

MRD-driven strategy in Ph-like ALL: does it work?

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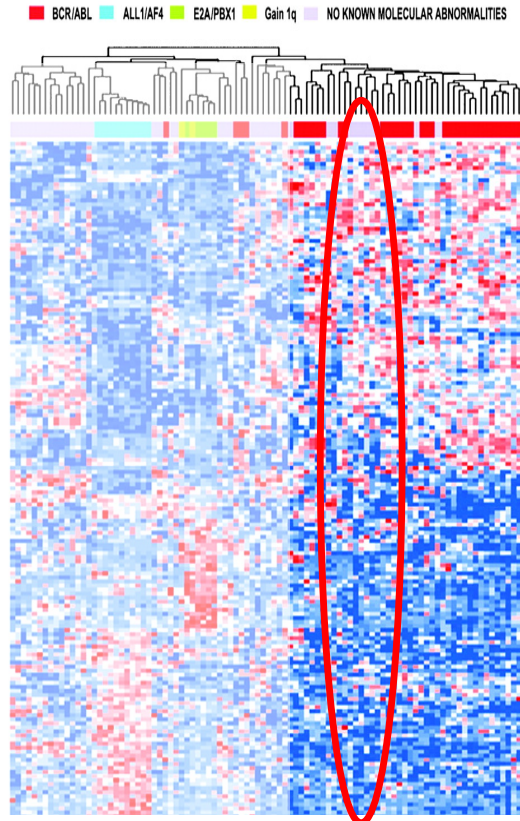
April 27th 2021



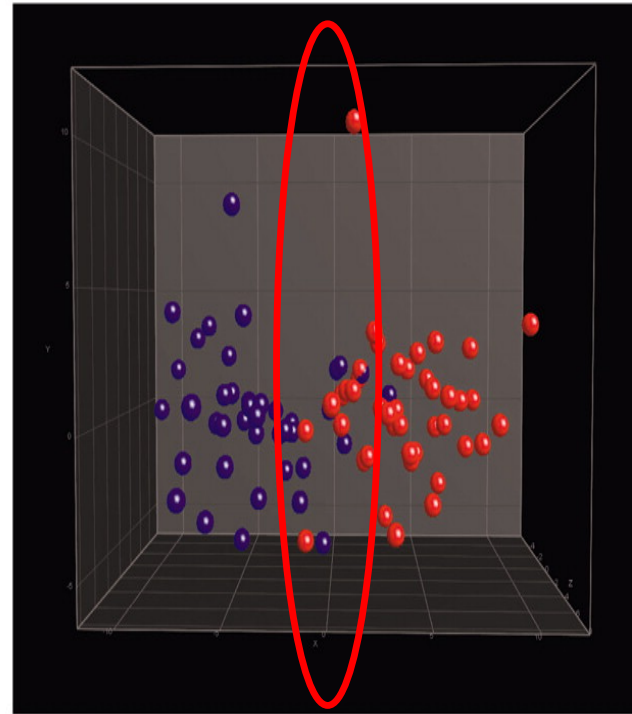
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Ph-like ALL: first report in adults

