



Precision Medicine In AML: The Goals of MyeloMATCH

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knowledge changing life



Disclosures

- Research Funding: Jazz Pharmaceuticals
- Pharmaceutical Consulting: Abbvie, Kura Oncology, MERCK, Curio Science, Celgene, Incyte Corporation, Nkarta Biotechnology
- I will be discussing the off-label use of medications and will indicate when that is the case

knowledge changing life

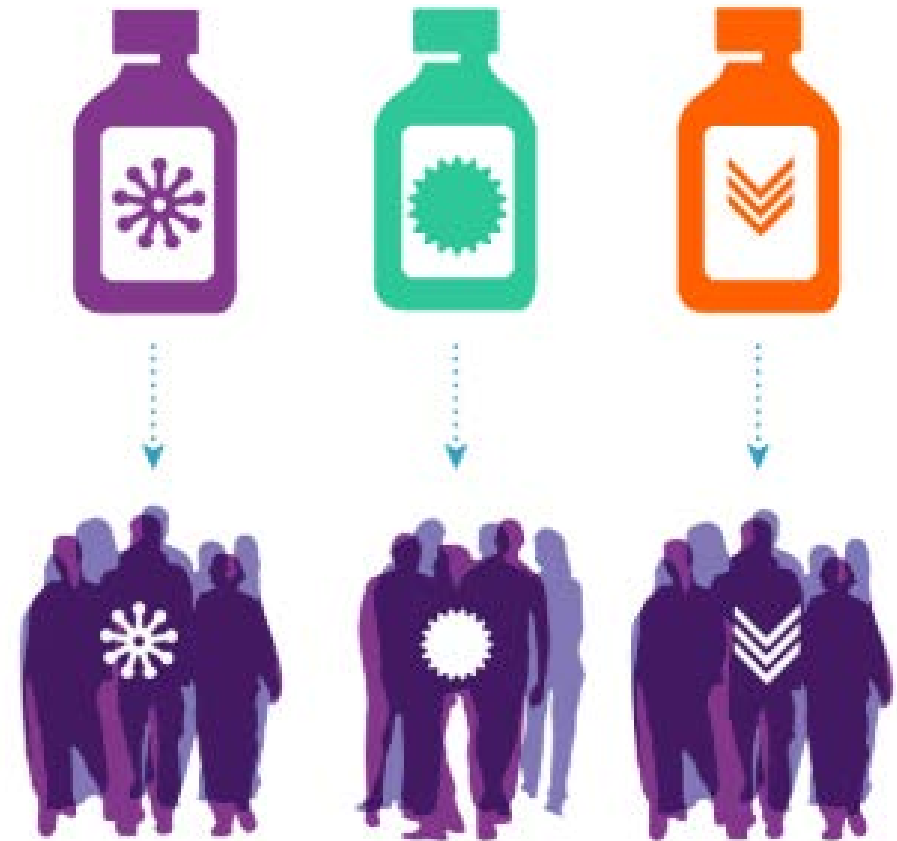


Topics of the Day

- Background
 - What is meant by “precision medicine?”
 - How should we think about Acute Myeloid Leukemia treatment?
 - What is MyeloMATCH?
- Current Standards of Care
 - Who, When and Why
- Where are we falling short?
- What can we do to address these unmet needs?

In the Ideal....

Precision Medicine is defined as:
“A form of medicine that uses information about a person’s own genes or proteins to prevent, diagnose, or treat disease”



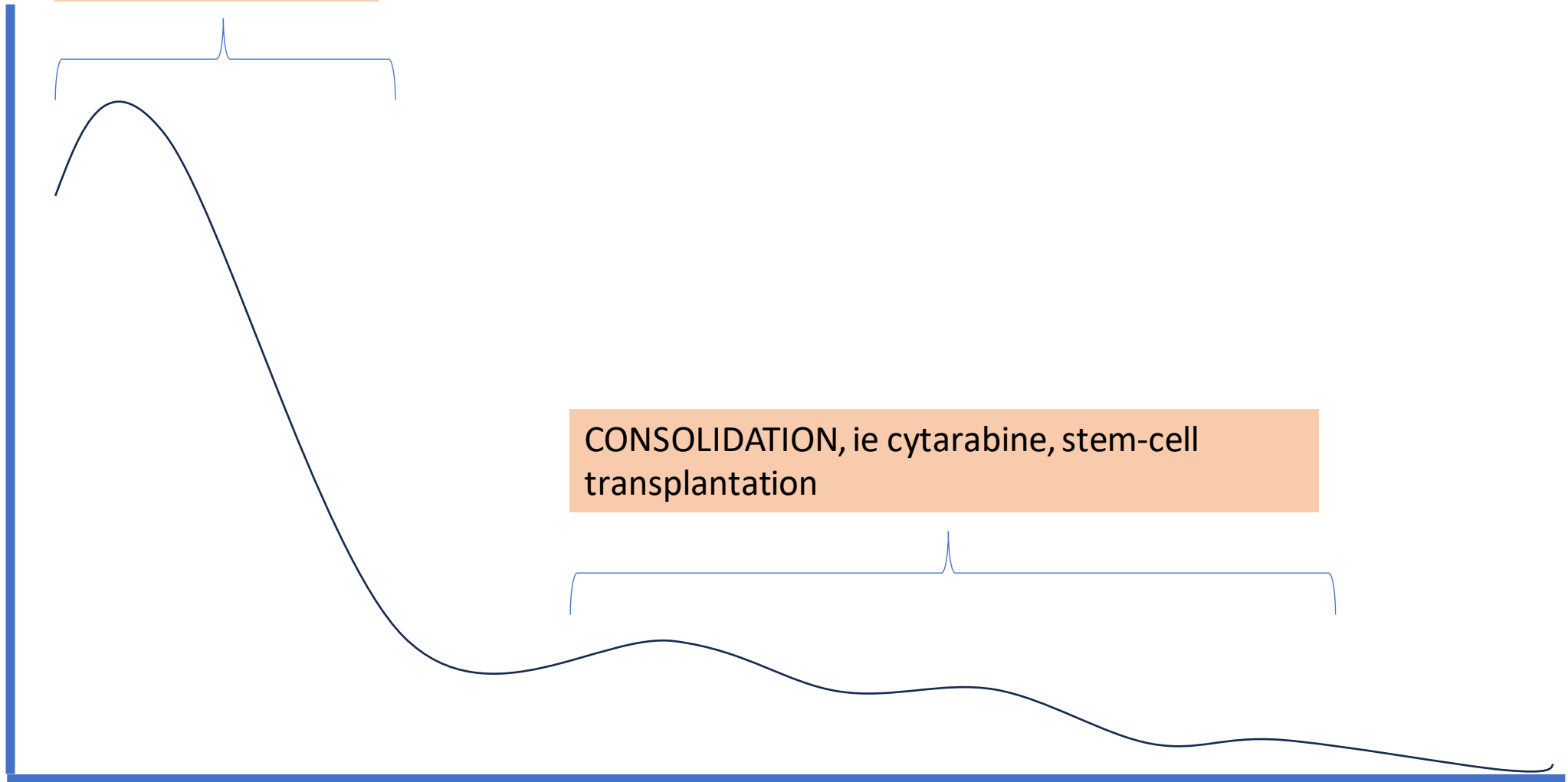
In Reality: for most of the history of hematology, the available therapeutic options, rather than disease or even patient characteristics, have dictated treatment for AML

The Traditional Framework for AML Therapy

- Induction – Eliminate the bulk of the leukemic clone
- Consolidation – Eliminate remaining leukemic cells
 - Cytotoxic Chemotherapy
 - Cellular Therapy in the form of a stem-cell transplant
- Measurable Residual Disease
 - Detectable disease present after Induction and Consolidation

LEUKEMIA BURDEN

INDUCTION, ie 7+3



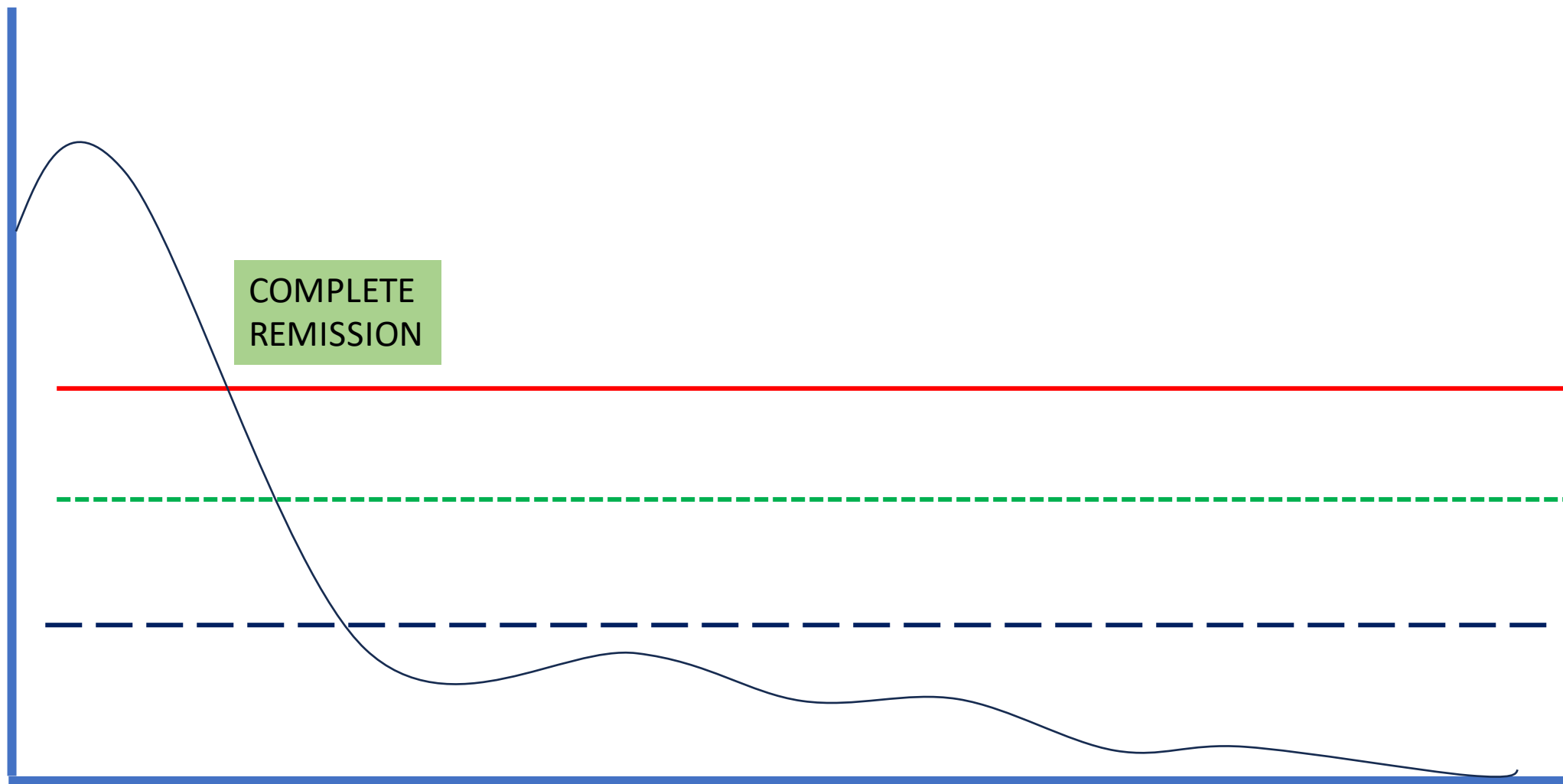
CONSOLIDATION, ie cytarabine, stem-cell transplantation

TIME

The Traditional Framework for AML Therapy

- Induction – Eliminate the bulk of the leukemic clone – Goal is
- Complete Remission
 - Morphologically with less than 5% leukemia cells in the marrow, count recovery
- Measurable Residual Disease
 - Detectable disease present after therapy using highly sensitive techniques
 - Generally measured after induction and first cycle of consolidation
 - Can be very informative prior to stem-cell transplant
 - Measured with molecular sequencing or flow cytometry

