

# LEUKEMIA2022

Rome, Hotel NH Collection - Vittorio Veneto

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AIL President: P. Toro

Coordinators: A.M. Carella, S. Amadori



## MRD Driven Strategy in Adult ALL

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UNDER THE AUSPICES OF:



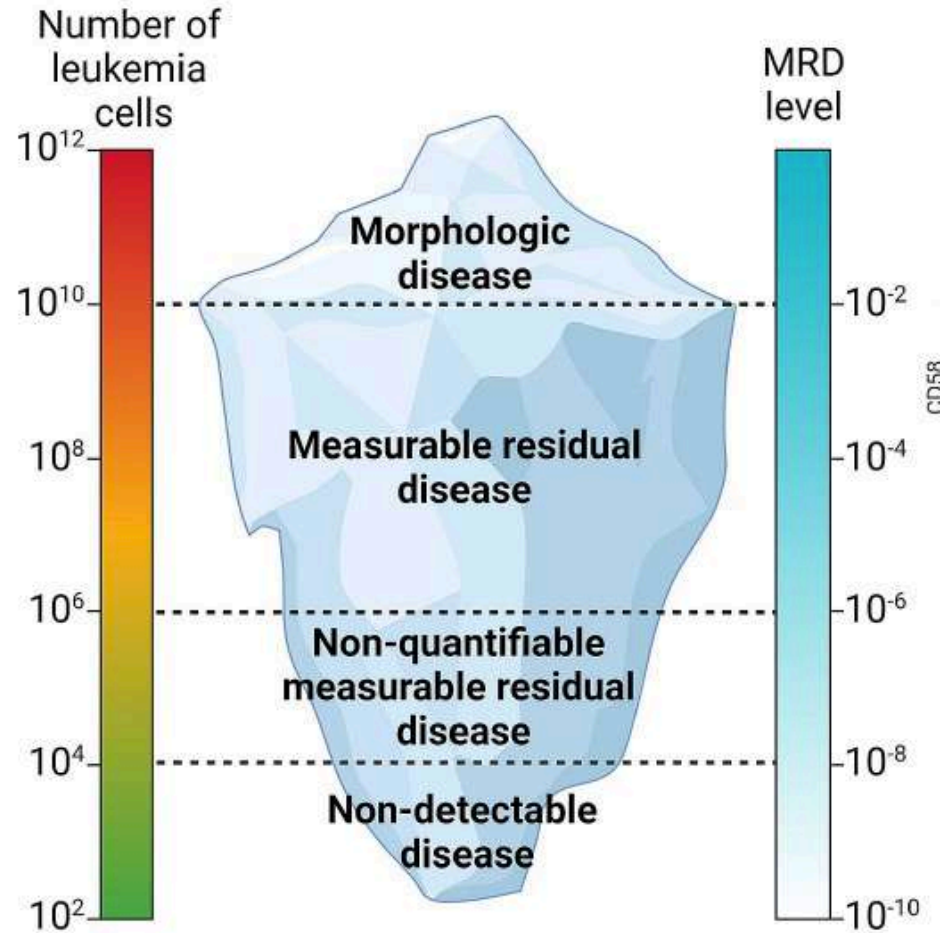
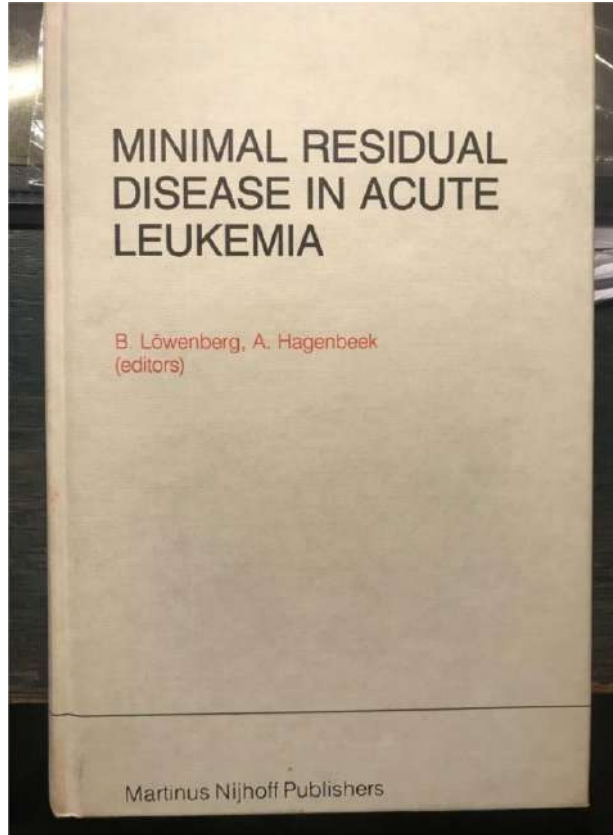
SIE - Società Italiana di Ematologia

## COI disclosures: Renato Bassan

- Advisory boards: Amgen, Novartis, Kite Pharma/Gilead
- Travel grants/honoraria/symposia: Amgen, Incyte, Servier, Jazz Pharmaceuticals, Pfizer

