

How should we monitor MRD in Ph+ ALL : BCR::ABL1 or IG/TR ?

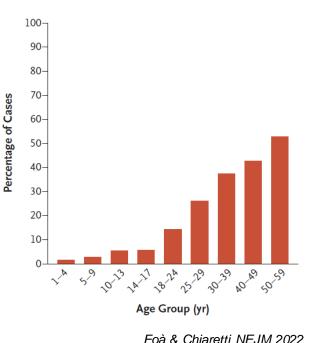
Rathana Kim

Leukemia institute Paris Saint-Louis, Inserm UMR1342 Laboratoire d'Hématologie - Hôpital Saint-Louis - AP-HP, Paris, France

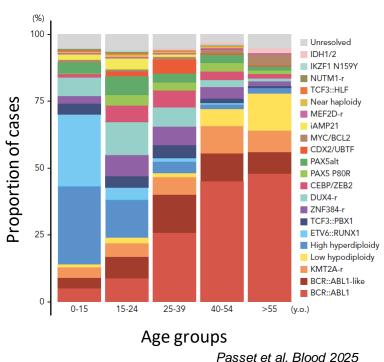
Disclosures Rathana KIM

Nothing to disclose

Philadelphia chromosome-positive acute lymphoblastic leukemia (Ph+ ALL)

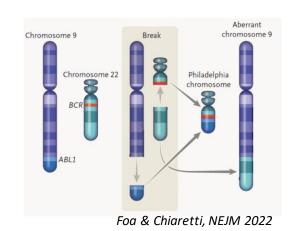


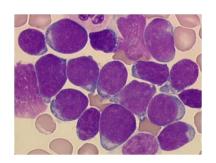
Foà & Chiaretti NEJM 2022



- The most prevalent B-ALL subtype in adults \rightarrow ~1/3 cases
- Proportion increase with age → up to 50% B-ALL cases after 55-60 years old

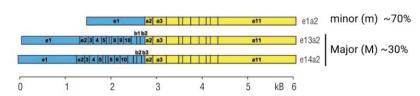
Ph+ ALL: two molecular targets for MRD monitoring





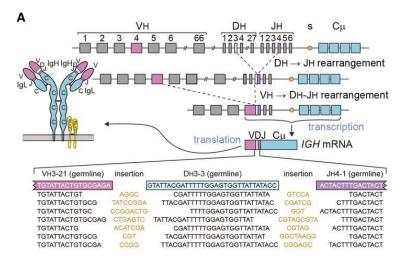
Proliferation of malignant

B-cell lymphoblasts in the bone marrow



Burmeister et al. Haematologica 2007

BCR::ABL1
Oncogenic fusion



van Dongen et al. Blood 2015

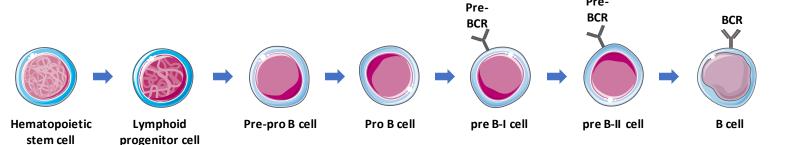
Lymphoid specific markers

Immunoglobulin/T-cell receptor

(IG/TR) gene rearrangements



Novel insights in Ph+ ALL pathophysiology revealed by MRD monitoring



CLP

CSH

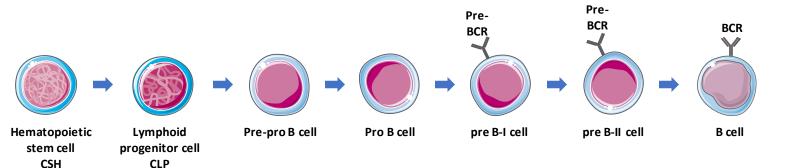
BCR::ABL1 fusion

RAG mediated recombinations (IG/TR and others targets)

Until recently

De novo Ph+ ALL was defined by the acquisition of BCR::ABL1 fusion in committed B-cell precursors

