

EMATOLOGIA DI LABORATORIO: percorsi diagnostici e obiettivi clinici.

Milano, 11-12 Novembre 2010

Aula Magna, Fondazione IRCCS Istituto Nazionale dei Tumori



**Citogenetica e genetica molecolare in
oncoematologia**

Francesco Cavazzini (Ferrara) 18:00-18:30

ARGOMENTI

Ruolo di: Citogenetica

**Citogenetica-Molecolare (ibridazione in situ
fluorescente, FISH)**

diagnosi prognosi monitoraggio della malattia minima

**Citogenetica e Citogenetica-Molecolare
come elementi di diagnosi**

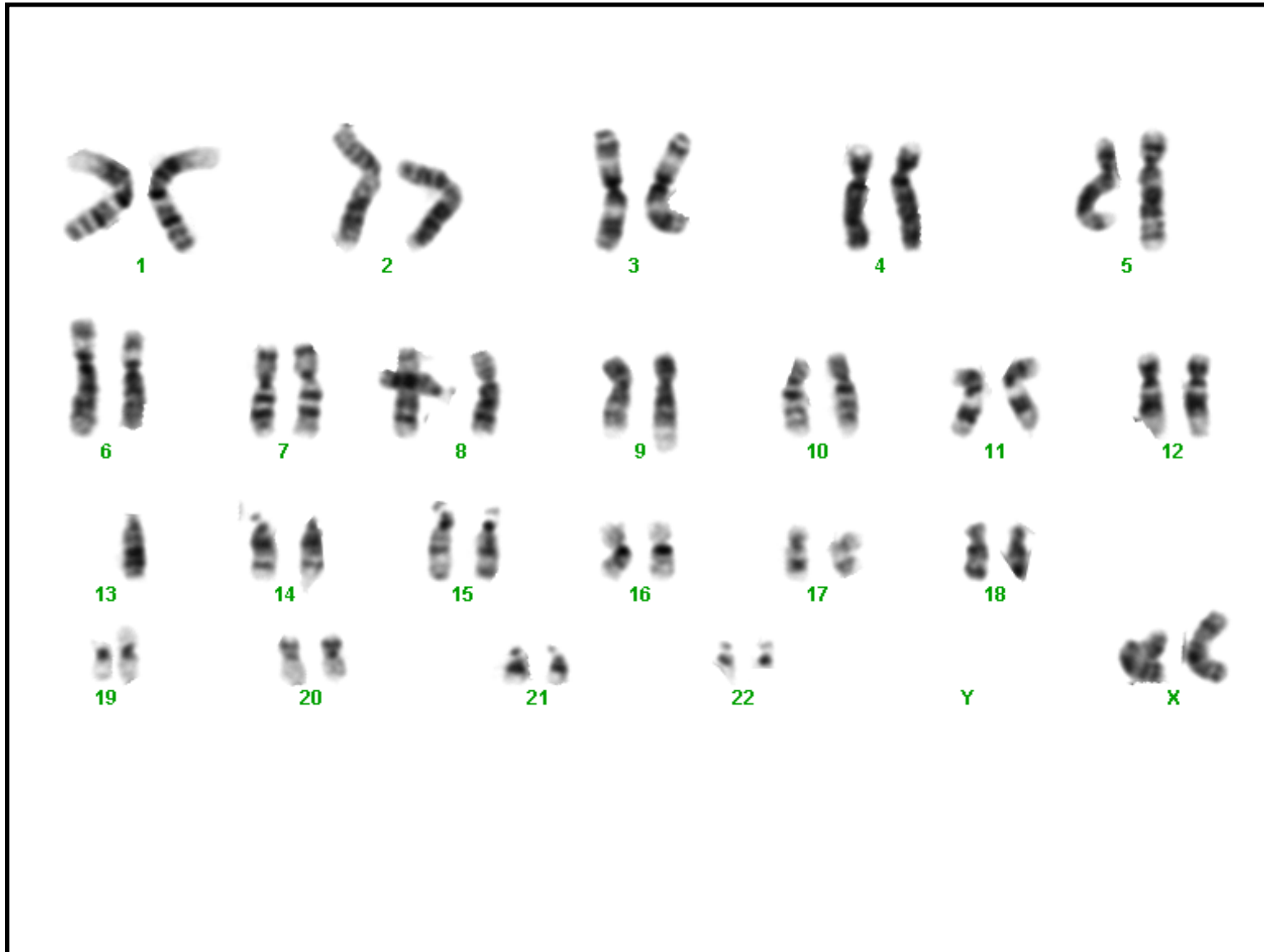
**Citogenetica e Citogenetica-Molecolare
come componenti integrativi di
sistemi diagnostico-prognostici**

Indagine citogenetica in ematologia

- Indagine di laboratorio che rileva la presenza di anomalie cromosomiche nel cariotipo delle cellule patologiche
 - Riconosce le anomalie “primarie” (presenti in tutte le cellule anomale), responsabili delle fasi precoci di trasformazione
 - Identifica le anomalie “secondarie” responsabili delle fasi di evoluzione clonale
 - Deve identificare le lesioni non rilevanti per la patogenesi della malattia in quanto semplice espressione della instabilità genetica
-

Cosa serve per l'indagine citogenetica in ematologia

- 5-20 X 10⁶ cellule vive, messe in coltura sterilmente entro 24 ore dal prelievo
 - Le cellule devono essere in grado di dividersi (midollo osseo, sangue periferico se contaminato da cellule immature, linfonodi, milza, biopsie tissutali patologiche)
 - Passaggi tecnici relativamente semplici (blocco delle metafasi, ipotonizzazione, allestimento dei vetrini, bandeggio)
 - Sistema di analisi automatizzata del cariotipo (economicamente vantaggioso per centri medio-grandi)
-



Ciccone M, Cavazzini F, Castoldi G.
Cytogenetics and Molecular cytogenetics in diagnosis and prognosis of hematologic malignancies. Trends in Medicine, Jan 2006, Vol 6, N 1