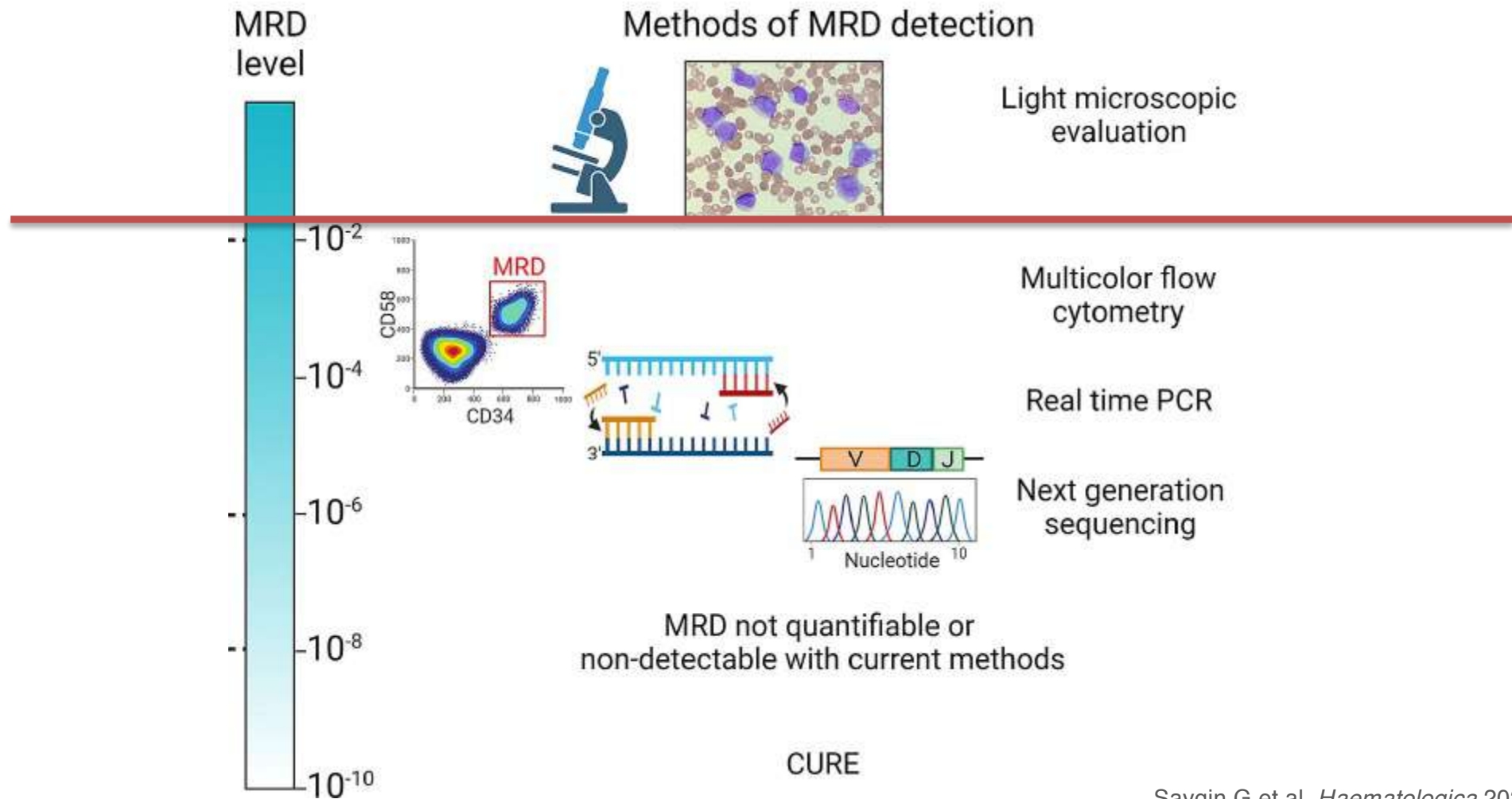
# Minimal or measurable ?

can't be  
seen



Saygin G et al, *Haematologica* 2022

# How to measure the minimal



- **Patients without MRD study**
  - *No or poor BM cell sampling*
  - *Technical failure/no IG/TCR molecular probe*  
*(less concerns with gene rearrangements and MFC)*

STUDY (N)	CR (%)	Key MRD timepoint(s)	MRD evaluable
NILG 09/00 (N 304) <sup>1</sup>	258 (85)	wk 16-22	77.5 %
NILG 10/07 (N 163) <sup>1</sup>	142 (87)	wk 10-22	77 %
GMALL 07/03 (N 2061) <sup>2</sup>	1857 (90)	wk 16	53 % (2003-2009) 69 % (2010-2016)

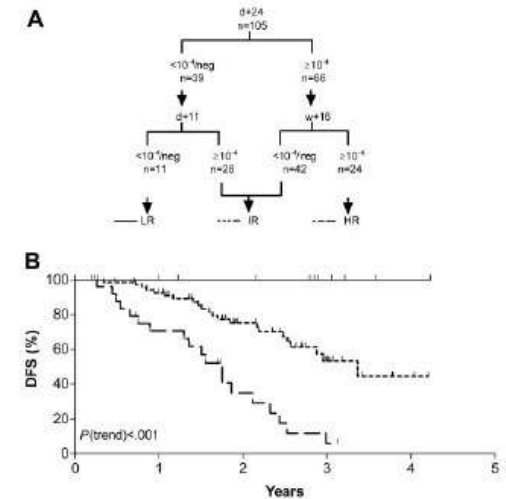
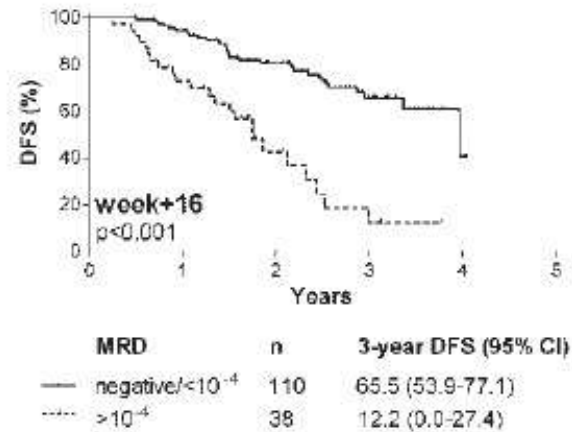
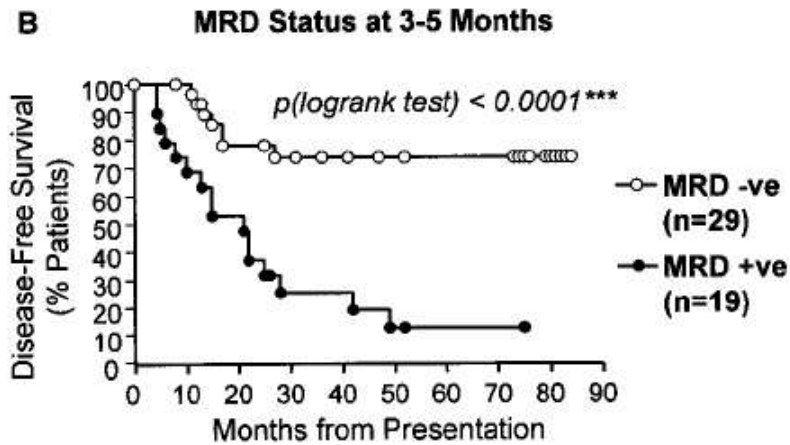
- **Maximum sensitivity  $10^{-5}$  (likely to improve)**
  - *25% MRD negative patients will relapse*

