Midostaurina: La primera terapia dirigida aprobada para el tratamiento de la LMA FLT3+

Dr. Jorge Sierra Servicio de Hematología Hospital de la Santa Creu i Sant Pau Universidad Autónoma de Barcelona

Puerto Varas, Chile, 9-10 de Agosto de 2018

Disclosures

- Research grants:
 - Celgene, Amgen, Novartis
- Advisory board:
 - Celgene, Janssen, AbbVie, Daiichi-Sankyo, Pfizer, Novartis, Jazz Pharmaceuticals

Outline

Molecular heterogeneity of AML

The FLT3 gene, the receptor and Midostaurin

The RATIFY trial

The AMLSG Phase II trial

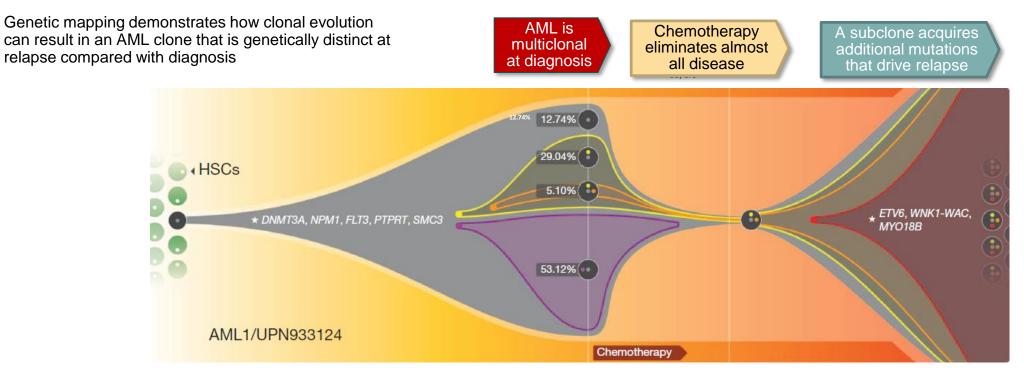
My experience with Midostaurin

Other FLT3 inhibitors under investigation

Future approaches

AML molecular development

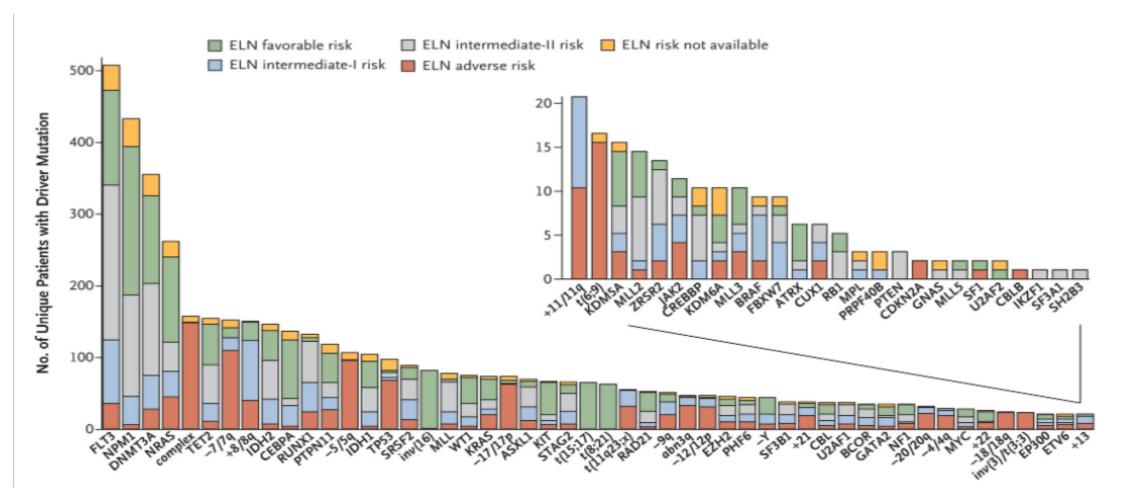
- Development of AML follows specific and ordered evolutionary trajectories
- Mutations in DNMT3A, ASXL1, IDH1/2 and TET2, genes that encode epigenetic modifiers, were often acquired earliest. They are not sufficient for overt leukaemia
- Mutations in receptor tyrosine kinase–RAS pathway genes typically occurred late and frequently happened more than once in the same patient
- Tumor sub-clones (with differing mutations) compete with each other and with normal cells



Adapted from Adapted from Ding L et al. Nature 2012;481:506–510, Hackl et al. Journal of Hematology & Oncology (2017)

Whole-Genome Sequencing in AML has revealed the diversity and frequency of mutations

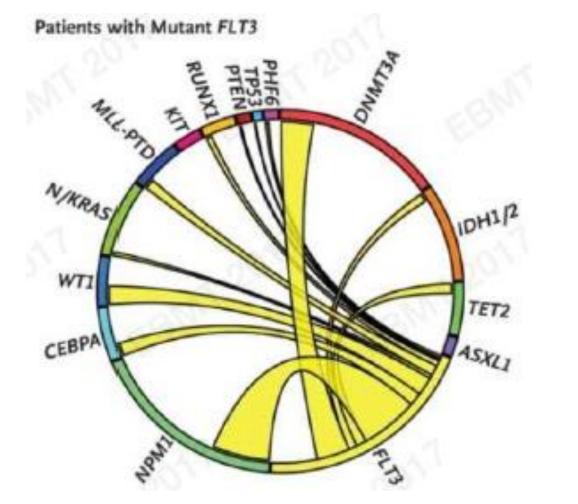
• 1540 patients with AML from three prospective trials of intensive therapy

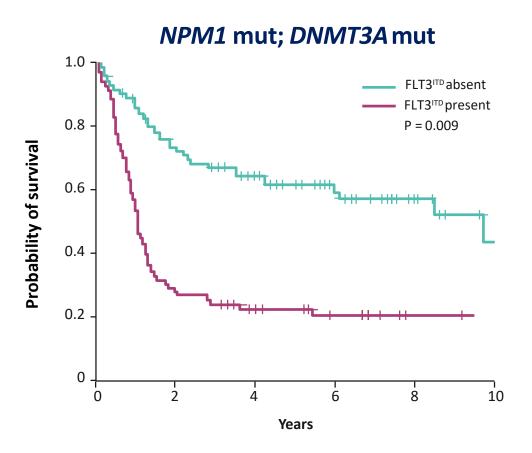


Recurrent Mutations Interfering with Intracellular Signalling

Gene	Key Points	Role/Function
FLT3 (receptor tyrosine kinase)	 FLT3-ITD or TKD mutations in ~30% of patients FLT-ITD common in nK AML FLT3-ITD wild-type ratio impacts prognosis Concurrent with many other common AML mutations 	 Constitutively active kinase domain Promotes RAS and AKT pathways
RAS oncogenes	 KRAS and NRAS mutations in ~14% of patients 	 Increased RAS signalling

FLT3: Concomitant mutations





European LeukemiaNet 2017

Review Article

S blood

Diagnosis and management of AML in adults: 2017 ELN recommendations from an international expert panel

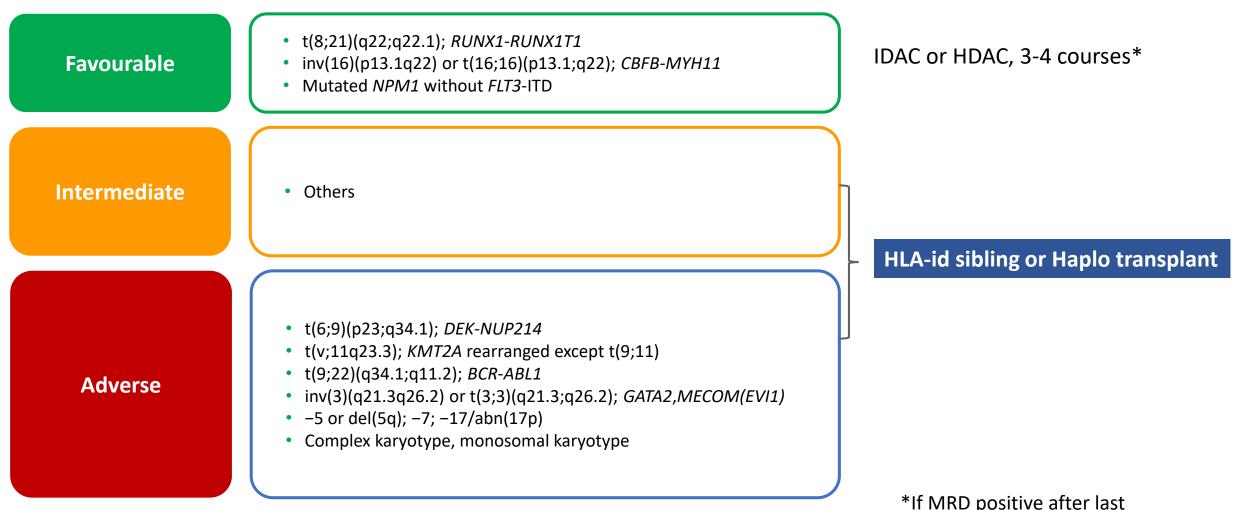
Hartmut Döhner,¹ Elihu Estey,² David Grimwade,³ Sergio Amadori,⁴ Frederick R. Appelbaum,² Thomas Büchner,⁵ Hervé Dombret,⁶ Benjamin L. Ebert,⁷ Pierre Fenaux,⁸ Richard A. Larson,⁹ Ross L. Levine,¹⁰ Francesco Lo-Coco,⁴ Tomoki Naoe,¹¹ Dietger Niederwieser,¹² Gert J. Ossenkoppele,¹³ Miguel Sanz,¹⁴ Jorge Sierra,¹⁵ Martin S. Tallman,¹⁰ Hwei-Fang Tien,¹⁶ Andrew H. Wei,^{17,18} Bob Löwenberg,¹⁹ and Clara D. Bloomfield²⁰