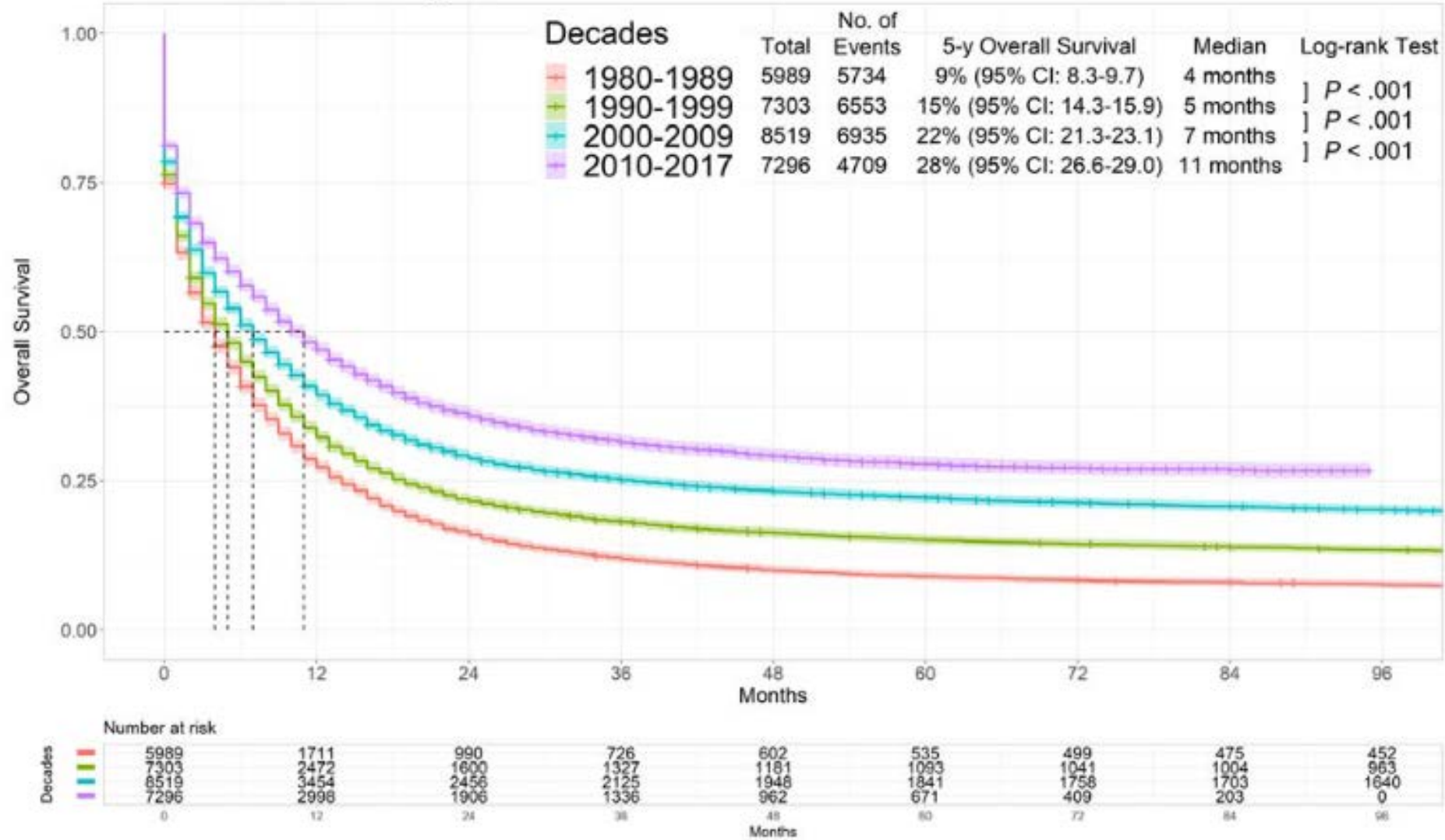
LEUKEMIA BURDEN



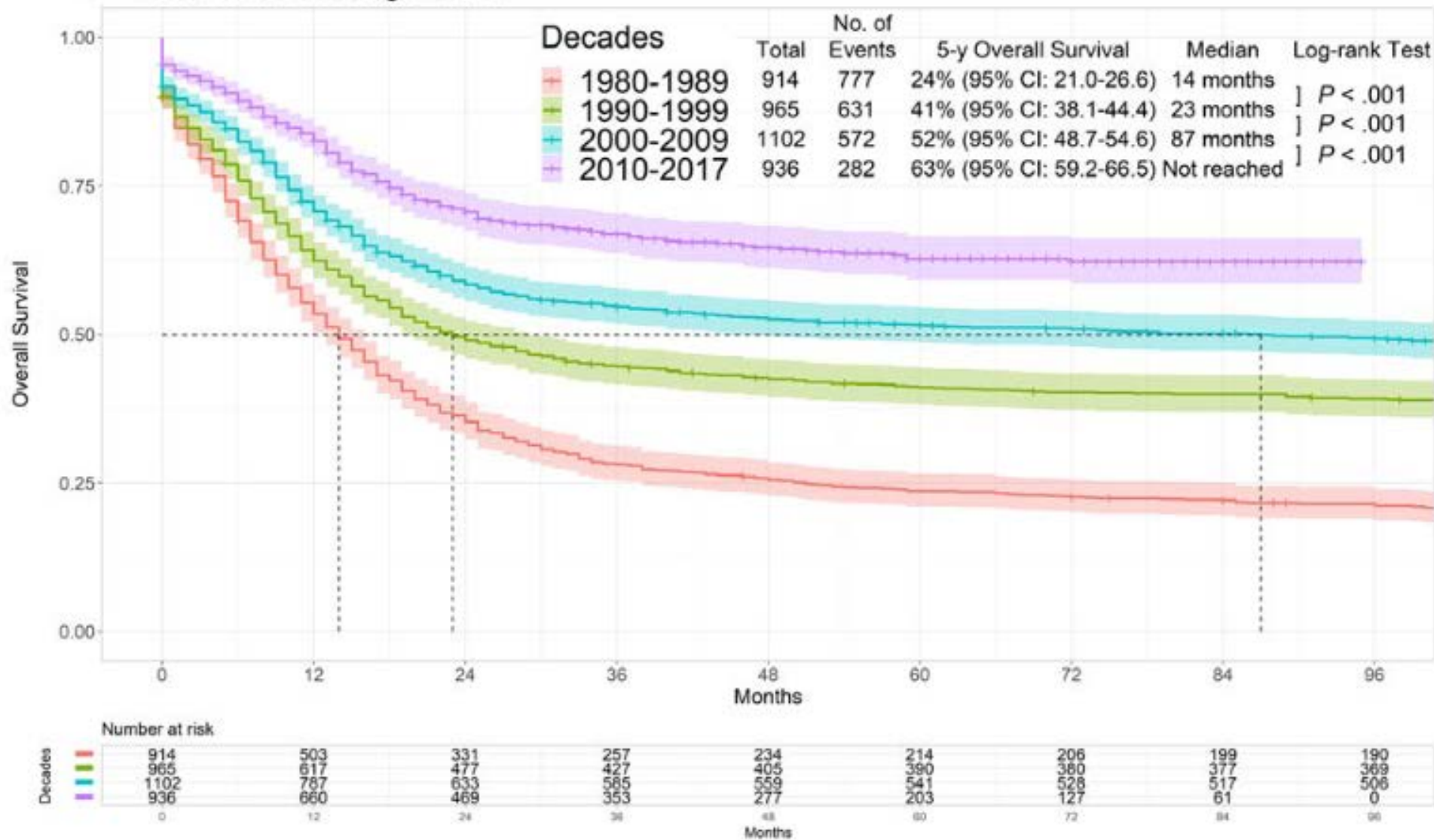
COMPLETE
REMISSION

TIME

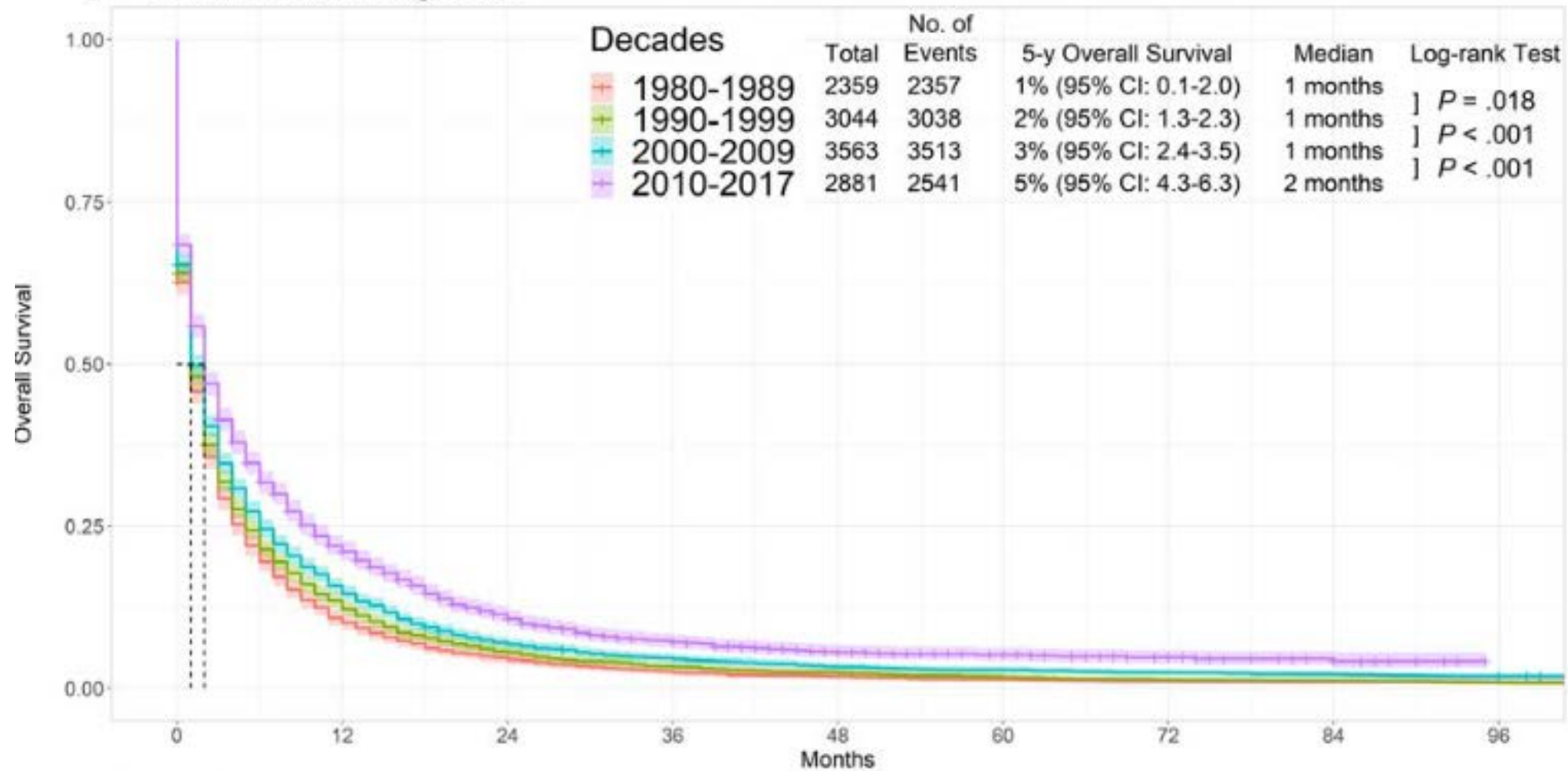
A SEER All AML: All Ages



C SEER All AML: Age 15-39



F SEER All AML: Age ≥70



Decades	0	12	24	36	48	60	72	84	96
1980-1989	2359	256	112	62	45	33	26	24	22
1990-1999	3044	411	174	104	77	55	42	37	28
2000-2009	3563	561	249	164	119	100	85	77	65
2010-2017	2881	540	236	129	71	45	26	12	0

ERA	REGIMEN: Newly-Diagnosed Patients	POPULATION	EXPECTATIONS
1970s-2000	Anthracycline – Based (7+3)	Fit	CR rates: 60%-70% Med OS: Highly Variable
	Hypomethylating Agent	Older, Less-Fit	CR rates 17% Med OS: 10 mo
2017	Midostaurin + 7 + 3	FLT-3mut + Fit + Young	CR rates: 59% Med OS: 74 months
	CPX-351	Older, High-Risk	CR rates: 47% Med OS: 9.5 mo
	Gemtuzumab + Cytotoxic Agents	Fit + Fav/Inter Cytogenetics	CR rates: 70% Med OS: 27 mo
2018-2023	Azacitadine + Venetoclax	Older, Less-Fit	CR/CRi Rates: 66% Med OS: 14 mo
	Cytarabine + Glasdegib	Older, Less-Fit	CR: 17% Med OS: 8.8 mo
	Cytarabine + Venetoclax	Older, Less-fit	CR/Cri: 54% Med OS: 10 mo

What is required to improve our outcomes?

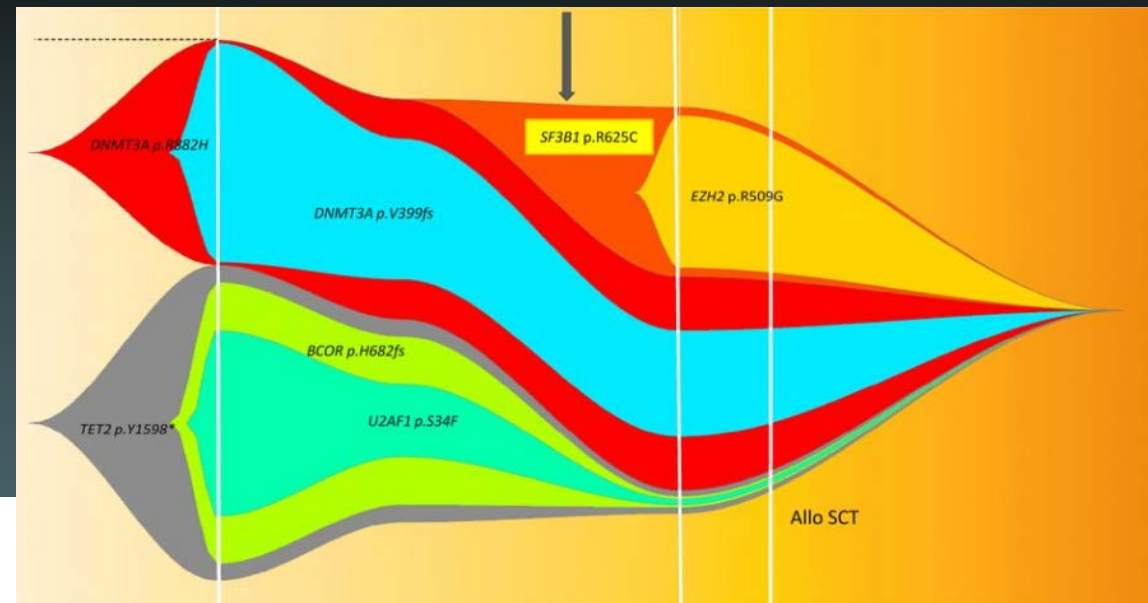
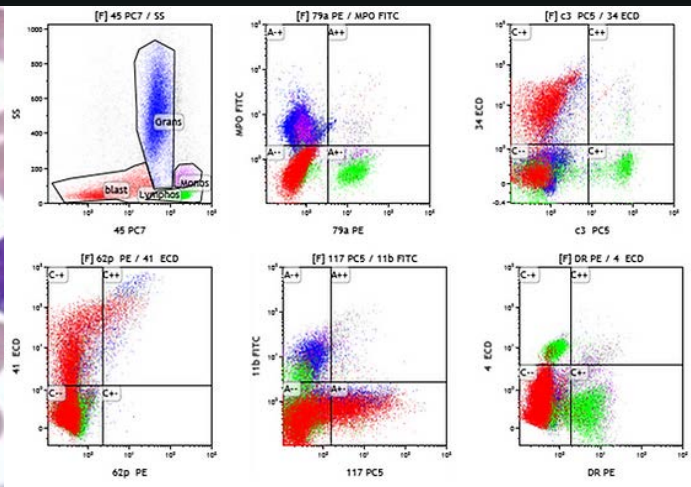
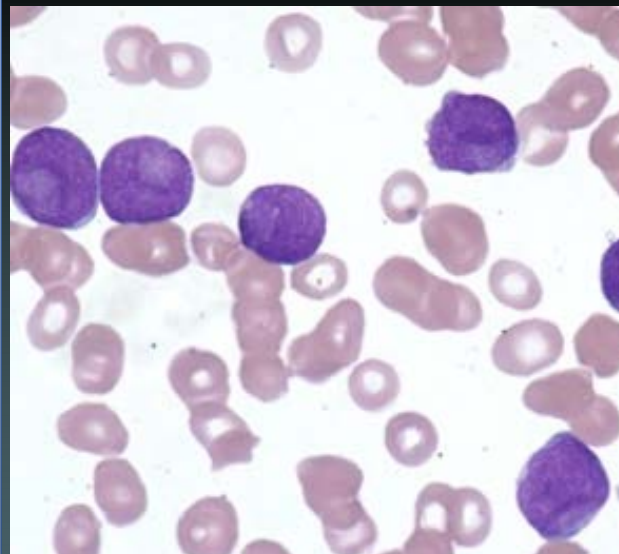
Disease Understanding

Accurate Disease Characterization

Rationally Designed Therapies

Reproducible Assessment of Fitness

Robust Platform for Efficacy Evaluation



It is not easy to see what should be done to prevent symptoms of the kind described in this paper. Faced by a patient with a duodenal ulcer who is known to be a moderate drinker and who has had a number of acute complications of his ulcer, the surgeon must choose whether or not to operate. If he does not do so he

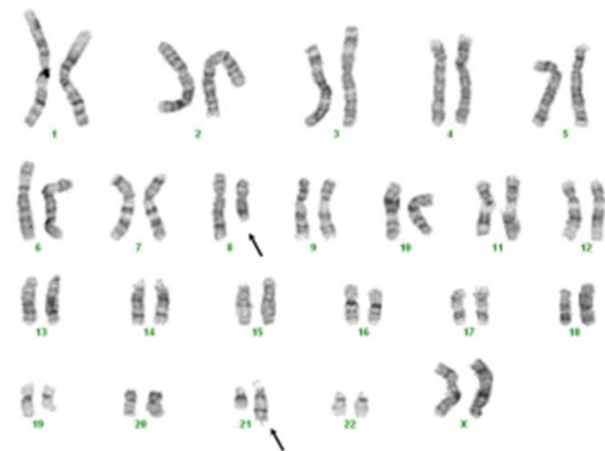
CYTOGENETIC STUDIES IN ACUTE LEUKAEMIA

BY
A. G. BAIKIE, M.B., Ch.B., M.R.C.P.Ed.
PATRICIA A. JACOBS, B.Sc.
J. A. McBRIDE, M.B., Ch.B., M.R.C.P.Ed.

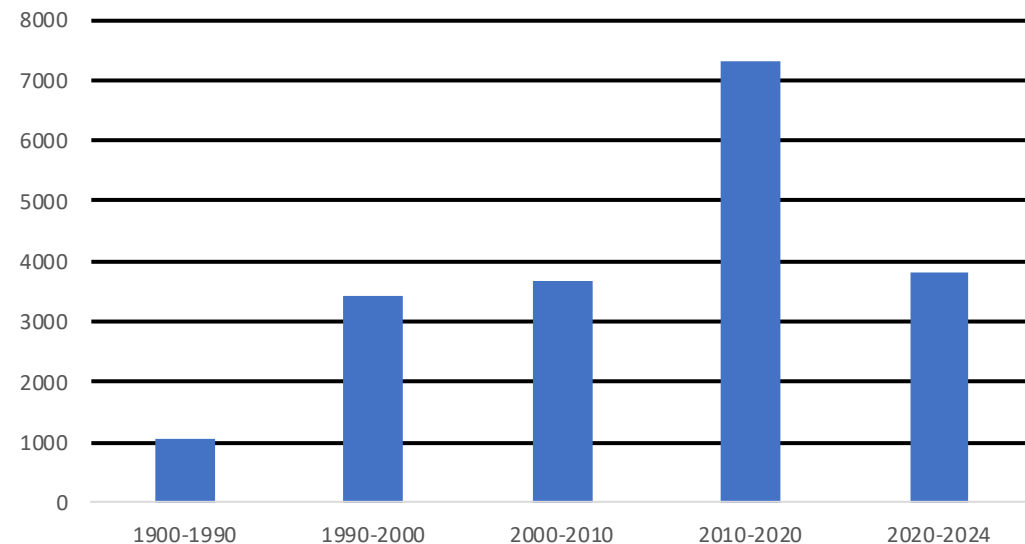
AND
ISHBEL M. TOUGH, B.Sc.

Medical Research Council Clinical Effects of Radiation Research Unit, Western General Hospital, Edinburgh

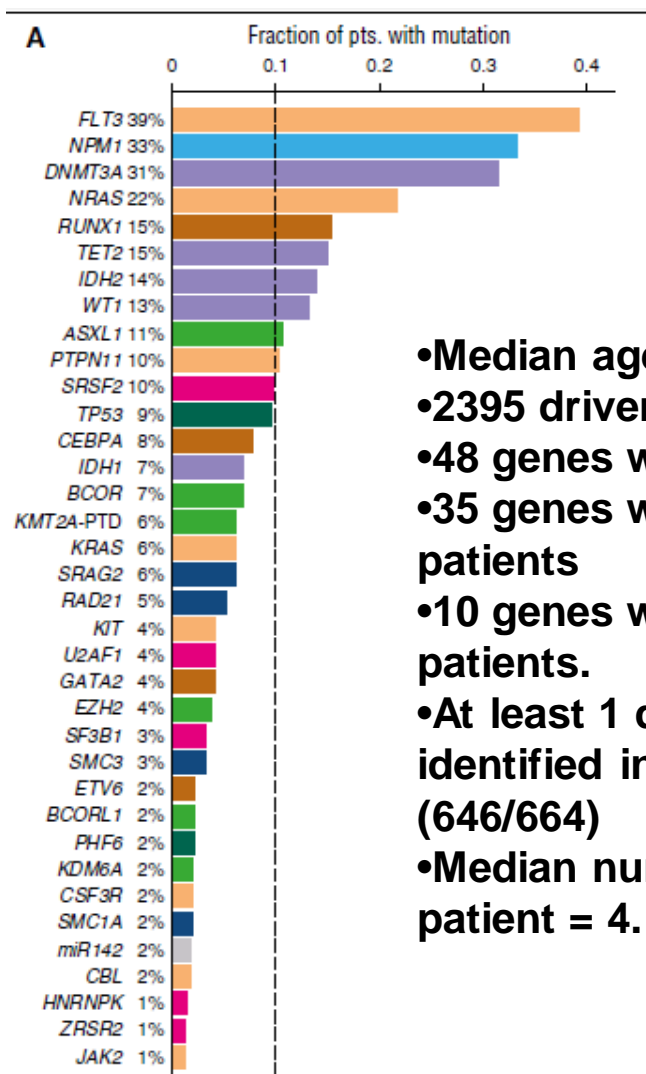
Abnormalities of the chromosomes of leukaemic cells in human acute leukaemia have now been the subject of several reports (Ford *et al.*, 1958b; Baikie *et al.*, 1959; Ford, 1960; Sandberg *et al.*, 1960). It would appear that in about half the cases reported some abnormality has been found, including aberrations of chromosome number, morphology, and sometimes both of these. Among the reported cases no two appear to have had an identical abnormality. In this respect, and as regards



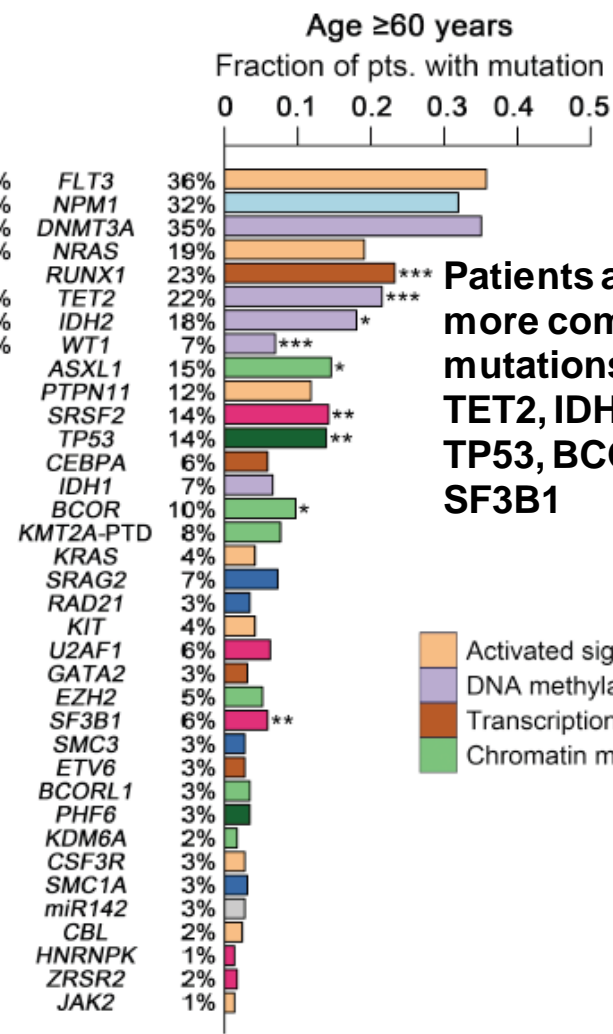
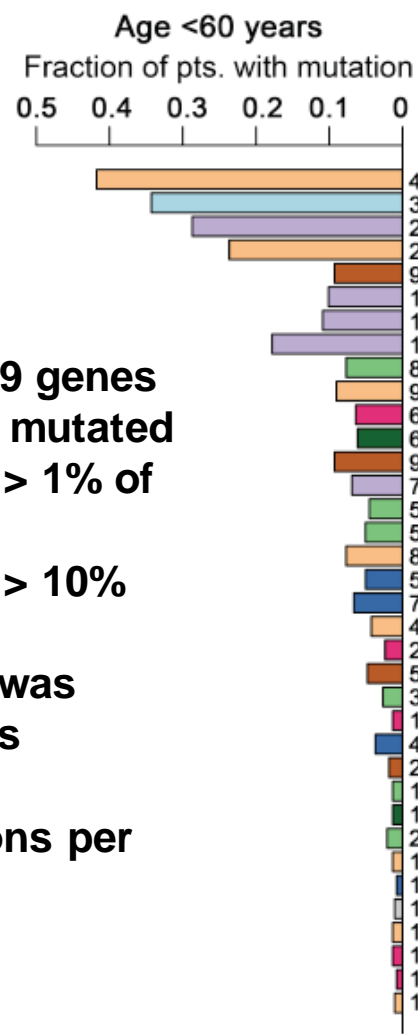
AML publications using the word "Molecular"



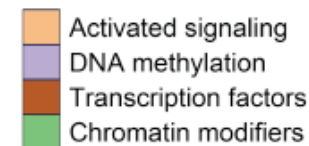
Mutational Profile in AML (664 Patients)