***SUGGESTING, «true» MRD negativity lower than commonly reported***

<sup>1</sup>Bassan R et al, *Clin Lymph Myeloma Leuk* 2017; <sup>2</sup>Goekbuget N et al, *Blood* 2017 [abstr]

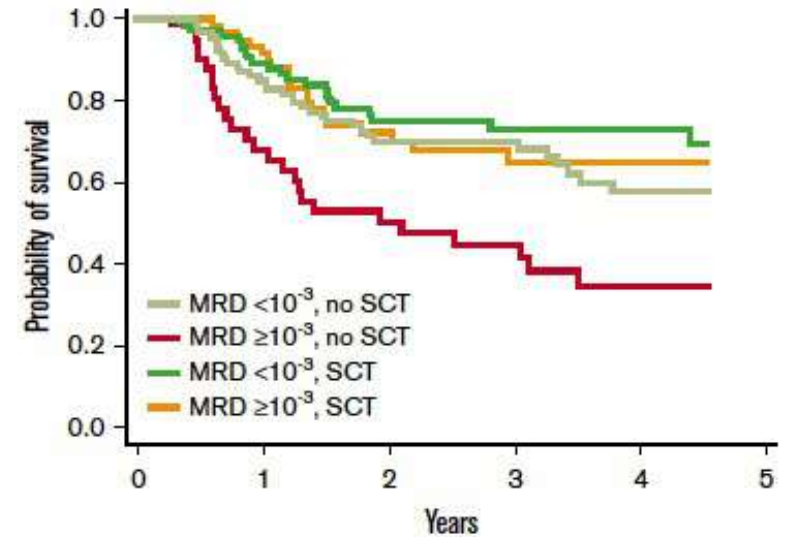
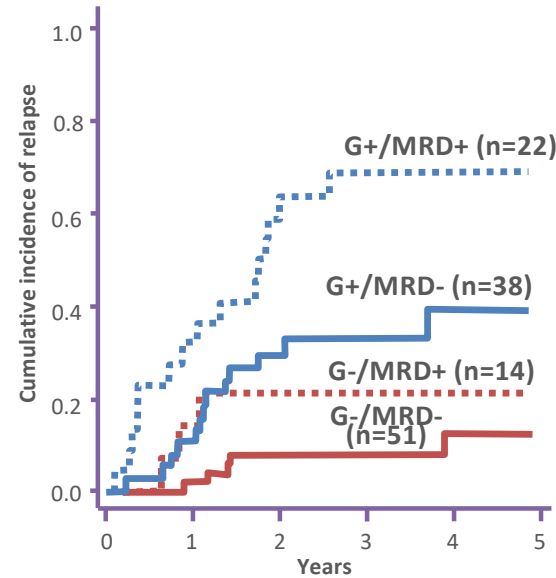
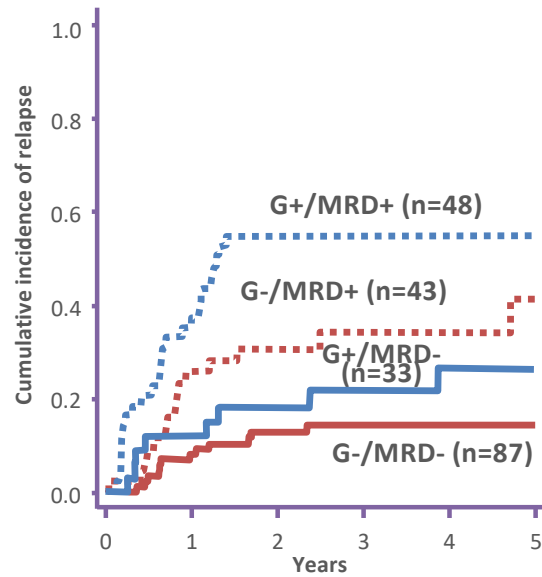
## UKALL XII B-ALL patients (cumulative chemo/SCT)

