

# **Case SH2017-0119: A 22-year-old male with a Ph-like mixed phenotype acute leukemia (B/myeloid)**

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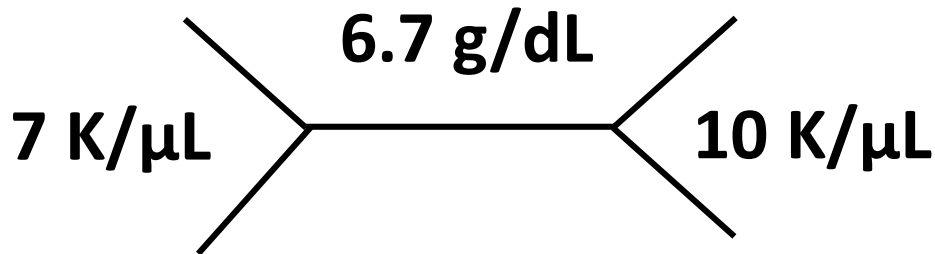
Lina Shao

University of Michigan

## Clinical history

- A 22-year-old male with a one-month history of worsening dyspnea on exertion and subsequent headaches, oral lesions, and gingival bleeding.

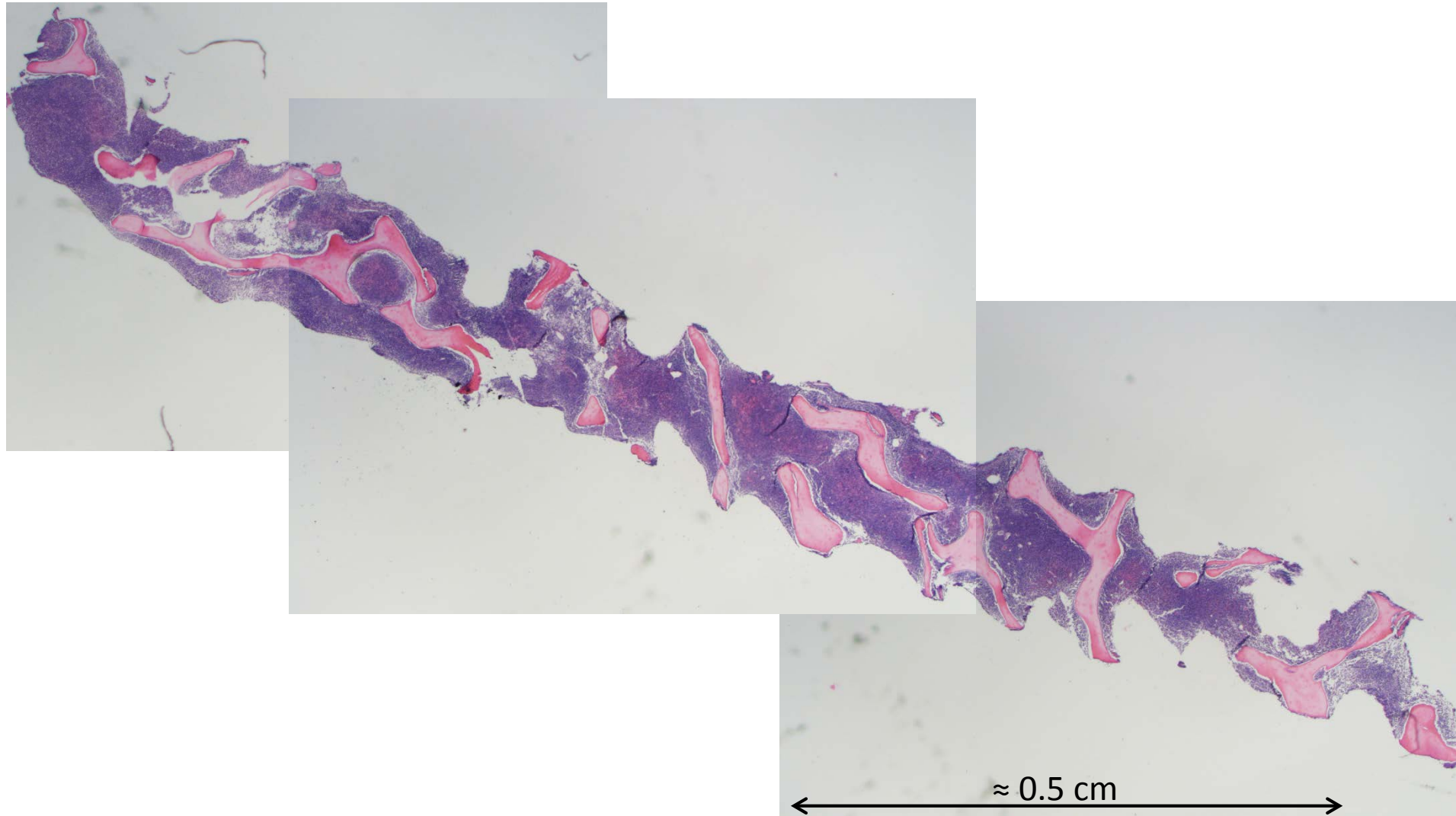
## Laboratory findings



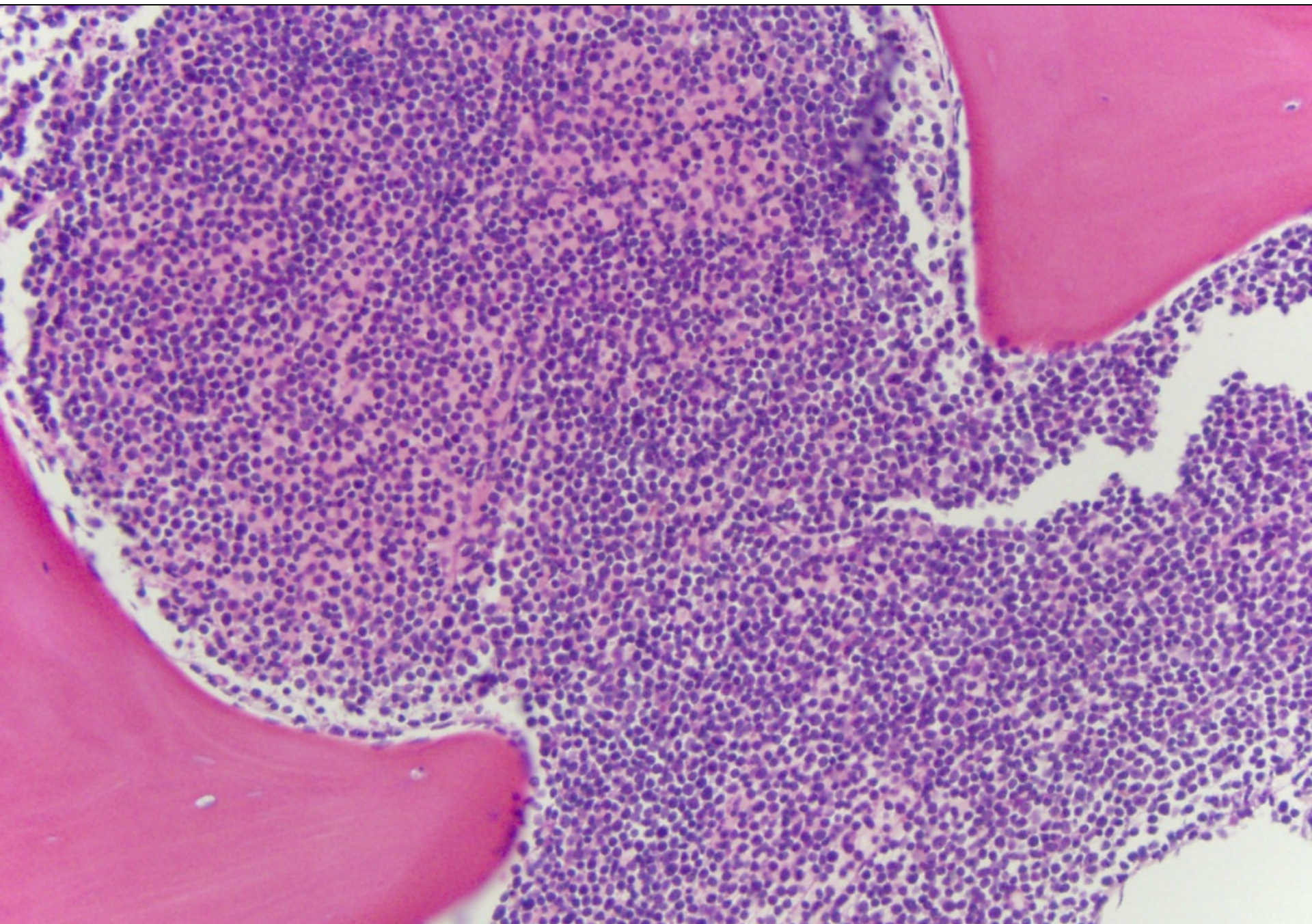
- Peripheral blood differential count:

Blasts 41%  
Neutrophils 19%  
Lymphocytes 36%  
Monocytes 3%  
Eosinophils 1%

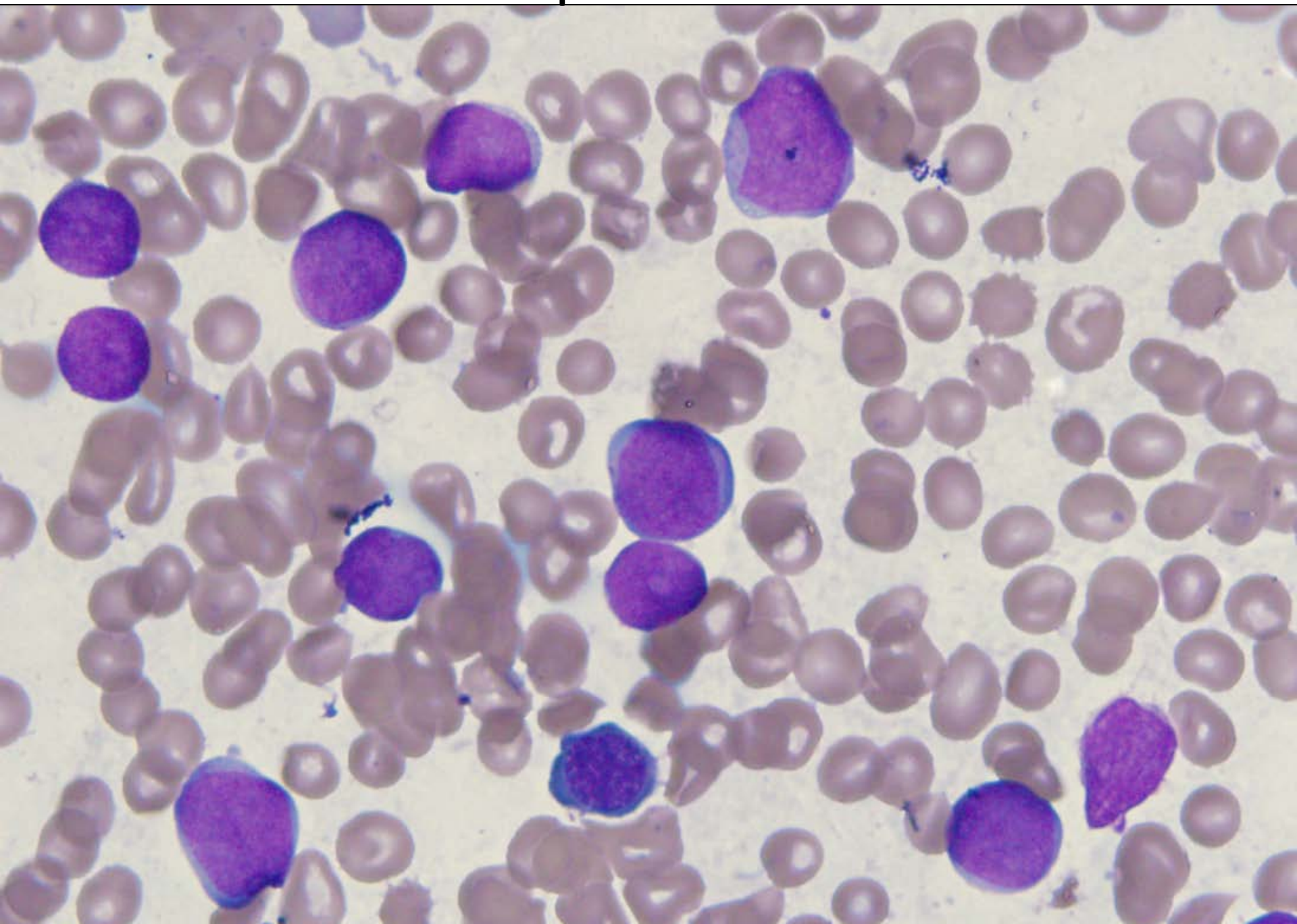
# Core biopsy



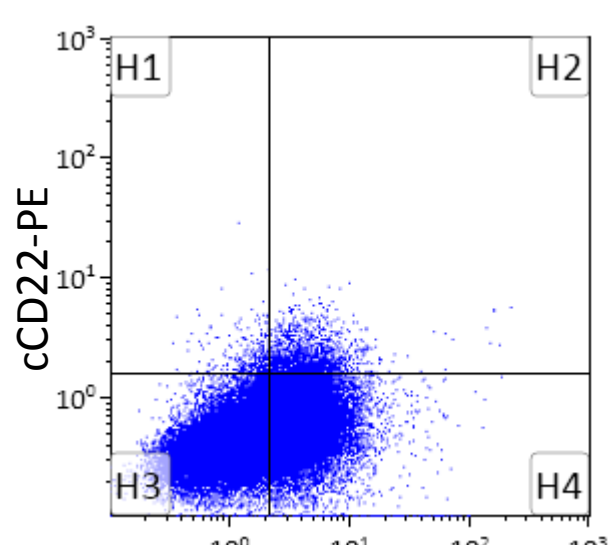
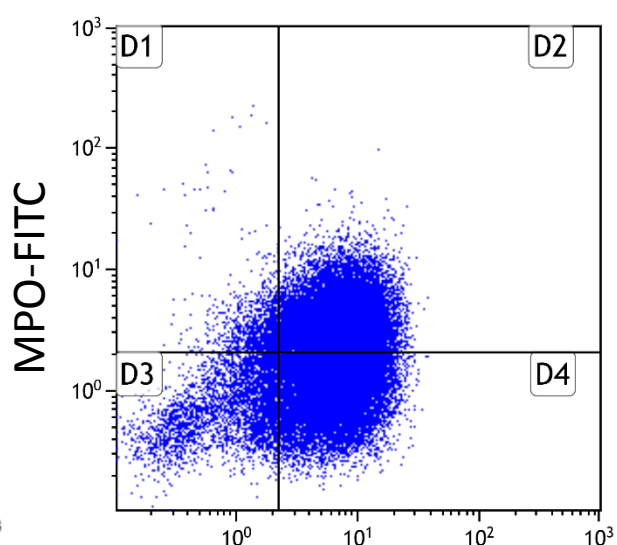
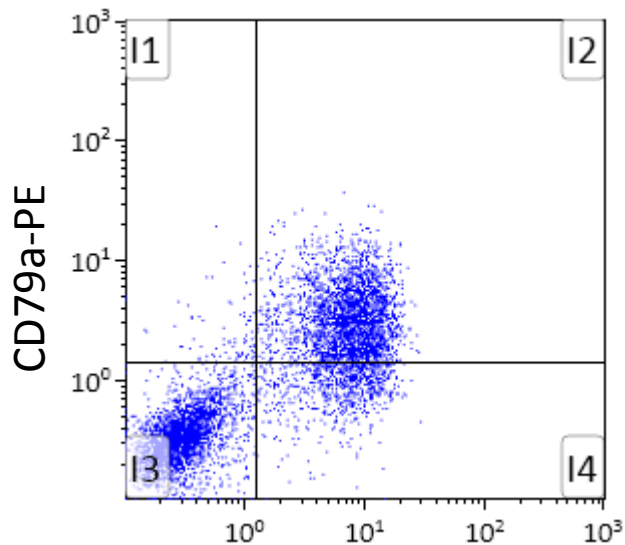
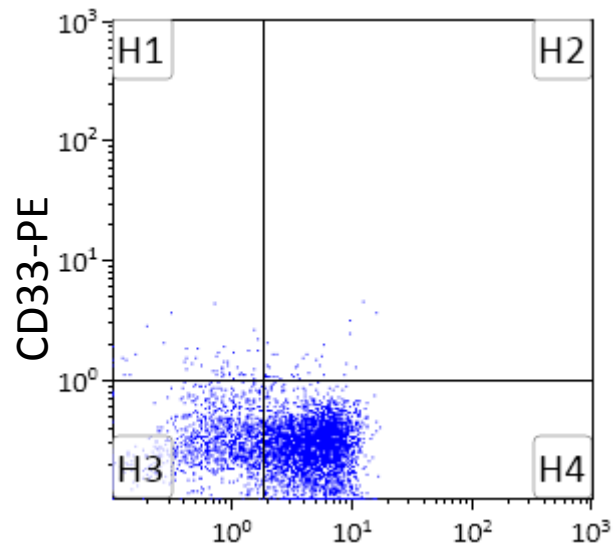
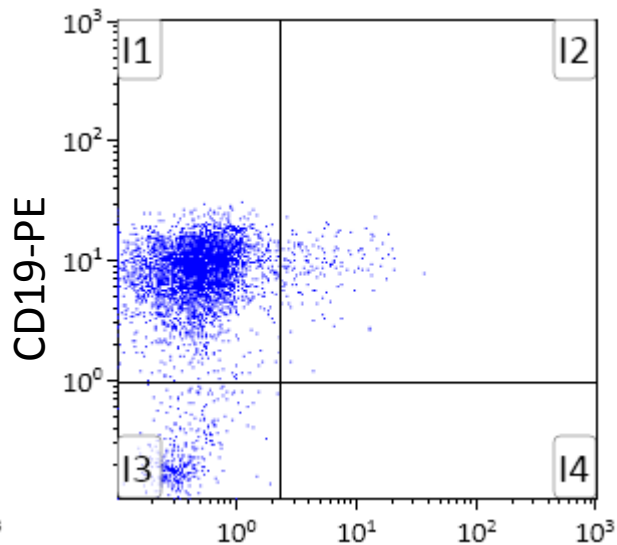
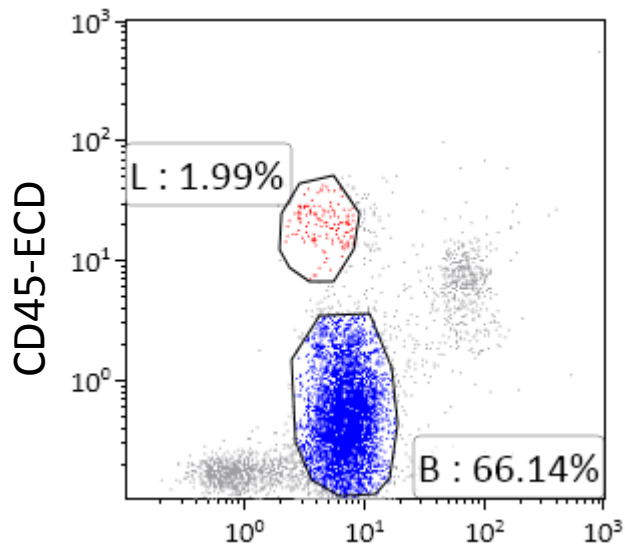
# Core biopsy