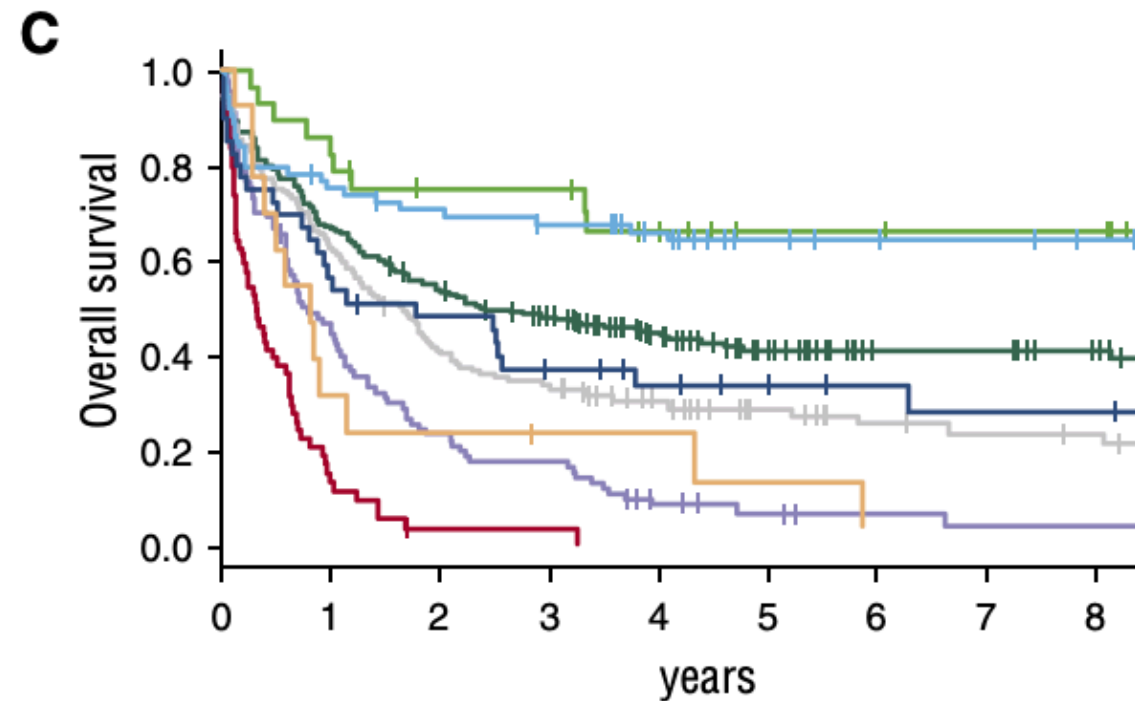
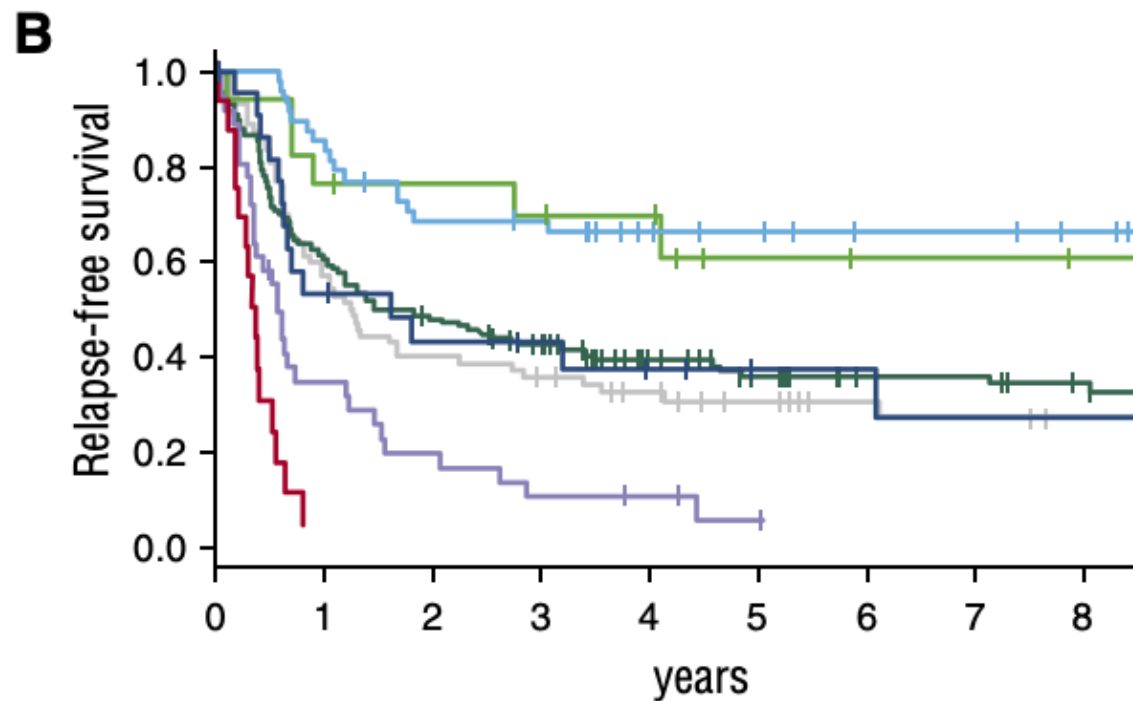
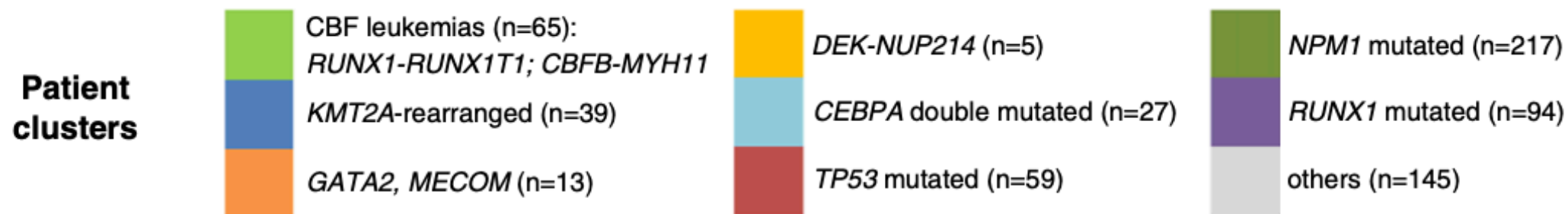
- Median age: 57 yrs
- 2395 driver mutations in 59 genes
- 48 genes were recurrently mutated
- 35 genes were mutated in > 1% of patients
- 10 genes were mutated in > 10% patients.
- At least 1 driver mutation was identified in 97% of patients (646/664)
- Median number of mutations per patient = 4.



Patients aged > 60 yr more commonly had mutations in RUNX1, TET2, IDH2, SRSF2, TP53, BCOR, and SF3B1



Outcomes by cytogenetic/molecular clusters



What is required to improve our outcomes?

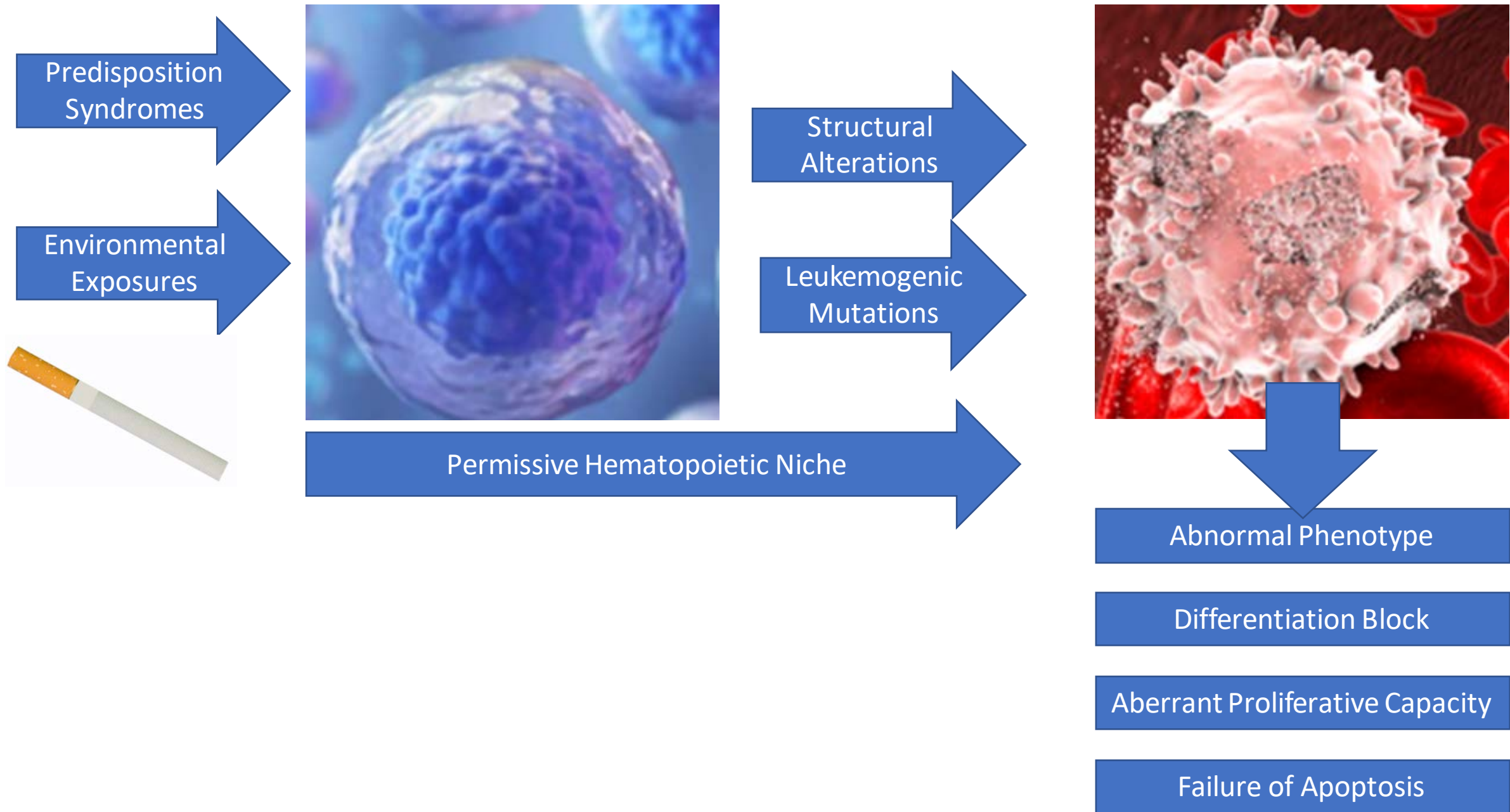
Disease Understanding

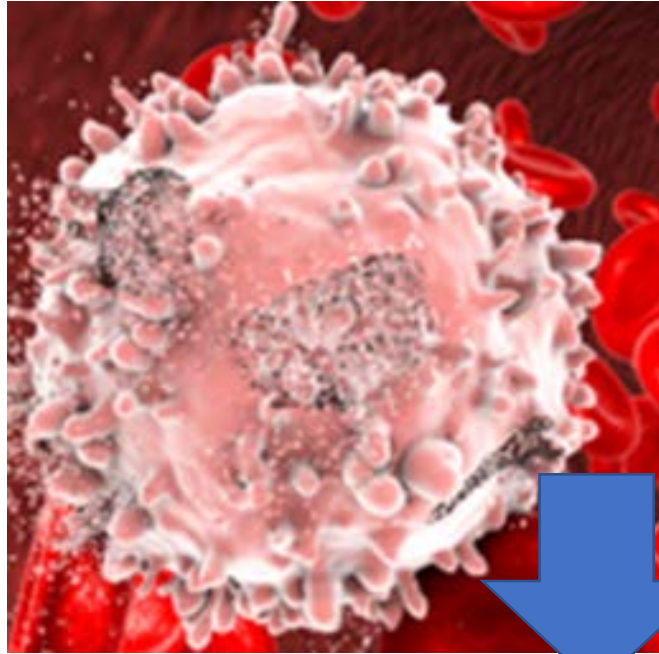
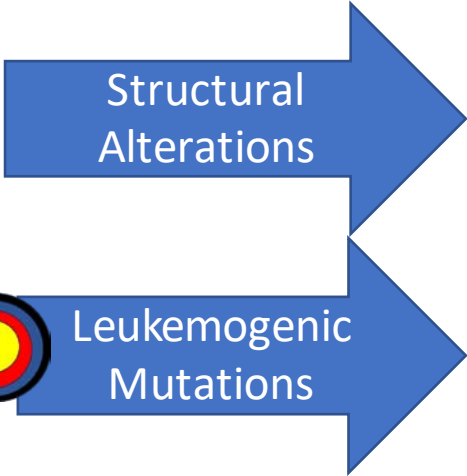
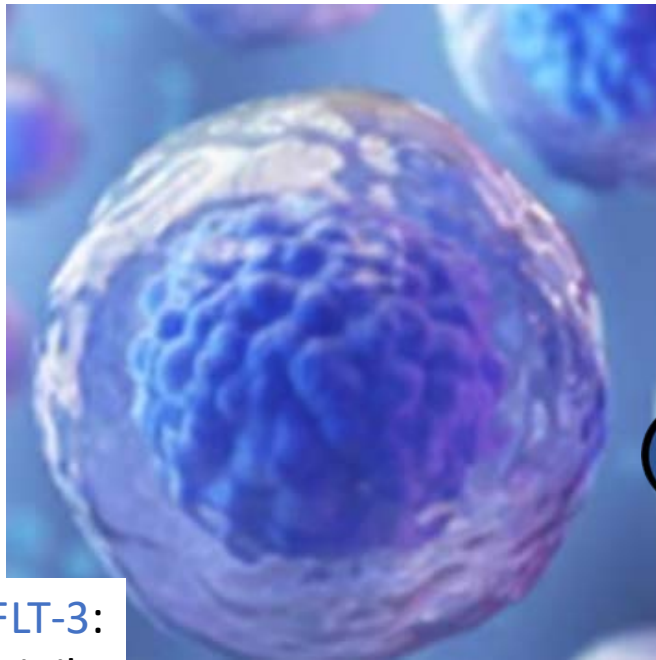
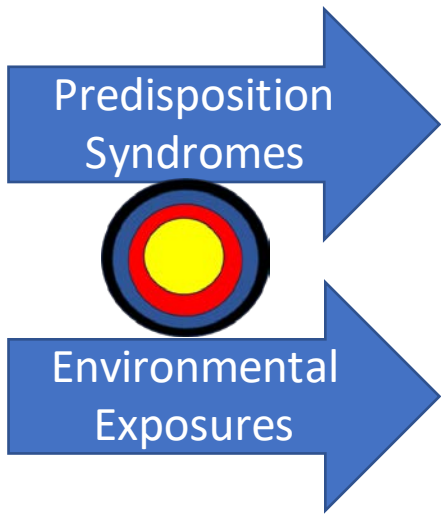
Accurate Disease Characterization

Rationally Designed Therapies

Reproducible Assessment of Fitness

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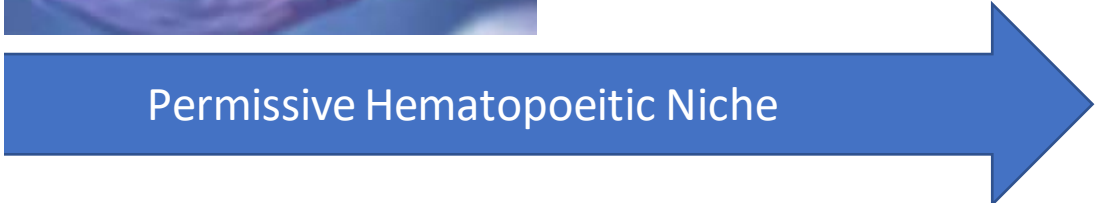


Inhibitors of FLT-3:
 Midostaurin, Gilteritinib,
 Quizartinib

Hedgehog Pathway Inhibitor:
 Glasdegib

Inhibitors of Isocitrate
 Dehydrogenase:
 Enasidenib, Ivosidenib,
 Olutasidenib

Inhibitors Menin Pathway:
 Revumenib, Ziftomenib,
 JNJ-75276617
 BN-104; DSP-5336



- Abnormal Phenotype
- Differentiation: ATO/ATRA
- Differentiation Block
- Cytotoxic Agents, i.e., 7+3 and CPX-351
- Aberrant Proliferative Capacity
- BCL2 Inhibition: Venetoclax
- Failure of Apoptosis

Challenges to Drug Development in AML

- Rare Disease
- Potential targets must be identified rapidly based on clinical presentation.
- Overall Survival has been the primary endpoint for full approval of new agents in AML.
- Lack of prospective data proving the predictive value of an MRD assay
- The definition of eligibility for curative intensive therapy has been elusive
- Absence of coordinated national approach
- Most patients, at most centers, are not offered investigational options or are ineligible for the trials that exist

myeloMATCH

**Myeloid Malignancies Molecular
Analysis for Therapy Choice**

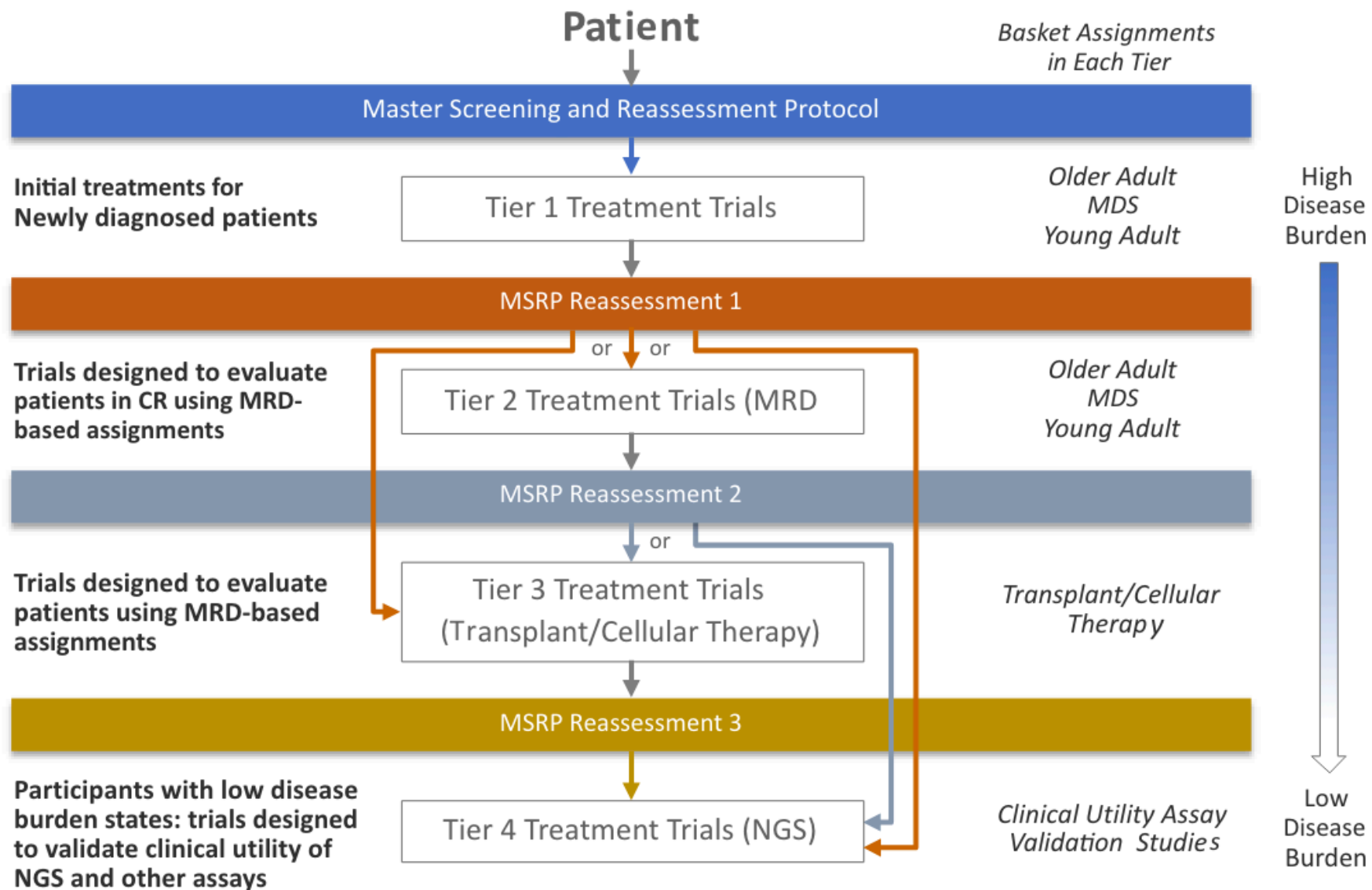
**NCI National Clinical Trials
Network**

**Leadership: Richard F. Little, M.D.
CTEP, NCI**

myeloMATCH Aims

- Create a portfolio of rationally designed treatment substudies
 - Patients enroll sequentially over their entire treatment journey
 - Scientific data is generated at every stage
- Create an efficient operational model which attracts
 - Industry partners
 - NCTN sites
- Develop the careers of young investigators
 - Promoting leadership throughout the clinical trial portfolio and laboratory program.
- Efficiently test and promulgate innovative standards of care for AML
 - Outcomes to provide therapeutic clarity

