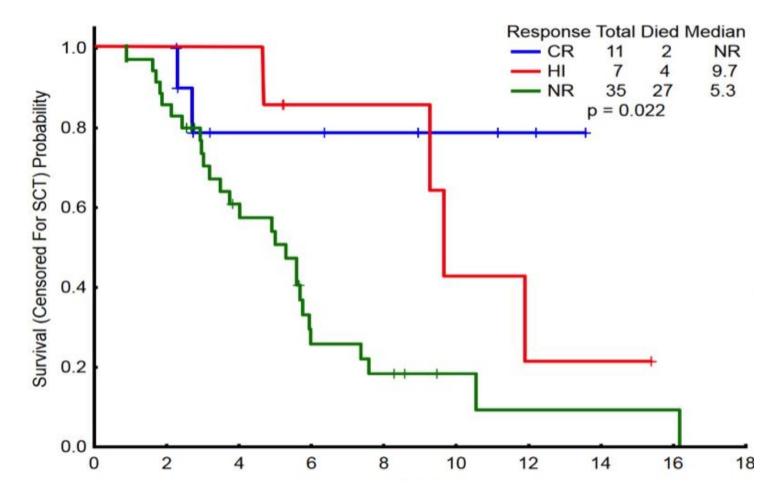
Phase IB/II study conducted at the MDACC



- Durable CR/CRi or HI in 18% and 15% respectively
- Median OS was 9.3 months (range, 1.8–14.3)
- Compares favorably to historical survival with AZA-based salvage protocols
- N. Daver et al. Abstract 763, ASH 2016



Nivolumumab + AZA



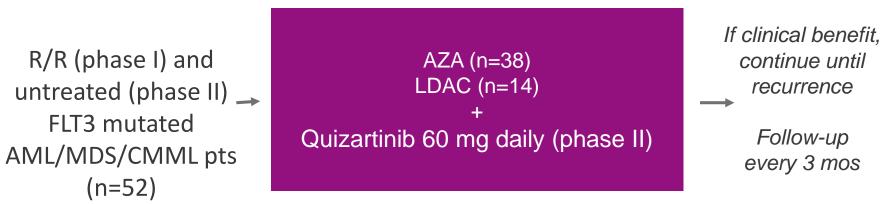
N. Daver et al. Abstract 763, ASH 2016

Quizartinib + HMA

Activating mutations of FLT3 occur in 30% of AML cases

Quizartinib is a potent, selective FLT3-ITD inhibitor, in vitro synergy with AZA/LDAC

• Phase I/II



- ➢ ORR of 67% (CR=8, CRp=7, CRi=18, PR=2); 23% for LDAC arm and 77% for AZA arm
- Median OS: 14.8 months (7.5 months for the LDAC arm and not reached for the AZA arm)
- Clinically significant QTcF prolongation was infrequent

Similar results with sorafenib + AZA (M. Ohanian et al. Abstract 1611. ASH 2016)

W. Abdelall et al. Abstract 1642, ASH 2016



CD123, the perfect myeloid target?

Immunotherapy works best when the disease load is reduced and is therefore particularly indicated to eliminate MRD

Leukemic stem cell (LSC) participates to MRD persistence and reappearance. The ideal target for immunotherapy has to be expressed on LSC surface

CD123 (IL-3 receptor α chain) has emerged as a very promising target for immunotherapeutic strategies in AML

- o aberrant expression frequently observed in a subset of leukemic disorders
- o increased CD123 expression is associated with a poor prognosis

N. Arai et al. Abstract 2887, ASH 2016



Blastic plasmacytoïd dendritic cell neoplasm

BPDCN is a rare myeloid entity

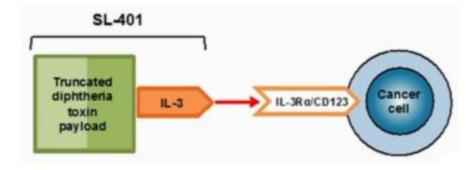
Caracterised by a particular CD4, CD56, CD123, TCL-1 expression profil, frequent skin involvement, sometimes as primary site

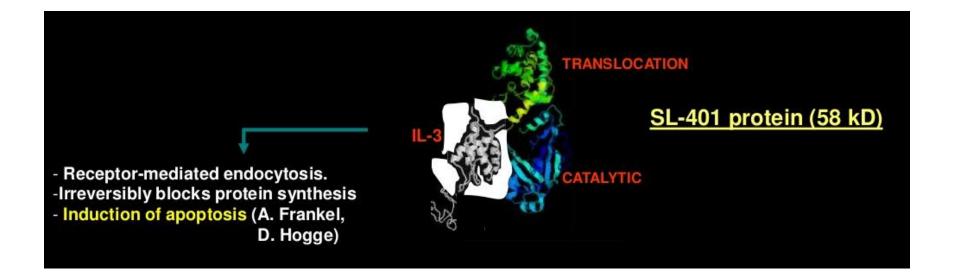
No standard of care. ALL type treatment usually proposed

OS is poor, arround 8-14 months.

