



Il trapianto allogenico integrato nella strategia terapeutica delle malattie ematologiche

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DEGLI STUDI
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Unità di Malattie del Sangue e Trapianto Midollo Osseo - Programma Terapie Cellulari e Ricerca in Ematologia
Unit of Blood Diseases and Bone Marrow Transplantation - Cell Therapies and Hematology Research Program

Sistema Socio Sanitario



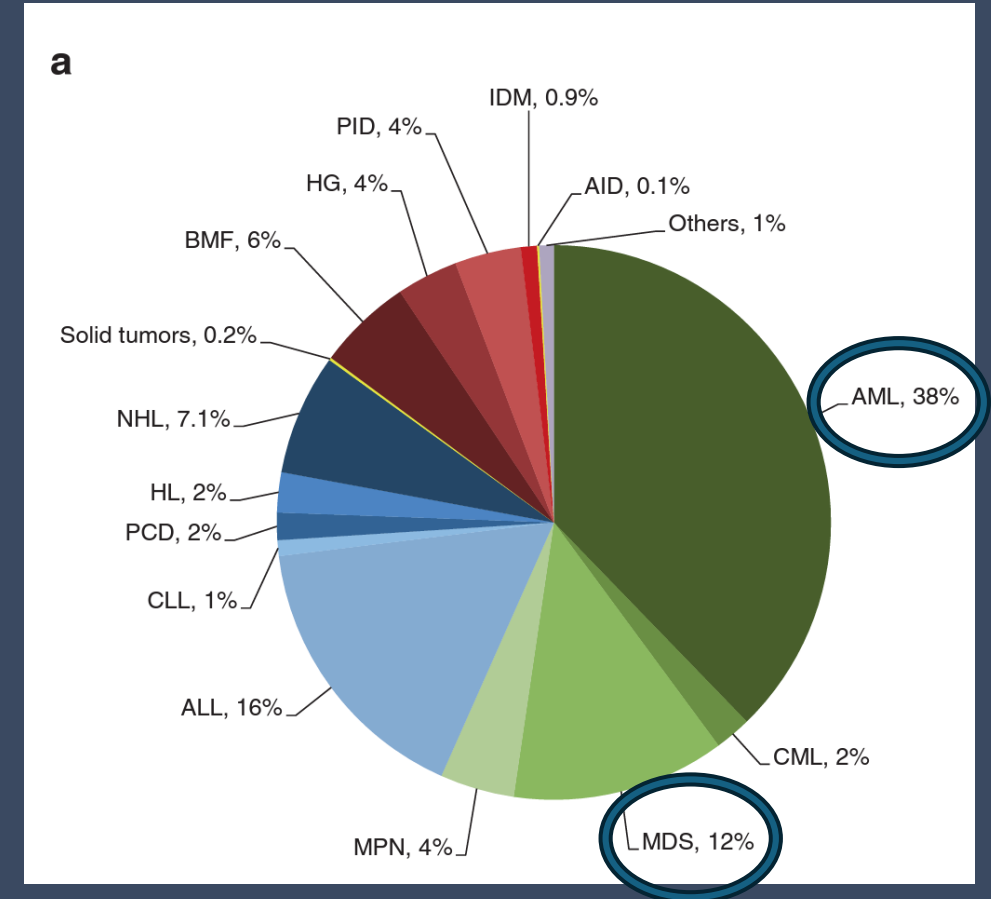
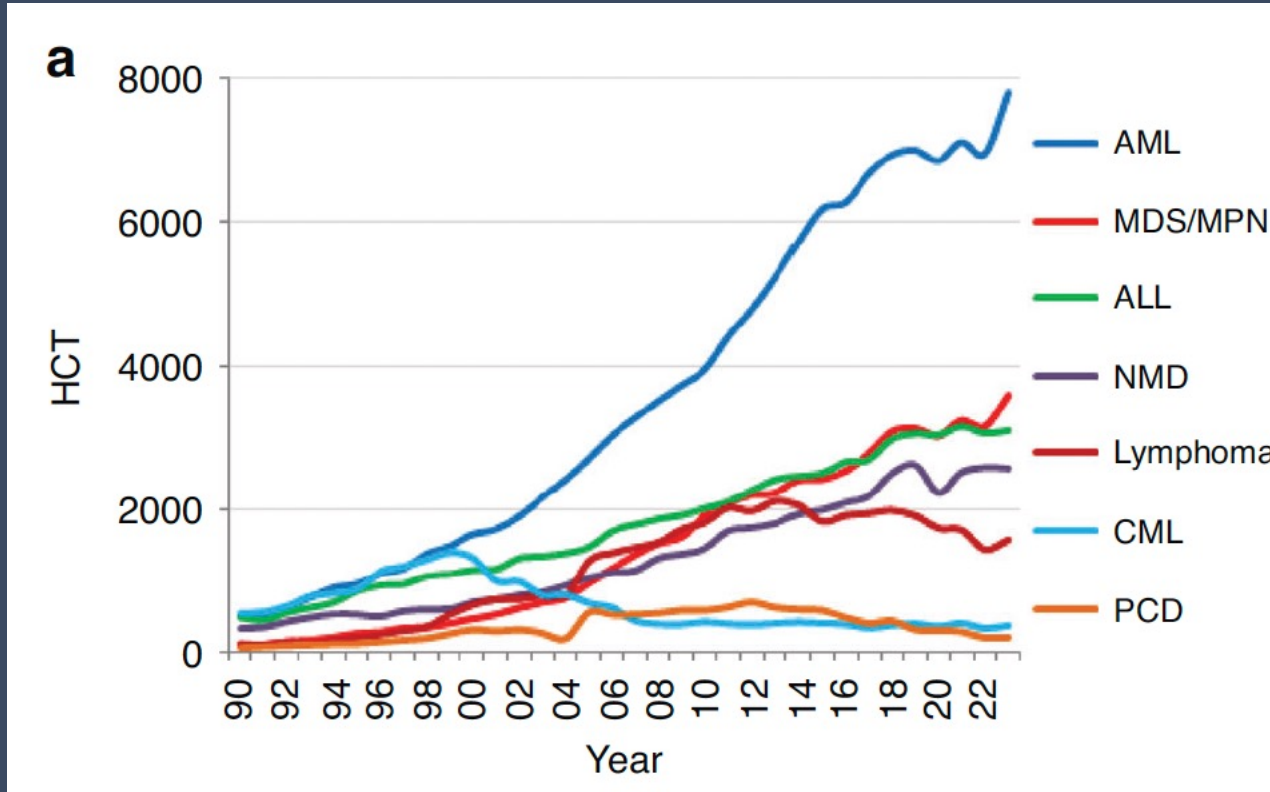
Regione
Lombardia

ASST Spedali Civili

Disclosures

	Jazz Pharmaceuticals	Pfizer	Mallinckrodt/ Therakos	Sanofi	Alexion	Novartis	Medac Pharma
Speaking	X	X	X	X	X		X
Research grant			X	X			
Advisory board		X		X	X	X	
Travel Grant	X			X			X

TRENDS IN TRANSPLANT PRACTICE IN EUROPE



Passweg JR et al Hematopoietic cell transplantation and cellular therapy survey of the EBMT: monitoring of activities and trends over 30 years. Bone Marrow Transplant. 2021 Jul;56(7):1651-1664.

Passweg JR et al The 2023 EBMT report on hematopoietic cell transplantation and cellular therapies. Increased use of allogeneic HCT for myeloid malignancies and of CAR-T at the expense of autologous HCT. Bone Marrow Transplant. 2025 Apr;60(4):519-528.

ALLOHCT IN ELDERLY PATIENTS – CIBMTR

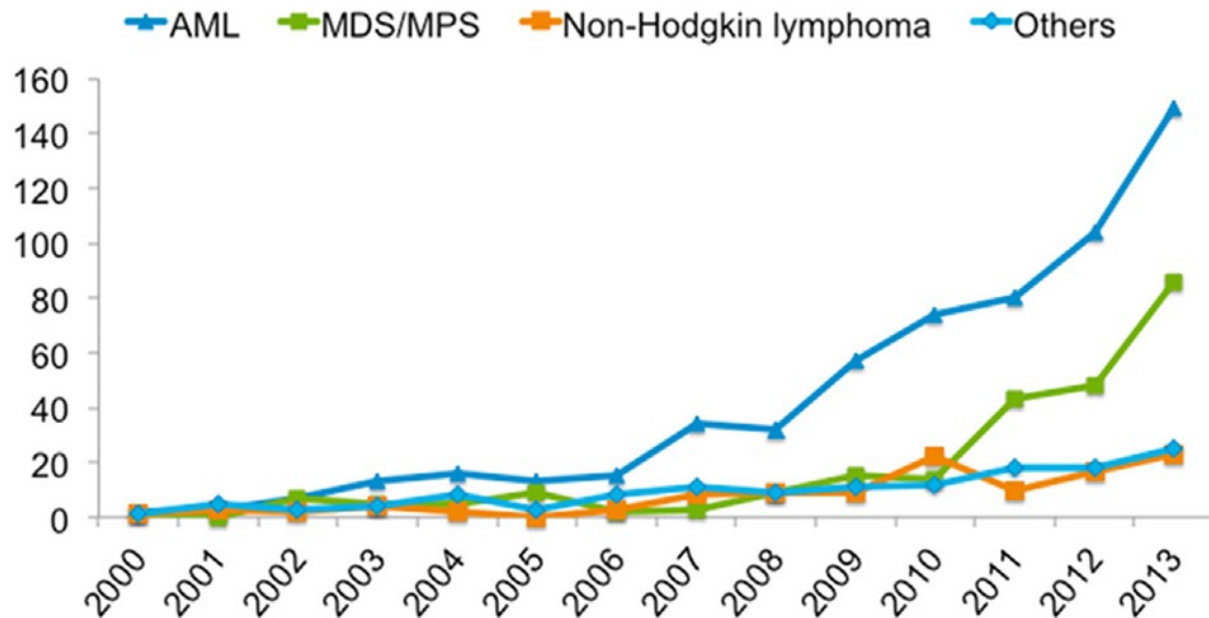


Figure 1. Absolute number of allogeneic transplants in recipients 70 years and older from 2000 to 2013 reported to the CIBMTR by disease category.

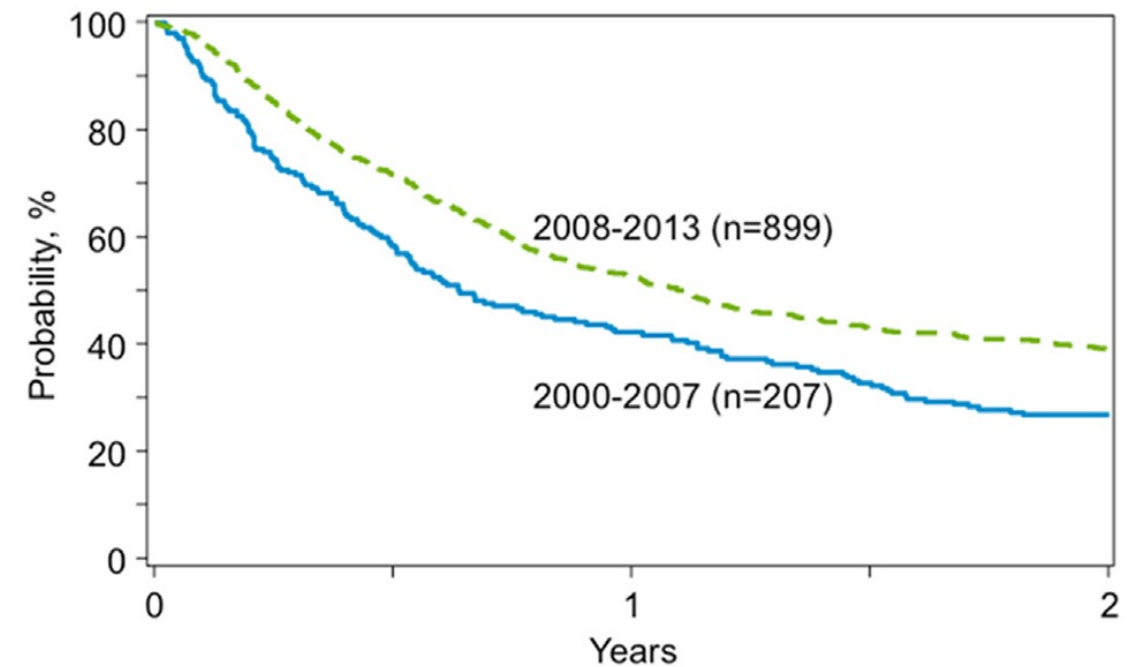
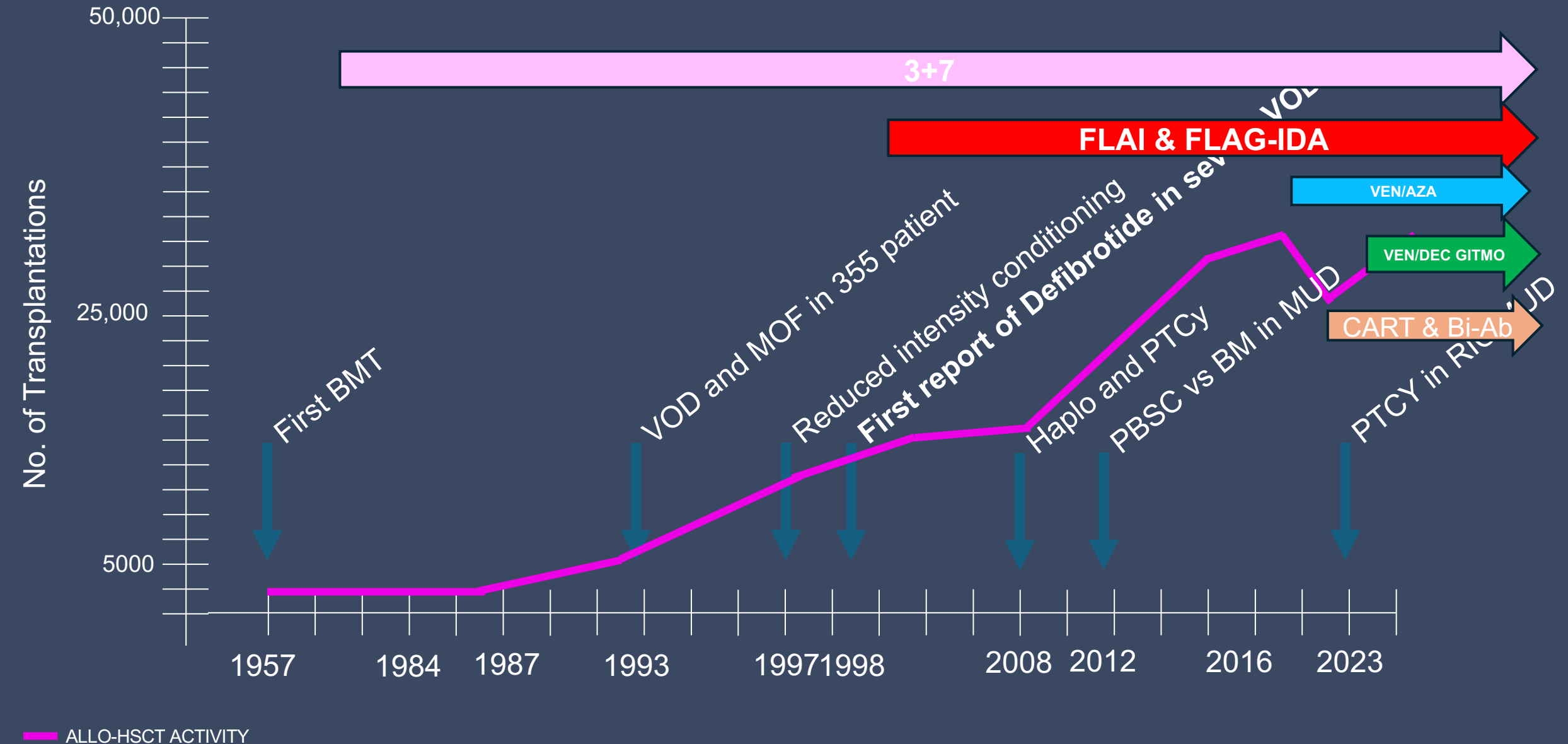


Figure 2. OS for allogeneic HCT recipients 70 years and older transplanted between 2000-2007 and 2008-2013.