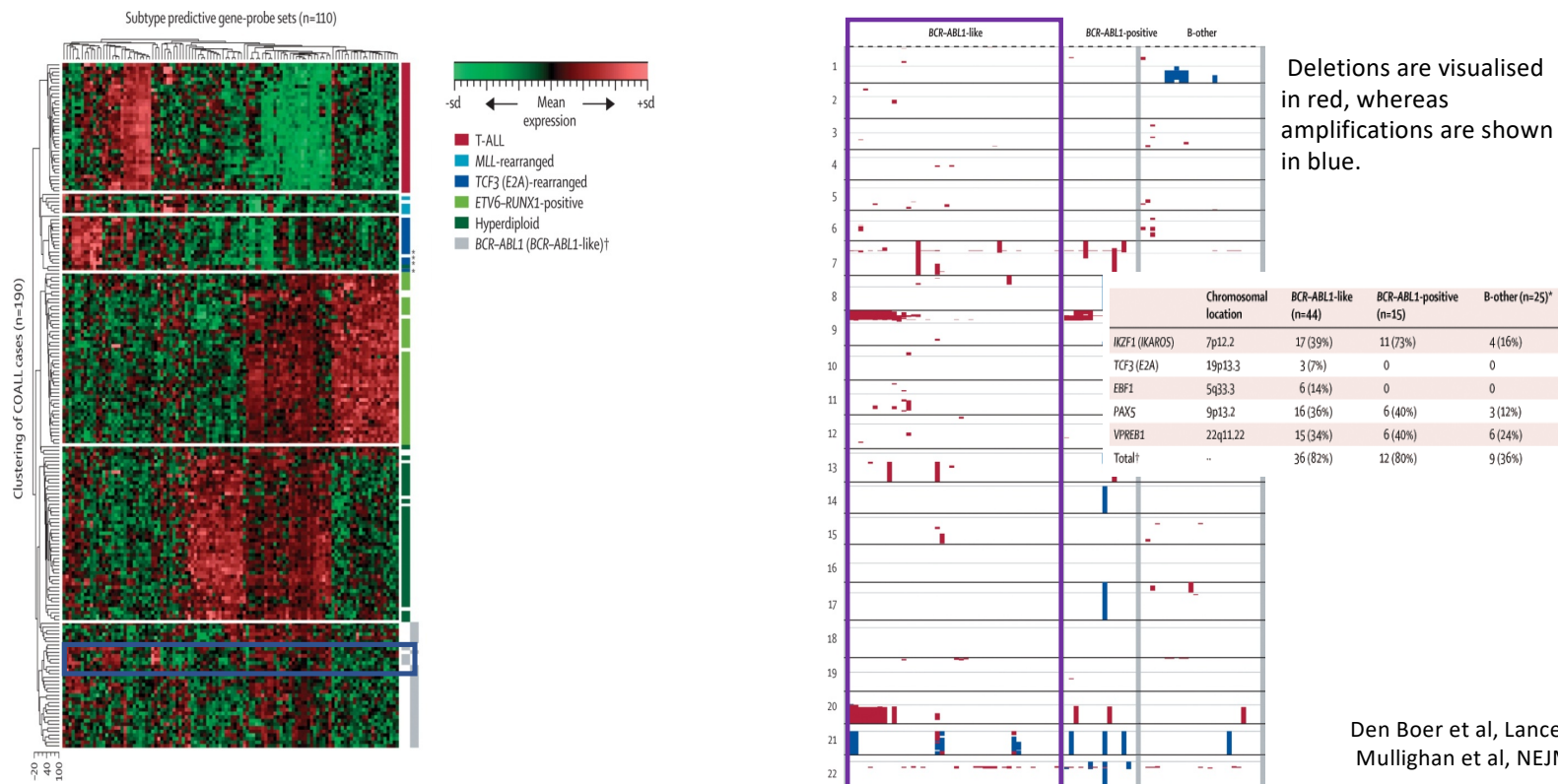


Chiaretti et al, CCR 2005



Haferlach et al, Blood 2005

Ph-like ALL: genetic characterization

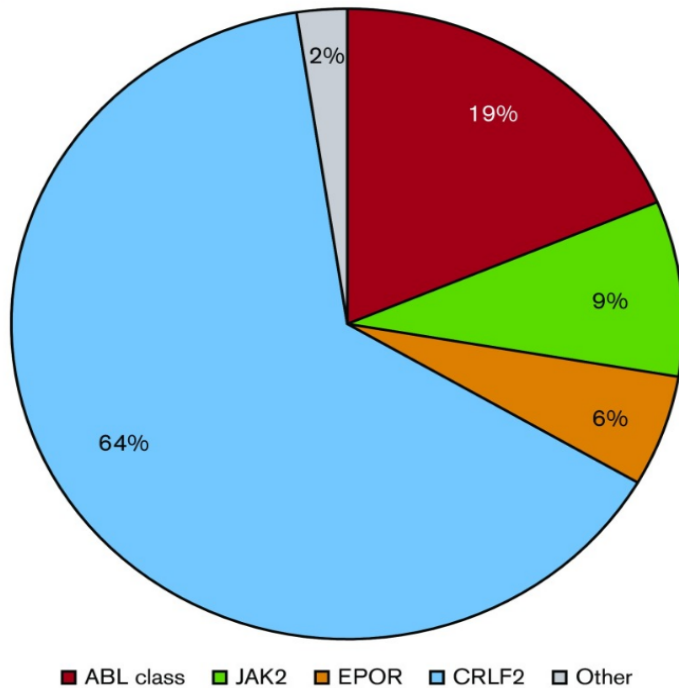


GEP: Identification, within children (n=297), of a subset with a transcriptional profile resembling that of *BCR/ABL1*+ cases (≈15-20%)

Clinical features: Hyperleukocytosis, poor response to VCR, ASP and DNM, poor prognosis (reduced DFS at 5 years and increased resistance to induction)

Array-CGH: *IKZF1*, *PAX5*, *TCF3* and *VPREB1* deletions, *CRLF2* deregulation

Ph-like ALL: genetic characterization



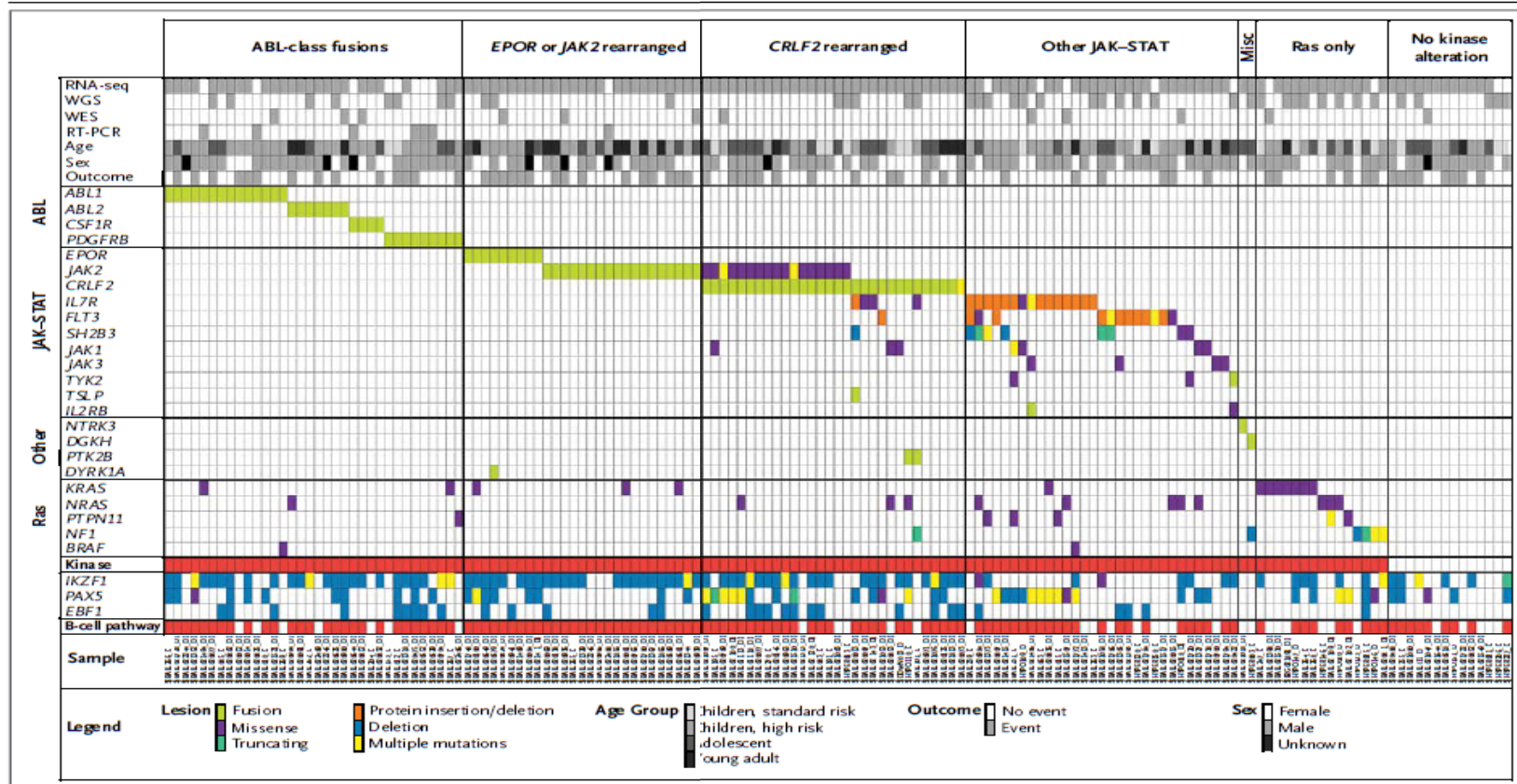
Harvey RC and Tasian SK. *Blood Adv*, 2020