FISH: Fluorescence In Situ Hybridization

Metodica di **citogenetica molecolare**

Indaga aspetti
dell'assetto genomico
della cellula

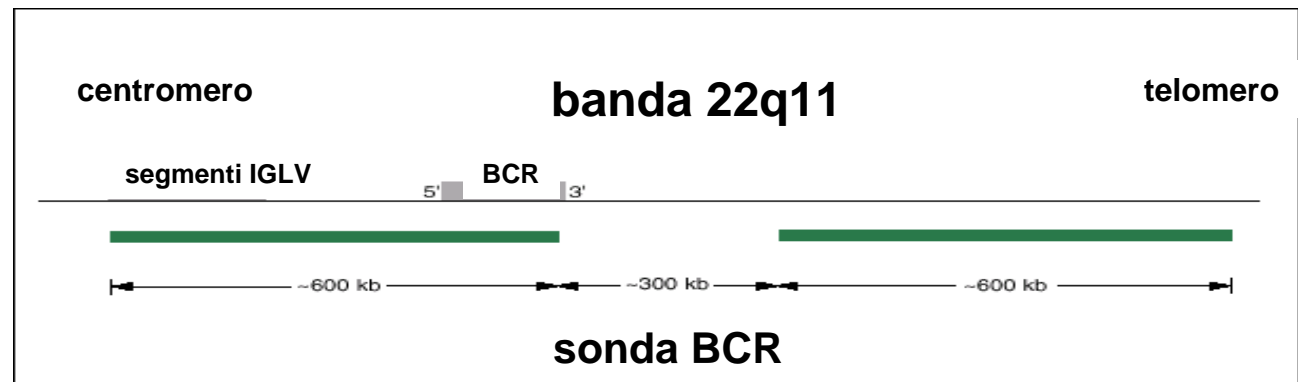
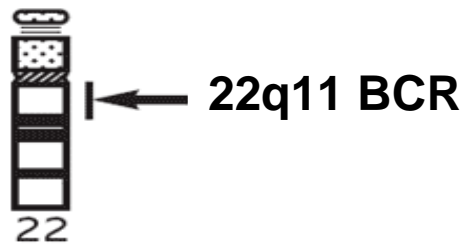
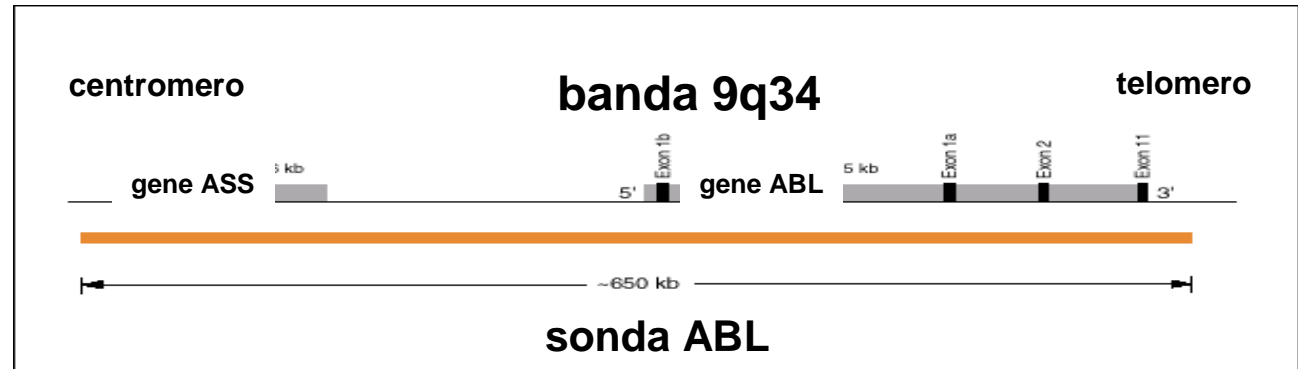
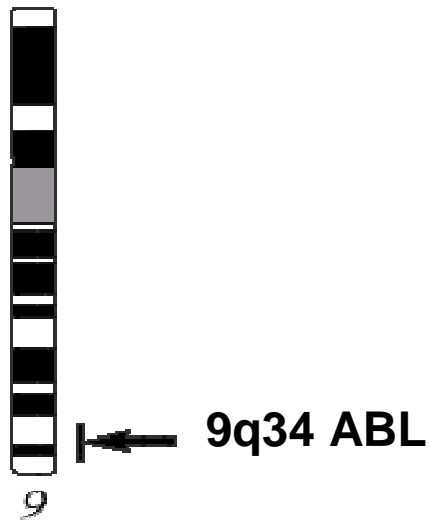
Utilizza sonde di acido nucleico
marcate con fluorocromi diversi

Il principio di base è l'ibridazione (hybridization, annealing) di un filamento di DNA denominato "sonda" (probe) con sequenze omologhe presenti nel genoma in studio

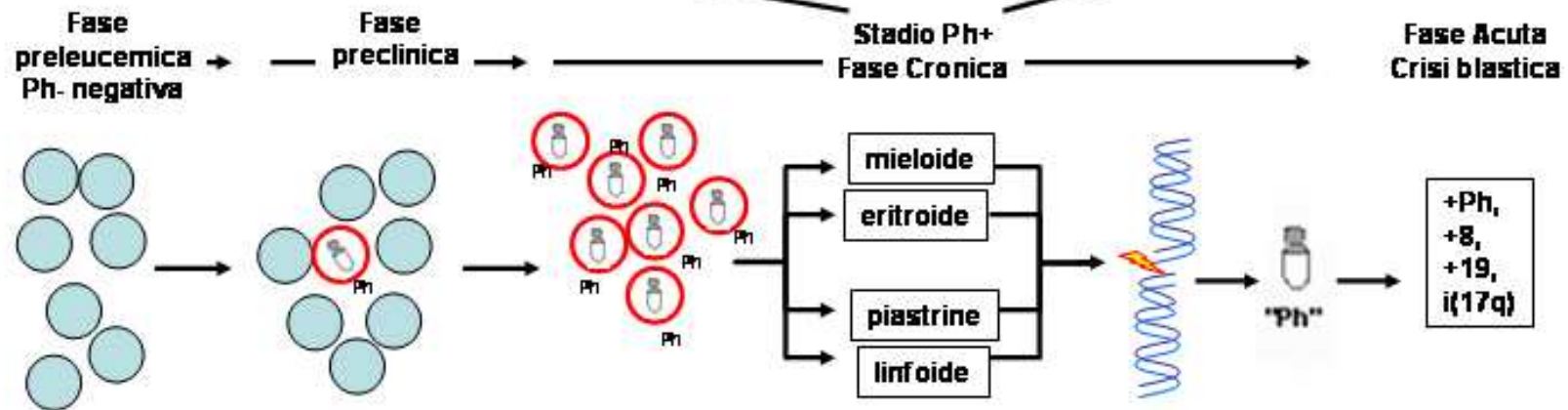
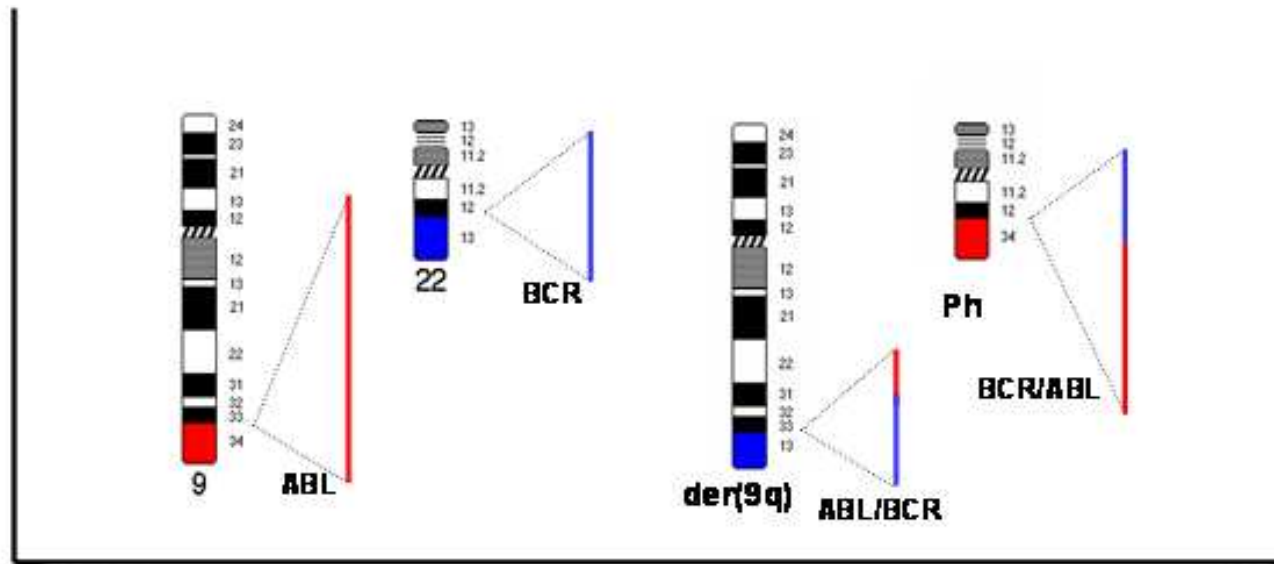