ALLO-HSCT activity across two centuries



ALLO-HSCT activity across two centuries

50,000

BLOOD, 21 JANUARY 2010 • VOLUME 115, NUMBER 3

Table 4. Standardized reporting for correlation of cytogenetic and molecular genetic data in AML with clinical data

Genetic group	Subsets
Favorable	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</i> -ITD (normal karyotype) Mutated <i>CEBPA</i> (normal karyotype)
Intermediate-I*	Mutated <i>NPM1</i> and <i>FLT3</i> -ITD (normal karyotype) Wild-type <i>NPM1</i> and <i>FLT3</i> -ITD (normal karyotype) Wild-type <i>NPM1</i> without <i>FLT3</i> -ITD (normal karyotype)
Intermediate-II	t(9;11)(p22;q23); <i>MLL3-MLL</i> Cytogenetic abnormalities not classified as favorable or adverse†
Adverse	inv(3)(q21q26.2) or t(3;3)(q21;q26.2); <i>RPN1-EVI1</i> t(6;9)(p23;q34); <i>DEK-NUP214</i> t(v;11)(v;q23); <i>MLL</i> rearranged -5 or del(5q); -7; abn(17p); complex karyotype‡

Table 5. 2017 ELN risk stratification by genetics

Risk category*	Genetic abnormality
Favorable	t(8;21)(q22;q22.1); <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</i> -ITD or with <i>FLT3</i> -ITD ^{low} † Biallelic mutated <i>CEBPA</i>
Intermediate	Mutated <i>NPM1</i> and <i>FLT3</i> -ITD ^{high} † Wild-type <i>NPM1</i> without <i>FLT3</i> -ITD or with <i>FLT3</i> -ITD ^{low} † (without adverse-risk genetic lesions) t(9;11)(p21.3;q23.3); <i>MLL3-KMT2A</i> ‡ Cytogenetic abnormalities not classified as favorable or adverse
Adverse	t(6;9)(p23;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> inv(3)(q21.3q26.2) or t(3;3)(q21.3;q26.2); <i>GATA2, MECOM(EVI1)</i> -5 or del(5q); -7; -17/abn(17p) Complex karyotype,§ monosomal karyotype Wild-type <i>NPM1</i> and <i>FLT3</i> -ITD ^{high} † Mutated <i>RUNX1</i> ¶ Mutated <i>ASXL1</i> ¶ Mutated <i>TP53</i> ##

Table 6. 2022 ELN risk classification by genetics at initial diagnosis*

Risk category†	Genetic abnormality
Favorable	- t(8;21)(q22;q22.1)/<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</i>-ITD - bZIP in-frame mutated <i>CEBPA</i>
Intermediate	- 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>MLL3::KMT2A</i>†,¶ - Cytogenetic and/or molecular abnormalities not classified as favorable or adverse
Adverse	- 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, MECOM(EVI1)</i> - t(3q26.2;q21.3)/<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>§§

25,000

5000

1957

1984

1987

1993

1997

1998

2008

2012

2016

2023

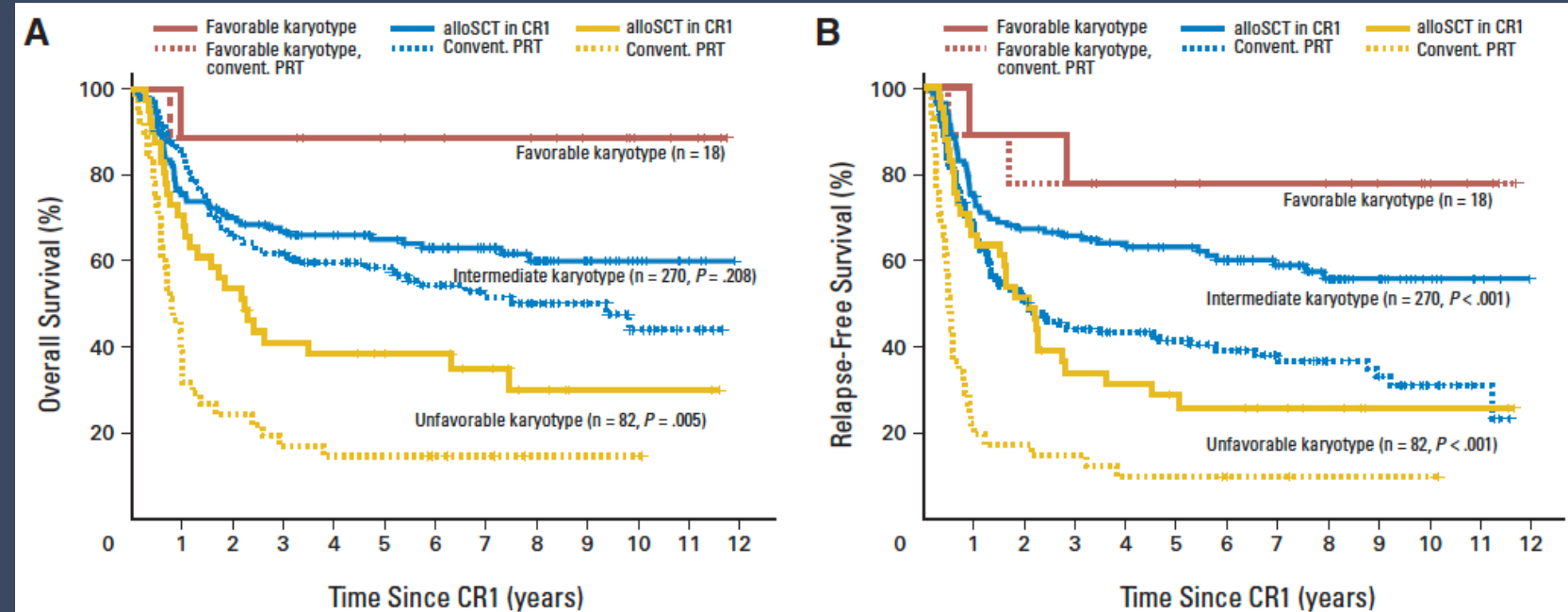
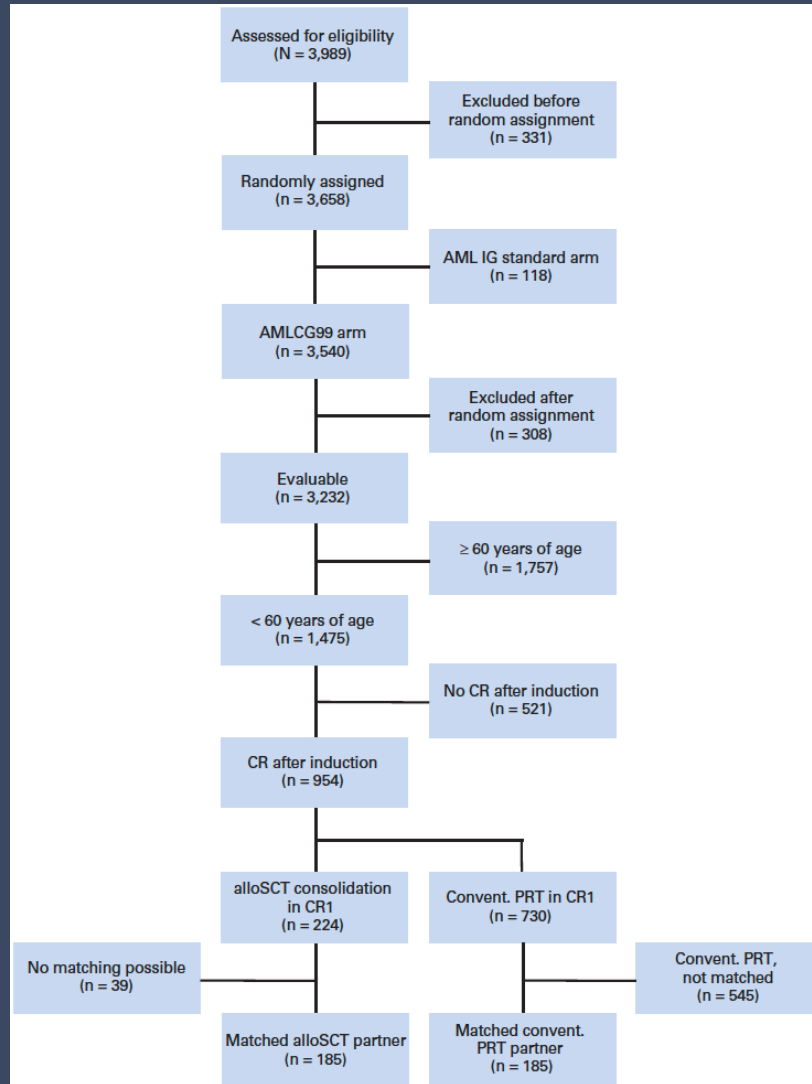
ALLO-HSCT ACTIVITY

Allogeneic Transplantation Versus Chemotherapy as Postremission Therapy for Acute Myeloid Leukemia: A Prospective Matched Pairs Analysis

Matthias Stelljes, Utz Krug, Dietrich W. Beelen, Jan Braess, Maria C. Sauerland, Achim Heinecke, Sandra Liggies, Tim Sauer, Petra Teschauer, Gabriela B. Thoenigsen, Birgitta Borning, Hans-J. Kolb