| Kinase gene | Fusion partners, n | Patients, n | 5' genes |
|-------------|--------------------|-------------|---|
| ABL1 | 6 | 14 | ETV6, NUP214, RCSD1, RANBP2, SNX2, ZMIZ1 |
| ABL2 | 3 | 7 | PAG1,* RCSD1,* ZC3HAV1* |
| CSF1R | 1 | 4 | SSBP2* |
| PDGF RB | 4 | 11 | EBF1, SSBP2,* TNIP1,* ZEB2* |
| CRLF2 | 2 | 30 | IGH, P2RY8 |
| JAK2 | 10 | 19 | ATF7IP,* BCR, EBF1,* ETV6, PAX5, PPFIBP1,* SSBP2, STRN3, TERF2,* TPR* |
| EPOR | 2 | 9 | IGH, IGK* |
| DGKH | 1 | 1 | ZFAND3* |
| IL2RB | 1 | 1 | MYH9* |
| NTRK3 | 1 | 1 | ETV6† |
| PTK2B | 2 | 1 | KDM6A,* STAG2* |
| TSLP | 1 | 1 | IQGAP2* |
| TYK2 | 1 | 1 | MYB* |

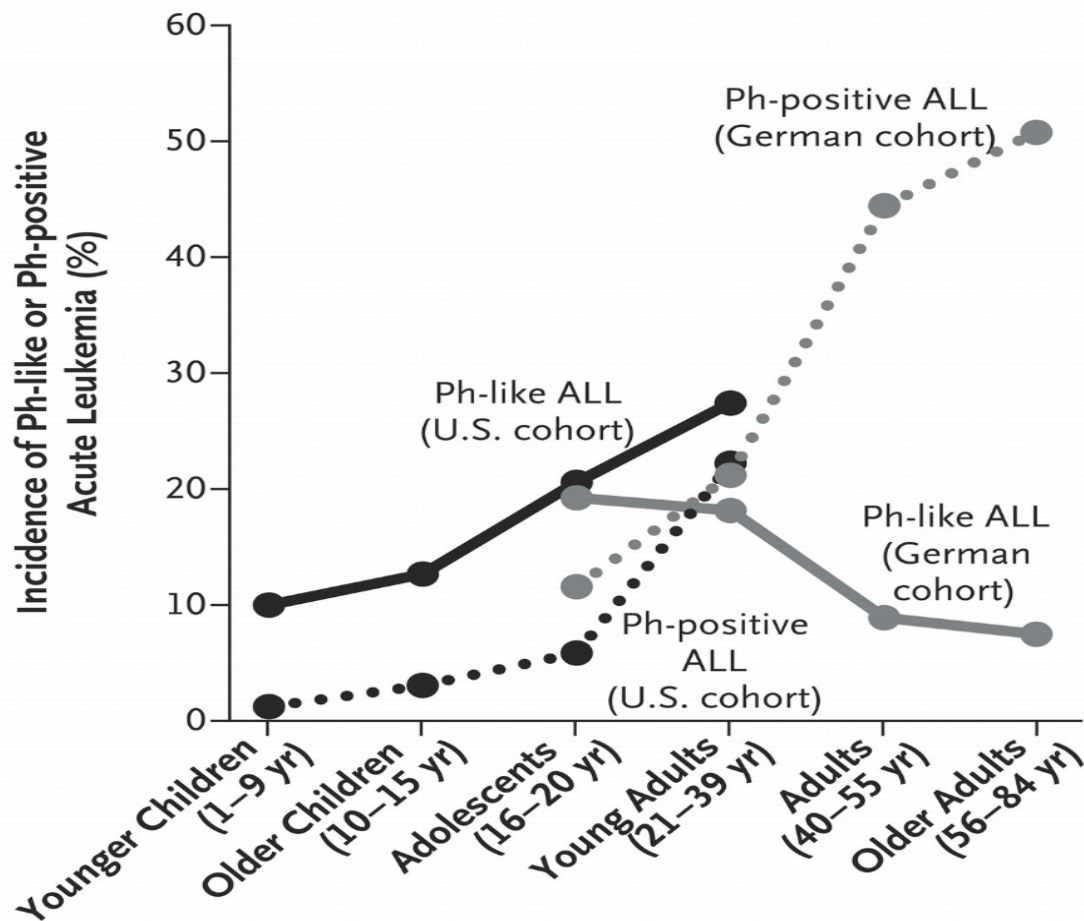
Roberts KG, et al. *N Engl J Med* 2014;371:1005–1015