La FISH è una metodica che prevede un "target" di studio

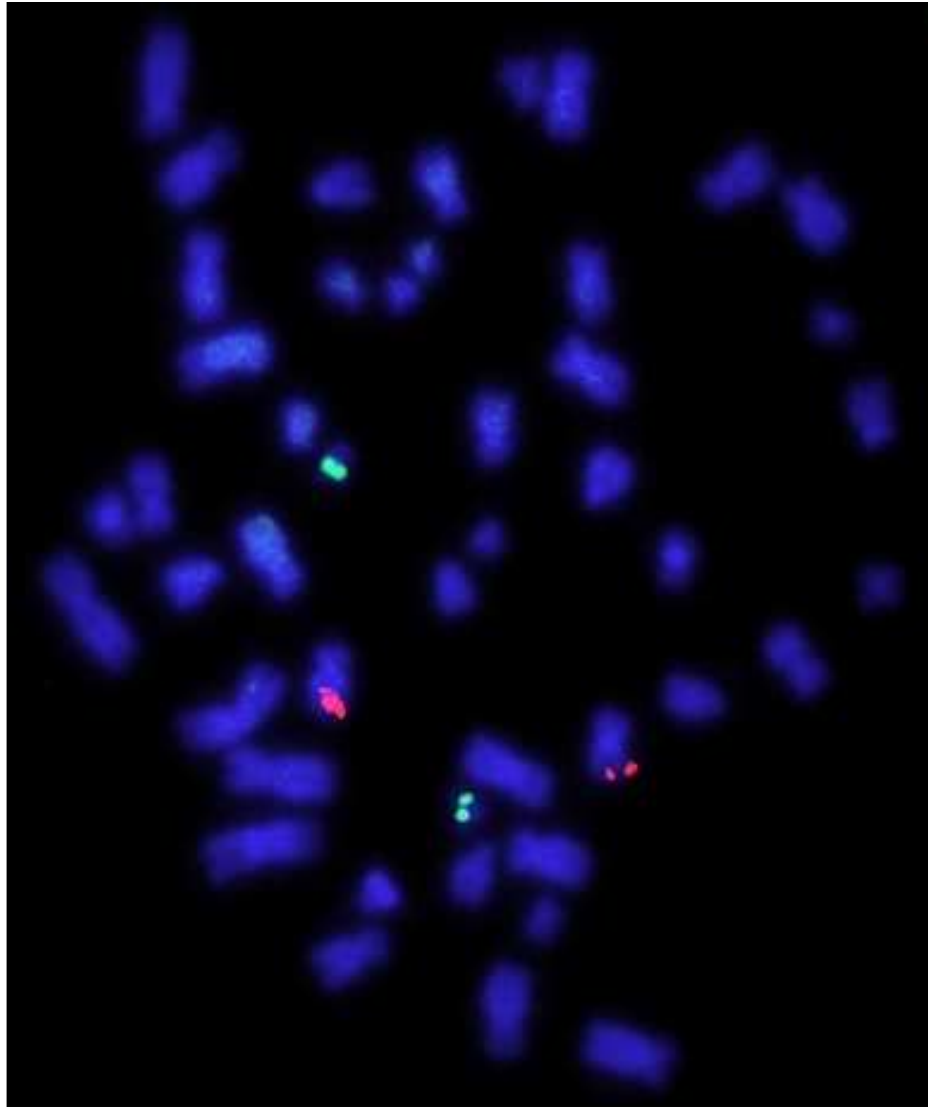
t(9;22)(q34;q11) SISTEMA COMBINATO SEGREGAZIONE-COLOCALIZZAZIONE, Dual Color Dual Fusion