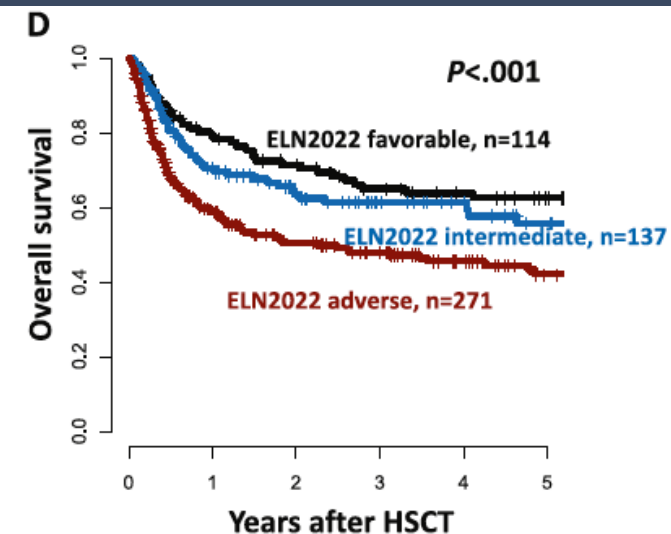
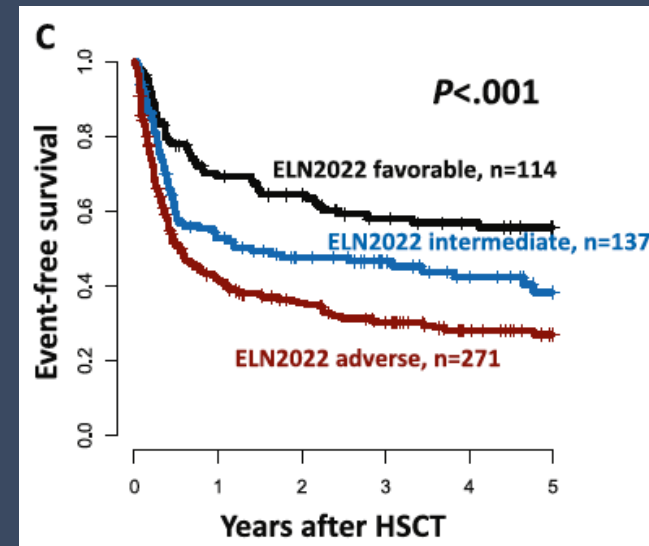
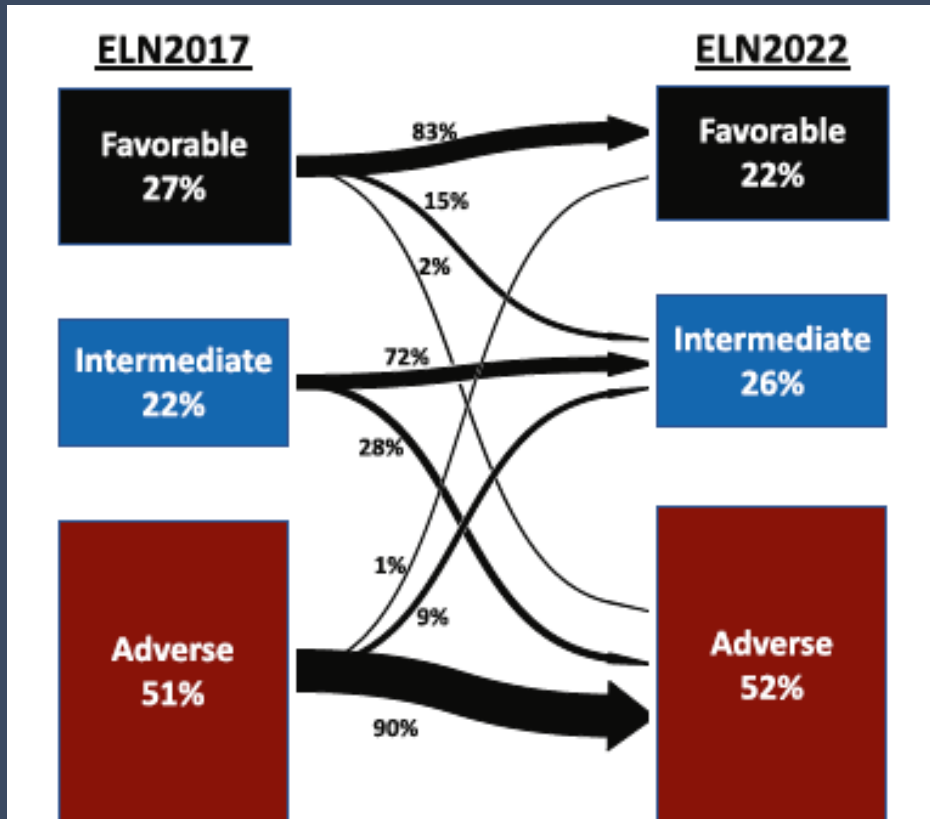
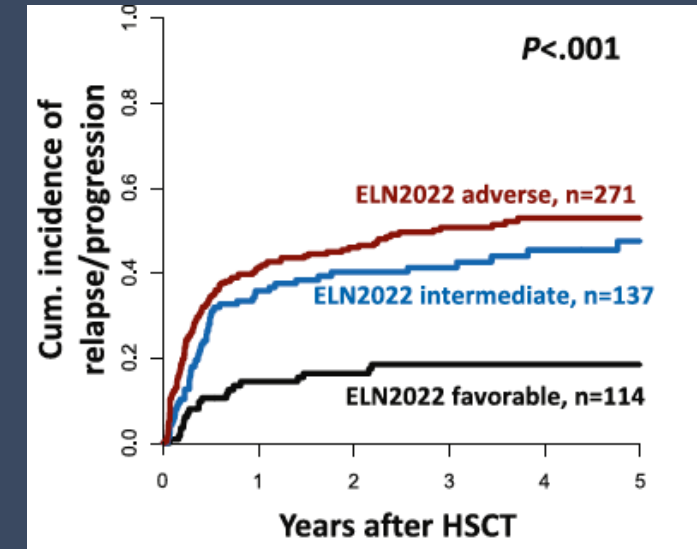
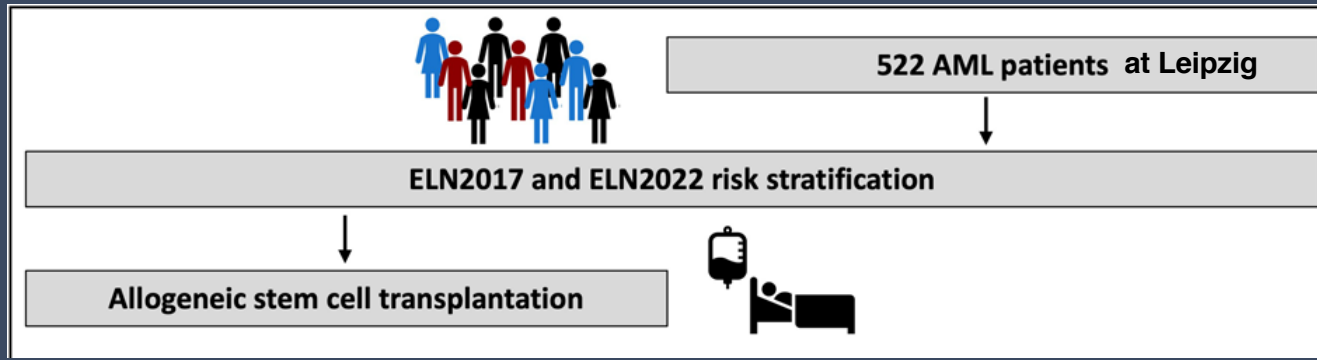
VOLUME 32 · NUMBER 4 · FEBRUARY 1 2014

JOURNAL OF CLINICAL ONCOLOGY



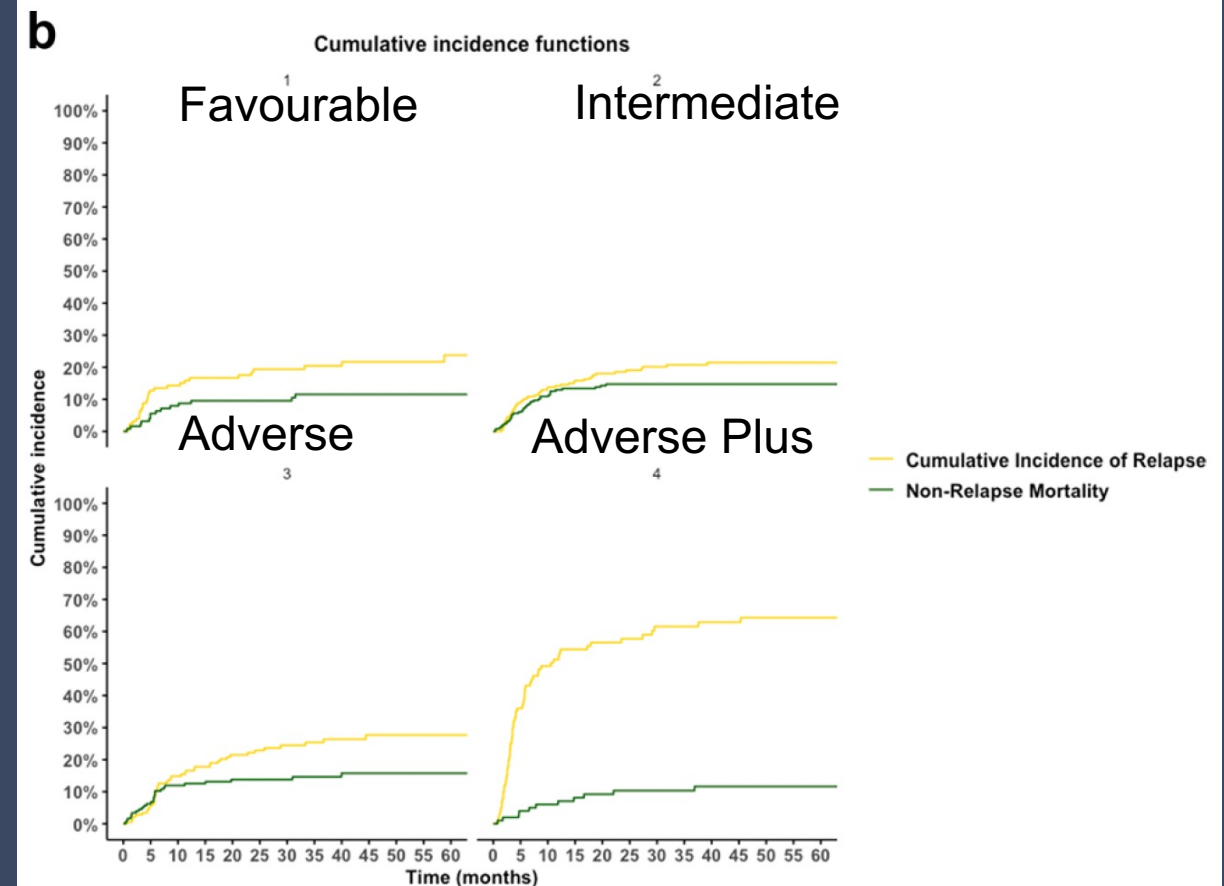
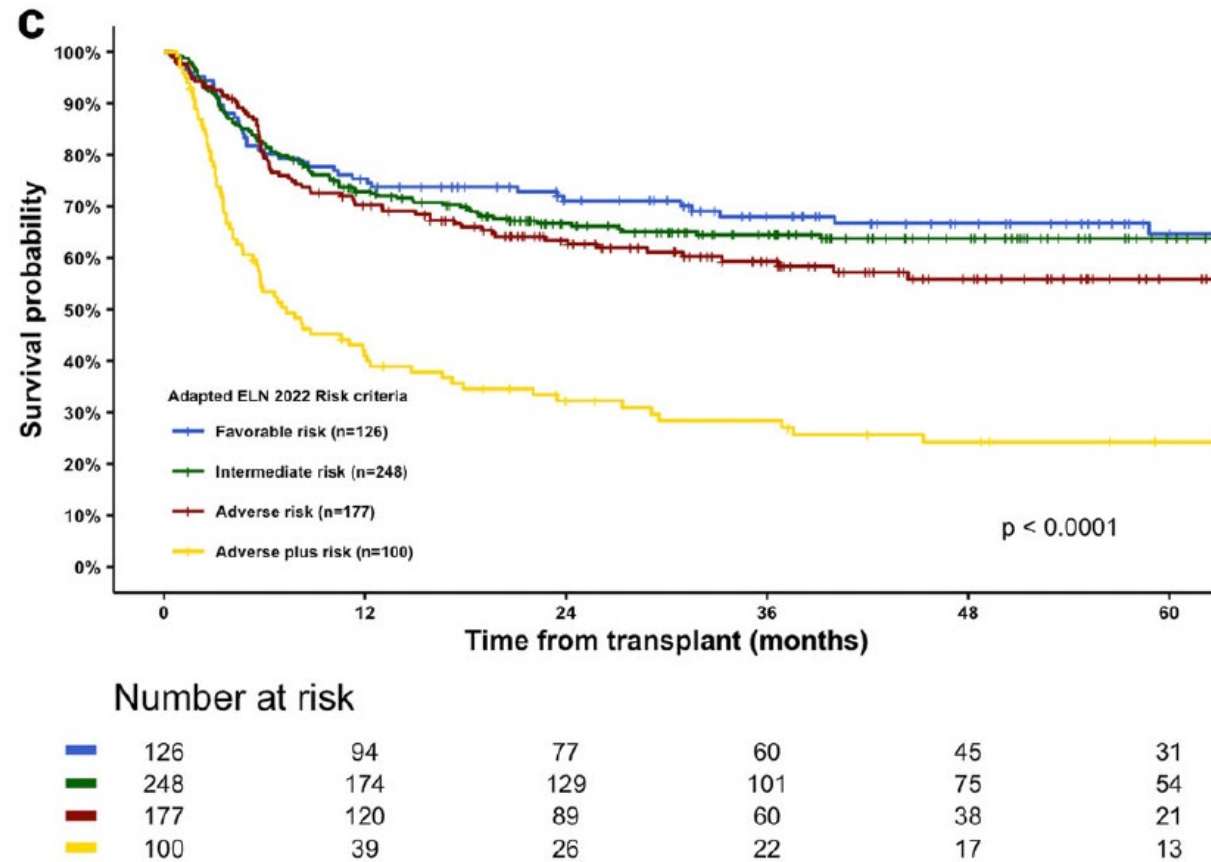
AlloSCT in CR1 (linea cont.) Vs. Chemotherapy (dotted line)

AML_ELN22 new classification_validation



Allo-HCT refined ELN 2022 risk classification: validation of the Adverse-Plus risk group in AML patients undergoing allogeneic hematopoietic cell transplantation within the Spanish Group for Hematopoietic Cell Transplantation (GETH-TC)

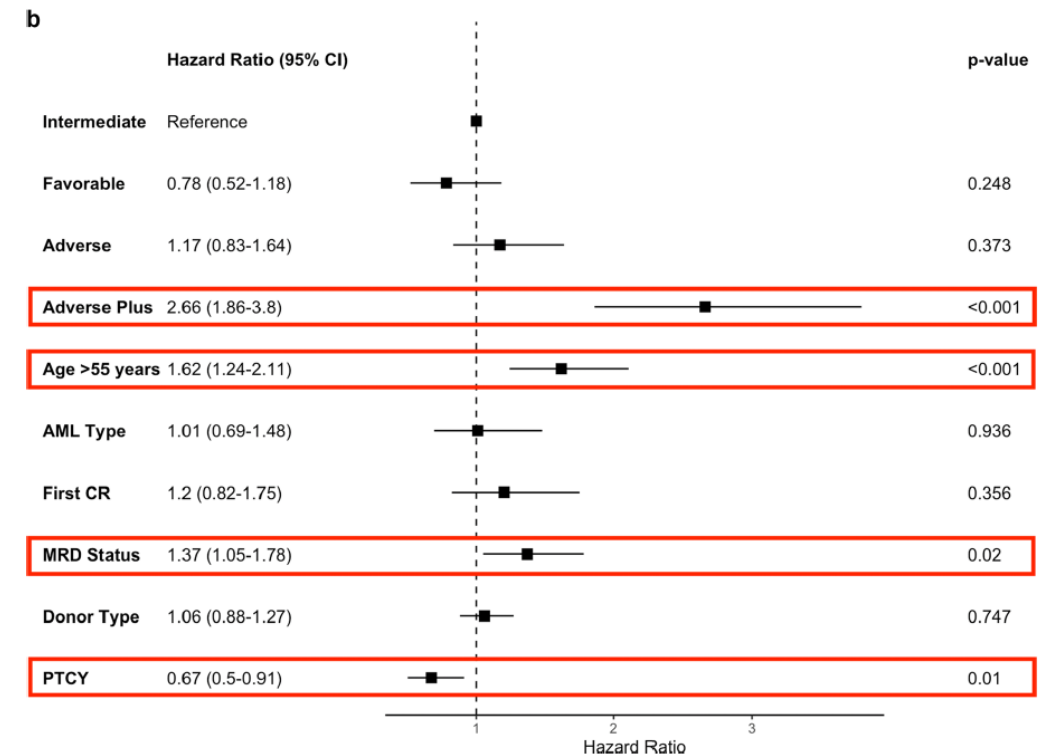
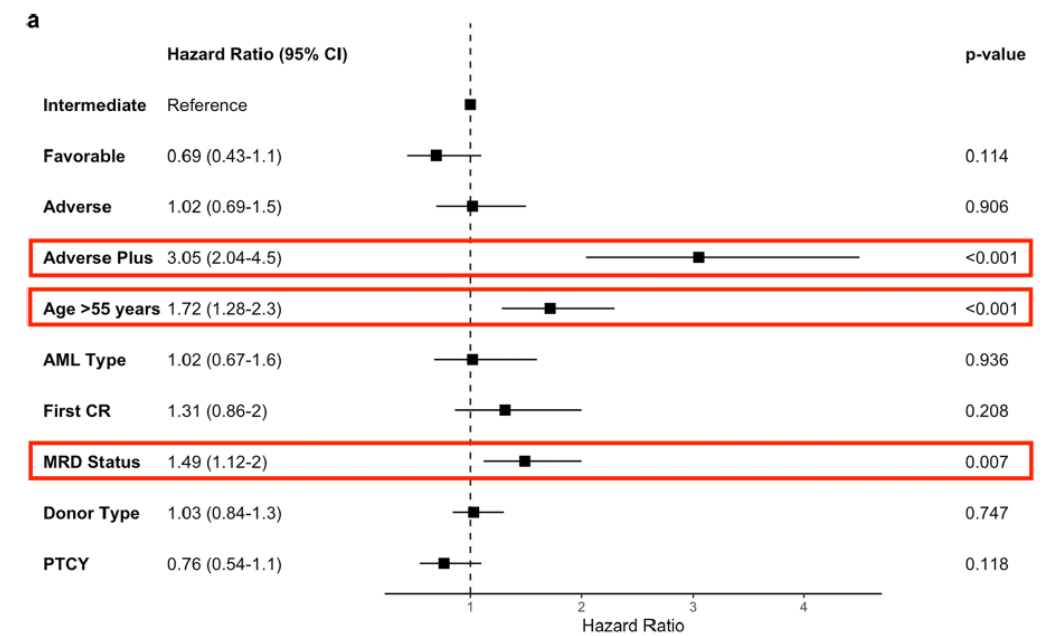
Carlos Jiménez-Vicente¹, Jordi Esteve¹, Mónica Baile-González², Estefanía Pérez-López², Carmen Martín Calvo³, Clara Aparicio³, Itziar Oñatividad Ormategi⁴, Albert Esquirol⁵, Felipe Peña-Muñoz⁶, Sara Fernández-Luis⁷, Inmaculada Heras Fernando⁸, Ana Pilar González-Rodríguez⁹, Alberto López-García¹⁰, Jose Luis López-Lorenzo¹⁰, Tamara Torrado¹¹, Adolfo Jesús Sáez Marín¹², Cynthia Acosta Fleytas¹³, Lucía García¹⁴, Sara Villar¹⁵, Silvia Filaferro¹⁶, Pascual Balsalobre¹⁶, María Jesús Pascual Cascón^{16,17} and María Queralt Salas¹