# Aspirate smear



# Flow cytometry



# Immunophenotypic summary

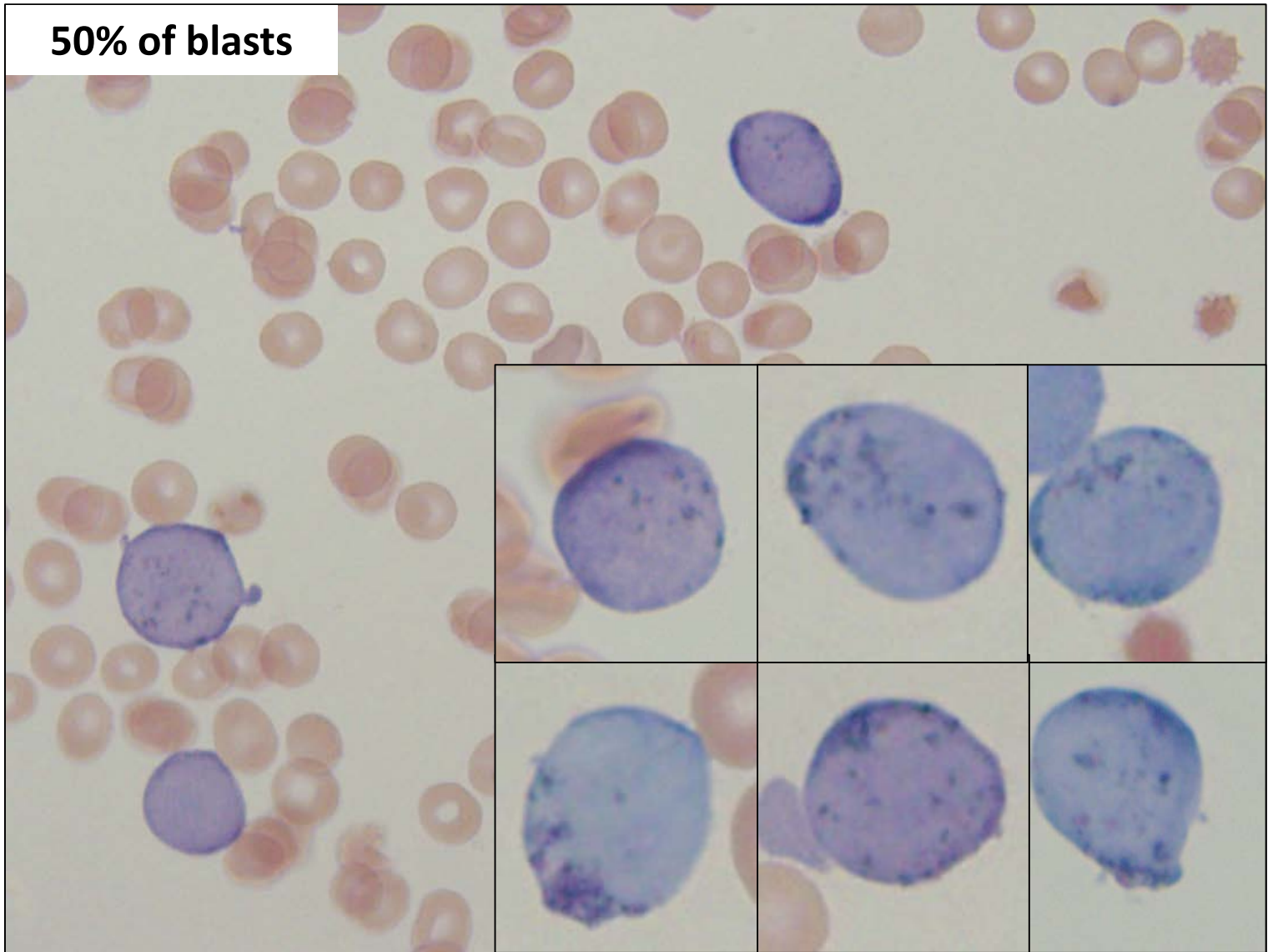
## Blasts (positive flow cytometry markers)

- CD19 (moderate)
- CD79a (moderate)
- CD10 (dim, very minor subset)
- CD20 (dim, minor subset)
- CD22 (surface/cytoplasmic, dim to negative)
- CD34 (dim, subset)
- CD45 (dim to negative)
- TdT (moderate, subset)
- MPO (dim, subset)

**Negative:** CD11c, CD13, CD14, CD33, CD38, CD56, CD117, T cell markers

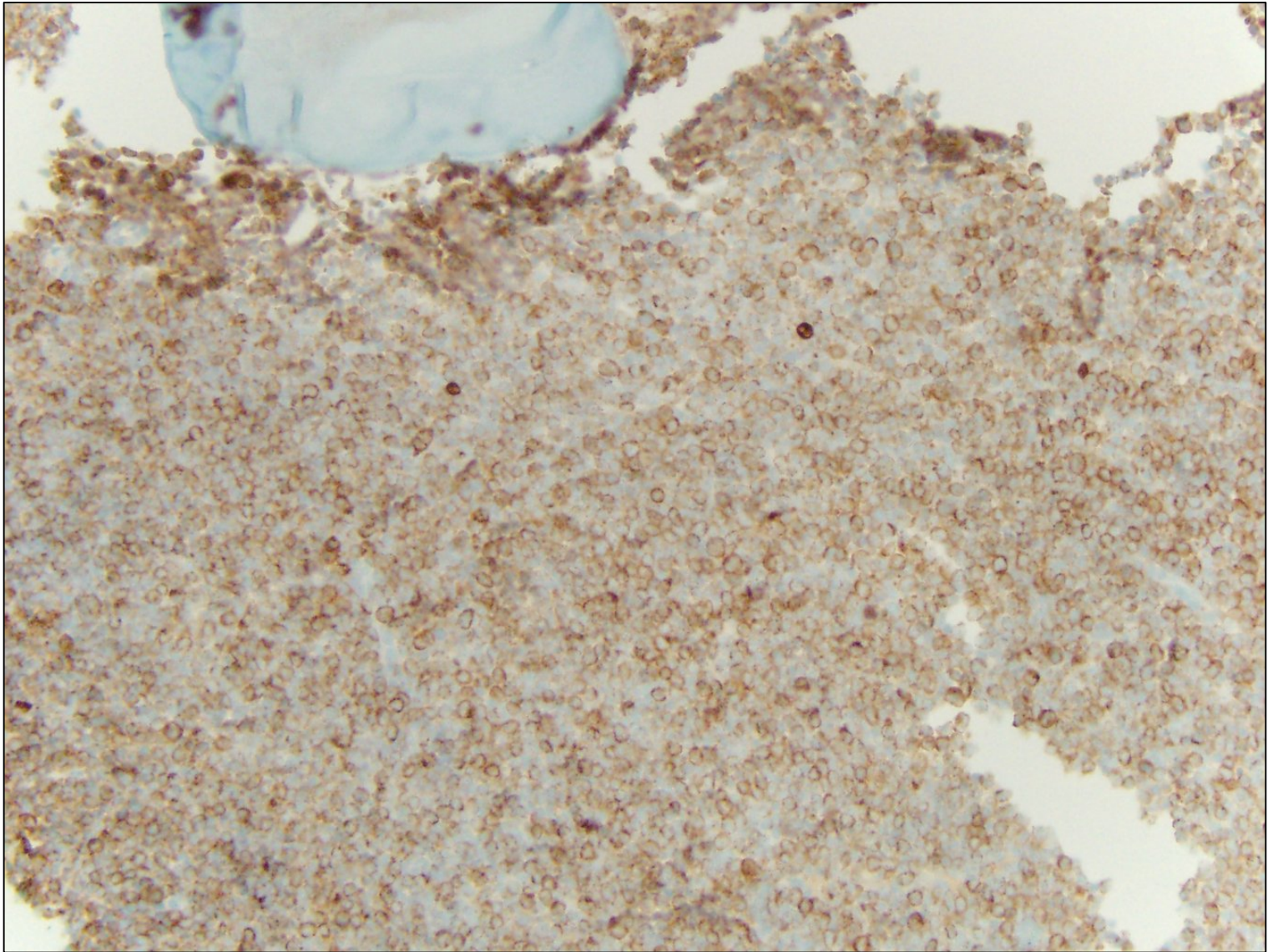
# Aspirate smear – Myeloperoxidase cytochemistry

50% of blasts





# Core biopsy – Myeloperoxidase immunohistochemistry



# Panel differential diagnosis

Mixed phenotype acute leukemia, B/myeloid

vs.

B-ALL (with isolated MPO expression)

# THE UPDATED WHO CLASSIFICATION OF HEMATOLOGICAL MALIGNANCIES

## The 2016 revision to the World Health Organization classification of myeloid neoplasms and acute leukemia

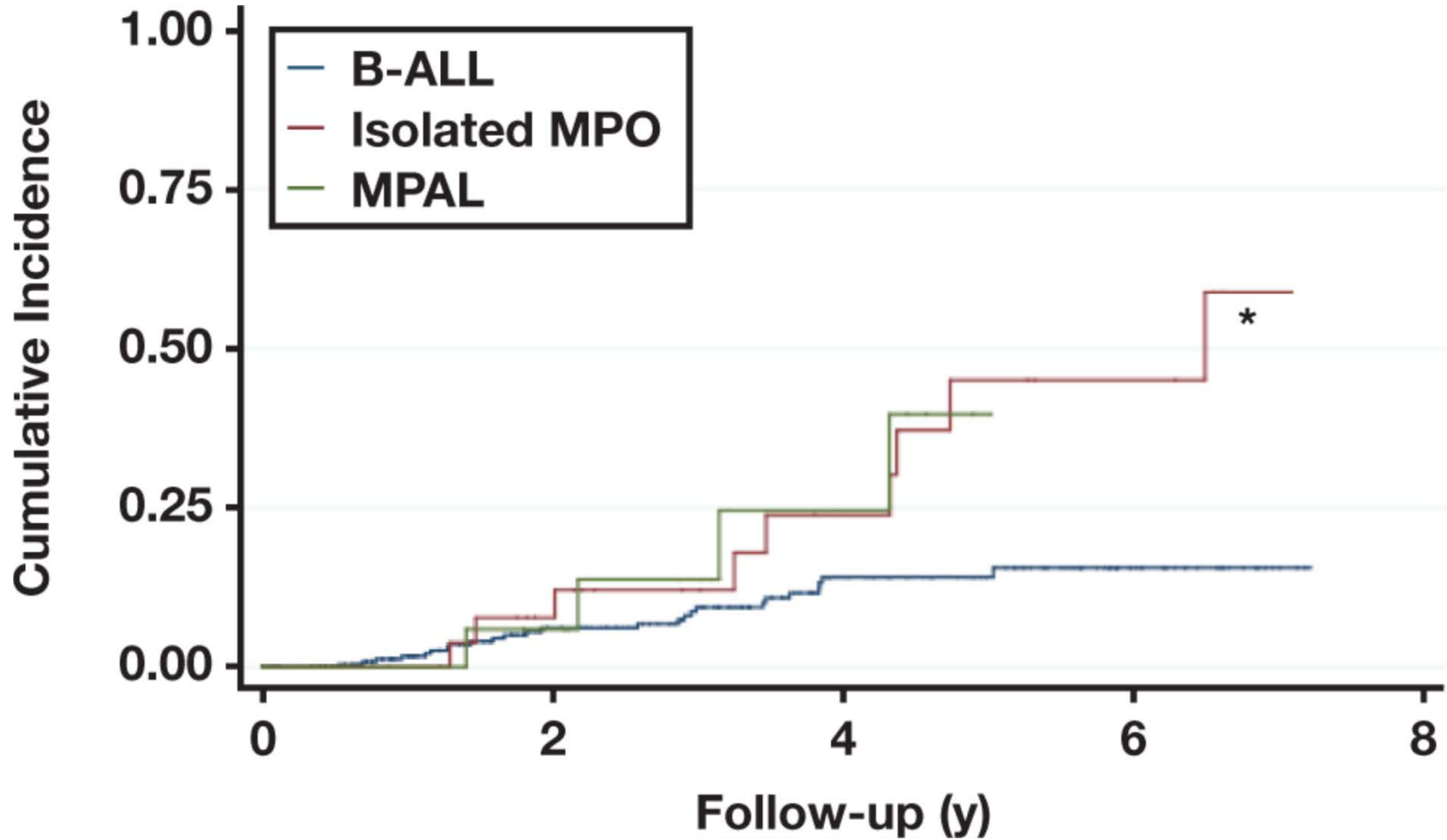
Daniel A. Arber,<sup>1</sup> Attilio Orazi,<sup>2</sup> Robert Hasserjian,<sup>3</sup> Jürgen Thiele,<sup>4</sup> Michael J. Borowitz,<sup>5</sup> Michelle M. Le Beau,<sup>6</sup> Clara D. Bloomfield,<sup>7</sup> Mario Cazzola,<sup>8</sup> and James W. Vardiman<sup>9</sup>