¹Department of Internal Medicine III, University of Ulm, Ulm, Germany; ²Clinical Research Division, Fred Hutchinson Cancer Research Center, Seattle, WA; ³Department of Medical and Molecular Genetics, Faculty of Life Sciences and Medicine, King's College London, London, United Kingdom; ⁴Department of Biomedicine and Prevention, Università di Roma "Tor Vergata," Rome, Italy; ⁵Department of Hematology/Oncology, University of Münster, Münster, Germany; ⁶Institut Universitaire d'Hématologie, Hôpital Saint-Louis, Assistance Publique-Hôpitaux de Paris, Paris, France; ⁷Division of Hematology, Department of Medicine, Brigham and Women's Hospital, Harvard Medical School, Boston, MA; ⁸Service d'Hématologie, Hôpital Saint-Louis, Paris, France; ⁹Department of Medicine, University of Chicago, Chicago, IL; ¹⁰Leukemia Service, Department of Medicine, Memorial Sloan Kettering Cancer Center, New York, NY; ¹¹National Hospital Organization Nagoya Medical Center, Nagoya, Japan; ¹²Department of Hematology, Oncology and Hemostasis, University of Leipzig, Leipzig, Germany; ¹³Department of Haematology, Vrije Universiteit University Medical Center, Amsterdam, The Netherlands; ¹⁴Department of Hematology, University Hospital La Fe, University of Valencia, Valencia, Spain; ¹⁵Hematology Department, Hospital de la Santa Creu i Sant Pau, Jose Carreras Leukemia Research Institute, Barcelona, Spain; ¹⁶Division of Hematology, Department of Internal Medicine, National Taiwan University Hospital, Taipei, Taiwan; ¹⁷Department of Clinical Hematology, The Alfred Hospital, Mebourne, Australia; ¹⁸Australian Centre for Blood Diseases, Monash University, Mebourne, Australia; ¹⁹Department of Hematology, Erasmus University Medical Center, Rotterdam, The Netherlands; and ²⁰The Ohio State University Comprehensive Cancer Center, Columbus, OH

Updated ELN prognostic stratification of AML

Favourable	 t(8;21)(q22;q22.1); RUNX1-RUNX1T1 inv(16)(p13.1q22) or t(16;16)(p13.1;q22); CBFB-MYH11 Mutated NPM1 without FLT3-ITD or with FLT3-ITD^{low} Biallelic mutated CEBPA
Intermediate	 Mutated NPM1 and FLT3-ITD^{high} Wild-type NPM1 without FLT3-ITD or with FLT3-ITD^{low} (without adverse-risk genetic lesions) t(9;11)(p21.3;q23.3); MLLT3-KMT2A Cytogenetic abnormalities not classified as favourable or adverse
Adverse	 t(6;9)(p23;q34.1); DEK-NUP214 t(v;11q23.3); 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 karyotype, monosomal karyotype Wild-type NPM1 and FLT3-ITD^{high} Mutated RUNX1 Mutated ASXL1 Mutated TP53

Simplified classification for middle income countries



*If MRD positive after last consolidation proceed to allogeneic transplantation

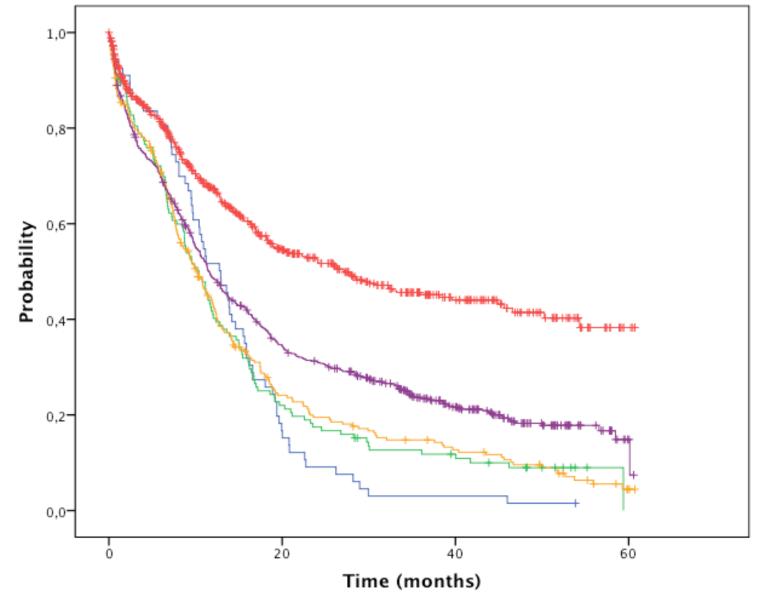
Risk-adapted CETLAM trials for primary AML

	AML-88 ¹	AML-94 ²	AML-99 ³	AML-03 ⁴	AML-12 ⁵
Patients	94	200	350	860	471
Upper age (yrs)	50	60	60	70	70
Induction	DCE (3x7x3)	ICE (3x7x3)	IDICE (3x8x3)	IDICE + G-CSF	IA + G-CSF
Consolidation	Mitox-HiDAC Amsa-IDAC	Mitox-IDAC	Mitox-IDAC	Mitox-IDAC + G-CSF	Hidac
Post-CR adapted therapy based on	HLA-id sibling	HLA-id sibling (age 50 yrs only Auto)	CBF, NK Courses CR HLA-id sibling	CBF, NK Courses CR FLT3 , MLL, MRD Allo Sibl, URD, CB	CBF, NPM1, CEBPA, FLT3 , MLL, MRD EBMT score, Sibl, URD, CB, Haplo
Favourable risk	Auto	Auto Auto	Hidac	HiDAC or Auto	Hidac
Intermediate risk	or Allo	or Allo	Auto	Auto	Auto – HLA-id
Adverse risk			Auto or Allo	GO Auto or Allo	Allo any type
Transplantation	Auto: BM Allo: Sib BM	Auto: BM Allo: Sib PB	Auto: PB Allo: Sib PB	Auto: PB Allo: PR or CB MAC or RIC	Auto: PB Allo: PB, CB, Haplo MAC or RIC
Report	JCO 1996 ¹ Haematologica 2004 ²		<i>Haematologica</i> 2004 ³		

Allo, allogeneic; AML, acute myeloid leukaemia; AMSA, amsacrine; Auto, autologous; BM, bone marrow; CB, cord blood; CBF, core binding factor; CR, complete remission; DCE, daunorubicin, cytarabine and etoposide; EBMT, European Group for Blood and Marrow Transplantation; G-CSF, granulocyte-colony stimulating factor; HLA-id, human leukocyte antigen id; Haplo, haploidentical; HiDAC, high-dose cytarabine; IDAC, intermediate dose cytarabine; IDICE, idarubicin, intermediate doses of ara-C and etoposide; MAC, myeloablative conditioning; Mitox, mitoxantrone; MRD, minimal residual disease; NK, natural killer; PB, peripheral blood; RIC, reduced intensity conditioning; Sib, sibling; URD, unrelated donor 1. Sierra J. et *al. J Clin Oncol.* 1996 Apr;14:1353-63; 2. Brunet *et al. Haematologica* 2004;89:940-9; 3. Oriol *et al. Haematologica* 2004;89:791-800; 4. Unpublished, courtesy of CETLAM; 5. clinicailtrials.gov (NCT01723657)