Novel insights in Ph+ ALL pathophysiology revealed by MRD monitoring

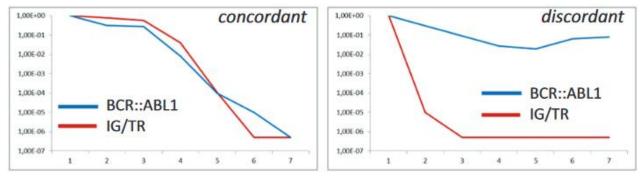


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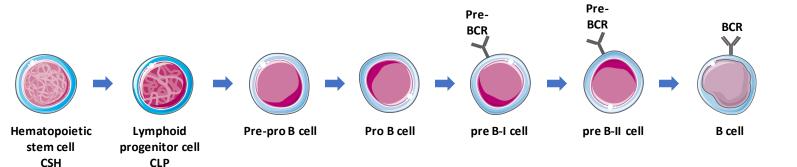


MRD monitoring of two illustrative cases

Zuna et al. Leukemia 2022 Kim et al. JCO 2024



Novel insights in Ph+ ALL pathophysiology revealed by MRD monitoring

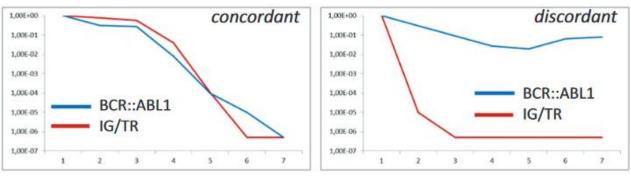


BCR::ABL1 fusion

RAG mediated recombinations (IG/TR and others targets)

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MRD monitoring of two illustrative cases



Evidence for multilineage involvement in de novo Ph+ ALL revealed by MRD assessed on both targets

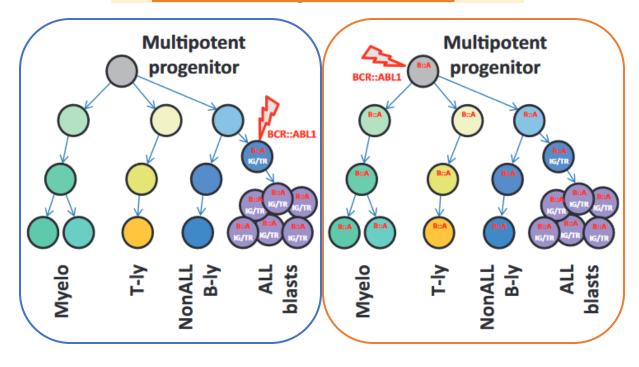
~35-40% of patients

Zuna et al. Leukemia 2022 Kim et al. JCO 2024

New pathophysiology model of Ph+ ALL

2022 ICC classification

B-ALLwith t(9;22)(q34.1;q11.2)/BCR::ABL1
with lymphoid only involvement
with multilineage involvement

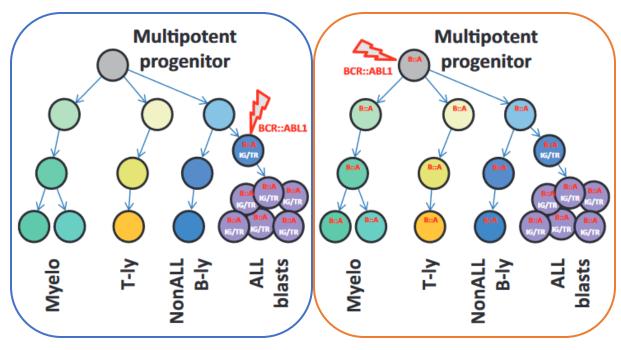


Zuna et al. Leukemia 2022 Kim et al. JCO 2024 Arber et al., Blood 2022

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New clinical questions

- Which MRD target is the most clinically relevant?
- Impact on new therapeutic strategies?

