


Evolving Practice in AML: A Case-based Guide to New and Emerging Treatment Options

**Evolving Practices in AML:
A Case-based Guide to New and Emerging Treatment Options**

**Wednesday, September 11, 2019
Houston, Texas**

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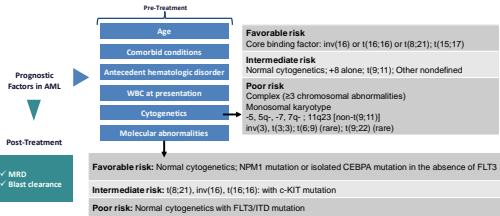
1

**The Impact of Cytogenetic and Molecular Abnormalities on Patient Management:
Diagnostic, Prognostic, and Therapeutic Importance**

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Associate Professor
Department of Leukemia
The University of Texas MD Anderson Cancer Center
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6

Traditional and New Prognostic Factors in AML^{1,2}



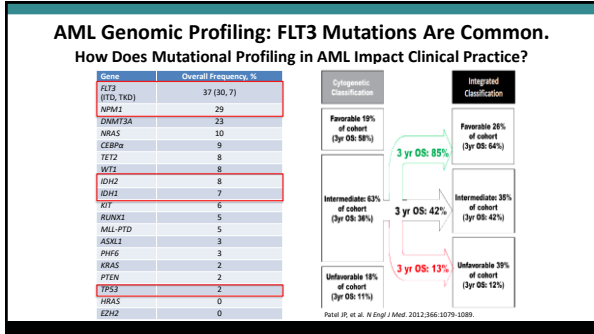
| Pre-Treatment | Post-Treatment | Risk Category |
|---------------------------------|-----------------|-------------------|
| Age | MRD | Favorable risk |
| Comorbid conditions | Blast clearance | Favorable risk |
| Antecedent hematologic disorder | | Favorable risk |
| WBC at presentation | | Favorable risk |
| Cytogenetics | | Favorable risk |
| Molecular abnormalities | | Favorable risk |
| | | Intermediate risk |
| | | Poor risk |

Favorable risk: Normal cytogenetics; NPM1 mutation or isolated CEBPA mutation in the absence of FLT3
Intermediate risk: t(8;21), inv(16), t(16;16); with c-KIT mutation
Poor risk: Normal cytogenetics with FLT3/ITD mutation

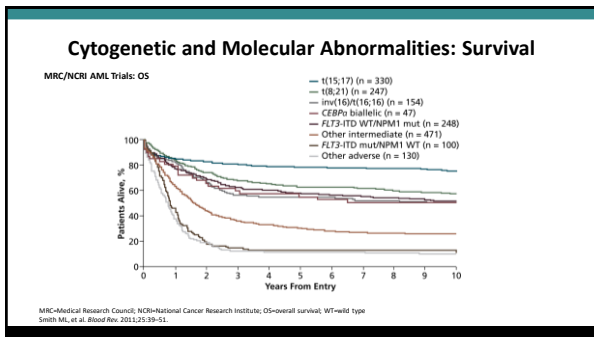
AML=acute myeloid leukemia; CEBPA=CCAAT/enhancer binding protein α; FLT3=FLT3-like tyrosine kinase 3; ITD=internal tandem duplications; MRD=minimal residual disease; NPM1=nucleophosmin; WBC=white blood cell
¹Grimwade D, Hills RK. Hematology Am Soc Hematol Educ Program. 2009:385-395; ²NCCN Clinical Practice Guidelines in Oncology. Acute myeloid leukemia. Version 1. 2016.

16

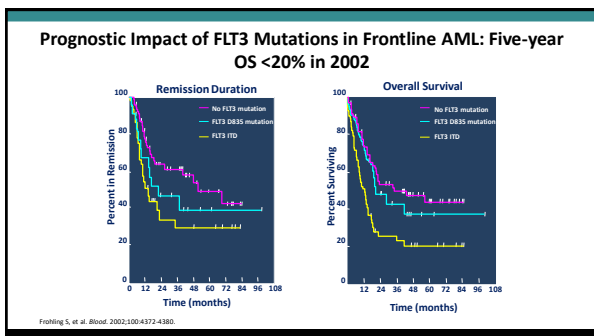
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17

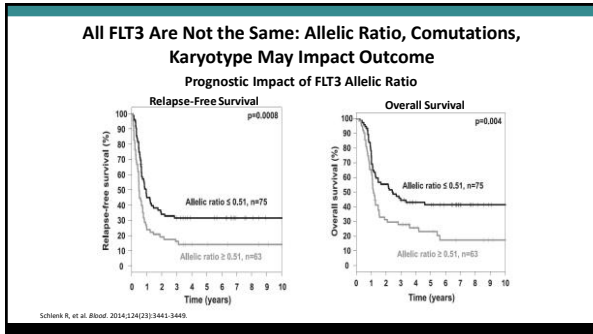


18

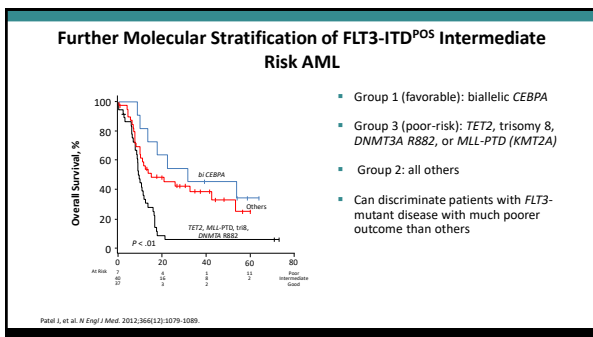


19

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20



21

| Genetic Risk Group | Frequency | Survival | ELN 2017 Subset |
|--------------------|-----------|----------|---|
| Favorable | 15% | 65-75% | <ul style="list-style-type: none"> t(8;21)(q22;q22); <i>RUNX1-RUNX1T1</i> inv(16)(p13.1q22) or t(16;16)(p13.1;q22); <i>CBFB-MYH11</i> Mutated <i>NPM1</i> without <i>FLT3-ITD</i> or <i>FLT3-ITD</i>^{low} Biallelic mutated <i>CEBPA</i> |
| Intermediate | 55% | 50-55% | <ul style="list-style-type: none"> Mutated <i>NPM1</i> and <i>FLT3-ITD</i>^{low} Wild-type <i>NPM1</i> without <i>FLT3-ITD</i> or <i>FLT3-ITD</i>^{low} (without adverse-risk genetic lesions) Any cytogenetics not classified as favorable or adverse t(9;11)(p22;q23); <i>MLL13-MLL</i> |
| Adverse | 30% | 20-25% | <ul style="list-style-type: none"> t(6;9)(q23;q34); <i>DEK-NUP214</i> t(v;11)(v;q23); <i>MLL (KMT2A)</i> rearranged inv(3)(q21q26.2) or t(3;3)(q21;q26.2); <i>RPN1-EVI1 (GATA2, MECOM (EVI1))</i> t(9;22)(q34.1;q11.2) <i>BCR-ABL1</i> Monosomy 5 or del(5q); monosomy 7; monosomy 17; abnormal 17p Complex karyotype (≥ 3 abnormalities) or monosomal karyotype Wild-type <i>NPM1</i> and <i>FLT3-ITD</i>^{high} Mutated <i>RUNX1</i> Mutated <i>ASXL1</i> Mutated <i>TP53</i> |

Döhner H, et al. Blood. 2017;129(4):424-447.