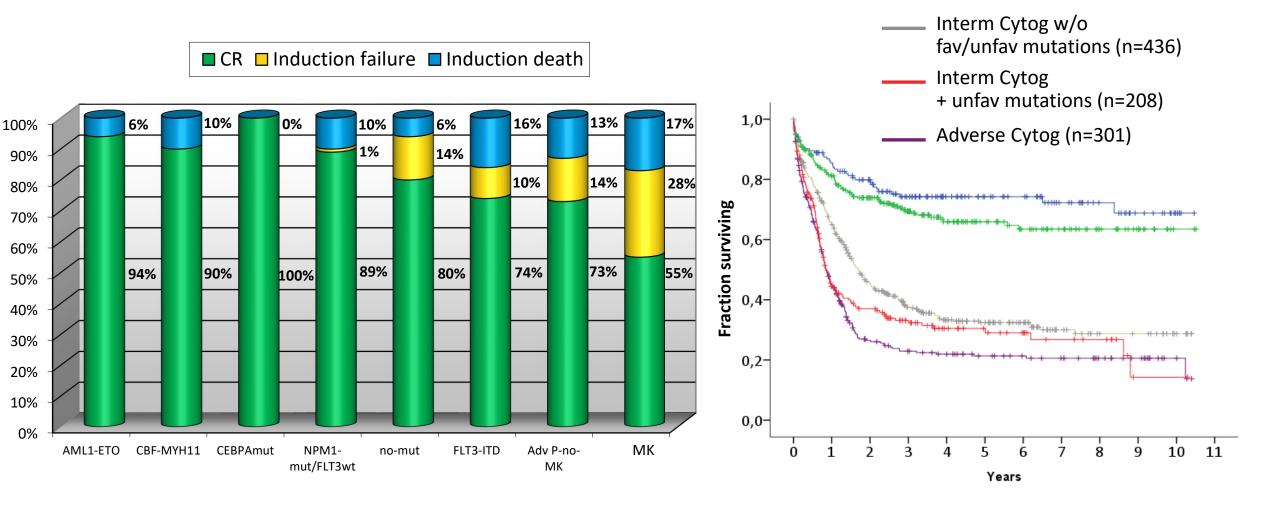
Overall survival in the consecutive CETLAM protocols

1988: 5±% n=67 1994: 15±6% n=134 1999: 18±5% n=253 2003: 28±3% n=682 2012: 48±5% n=603

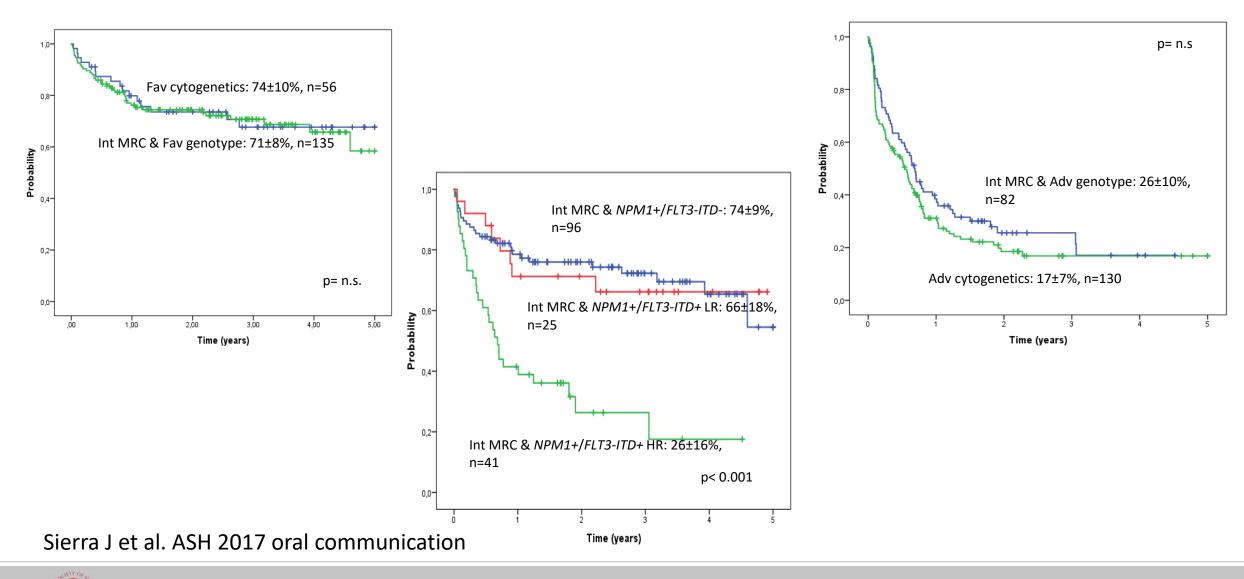


Data on file, May 2018

AML: Genetics and outcome of adults intensively treated (Spanish CETLAM) ---- CBF AML (n=153) ----- Interm Cytog + Fav mut (n=281)

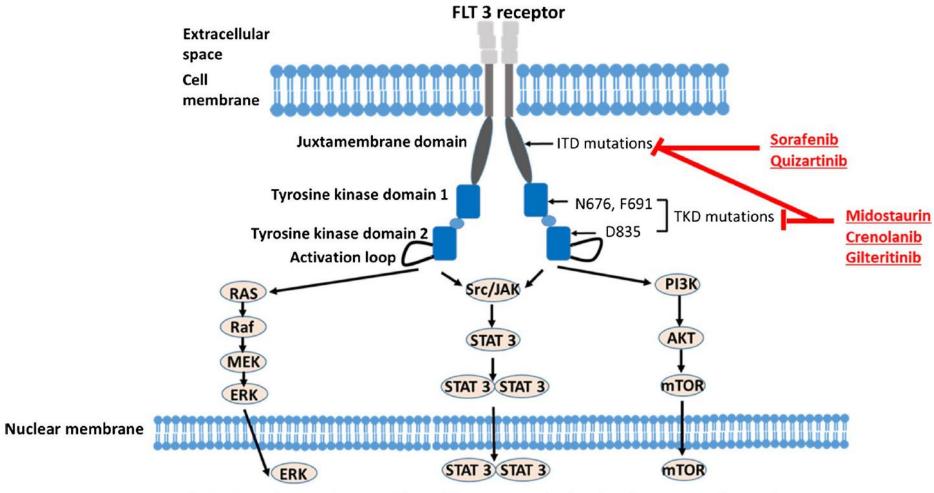


Results: EFS according to genetics at Dx



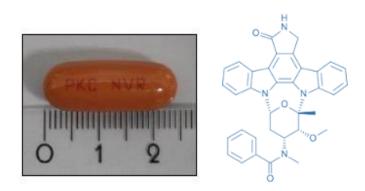
American Society of Hematology

FLT3 inhibitors



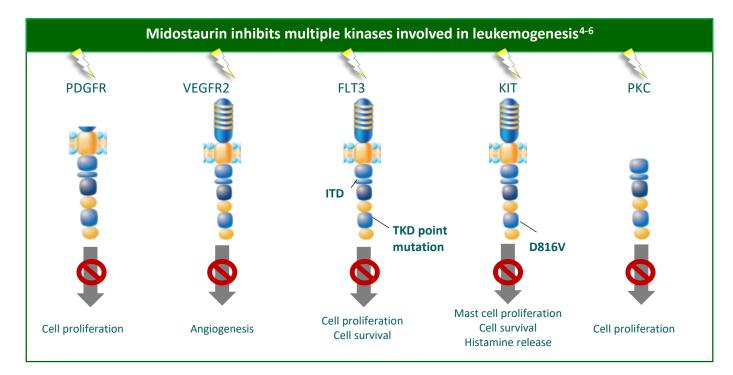
Activation of transcription with proliferation, survival and malignant transformation

Midostaurin: Multikinase inhibitor



- Midostaurin is rapidly absorbed after oral administration¹
- Midostaurin is predominantly metabolized by CYP3A4 to form 2 major active metabolites (CGP62221 and CGP52421)²
- The mean terminal half-lives of midostaurin, CGP62221, and CGP52421 are approximately 20.5 hours, 32.3 hours, and 471 hours, respectively³
 - The metabolite CGP52421 accumulates continuously through day 28

Midostaurin inhibits FLT3-WT as well as FLT3-ITD and FLT3-TKD (D835H and D835Y)



KIT, also known as CD117 and stem cell growth factor receptor (SCGFR); PDGFR, platelet derived growth factor; PKC, protein kinase C.