MyeloMATCH MSRP Schema

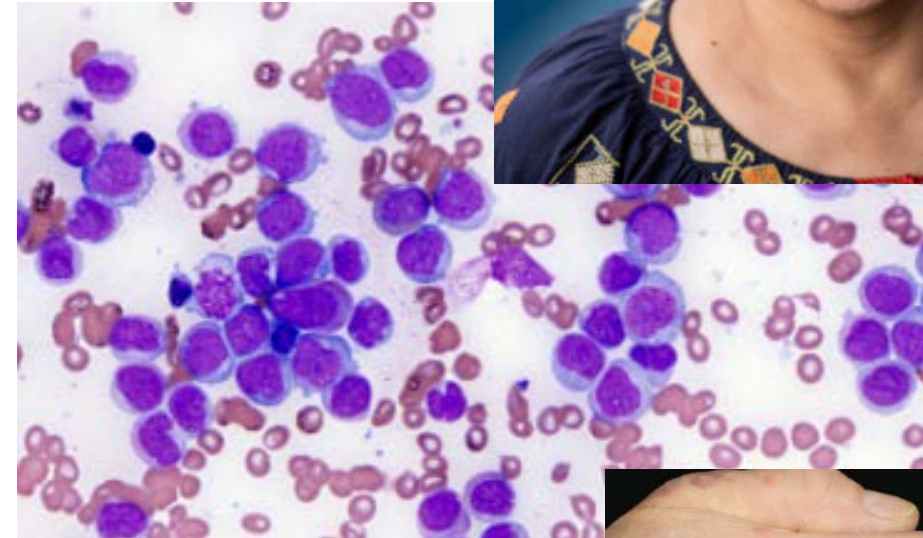


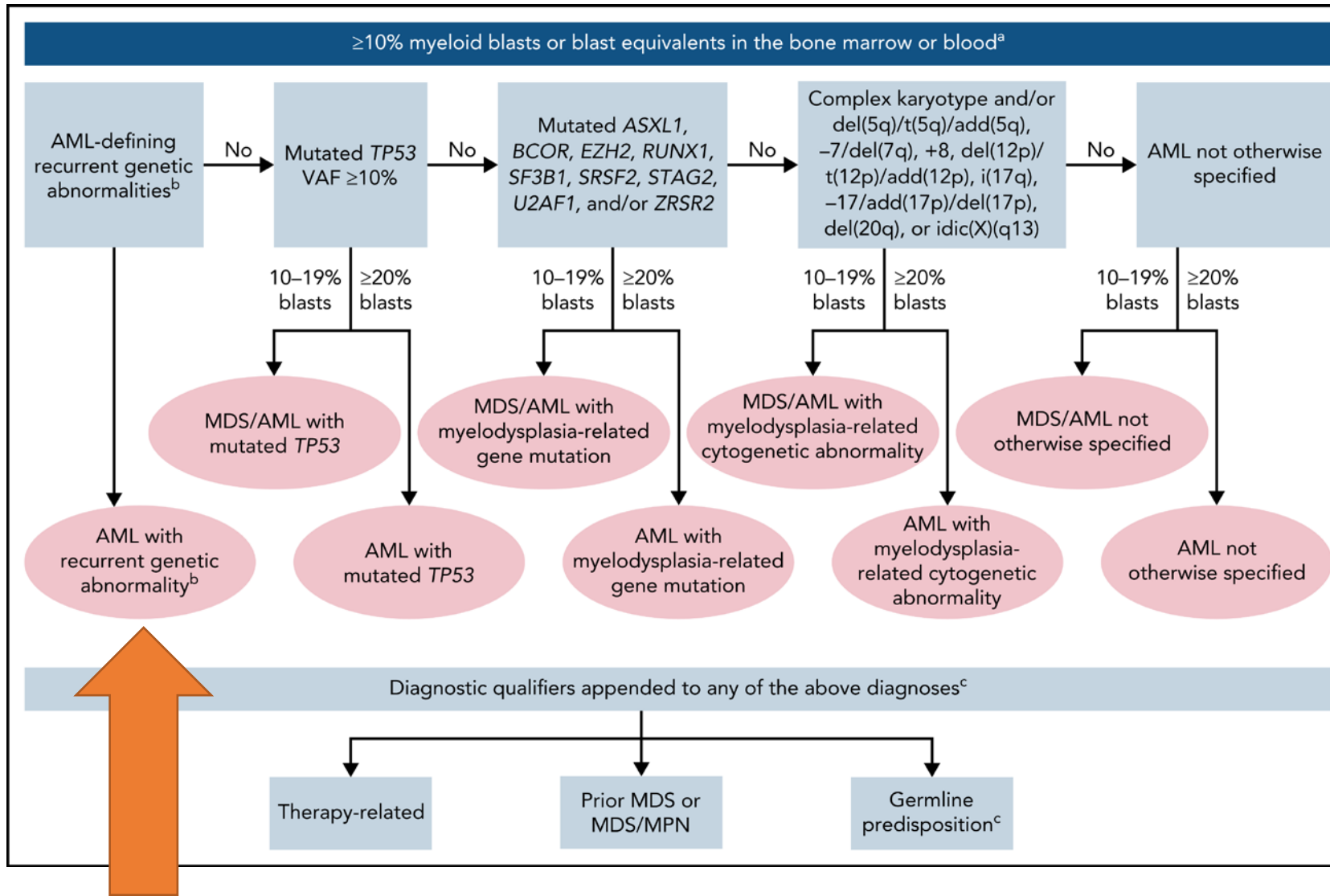
Topics for the Day

- Definitions
 - Precision Medicine
 - myeloMATCH
- Current State of Therapeutic Options
 - Younger, Fit
 - Older or Frail
- Hurdles and Potential for this Strategy
- Discussion

- 58-year-old woman
- History of well-controlled hypothyroidism
- Presents with night sweats, mild cough, new rash

WBC	24 X 10 ³ /uL
HGN	9.3 g/dL
PLTS	67 X 10 ³ /uL
Circulating Blasts	70%
Chromosomes	t(9;11)(p21;q23) KMT2A/MLLT3
Molecular Mutation	KRAS c.436G>A, p.Ala146Thr VAF: 10.2% KRAS c.38G>A, p.Gly13Asp VAF: 7.0%





Döhner et al., Diagnosis and management of AML in adults: Blood, 2022

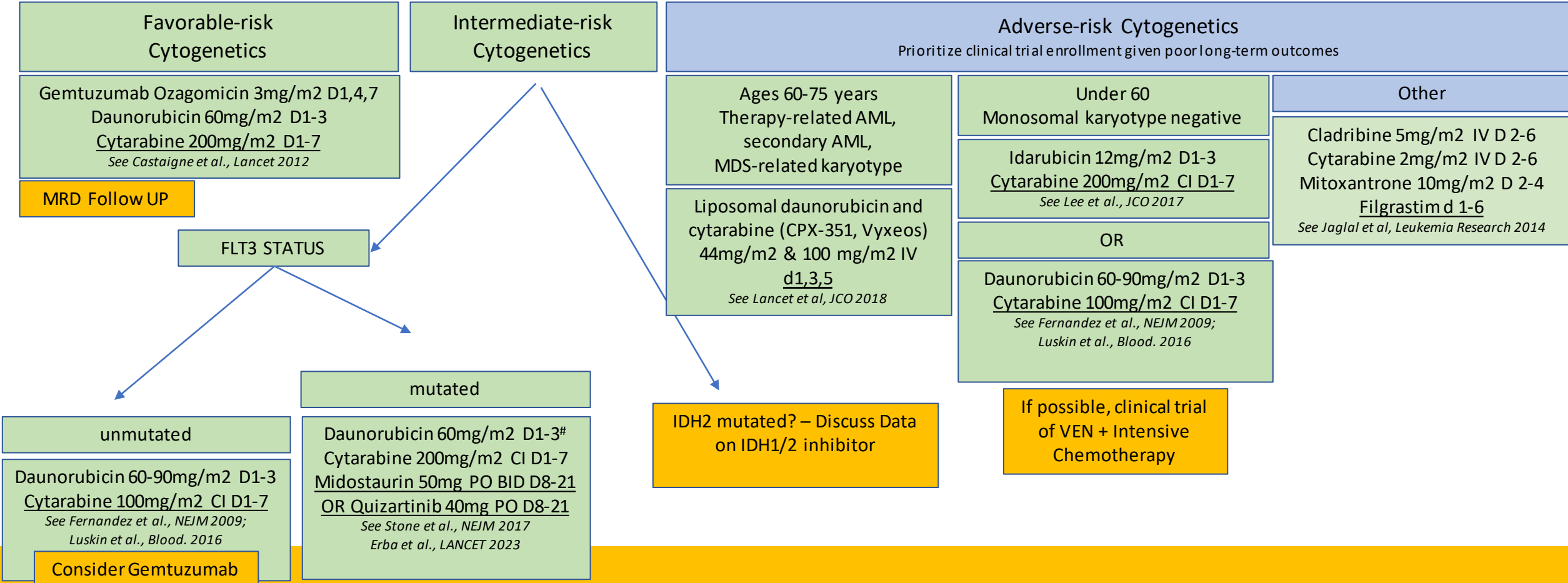
2022 ELN risk classification by genetics at initial diagnosis*

Risk category†	Genetic abnormality
Favorable	<ul style="list-style-type: none"> t(8;21)(q22;q22.1)/<i>RUNX1::RUNX1T1</i>†,‡ inv(16)(p13.1;q22) or t(16;16)(p13.1;q22)/ <i>CBFB::MYH11</i>†,‡ Mutated <i>NPM1</i>†,§ without <i>FLT3</i>-ITD bZIP in-frame mutated <i>CEBPA</i>
Intermediate	<ul style="list-style-type: none"> Mutated <i>NPM1</i>†,§ with <i>FLT3</i>-ITD Wild-type <i>NPM1</i> with <i>FLT3</i>-ITD (without adverse-risk genetic lesions) t(9;11)(p21.3;q23.3)/<i>MLLT3::KMT2A</i>†,¶ ← Cytogenetic and/or molecular abnormalities not classified as favorable or adverse
Adverse	<ul style="list-style-type: none"> t(6;9)(p23.3;q34.1)/<i>DEK::NUP214</i> t(v;11q23.3)/<i>KMT2A</i>-rearranged# t(9;22)(q34.1;q11.2)/<i>BCR::ABL1</i> t(8;16)(p11.2;p13.3)/<i>KAT6A::CREBBP</i> inv(3)(q21.3q26.2) or t(3;3)(q21.3;q26.2)/ <i>GATA2</i>, <i>MECOM(EVI1)</i> t(3q26.2;v)/<i>MECOM(EVI1)</i>-rearranged -5 or del(5q); -7; -17/abn(17p) Complex karyotype,** monosomal karyotype†† Mutated <i>ASXL1</i>, <i>BCOR</i>, <i>EZH2</i>, <i>RUNX1</i>, <i>SF3B1</i>, <i>SRSF2</i>, <i>STAG2</i>, <i>U2AF1</i>, and/or <i>ZRSR2</i>‡‡ Mutated <i>TP53</i>§§

MCW Acute Myeloid Leukemia:2024

Fit, Induction-eligible candidate

Clinical trial available? Appropriate? If not, then



Intermediate/Adverse-risk Cytogenetics and NO transplant: Discuss MRD follow up and /or Maintenance therapy X 2 years

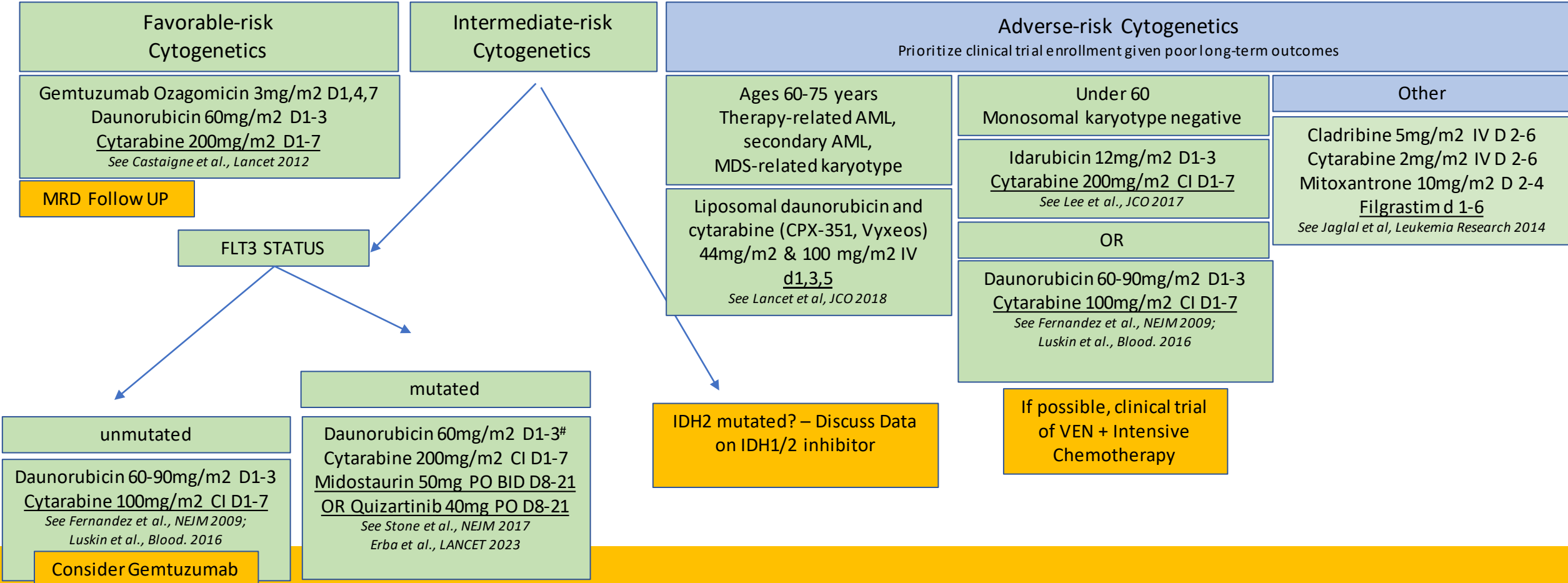
Study	Arms				Comments
E1900 Fernandez et al., NEJM 2009; Luskin et al., Blood. 2016 Mar 24;127(12):1551-8	Daunorubicin 45mg/m2 d1-3 + 100mg/m2 Cytarabine CI d1-7		Daunorubicin 90mg/m2 d1-3 + 100mg/m2 Cytarabine CI d1-7		Ages 17-60 yrs Data after Median F/U: 80.1 mo for survivors 90mg/m2 of daunorubicin vs 45mg/m2 benefits AML patients with favorable and intermediate cytogenetics and with FLT3-ITD, NPM1, and DNMT3A mutations.
	n:330		n:327		
	CR:	4-y OS:	CR:	4-y OS:	
All Patients	59%	31%	71%	39%	
Karyotype					
Favorable Risk	84%	46%	80%	64%	
Intermediate Risk	56%	35%	77%	45%	
Adverse Risk	44%	14%	57%	19%	

Study	Arms		Arms		Comments
Lee et al. JCO 2017	Idarubicin (12mg/m2 d1,2,3) + cytarabine (200 mg/m2) d1-7 (Control arm)		Daunorubicin 90mg/m2 (d1,2,3) + cytarabine (200 mg/m2) d1-7		Ages: 15-65 years Powered for non-inferiority No difference between Idarubicin and Daunorubicin arms with regard to CR, OS. High-dose daunorubicin was more effective than idarubicin for patients with FLT-ITD mutation
	N=149		N=150		
	CR/CRi	4-y OS	CR/CRi	4-yr OS	
All Patients	80.5%	51.1%	74.7%	54.7%	
Cytogenetic Risk Group					
Good	88.9%	85.2	100%	90.7	
Intermediate	84.5%	53.1	75.3%	49	
Poor, monosomal karyotype neg	58.3%	25.0	61.3%	40.8	
Poor, monosomal karyotype pos	44.4%	0	30.0%	24.0	

MCW Acute Myeloid Leukemia:2024

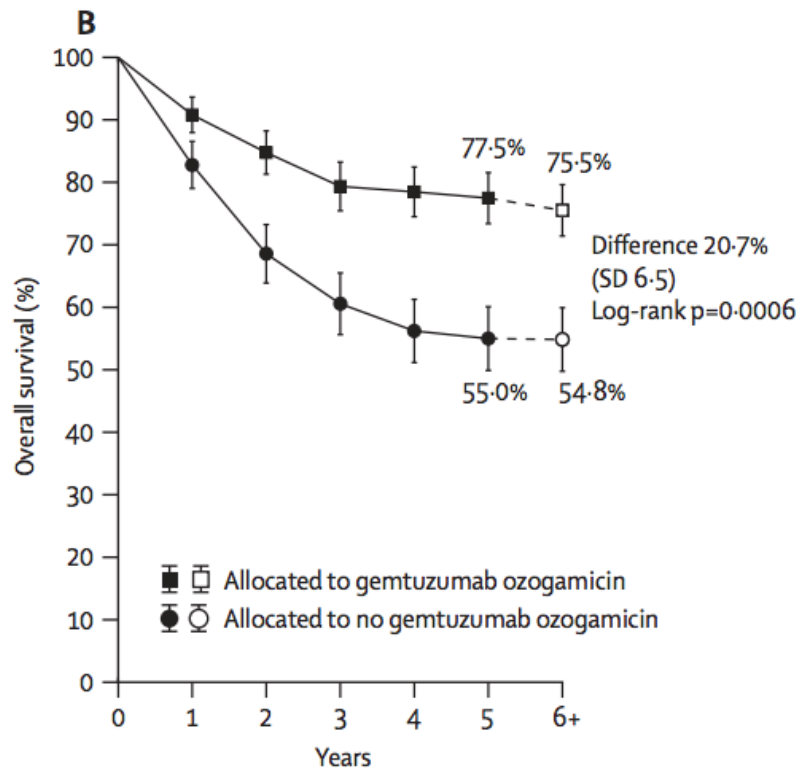
Fit, Induction-eligible candidate

Clinical trial available? Appropriate? If not, then

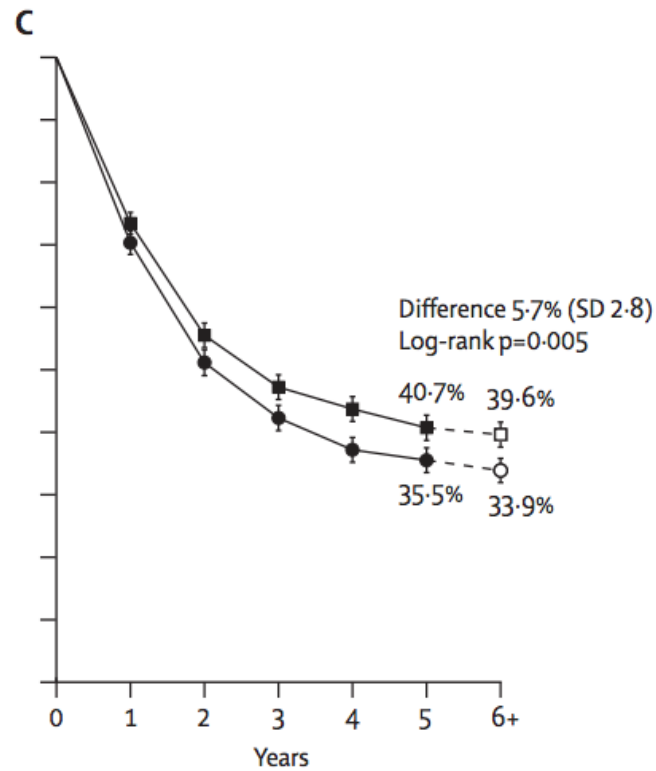


Intermediate/Adverse-risk Cytogenetics and NO transplant: Discuss MRD follow up and /or Maintenance therapy X 2 years

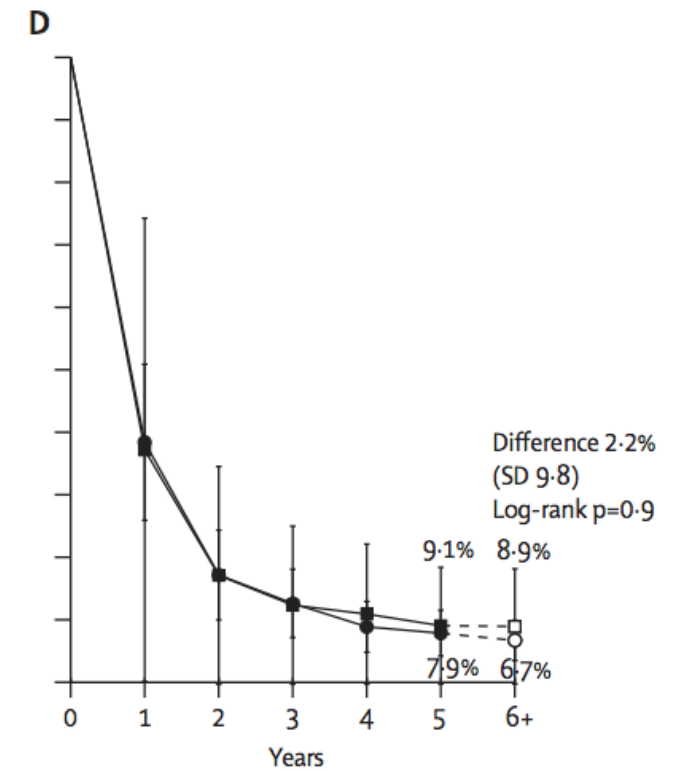
Meta Analysis of Gemtuzumab combination therapy



Annual event rates	Years 1-5	Years 6+
Gemtuzumab ozogamicin	5.8% SD 1.1	2.3% SD 1.3
No gemtuzumab ozogamicin	14.1% SD 1.9	0.0% SD 0.0



Annual event rates	Years 1-5	Years 6+
Gemtuzumab ozogamicin	22.4% SD 1.0	2.7% SD 0.9
No gemtuzumab ozogamicin	26.2% SD 1.1	4.9% SD 1.3