22

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Major Targets of Past and Future Therapeutic Development

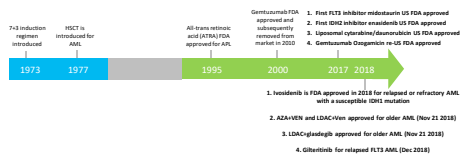
| | |
|---|---|
| AML with <i>FLT3</i> internal tandem duplication | <ul style="list-style-type: none"> Impact on both therapy and prognosis Many <i>FLT3</i> kinase inhibitors explored in recent years...now several next-generation agents in development |
| <i>KIT</i> mutations in CBF AML | <ul style="list-style-type: none"> <i>KIT</i> mutations found in 30%-35% of CBF AML cases, but rare in other AML subgroups In CBF AML, mutations cluster mainly in exons 8 and 17 |
| <i>IDH1/2</i> mutations in AML | <ul style="list-style-type: none"> <i>IDH1/2</i> mutations confer a gain-of-function, including increased histone and DNA methylation and impaired cellular differentiation |
| <i>Bcl-2</i> as a therapeutic target in AML | <ul style="list-style-type: none"> <i>Bcl-2</i> binds and sequesters pro-apoptotic molecules; inhibition of <i>Bcl-2</i> primes cancer cells for death |
| Epigenetic targets (EZH2, LSD1, BRD, PRMT5, others) | <ul style="list-style-type: none"> Novel agents on the horizon that target specific epigenetic pathways These are in early clinical trial development |
| TP53, C-CBL, MLL-Menin | <ul style="list-style-type: none"> Phase 1 clinical trials |

23

Treatment of AML (Accelerated Progress 2017-2019): History

Since its introduction in the early 1970s, 7+3 therapy (Cytarabine for 7 days + Anthracycline for 3 days) has been the standard of care for AML

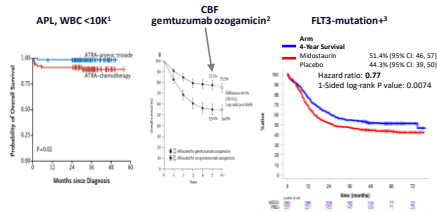
US FDA approvals



| | | | | | | | | | |
|-----------------|------|------|-------|-------|-------|-------|-------|------|------|
| Year | 1975 | 1980 | 1990 | 1995 | 2000 | 2005 | 2009 | 2013 | 2018 |
| 5-year survival | 6.3% | 6.8% | 11.4% | 17.3% | 16.8% | 25.7% | 28.1% | 27% | ?? |

24

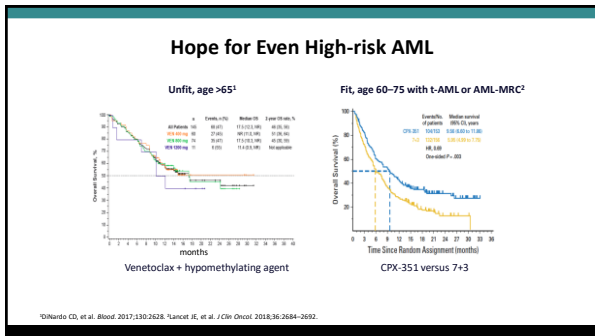
Data-driven Therapy for Molecular Subtypes



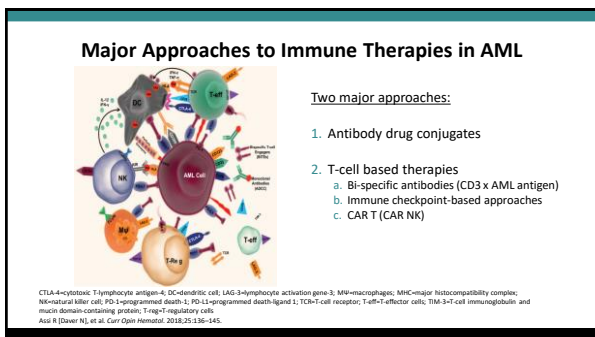
¹Lo Coco F, et al. N Engl J Med. 2013;369:111-121. ²Hibi RK, et al. Lancet Oncol. 2014;15:986-996. ³Stone RM, et al. N Engl J Med. 2017;377:454-464.

25

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26



27

Current and Emerging Therapies in Newly Diagnosed AML: Adapting Treatment to Meet Needs of the Individual Patient

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28

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Case 1: Presentation

- 68-year-old woman with prior history of polycythemia vera diagnosed 5 to 6 years ago and treated with intermittent phlebotomy only. Over time, her hemoglobin has normalized; however her WBC has started to rise. Also has atrial fibrillation s/p CVA on coumadin therapy, Graves disease, HTN
 - June 2019: CBC showed WBC of 30K, normal hgb and platelet count
 - Early August 2019: WBC 108K with 50% blasts
 - Mid-August 2019: Referred to academic center for new leukemia diagnosis
- Complains of increased shortness of breath for weeks prior to admission, intermittent blurry vision, morning nausea, and blood-stained stools

34

Case 1 (continued)

- Exam: ECOG 1, BP 164/69, 92% oxygen saturation on room air, unremarkable exam except decreased breath sounds in both lungs, no organomegaly
 - CBC: WBC 139K, hgb 11.7, plts 208, 66% peripheral blasts, INR 2.46
 - Chemistry: BUN 35, Cr 1.65, uric acid 12.9, LDH 1361, Ph 4.9
 - Cardiac echo: Normal LVEF 65-70%
- Peripheral blasts: MPO+
- Flow cytometry: 100% of blasts are CD33+ bright
- Cytogenetics: Normal female karyotype
- Molecular: FLT3^{mut}, IDH1/2^{wt}, DNMT3A^{mut}, TET2^{mut}, JAK2V617F^{mut}, TP53^{mut}

35

Case 2: Presentation

- 87-year-old woman with PMHx COPD, HTN, PMR presents for second opinion for AML
 - March 2018: Counts noted to be slowly dropping
 - Dec 2018: Pancytopenic, referred to hematologist
 - BMBx: Hypercellular BM with 10% blasts, normal karyotype -> Dx MDS
 - Started darbepoetin alfa growth factor therapy -> no response
 - April 2019: azacitidine SQ 5 days every 28 days (now cycle 4 day 13)
 - July 2019: Neutropenic fever, RLL pneumonia treated with IV antibiotics
- Repeat BMBx: AML with 20% blasts, MDS related changes

38

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Case 2 (continued)

- Complaints of chronic fatigue improved after transfusions, mouth sore, easy bruising but remains very active and mobile with family support present
 - Exam: ECOG 1, one oral ulcer, scattered bruises, independent ambulation
 - CBC: WBC 0.87, hgb 6.6 gm/dl, plts 16K, 1% blasts, 37% neutrophils, 55% lymphs
- Repeat BMBx: AML with 21% blasts, MDS related changes
- Flow cytometry: 86% of blasts are CD33+
- Cytogenetics: normal karyotype
- Molecular panel: NPM^{wt}, FLT3^{wt} (full NGS panel is pending)

39

Please Pick up Your Key Pad to Answer
the Following Question

40

AML Therapy: An Embarrassment of Riches



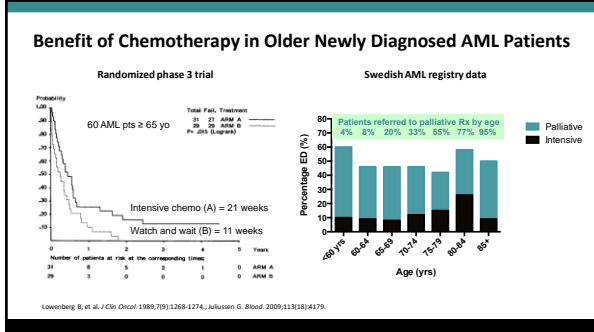
Nine Drugs FDA approved for AML since 2017

1. Midostaurin (April 2017)
2. Liposomal cytarabine/daunorubicin (Aug 2017)
3. Enasidenib (Aug 2017)
4. Gemtuzumab ozogamicin (Sept 2017)
5. Ivosidenib (July 2018)
6. Gilteritinib (Nov 2018)
7. Glasdegib (Nov 2018)
8. Venetoclax (Nov 2018)
9. Tagraxofusp-erzs* (Dec 2018)

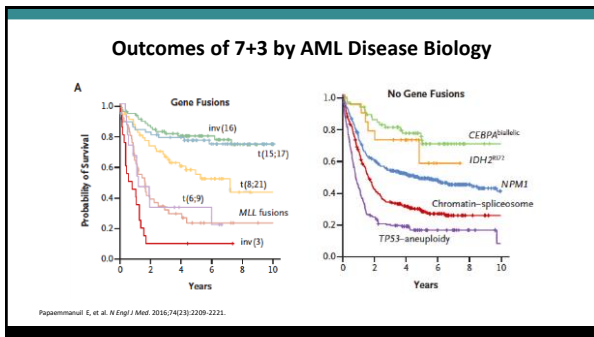
*Tagraxofusp-erzs FDA approved for the treatment of BPOCN

42

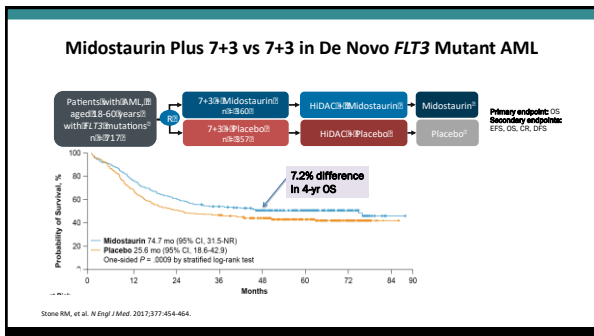
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45