Midostaurin FDA and EMA approved indications

	Indications
FDA ¹	Midostaurin is a kinase inhibitor indicated for the treatment of (1) adult patients with newly diagnosed AML that is <i>FLT3</i> -mutation positive as detected by an FDA-approved test, in combination with standard cytarabine and daunorubicin induction and cytarabine consolidation, and (2) adult patients with ASM, SM-AHN, or MCL. Limitations of use: midostaurin is not indicated as a single-agent induction therapy for the treatment of patients with AML.
EMA ²	Midostaurin is indicated (1) in combination with standard daunorubicin and cytarabine induction and high-dose cytarabine consolidation chemotherapy and for patients in complete response followed by midostaurin single-agent maintenance therapy, for adult patients with newly diagnosed AML who are <i>FLT3</i> -mutation positive; and (2) as monotherapy for the treatment of adult patients with ASM, SM-AHN, MCL.

ASM, aggressive systemic mastocytosis; EMA, European Medicines Agency; FDA, US Food and Drug Administration; MCL, mast cell leukemia; SM-AHN, systemic mastocytosis with associated hematological neoplasm.

Stone R et al. *Blood Adv.* 2018;2(4):444-453



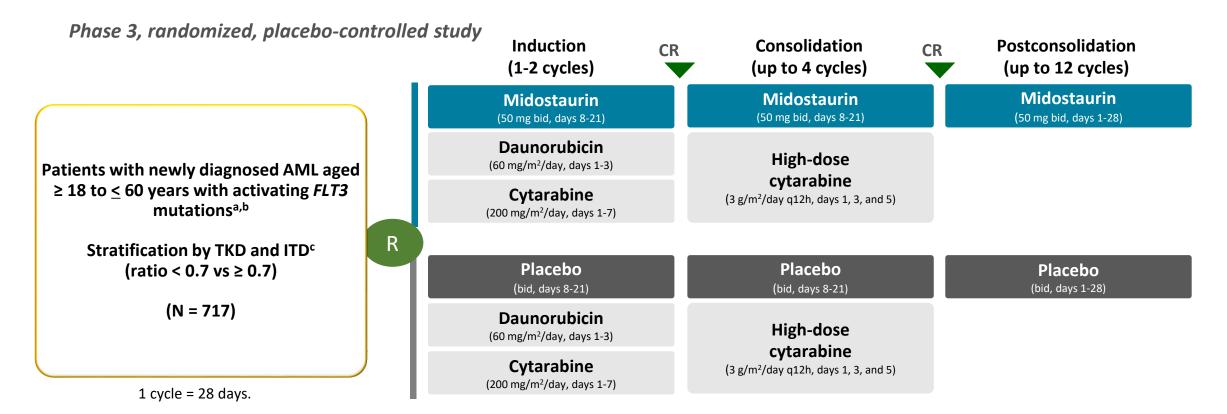
ORIGINAL ARTICLE

Midostaurin plus Chemotherapy for Acute Myeloid Leukemia with a FLT3 Mutation

R.M. Stone, S.J. Mandrekar, B.L. Sanford, K. Laumann, S. Geyer, C.D. Bloomfield,
C. Thiede, T.W. Prior, K. Döhner, G. Marcucci, F. Lo-Coco, R.B. Klisovic, A. Wei,
J. Sierra, M.A. Sanz, J.M. Brandwein, T. de Witte, D. Niederwieser, F.R. Appelbaum,
B.C. Medeiros, M.S. Tallman, J. Krauter, R.F. Schlenk, A. Ganser, H. Serve,
G. Ehninger, S. Amadori, R.A. Larson, and H. Döhner

N Engl J Med. 2017 Aug 3;377(5):454-464.

RATIFY: Schema



First patient enrolled July 2008, Last patient enrolled October 2011 3270 screened, 717 randomized

Primary endpoint: OS **Key secondary endpoint:** EFS

APL, acute promyelocytic leukemia; EFS, event-free survival; OS, overall survival; q12h, every 12 hours.

^a Documented AML (no APL or therapy-related AML).

^b Hydroxyurea therapy allowed \leq 5 days prior to start of study treatment.

^c As determined using BM or PB samples using an assay that was later developed into the LeukStrat CDx FLT3 Mutation Assay.

Stone RM, et al. N Engl J Med. 2017;377:454-464.

RATIFY: Revised analysis plan

Statistical Plan	No. of Events	No. of Patients Accrued	Power	Alphaª	HR to Be Detected
Original	374	514	90%	0.025	0.71
2010 amendment	509	714	84%	0.025	0.78
2015 amendment	357	717	84%	0.0239	0.78

- The event rate (ie, deaths) reached a plateau in 2014 (6 events in 2014 and 4 events by May 2015). Alliance DSMB decided to perform the final analysis with a data cutoff of May, 2015, without achieving the required 509 OS events
 - The SCT rate was higher than expected (25% at the time of amendment vs 15% anticipated)
- EFS was promoted to a key secondary endpoint to be tested in a hierarchical manner if OS was significant

^a Critical value to declare statistical significance (1-sided).

Stone RM, et al. N Engl J Med. 2017;377:454-464.

CTEP, Cancer Therapy Evaluation Program; DSMB, Data Safety Monitoring Board.

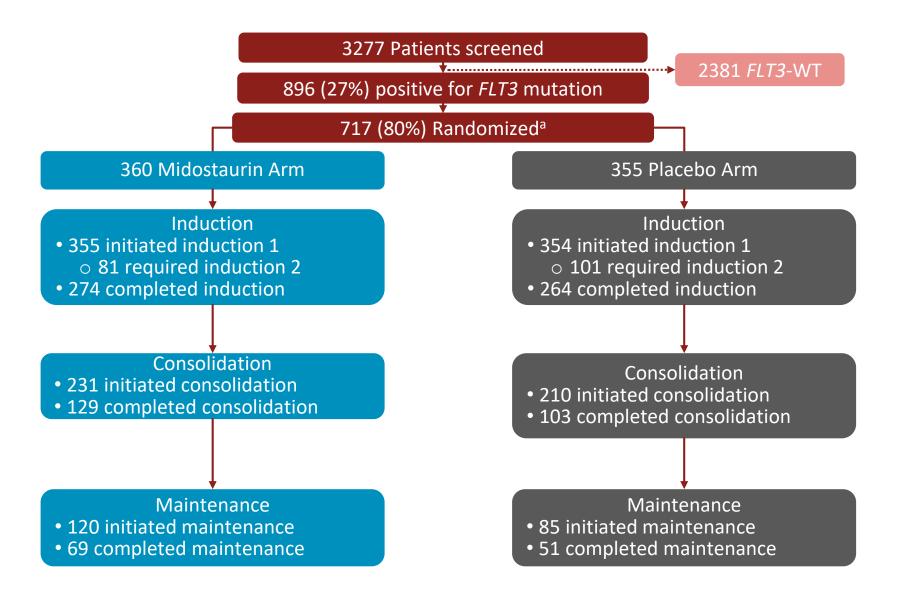
RATIFY: Patient's characteristics

	All Patients (N = 717)	Midostaurin (n = 360)	Placebo (n = 357)	P Value ^a
Median age (range), years	47.9	47.1	48.6	.22
	(18.0-60.9)	(19.0-59.8)	(18.0-60.9)	.22
Female sex, n (%)	398 (55.5)	186 (51.7)	212 (59.4)	.04
FLT3 stratification groups, n (%)				1.00
FLT3-TKD	162 (22.6)	81 (22.5)	81 (22.7)	
FLT3-ITD allelic ratio < 0.7 ^b	341 (47.6)	171 (47.5)	170 (47.6)	
FLT3-ITD allelic ratio ≥ 0.7 ^b	214 (29.8)	108 (30.0)	106 (29.7)	
Madian M/PC (range) $\times 10^{3}/m$	34.9	35.6	33.0	70
Median WBC (range), × 10 ³ /μL	(0.6-421.8)	(0.6-421.8)	(0.8-329.8)	.72
Madian platalat count (ranga) x 103/ul	50.0	50.0	50.0	FO
Median platelet count (range), × $10^3/\mu$ L	(2.0-461.0)	(2.0-461.0)	(8.0-444.0)	.58
Madian ANC (ranga) nor mm ³	2.2	2.2	2.3	CE
Median ANC (range), per mm ³	(0-55.9)	(0-55.9)	(0-128.7)	.65

From Stone RM, et al. *N Engl J Med*. 2017;377:454-464. Copyright © 2017 Massachusetts Medical Society. Reprinted with permission from Massachusetts Medical Society.