Allo-HCT refined ELN 2022 risk classification: validation of the Adverse-Plus risk group in AML patients undergoing allogeneic hematopoietic cell transplantation within the Spanish Group for Hematopoietic Cell Transplantation (GETH-TC)

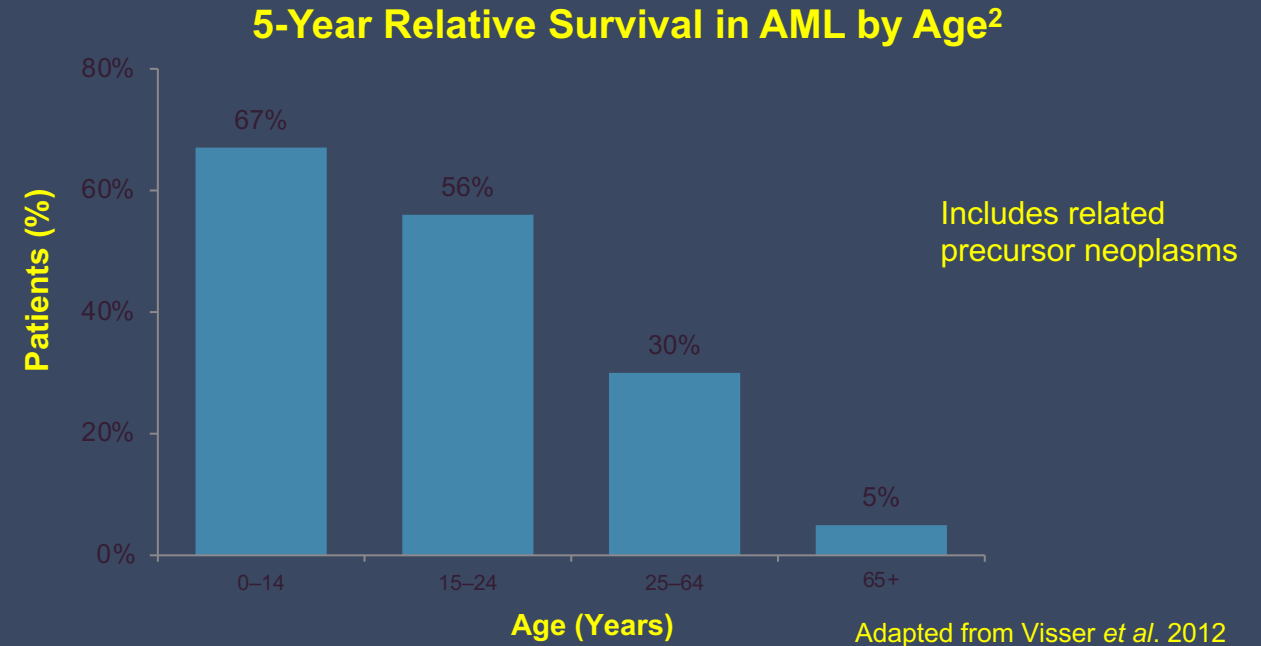
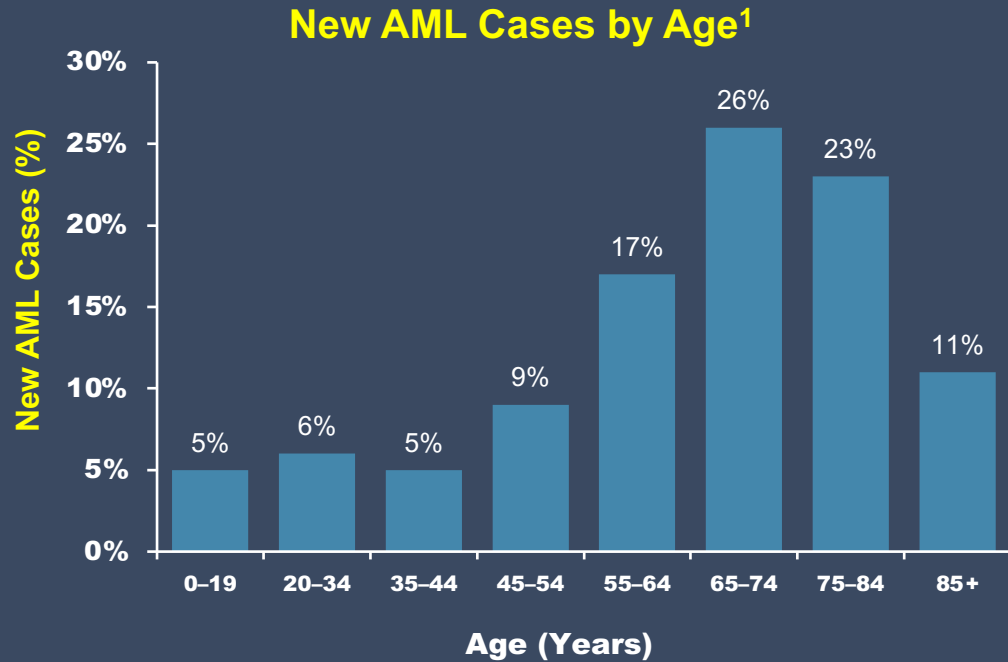
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- The Allo-HCT–Refined ELN 2022 classification further stratifies adverse-risk AML into Adverse* and Adverse-Plus (AdvP), identifying a subgroup with significantly inferior outcomes after allo-HCT.
- Patients classified as AdvP (complex karyotype, MECOM(EVI1) rearrangement, and/or TP53 mutation/deletion) showed markedly reduced 5-year OS (≈28–32%) and LFS (≈24%) with a very high cumulative incidence of relapse (≈64%).
- In multivariable analysis, AdvP status, age >55 years, and pre-transplant MRD positivity were independently associated with worse OS and LFS, while PTCY-based GVHD prophylaxis was associated with improved LFS.



AML IS MOST FREQUENTLY DIAGNOSED IN OLDER ADULTS¹

- Prognosis worsens with increasing age²



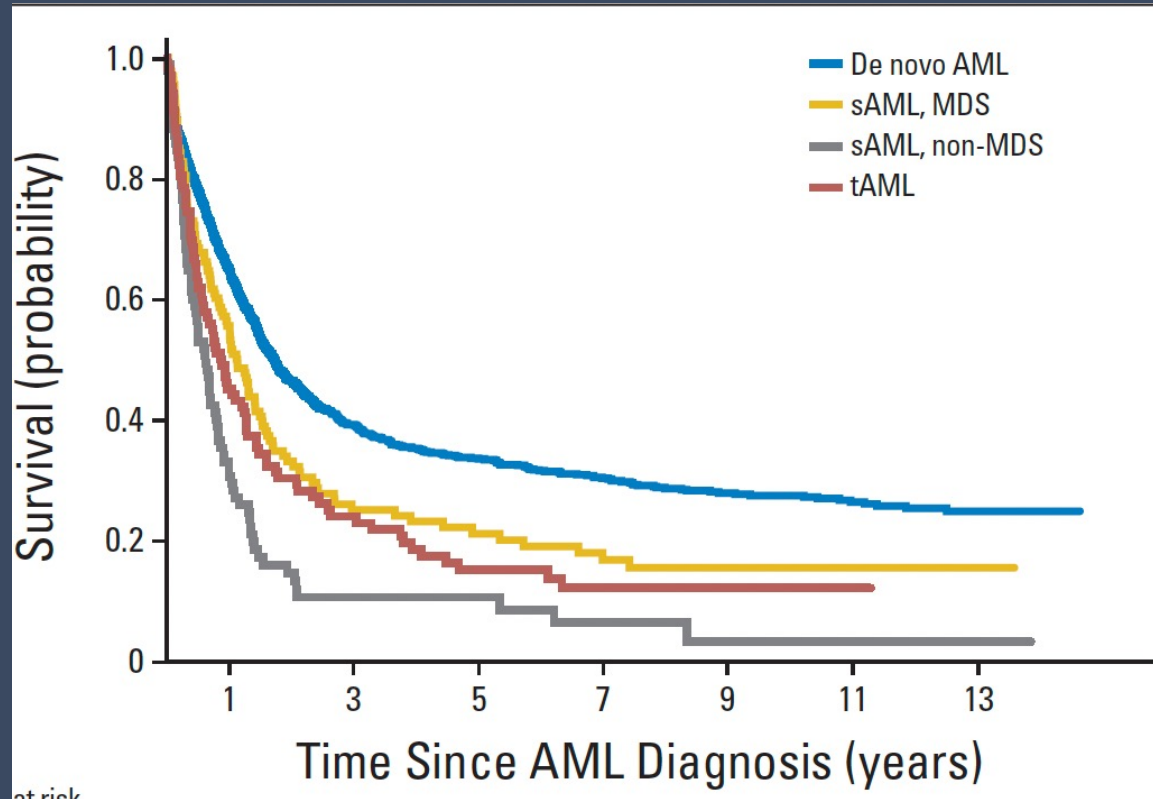
Adapted from Visser *et al.* 2012

Older patients can be challenging to treat³

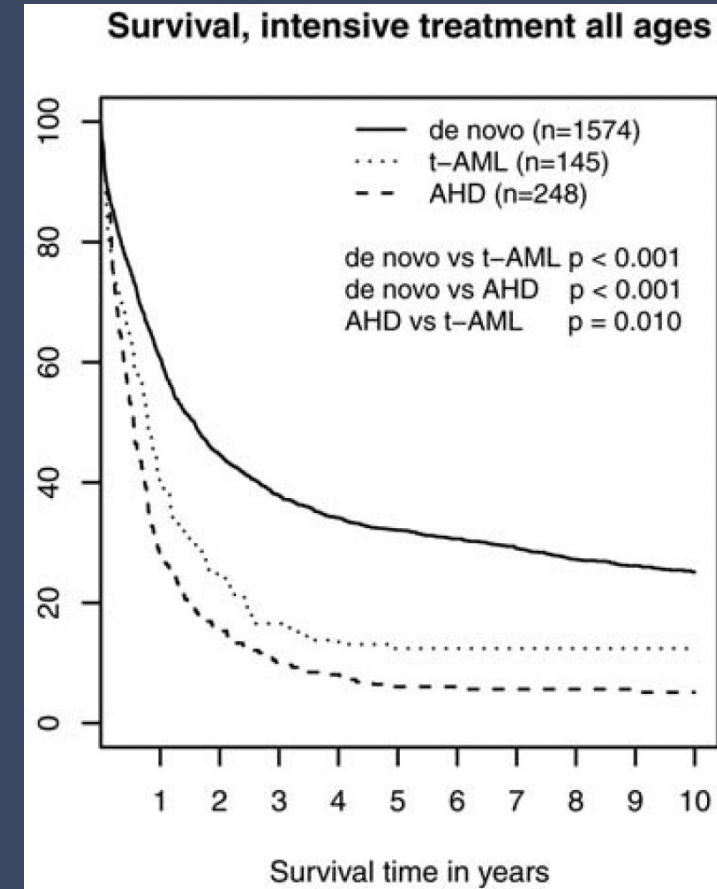
- Significant comorbidities
- Increased likelihood of prior treatment for malignancies or haematological disorders
- Complex genetic and molecular leukaemia biology