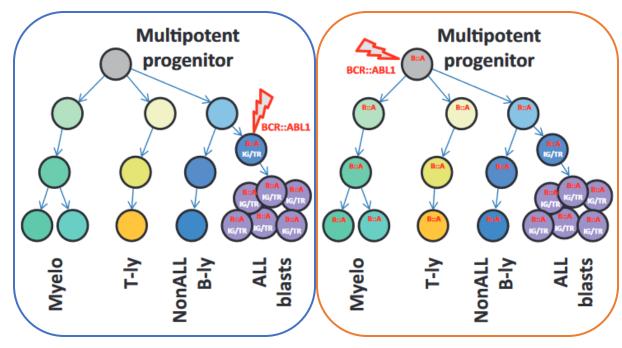
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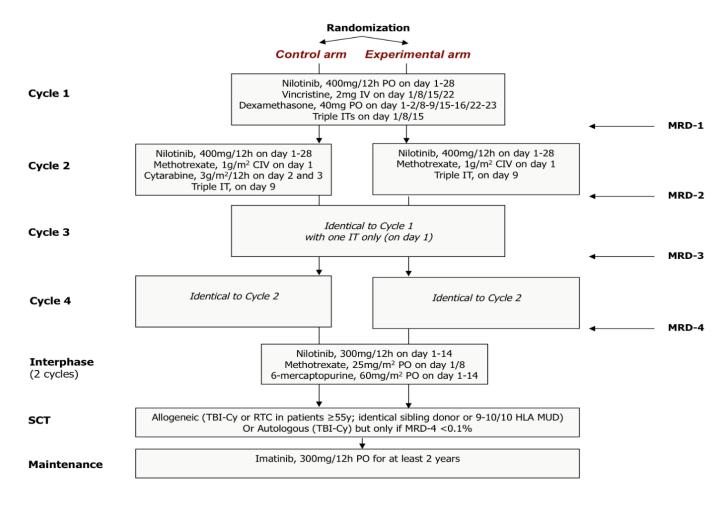
Zuna et al. Leukemia 2022 Kim et al. JCO 2024 Arber et al., Blood 2022



The GRAAPH-2014 study

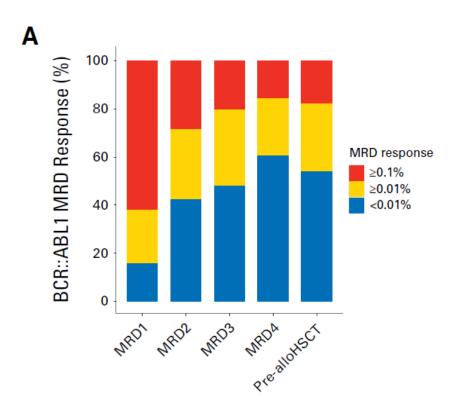
Nilotinib combined with lower-intensity chemotherapy for front-line treatment of younger adults (18-59 years) with Ph+ ALL





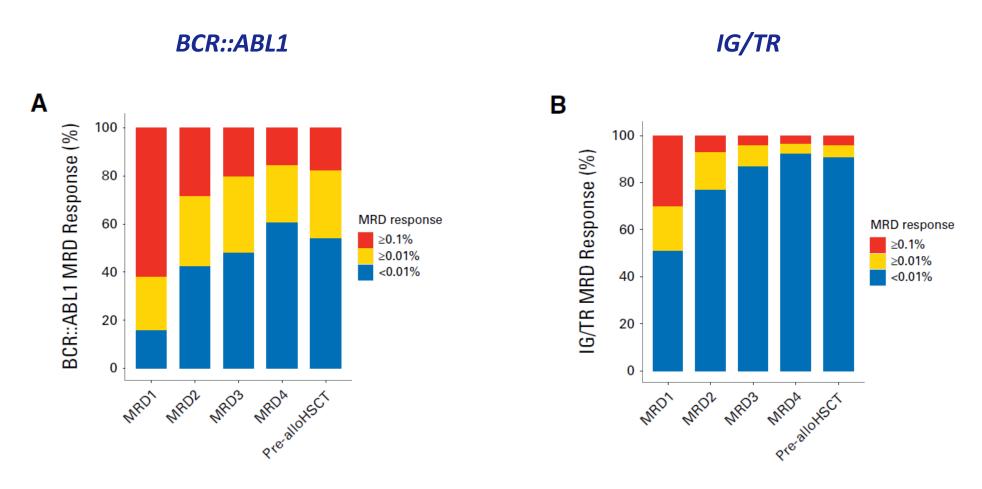
MRD responses during the early phases of treatment





BCR::ABL1 MRD response reaches a plateau

MRD responses during the early phases of treatment



BCR::ABL1 MRD response reaches a plateau
Better proportion of IG/TR MRD clearance during early phase of treatment