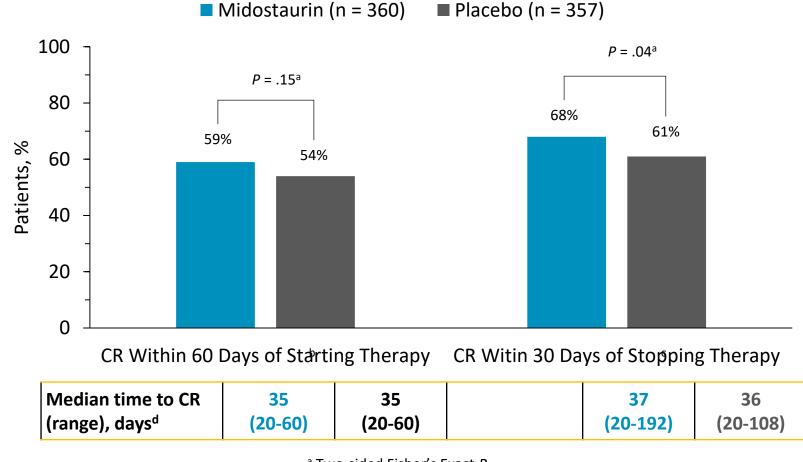
^a Two-sided; Kruskal-Wallis P values are presented for continuous measures (eg, age) and Chi-square values are presented for categorical counts (eg, sec).^b Patients could also have *FLT3*-TKD.

RATIFY: Consort diagram



RATIFY: CR rates by arm

Secondary endpoint

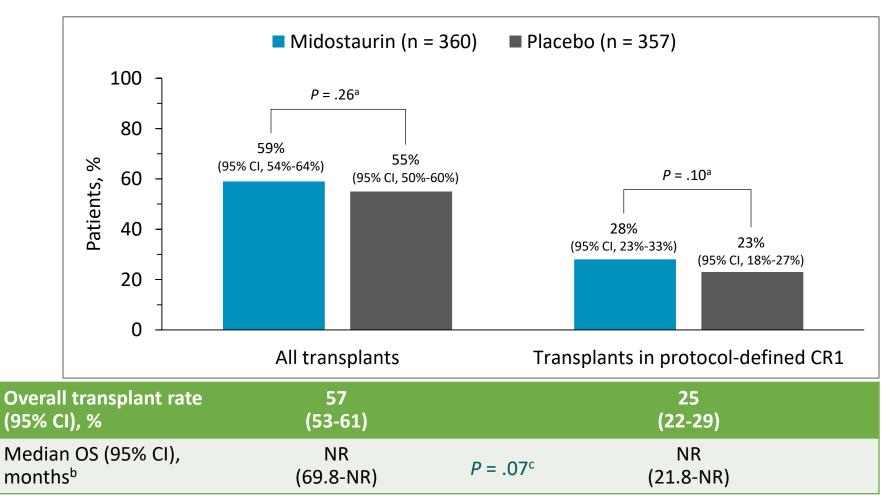


^a Two-sided Fisher's Exact P.

^b Per protocol, CRs were defined as occurring within 60 days of initiating study protocol therapy. ^c An expanded definition of CR Included all CRs reported within 30 days of ending protocol therapy. ^d Kaplan-Meier estimates.

RATIFY: AlloSCT rates by arm

Secondary endpoint



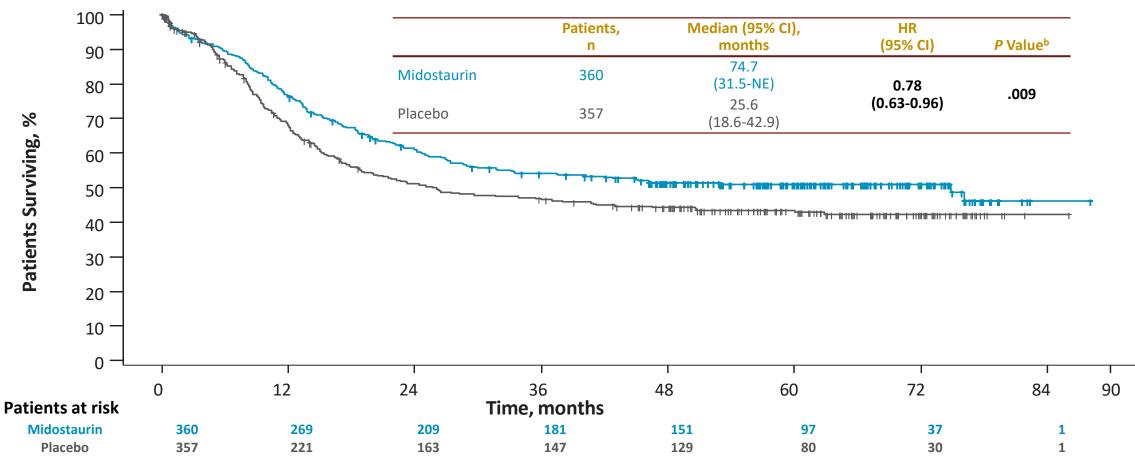
^a Two-sided Fisher's Exact P.

^b For patients who received an SCT in CR1 (101 in the midostaurin arm and 81 in the placebo arm). ^c Two-sided long-rank test.

RATIFY: OS, non-censored for SCT

Primary endpoint

22% reduced risk of death in the midostaurin arm (vs placebo)



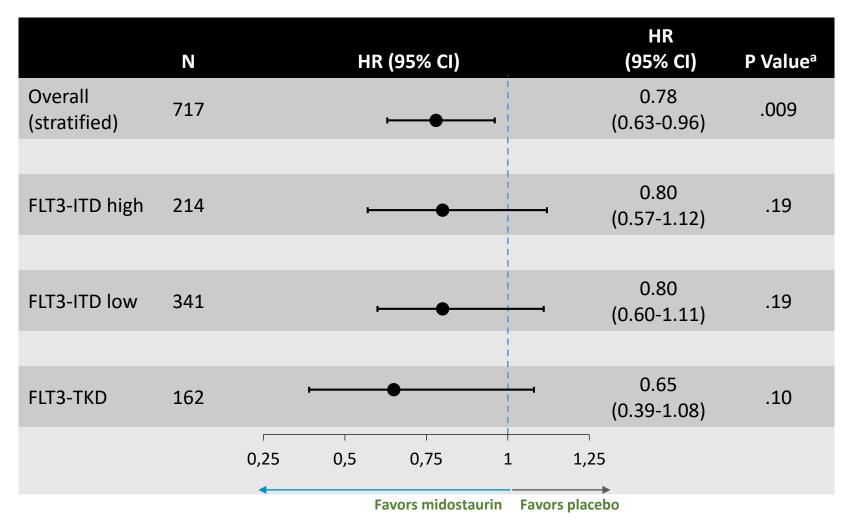
From Stone RM, et al. N Engl J Med. 2017;377:454-464. Copyright © 2017 Massachusetts Medical Society. Reprinted with permission from Massachusetts Medical Society.

NE, not estimable.

^a Cox model stratified on FLT3 subtype. ^b Stratified on FLT3 subtype; one-sided, log-rank P value.

RATIFY: Consistent effect on OS by FLT3 status

Secondary endpoint

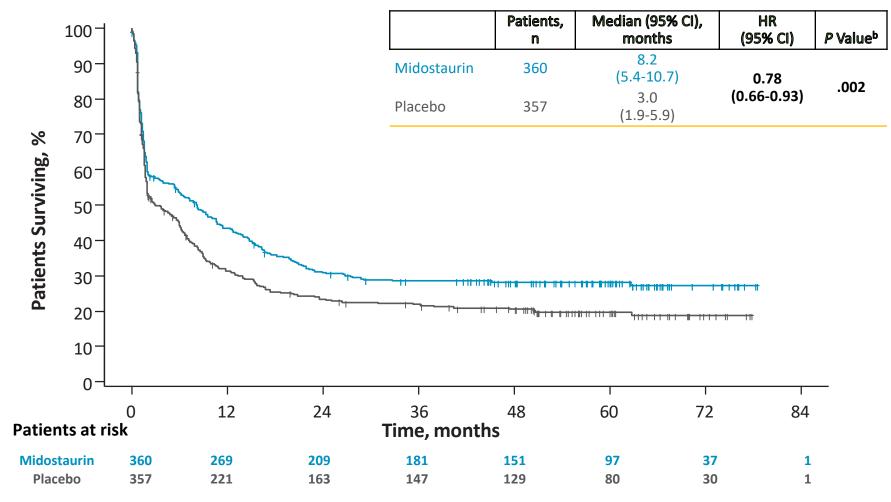


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^a *P* value is one-sided for the overall (stratified) analysis; *P* values are two-sided for the analyses by FLT3 subgroup. *FLT3*-ITD-low, *FLT3*-ITD/-WT allelic ratio < 0.7; *FLT3*-ITD-high, *FLT3*-ITD/-WT allelic ratio \ge 0.7.