NOT ALL THE AML ARE THE SAME

Granfeldt Østgård et al – Sweden group

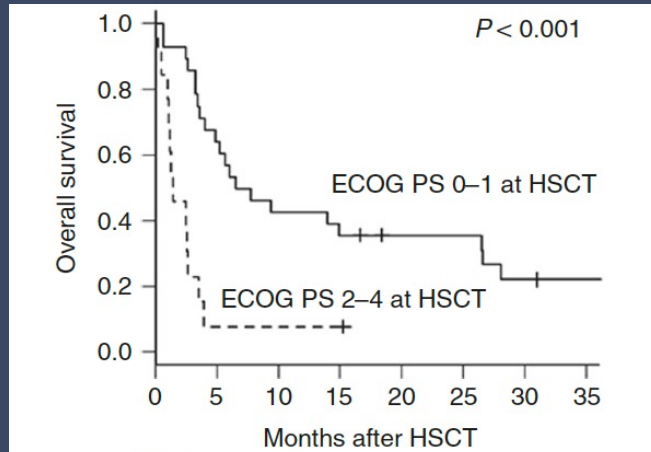


Erik Hulegårdh et al - Danish group

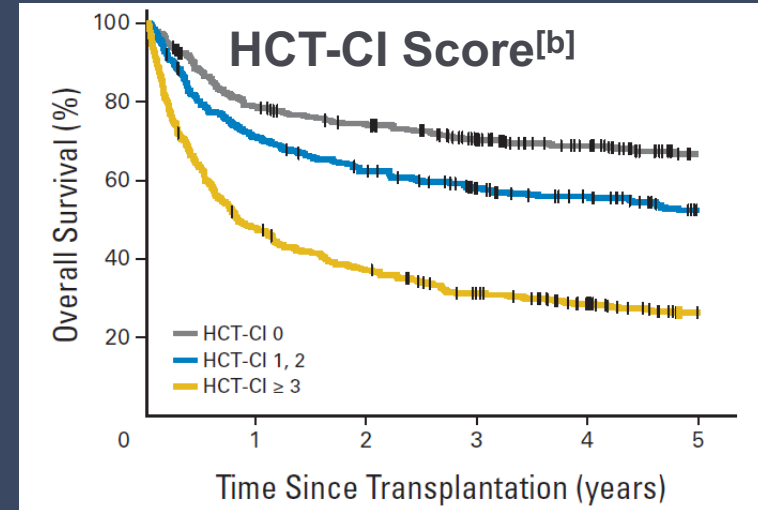


Eligibility for Allo-SCT in Older Adult Patients Is Challenging

Performance Status^[a]

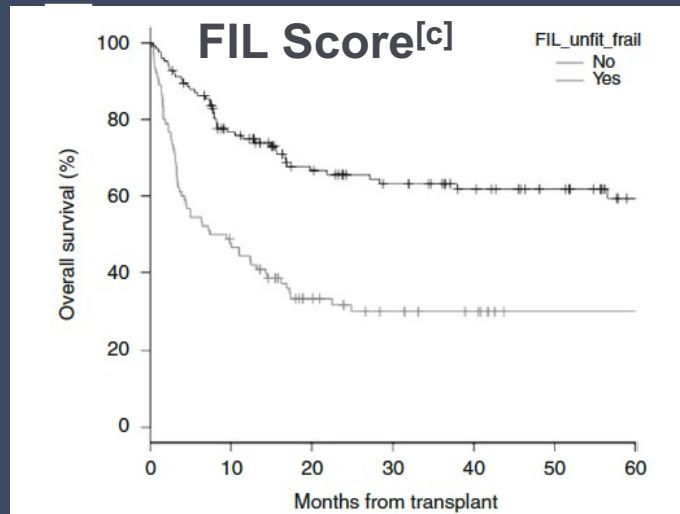


✓ Simple
X Too limited
X Functional status



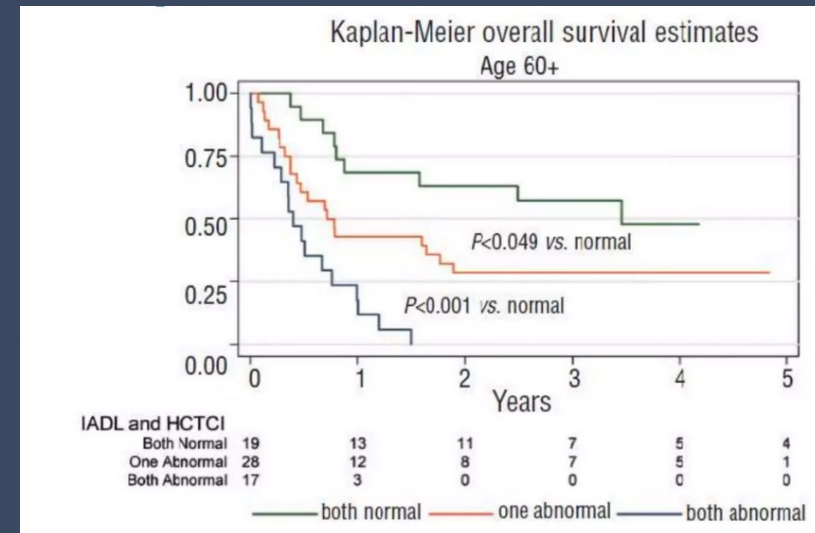
✓ Comorbidity
✓ Choice conditioning regimen
X Frailty

FIL Score^[c]



Frailty
Comorbidity

Geriatric Assessment^[d]



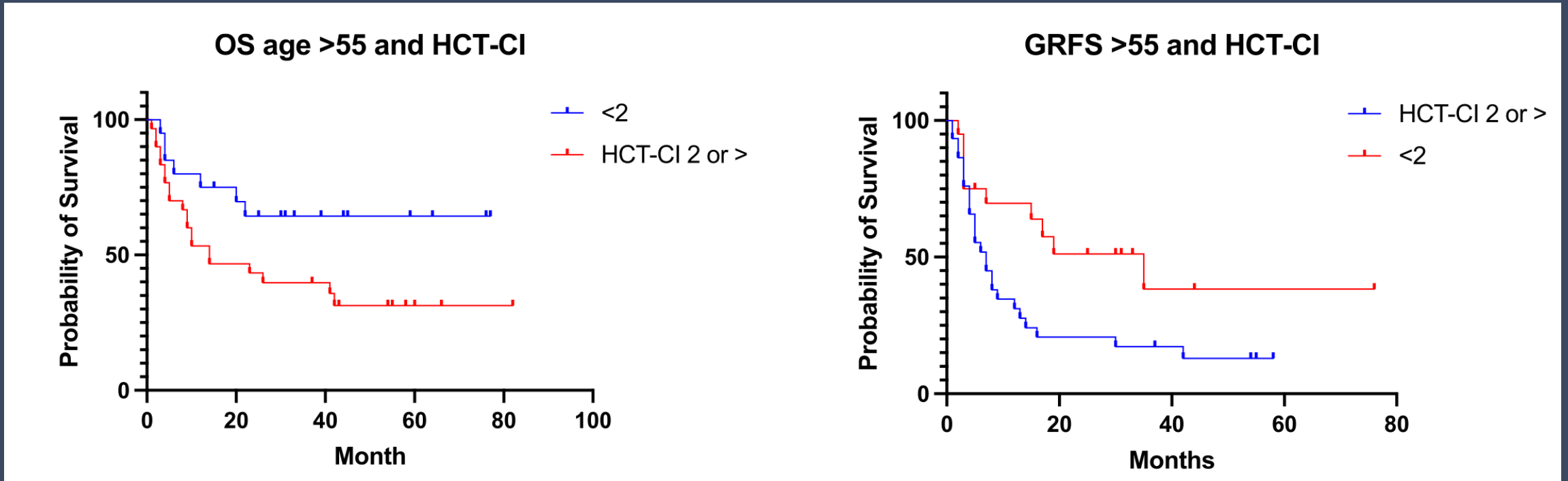
✓ Simple
✓ Frailty
X Comorbidity

FIL, Fondazione Italiana Linfomi; HSCT, hematopoietic stem cell transplant; IADL, instrumental activities of daily living.

a. Muffy LS, et al. Haematologica. 2014;99:1373-1379; b. Sorror ML, et al. J Clin Oncol. 2014;32:3249-3256; c. Poverelli N, et al. Bone Marrow Transplant. 2020;55:2224-2233; d. Tomori S, et al. Bone Marrow Transplant. 2020;55:233-241.

A deeper T-Cell depletion is the right answer?

103 AML and MDS-EB after FB4-CAMPATH



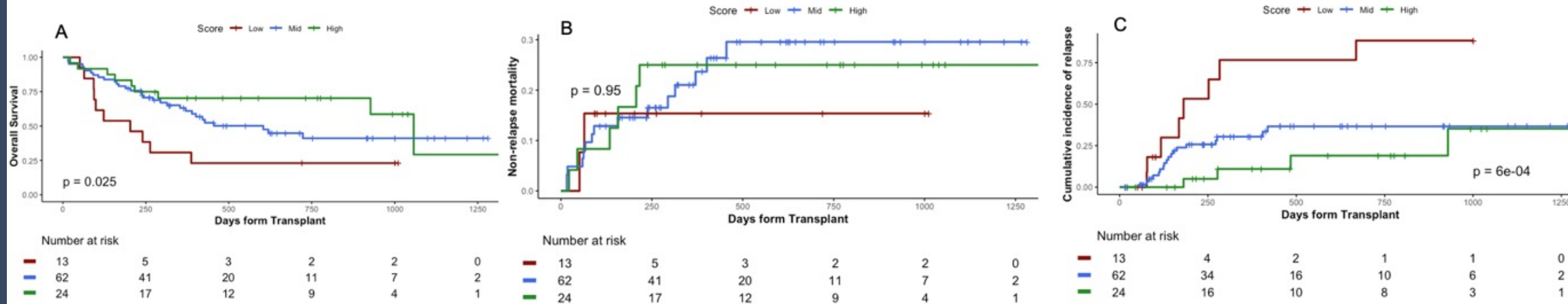
Moderate and severe chronic GVHD are only 16% and 4%, respectively.

Relapse and TRM 30% and 26%

Causes of TRM: Infections (65% of total TRM), GVHD (5%), CNS bleeding (5%), and stroke (5%).

Development of a Simplified Geriatric Score-4 (SGS-4) to Predict Outcomes After Allogeneic Hematopoietic Stem Cell Transplantation in Patients Aged over 50

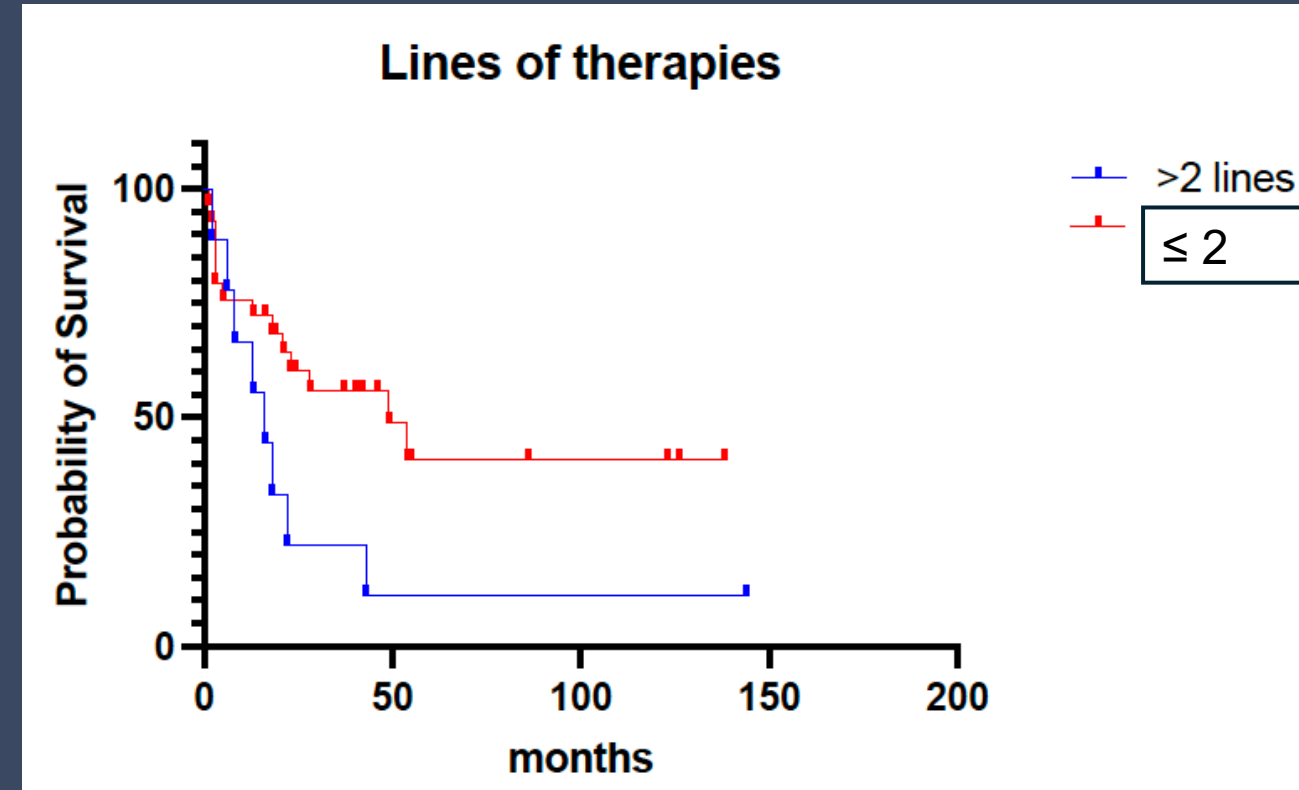
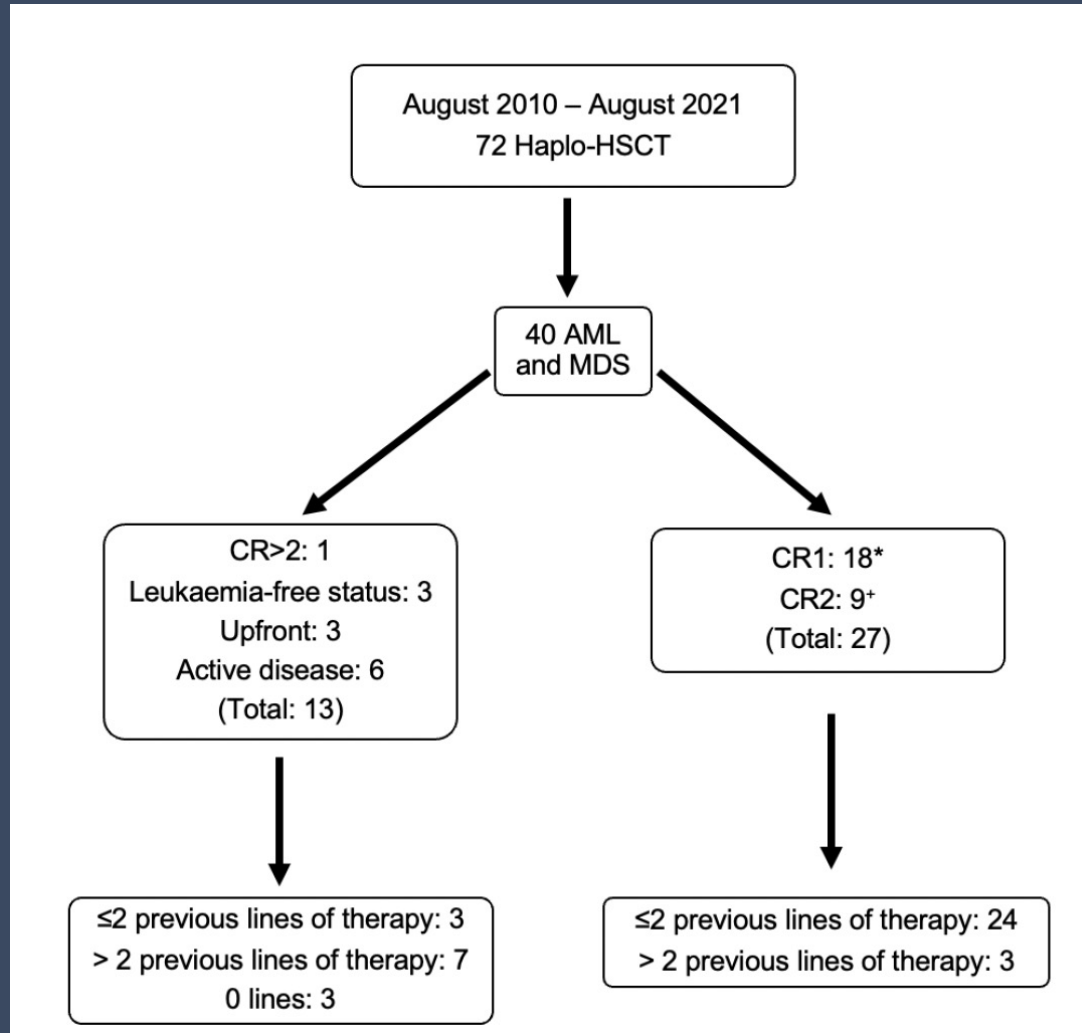
Eugenia Accorsi Buttini ^{1,2,*} , Alberto Zucchelli ^{1,3}, Paolo Tura ¹, Gianluca Bianco ¹ , Daniele Avenoso ^{1,4} , Giovanni Campisi ⁵, Mirko Farina ^{1,4} , Gabriele Magliano ⁴, Enrico Morello ⁴ , Vera Radici ^{1,4} , Nicola Polverelli ⁶ , Domenico Russo ^{1,4}, Alessandra Marengoni ^{1,3,†}  and Michele Malagola ^{1,4,†} 