CD123 is overexpressed in nearly 100% of pts with BPDCN

New strategies targeting CD123: SL-401







New strategies targeting CD123: SL-401

In pts with BPDCN

• Phase 2 trial ongoing expension stage of SL-401

Daily IV infusion dose of 12 μg/kg CR/CRc: 81% (13/16) when used in 1ste line and 31% (4/13) in R/R setting 7/32 pts underwent allo or auto

In pts with AML in CR1 or CR2 with high risk of relapse

Multicenter, single-arm phase 2 trial

Lead-in stage 1 completed without DLT or MTD, and stage 2 (expansion) still open Safety profile similar to that observed in other SL-401 clinical studies Potential to reduce this chemo-resistant cell population and to offer improved long-term outcomes

> N. Pemmaraju et al. Abstract 342, ASH 2016 AA. Lane et al. Abstract 215, ASH 2016



Preclinical studies targeting CD123

• IMGN632

- another CD123-directed antibody-drug conjugate
- o conjugation with indolinobenzodiazepine dimers for DNA alkylation

o JNJ-63709178

- anti-CD3/CD123 bispecific humanized lgG4
- recruits CD3+ T cells to CD123+ tumor cells and induces their killing

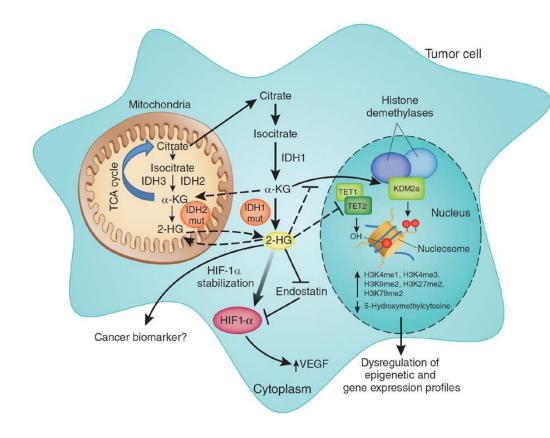
$\,\circ\,$ Allogeneic TCR α/β deficient CAR T-cell targeting CD123

prolongs OS of AML and BPDCN patient-derived xenografts

Y. Kovtun et al. Abstract 768, ASH 2016 F. Gaudet et al. Abstract 2824, ASH 2016 R. Mani et al. Abstract 580, ASH 2016 T. Cai et al. Abstract 4039, ASH 2016



IDH1/2 inhibitors



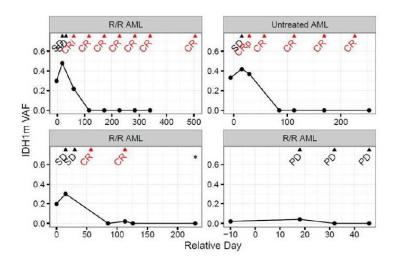
- IDH mutations in ~ 20% of AML, in founding clone
 - 6-10% IDH1-R132
 - 9-15% IDH2-R140 or IDH2-R172
 - Associate with NPM1 and FLT3 mutations, CN-AML
- Accumulation of 2HG competitively inhibits αKG
 - Hypermethylated phenotype via silencing of TET family
 - Block in differentiation via inhibition of histone demethylases
 - Altered hypoxic response
 - BCL2 dependence



AG-120

AG-120, a first-in-class, oral, potent, reversible, selective inhibitor of the IDH1 mutant enzyme is under evaluation in multiple ongoing single agent and combination clinical trials