Targetable Kinase-Activating Lesions in Ph-like Acute Lymphoblastic Leukemia

K.G. Roberts, Y. Li, D. Payne-Turner, R.C. Harvey, Y.-L. Yang, D. Pei, K. McCastlain, L. Ding, C. Lu, G. Song, J. Ma, J. Becksfort, M. Rusch, S.-C. Chen, J. Easton, J. Cheng, K. Boggs, N. Santiago-Morales, I. Iacobucci, R.S. Fulton, I. Wen, M. Valentine, C. Cheng



BCR-ABL1-like. Incidence



Incidence is higher in AYA (10% in children; 27% in AYA). NEVER detected in cases positive for known fusion transcripts (*BCR/ABL1*, *KMT2A-r*, *TCF3/PBX1*)

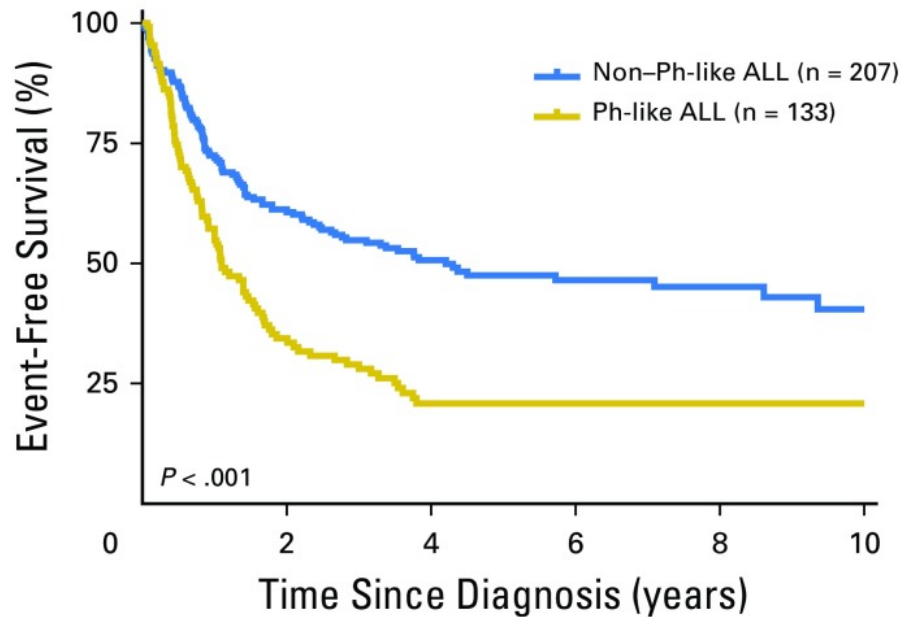
However:

- It highly depends on the denominator (all B-lineage ALL or B-neg ALL) and
- on the assay used for *BCR/ABL1*-like identification
- More adult cases are being evaluated → incidence in adults is almost equal to AYA ≥ 25%

Survival in adults

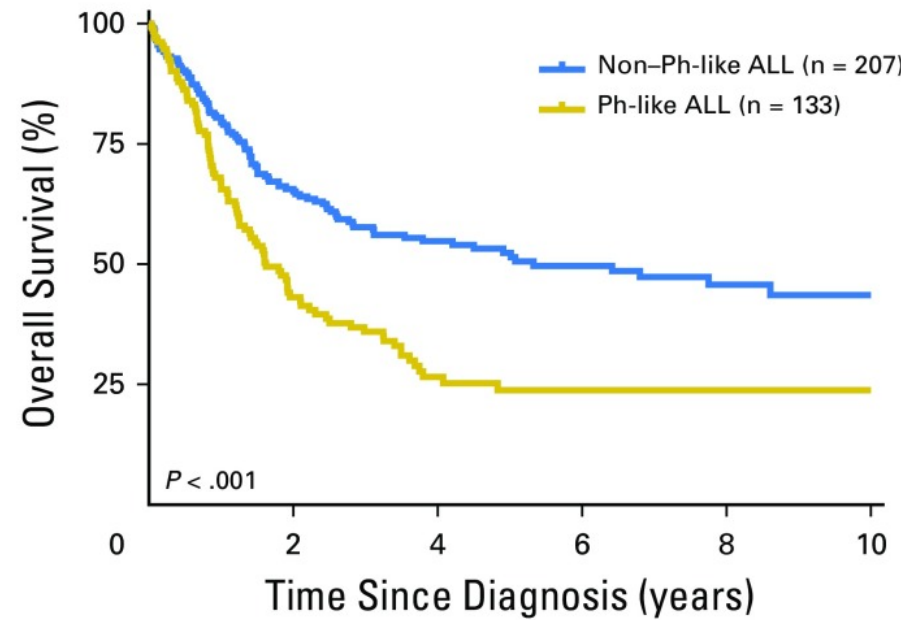
- Significantly inferior survival (EFS, DFS, OS) in all reported studies