## GMALL 05-06 Standard risk B/T-ALL patients (chemo – 1-year maintenance)



High relapse rate with 'limited' chemo

## GRAALL 2003-2005



**Four gene prognostic classifier (adverse, G+) and MRD**

B-ALL: *MLL* rearrangement and/or *IKZF1* deletion

T-ALL: Unmutated *NOTCH1/FBXW7* and/or *RAS/PTEN* abnormalities

**Allogeneic SCT vs. MRD in HR ALL**

**Risk of very early relapse in allo-SCT eligible patients**

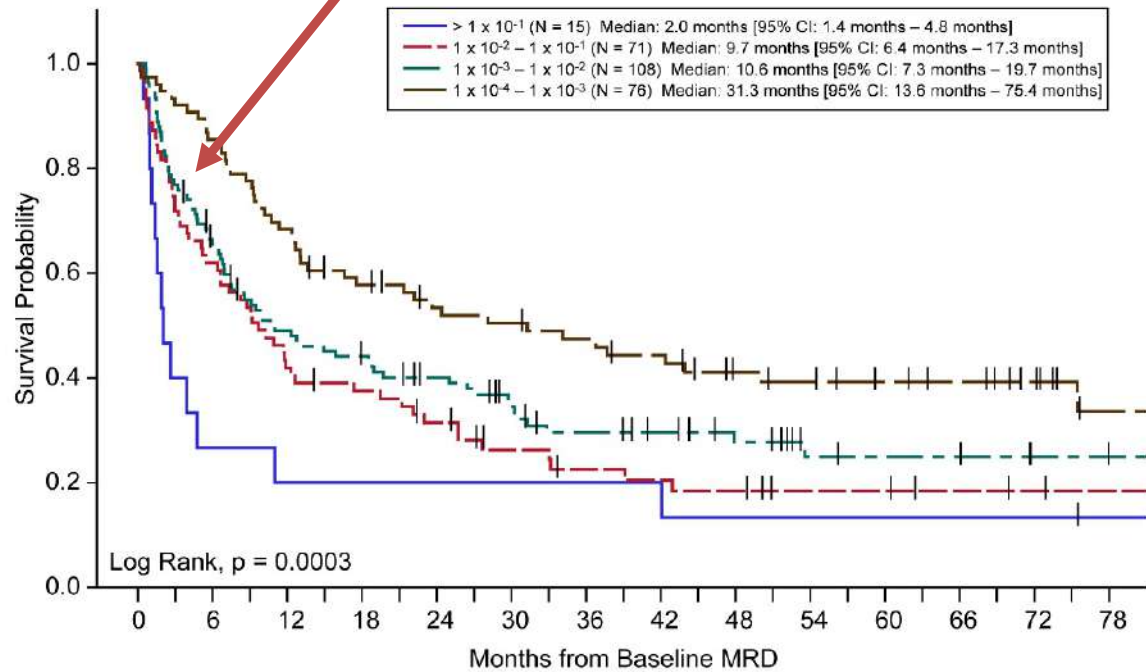
HEMATOLOGY  
 2019, VOL. 24, NO. 1, 337-348  
<https://doi.org/10.1080/16078454.2019.1567654>

Taylor & Francis  
 Taylor & Francis Group

OPEN ACCESS Check for updates

## Minimal residual disease level predicts outcome in adults with Ph-negative B-precursor acute lymphoblastic leukemia

Nicola Gökbüget<sup>a</sup>, Hervé Dombret<sup>b</sup>, Sebastian Giebel<sup>c</sup>, Monika Brüggemann<sup>d</sup>, Michael Doubek<sup>e</sup>, Robin Foà<sup>f</sup>, Dieter Hoelzer<sup>g</sup>, Christopher Kim<sup>g</sup>, Giovanni Martinelli<sup>h</sup>, Elena Parovichnikova<sup>i</sup>, Alessandro Rambaldi<sup>j</sup>, Josep-Maria Ribera<sup>k</sup>, Marieke Schoonen<sup>l</sup>, Julia M. Stiglmair<sup>m</sup>, Gerhard Zugmaier<sup>m</sup> and Renato Bassan<sup>n</sup>

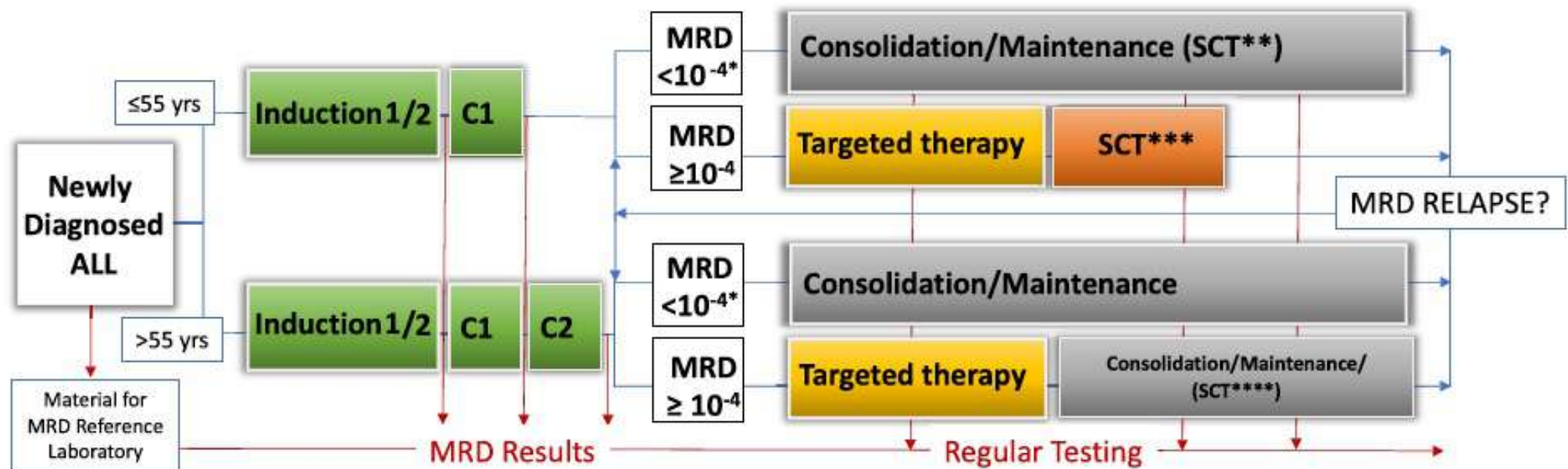


Number of Subjects at Risk:

1:	15	4	3	3	3	3	3	2	2	2	2	2	1
2:	71	44	29	25	20	14	11	10	9	6	6	4	3
3:	108	69	50	44	37	29	23	20	15	9	8	7	5
4:	76	65	52	42	36	34	31	28	22	20	17	15	11

Censor indicated by vertical bar |.