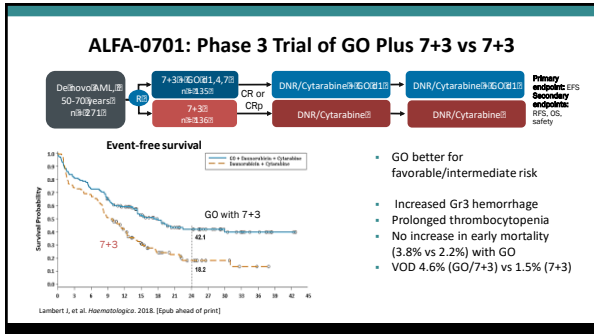


46

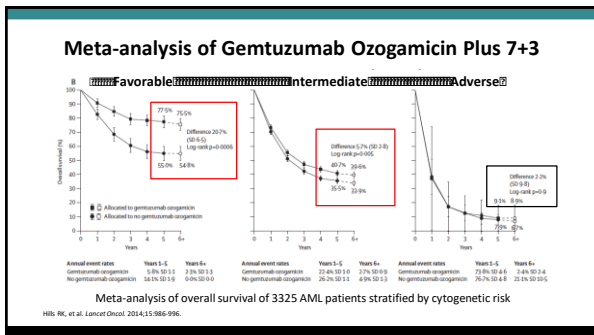


47

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48



49

AML with Myelodysplasia-related Changes (AML-MRC)

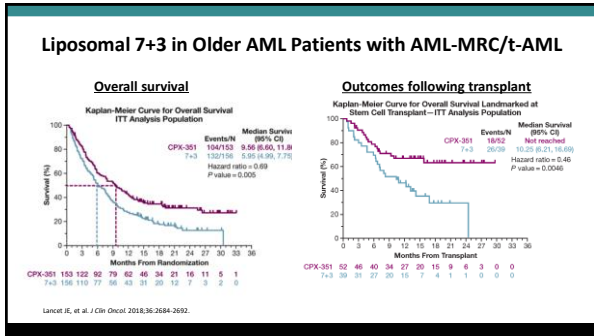
Definition: AML with a history of MDS or MDS-related cytogenetic findings

- 20% or more blasts in the peripheral blood or marrow AND
- Any of the following:
 - Previously documented MDS or MDS/MPN
 - Myelodysplasia-related cytogenetic abnormalities^a
 - Morphologic detection of multilineage dysplasia^b

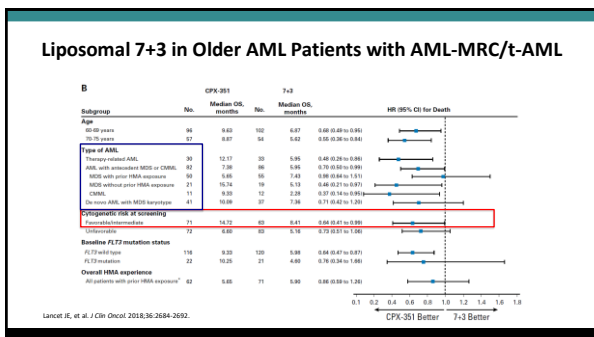
a. Complex karyotypes, chromosome 5 and 7 aberrations, multiple others
b. In the absence of CEP/alpha mutations
Arber DA, et al. Blood. 2016;127(20):2391-2405; Varadhan JK, et al. Blood. 2009;114(5):937-951.

50

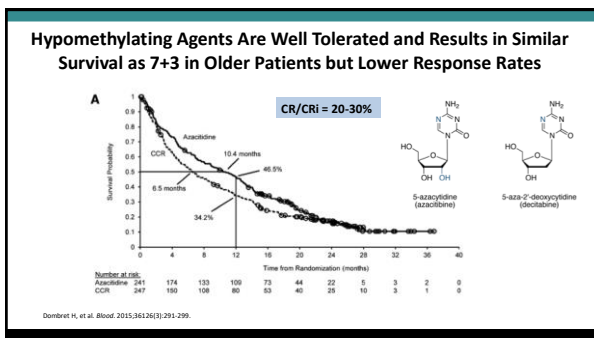
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51



52



53



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Decitabine Monotherapy in *TP53* Mutant AML Patients

- 116 patients with AML or MDS
- Decitabine 20 mg/m² qd x 10d
- 46% (53/116) responses
- High responses in unfavorable (67%) vs favorable/intermediate (34%) cytogenetics
- 100% response in *TP53* mutant (21/21) vs wild-type (32/78, 41%) patients

| No. at Risk | 0 | 200 | 400 | 600 | 800 | 1000 |
|-------------|----|-----|-----|-----|-----|------|
| CR/CR/mCR | 53 | 44 | 20 | 9 | 4 | |
| PR/SD | 36 | 17 | 10 | 8 | 4 | |
| PD/NA | 27 | 15 | 11 | 5 | 3 | |

Welch JS, et al. *N Engl J Med*. 2016;375 (21):2027-2036.

54

Venetoclax + Hypomethylating Agents - Phase 1b

| Baseline Characteristics | N = 145 |
|-----------------------------|---------|
| Median age, y | 74 |
| Age ≥75 years, % | 36 |
| Baseline BM blasts > 50%, % | 38 |
| Poor-risk cytogenetics, % | 49 |
| Baseline hydroxyurea use, % | 12 |

- CR + CRi = 97 (67%)
- Median duration of response = 11.3 mos (8.9 – not reached)
- Median overall survival = 17.5 mos (12.3 – not reached)

D'Nardo CD, et al. *Blood*. 2019;133:7-17.

55

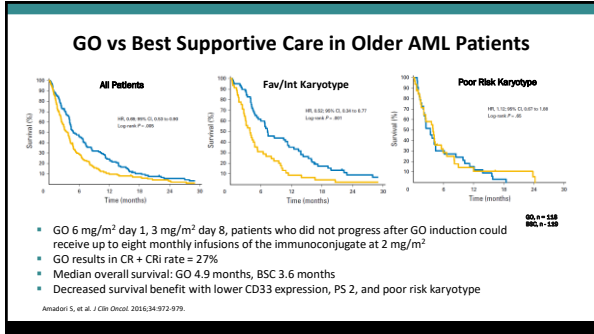
Venetoclax + Azacitidine/Decitabine: Phase 1b Subgroup Results

| Subgroup | Evaluable for responses/OS, n (%) | CR + CRi, n (%) | n for Median duration of CR + CRi | Median duration of CR + CRi (95% CI) | Median OS, mo (95% CI) |
|-------------------------|-----------------------------------|-----------------|-----------------------------------|--------------------------------------|------------------------|
| All patients | 145 | 97 (67) | 97 | 11.3 (8.9, NR) | 17.5 (12.3-NR) |
| Cytogenetic risk | | | | | |
| Intermediate | 74 (51) | 55 (74) | 55 | 12.9 (11, NR) | NR (17.5-NR) |
| Poor | 71 (49) | 42 (60) | 42 | 6.7 (4.1, 9.4) | 9.6 (7.2-12.4) |
| Age | | | | | |
| ≥75 y | 62 (43) | 40 (65) | 40 | 9.2 (6.4, 12.5) | 11 (9.3-NR) |
| <75 y | 83 (57) | 57 (69) | 57 | 12.9 (9.2, NR) | 17.7 (14.2-NR) |
| AML | | | | | |
| De novo | 109 (75) | 73 (67) | 73 | 9.4 (7.2, 11.7) | 12.5 (10.3-24.6) |
| Secondary | 36 (25) | 24 (67) | 24 | NR (12.5, NR) | NR (14.6-NR) |

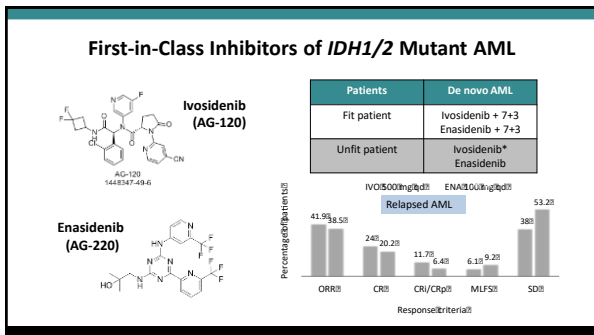
D'Nardo CD, et al. *Blood*. 2019;133:7-17.