A



| No. at risk: | | 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 |
|-----------------|-----|-----|-----|-----|----|----|----|----|----|----|----|----|
| Non-Ph-like ALL | 207 | 146 | 117 | 102 | 73 | 53 | 47 | 35 | 28 | 20 | 13 | |
| Ph-like ALL | 133 | 70 | 39 | 32 | 19 | 15 | 14 | 11 | 9 | 5 | 3 | |

B



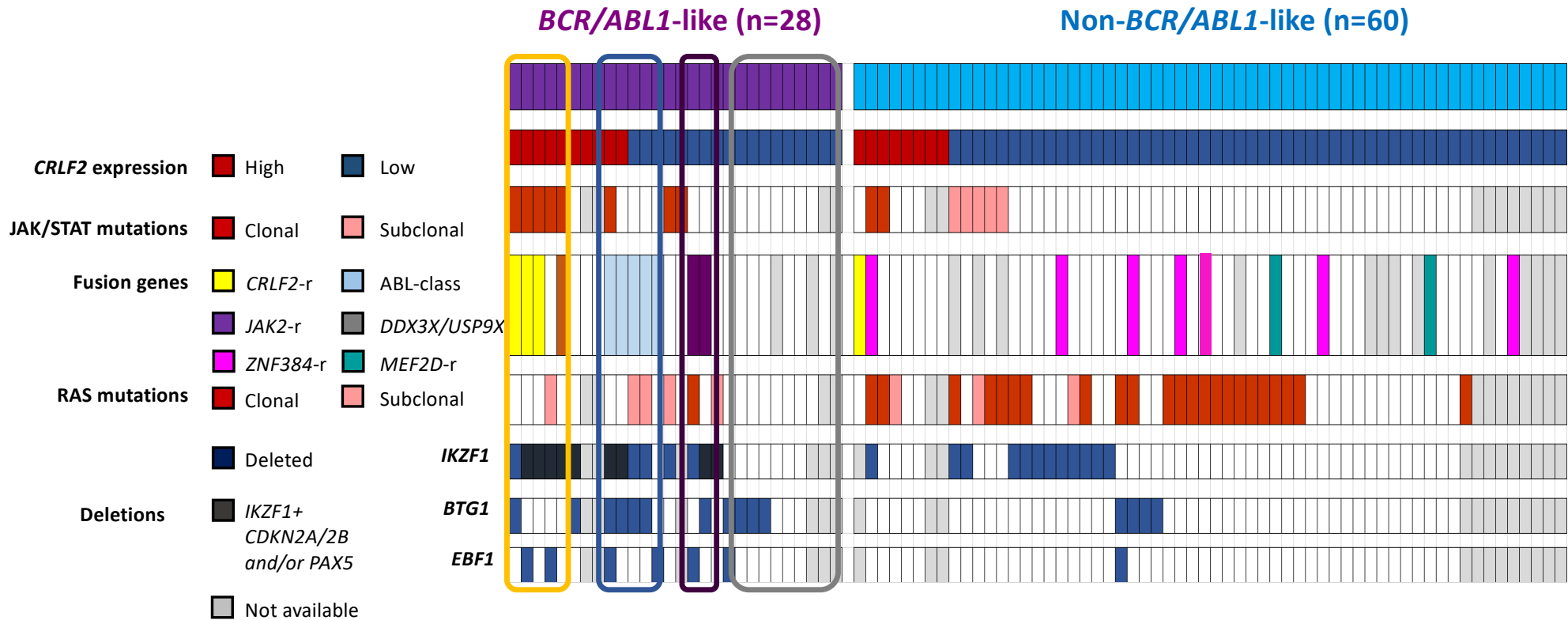
| No. at risk: | | 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 |
|-----------------|-----|-----|-----|-----|----|----|----|----|----|----|----|----|
| Non-Ph-like ALL | 207 | 162 | 127 | 107 | 80 | 60 | 51 | 37 | 29 | 20 | 14 | |
| Ph-like ALL | 133 | 82 | 49 | 40 | 23 | 17 | 16 | 12 | 9 | 5 | 3 | |

Ph-like ALL: diagnosis

| | |
|-----------------------|--|
| LDA | Quantification of expression of 15 transcripts (Kang BW et al, ASH 2013) |
| Q-RT-PCR | Quantification of expression of 10 transcripts (Chiaretti S et al, BJH 2018) |
| Integrated algorithms | LDA, FISH, RT-PCR, NGS (RNA-seq, WGS, WES) (Roberts KG et al, NEJM 2014) |
| | FISH, RT-PCR, Q-RT-PCR, NGS (Fasan A et al, ASH 2015) |
| | Known fusion transcripts, <i>JAK2</i> mutations, <i>CRLF2-r</i> (Herold T et al, Haematologica 2016) |
| | <i>CRLF2</i> expression (FC), <i>JAK2</i> mutations, <i>FISH</i> for TK-rearrangements and <i>CRLF2-r</i> (Jain N et al, ASH 2017) |