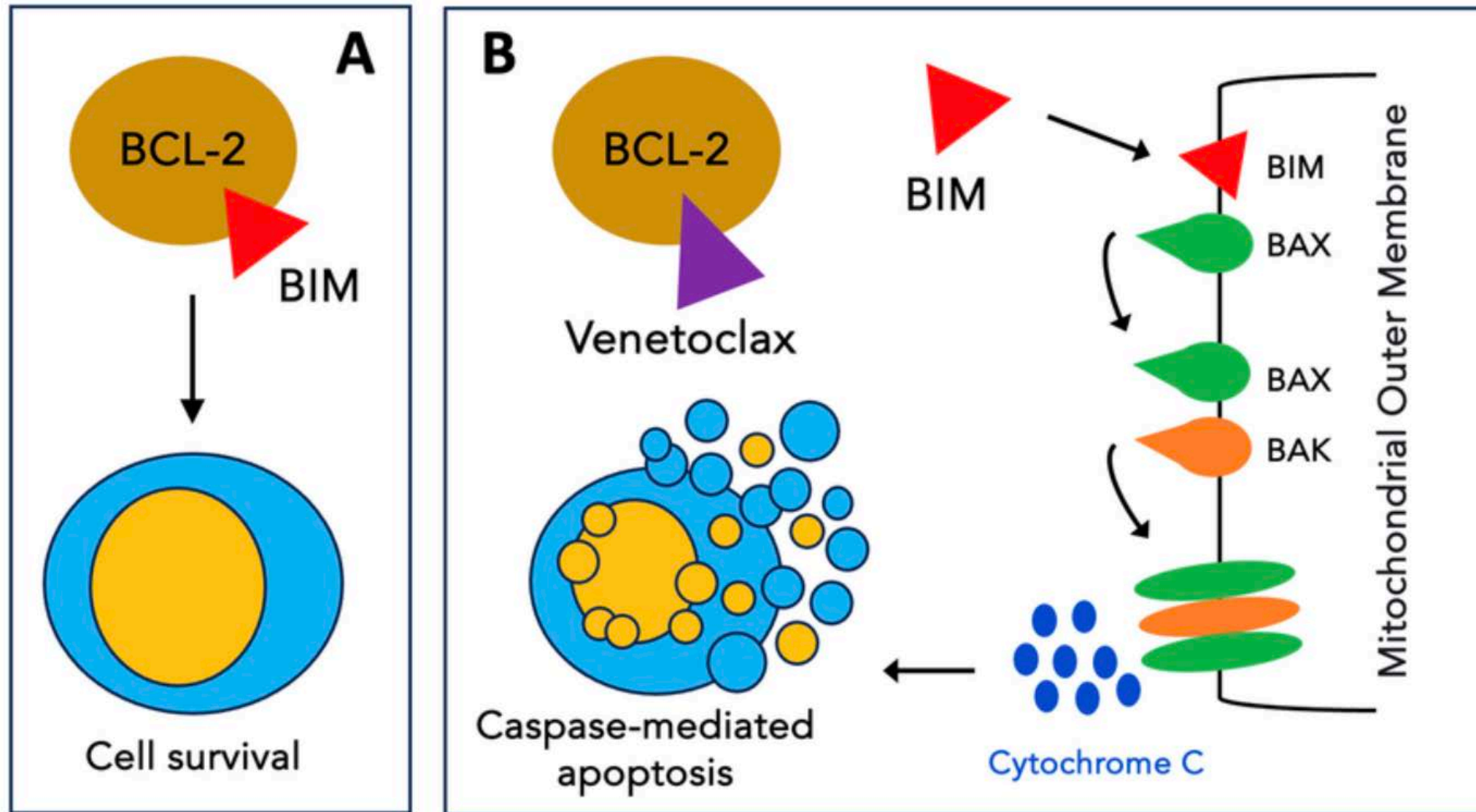


Annual event rates	Years 1-5	Years 6+
Gemtuzumab ozogamicin	73.8% SD 4.6	2.4% SD 2.4
No gemtuzumab ozogamicin	76.7% SD 4.8	21.1% SD 10.5

Unanswered Questions

- What about adding venetoclax to cytotoxic induction?
- Could a menin inhibitor be utilized here alongside the cytotoxic backbone?
- How best follow her MRD? What do we do with persistent MRD?
- Is the cytotoxic backbone even necessary?





Why BCL2 inhibition in AML?

Preclinical data

- Upregulated BCL-2 → evasion of apoptosis
- Increased BCL-2 expression → worse prognosis

CALBG Study 2000s

- BCL-2 antisense oligonucleotide + intensive chemotherapy

Venetoclax

- BH3 mimetic, BCL-2 selective inhibitor
- Monotherapy in R/R AML: Overall response rate of 19%

Regimen	Trial	Citation
Daunorubicin Cytarabine + Venetoclax	Phase II; Conducted in China N=33; CCR: 91% Ages 18-60 years No TRM	Wang et al., Lancet Haematol. 2022 Jun;9(6):e415-e424. doi: 10.1016/S2352-3026(22)00106-5. Epub 2022 May 2. PMID: 35512726
Cladribine, Idarubicin, Cytarabine + Venetoclax	Phase II; MD Anderson N=50; Up to age 65 years CCR: 94% TRM 1/50	Kadia et al., Lancet Haematol. 2021 Aug;8(8):e552-e561. doi: 10.1016/S2352-3026(21)00192-7.
Fludarabine Cytarabine Idarubicin + Venetoclax	Phase I; MD Anderson ND AML, N=29 Up to age 65; CCR: 96% 60-day mortality: 4%	DiNardo et al., J Clin Oncol. 2021 Sep 1;39(25):2768-2778. doi: 10.1200/JCO.20.03736. Epub 2021 May 27.
Daunorubicin Cytarabine + Venetoclax	Phase Ib Conducted in Australia ND AML, Ages 63-80 CR/CRI 97% in denovo AML	Chua et al., J Clin Oncol . 2020 Oct 20;38(30):3506-3517. doi: 10.1200/JCO.20.00572. Epub 2020 Jul 20.

Multiagent Chemotherapy + Venetoclax

- MD Anderson team investigated Fludarabine, Cytarabine, Idarubicin backbone with Venetoclax
- Newly Diagnosed AML
- Phase II, single-arm study
- Endpoint: ORR
- N=45 patients

TABLE 2 Response outcomes

Parameter ^a	All patients (N = 45)	ELN favorable ^c (N = 8)	ELN intermediate (N = 18)	ELN adverse (N = 19)
Overall response ^b	98 (44 [90–100])	100 (8 [74–100])	94 (17 [77–99])	100 (19 [90–100])
Composite CR	89 (40 [75–96])	88 (7 [32–99])	89 (16 [61–98])	89 (17 [63–98])
Complete response	73 (33)	63 (5)	72 (13)	79 (15)
CRh	11 (5)	25 (2)	11 (2)	5 (1)
CRi	4 (2)	-	6 (1)	5 (1)
MLFS	9 (4)	13 (1)	6 (1)	11 (2)
No response	1 (2)	-	1 (6)	-
MRD-negative (MFC)	93 (37/40 [78–98])	100 (7/7 [NA])	88 (14/16 [57–97])	94 (16/17 [63–99])
Duration of response	NR (-)	NR (-)	NR (17-NR)	NR (11-NR)

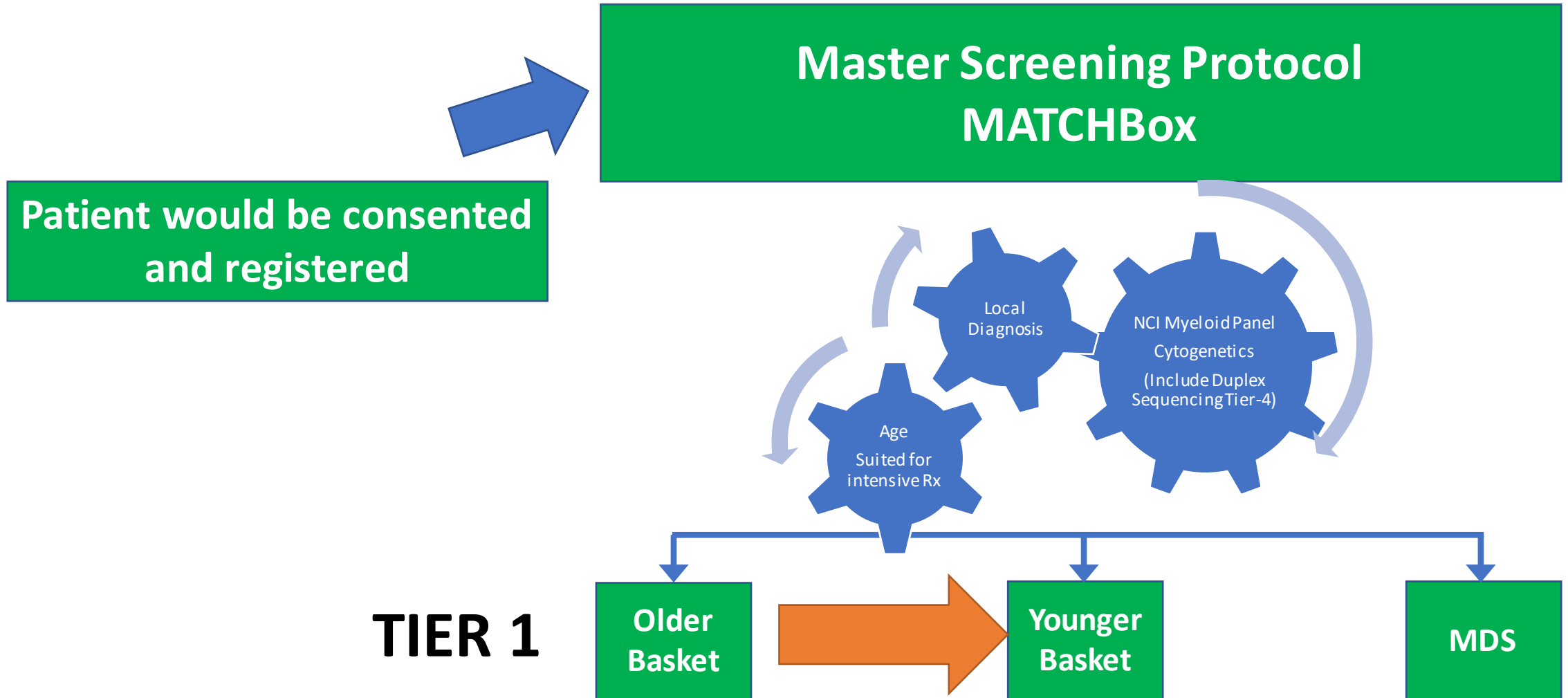
^aAll variables presented as median (range) or % (N).

^b95% Credible intervals per Protocol-defined primary efficacy outcome (95% credible interval estimation assumed ORR follows a prior distribution of beta of (1.4, 0.6). 95% Exact confidence intervals presented for other response outcomes. Response variables reported as % (N [95%CI]) when credible interval or confidence interval included.

^cELN favorable risk patients were composed of *NPM1* or biallelic *CEBPA* mutations, no patients with favorable risk cytogenetics were enrolled on study.

What would this patient's
treatment look like on
myeloMATCH?

myeloMATCH option



MyeloMATCH: Menu of Assays (<72 hours TAT)

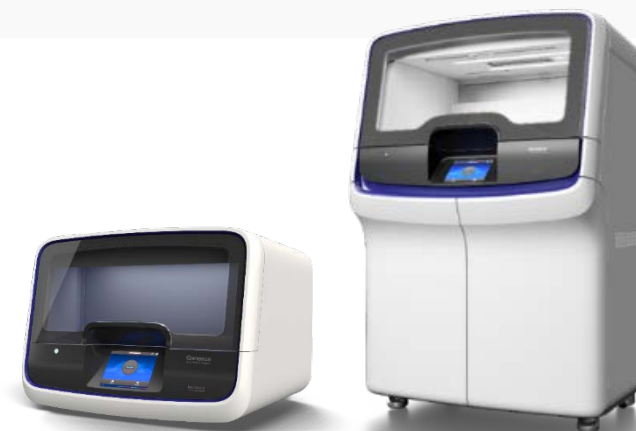
- At diagnosis (risk classification, therapy targets)
 - Rapid cytogenetics/FISH/CGAT
 - Flow cytometry
 - Rapid NGS (Ion Torrent™)
- MRD assessment
 - Multicolor flow cytometry (outcome, MRD eraser assignment)
 - Duplex sequencing-potentially better than flow?
- Clonal heterogeneity (understanding clonal selection and response)
 - Single cell DNA, RNA, protein

Integral studies
Integrated studies

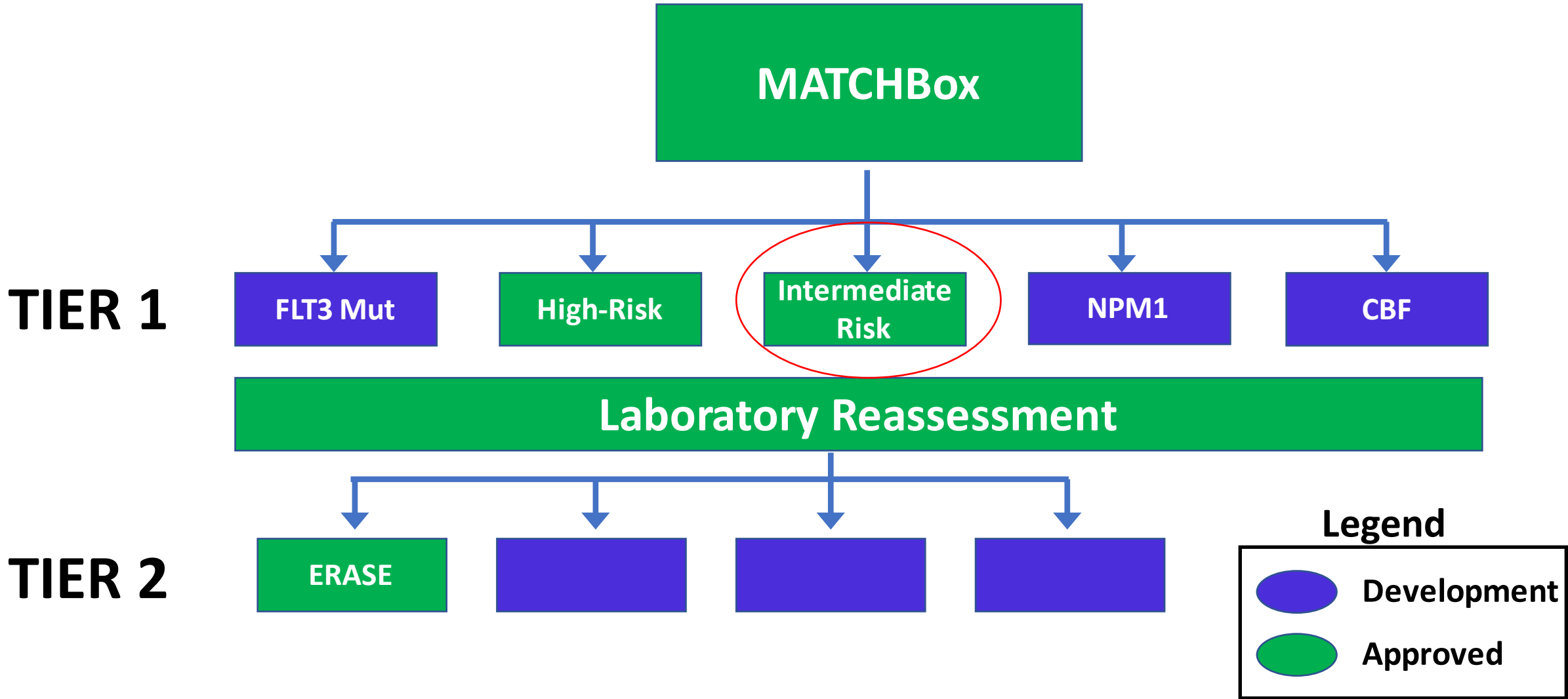
MyeloMATCH: NCI Myeloid Gene Assay version 2

DNA hotspots				
<i>ABL1</i>	<i>ANKRD26</i>	<i>BRAF</i>	<i>CBL</i>	<i>CSF3R</i>
<i>DDX41</i>	<i>DNMT3A</i>	<i>FLT3</i>	<i>GATA2</i>	<i>HRAS</i>
<i>IDH1</i>	<i>IDH2</i>	<i>JAK2</i>	<i>KIT</i>	<i>KRAS</i>
<i>MPL</i>	<i>MYD88</i>	<i>NPM1</i>	<i>NRAS</i>	<i>PPM1D</i>
<i>PTPN11</i>	<i>SETBP1</i>	<i>SF3B1</i>	<i>SMC1A</i>	<i>SMC3</i>
<i>SRSF2</i>	<i>U2AF1</i>	<i>WT1</i>		
DNA Full Gene				
<i>ASXL1</i>	<i>BCOR</i>	<i>CALR</i>	<i>CEBPA</i>	<i>ETV6</i>
<i>EZH2</i>	<i>IKZF1</i>	<i>NF1</i>	<i>PHF6</i>	<i>PRPF8</i>
<i>RB1</i>	<i>RUNX1</i>	<i>SH2B3</i>	<i>STAG2</i>	<i>TET2</i>
<i>TP53</i>	<i>ZRSR2</i>			
RNA Fusion Driver Genes				
<i>ABL1</i>	<i>ALK</i>	<i>BCL2</i>	<i>BRAF</i>	<i>CCND1</i>
<i>CREBBP</i>	<i>EGFR</i>	<i>ETV6</i>	<i>FGFR1</i>	<i>FGFR2</i>
<i>FUS</i>	<i>HMGA2</i>	<i>JAK2</i>	<i>KMT2A</i>	<i>MECOM</i>
			(<i>MLL</i>) +PTDs	
<i>MET</i>	<i>MLLT10</i>	<i>MLLT3</i>	<i>MYBL1</i>	<i>MYH11</i>
<i>NTRK3</i>	<i>NUP214</i>	<i>NUP98</i>	<i>PDGFRA</i>	<i>PDGFRB</i>
<i>RARA</i>	<i>RBM15</i>	<i>RUNX1</i>	<i>TCF3</i>	<i>TFE3</i>
<i>BAALC</i>	<i>MECOM</i>	<i>MYC</i>	<i>SMC1A</i>	<i>WT1</i>

- Platform: Ion Torrent™ Genexus™ System
 - 45 DNA genes and 35 fusion driver genes
 - Includes 28/30 (93.3%) genes mutated with $\geq 3\%$ frequency in AML.
 - Includes 36/50 (72%) genes mutated with $> 1\%$ frequency in AML.
 - Includes 779 unique fusions reported in AML
 - Can detect FLT3-ITD up to 120bp
- Can detect all genetic alterations needed for
 - WHO classification of AML, except inv 3
 - NCCN/ELN risk stratification, except inv 3

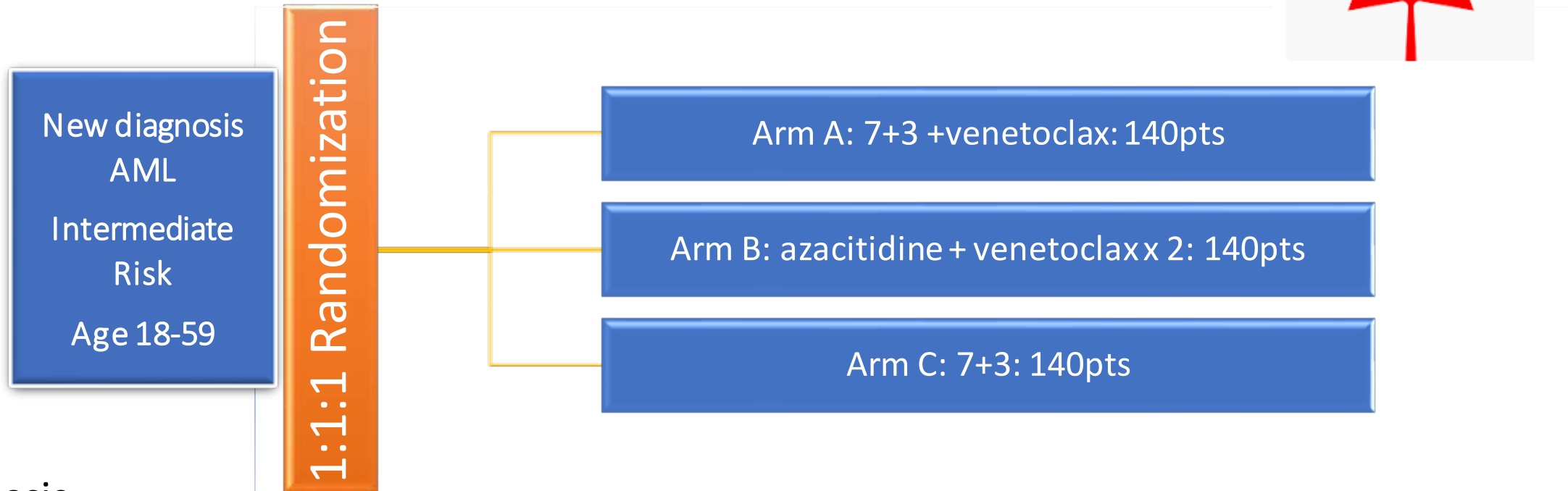


First Generation Studies in the Younger Basket



MM1YA-CTG01

Study co-chairs: Sarit Assouline and Lynn Savoie



Hypothesis

- The addition of venetoclax to chemotherapy will increase MRD negative complete response (CR) in newly diagnosed AML patients fit for induction chemotherapy with intermediate risk disease