Outcome	Low (%)	Intermediate (%)	High (%)	p-value
OS @ 1 yr	32.7	62.7	70.3	0.025
NRM @ 1 yr	16.1	20.6	25.0	0.95
CIR @ 1 yr	69.2	30.4	10.9	<0.01

- **Score Components**
- Developed via Factor Analysis: **Gait speed, Hand grip, G8, Sex**
- Score ranges stratified patients into:
 - **Low fitness (≤ 13):** n=13
 - **Intermediate fitness ($>13-22.5$):** n=62
 - **High fitness (>22.5):** n=24

If no HLA-identical donor available, go for HAPLO. The sooner, the better.



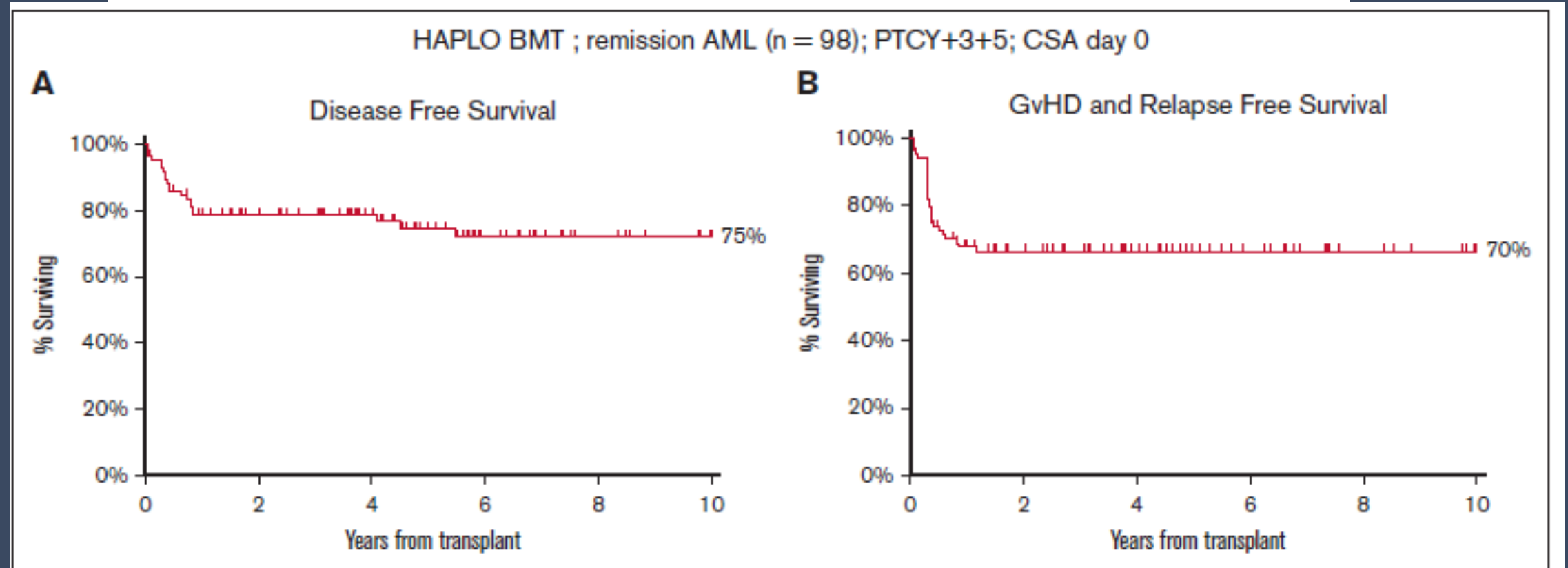
OS / PFS: 1- and 3-year OS were 62% and 43% (median OS 22 months). Patients with ≤ 2 prior therapy lines had superior OS (3-year: 55% vs 22%, $p=0.04$). One- and 3-year PFS were 57% and 46%.

CIR: Overall relapse incidence was 25% (median time to relapse 5 months). Patients with >2 prior therapy lines had higher 3-year CIR than those with ≤ 2 (49% vs 17%, $p=0.05$).

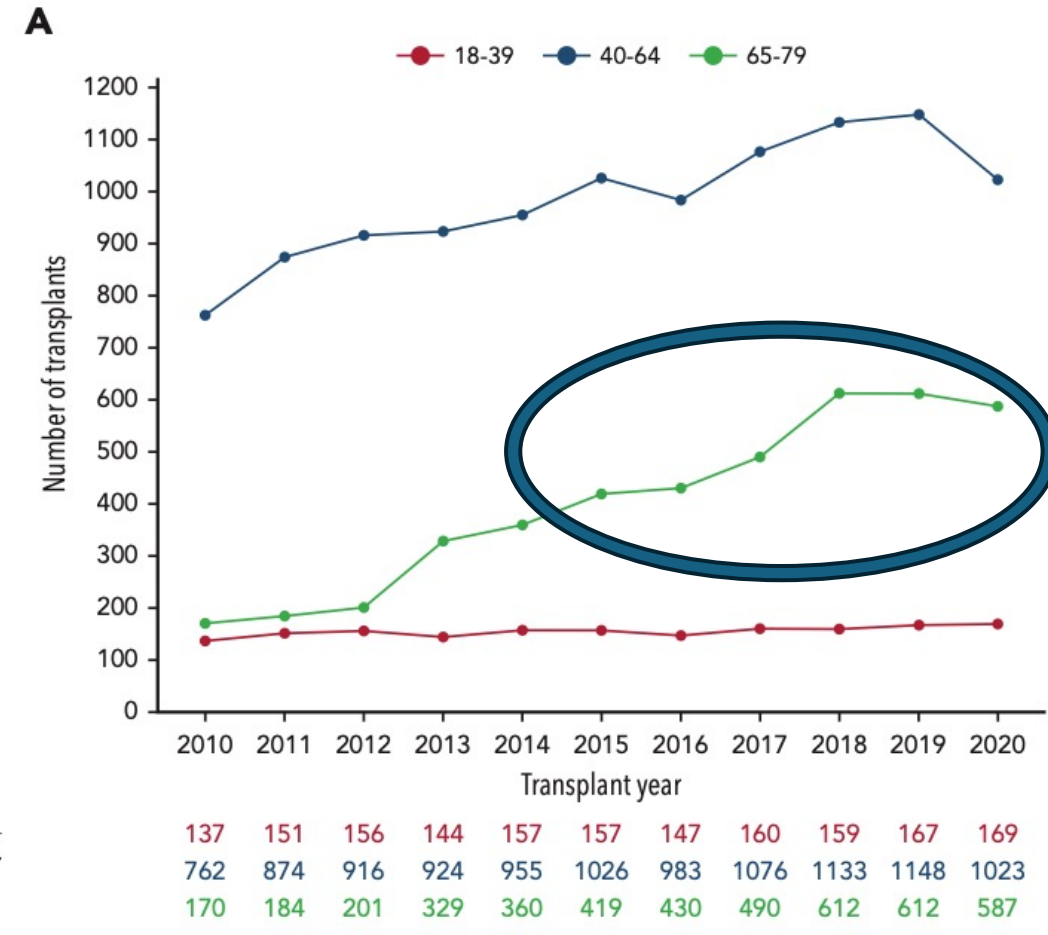
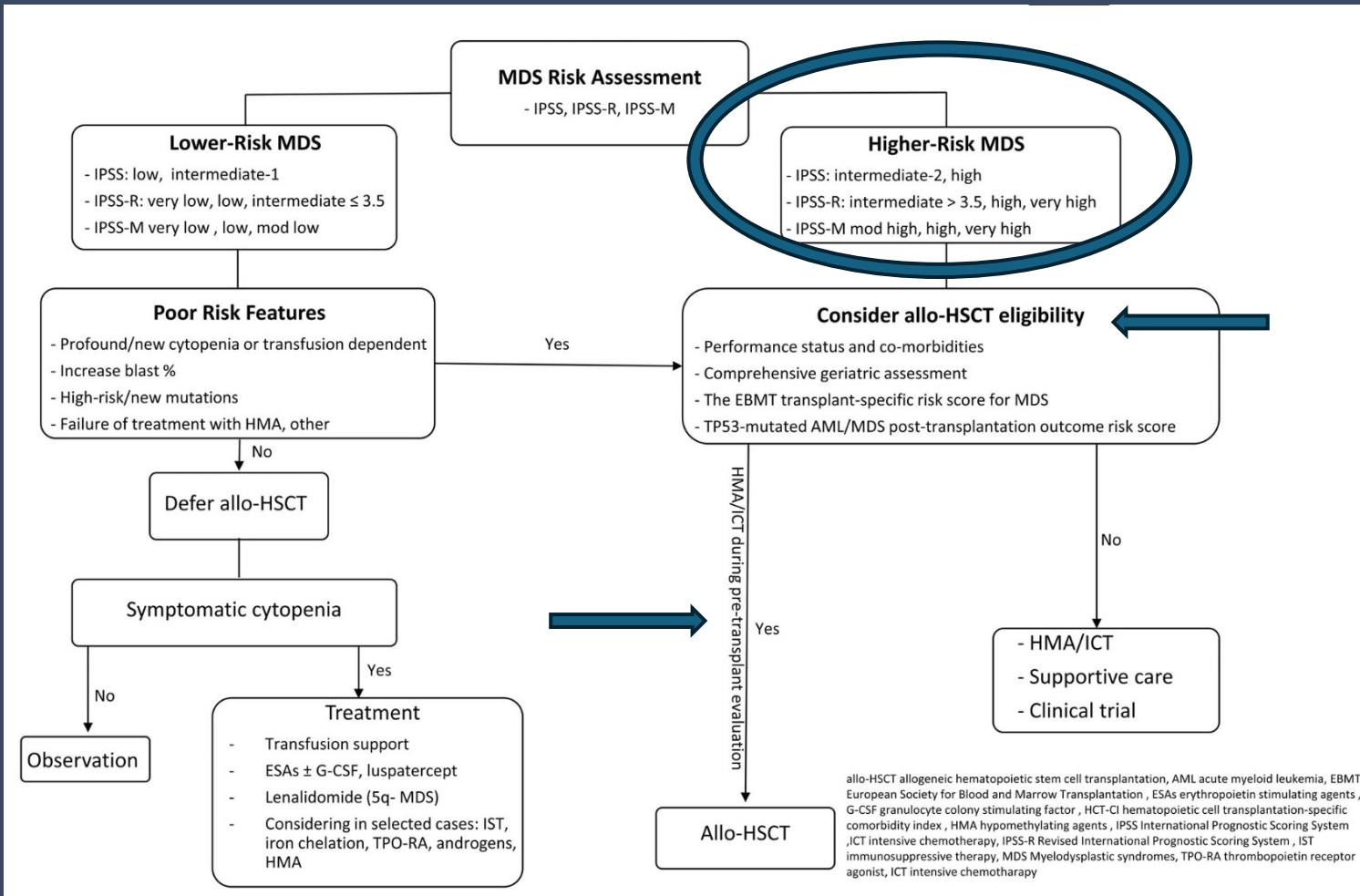
Haploidentical bone marrow transplantation for AML in remission after TBF conditioning: a long-term follow-up

Anna M. Raiola,¹ Carmen Di Grazia,¹ Alida Dominiotto,¹ Stefania Bregante,¹ Sabrina Giammarco,² Riccardo Varaldo,¹ Federica Sorà,^{2,3} Elisabetta Metafuni,² Maria A. Limongiello,² Antonella Laudisi,¹ Monica Passannante,¹ Eugenio Galli,² Massimiliano Gambella,¹ Simona Sica,^{2,3} Andrea Bacigalupo,^{2,3} Emanuele Angelucci,¹ and Patrizia Chiusolo^{2,3}