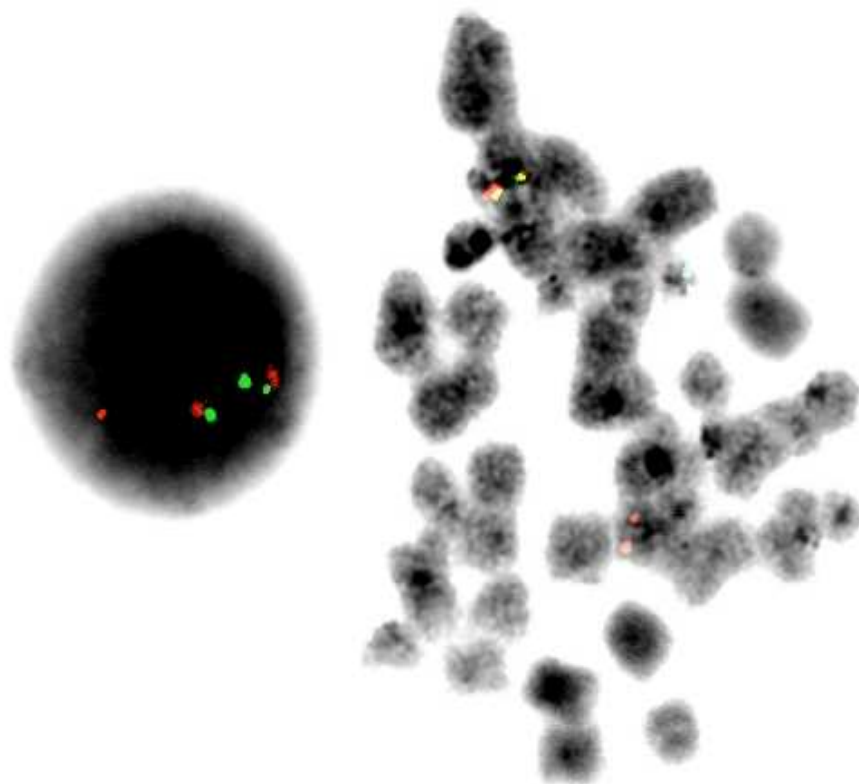
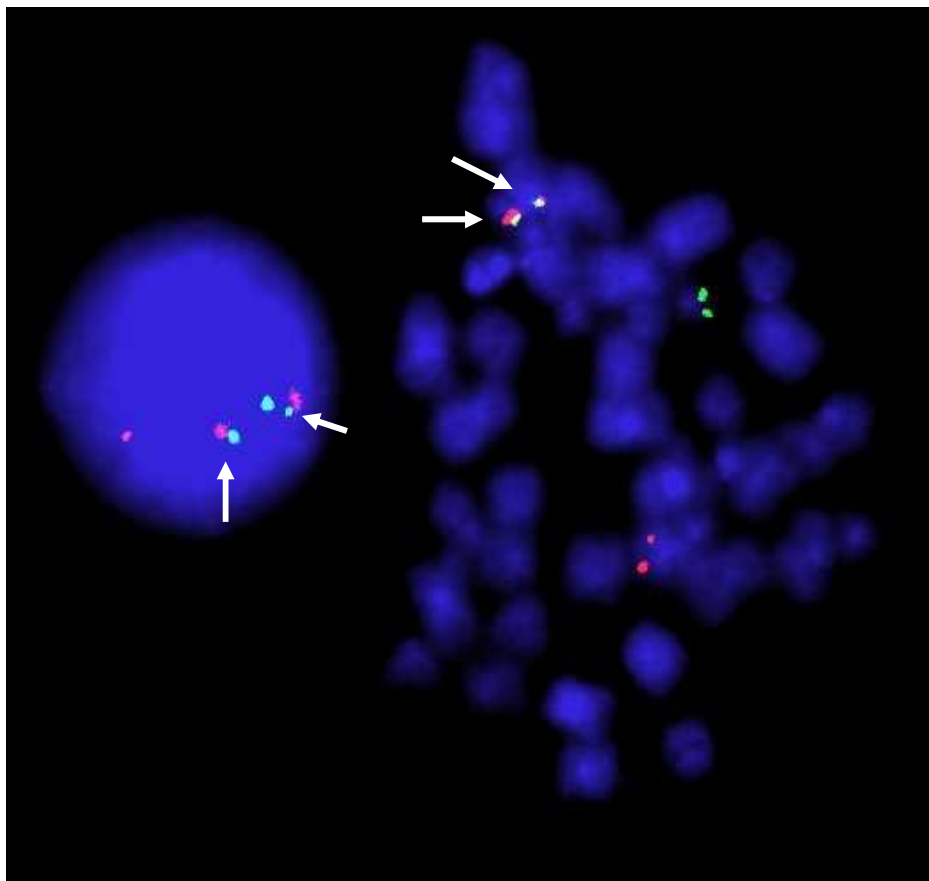
LA LEUCEMIA MIELOIDE CRONICA COME MODELLO DI LESIONE DIAGNOSTICA