Table 19. Criteria for lineage assignment for a diagnosis of MPAL

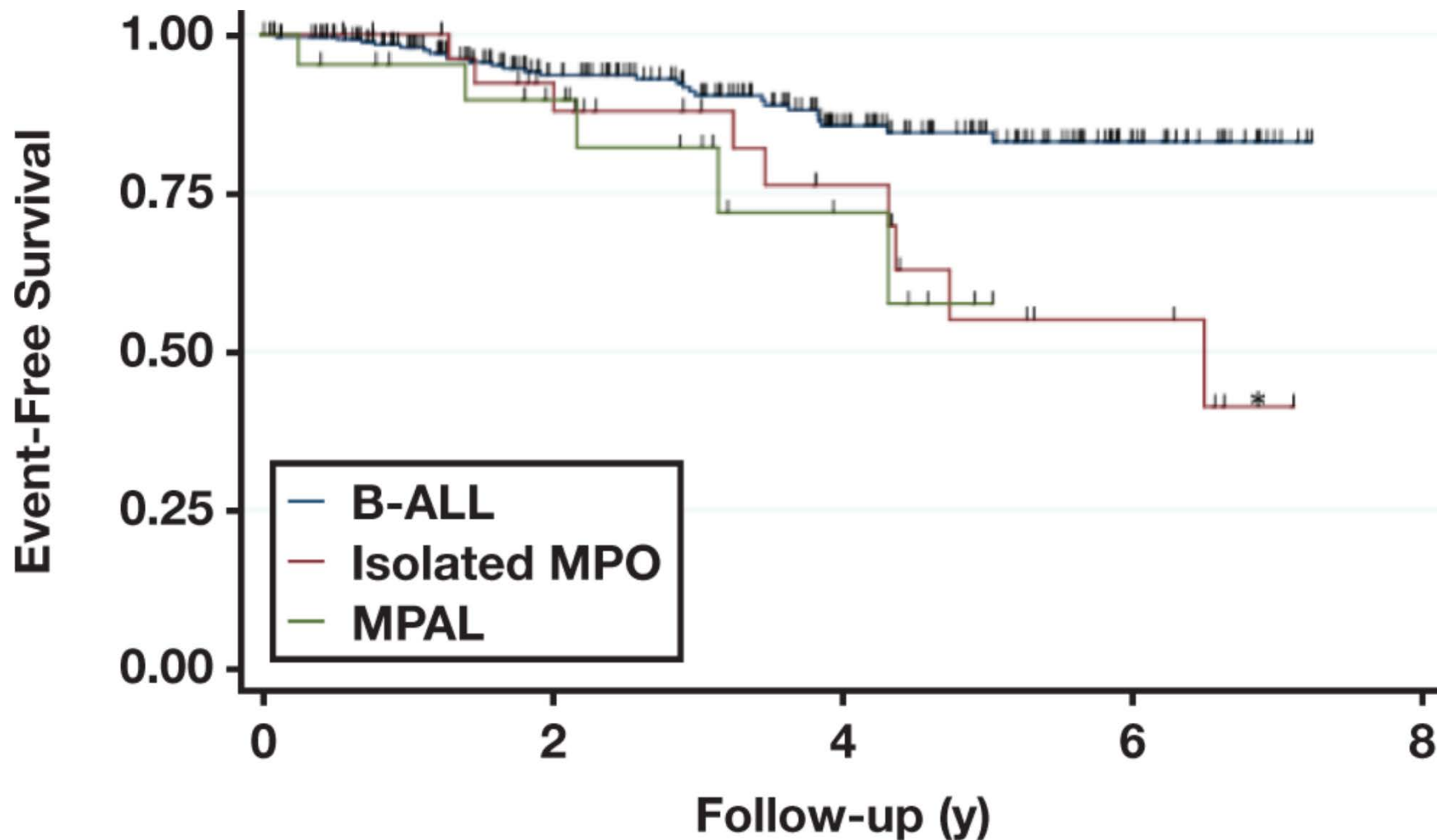
ALL, but only for MPAL. It is also now recognized that some cases of otherwise typical B-ALL with homogeneous expression of lymphoid markers on a single blast population may express low-level myeloperoxidase using immunophenotypic methods without other evidence of myeloid differentiation. Because the clinical significance of this finding has not yet been established, it is recommended that care be taken before making a diagnosis of B/myeloid MPAL when low-intensity myeloperoxidase (MPO) is the only myeloid-associated feature. Multiparameter flow cytometry is the method of choice for recognizing MPAL; even when there

cytoplasmic CD22, or CD10

# B-ALL with isolated MPO expression shows higher cumulative incidence of relapse



# B-ALL with isolated MPO expression shows worse event-free survival



# Cytogenetics/Molecular analysis

## Karyotype:

46,XY[20] (normal male karyotype)

## FISH:

Negative for *BCR/ABL1* fusion and *KMT2A (MLL)* rearrangement.

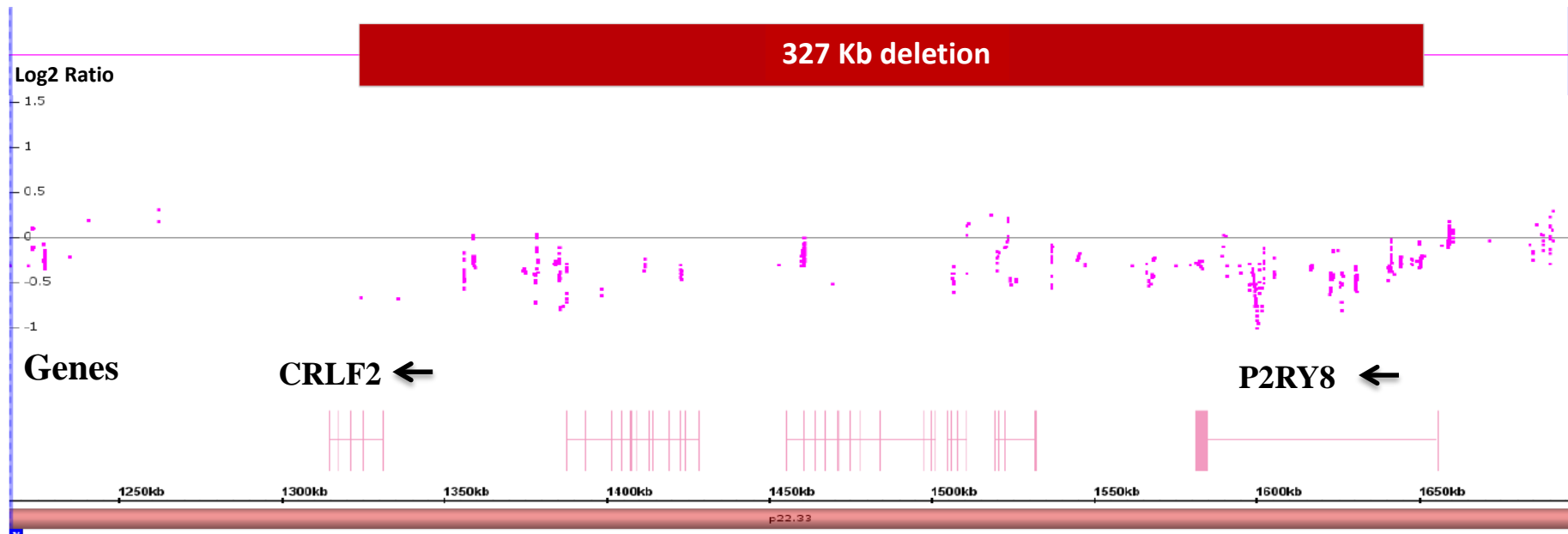
## Molecular genetics:

Negative for *JAK2* V617F mutation.