- \* In case of low-positive MRD more frequent controls
- \*\* SCT independent of MRD in defined high-risk patients; compatible donor; age adapted conditioning
- \*\*\* compatible donor; age adapted conditioning
- \*\*\*\* SCT in selected points depending on donor availability and general condition

Figure 1. Flow of MRD surveillance and treatment decisions (GMALL strategy).

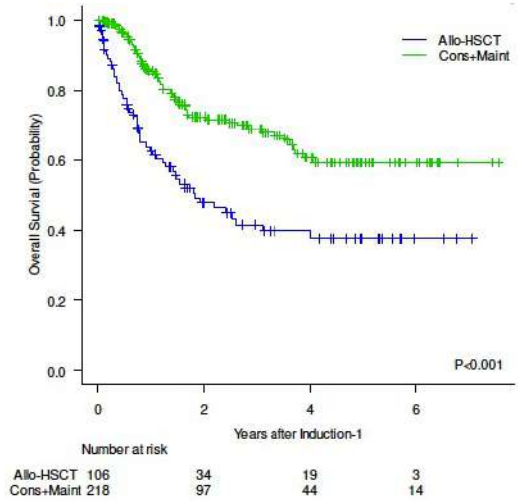
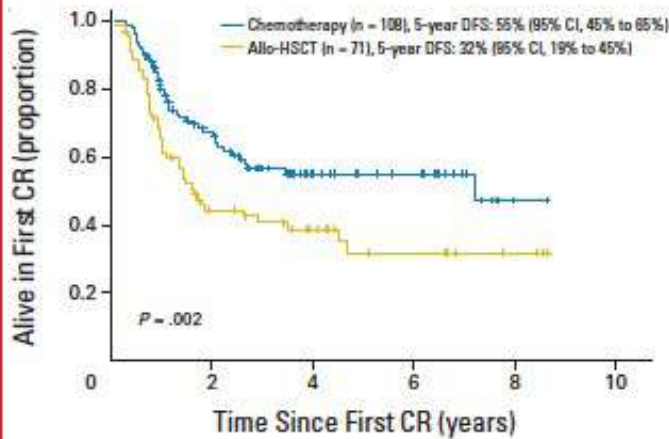
## PETHEMA ALL-AR-03 and HR-11

HR patients

RISK ORIENTED:

d14 BM blasts and MRD

MRD

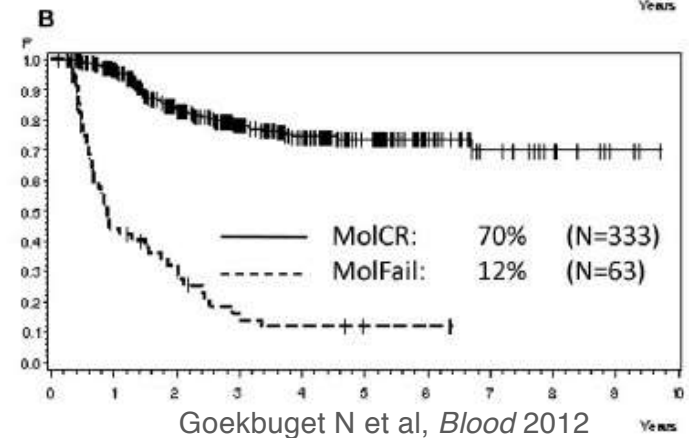
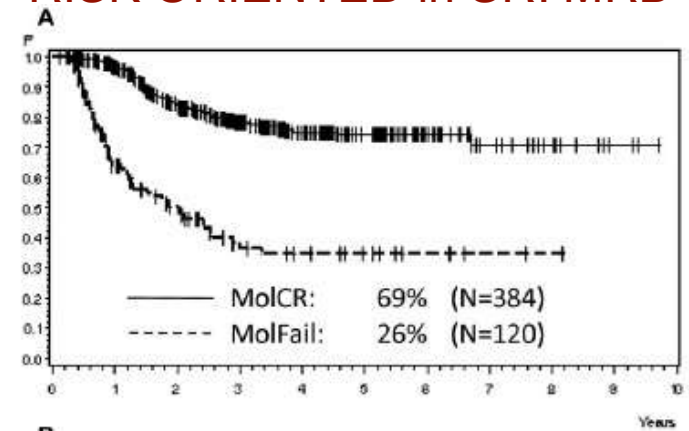


Ribera JM et al, *J Clin Oncol* 2014 and *Blood* 2020

## GMALL 06-07

SR B/T-ALL patients (HR to allo-SCT)