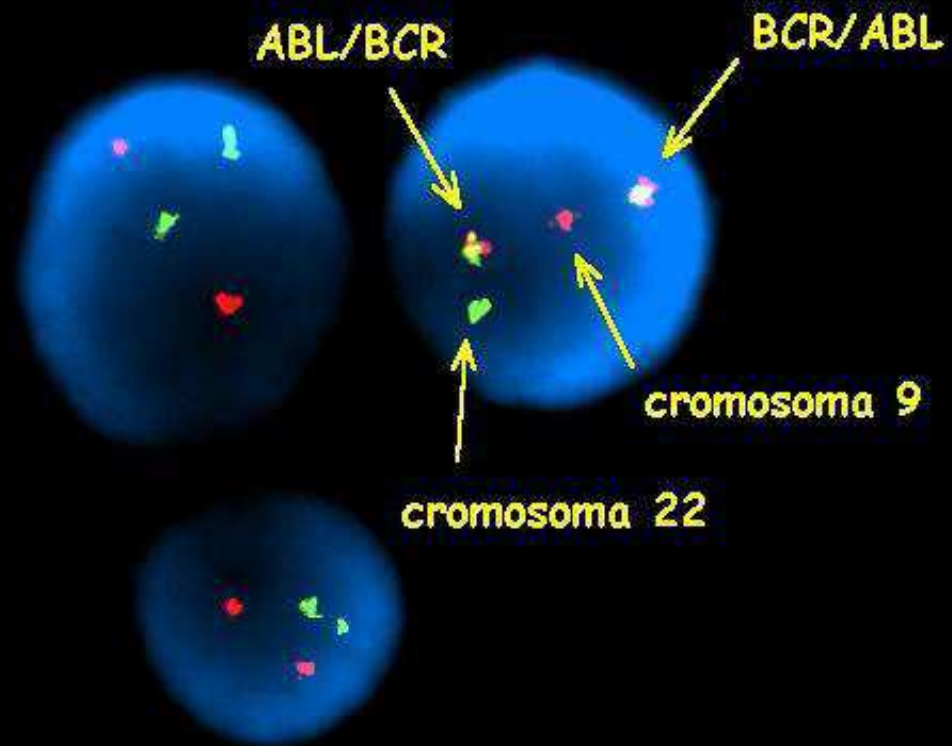
Sonda BCR/ABL DCDF – metafase normale



Sonda BCR/ABL DCDF – metafase Ph+



Fusione BCR/ABL nella Leucemia Mieloide Cronica

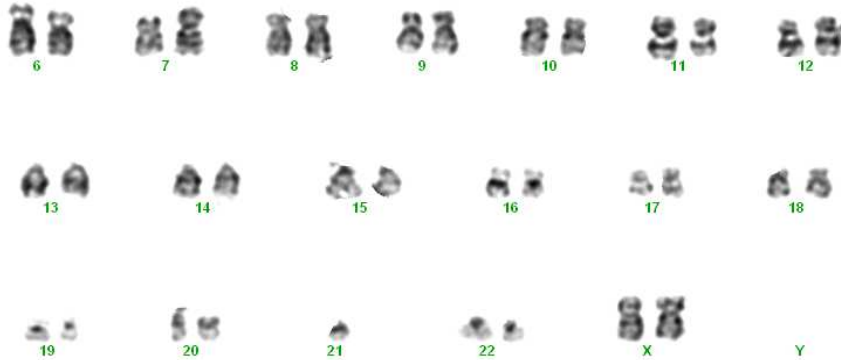


VANTAGGI INTERPRETATIVI DELLA FISH

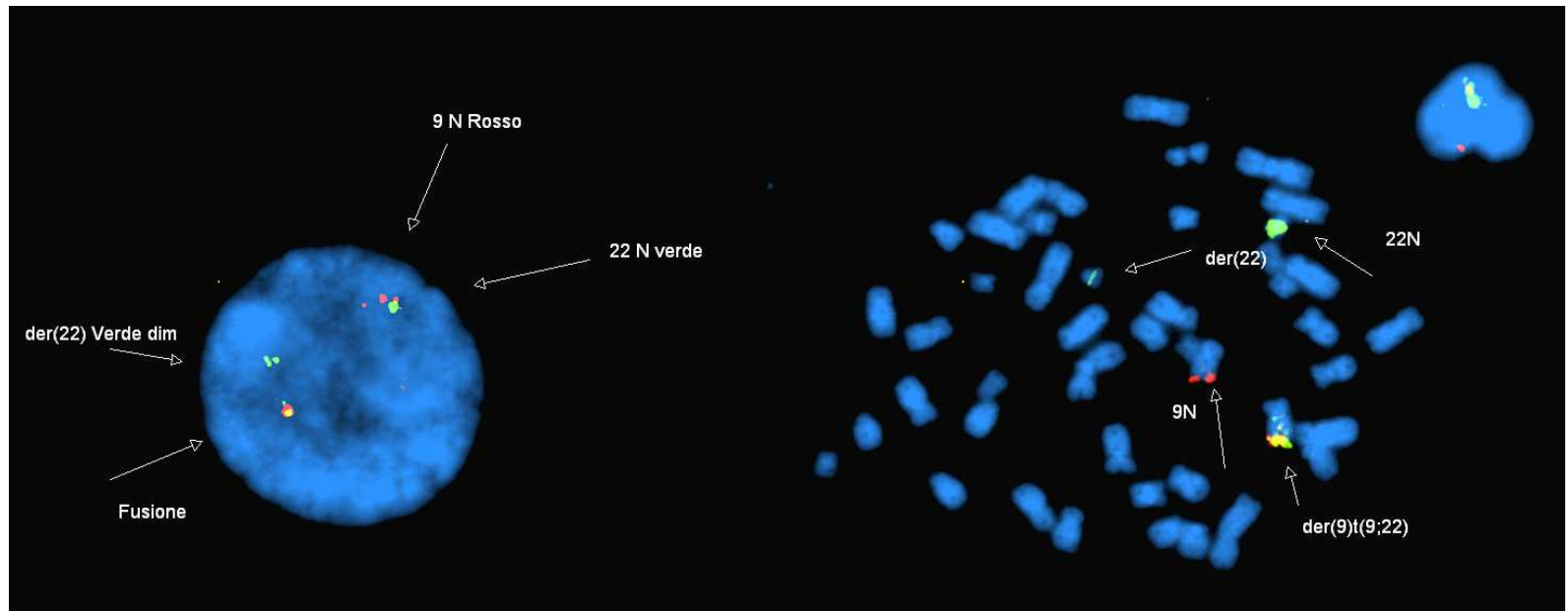
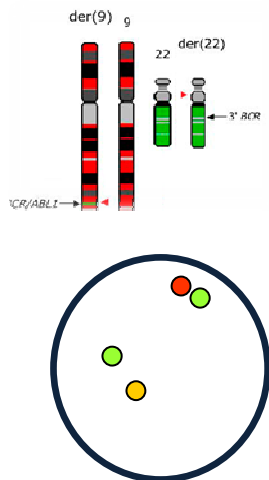
Inserzione di BCR su ABL: CASO 1



L'analisi del cariotipo non evidenzia t(9;22)

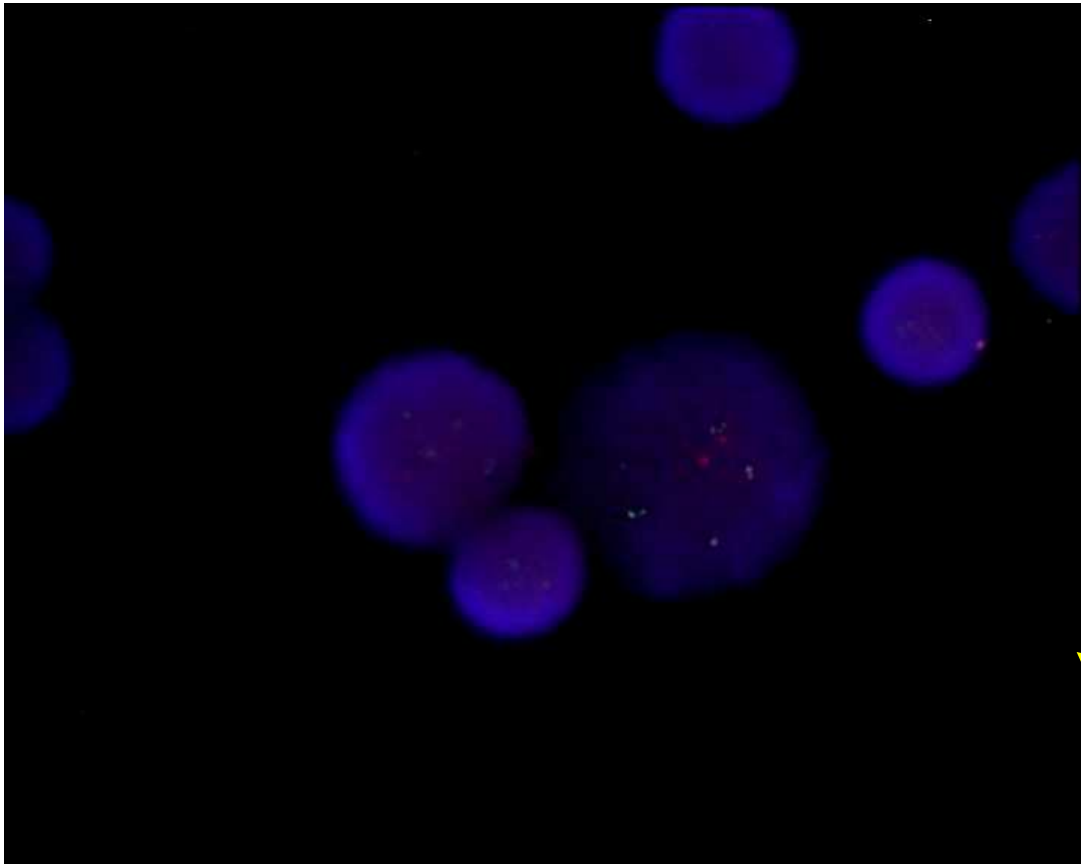


All'analisi I-FISH il pattern trovando riscontro nel **76% 1F2V1R** → conferma nel pattern di 15/15 metafasi che rivela **1R** sul **9N**, **1V** sul **22N** e **1V** sul **22q**. Un **segnale di fusione** sul **9q**. Non esiste evidenza di **fusione** sul **der(22)**, ma il pattern è compatibile con l'**inserzione di BCR su cromosoma 9q-**.