56

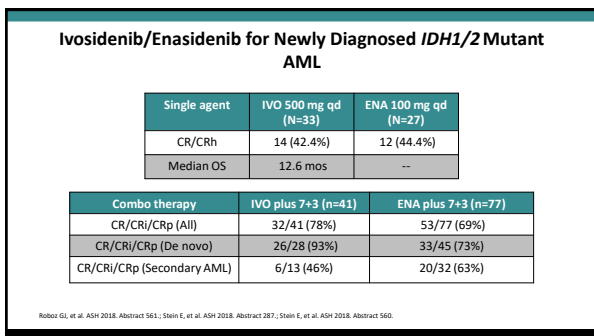
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60



61



62

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Second-generation FLT3 Inhibitors vs Midostaurin

| | Other Kinases | IC ₅₀ (plasma) |
|--------------|-------------------------|---------------------------|
| Lestaurtinib | JAK2, TrkA | 700 nM |
| Midostaurin | cKIT, PKC, PDGFR, VEGFR | 1000 nM |
| Sorafenib | cKIT, PDGFR, RAF, VEGFR | 265 nM |
| Quizartinib | cKIT, PDGFR, RET | 18 nM |
| Crenolanib | PDGFR | 48 nM |
| Gilteritinib | AXL | 43 nM |

Pratz KW, et al. Blood. 2010;115(7):1425-1432. Zarinnikar P, et al. Blood. 2009;114(14):2884-2892. Galanis A, et al. Blood. 2014;123(1):94-100. Levin MI, et al. ASCO 2015. Abstract 7003.

63

FLT3 TKI Plus 7+3 for Younger FLT3 Mutant AML Patients

| FLT3 TKI | No. patients | CR/CRi/CRh | 2-year OS | Ongoing |
|--------------------|--------------|------------|------------------------|------------------------|
| Midostaurin + 7+3 | 717 (ph 3) | 59% | 60% | AraC/DNR vs AraC/Ida |
| Quizartinib + 7+3 | 16 (ph 1) | 84% | Ph 3 ongoing | Phase 3 (7+3) ongoing |
| Crenolanib + 7+3 | 38 (ph 2) | 88% | 79% Ph 3 ongoing | Phase 3 (mido) ongoing |
| Gilteritinib + 7+3 | 30 (ph 1) | 93% | Not known Ph 3 planned | Phase 3 (mido) planned |

Stone RM, et al. N Engl J Med. 2017;377:454-464. Wang F, et al. ASH 2017; Abman L, et al. Am J Hematol. 2018;9(2):213-221; Pratz K, et al. ASH 2018.

64

AML: Take Home Points

- Timely knowledge of cytogenetics, mutations (*FLT3*, *IDH1/2*), and antigen expression (CD33) is required to select best therapy in this new era
- Choice of therapy should be individualized based on:
 - Patient age, fitness, and goals of care
 - Disease biology
 - Transplant eligibility
- Novel regimens offer survival benefits in older individuals without the need for intensive chemotherapy. Future combination regimens promise to further improve outcomes

70

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Optimizing Outcomes in Relapsed/Refractory AML: Incorporating New Treatment into Practice

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Department of Leukemia
Division of Cancer Medicine
The University of Texas MD Anderson Cancer Center
Houston, Texas

71

How Do We Best Incorporate Novel Therapies?

72

Case #1

- 58-year-old male with gingival swelling, myalgias, fevers and epistaxis. WBC 44K, Hgb 6.8 g/dl, Pts 22K. Bone marrow with 72% MPO+, CD33+ and CD123+ blasts. Diploid cytogenetics. ECOG PS 0. NPM1, IDH1, and FLT3-ITD (AR 0.49) mutations
- Started 7+3 + midostaurin, attained CR. Received 4 consolidation courses. No maintenance midostaurin
- Noted to have WBC 27K with 37% circulating blasts 5 months after last consolidation

- How would you treat this patient?...

73

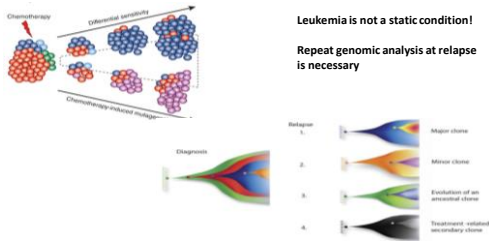
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Case #2

- 68-year-old female with history of coronary artery disease and CHF. Noted to have pancytopenia on pre-op labs for knee repair. WBC 2.8K, Hgb 8.7g/dl, Plts 128K. Bone marrow with 27% MPO+ CD33+ CD123+ blasts with MDS-related changes. Trisomy 8 cytogenetics. Received azacitidine for 8 cycles with transfusion independence and CR documented after 3 cycles, however progressive anemia noted and repeat marrow with 32% blasts. +8 and new del(20q) detected on cytogenetics. NGS panel with DNMT3A, IDH2 and SRSF2 mutations
- How would you treat this patient?

76

Individualizing Therapy at Relapse



79

Targeting FLT3 Mutations

80

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Characteristics of FLT3 Mutations in AML

- *FLT3-ITD* in ~25% and *FLT3-TKD* in ~10% AML
- More frequent in younger patients, *de novo* AML and diploid cytogenetics
- Leads to constitutive activation of FLT-3 receptor
- *FLT3-ITD* independent predictor of poor prognosis

Utzew MR, et al. *Blood*. 2015;126(7):833-841.

81

FLT3 Inhibitors in Development¹⁻⁵

¹Owens R, et al. *Expert Rev Hematol*. 2016;9:433-445. ²Cortez J, et al. *ASH* 2018. ³Adapted from Zarinikar PP, et al. *Blood*. 2009;114:2984-2992. ⁴Peri AJ, et al. *Lancet Oncol*. 2017;18:1061-1075. ⁵Kawanishi A, et al. *Blood*. 2012;120:1341.

82

Gilteritinib: ADMIRAL Phase III Trial in RR *FLT3*^{mut} AML

Key Eligibility Criteria:

- Refractory to initial induction or untreated first relapse after prior Chx (defined as CR plus CRi plus Chx)
 - Prior frontline midostaurin or sorafenib allowed
 - Prior gilteritinib or other FLT3 inhibitors excluded
- Central laboratory-confirmed *FLT3-ITD* or *FLT3-TKD* (D835/836) by PCR
- ECOG performance status ≥ 2
- Normal liver, renal function
- QTcF ≤ 450 msec by central ECG reading

Co-Primary Endpoints: OS, CR/CRh rate
Key Secondary Endpoints: EFS, CR rate

***Salvage chemotherapy regimen was selected prior to randomization**
MEC (mitoxantrone, etoposide, and cytarabine) } High intensity (1-2 cycles)
FLAG-IDA (fludarabine, cytarabine, idarubicin, and G-CSF)
Low-dose cytarabine } Low intensity (given until disease progression or intolerance)
Azacitidine

Peri A, et al. *ASCO* 2019.

84

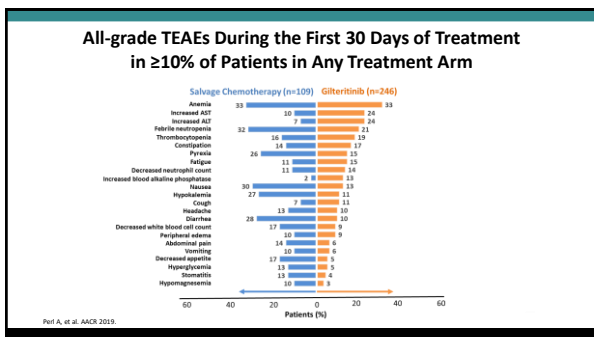
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Response Outcomes (ITT Population: N=371)

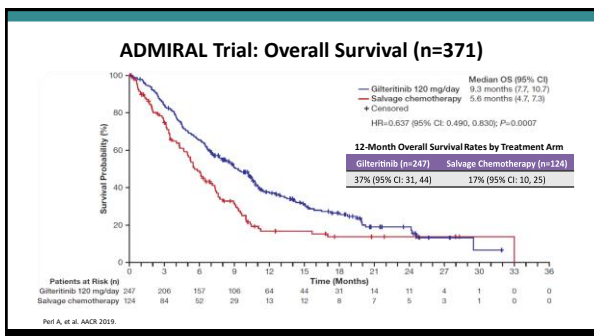
| Parameter* | Giilteritinib (n=247) | Salvage Chemotherapy (n=124) |
|--|-----------------------|------------------------------|
| CR, n (%) | 52 (21) | 13 (11) |
| CRh, n (%) | 32 (13) | 6 (5) |
| CRi, n (%) | 63 (26) | 14 (11) |
| CRp, n (%) | 19 (8) | 0 (0) |
| CRc, n (%) | 134 (54) | 27 (22) |
| CR/CRh, n (%) | 84 (34) | 19 (15) |
| PR, n (%) | 33 (13) | 5 (4) |
| ORR, n (%) | 167 (68) | 32 (26) |
| NR, n (%) | 66 (27) | 43 (35) |
| Median duration of drug exposure (range), months | 4.1 (0.1-29.1) | 0.9 (0.2-7.1) |
| Median time to achieve CR (95% CI), months | 1.8 (0.9, 9.5) | 1.1 (0.8, 2.9) |
| Median DoR* (95% CI), months | 11.0 (4.6, NE) | 1.8 (NE, NE) |
| Allogeneic HSCT, n (%) | 63 (26) | 19 (15) |