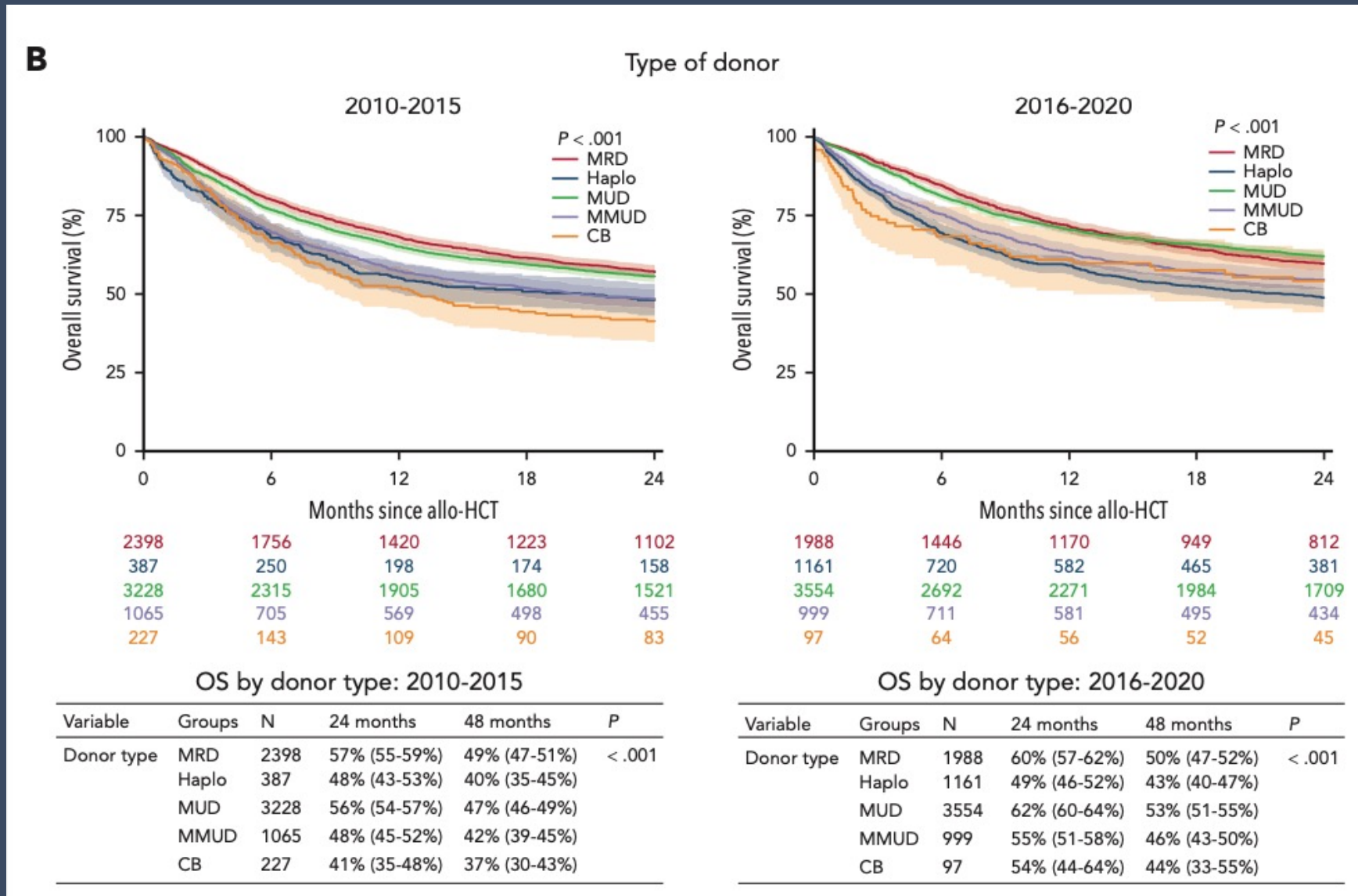
¹Divisione Ematologia e Terapia Cellulare, IRCCS Policlinico San Martino IST, Genova, Italy; ²Dipartimento di Diagnostica per Immagini, Radioterapia Oncologica ed Ematologia, Fondazione Policlinico Universitario A. Gemelli IRCCS, Rome, Italy; and ³Sezione di Ematologia, Dipartimento di Scienze Radiologiche ed Ematologiche, Università Cattolica del Sacro Cuore, Rome, Italy



MDS state of art



Allo-HSCT as a curative strategy



**ACUTE
LEUKEMIA**



**3+7? VENETOX?
RIC MAC PTCY
TRANSPLANT
FEASIBLE
REMISSION
TOTAL CARE**

The optimal AML/MDS patient journey: Highly effective and Less Toxic and Tailored.

Diseases (High/Int)

AML > MDS > MPN > ALL > NHL

Age

> 50yy – 75yy

MRD

CR1 mrd-/mrd+ > CR2

Fitness

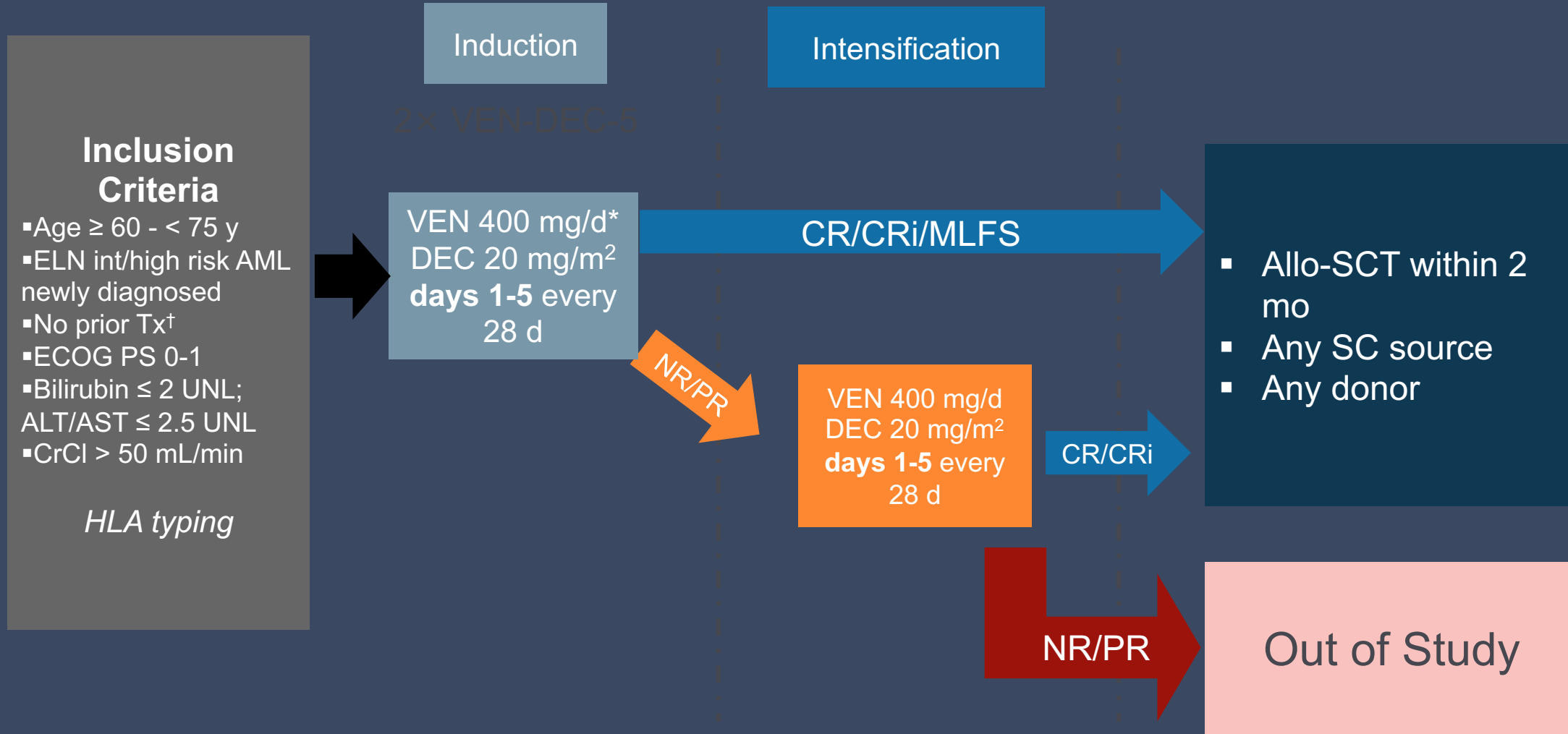
FIT > Frail

Pre-Transplant Treat-Prog
With CHT (3+7 or similar)

Highly effective and Less Toxic
no more than 15% of PTS get a Transplant



Phase 2 VEN-DEC GITMO Trial as “Bridge” to Allo-SCT *Study Design*



*VEN ramp up: 100 mg on day 1, 200 mg on day 2, and 400 mg from day 3 (only for first cycle). †HU is allowed in case of WBC count $> 25,000$.

ClinicalTrials.gov. Accessed March 28, 2023. <https://clinicaltrials.gov/study/NCT04476199>.

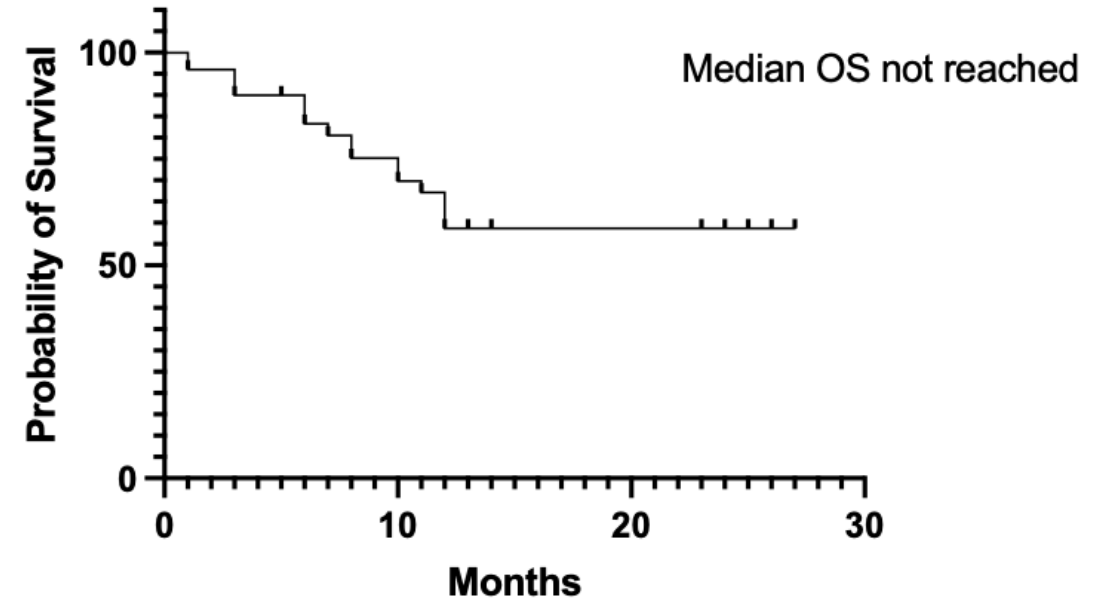
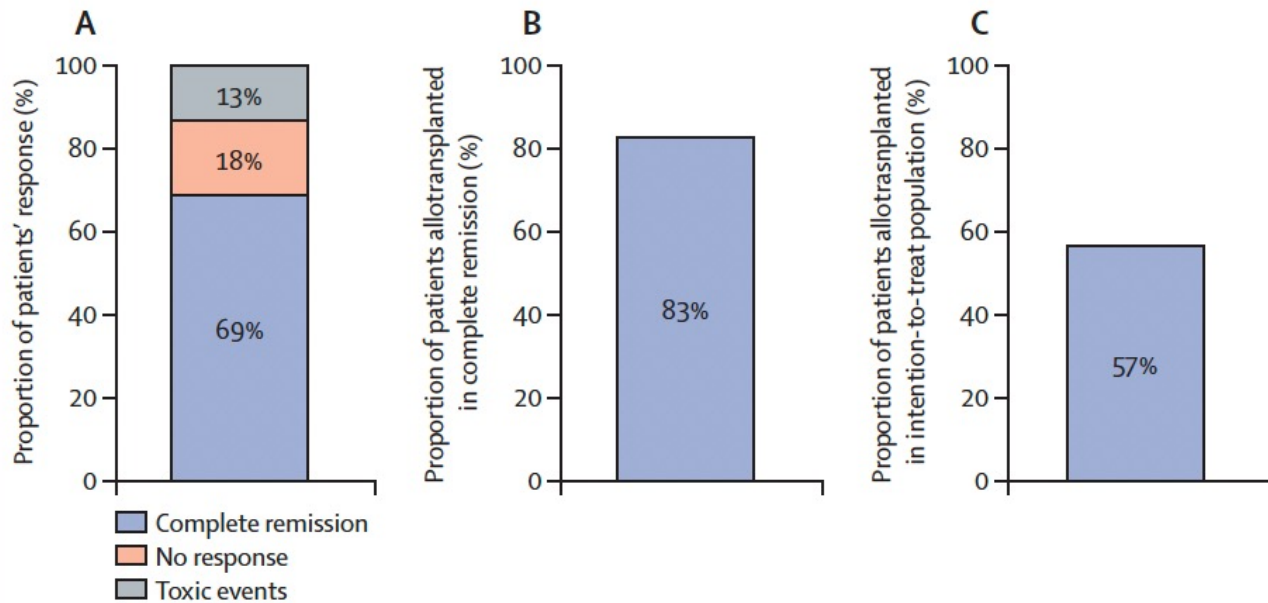


Transplant Feasible Remission (TFR) in AML

65/93 (=69%) achieved CR

53/65 CR patients (=83%)
underwent allo-SCT

53/93 treated patients (=57%)
underwent allo-SCT



REFRACTORY ACUTE MYELOID LEUKEMIA



Hypothesis: salvage chemotherapy prior to alloHCT would not provide a net benefit.