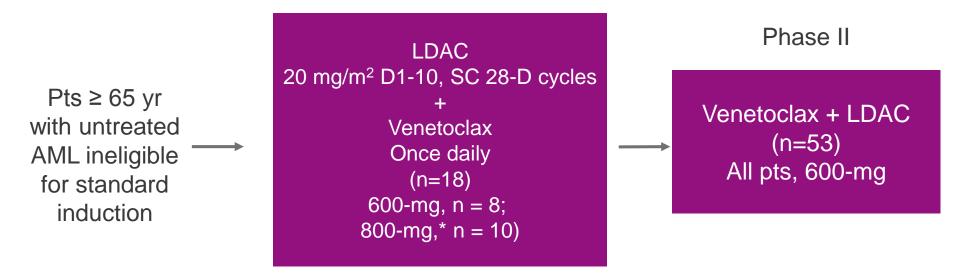
• Courtney D. DiNardo presented the first demonstration that treatment with single agent AG-120 can result in mutated IDH1 clearance as determined by NGS



 Studies with the IDH2 inhibitor, enasidenib (AG221) and the pan IDH1/IDH2 inhibitor AG881 are also ongoing
 CD. DiNardo et al. Abstract 1070, ASH 2016



• Phase I (3+3 design)



*2 pts had dose-limiting toxicity at the 800-mg dose level.

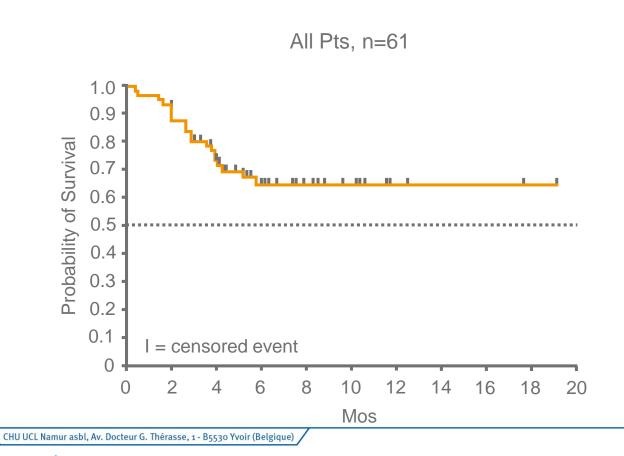


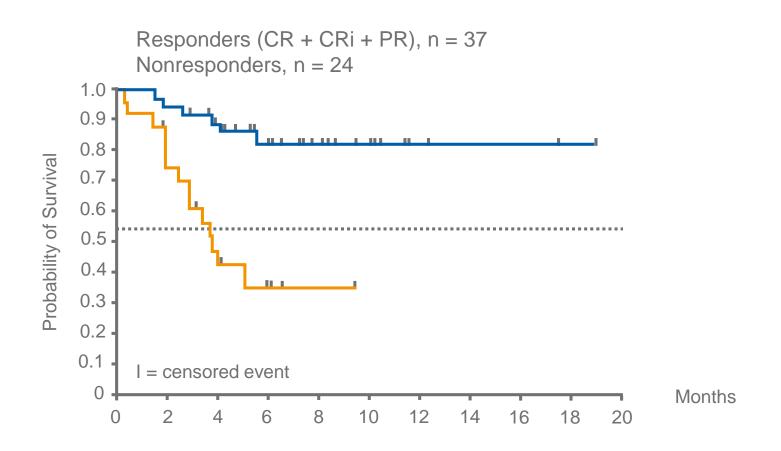
Overall Response, %	Venetoclax 600 mg (n=61)
CR	21
CRi	33
CR + CRi*	54
PR	7
ORR (CR + CRi + PR)	61
Resistant/progressive disease	38

*70% of CR + CRi achieved during cycle 1 or cycle 2.

- Most frequent TRAEs (any grade) included nausea (72%), hypokalemia (46%), diarrhea (44%), fatigue (43%), decreased appetite (41%)
- Most frequent grade 3/4 TRAEs included febrile neutropenia (34%), hypokalemia (15%), hypophosphatemia (13%), hypertension (10%)







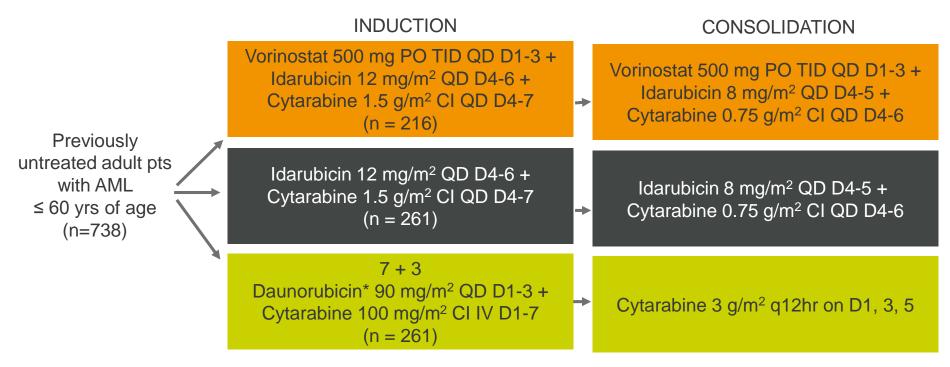
NAMUR



Induction therapy : less is more ?