MRD responses during the early phases of treatment



BCR::ABL1 MRD Response (%) MRD response 60 ≥0.1% ≥0.01% <0.01% 40 20

Blinatumomab + Dasatinib D-ALBA study

Assessment	No Molecular Response	Complete Molecular Response	Positive Nonquantifiable Response	Overall Molecula Response			
		number of patients/total number (percent)					
Induction period							
Day 22	48/58 (83)	3/58 (5)	7/58 (12)	10/58 (17)			
Day 45	43/60 (72)	9/60 (15)	8/60 (13)	17/60 (28)			
Day 57	38/56 (68)	11/56 (20)	7/56 (12)	18/56 (32)			
Day 85	42/59 (71)	6/59 (10)	11/59 (19)	17/59 (29)			
Blinatumomab cycle							
After cycle 1	20/55 (36)	19/55 (35)	16/55 (29)	35/55 (64)			
After cycle 2	22/55 (40)	23/55 (42)	10/55 (18)	33/55 (60)			
After cycle 3	12/40 (30)	20/40 (50)	8/40 (20)	28/40 (70)			
After cycle 4	7/36 (19)	17/36 (47)	12/36 (33)	29/36 (81)			
After cycle 5	8/29 (28)	16/29 (55)	5/29 (17)	21/29 (72)			

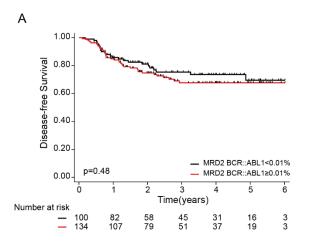
BCR::ABL1 MRD response reaches a plateau
Similar findings in immunotherapy + TKI combination (D-ALBA study)

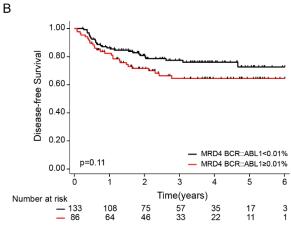
Kim et al. JCO 2024 Foà et al., NEJM 2020



Prognostic impact of MRD

BCR::ABL1

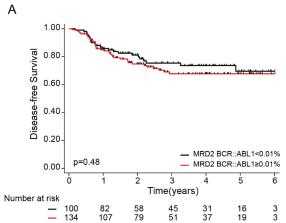


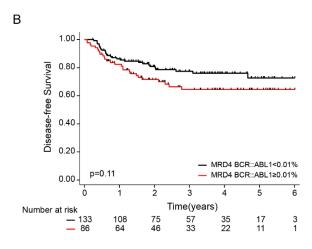


No prognosis value of BCR::ABL1 in a chemotherapy + TKI + alloHSCT regimen (GRAAPH-2014 study)

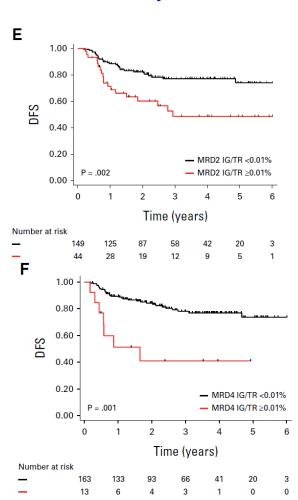
Prognostic impact of MRD







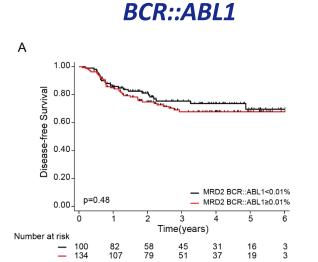
IG/TR

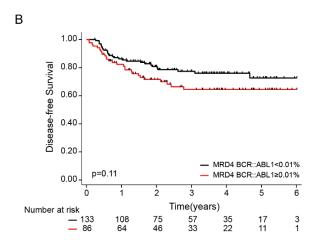


IG/TR MRD is a better predictor of outcome

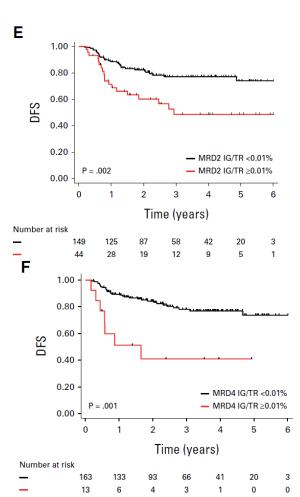


Prognostic impact of MRD

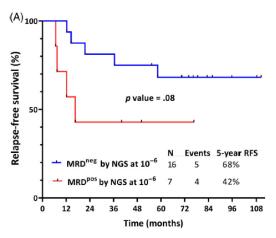


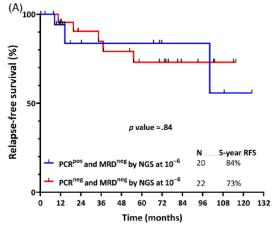


IG/TR



MD Anderson study IGH MRD by NGS (clonoSEQ)