RISK ORIENTED in SR: MRD



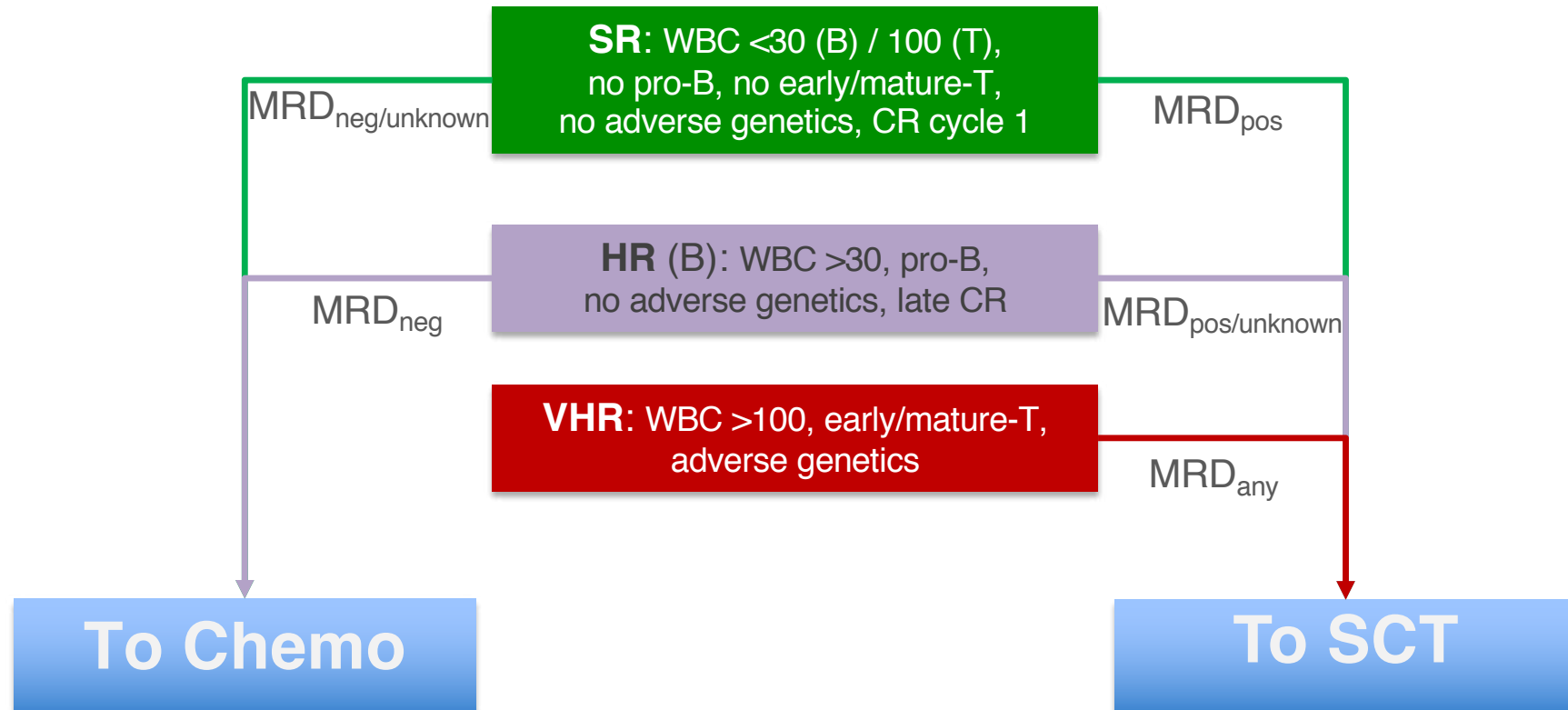
Goekbuget N et al, *Blood* 2012

# NILG and GIMEMA strategy



**MRD<sub>neg</sub>** < 10<sup>-4</sup> @ w10-16, negative @ w22

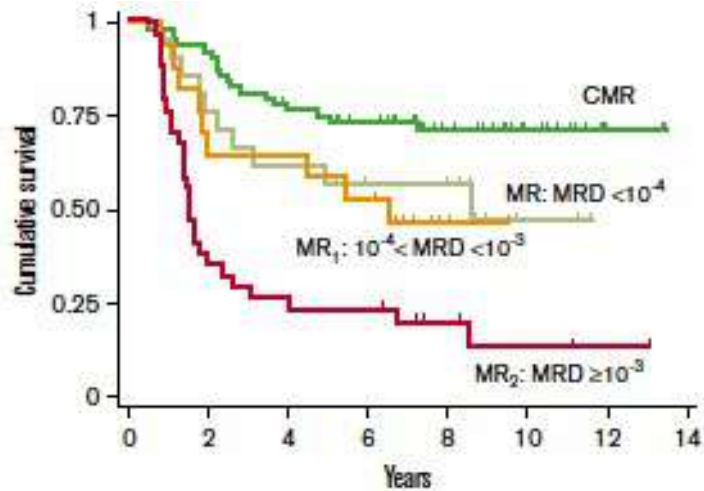
**MRD<sub>pos</sub>** ≥ 10<sup>-4</sup> @ w10-16, positive @ w22



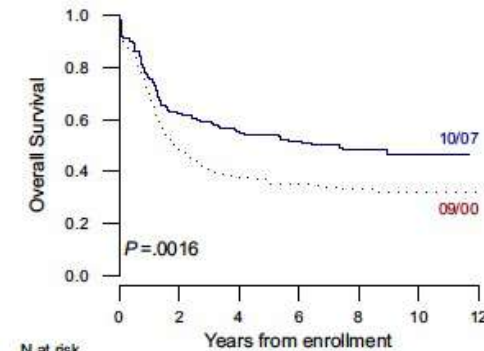
## NILG 09/00 and 10/07; GIMEMA LAL 1913 Age 18-65, SR and HR patients (VHR to allo-SCT)

### RISK ORIENTED: MRD

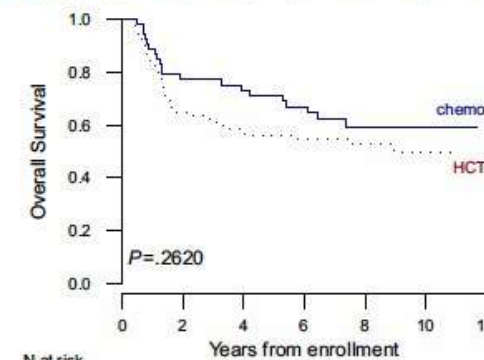
58.6% with  
end of induction MRD  $<10^{-4}$