Primary objective

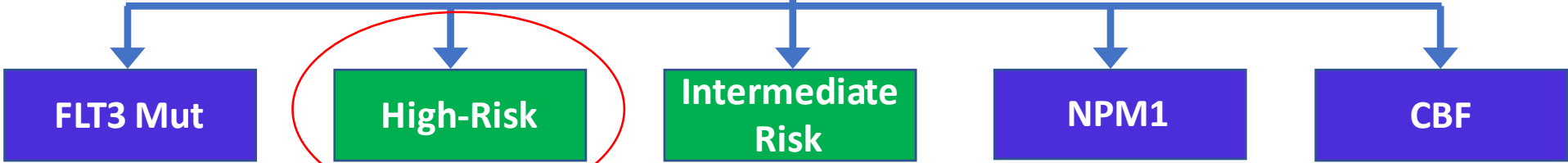
- MRD negative (by FLOW Cytometry) complete response rate (CR) after standard induction, standard induction with venetoclax, or two cycles of venetoclax + azacitidine

First Generation Studies in the Younger Basket

Chromosomes	48,XX,+8,+8[20] Tetrasomy 8
Molecular Mutation	SF3B1 c.1998G>C, p.Lys666Asn (NM_012433.4) VAF: 40.7%

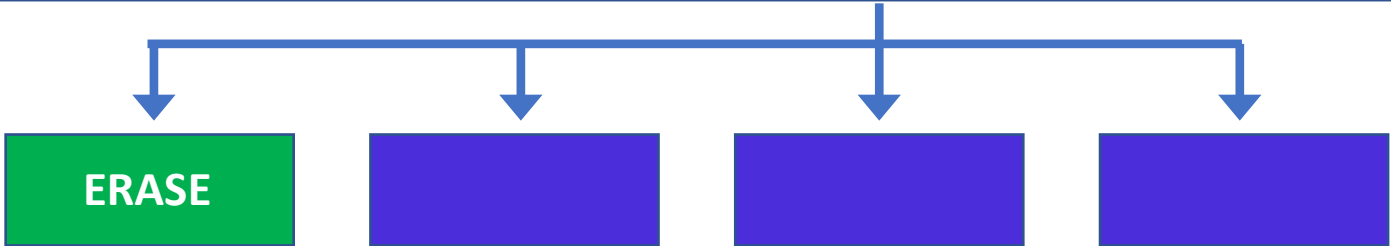
MATCHBox

TIER 1



Laboratory Reassessment

TIER 2

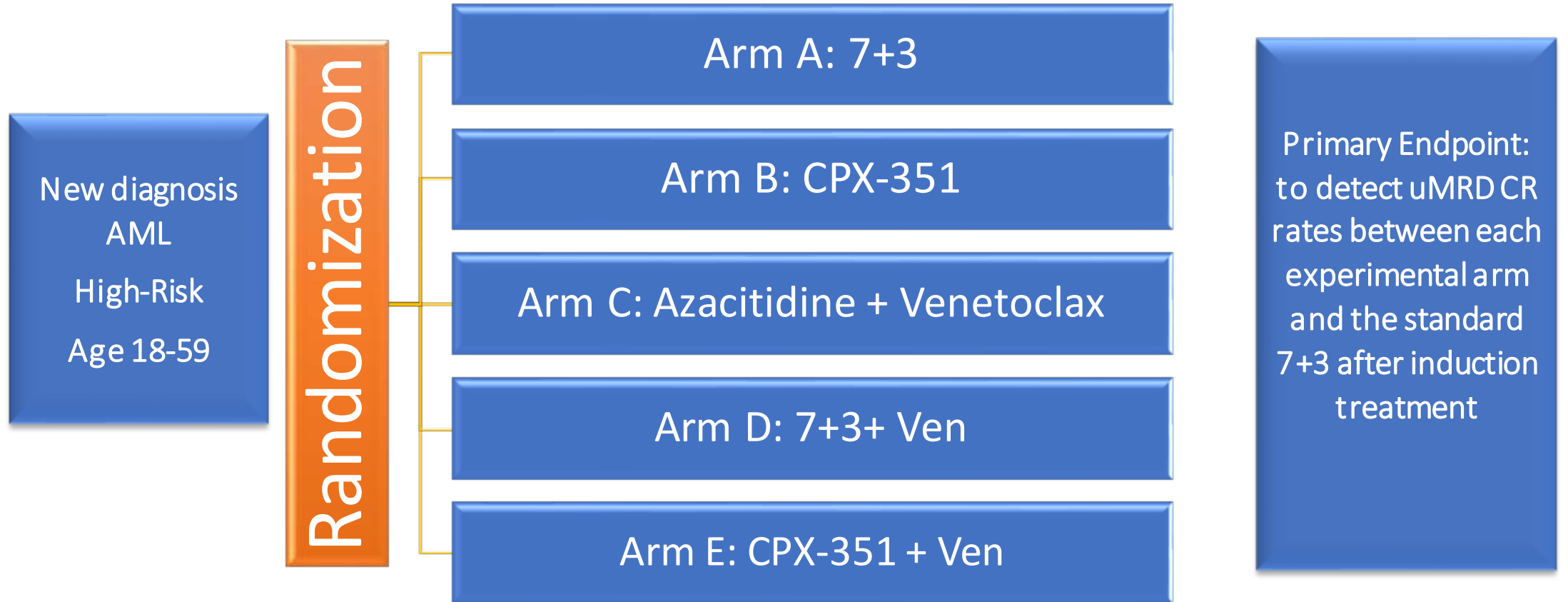


Legend

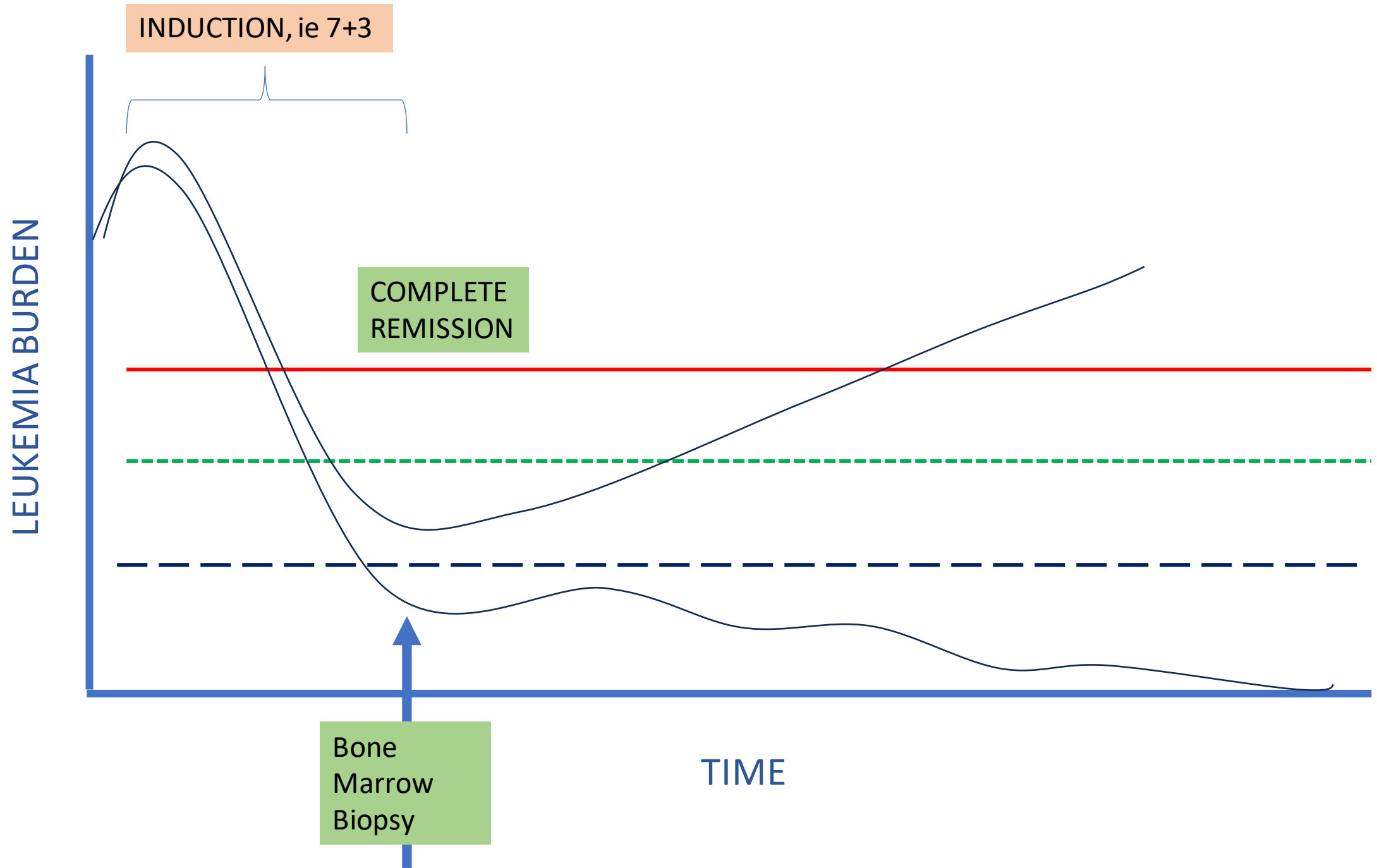
- Development (Blue oval)
- Approved (Green oval)

MM1YA-S01

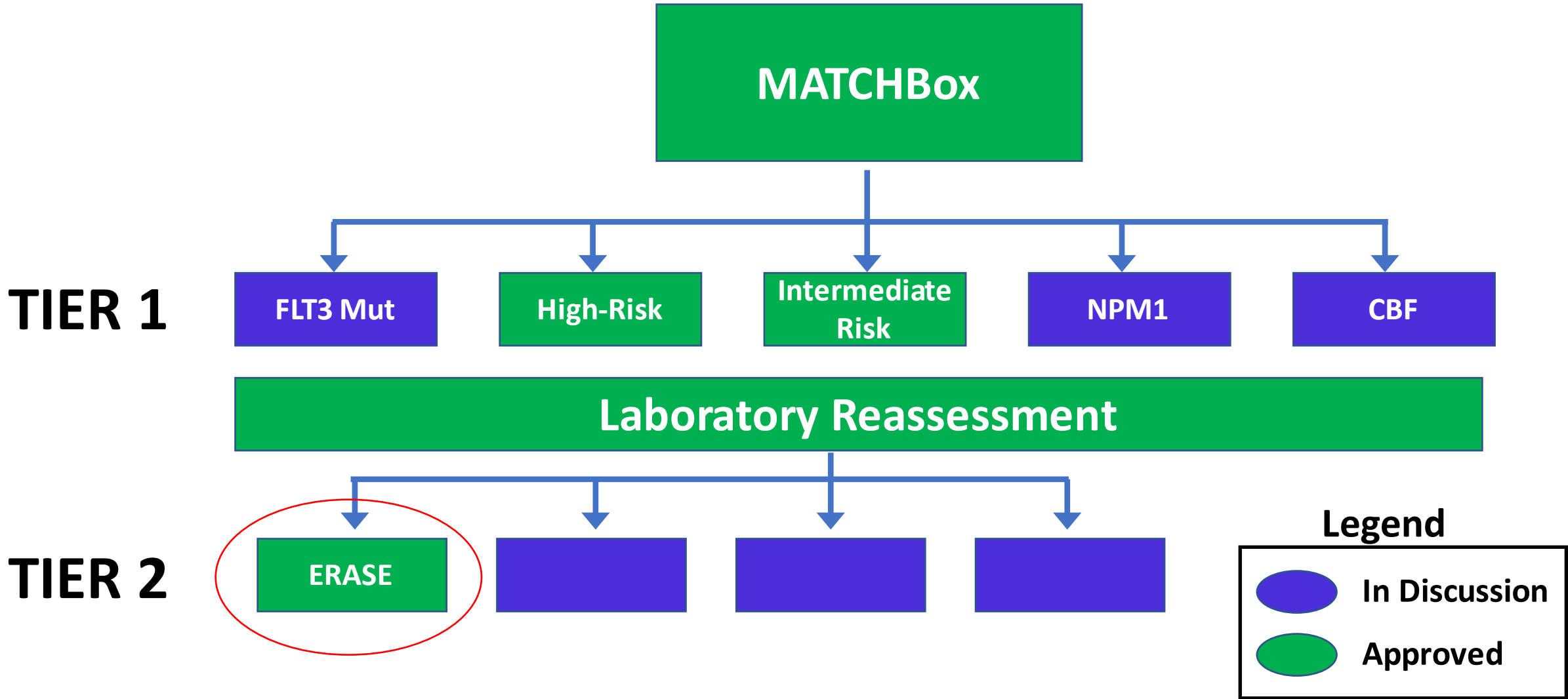
Study co-chairs: Paul Shami and Tara Lin



Sample size: 268 (60 +7 per arm)

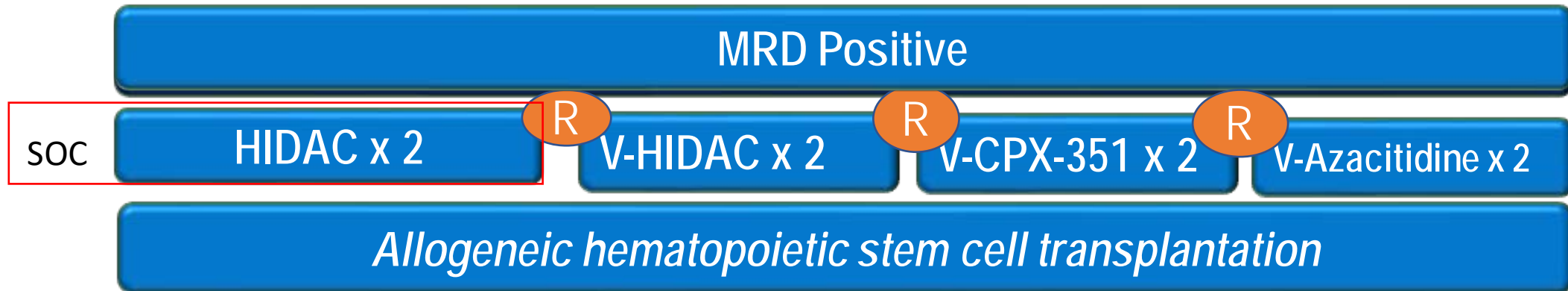


First Generation Studies in the Younger Basket



MM2YA-EA01

Study co-chairs: Ehab Atallah and Rita Assa



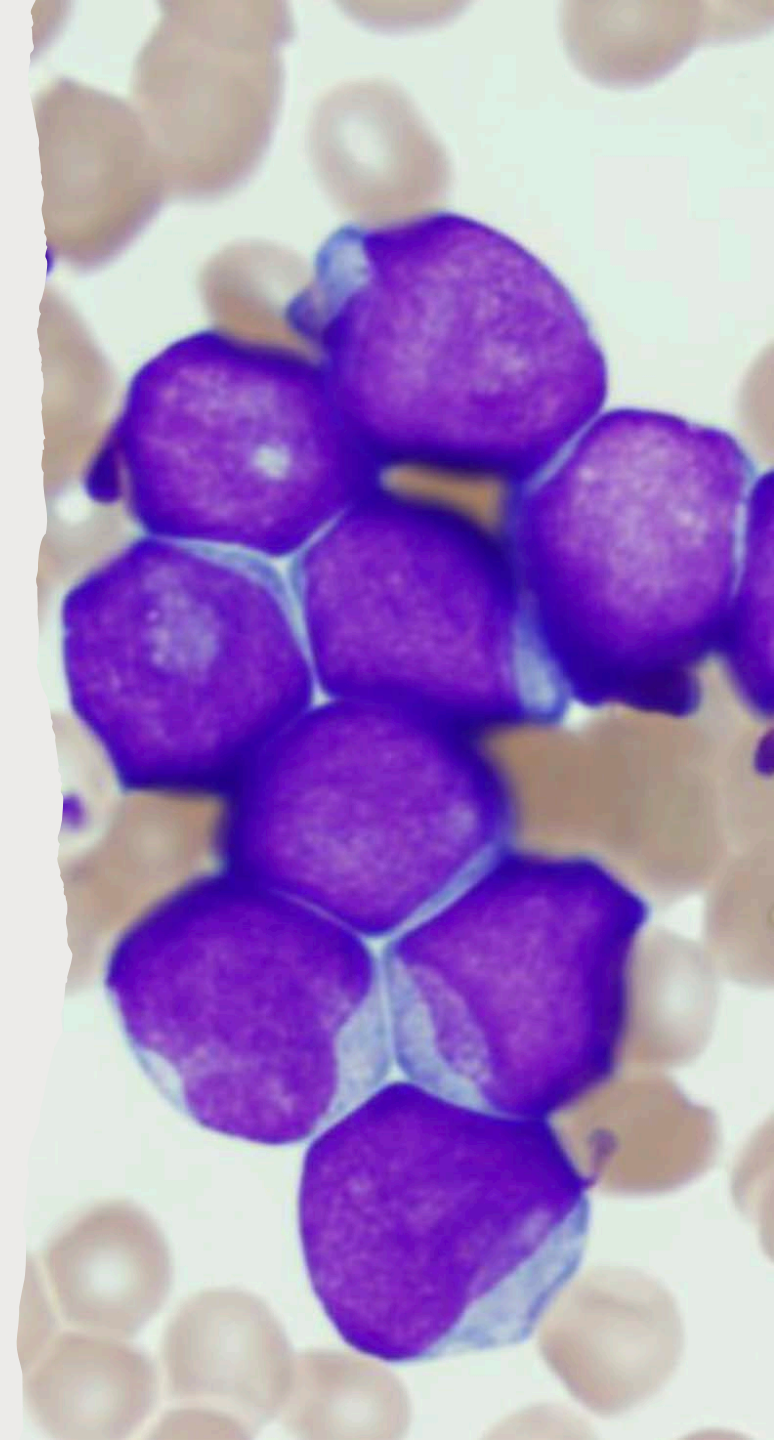
Specific Hypothesis: The inclusion of venetoclax in post remission therapy will lead to better MRD –ve CR rates amongst patients with MRD+ve disease after initial induction therapy compared to standard post- remission treatment.

Topics for the Day

- Definitions
 - Precision Medicine
 - myeloMATCH
- Current State of Therapeutic Options
 - Younger, Fit
 - Older or Frail
- Hurdles and Potential for this Strategy
- Discussion



- 74-yo man referred from emergency department of hospital in Appleton Wisconsin
- Hypertension, hyperlipidemia
- Smoked: ages 20-50
- H/O renal cell carcinoma with left-sided nephrectomy 19 years ago
- Presents with fatigue, night sweats, cough
- Complete blood count concerning for leukocytosis, anemia, thrombocytopenia and circulating blasts
- History of Chronic Obstructive Pulmonary Disease (COPD)





- Peripheral Blood:
 - WBC: 23.20×10^3 /uL
 - Hgn: 7.5 gm/dL
 - Platelet Count: 36×10^3 /uL
 - 37% circulating myeloid blasts
- Flow-cytometry with aberrant immunophenotype
- Bone Marrow Biopsy confirms Acute Myeloid Leukemia
- FISH: No recurrent cytogenetic mutations
- NPM-1 mut, FLT3 ITD

Fit for Intensive therapy?

Acute Myeloid Leukemia : 2024

Clinical trial available? Appropriate? If not, then...

Likely To Respond to cytotoxic therapy? Fit for Intensive therapy? Curable with cytotoxic therapy alone? If not, then...