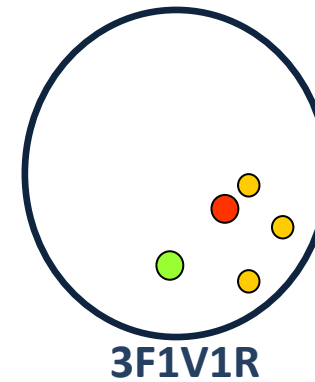


VANTAGGI INTERPRETATIVI DELLA FISH

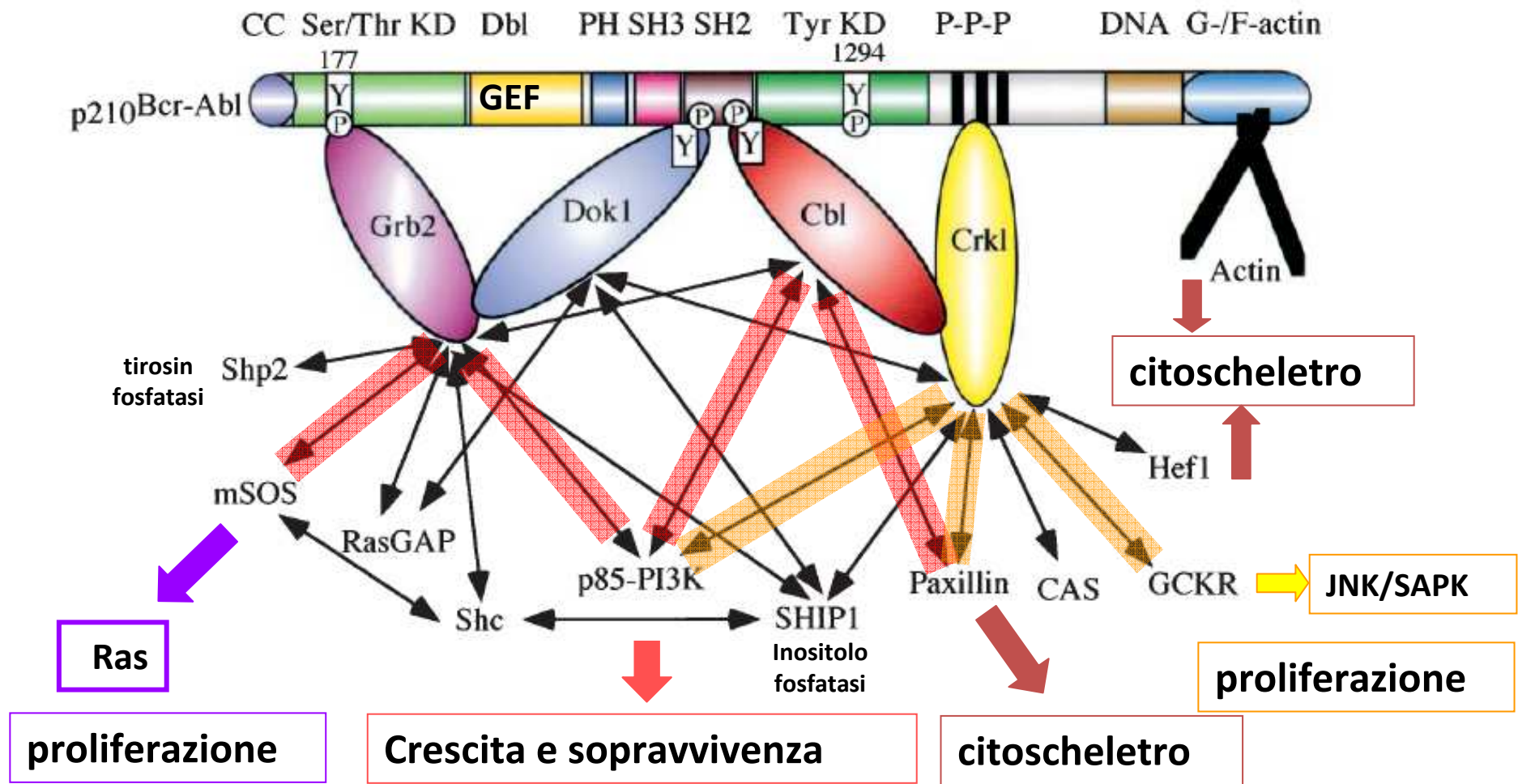
Doppio Cromosoma Philadelphia: CASO 2



Co-esistenza del clone neoplastico convenzionale (Ph+) e di un assetto di **3 segnali gialli di fusione, 1 segnale verde e 1 rosso**, ad evidenza della presenza di un **clone con Ph''**

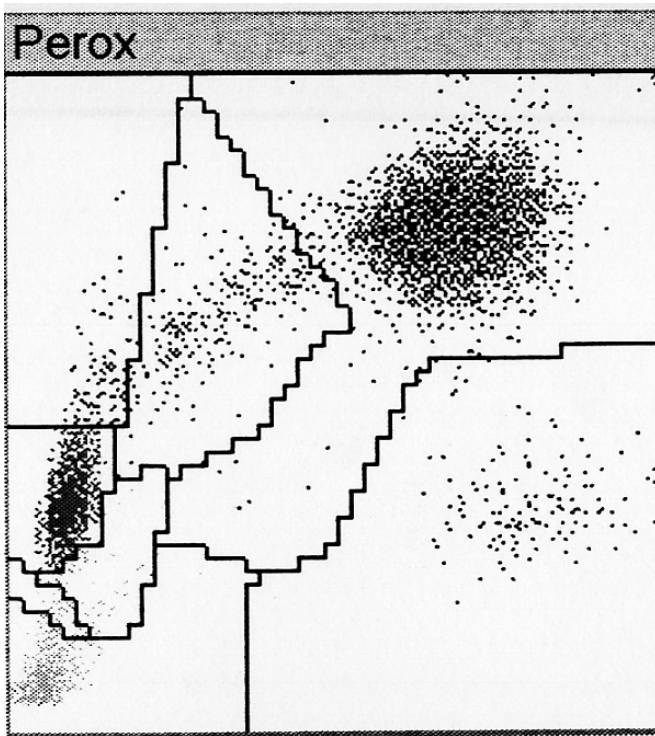


Oncogenesi *BCR/ABL*-mediata: interazioni molecolari

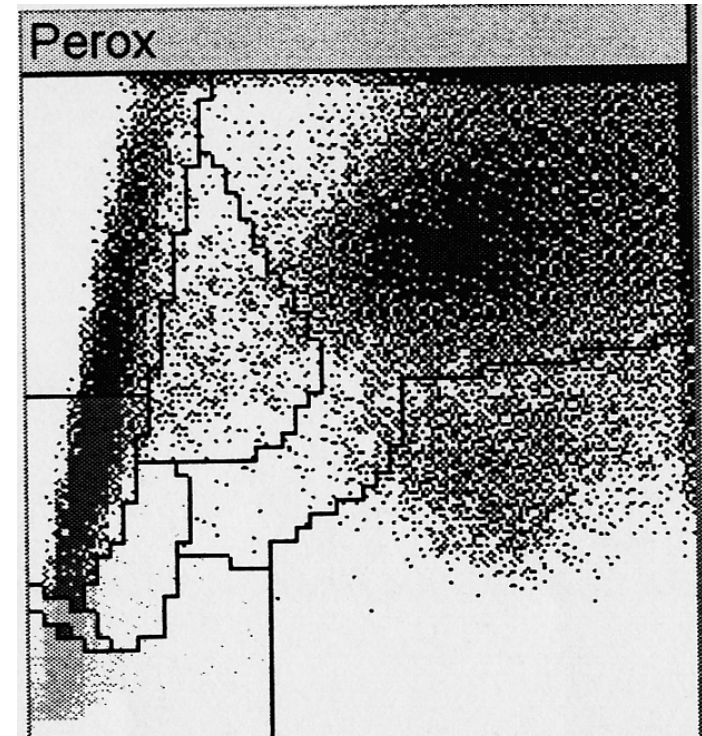


Sospetto Diagnostico Il Laboratorio

WBC		7.40	$\times 10^3 / \mu\text{L}$
Neutro	54.4	4.03	$\times 10^3 / \mu\text{L}$
Linfo	34.0	2.52	$\times 10^3 / \mu\text{L}$
Mono	4.8	0.35	$\times 10^3 / \mu\text{L}$
Eos	2.6	0.19	$\times 10^3 / \mu\text{L}$
Baso	H 2.2	0.16	$\times 10^3 / \mu\text{L}$
LUC	2.0	0.15	$\times 10^3 / \mu\text{L}$