High response rate of idarubicin (ida) and high-dose ara-C (IA) in combination with the HDAC inhibitor vorinostat (IA+V) compared to IA or 7+3 (Garcia-Manero et al. JCO 2011)

• Randomized phase III superiority trial of the added value of vorinostat at induction



G. Garcia-Manero et al. Abstract 901, ASH 2016



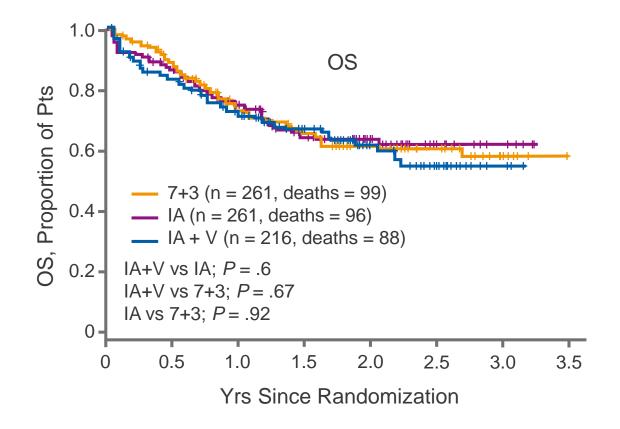
No added value of vorinostat at induction

Outcome, %	Vorinostat + Ida + Ara-C (n = 216)	lda + Ara-C (n = 261)	7 + 3 (n = 261)	All Pts (N = 738)	<i>P</i> Value
CR	60	64	63	62	.58
CRi	17	16	13	15	
Failure	23	21	25	23	
Mortality					
30 day	4	6	3	4	.013
60 day	9	9	5	7	.097

G. Garcia-Manero et al. Abstract 901, ASH 2016



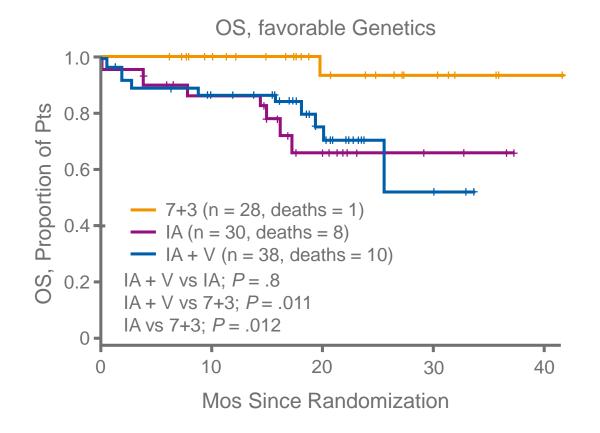
No added value of vorinostat at induction



G. Garcia-Manero et al. Abstract 901, ASH 2016



No added value of vorinostat at induction



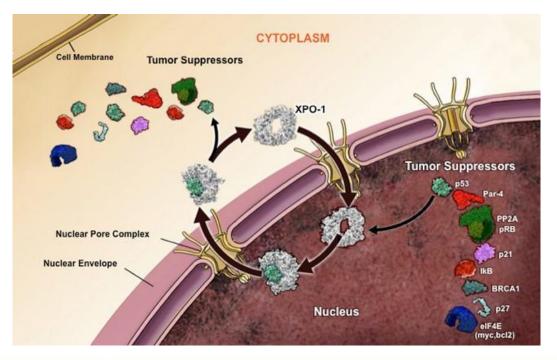
G. Garcia-Manero et al. Abstract 901, ASH 2016



Selinexor combined with 7+3

By inhibiting the primary export protein, XPO1, selinexor maintains tumor suppressor proteins to the nucleus leading to their activation and an anti-leukemic effect as a single agent

It inhibits DNA damage repair, rationalizing its use in combination with DNA damaging agents