*Response was not evaluable in 14 patients (6%) in the giilteritinib arm and in 49 patients (40%) in the salvage chemotherapy arm
*Duration of remission was defined as the duration of CR/CRh

85

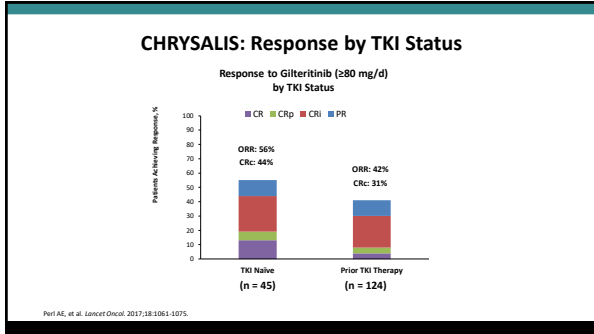


86



87

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88

Molecular Response: Chrysalis

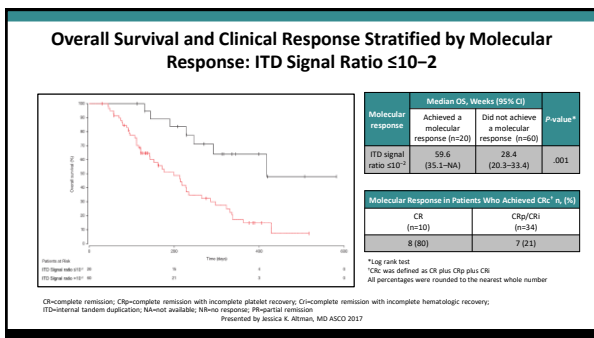
| Response Outcomes in All FLT3-ITD ^{mut} Patients | N=80 |
|---|------------------|
| Patients who had a molecular response (ITD signal ratio $\leq 10^{-2}$), n (%) | 20 (25) |
| Patients who had MMR (ITD signal ratio $\leq 10^{-3}$), n (%) | 18 (23) |
| Patients who had MRD negative status (ITD signal ratio $\leq 10^{-4}$), n (%) | 13 (16) |
| Median time to achieve minimum ITD signal ratio, weeks (range) | 8.2 (3.7–64) |
| Median OS, weeks (95% CI) | 32.6 (25.1–42.4) |

- Three patients with a molecular response underwent allogeneic HSCT

CI=confidence interval; FLT3=FLT3-like tyrosine kinase 3; ITD=internal tandem duplication; MRD=minimum residual disease; MMR=minor molecular response; OS=overall survival

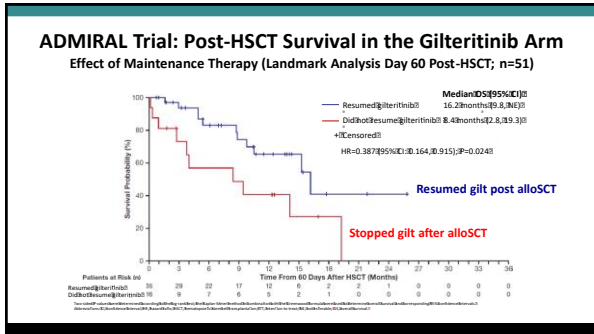
Presented by Jessica K. Altman, MD ASCO 2017

89

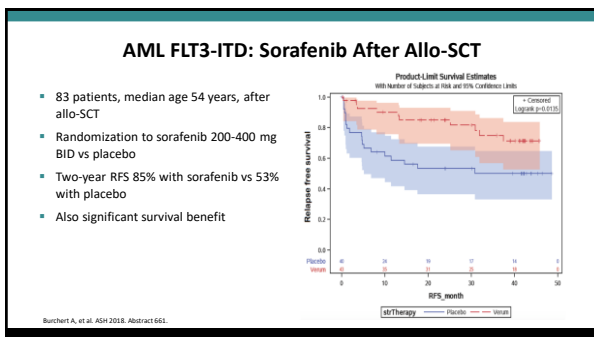


90

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91



92

Targeted Therapy: FLT3 Inhibitors

- FLT3-inhibitors are safe and effective targeted therapeutics
 - Well tolerated
 - Improved outcomes
 - Responses short-lived as single agents
 - Resistance is common with single agents and depends on the unique FLT3i
- FLT3 inhibitors improve outcomes in newly diagnosed and R/R patients, rational combinations to prevent resistance will further improve upon FLT3 outcomes

93

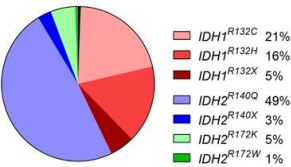
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Targeting IDH1 and IDH2 Mutations

96

Characteristics of mIDH AML

- IDH mutations occur in ~20% of AML**
 - Most (~85%) occur in *de novo* diploid or +8 AM
 - IDH1 in ~8% AML, IDH2 in ~12% AML
 - ↑ prevalence with ↑ patient age
- Hot-spot mutations in enzymatic active site**
 - IDH1-R132, IDH2-R140 or IDH2-R172
- Often early mutational events**
 - Ancestral in 20% IDH1 and 35% IDH2 cases
- Can be acquired at progression**
 - ~10-15% of AML from MDS
 - ~20-25% of AML from MPN



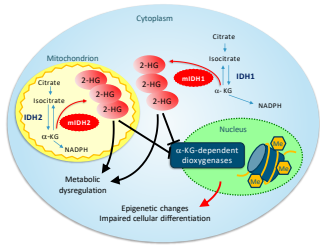
| Mutation | Percentage |
|-----------------------|------------|
| IDH1 ^{R132C} | 21% |
| IDH1 ^{R132H} | 16% |
| IDH1 ^{R132X} | 5% |
| IDH2 ^{R140Q} | 49% |
| IDH2 ^{R140X} | 3% |
| IDH2 ^{R172K} | 5% |
| IDH2 ^{R172W} | 1% |

Dang L, et al. Trends Mol Med. 2010;16(9):387-397; Chou WC, et al. Leukemia. 2011;25(2):246-253; Molenaar RJ, et al. Leukemia. 2015;29(11):2134-2142.

97

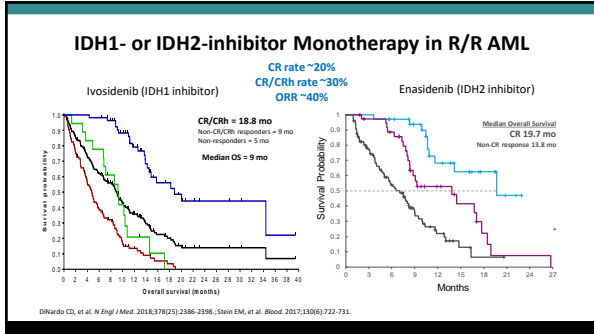
Pathophysiology of IDH Mutations

- mIDH results in accumulation of the oncometabolite 2-HG which competitively inhibits αKG-dependent reactions**
- 2HG leads to DNA and histone hypermethylation, and a resultant block in differentiation**

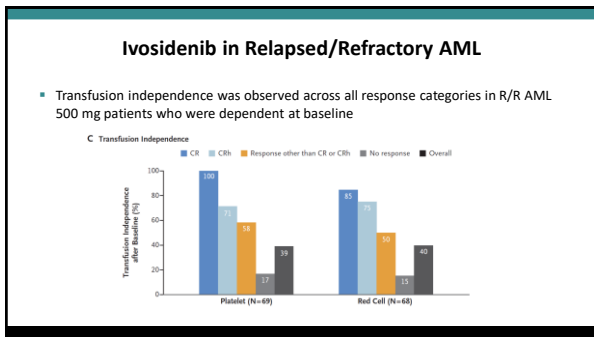


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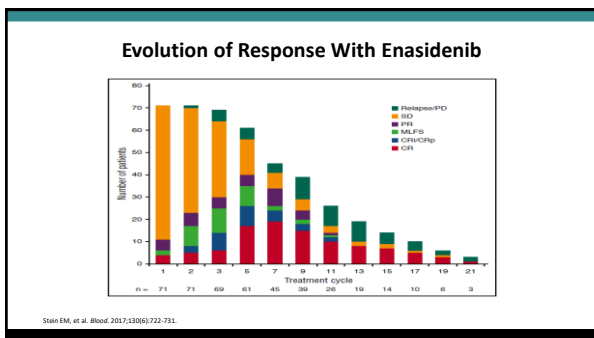
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99