RIST

Salvage CT (HAM):
AraC 2 x 3 g/m² d 1-3¹
Mitox 10 mg/m² d 3-5



R

281 patients, rando 1:1

DisC

watchful waiting
or
low-dose AraC
or
Mitoxantron 10 mg/m²



Conditioning Intensity
tailored to amount of
residual AML

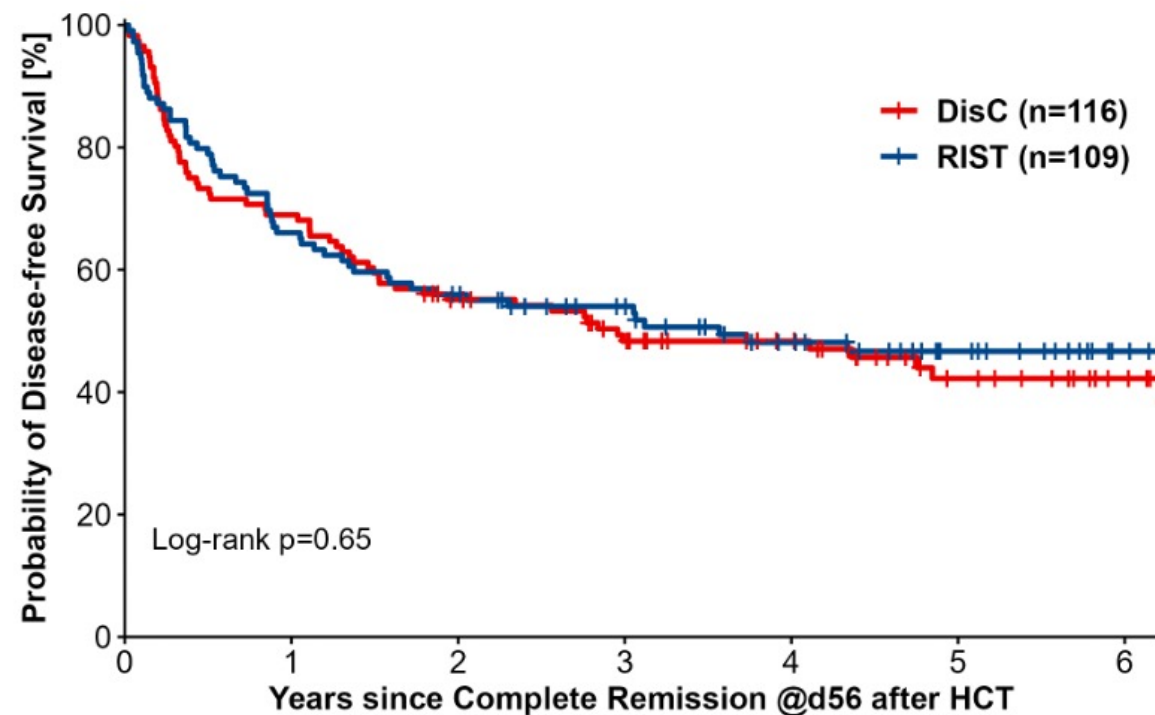


Primary Endpoint:
Treatment success
OS from Rando

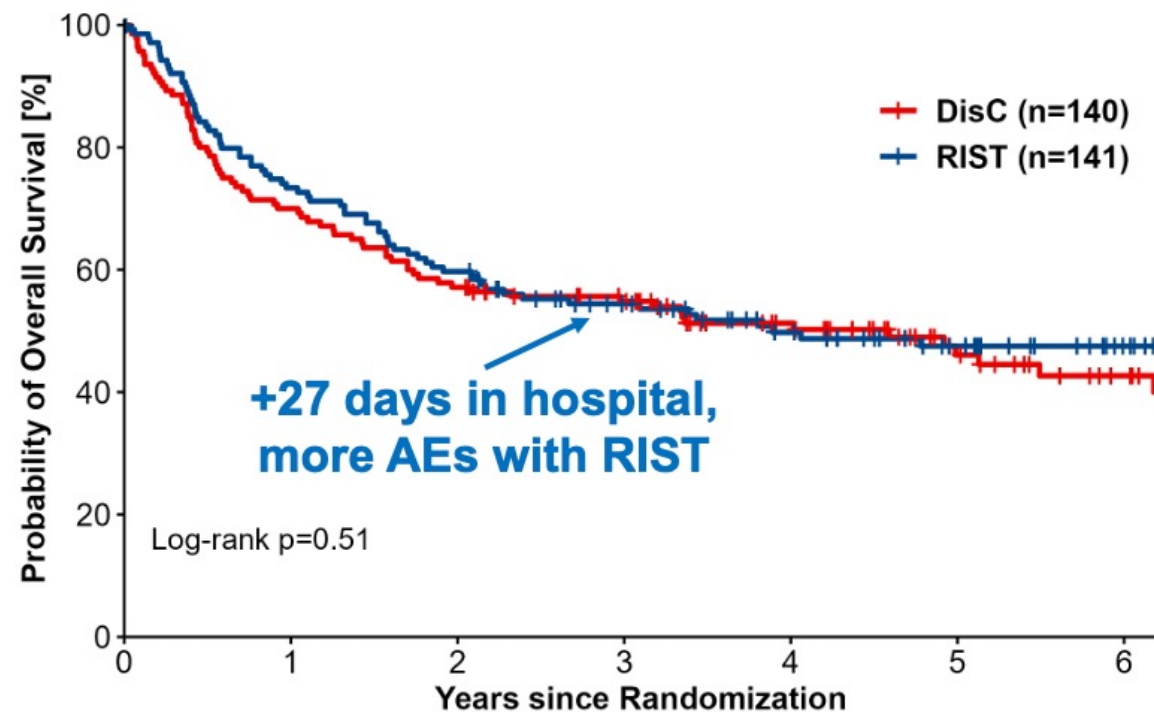
Sequential conditioning
FLAMSA-RIC or
MeI/Flu/TBI

No difference in DFS! No difference in OS from randomization!

Disease-free Survival from d56



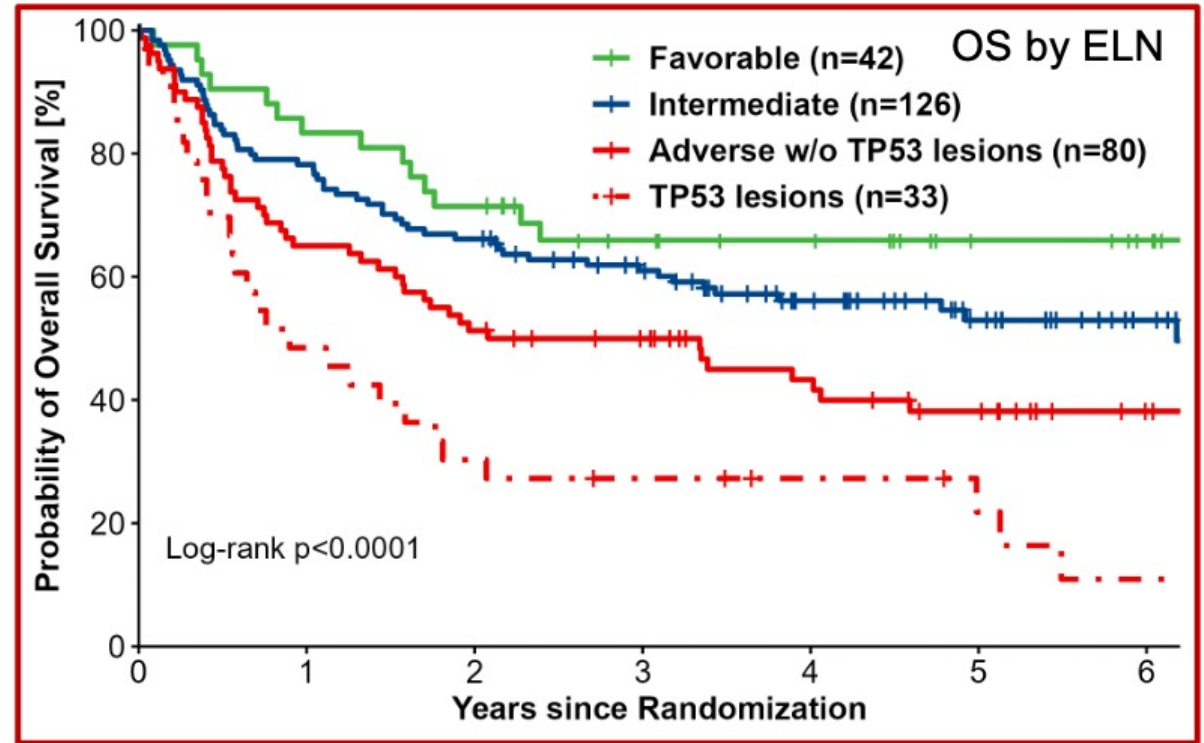
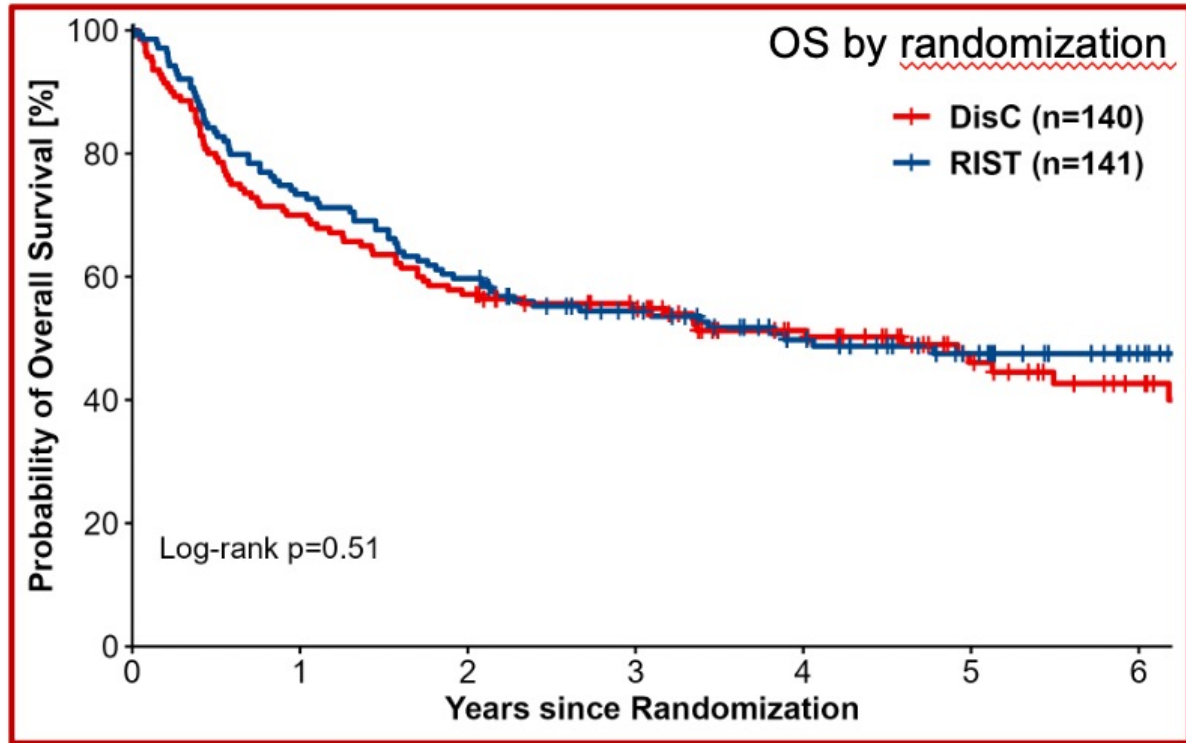
Overall Survival from randomization



Median follow-up from randomization: 60 months

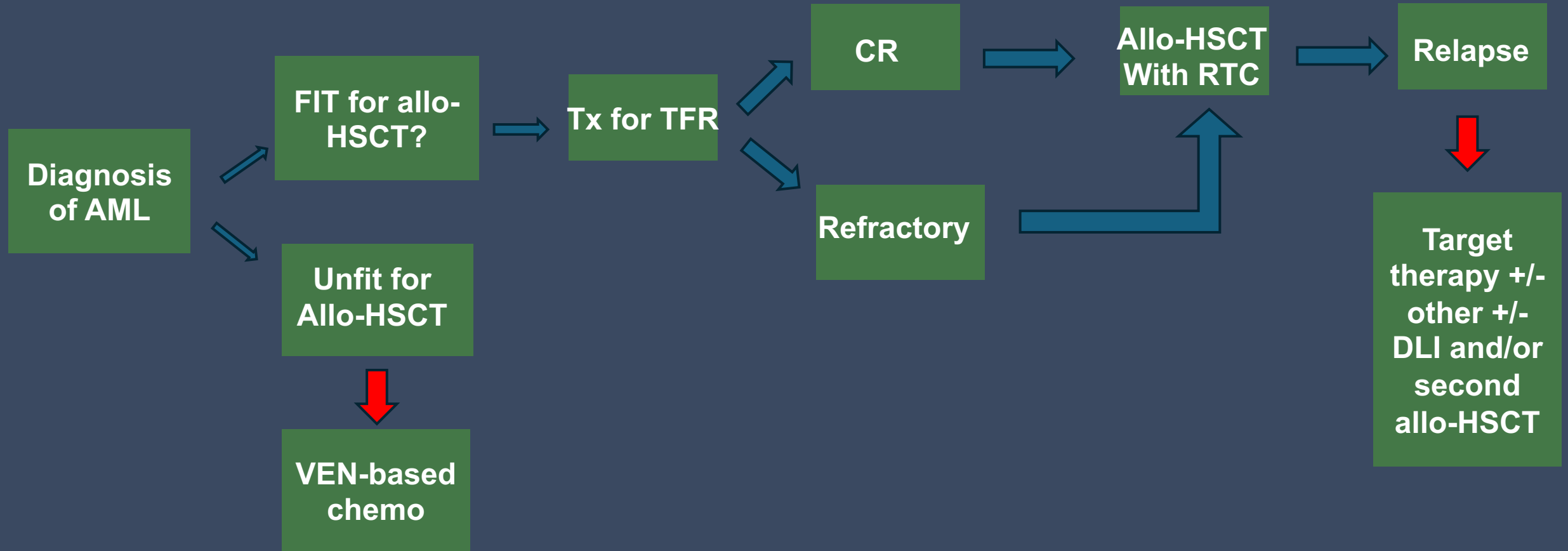


Summary: AML biology matters, but not: depth of remission



- Long-term follow-up of ASAP trial shows no advantage of remission induction prior to alloHCT
- Sequential conditioning for patients with AML not in remission needs to be improved in frame-work of RCT
- Especially for adverse risk AML bridging, conditioning, and maintenance after HCT need to be improved.

The AML pathway in 2030?



Conclusions and open questions

- Allo-HSCT remains a central curative strategy in AML, with growing use in older patients and in high-risk disease, supported by refined risk stratification (ELN 2022 AdvP) and by the emerging concept of Transplant Feasible Remission within a comprehensive treatment journey.
- In older and high-risk AML, venetoclax-based therapies (e.g. VEN-DEC GITMO) enable a higher proportion of patients to reach transplant and may redefine the concept of remission as “transplant-oriented” rather than purely morphological.
- PTCY platforms, haploidentical donors, and reduced-intensity conditioning have expanded transplant eligibility, but relapse and non-relapse mortality remain major limitations, especially in AdvP and heavily pretreated patients.

Open questions: Can TFR become a standardized endpoint in AML? How should we integrate MRD, fitness scores and biological risk to personalize conditioning, donor choice and post-transplant strategies (maintenance, DLI, second allo-HSCT) in the 2030 AML pathway?