Selinexor combined with 7+3

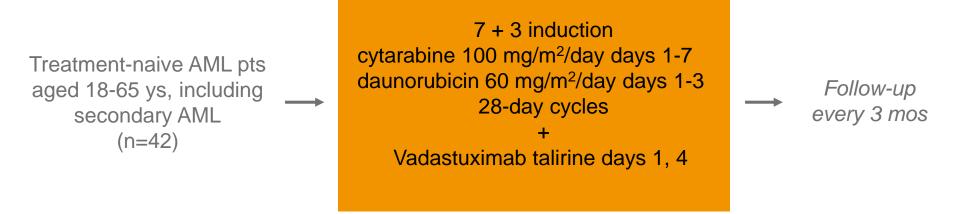
- Phase I of Selinexor in combination with daunorubicin and cytarabine in newly diagnosed poor-risk AML patients
- Phase II of Ara-C and Idarubicin in combination with Selinexor (KPT-330) in R/R AML
- Both indicate that the combination can be safely administered. The encouraging activity presents as an interesting bridge to transplant

KL. Sweet et al. Abstract 4040, ASH 2016 W. Fiedler et al. Abstract 34, ASH 2016



Vadastuximab talirine combined with 7+3

Phase Ib dose escalation/expansion study



dose escalation/expansion from 10 + 10 μ g/kg to 20 + 10 μ g/kg per SMC

HP. Erba et al. Abstract 211, ASH 2016



7+3 combined with vadastuximab talirine

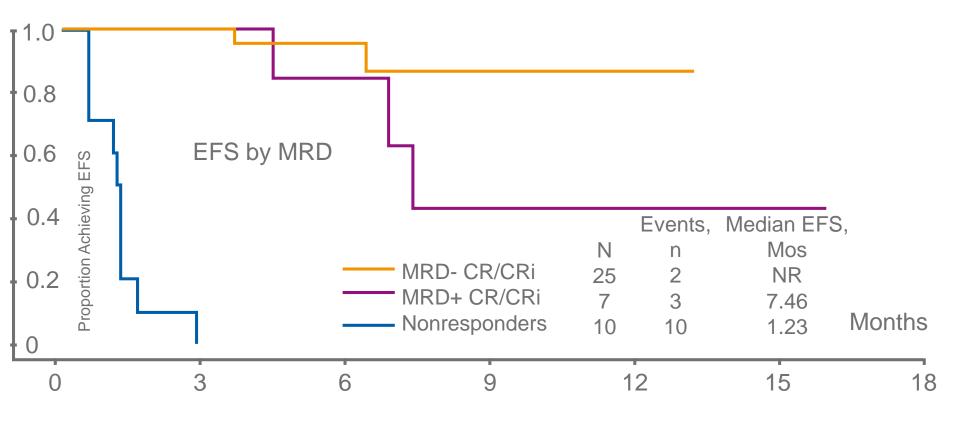
Response, %	CR	CRi*	CRc (CR + CRi)
Evaluable pts (n=42)	60	17	76
Cytogenetic risk by MRC Favorable (n = 5) Intermediate (n = 21) Adverse (n = 15)	100 67 40	0 19 20	100 86 60

*All pts with CRi had CR with ANC \geq 1000/uL, incomplete platelet recovery.

- > 78% (25/32) of pts in CRc were MRD negative
- Similar AE rate to 7 + 3 alone
- > 2% 30-day mortality rate
- > 50% (21/42) of pts received allo-SCT

HP. Erba et al. Abstract 211, ASH 2016

7+3 combined with vadastuximab talirine



HP. Erba et al. Abstract 211, ASH 2016

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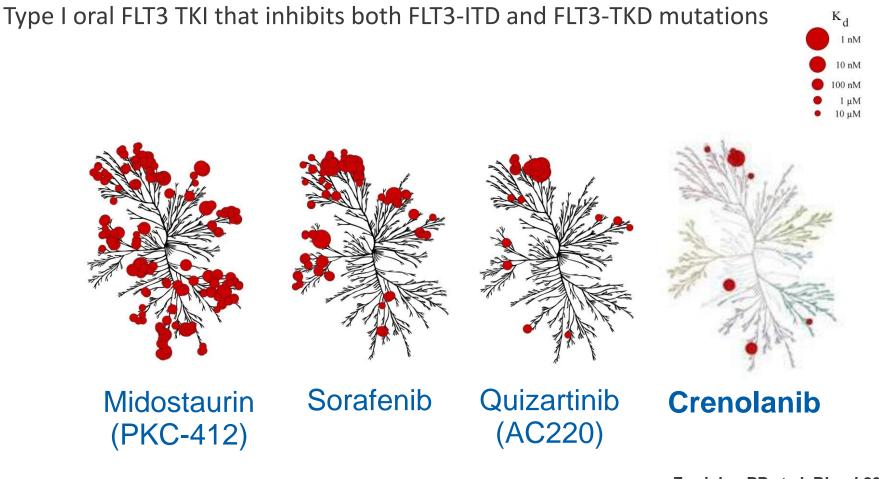
7+3 combined with vadastuximab talirine