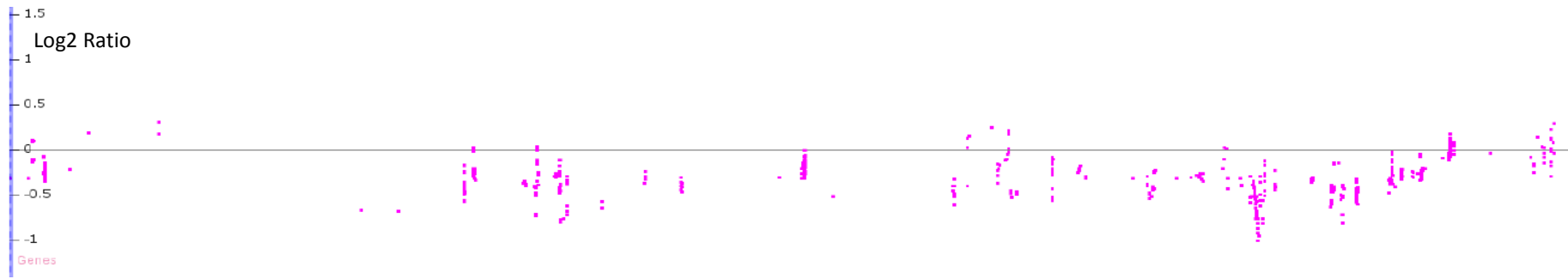
**Positive for *JAK1* c.1972G>T;p.V658F mutation.**

**Positive for *CDKN2A* c.151-6delTinsCCAGGGG mutation.**

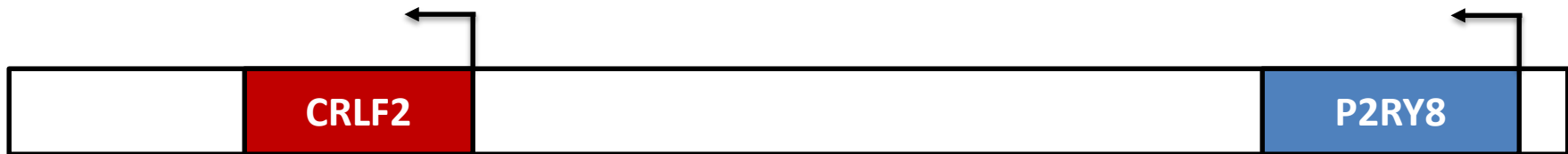
## Cytogenomic array:



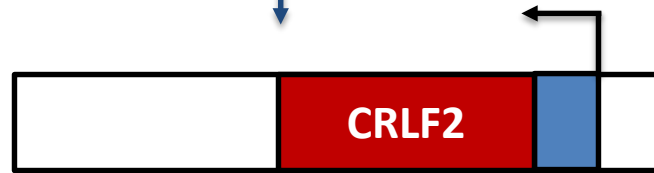
# Formation of *P2RY8-CRLF2* fusion in pseudoautosomal region 1 (PAR1)



**327 Kb deletion**



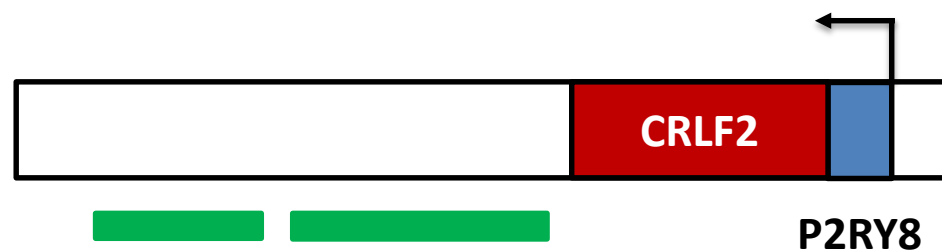
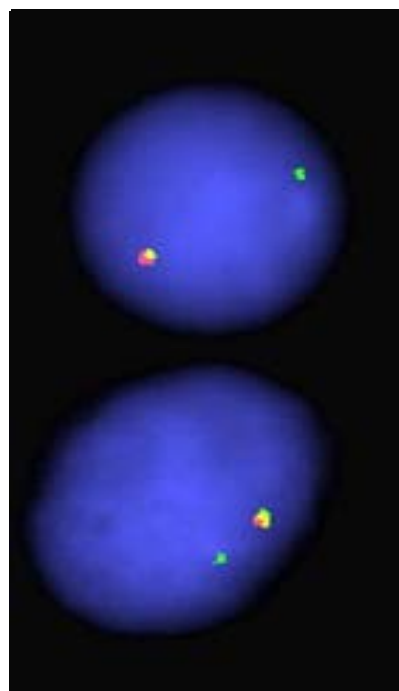
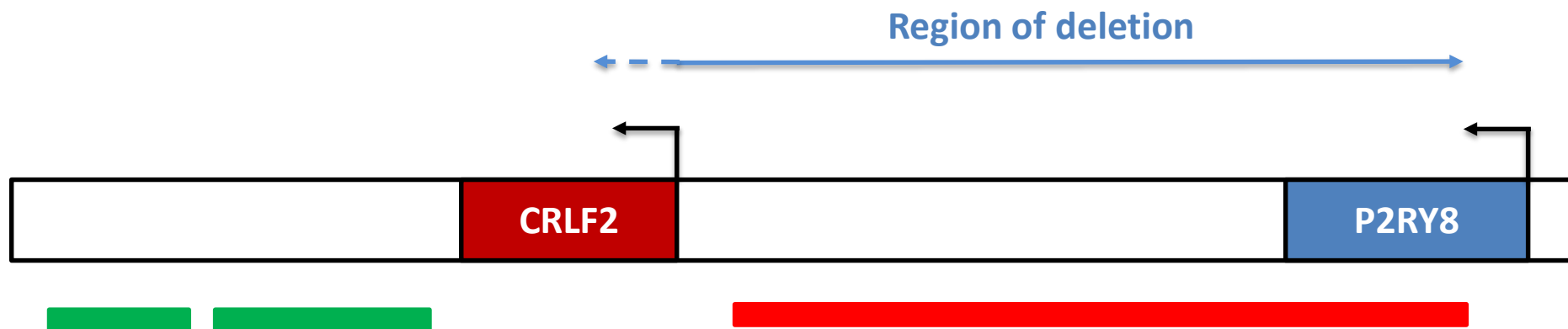
**Resultant fusion**



**P2RY8**

# Formation of *P2RY8-CRLF2* fusion in pseudoautosomal region 1 (PAR1)

FISH *CRLF2* breakapart probe



Loss of red-orange signal



## Historical context of BCR-ABL1-like vs. Ph-like B-ALL

Den Boer et al, The Lancet  
February 2009 Vol 10

COALL discovery cohort  
DCOG validation cohort

**Hierarchical clustering (HC)** of 110 gene probe sets identified to predict the major pediatric ALL subtypes (T-cell ALL, *ETV6-RUNX1*, high-hyperdiploidy, *TCF3* or *MLL*-rearranged, *BCR-ABL1*). “*BCR-ABL1*-like”

Mullighan et al, N Engl J  
Med 2009; 360:470-480

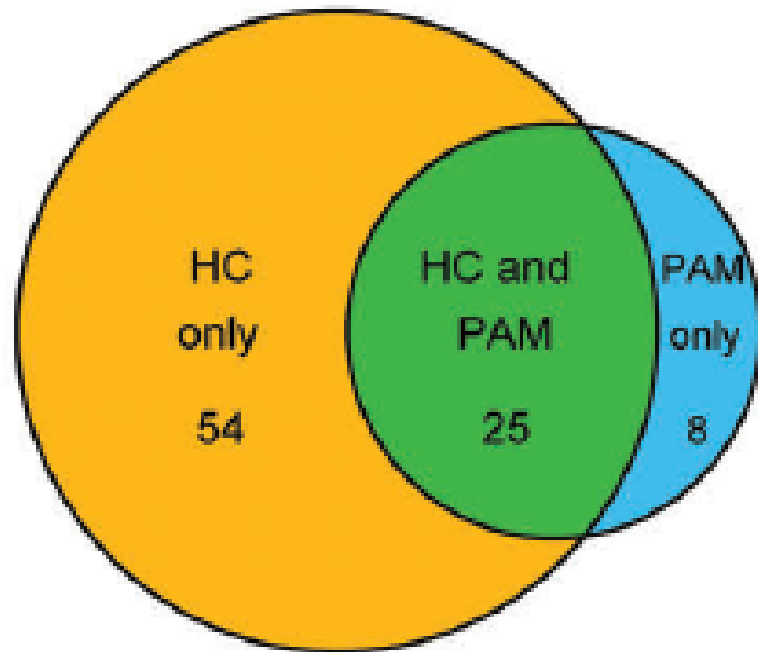
COG P9906 discovery cohort  
St. Jude validation cohort

“Ph-like signature” is based on the **prediction analysis of microarrays (PAM)** classifier consisting of 257 gene probe sets trained on *BCR-ABL1*-positive cases.

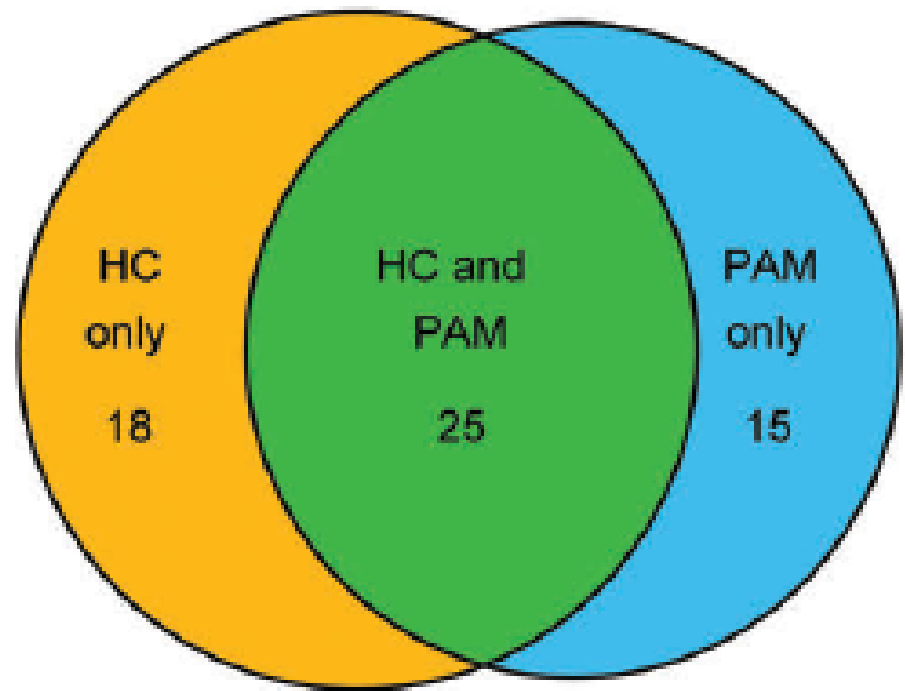
# **BCR-ABL1-like cases in pediatric acute lymphoblastic leukemia: a comparison between DCOG/Erasmus MC and COG/St. Jude signatures**

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## **Total DCOG/COALL**



## **COG P9906**



## ORIGINAL ARTICLE

# 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. Rowe, N. Santiago-Morales, I. Iacobucci, P.S. Fulton, J. Wen, M. Valentine, C. Cheng