Not yet available a gold standard

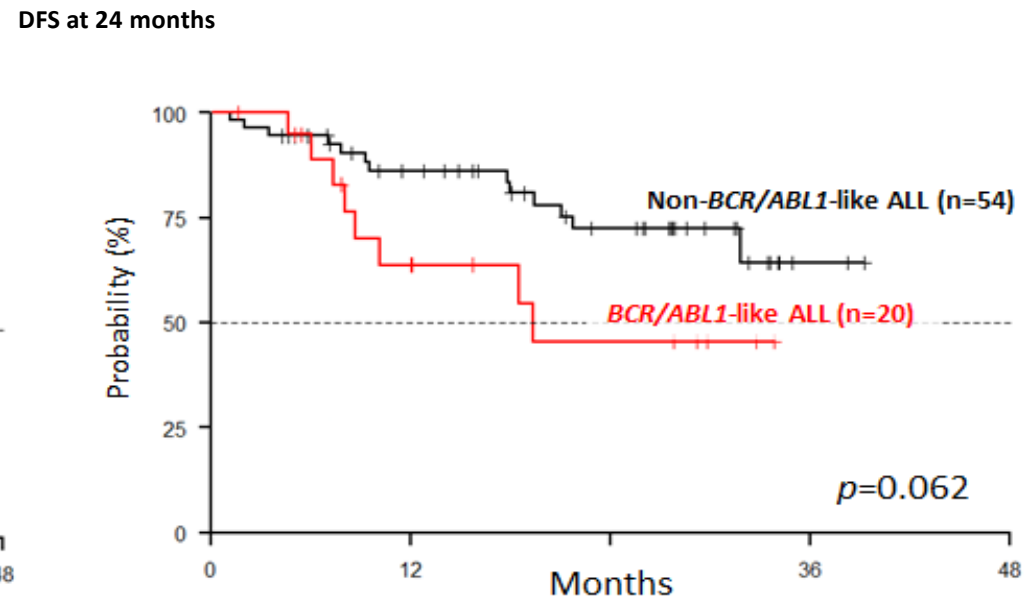
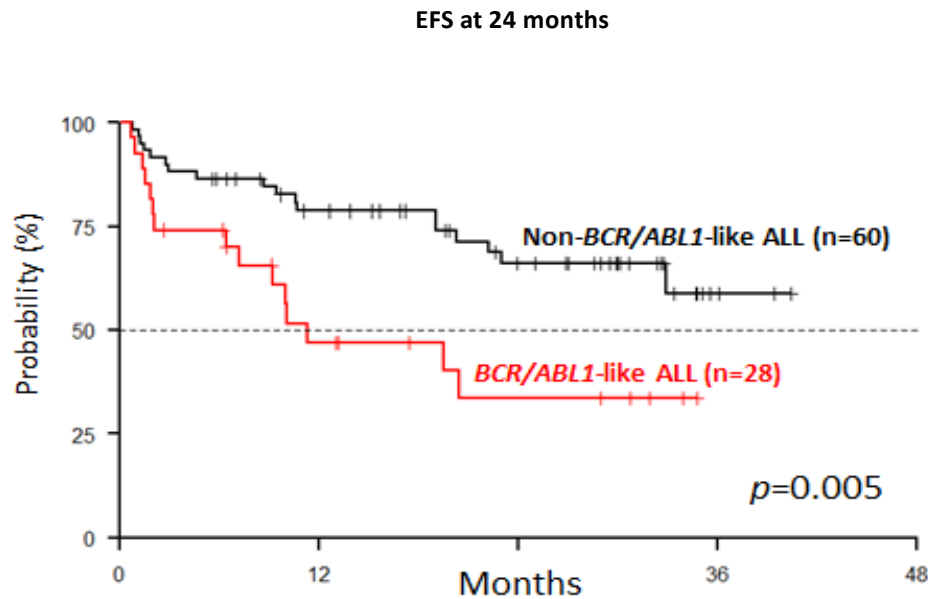
Biological features of GIMEMA LAL1913 patients according to the *BCR/ABL1*-like status