RATIFY: EFS, noncensored for SCT^a

Key secondary endpoint



From Stone RM, et al. N Engl J Med. 2017;377:454-464. Copyright © 2017 Massachusetts Medical Society. from Massachusetts Medical Society. ^a Event defined as first of no CR within 60 days of initiating study therapy, relapse, or death.

^b Stratified on *FLT3* subtype; one-sided, log-rank *P* value.

RATIFY: Consistent effect on EFS by FLT3 status

Secondary endpoint

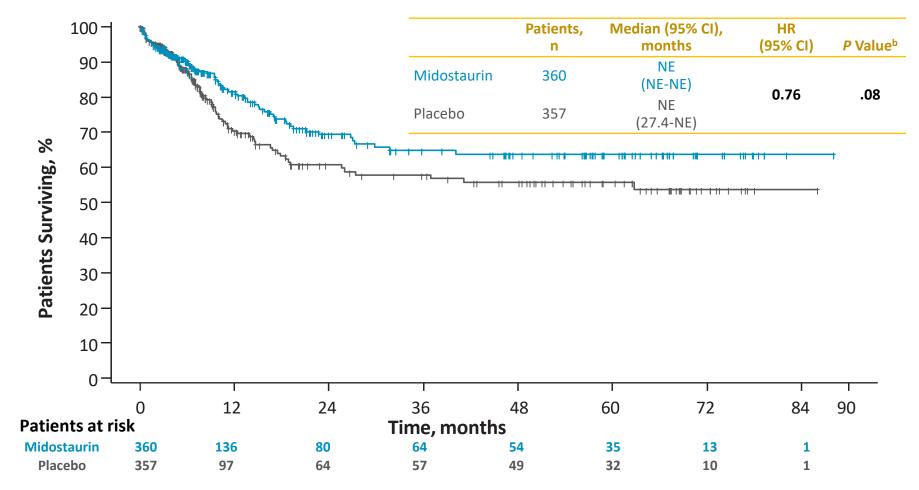
	N	HR (95% CI)	HR (95% CI)	P Value ^a
Overall (stratified)	717	▶ 	0.78 (0.66-0.96)	.002
FLT3-ITD high	214	••	0.77 (0.57-1.04)	.08
FLT3-ITD low	341	•	0.5 (0.67-1.09)	.21
FLT3-TKD	162	••	0.64 (0.43-0.96)	.03
		0,25 0,5 0,75 1 1,25		
		Favors midostaurin Favors placebo		

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FLT3-ITD-low, *FLT3*-ITD/-WT allelic ratio < 0.7; *FLT3*-ITD-high, *FLT3*-ITD/-WT allelic ratio \ge 0.7.

^a *P* value is one-sided for the overall (stratified) analysis; *P* values are two-sided for the analyses by FLT3 subgroup.

RATIFY: OS, censored at time of SCT Secondary endpoint

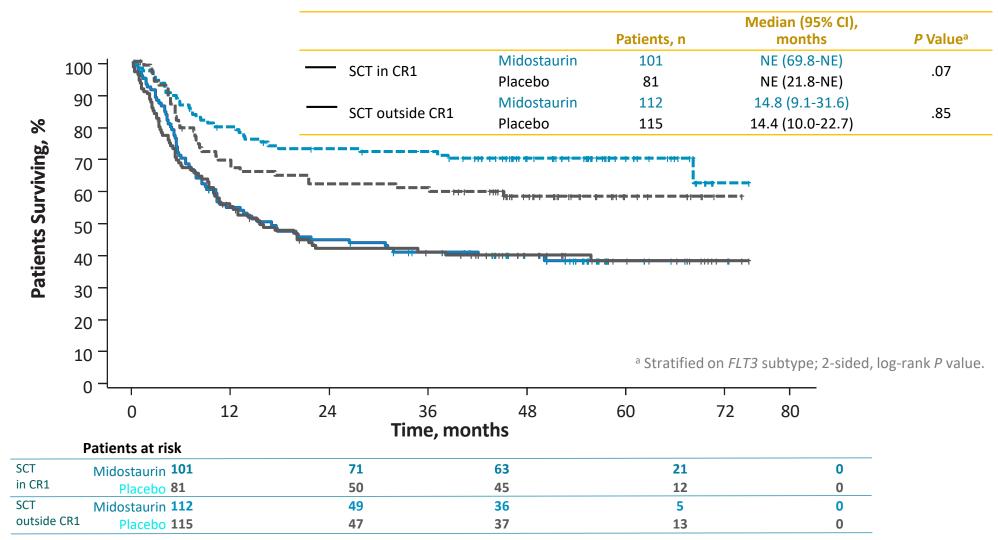


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* Stratified on FLT3 subtype; two-sided, long-rank P value.

RATIFY: OS by timing of SCT

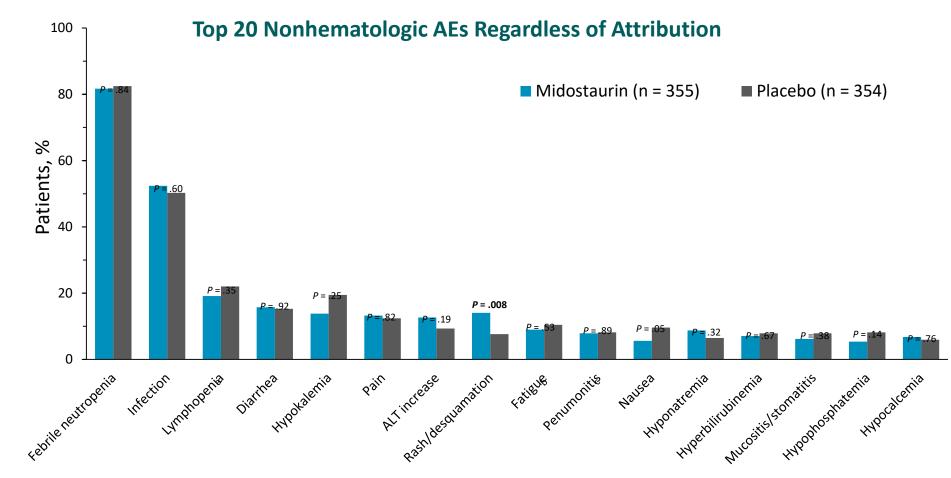
Secondary endpoint



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RATIFY: Safety

Most common nonhematologic grade \geq 3 AEs



^a Infection includes the following terms: infection with grade 3/4 neutrophils (ANC < 1.0 × 10³/µL), infections with normal ANC or grade 1/2 neutrophils, infection with unknown ANC, opportunistic infection associated with grade ≥ 2 lymphopenia, and other infections.

^b Fatigue includes asthenia, lethargy, and malaise.

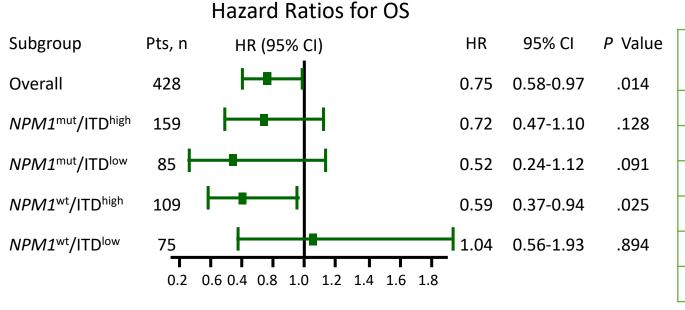
^c Includes pneumonitis and pulmonary infiltrates.

Stone RM, et al. N Engl J Med. 2017;377:454-464.