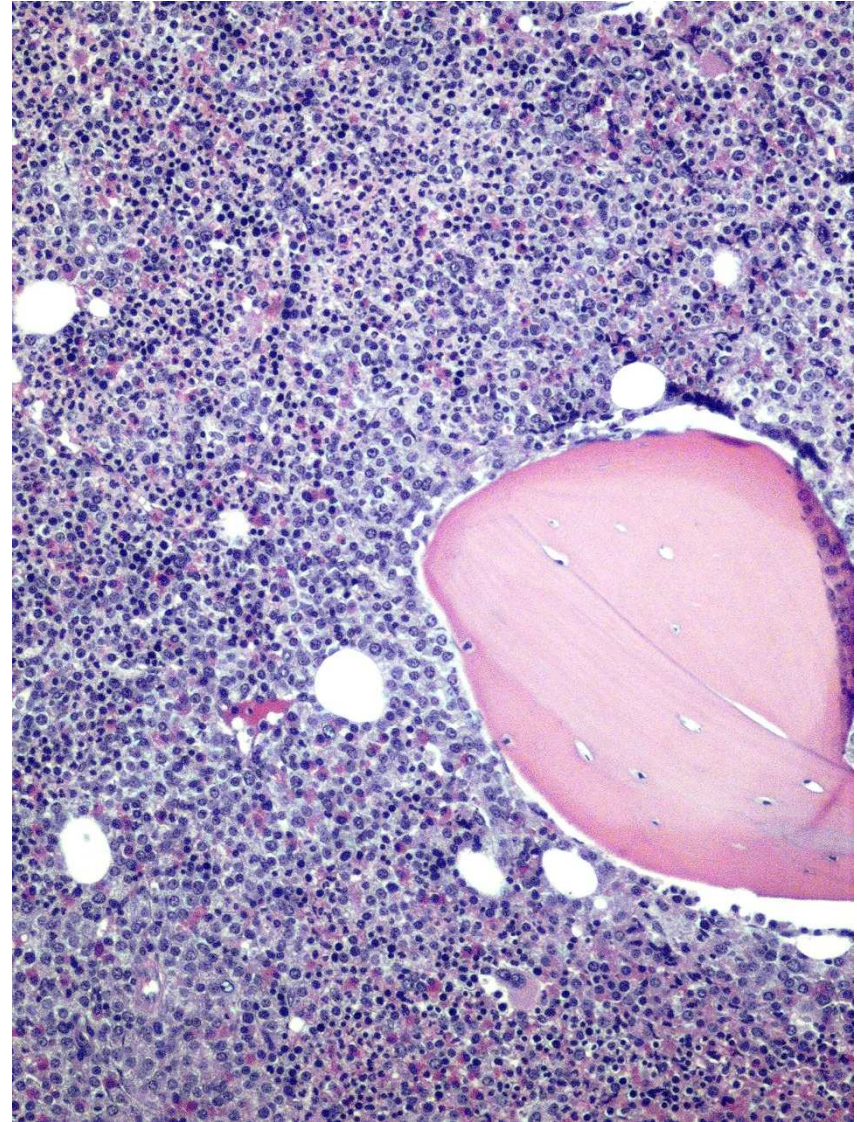
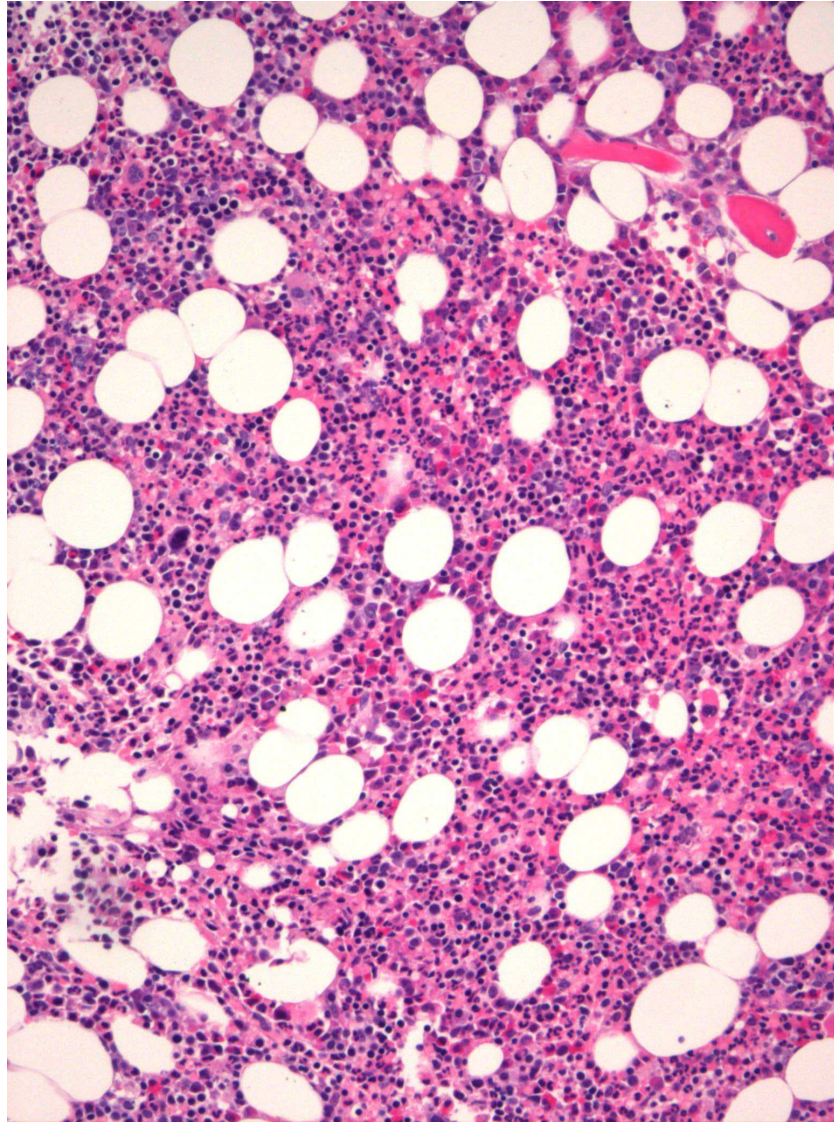


WBC		H 259.19	$\times 10^3 /$
Neutro	64.8	* H 168.05	$\times 10^3 /$
Linfo	19.3	* H 49.95	$\times 10^3 /$
Mono	L 1.0	* H 2.48	$\times 10^3 /$
Eos	4.7	* H 12.10	$\times 10^3 /$
Baso	H 13.6	* H 35.19	$\times 10^3 /$
LUC	H 10.3	* H 26.60	$\times 10^3 /$