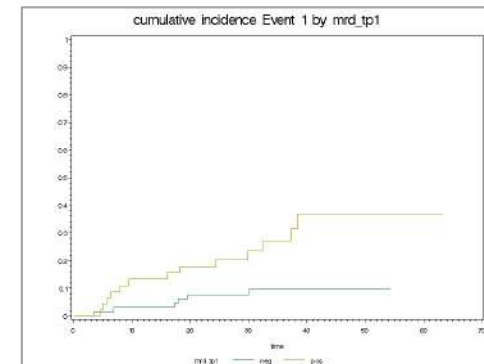
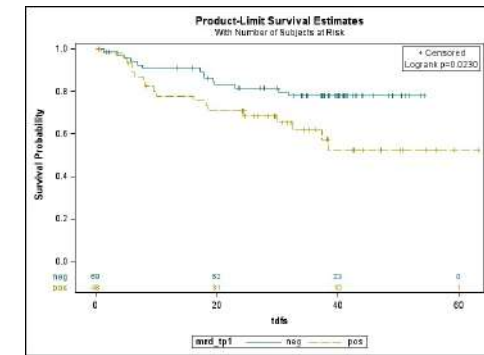
Bassan R et al, *Blood* 2009; *Blood Cancer J* 2014 and 2019



N at risk	0	2	4	6	8	10	12
Ph- 10/07	161	97	84	73	34	15	6
Ph- 09/00	304	146	113	97	62	25	6



N at risk	0	2	4	6	8	10	12
HCT	85	54	47	43	21	9	
chemo	55	40	37	30	13	6	



Bassan R et al, *EHA* 2022

# Improving risk stratification - II



**Prognostic Index<sub>UKALL</sub>**  
 $T(MRD^*) \times -0.218$   
 $+ CYTO-GR \times -0.440$   
 $+ CYTO-HR \times 1.066$   
 $+ \log(WCC^*) \times 0.138$

*\*continuous risk variables*

**Relapse 42%** **MAC**

**EFS 31%** **RIC**

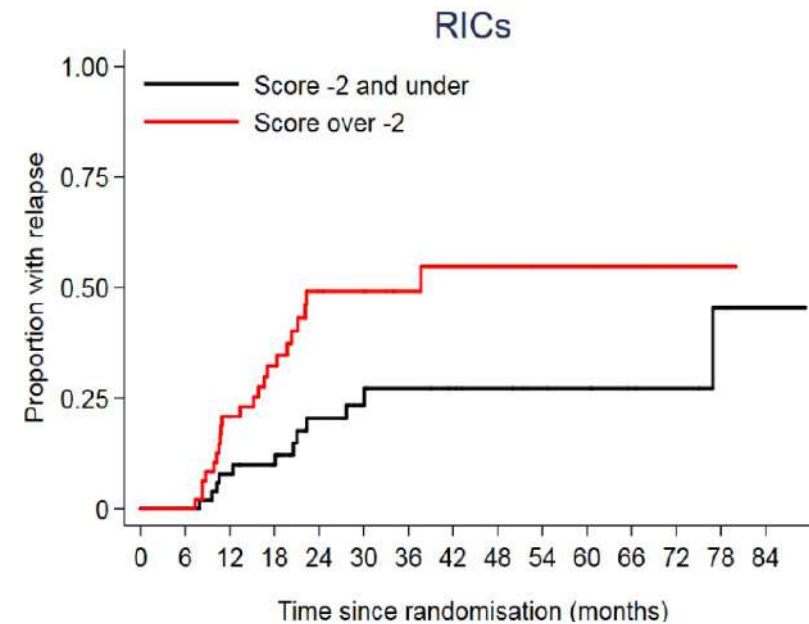
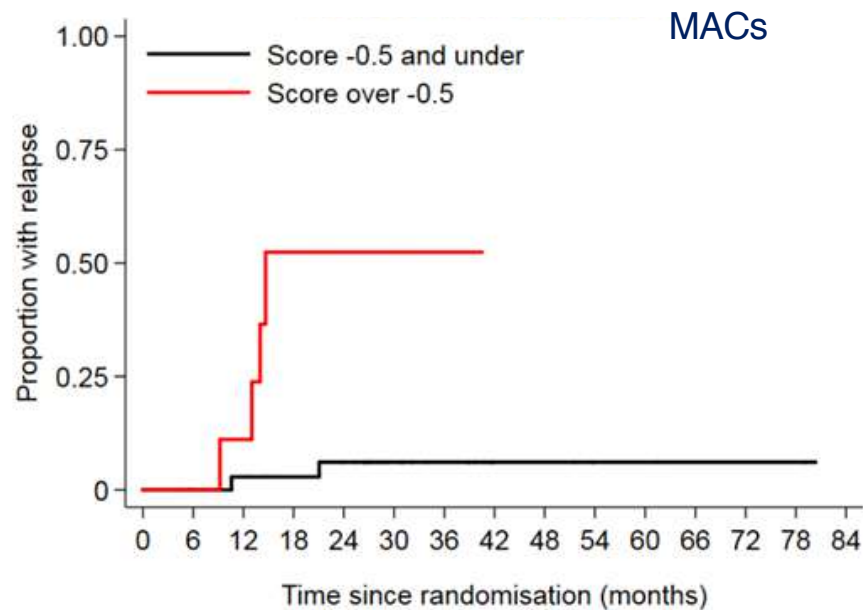
**EFS 90%** **chemo**

**Selected examples of how PI<sub>UKALL</sub> can be used to identify patients on the same treatment pathway who have differential outcomes**

	Hazard ratio (95% CI)	3 years rates (95% CI)	p value
<b>Risk of relapse after myeloablative alloSCT (n=53)</b>			
PI2 score ≤ -1.5	1	5% (1-19)	0.006
PI2 score > -1.5	11.1 (2-62)	42% (18-78)	
<b>Event free survival (EFS) after RIC alloSCT (n=105)</b>			
PI2 score ≤ -2.0	1	62% (45-75)	0.004
PI2 score > -2.0	2.3 (1.3-4.1)	31% (17-46)	
<b>EFS of standard risk patients after maintenance chemotherapy (n=51)</b>			
PI1 score ≤ -2.25	1	90% (66-98)	0.041
PI1 score > -2.25	5.1 (1.1-24.3)	71% (45-86)	

Moorman A, et al. HemaSphere 2019;3:748-9 (abstr #S1621).