VOLUME 35 • NUMBER 4 • FEBRUARY 1, 2017

JOURNAL OF CLINICAL ONCOLOGY

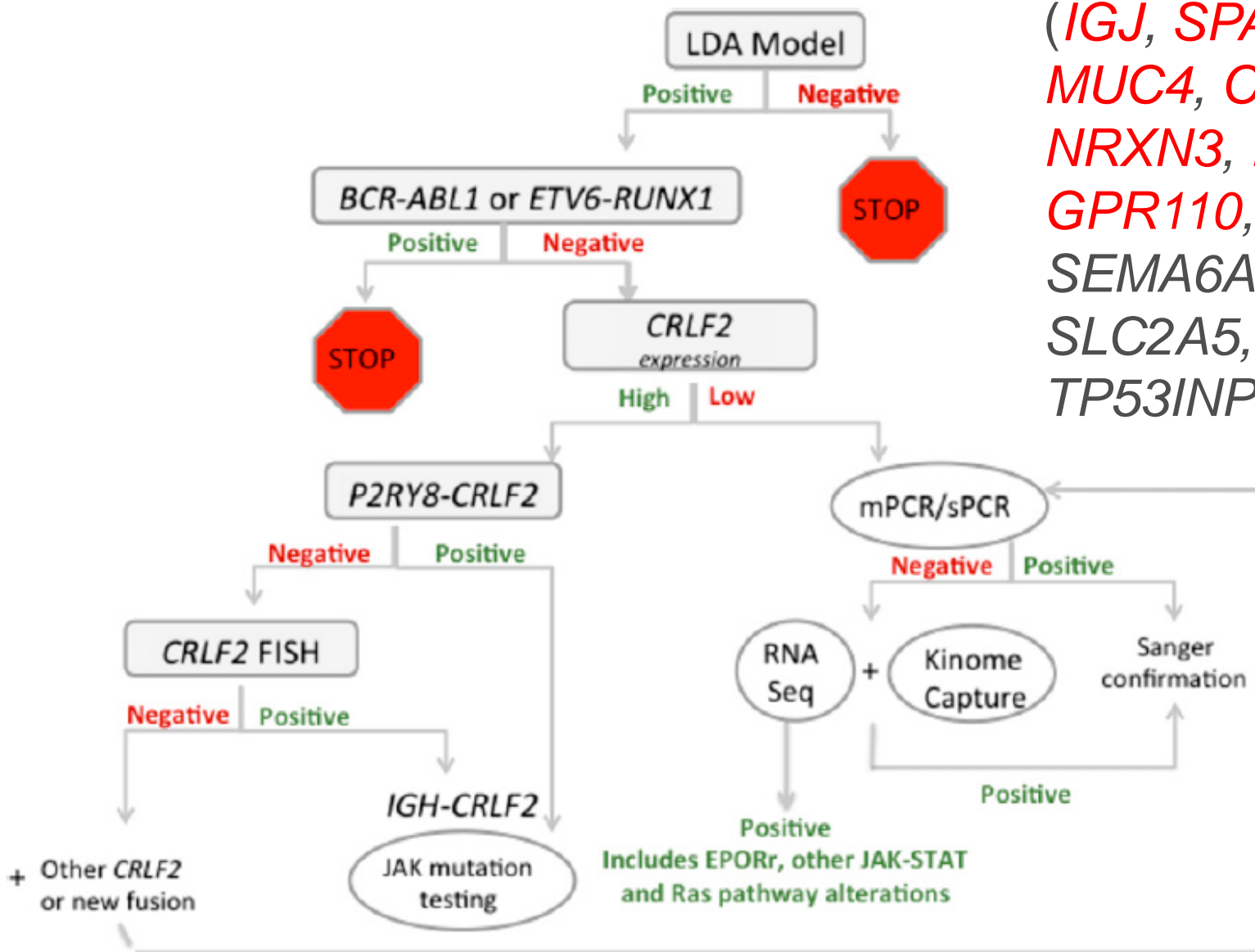
ORIGINAL REPORT

## High Frequency and Poor Outcome of Philadelphia Chromosome–Like Acute Lymphoblastic Leukemia in Adults

*Kathryn G. Roberts, Zhaohui Gu, Debbie Payne-Turner, Kelly McCastlain, Richard C. Harvey, I-Ming Chen, Deqing Pei, Ilaria Iacobucci, Marcus Valentine, Stanley B. Pounds, Lei Shi, Yongjin Li, Jinghui Zhang, Cheng Cheng, Alessandro Rambaldi, Manuela Tosi, Orietta Spinelli, Jerald P. Radich, Mark D. Minden, Jacob M. Rowe, Selina Luger, Mark R. Litzow, Martin S. Tallman, Peter H. Wiernik, Ravi Bhatia, Ibrahim Aldoss, Jessica Kohlschmidt, Krzysztof Mrózek, Guido Marcucci, Clara D. Bloomfield, Wendy Stock, Stephen Kornblau, Hagop M. Kantarjian, Marina Konopleva, Elisabeth Paietta, Cheryl L. Willman, and Charles G. Mullighan*

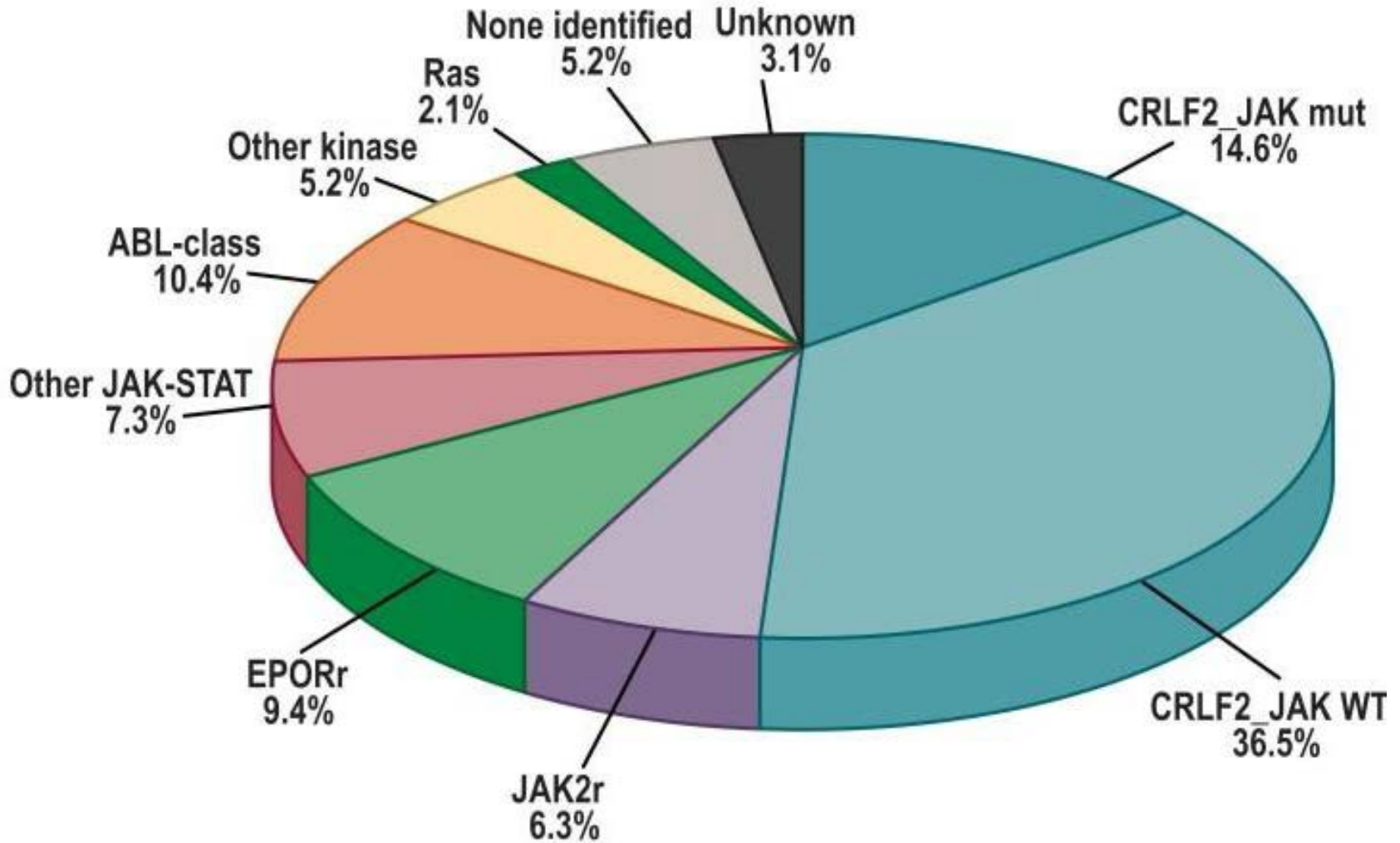
# Testing algorithm for Ph-like ALL in COG trials

(*IGJ, SPATS2L, MUC4, CRLF2, CA6, NRXN3, BMPR1B, GPR110, CHN2, SEMA6A, PON2, SLC2A5, S100Z, TP53INP1, IFITM1*).