RATIFY: Summary

- The RATIFY trial met its primary endpoint of OS
 - Midostaurin reduced the risk of death vs placebo plus standard chemotherapy by 22%
- Despite high rates of SCT, OS and EFS benefits were consistent in censored and uncensored analyses
 - OS uncensored HR = 0.78 (0.63-0.96), censored HR = 0.76
 - EFS uncensored HR = 0.78 (0.66-0.93)
- The most common non-hematologic grade 3/4 AEs (midostaurin vs placebo) were febrile neutropenia (82% vs 82%), infection (52% vs 50%), and diarrhea (16% vs 15%)
- Rates of grade 3/4 AEs were similar between midostaurin and placebo arms with the exception of rash/desquamation (14% vs 8%, respectively)

Prognostic Analysis of NPM1/FLT3-ITD Genotypes in RATIFY: Multivariate OS Analysis



Midostaurin Better

Placebo Better

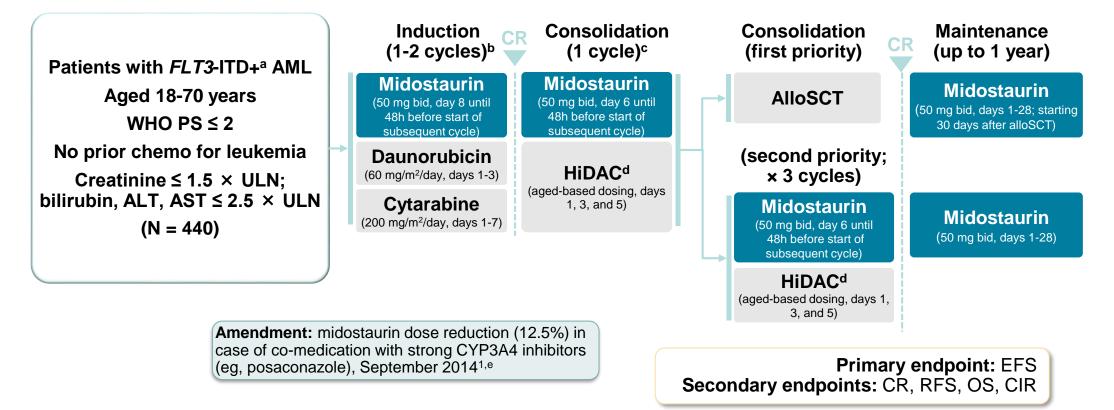
Multivariate Analysis for OS

ue	Characteristic	2-Sided P Value
1	ELN subgroup (NPM1/FLT3-ITD)	< .001
3	Treatment (midostaurin vs placebo)	.012
-	Allogeneic HCT	< .001
5	WBC (≥ vs < 50 x 10 ⁹ /L)	.028
Ļ	Age (difference of 10 yrs)	.335
	Sex	.689

Slide credit: clinicaloptions.com

DE02T/AMLSG 16-10: study design^{1,2}

Phase 2, open-label, uncontrolled, single-arm trial



CIR, cumulative incidence of relapse; CR, complete remission; EFS, event-free survival; OS, overall survival; RFS, relapse-free survival.

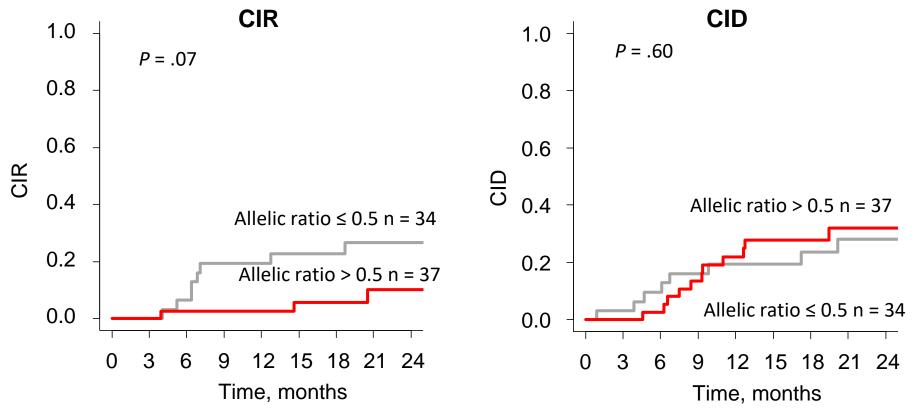
- ^a FLT3 screening results within 48 hours; FLT3-ITD/-WT ratio > 0.05 by Genescan-based fragment length analysis required to be FLT3-ITD positive.
- ^b During induction, patients achieving PR after cycle 1 can receive an optional cycle 2.
- ^c For patients eligible for alloSCT, 1 course of HiDAC is optional before alloSCT.
- ^d Age-appropriate cytarabine dose on days 1, 3, and 5: 18-65 years, 3 g/m² q12h (total dose 18 g/m²); > 65 years ,1 g/m² q12h (total dose 6 g/m²).

e Dose reduction of 7-fold (12.5%) was based on the lower bound of the confidence interval of the ratio [AUC midostaurin with ketoconazole/AUC midostaurin without ketoconazole]; midostaurin 25 mg every other day.

1. Schlenk RF, et al. Blood. 2016;128(22):[abstract 449]. 2. ClinicalTrials.gov.

DE02T/AMLSG 16-10: CIR and CID

For patients in CR1 per protocol



- Patients who received maintenance therapy (n = 52; 41 after alloSCT and 11 after high-dose cytarabine) received a median number of 6 cycles (range, 1-12)
 - Maintenance after alloSCT was prematurely terminated in 33 patients (80%)

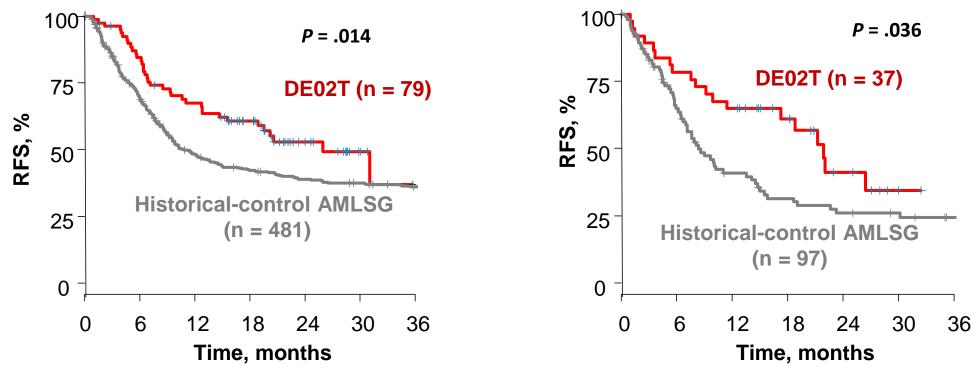
From Schlenk RF, et al. In: Proceedings from the American Society of Hematology; December 5-8, 2015; Orlando, FL [abstract 322]. Reprinted with author's permission.

DE02T/AMLSG 16-10: RFS

Patients Aged 18 to < 60 years

Patients Aged 60-70 years

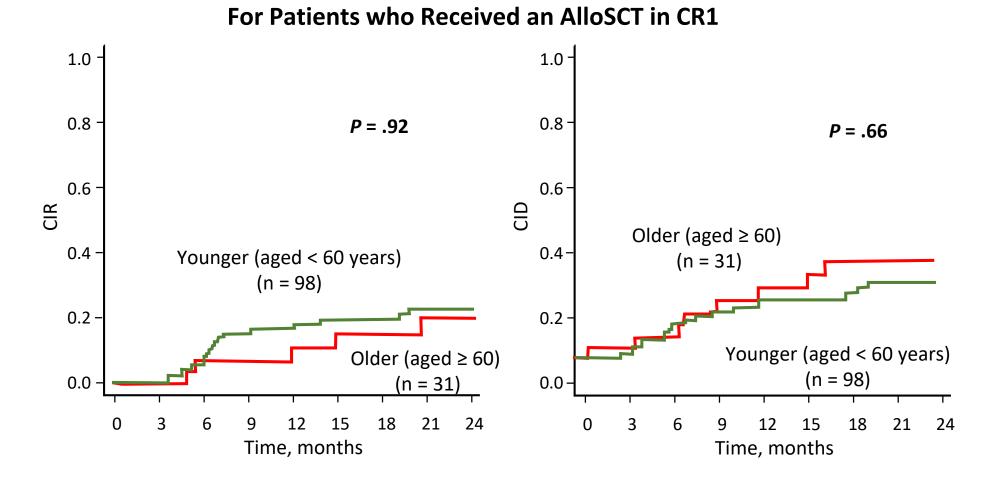
• Patients in the DE02T/AMLSG 16-10 study had improved RFS vs historical control



From Schlenk RF, et al. In: Proceedings from the American Society of Hematology; December 5-8, 2015; Orlando, FL [abstract 322]. Reprinted with author's permission.