FLT-3 Mutated

Venetoclax + Azacitidine or
Decitabine
DiNardo, NEJM 2020

Azacitidine + Sorafenib
Ohanian et al., Am J Hematology
2018; 93:1136-1141

IDH1 or IDH2 mutated

Venetoclax + Azacitidine or
Decitabine
DiNardo, NEJM 2020

Ivosidenib or Enasidenib
monotherapy
Dinardo et al., Blood 2017; Stein et
al, Blood 2015

Ivosidenib + Azacitidine
Montesinos et al., NEJM 2022

No targetable Mutations

Venetoclax + Azacitidine or Decitabine
DiNardo, NEJM 2020

HMA Exposed/Refractory

Venetoclax + Low-Dose Cytarabine
Wei et al., JCO, 2020

Glasdegib + Low-Dose Cytarabine – over 75 years of age
Cortes, Blood 2016

Complete Remission (+/- MRD) AND No significant comorbidities: Discuss Reduced-Intensity Allogeneic Stem Cell Transplant

Why BCL2 inhibition in AML?

Preclinical data

- Upregulated BCL-2 → evasion of apoptosis

ORIGINAL ARTICLE

Azacitidine and Venetoclax in Previously Untreated Acute Myeloid Leukemia

Courtney D. DiNardo, M.D., Brian A. Jonas, M.D., Ph.D., Vinod Pullarkat, M.D., Michael J. Thirman, M.D., Jacqueline S. Garcia, M.D., Andrew H. Wei, M.B., B.S., Ph.D., Marina Konopleva, M.D., Ph.D., Hartmut Döhner, M.D., Anthony Letai, M.D., Ph.D., Pierre Fenau, M.D., Ph.D., Elizabeth Koller, M.D., Violaine Havelange, M.D., Ph.D., *et al.*

CAL

VENETOCLAX

- Monotherapy in R/R AML: Overall response rate of 19%

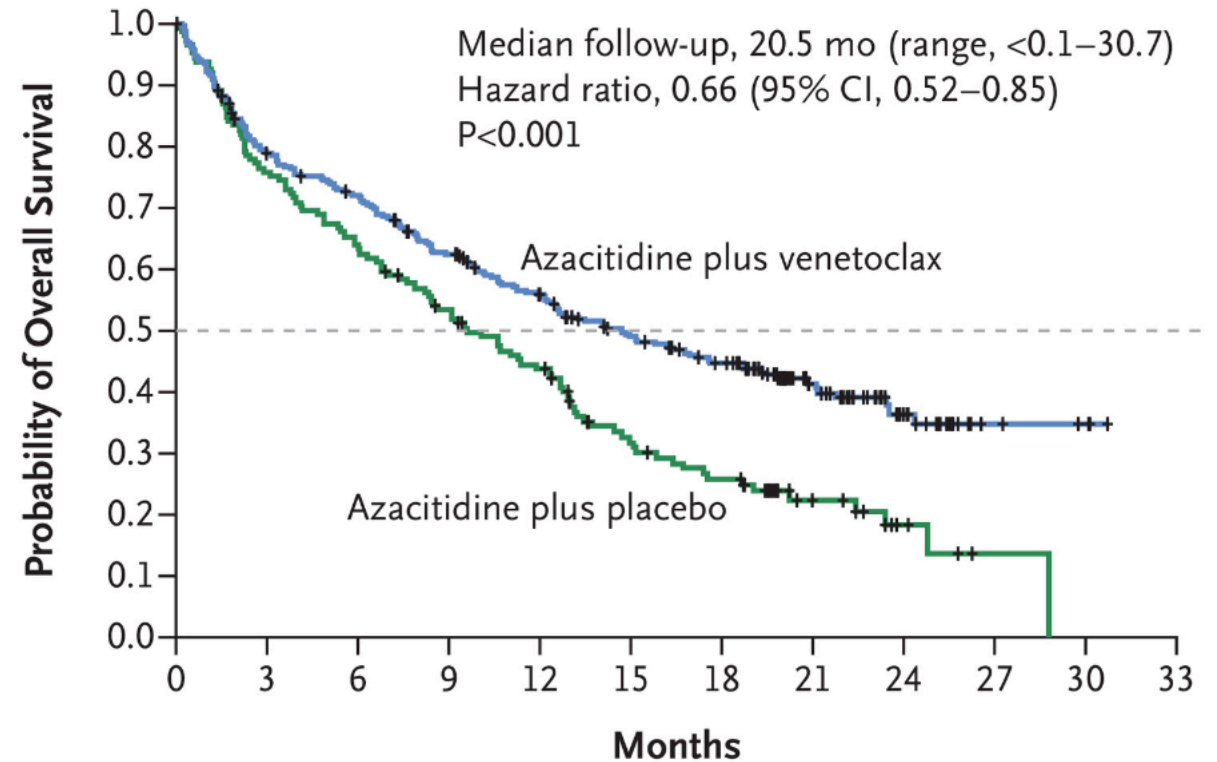
Azacitidine

- Non-specific, but some anti-MCL-1 activity

- Median Follow-up 20.5 months
- Median OS
 - Interventional Arm: 14.7 months
 - Control Arm: 9.6 months
 - HR for death, 0.66; 95% CI, 0.52 to 0.85; P<0.001
- Complete Remission Rate/Composite Complete Remission Rate (CR + CRi)
 - Interventional Arm: 36.7%/66.4%
 - Control Arm: 17.9%/28.3%

TAKE HOME MESSAGES

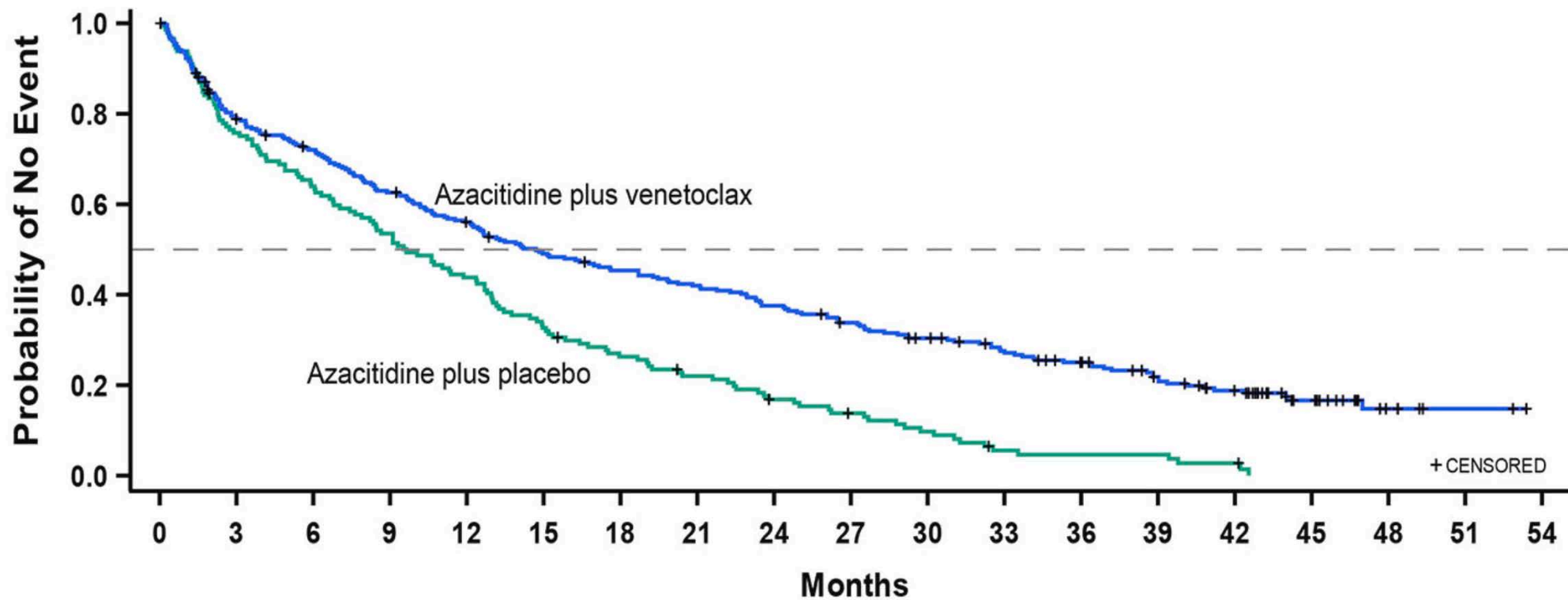
- Excellent New Option
- Safe, but not risk-free
- Unknown effectiveness compared to cytotoxic therapy



No. at Risk

Azacitidine plus venetoclax	286	219	198	168	143	117	101	54	23	5	3	0
Azacitidine plus placebo	145	109	92	74	59	38	30	14	5	1	0	0

Figure 1. Overall Survival



Patients at Risk

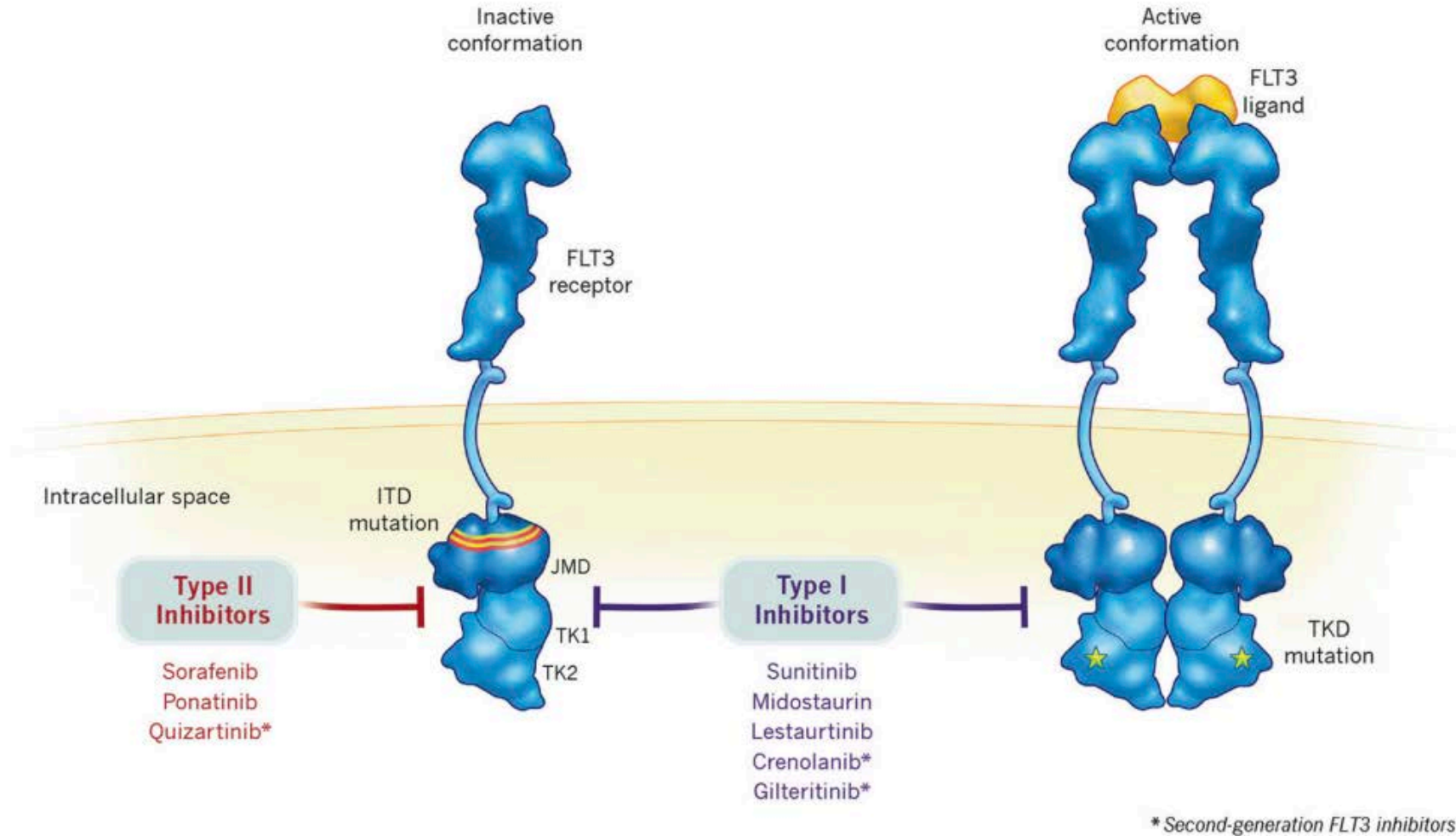
Azacitidine plus placebo

145 109 92 77 63 47 37 30 22 17 12 6 5 5 3 0

Azacitidine plus venetoclax

286 220 199 173 153 133 122 113 101 89 78 67 57 45 34 18 6 2 0

FLT3 inhibition



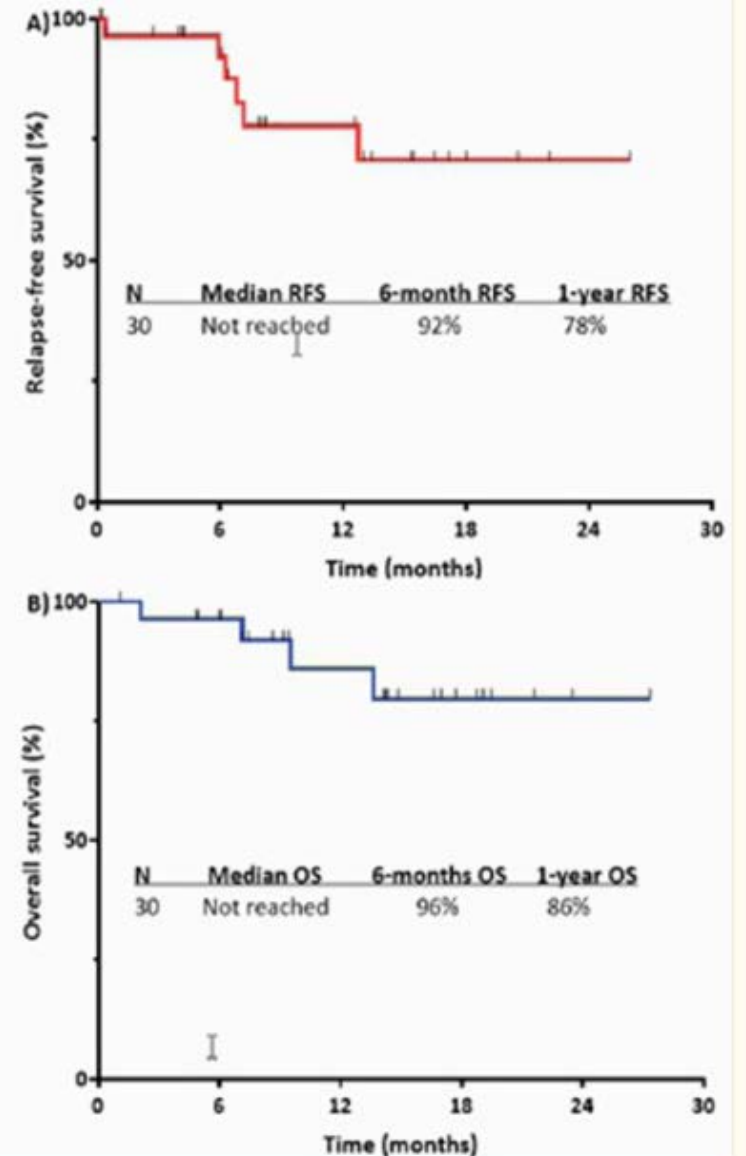
What about triplet therapy?

- Gilteritinib plus Aza-Ven N=30
- ND FLT3mut AML
- MD Anderson team
- Phase II, FLT3mut (ITD and TKD)
- Newly Diagnosed AML, ages 18-60

CR/CRi	29/30 (96%)
MRDneg by flow	26/30 (87%)
MRDneg by PCR	27/30 (90%)

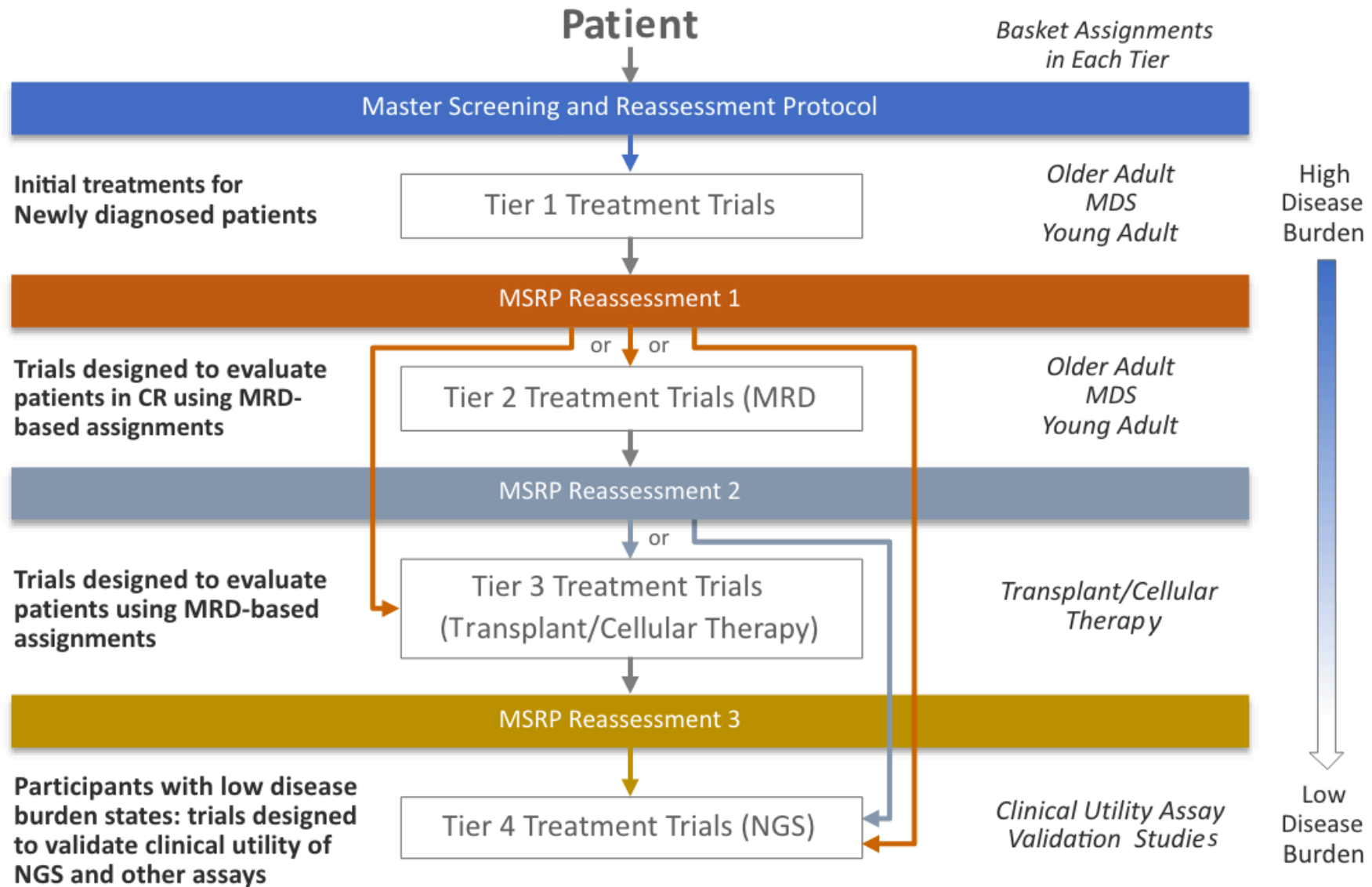
Figure 1 – Outcomes for patients with newly diagnosed FLT3-mutated AML.

(A) Relapse-free survival and (B) overall survival.