100



101

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Identification and Treatment of Clinical IDH-DS

Table 1. Frequency of Signs and Symptoms Consistent With IDH-DS*

| Sign or Symptom | Patients With IDH-DS, No. (%) |
|--|-------------------------------|
| Dyspnea | 28 (85) |
| Unexplained fever (body temperature of 38.0°C for ≥ 3 d) | 26 (79) |
| Pulmonary infiltrates | 24 (71) |
| Hypoxia | 19 (58) |
| Acute kidney injury (CTCAE grade ≥ 2) | 14 (42) |
| Pleural effusion | 14 (42) |
| Bone pain or arthralgia | 9 (27) |
| Lymphadenopathy | 8 (24) |
| Edema | 8 (24) |
| Disseminated intravascular coagulopathy | 7 (21) |
| Edema or weight gain of ≥ 5 kg from screening | 7 (21) |
| Pericardial effusion | 5 (15) |

Flowchart: Suspicion of DS (≥ 1 symptom or sign*) → Desamethasone 10-mg twice daily** → (If response) Continue until symptom improvement → Start tapering after resolution of symptoms; (If no response) Stop enasidenib*** → Resume after resolution of symptoms.

Notes:
 *Main symptoms and signs: Dyspnea, fever, rash, pulmonary infiltrates, pleural effusion, renal dysfunction.
 **With other supportive measures: Broad spectrum antibiotics for possible underlying infection; Cytoreductive therapy if leukocytosis; Hypernatremia management if tumor lysis syndrome.
 ***Symptoms will last several days until improvement due to the long half-life of enasidenib.

Abbreviations: CTCAE, Common Terminology Criteria for Adverse Events™; IDH-DS, isocitrate dehydrogenase differentiation syndrome.
 *Signs and symptoms included in this table are based on retrospective differentiation syndrome review committee review of clinical records.
 **Patients may have had multiple symptoms.
 Fathi A, et al. *JAMA Oncol*. 2018;4(8):1106-1110; Abou Dalil I, et al. *Ther Adv Hematol*. 2018;9(7):163-173.

102

Impact of IDH1 Mutation-clearance Status with AG120

Duration of CR+CRh

Overall survival

- Patients with "deep IDH1 mutational clearance" had improved DOR and OS vs those in CR/CRh with persistent IDH1 mutation detected
- Defined as a reduction in mIDH1 VAF to below the limit of detection by digital PCR (0.02-0.04%)

Polyva D, et al. *EMO*. 2018.

103

Impact of Co-occurring Mutations at Start of Enasidenib

A Co-Mutational Burden vs Response ($p < 0.0001$)

B Low Mutational Burden (n=27, CRh=70.4%) vs High Mutational Burden (n=22, CRh=27.3%)

C mNRAS (G15, G19, G45) Response (n=14)

D Co-Mutational Burden vs mNRAS ($p < 0.0001$)

E mNRAS vs mIDH2 VAF ($p < 0.0001$)

Amatangelo MD, et al. *Blood*. 2017;130(6):732-741.

104

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IDH Inhibitor Therapy Take-Aways

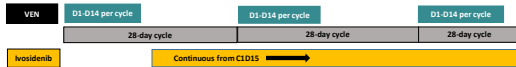
- Ivosidenib and enasidenib are safe and effective oral targeted therapies for patients with IDH1 or IDH2 mutant AML
 - Well-tolerated oral therapies
 - Durable responses
 - Improved outcomes
- Ivosidenib and enasidenib in combination therapies will further enhance responses
 - In combination with standard agents; ie, 7+3 and azacitidine
 - With other small molecule and targeted therapies (FLT3i, MEKi, VEN, others)

105

Phase Ib/II Open-label Study of Ivosidenib in Combination with Venetoclax

Phase 1b: To determine the safety and tolerability, maximum tolerated dose (MTD) and recommended phase 2 dose (RP2D) of the combination of ivosidenib and venetoclax +/- azacitidine

DLT Period = 2 cycles to ensure adequate safety evaluation of the combination



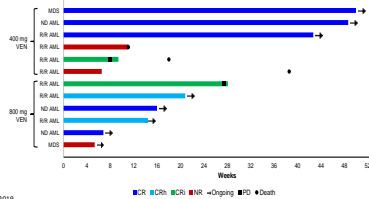
| Phase 1 Cohorts | Venetoclax | Ivosidenib | Azacitidine |
|--------------------------|-------------------|-------------------|-------------------------------|
| +2 (Target dose) | 400 mg once daily | 500 mg once daily | 75 mg/m ² days 1-7 |
| +1 (Target dose) | 800 mg once daily | 500 mg once daily | |
| 0 (Starting dose) | 400 mg once daily | 500 mg once daily | |

©Naraino CD, et al. EHA 2019

106

Response and Duration

- ORR was 75%, which included CR (42%), CRh (17%), and CRi (17%)
- Median time to first response was one month (range 1-3 months)



©Naraino CD, et al. EHA 2019

108

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Case #1

- 58-year-old male with gingival swelling, myalgias, fevers and epistaxis. WBC 44K, Hgb 6.8 g/dl, Plts 22K. Bone marrow with 72% MPO+, CD33+ and CD123+ blasts. Diploid cytogenetics. ECOG PS 0. NPM1, IDH1, and FLT3-ITD (AR 0.49) mutations
 - Started 7+3 + midostaurin, attained CR. Received 4 consolidation courses. No maintenance midostaurin
 - Noted to have WBC 27K with 37% circulating blasts 5 months after last consolidation
- How would you treat this patient?...

109

Case #2

- 68-year-old female with history of coronary artery disease and CHF. Noted to have pancytopenia on pre-op labs for knee repair. WBC 2.8K, Hgb 8.7g/dl, Plts 128K. Bone marrow with 27% MPO+ CD33+ CD123+ blasts with MDS-related changes. Trisomy 8 cytogenetics. Received azacitidine for 8 cycles with transfusion independence and CR documented after 3 cycles, however progressive anemia noted and repeat marrow with 32% blasts. +8 and new del(20q) detected on cytogenetics. NGS panel with DNMT3A, IDH2 and SRSF2 mutations
- How would you treat this patient?

112

Conclusions

- **Targeted therapy and precision medicine:** transitioning from population-based → individual "genotype-phenotype" based treatment
- Important for disease classification and prognostication
- Can now be used to guide and inform clinical practice
 - Optimal treatment strategies (at diagnosis and relapse)
 - Addition of target-specific drugs
 - *Improved treatment options for poor-risk patients still needed*
- NGS-panel testing for prognostic and therapeutically informative mutations should be performed at diagnosis and relapse time-points to optimize informed decision-making

115

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**Evolving Treatment Paradigms in AML:
New Data and Clinical Trials That Could Change
Clinical Practice**

Naval Daver, MD
Associate Professor
Department of Leukemia
The University of Texas MD Anderson Cancer Center
Houston, Texas

121

New Therapies for AML...and Novel Combinations

...for many populations, including those at high risk or defined by molecular abnormalities

AML with actionable mutations, molecular, or high-risk features (eg, age)
Combination therapy with HMAs for older patients

Novel cytotoxics (CPX-351, vosaroxin)
Emerging/next-generation HMAs (BGI-110, CC-885, ASTX727)

Immune checkpoint inhibitors (Nivolumab, Pembrolizumab)

Targeted therapies (FLT3, IDH, Bcl-2, MDM2, MCL-1, APR253)

Novel antibodies (Gemtuzumab, IMGN 33 and 123, AMG-330, MGD, CD33, CD47)