***BCR/ABL1*-like associated lesions identified in 69.6%**

CRLF2 deregulation
 ABL-class genes + other TKs
 No fusions

Ph-like ALL in GIMEMA LAL1913: outcome

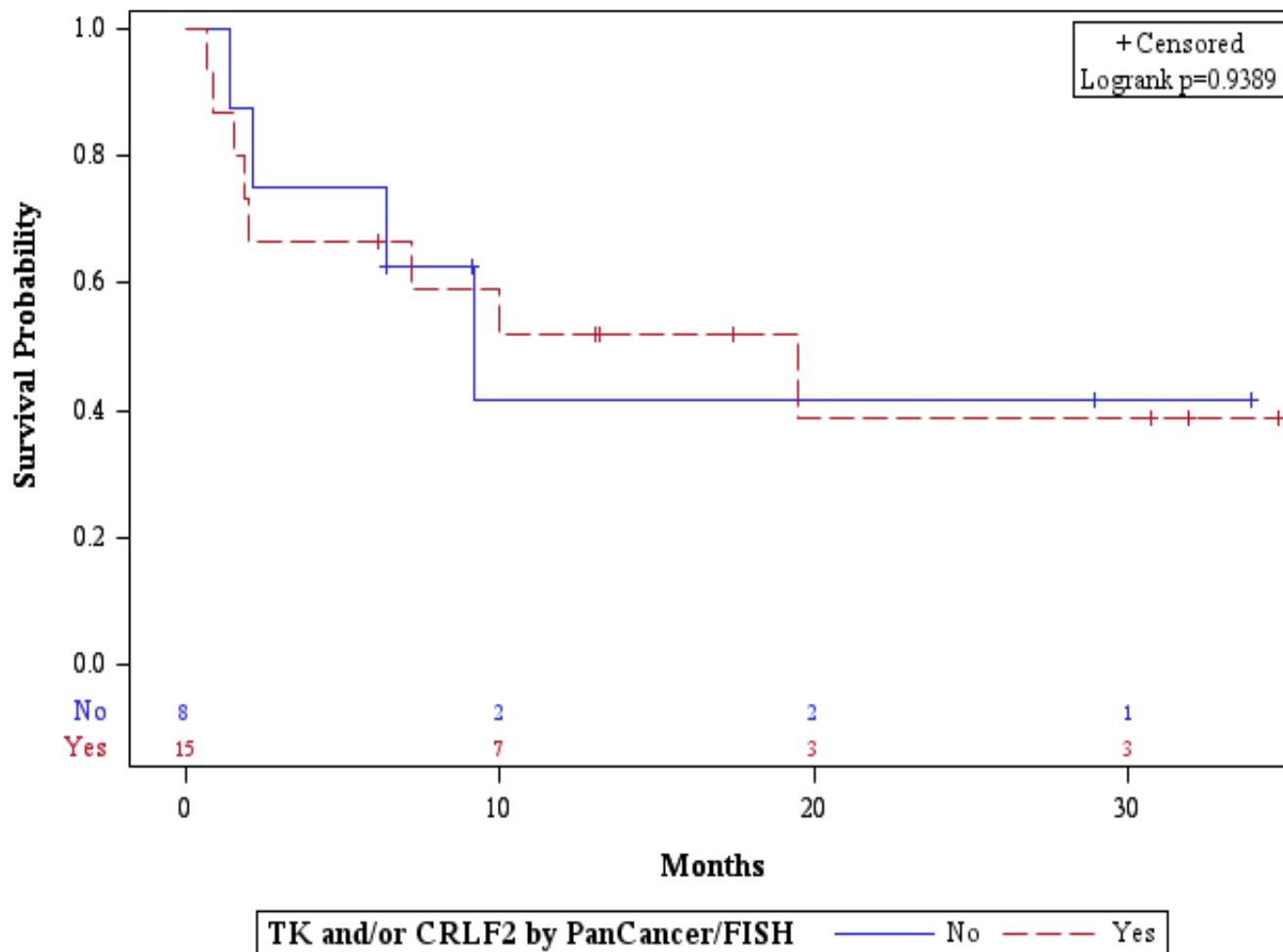


| | HR (95% CI) | p-value |
|------------------------------------|------------------|---------|
| BCR/ABL1-like vs non-BCR/ABL1-like | 2.3 (1.124–4.92) | 0.023 |

Chiaretti S, et al. *Haematologica* 2020; ePub ahead of print.

Clinical outcome of GIMEMA LAL1913 patients according to the presence/absence of well-defined molecular lesions in BCR/ABL1-like cases (I)

EFS



All cases with ABL-class lesions experienced an event within 10 months from diagnosis

Ph-like ALL in GIMEMA LAL1913 : MRD

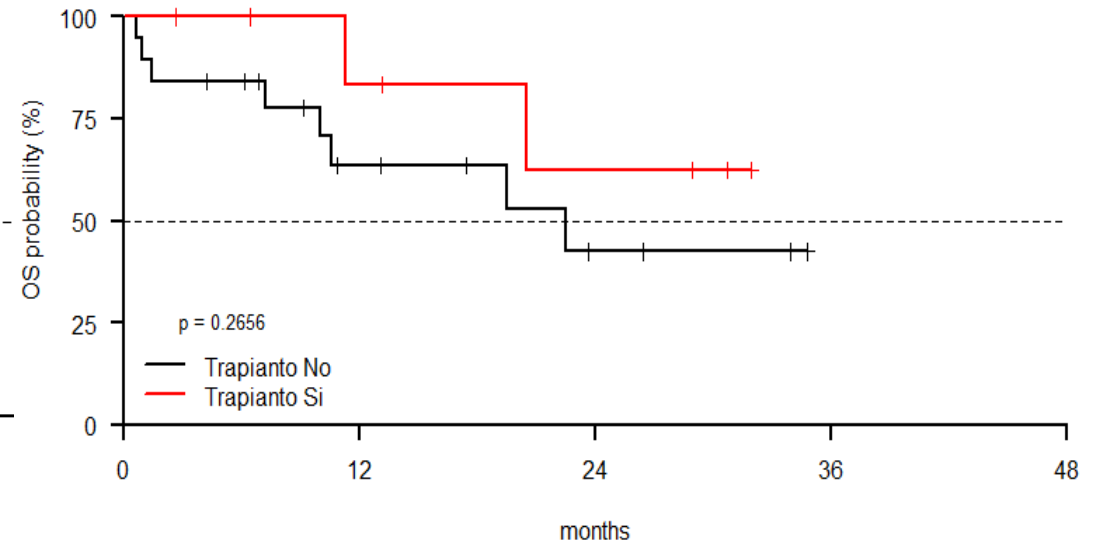
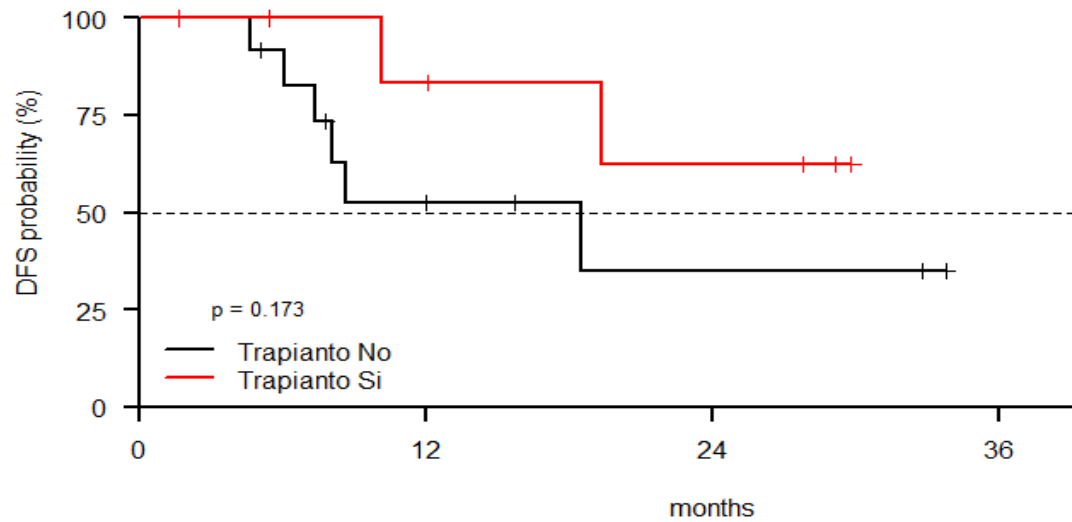
| 28/88 (31.8%) <i>BCR/ABL1</i> -like cases | | <i>BCR/ABL1</i> -like | Non- <i>BCR/ABL1</i> -like | p-value |
|---|------------------|-----------------------|----------------------------|---------|
| N | | 28 | 59 | |
| CR (%) | No CR | 7 (25.9) | 5 (8.5) | 0.044 |
| | CR | 20 (74.1) | 54 (91.5) | |
| TP1_MRD (%) | TP1 MRD positive | 14 (77.8) | 19 (41.3) | 0.012 |
| TP2_MRD (%) | TP2 MRD positive | 9 (52.9) | 9 (20.0) | 0.029 |
| TP3_MRD (%) | TP3 MRD positive | 5 (41.7) | 5 (13.5) | 0.05 |