Characteristics	Vadastuximab talirine + 7 + 3 SWOG Eligible* (n=30)	7 + 3 ± GO SWOG 0106 [†] (n=595)
Median age, yrs	45	48
Adverse cytogenetic risk, %	27	23
CRc rate (CR + CRi), %	80	75
CRc with 1 cycle, %	77	60
MRD-negative CRc, %	73	~ 54‡

*SWOG eligibility: 60 yrs of age or younger, de novo only. [†]Aggregate data from Othus M, et al. Leukemia. 2016;30:2080-2083. [‡]Calculated MRD-negative rate

HP. Erba et al. Abstract 211, ASH 2016

7+3 combined with crenolanib



Zarrinkar PP et al, Blood 2009 Galanis A et al, Cancer Res 2012



7+3 combined with crenolanib

Patients with newly diagnosed FLT3 mutant AML

Induction Chemotherapy Regimen	# Evaluable Patients	CR after 1 Cycle of Induction	CR after Additional Cycle of Chemotherapy*	Overall Complete Response
Cytarabine + Daunorubicin (n=18)	18	16	1	17/18 (94%)
Cytarabine + Idarubicin (n=8)	7	6	1	7/7 (100%)
TOTAL (n=26 pts)	25	22	2	24/25(96%)

*1 pt re-induced with cytarabine/idarubicin and 1 pt received HiDAC

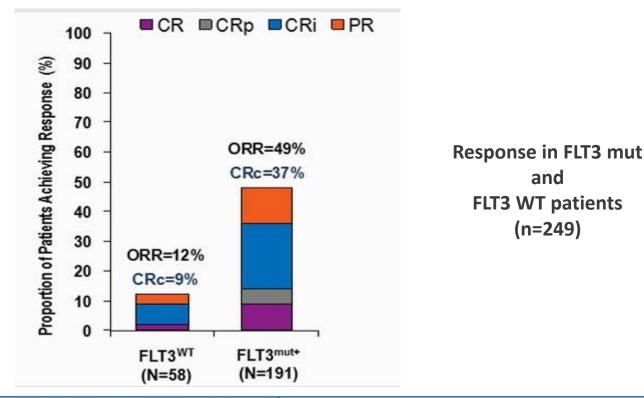


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Gilteritinib (ASP2215): the winner ?

Novel, highly potent and selective oral FLT3 inhibitor with preclinical activity against FLT3-ITD activating and FLT3-D835 resistance mutations

 First-in-human phase 1/2 study (CHRYSALIS) of once-daily oral gilteritinib given to pts with R/R AML irrespective of FLT3 mutation status

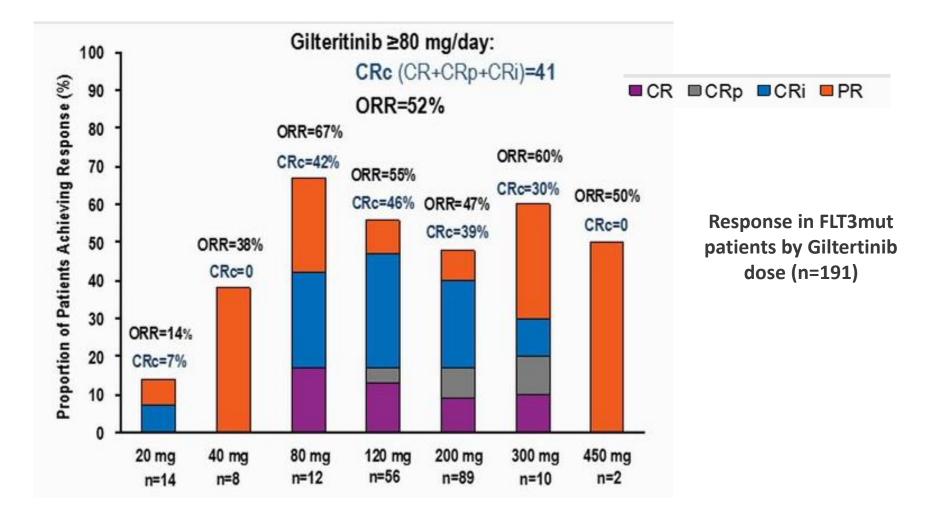


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Gilteritinib (ASP2215): the winner ?