IG/TR MRD is a better predictor of outcome

Kim et al. JCO 2024 Short et al. Am J Hematol 2023



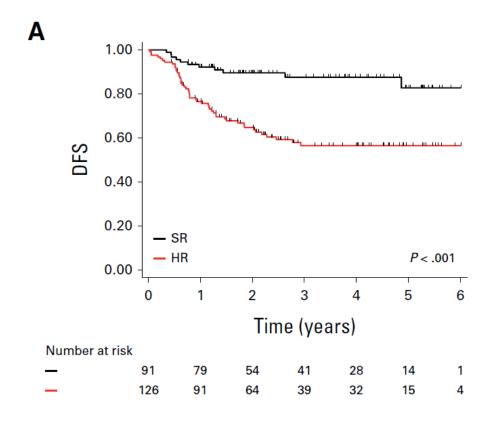
Incorporating IG/TR MRD into a prognostic model

TABLE 3. Univariable and Multivariable Analyses for DFS

No.	Univariable		Multivariable ^a	
	HR (95% CI)	P	HR (95% CI)	P
259	0.99 (0.98 to 1.02)	.70	0.99 (0.97 to 1.03)	.84
259	1.23 (1.05 to 1.44)	.01	_	_
259	1.84 (1.17 to 2.89)	.008	2.95 (1.44 to 6.03)	.003
248	1.16 (0.89 to 1.50)	.27	1.17 (0.92 to 1.49)	.21
259	1.05 (0.64 to 1.72)	.84	1.69 (0.79 to 3.57)	.17
259	0.90 (0.75 to 1.09)	.28	1.02 (0.50 to 2.05)	.97
228	0.83 (0.49 to 1.41)	.50	0.77 (0.40 to 1.50)	.44
193	2.49 (1.40;4.40)	.002	2.58 (1.34 to 4.96)	.004
259	1.59 (1.00;2.51)	.049	1.61 (0.86 to 3.02)	.14
	259 259 259 248 259 259 228 193	No. HR (95% CI) 259 0.99 (0.98 to 1.02) 259 1.23 (1.05 to 1.44) 259 1.84 (1.17 to 2.89) 248 1.16 (0.89 to 1.50) 259 1.05 (0.64 to 1.72) 259 0.90 (0.75 to 1.09) 228 0.83 (0.49 to 1.41) 193 2.49 (1.40;4.40)	No. HR (95% CI) P 259 0.99 (0.98 to 1.02) .70 259 1.23 (1.05 to 1.44) .01 259 1.84 (1.17 to 2.89) .008 248 1.16 (0.89 to 1.50) .27 259 1.05 (0.64 to 1.72) .84 259 0.90 (0.75 to 1.09) .28 228 0.83 (0.49 to 1.41) .50 193 2.49 (1.40;4.40) .002	No. HR (95% CI) P HR (95% CI) 259 0.99 (0.98 to 1.02) .70 0.99 (0.97 to 1.03) 259 1.23 (1.05 to 1.44) .01 — 259 1.84 (1.17 to 2.89) .008 2.95 (1.44 to 6.03) 248 1.16 (0.89 to 1.50) .27 1.17 (0.92 to 1.49) 259 1.05 (0.64 to 1.72) .84 1.69 (0.79 to 3.57) 259 0.90 (0.75 to 1.09) .28 1.02 (0.50 to 2.05) 228 0.83 (0.49 to 1.41) .50 0.77 (0.40 to 1.50) 193 2.49 (1.40;4.40) .002 2.58 (1.34 to 4.96)