HMA-hypomethylating agent

122

ASTRAL-1: Phase 3 Study Design

Treatment-naïve AML ineligible for intensive induction¹
N = 800 randomized

1:1 randomization

Guadecitabine (5GI-110)
• 60 mg/m² SC x 5 days

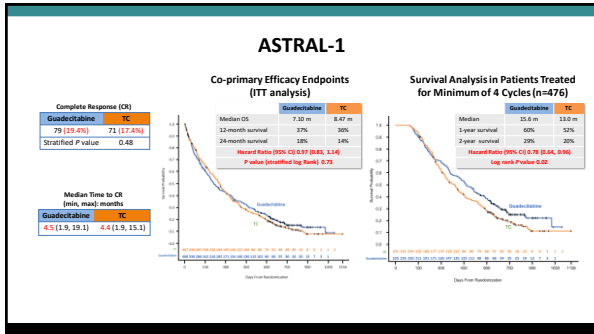
Treatment Choice (TC)
• Decitabine (DEC)
• Azacitidine (AZA)
• Low Dose Ara-C (LDAC)

Co-Primary Endpoints:
• Complete Response (CR) rate
• Overall Survival (OS)

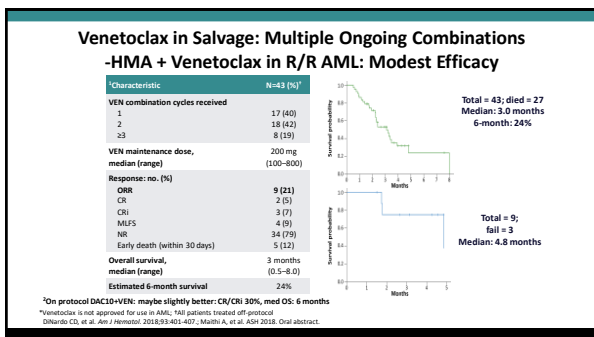
¹Age ≥ 75 years or older, or major organ comorbidities, and poor Eastern Cooperative Oncology Group (ECOG) Performance Status 2-3
Penaud P, et al. BHA 2019.

123

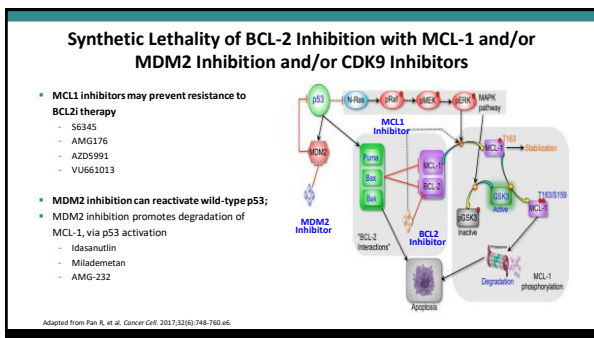
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124

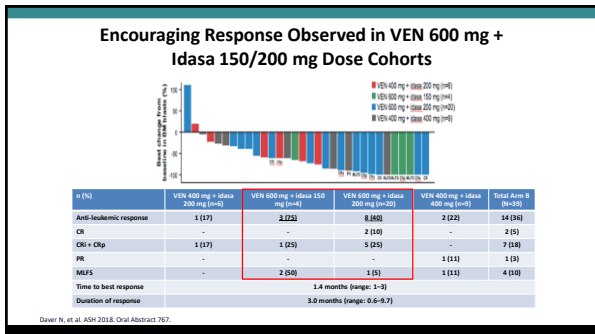


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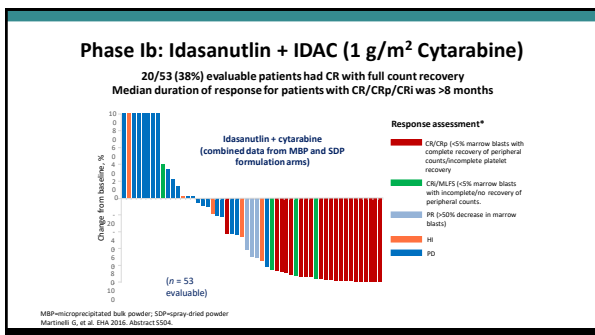


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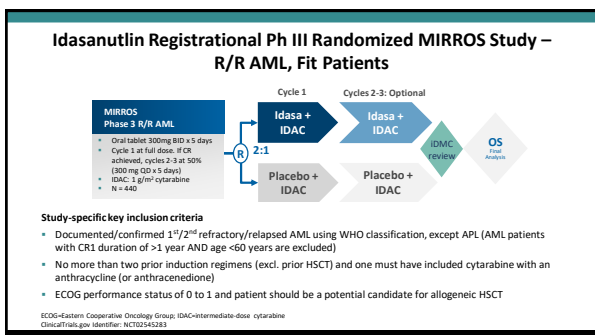
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127

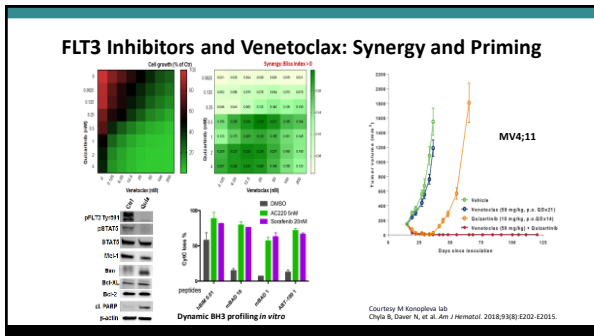


128

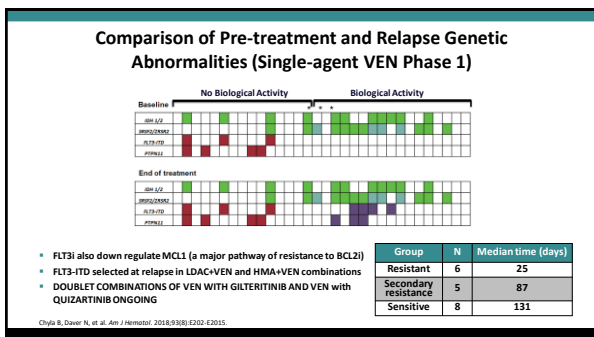


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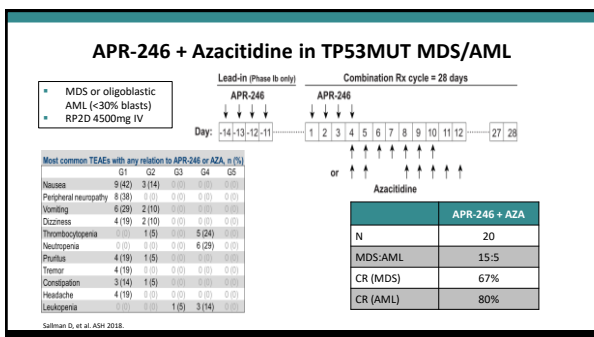
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130



131



132

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Major Approaches to Immune Therapies in AML

Two major approaches:

1. Antibody drug conjugates
2. T-cell based therapies
 - a. Bi-specific antibodies (CD3 x AML antigen)
 - b. Immune checkpoint-based approaches
 - c. CAR T (CAR NK)

CTLA-4—cytotoxic T lymphocyte antigen-4; DC—dendritic cell; LAG-3—lymphocyte activation gene-3; MH—macrophage; MHC—major histocompatibility complex; NK—natural killer cell; PD-1—programmed death-1; PD-L1—programmed death-ligand 1; TCR—T cell receptor; T-eff—T-effector cells; TIM-3—T cell immunoglobulin and mucin domain-containing protein; T-reg—T-regulatory cells.
Arai H (Dover NJ, et al. Curr Opin Hematol. 2018;25:136-145.

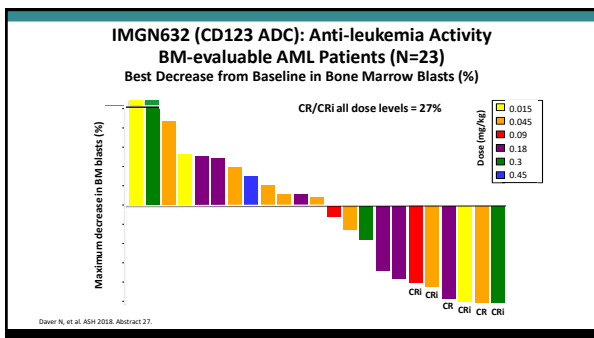
133

IMGN632: A Novel CD123-Targeting ADC

- Novel anti-CD123 antibody
 - Higher affinity binding to CD123
 - Unique epitope in extracellular domain
- Novel IGN payload (DGN549)
 - DNA-alkylating activity, single strand DNA breaks (vs. double strand)
 - 10-20x more potent than the IGN in IMGN779
 - Uniform loading of 2 IGN molecules per antibody
- Stable peptide linker
 - Protease cleavable
 - Confers stability in circulation, and controlled intracellular payload release

Korvan Y, et al. Blood Adv. 2018;2:848-858; Dover N, et al. ASH 2018; Abstract 27.