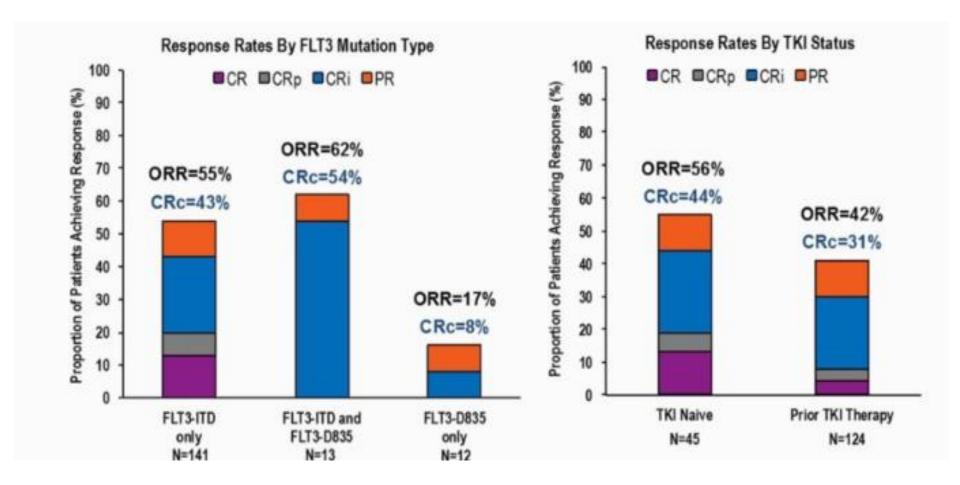


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Gilteritinib (ASP2215): the winner ?



CHU



The good place for ponatinib in Ph+ ALL?

T315I mutations commonly associated with relapse to TKI Potent BCR-ABL inhibitor effective against T3151 mutant ALL

Single-center, open-label phase II trial \bigcirc

8 x 21-day cycles

Pts ≥ 18 yrs with untreated (n=50) or previously treated (n=8*) Ph+ ALL *2 non responders, 6 in CR	Hyper-CVAD: cycle 1-3-5-7 cyclophosphamide (300 mg/m ² x 6; D1-3) vincristine (2 mg, D1, 11), doxorubicin (50 mg/m ² . D4), dexamethasone (40 mg, D1-4, D11-14) MTX-Ara-C: cycle 2-4-6-8 methotrexate (12 mg), cytarabine (3 mg/m ² , D2-3) Ponatinib: cycle 1: 45 mg for d1-14 cycle 2-8: 45 mg daily Due to vascular toxicity at 45 mg, protocol amended to allow 30 mg daily beyond induction, then 15 mg daily in pts achieving CMR !!!!!!! Rituximab given for first 4 courses, if CD20+!!!!!!!	Pts with CR received maintenance therapy of daily ponatinib + monthly vincristine and prednisone or hyper-CVAD for 2 yrs
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Sasaki K, et al. ASH 2016. Abstract 757

CHU UCL Namur asbl, Av. Docteur G. Thérasse, 1 - B5530 Yvoir (Belgique)



The good place for ponatinib?

Response, n/N (%)	Pts
CR*	52/52 (100)
CCyR [†]	48/48 (100)
CyR after induction Complete Minor Not done 	44/48 (92) 2/48 (4) 2/48 (4)
CyR after second cycle Complete 	48/48 (100)
MMR	56/58 (97)
MMR after induction	31/48 (65)
CMR	46/58 (79)

*6 pts in CR at enrollment

Sasaki K, et al. ASH 2016. Abstract 757

- No early deaths on study
- Median time to
 - MMR: 3 wks (2-14)
 - o CMR: 10 wks (2-96)
- Median follow-up: 39 mos (range: 2-59)
 - o 10 pts received alloSCT
 - 21 pts (47%) remain on ponatinib
 - 11 pts (24%) switched to other TKI



The good place for ponatinib?