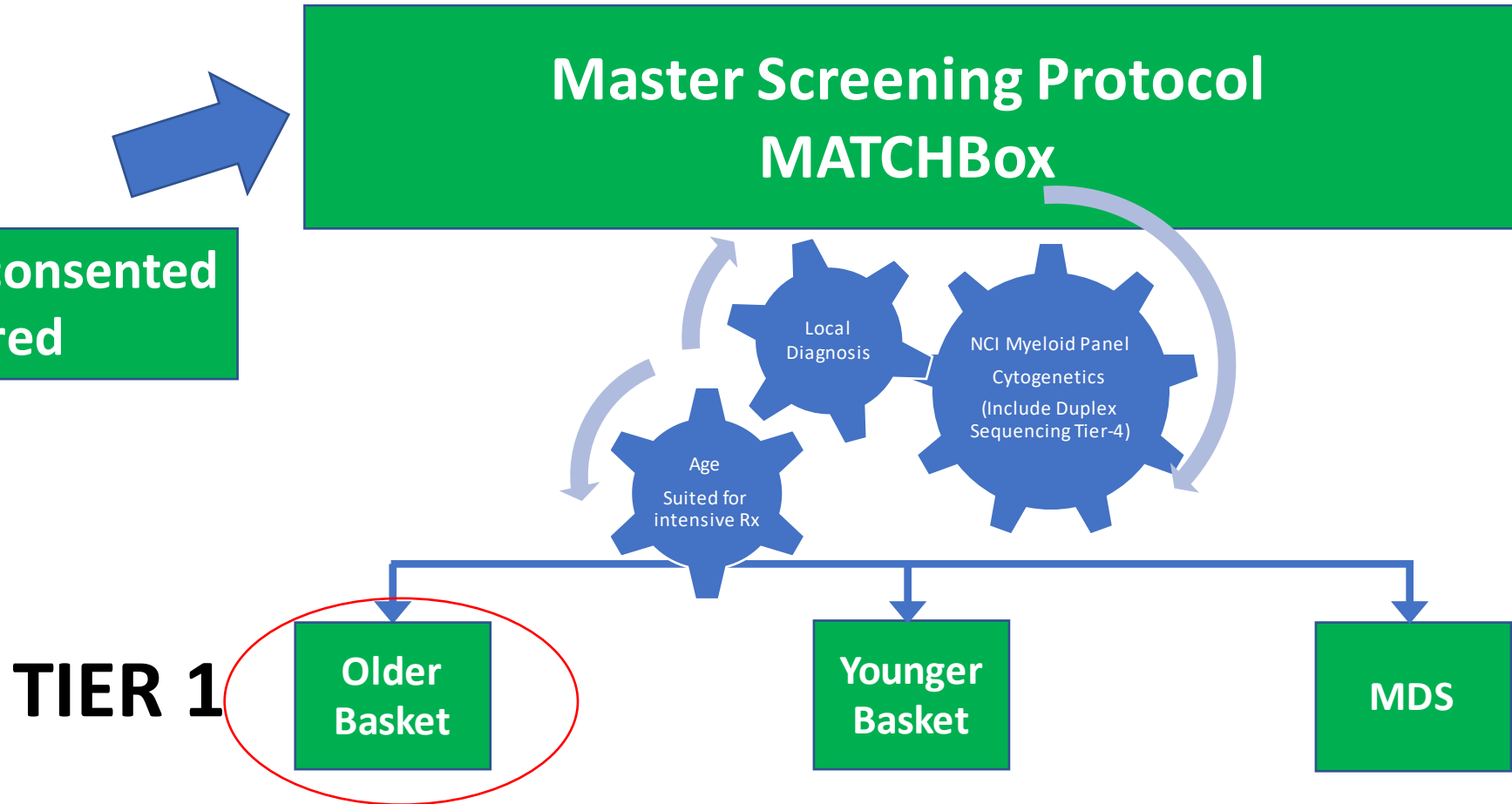
What would this patient's
treatment look like on
myeloMATCH?

MyeloMATCH MSRP Schema

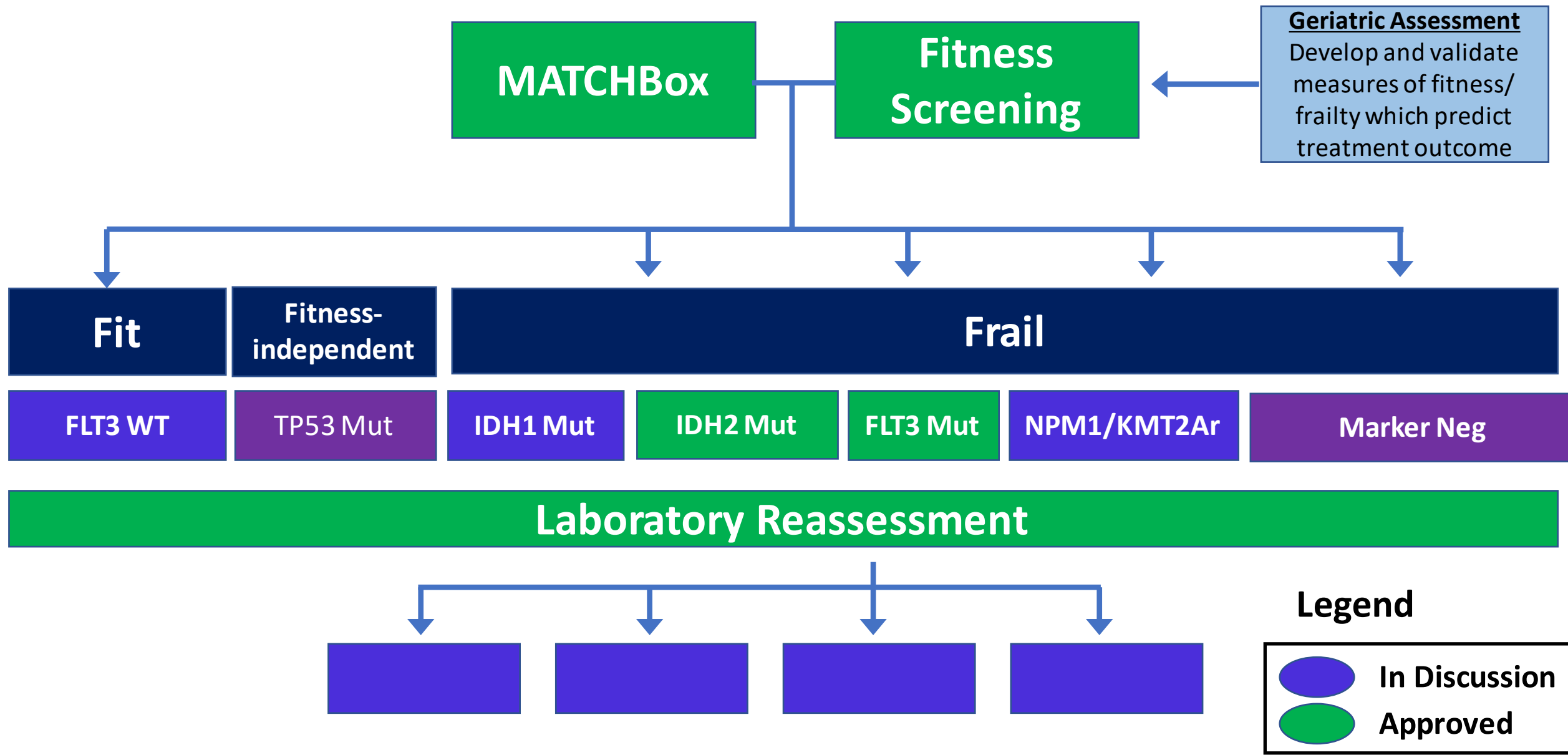


myeloMATCH option

Patient would be consented and registered

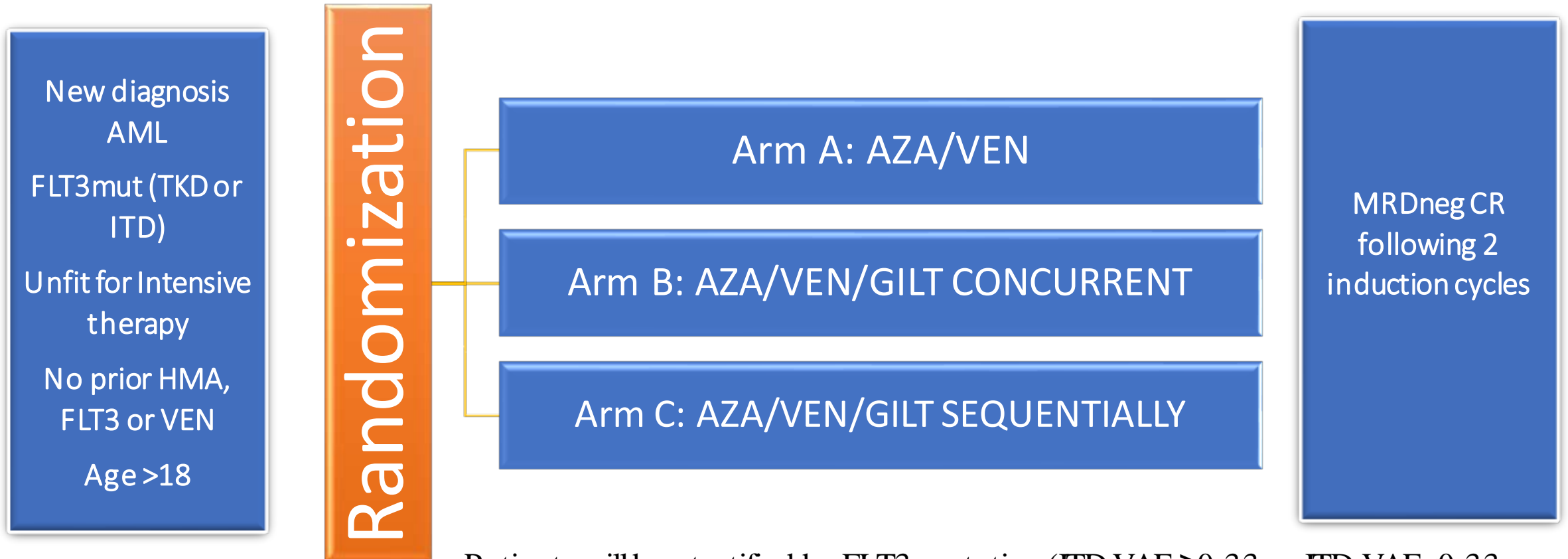


First Generation Studies in the Older Basket



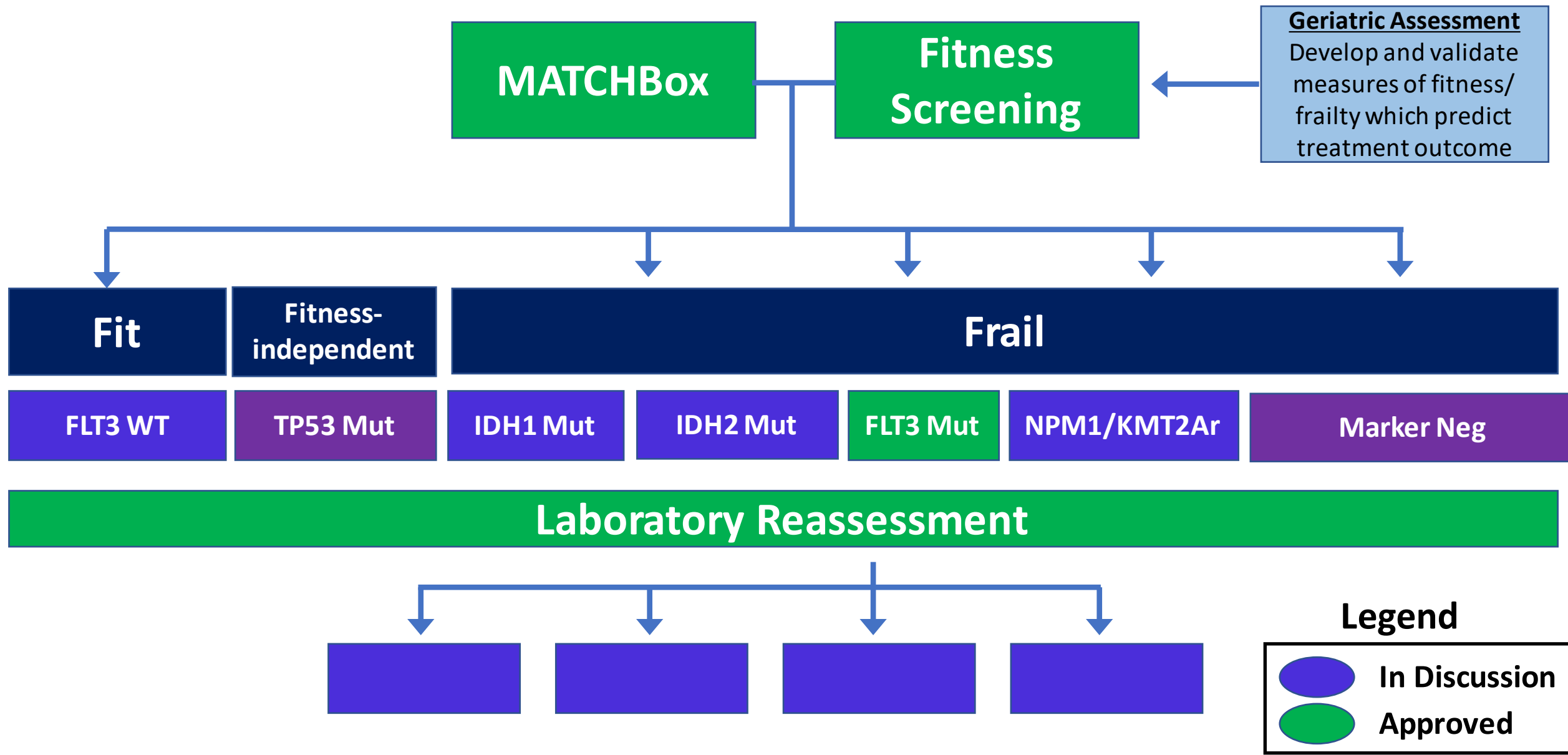
MM10A-EA02

Study co-chairs: Jessica Altman, Alexander Perl



Patients will be stratified by FLT3 mutation (ITD VAF ≥ 0.33 vs ITD VAF < 0.33 , or TKD) and age (< 70 , ≥ 70 years)
N=147

First Generation Studies in the Older Basket



What about the patients for
whom there is no trial?

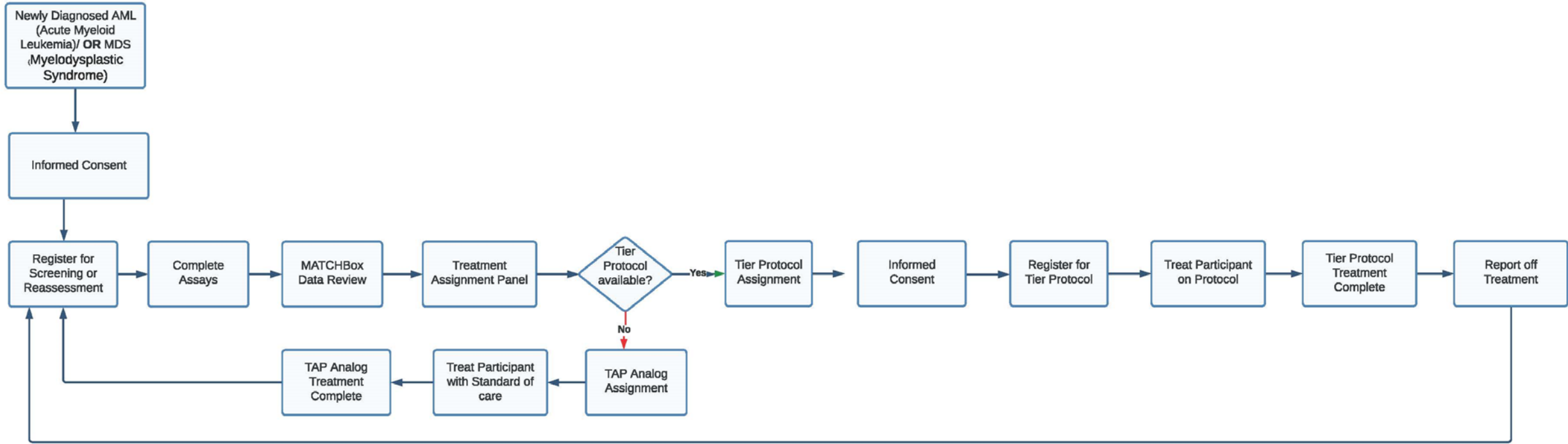
Tier Advancement Pathway (TAP)

Screening and Reassessment (MSRP): To evaluate the feasibility of MATCHBox generating all data needed for assignment to a myeloMATCH clinical trial or Tier Advancement Pathway (TAP) within 72 hours of MDNet receipt of all required specimens for initial therapy and within 10 days for subsequent therapy.

Tier Advancement Pathway (TAP): To enable participants who are not matched to an investigational myeloMATCH treatment substudy to receive standard of care while remaining on the MSRP to maintain access to later tiers of treatment substudies.

MyeloMATCH Treatment Options

Updated Aug 03, 2023



What is the current status of the platform?

MyeloMATCH: Current Status

- 20 MyeloMATCH-specific and CTEP-wide CRADAs with industry collaborators in place to support clinical sub-studies
- FDA CDRH approved MyeloMATCH MSRP IDE on 14 DEC 2023.
- FDA CDER gave the IND “safe to proceed” on 10 JAN 2024.
- Activated May of 2024
- More than 200 people currently enrolled
- 11 additional concepts are currently in development
 - 3-Older AML basket; 3-MDS basket; 3-Younger AML basket; 2-Transplant basket

What Can We Learn...

- Is the platform feasible?
- Which induction arm leads to highest rate of MRDu CRs?
- What are the comparative toxicities?
- Can we confirm MRDneg CR as a reliable surrogate for event-free survival and overall survival?
- What disease features, other than molecular signature and chromosome, impact outcome? What patient features?
- Can we accrue at sites outside of large leukemia centers?



MyeloMATCH Hurdles

- Immense Logistical Feat
 - Collaboration amongst the cooperative groups
 - Collaboration with pharmaceutical partners
 - Working with FDA
- Challenges facing the sites
- Dealing with a dynamic environment

- Even if the logistics all go perfectly...
 - Not enough drugs for the targets
 - Not enough studies for the patients
 - Not enough patients enrolled on the trials

MyeloMATCH Potential

Understand resistance pathways
Prevent the emergence of subclones
Be aware of predisposition syndromes

● Safely
Debulk
Disease

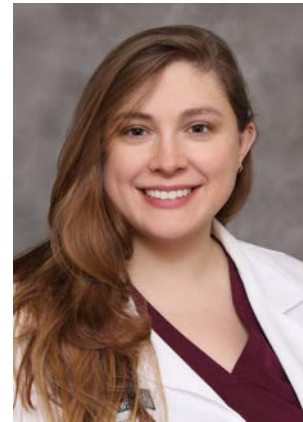
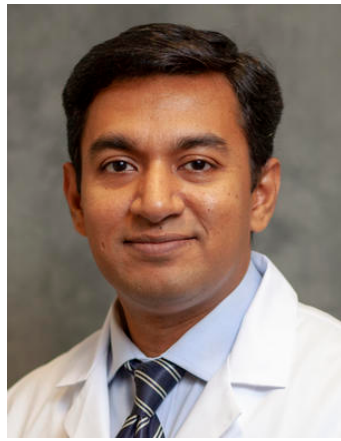
Mitigate toxicity
Target the most primitive clones
Derive meaningful comparison data amongst regimens
Allow widespread participation

● Eradicate
Measurable
Residual
Disease

● Prevent
Relapse

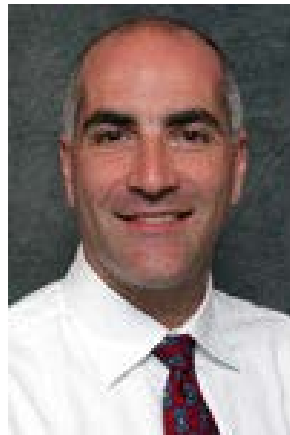
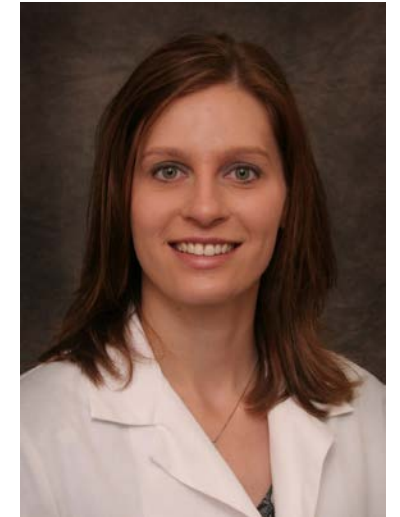
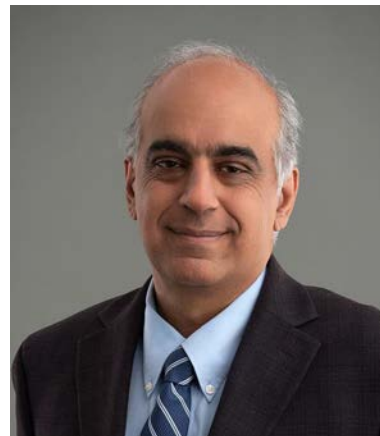
Avoid additional toxicities
Capitalize on immunotherapy
Prevent need for long-term therapy
Cure vs. Control

Leukemia Faculty and APPs



Malignant Hematology Faculty and Staff

- Transplant and Cellular Therapy





Thanks so much for the
invitation to speak

Very happy to take any
questions

NCI myeloMATCH Leadership Team

Rich Little; Jerry Radich; Percy Ivy; Rich Stone; Mark
Litzhow; Harry Erba; Geoff Uy; Ehab Atallah; Toyosi Odenike;
Selina Lugar; Patrick Stiff; Steve Gore and many many many
others

knowledge changing life