DE02T/AMLSG 16-10: CIR and CID by age

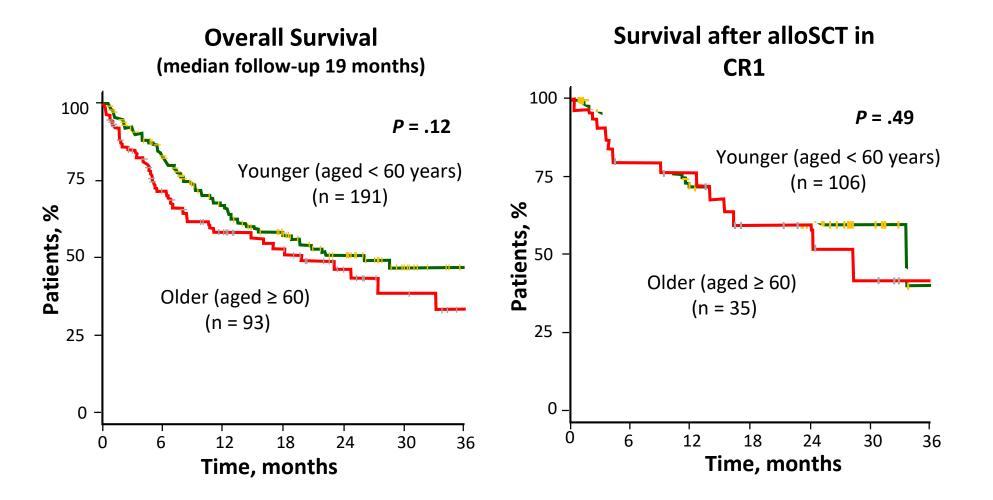
 $(< 60 \text{ years } vs \ge 60 \text{ years})$



From Schlenk RF, et al. In: Proceedings from the American Society of Hematology; December 3-6, 2016; San Diego, CA [abstract 449]. Reprinted with author's permission.

DE02T/AMLSG 16-10: Survival by age

 $(< 60 \text{ years } vs \ge 60 \text{ years})$

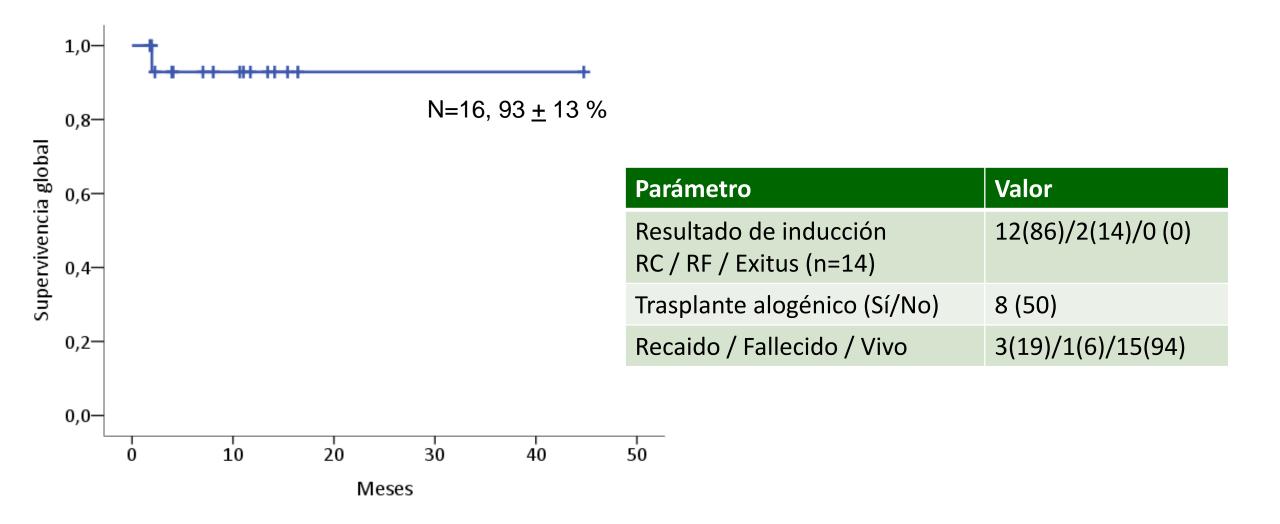


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Midostaurina en LMA de novo con FLT3-ITD): CETLAM (n=16)

Característica	Valor
Edad (mediana, extremos)	51 (24-64)
<60 años / >60 años	13 (81) / 3 (19)
Sexo (varón/mujer)	10 (63) / 6 (37)
Leucocitosx10 ⁹ /l	95 (66-122)
Plaquetas x10 ⁹ /l	53 (13-155)
Citogenética (MRC)	
Favorable/Intermedia/Adversa	1 (7) / 13 (81) / 2 (12)
Ratio FLT3 (<0,5 / <u>></u> 0,5)	4 (25) / 12 (75)
NPM1 mutado	8 (50)

Midostaurina en LMA de novo con FLT3-ITD): CETLAM (n=16)

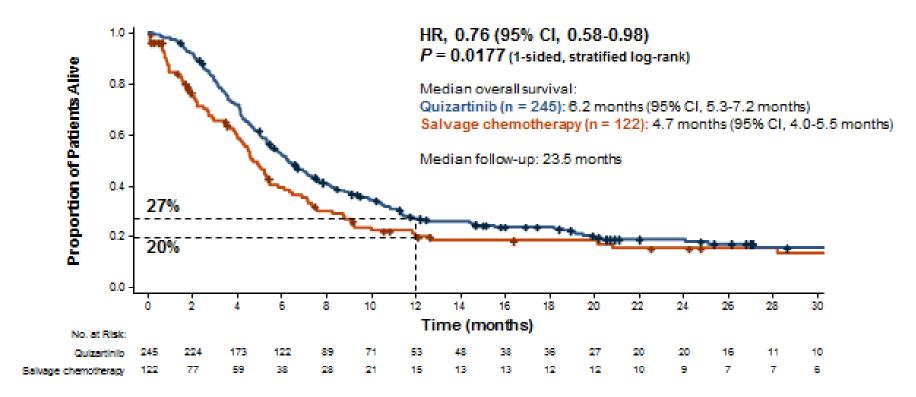


Other FLT3 inhibitors (I)

	Specificity	Available data	Ongoing or recently completed trials
Sorafenib	Non-specific but potent inhibitor of FLT3/ITD RTK	Leads to transient reductions in marrow and circulating myeloblasts Improves EFS in combination with chemotherapy in younger patients, regardless of FLT3 mutational status	Phase II sorafenib with HMA therapy as upfront approach for FLT3/ITD- mutant patients (NCT02196857) Maintenance sorafenib after HSCT for FLT3/ITD AML (EudraCT 2010- 018539-16)
Quizartinib	Selective and potent inhibitor of FLT3/ITD RTK	As monotherapy, CR and CRi in sizeable proportion of relapsed FLT3/ITD-mutant patients	Combined with HMA QuANTUM-R: quizartinib vs conventional salvage QuANTUM-First: Phase III quizartinib with conventional CT in FLT3/ITD (NCT02668653)

Quizartinib vs Salvage chemotherapy in Rel/Refr AML with FLT3-ITD: Quantum-R

Primary Endpoint: Overall Survival



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Cortes et al. 23rd EHA 2018, LB 2600;

Other FLT3 inhibitors (II)

	Specificity	Available data	Ongoing or recently completed trials
Crenolanib	Active against both FLT3/ITD and FLT3-TKD mutation variants	Multiple ongoing trials of this agent are in progress, including in the frontline and relapsed/ refractory settings	Safety study of crenolanib combined with upfront induction in FLT3-mutant patients (NCT02283177) Phase I/II trial of crenolanib combined with re-induction regimens for R/R FLT3-mutant patients (NCT02626338) Pilot study of crenolanib combined with salvage regimens for R/R patients, regardless of FLT3-mutation status (NCT02400281) Crenolanib maintenance following stem cell transplant (NCT02400255
Gilteritinib	Selective FLT3 inhibitor, which can target both FLT3/ITD and FLT3-TKD	Remains under study in clinical trials. Composite remission rate of 46% among relapsed /refractory FLT3-mutant patients	Phase III randomized study of gilteritinib vs conventional salvage CT among FLT3-mutant Phase II/III study of gilteritinib combined with HMA therapy Phase I study of gilteritinib combined with conventional induction CT Phase III study of gilteritinib maintenance among FLT3- mutant patients in first remission Phase III randomized study of maintenance gilteritinib after HSCT for patients with FLT3/ITD AML CR1

FLT3-ITD AML: Approaches to further investigate

Mechanisms of Resistance to FLT3 Inhibitors and the Role of the Bone Marrow Microenvironment

Ghiaur G, Levis M. Hematol Oncol Clin North Am. 2017Aug;31(4):681-692. doi: 10.1016/j.hoc.2017.04.005. PMID: 28673395.

Combination with immunotherapy

Tyrosine kinase inhibition increases the cell surface localization of FLT3-ITD and enhances FLT3-directed immunotherapy of acute myeloid leukemia. Reiter K, et al. Leukemia. 2017 Aug 14. doi: 10.1038/leu.2017.257. PMID: 28895560.

Intensification of therapy resulted in a reduced relapse risk

Gemtuzumab Ozogamicin in induction therapy (Castaigne S, et al. Lancet 2012; 379: 1508–16.) High-dose daunorubicin in induction therapy (Burnett AK, et al. Blood. 2016;128(3):449-52.)

¡Gracias por la invitación!