Outcome, %	Pts (N = 58)
3-yr CRD	78
3-yr OS	75
Landmark analysis at 4 mos by ASCT	
3-yr CRD* ■ ASCT ■ No ASCT	88 75
3-yr OS [†] ■ ASCT ■ No ASCT	79 86
*P = .36 †P = .81 Sasaki K, et al. ASH 2016. Abstract 757	

The good place for ponatinib?

AE, % (N = 58)
Infections during induction 52
ALT/AST increase 31
Bilirubin increase 17
Pancreatitis 17
Skin rash 16
Amylase/lipase 16
Hypertension 14
Hemorrhage 10
Mucositis 9
Abdominal pain 7
Thrombotic events 7
Myocardial infarction* 5

%

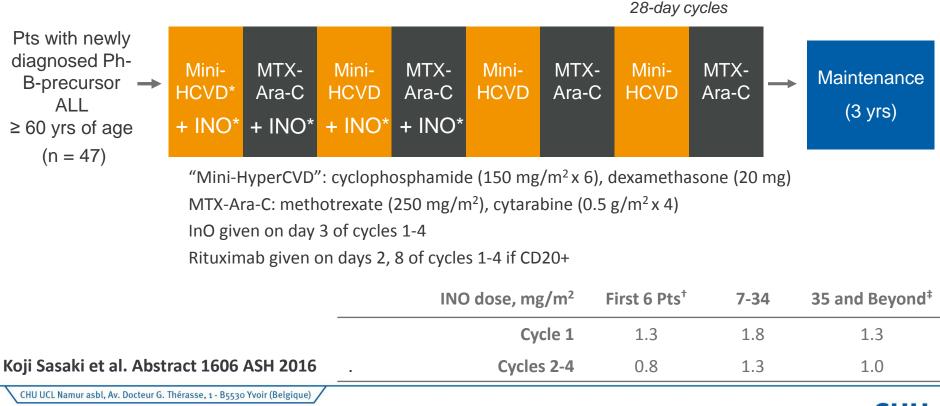
Sasaki K, et al. ASH 2016. Abstract 757

*Since protocol amendment, 1 pt with grade 2 angina possibly related to treatment



Inotuzumab Ozogamicin (InO), an anti-CD22 antibody-calicheamicin conjugate Superior clinical activity versus standard of care (SOC; intensive chemotherapy) R/R B-ALL in the phase 3 INO-VATE trial (Kantarjian. *N Engl J Med*. June 12, 2016)

• Single-center, open-label phase I/II trial



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Dinant • Godinne • Sainte-Elisabeth



Response, %	INO + Mini- HCVD (N = 43)
ORR CR CRp CRi	98 84 12 2
Cytogenetic CR*	100
 MRD negativity D21[†] Overall[‡] 	76 96

No early deaths on study

Median follow-up: 24 mos (range: 1-55)

• 3-yr CRD: 72%

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Grade 3/4 AEs in ≥ 10% of Pts, %	INO + Mini- HCVD (N = 47)
Prolonged thrombocytopenia	79
Infection (consolidation)	74
Infection (induction)	53
Hyperglycemia	53
Hypokalemia	34
ALT/AST elevation	19
Bilirubin elevation	17
Hemorrhage	15

- Thrombocytopenia > 6 wks in 37 pts (79%), 8 in induction, 36 in subsequent courses
- 4 pts experienced VOD after 2-4 cycles of INO
 - All received 1.8 mg/m² dose in cycle 1

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- Combination of inotuzumab ozogamicin with reduced-intensity mini-HCVD highly effective as frontline treatment in older pts with ALL
 - ORR: 98%
 - Low early mortality rate
- 54% 3-yr OS compares favorably with 31% 3-yr OS reported with HCVAD ± rituximab
- Manageable safety profile with inotuzumab ozogamicin plus mini-HCVD
 - VOD events in 4 pts led to inotuzumab ozogamicin dose reduction
- Investigators suggest inotuzumab ozogamicin + mini-HCVD may become a new frontline standard of care for older pts with ALL

Koji Sasaki et al. Abstract 1606 ASH 2016



Conclusion

- The effective translation of research-based innovations into the treatment of our AML patients will require collaborative efforts of accademics and companies to organise big trials that provide personalized treatment decisions, based on genomic informations
- It'll imply the characterisation of all known mutations in AML at the time of diagnosis and their monitoring during therapy
- NGS open new perspectives for clono specific evaluation of MRD

P. Hirsch et al. Abstract 1208, ASH 2016