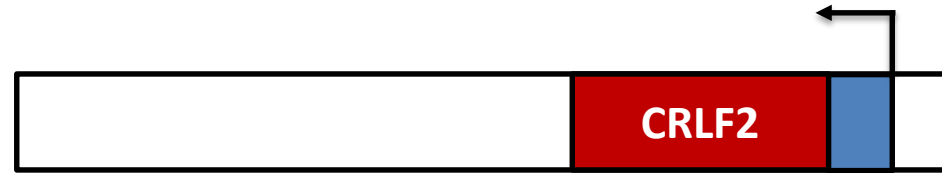


# CRLF2 is a frequent rearrangement seen in Ph-like B-ALL

Young Adult (21-39 yrs; n=96)



# *P2RY8-CRLF2* fusion leads to CRLF2 overexpression

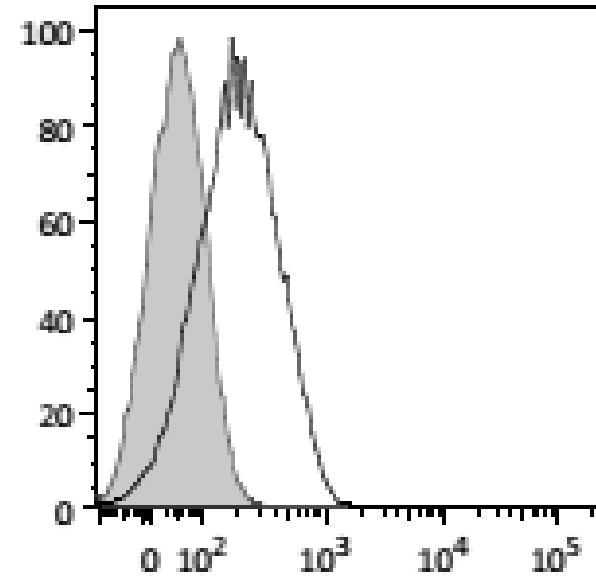
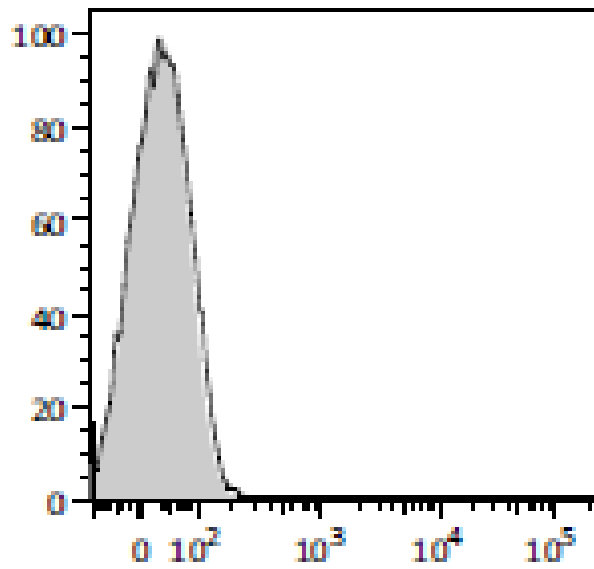