A *BCR/ABL1*-like status is characterized by a lower CR rate, MRD persistence and shorter survival also in a pediatric-oriented and MRD-driven clinical trial.

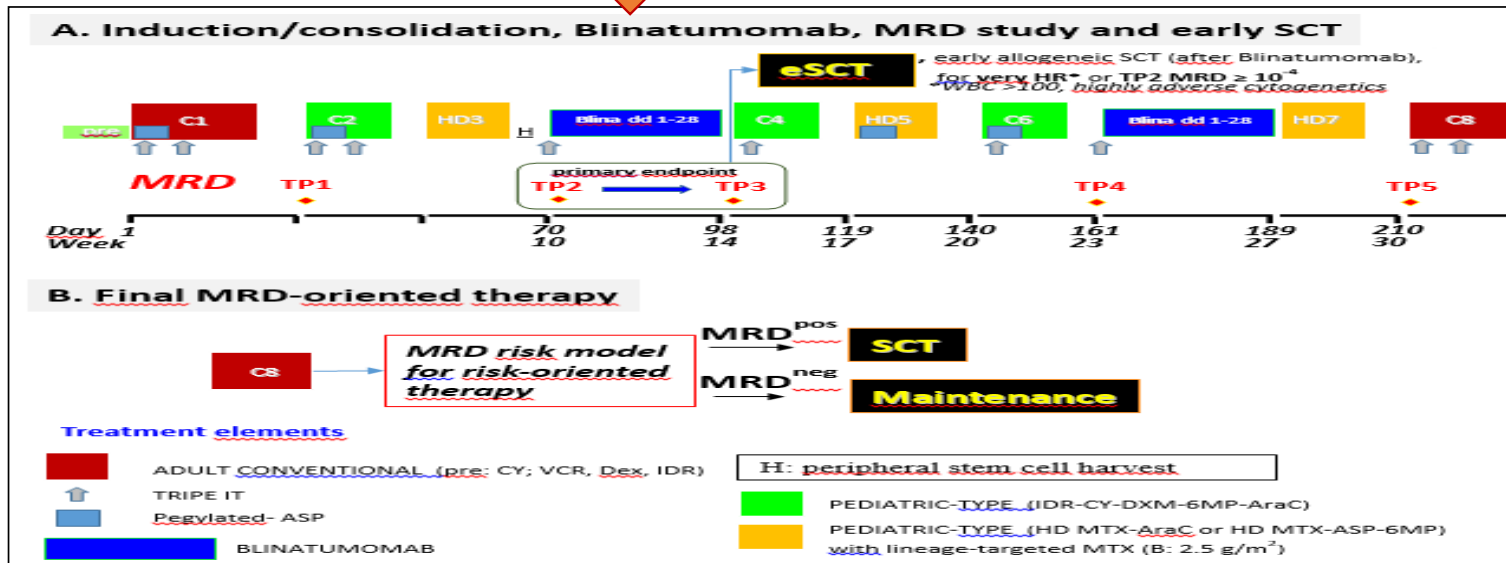
The prognostic role of the *BCR/ABL1*-like status is independent from the other clinico-biologic and genetic features.

Role of transplant within GIMEMA 1913 in BCR/ABL1-like patients

| | Transplant | No transplant | p |
|-------------------|------------|---------------|-------|
| BCR/ABL1-like | 9 | 11 | 0.003 |
| Non BCR/ABL1-like | 6 | 48 | |



Ph-like ALL, MRD and monoclonal antibodies



Blinatumomab effective in eradicating MRD: 10/25 patients MRD-positive after early consolidation and all became MRD-negative (Bassan et al, EHA 2021)

Inotuzumab reported to be effective in these patients

Conclusions

- Early recognition should be always carried out (possibly, at diagnosis)
- MRD often positive; no enrichment among various subgroups
- Allo-SCT should be carried out
- TKIs might/should be incorporated, at least, in MRD+ positive patients

Acknowledgments

Monica Messina

Alessia Lauretti

Alfonso Picicocchi

Akram Taherinasab

Martina Canichella

Antonella Vitale

Anna Guarini

Robin Foà