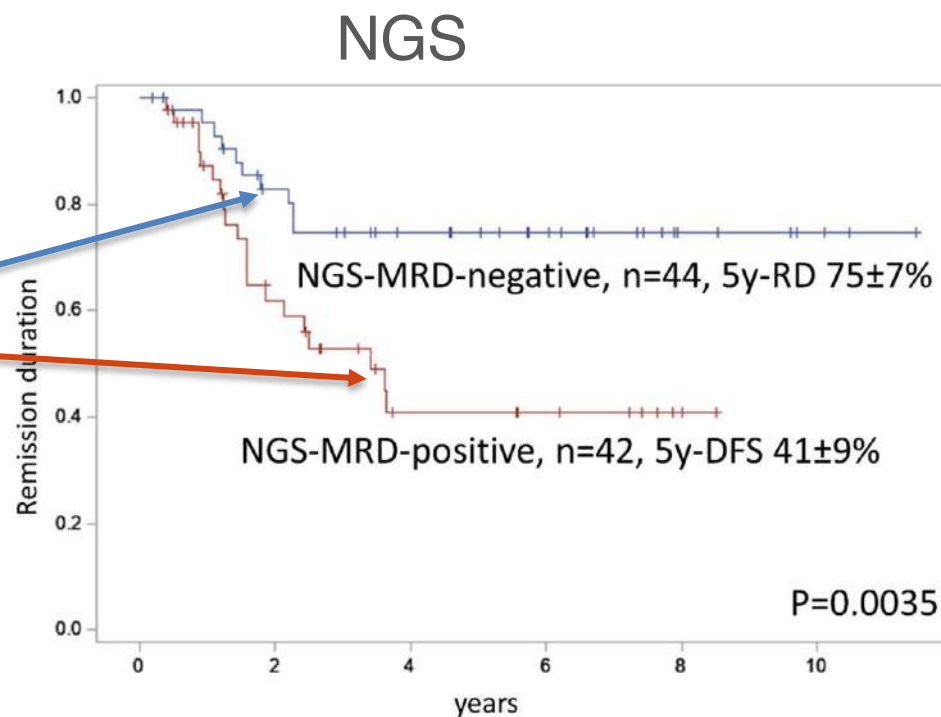
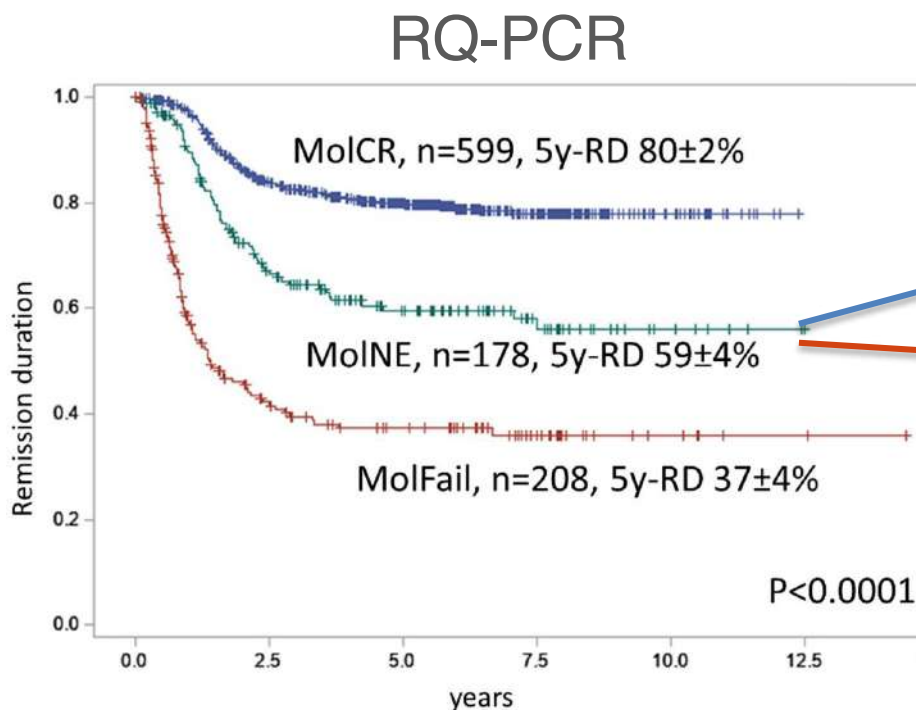
# PI<sub>UKALL</sub> vs. SCT



PI2 score	P-value	Relapse rate (3-year)	EFS (3-year)
≤-0.5		6% (2-22)	65% (46-79)
>-0.5	0.011	52% (23-88)	38% (10-66)

PI2 score	p value	Relapse rate (3 yr)	EFS (3 yr)
≤-2		27% (16-45)	62% (45-75)
>-2	0.011	49% (35-65)	31% (17-46)

# Improving MRD analysis: NGS



Uncertain prognostic effects of molecular MRD Not Evaluable (**MoINE**) by RQ-PCR with probe sensitivity at least  $10^{-4}$

- **MoICR**: negative
- **MoIFail**:  $\geq 10^{-4}$

- **MRD** as key prognostic factor and determinant of allo-SCT choice
- **MRD** interacts with other (WBC / GENETICS)  
*toward integrated risk models*
- **MRD** analysis can be improved (NGS, ddPCR)
- **MRD is a therapeutic target and supports progress in ALL therapy**
  - New MRD-oriented therapies



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