P2RY8

*CRLF2* wild-type

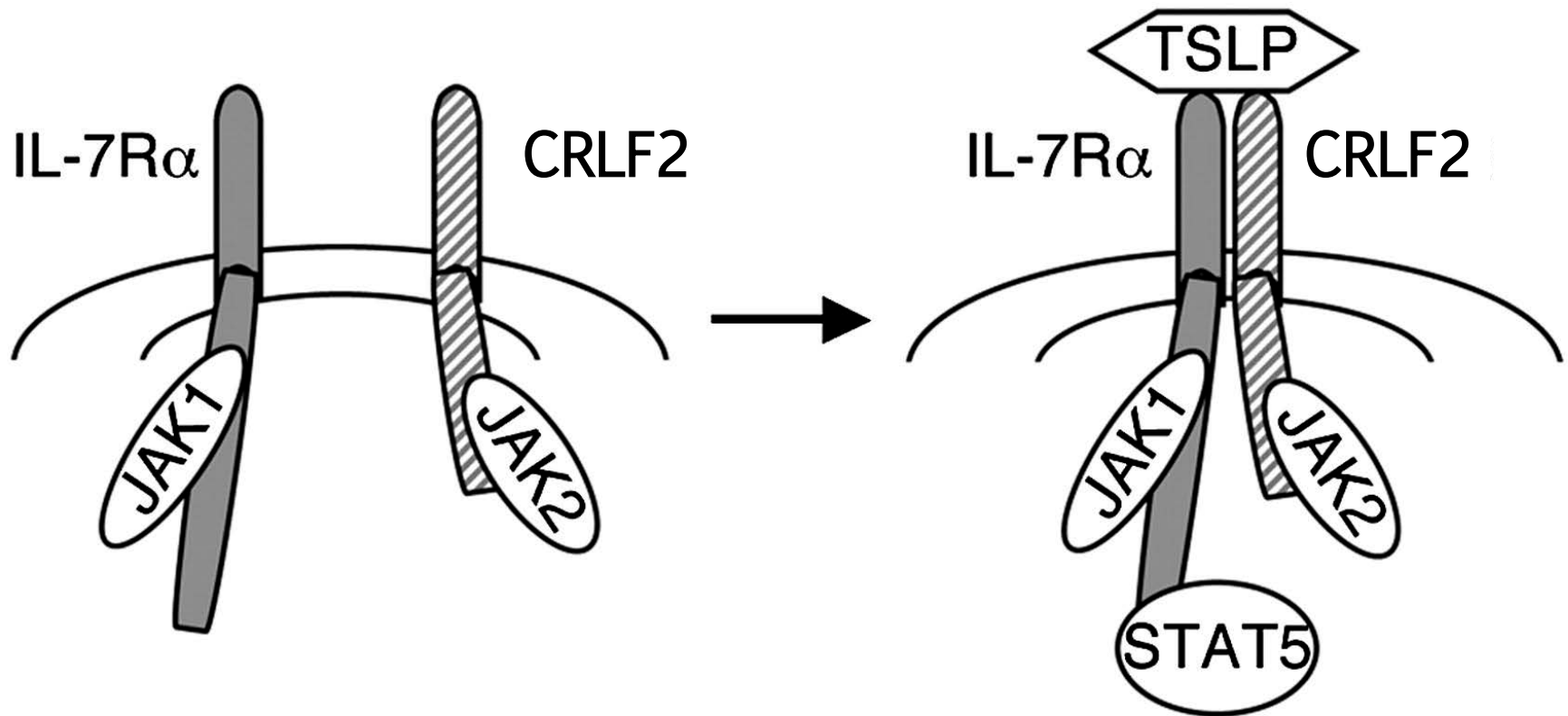
*IGH@-CRLF2*

Maximal Intensity

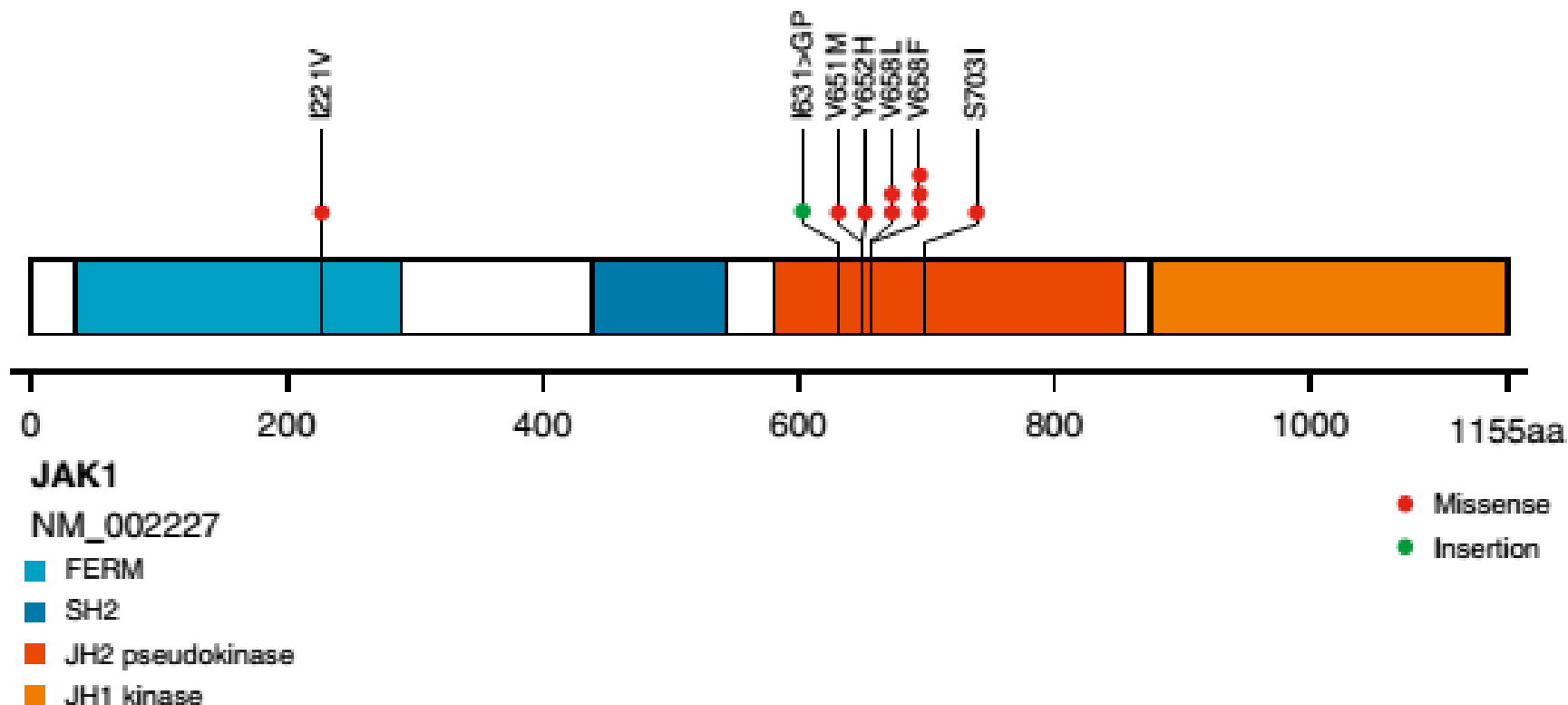


CRLF2

# CRLF2 acts upstream of JAK-STAT signaling

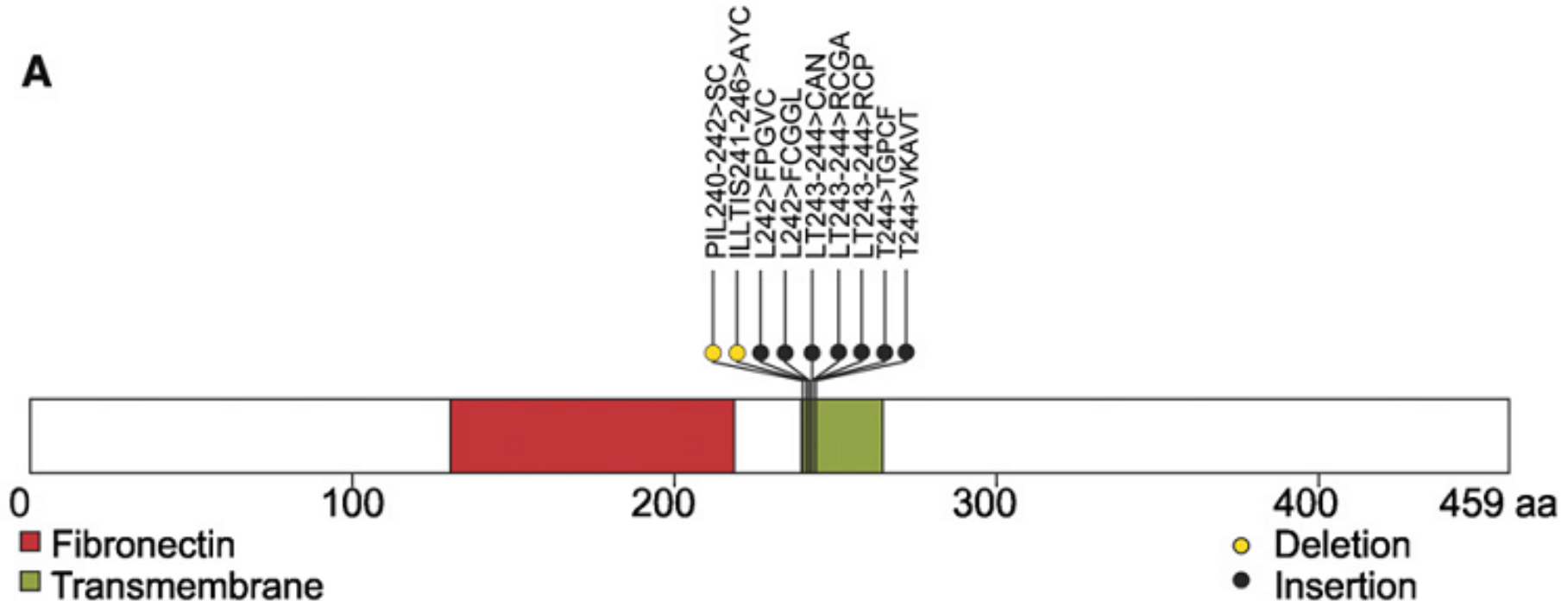


# CRLF2 rearranged cases have frequent concomitant JAK1 and JAK2 mutations





# IL-7R transmembrane domain mutations have also been reported in Ph-like B-ALLs



Co-occurrence with *CRLF2* rearrangement and *SH2B3* deletion has been reported.

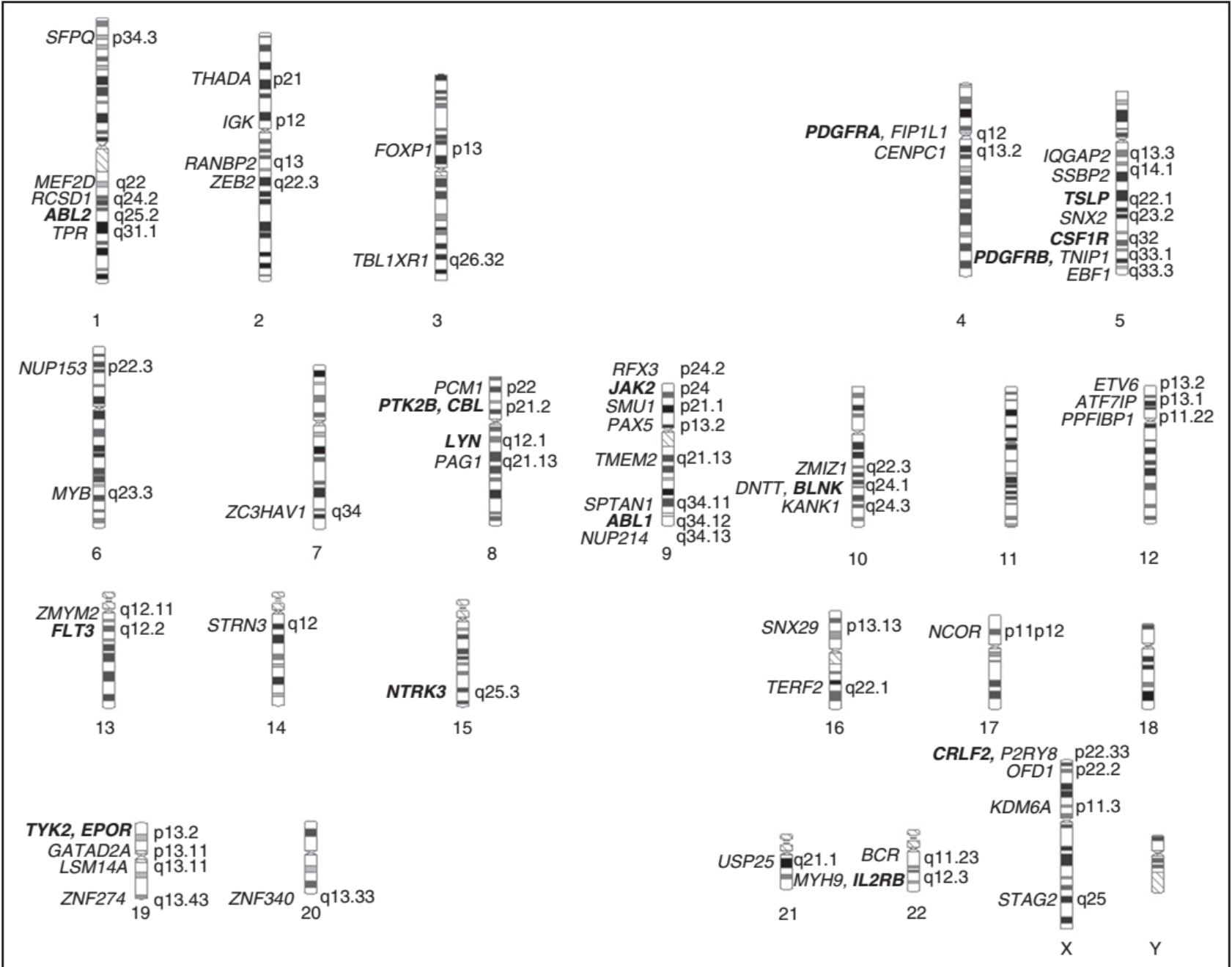
**Table 1. Kinase Fusions Identified in Ph-like Acute Lymphoblastic Leukemia.**

Kinase Gene	Tyrosine Kinase Inhibitor	Fusion Partners	Patients	5' Genes
				<i>number</i>
<i>ABL1</i>	Dasatinib	6	14	<i>ETV6</i> , <sup>11</sup> <i>NUP214</i> , <sup>11</sup> <i>RCSD1</i> , <sup>11</sup> <i>RANBP2</i> , <sup>11</sup> <i>SNX2</i> , <sup>19</sup> <i>ZMIZ1</i> <sup>20</sup>
<i>ABL2</i>	Dasatinib	3	7	<i>PAG1</i> ,* <i>RCSD1</i> ,* <i>ZC3HAV1</i> *
<i>CSF1R</i>	Dasatinib	1	4	<i>SSBP2</i> *
<i>PDGFRB</i>	Dasatinib	4	11	<i>EBF1</i> , <sup>11-13</sup> <i>SSBP2</i> ,* <i>TNIP1</i> ,* <i>ZEB2</i> *
<i>CRLF2</i>	JAK2 inhibitor	2	30	<i>IGH</i> , <sup>21</sup> <i>P2RY8</i> <sup>22</sup>
<i>JAK2</i>	JAK2 inhibitor	10	19	<i>ATF7IP</i> ,* <i>BCR</i> , <sup>11</sup> <i>EBF1</i> ,* <i>ETV6</i> , <sup>23</sup> <i>PAX5</i> , <sup>11</sup> <i>PPFIBP1</i> ,* <i>SSBP2</i> , <sup>24</sup> <i>STRN3</i> , <sup>11</sup> <i>TERF2</i> ,* <i>TPR</i> *
<i>EPOR</i>	JAK2 inhibitor	2	9	<i>IGH</i> , <sup>11</sup> <i>IGK</i> *
<i>DGKH</i>	Unknown	1	1	<i>ZFAND3</i> *
<i>IL2RB</i>	JAK1 inhibitor, JAK3 inhibitor, or both	1	1	<i>MYH9</i> *
<i>NTRK3</i>	Crizotinib	1	1	<i>ETV6</i> <sup>25-27</sup> †
<i>PTK2B</i>	FAK inhibitor	2	1	<i>KDM6A</i> ,* <i>STAG2</i> *
<i>TSLP</i>	JAK2 inhibitor	1	1	<i>IQGAP2</i> *
<i>TYK2</i>	TYK2 inhibitor	1	1	<i>MYB</i> *

\* The gene is a previously unreported fusion partner.

† *ETV6-NTRK3* has been reported in multiple cancers, including congenital fibrosarcoma<sup>25,26</sup> and secretory breast carcinoma,<sup>27</sup> but it has not previously been described in acute lymphoblastic leukemia.<sup>28,29</sup>

# There are MANY “actionable” gene fusions in Ph-like B-ALL



# Potential methodologies for identifying Ph-like leukemia

Low density array: currently used primarily in the research setting to initially screen for Ph-like ALL

FISH: breakpoint probes for known kinase genes would be simple, cost effective but limited in scope

RT-PCR: can screen for or confirm known fusions

Digital molecular barcoding platform

Capture-based RNA sequencing

Next-generation sequencing

## Clinical course and response to therapy

- C8811 (Larson) protocol with rituximab
  - 'B' arm of hyperCVAD together with ruxolitinib
  - Blinatumomab (1 cycle)
  - CLOVE chemotherapy (clofarabine, cyclophosphamide, etoposide).
  - Inotuzumab (1 cycle)
  - At various time points, considered for clinical trials that either closed, where he did not respond, or he was unable to enroll due to either diagnosis or condition (eg infection).
- Persistent disease (34% blasts)
  - Persistent disease (85% blasts)
  - Persistent disease (96% blasts)
  - Persistent disease (70% blasts)
  - Persistent disease (50% blasts)

# Summary

- Ph-like B-ALL with isolated MPO vs “Ph-like” MPAL
  - Potential for further clarification of diagnostic classification
- Ph-like B-ALL is characterized by kinase rearrangements of which *CRLF2* rearrangement is the most prevalent and is often associated with *JAK1/2* mutations.
  - Potential for targeted inhibition of JAK/STAT signaling pathways
- Optimal detection of Ph-like B-ALL? Multiple modalities are possible.
- Need for additional clinical trials investigating therapeutic efficacy of kinase inhibitors and optimization of treatment regimens in Ph-like B-lymphoblastic leukemia
  - Consider inclusion of cases of Ph-like MPAL

Final panel diagnosis:

B-lymphoblastic leukemia, *BCR-ABL1*-like

versus

Mixed phenotype acute leukemia,  
B/myeloid, not otherwise specified