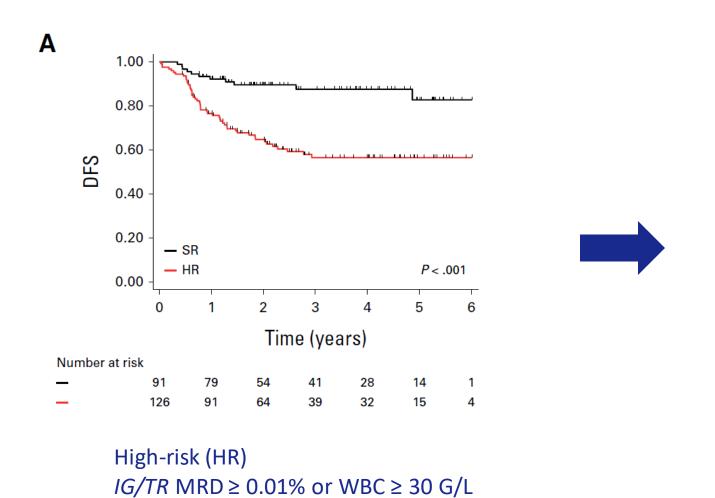
IG/TR MRD and high WBC were independently associated with poorer DFS

Incorporating IG/TR MRD into a prognostic model

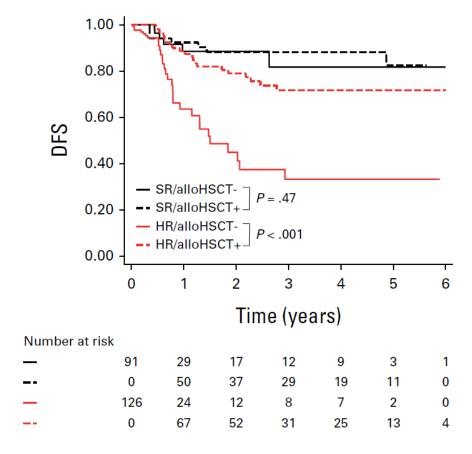


High-risk (HR) IG/TR MRD $\geq 0.01\%$ or WBC ≥ 30 G/L

Incorporating IG/TR MRD into a prognostic model



Simon-Makush plots



Patients with good *IG/TR* MRD response have excellent outcomes regardless of allo-HSCT



Summary

➤ A large fraction of Ph+ ALL have non-lymphoblastic *BCR::ABL1*-positive cells that persist under treatment

> BCR::ABL1 multilineage involvement and BCR::ABL1 persistence are not associated with poorer outcome

> IG/TR MRD becomes the new standard for disease monitoring and treatment stratification in Ph+ ALL

Discussion

Is BCR::ABL1 still a relevant MRD marker in Ph+ ALL management?



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- **√** When no diagnostic sample is available for *IG/TR* target identification
- ✓ For *ABL1* TKD mutation screening

Is BCR::ABL1 still a relevant MRD marker in Ph+ ALL management?



- **√** When no diagnostic sample is available for *IG/TR* target identification
- ✓ For ABL1 TKD mutation screening
- ✓ In case of MRD negativity with both *IG/TR* and *BCR::ABL1* MRD

maintenance or post allo-HSCT

Is BCR::ABL1 still a relevant MRD marker in Ph+ ALL management?



- **√** When no diagnostic sample is available for *IG/TR* target identification
- ✓ For ABL1 TKD mutation screening
- ✓ In case of MRD negativity with both *IG/TR* and *BCR::ABL1* MRD maintenance or post allo-HSCT
- ✓ In case of sustained *IG/TR* MRD negativity and stable *BCR::ABL1* MRD ?

Follow-up stable disease, indicate marrow aspirate with IG/TR MRD if molecular progression



Perspectives

- ✓ In the context of increased survival and decreased rates of allo-HSCT on Ph+ ALL, need to assess the long-term significance of persisting BCR::ABL1-positive cells
 - May be associated with TKI potency?
 - Long-term outcome? potential for CML-like disease or B-ALL recurrence?
 - Is treatment-free remission an option for patients with multilineage Ph+ ALL?

Aknowledgments







<u>AP-HP Saint-Louis</u>

<u>UMR1342</u> <u>Equipe Génome et Cancers</u> All investigators and biologists
from the GRAALL





GRAALL LALA GOELAMS SAKK

Emmanuelle Clappier

Jean-Michel Cayuela Marie Passet Jean Soulier GRAAPH-2014 Yves Chalandon Philippe Rousselot

Véronique Lhéritier Hervé Dombret Nicolas Boissel