134



135

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Flotetuzumab (CD3 x CD123) Phase 1 Study Design

Dose Escalation

Single Patient
Dose Escalation
N=14

**3 + 3 Multi-patient
Dose Escalation**

N=33

Expansion Cohort

R/R AML
Recommended Phase 2
Dose
(RP2D)
N=11

Key entry criteria

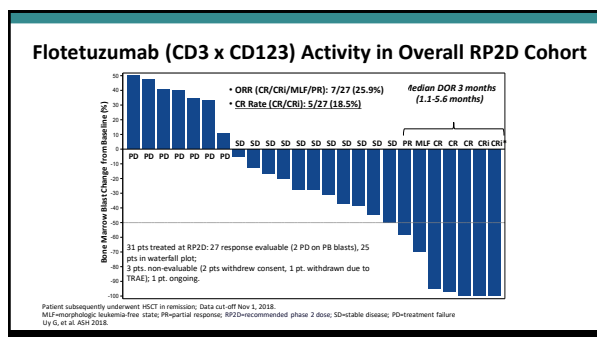
- Relapsed/refractory AML unlikely to benefit from cytotoxic chemotherapy
 - Refractory to ≥2 induction attempts
 - First relapse with initial CR duration of <6 months or any prior unsuccessful salvage
 - Second relapse or higher
 - HMA failure
- No prior allogeneic hematopoietic cell transplant

Study objectives

- Safety and preliminary clinical activity
- Optimize delivery and supportive care (manage CRS while minimizing corticosteroid use)
- Define PK, PD and PK/PD relationships

CR=cytotoxic relapse syndrome
UY G, et al. ASH 2018.

136



137

AZA + Nivo in Relapsed AML: Response (N = 70)¹

| Best response | N (%), Median (Range) | |
|---|-----------------------|---------------|
| | Azacitidine/Nivolumab | Control |
| Overall Response Rate | 23 (33) | 35 (20) |
| CR | 4 (6) | 17 (10) |
| CRi/CRp | 11 (16) | 15 (9) |
| PR | 1 (1) | 1 (1) |
| HI* (6 months+) | 7 (10) | 2 (1) |
| Stable disease (6 months+)[†] | 6 (9) | NA |
| Non responders | 41 (58) | 131 (76) |
| Median cycles to response | 2 [1-13] | 2 [1-6] |
| Median follow up, in months | 13.3 [8.2-25.5] | 51 [0.1-84.8] |
| ORR in prior HMA-naïve (N=25) | 52% | 19% (P<0.001) |
| CR/CRi in prior HMA-naïve | 28% | 16% (P=0.18) |
| ORR in prior HMA-exposed (N=45) | 22% | 23% |
| CR/CRi in prior HMA-exposed | 18% | 23% |

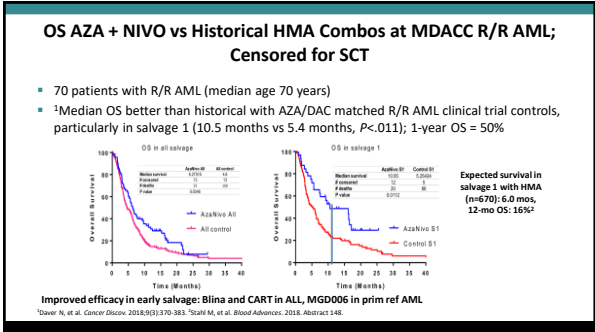
How does this compare to other HMA Rx in R/R AML??

- ²Single-agent AZA/DAC (n = 670) in prior HMA-naïve retro analysis, ORR = 23%, CR/CRi rate = 16%
- ^{3,4}AZA/DAC + VEN in prior HMA-naïve: CR/CRi: 30-35%

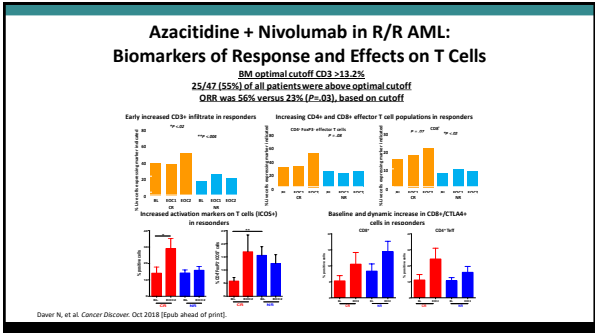
*HI: Response maintained ≥6 months
¹Oliver N, et al. Cancer Discov. 2018;8(9):1370-1383. ²Stathi M, et al. Blood Advances. 2018 (Epub). ³Marino C, et al. Am J Hematol. 2016;91(9):401-407. ⁴Goldberg A, et al. ASH 2018. Abstract 1333.

138

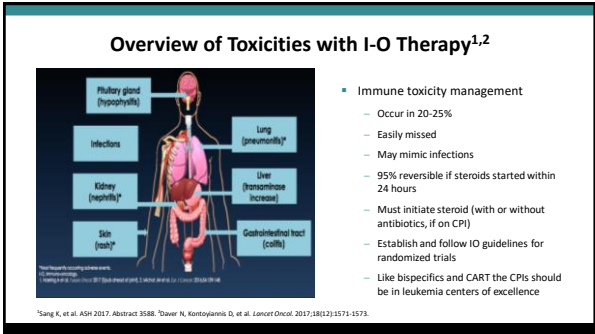
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139



140

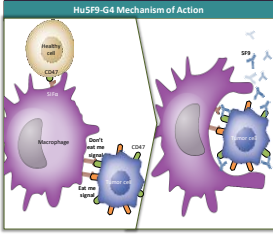


141

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Background on CD47 in AML

- Macrophages are a key part of the innate immune system
 - Engulf cancer cells
 - Recruit, activate, and present cancer cell antigens to T cells
- CD47 is a "do not eat me" signal on cancer cells that enables them to evade macrophages
- Increased CD47 expression predicts worse outcomes for AML patients
- Hu5F9-G4 (5F9) targets CD47 on tumor cells, inducing macrophage phagocytosis

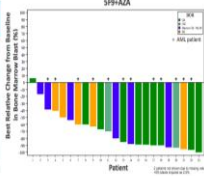


AML-associated myeloid leukemias: CD47-miR145- associated protein
Sallman DA, et al. 2019 ASCO. Abstract 7005

142

Anti-leukemic Activity is Observed with 5F9 Monotherapy and in Combination with AZA in AML and MDS

| Best Overall Response | RR AML/MDS SP9 mono n=10 | TL AML SP9+AZA n=14 | ORR MDS SP9+AZA n=17 |
|---------------------------------|-----------------------------------|------------------------------|-------------------------------------|
| ORR | 1 (10%) | 9 (64%) | 11 (100%) |
| CR | 0 | 5 (36%) | 6 (35%) |
| CR | 0 | 2 (14%) | - |
| PR | 0 | 0 | 0 |
| MRFU Median CR | 1 (10%) | 2 (14%) | 4 (36%) 2 with response Chrom |
| Transfusions Requirement (d) | - | - | 1 (9%) |
| ID | 7 (70%) | 5 (36%) | 0 |
| PD | 2 (20%) | 0 | 0 |



Response rate of best overall response (ORR) and CR. CR=complete remission. MRFU=median relapse-free survival. ID=intermediate delinquency. PD=partial duration. PR=partial response. TL=total leukemic count. AZA=azacitidine. SP9=5F9. n=number of patients. n/N=number of patients with response/total number of patients. n/N(%)=number of patients with response/total number of patients (%).

- 5F9 monotherapy has an ORR of 10% in r/r AML/MDS
- 5F9+AZA has an ORR of 100% in MDS, 64% in AML, which compares favorably to AZA monotherapy ORR. Median time to response is more rapid (1.9 months) than AZA alone

143

Conclusions

- HMA+VEN outstanding in frontline unfit AML but response and OS in R/R AML modest. Novel combinations VEN+MDM2, VEN+MCL1, VEN+FLT/IDH appear encouraging. Triplets ongoing/planned
- MDM2i (Idasanutlin) and E-selectin-I (GMI-1271) in phase 3, registrational trials
- Immune therapies (novel and safer ADCs, bispecific Ab, CPI based) may be next major wave of development in AML, multiple trials ongoing
- AML may be following MM paradigm, effective and safe doublets/triplets to improve PFS/OS with lower toxicity/mortality

145