FISIOPATOLOGIA DI ABL

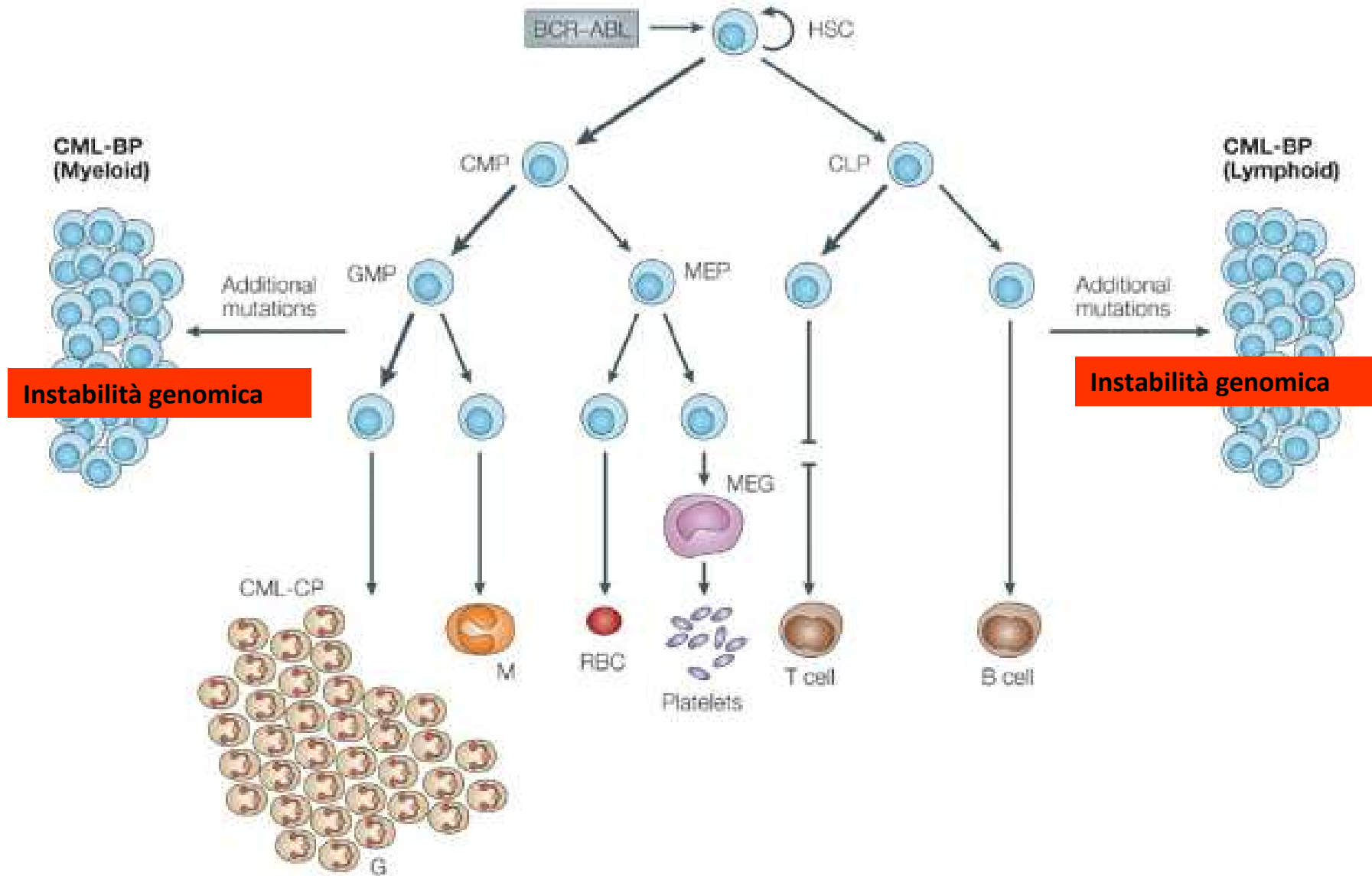
La Leucemia Mieloide Cronica



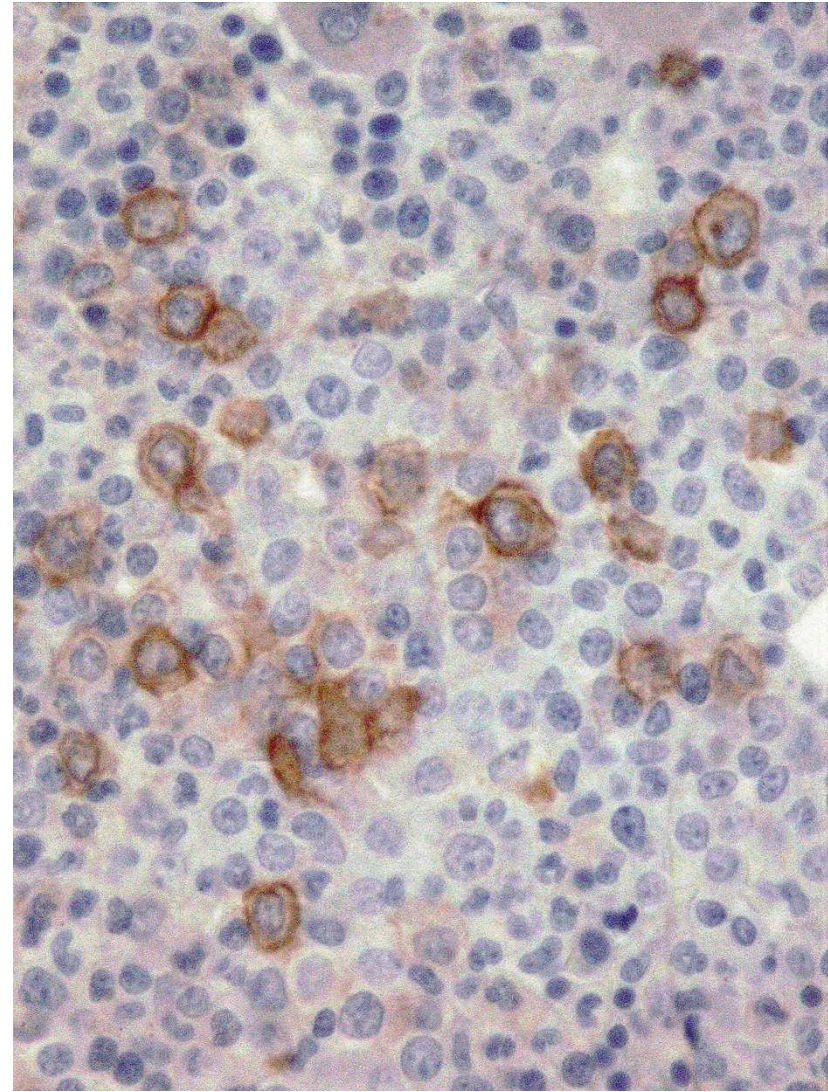
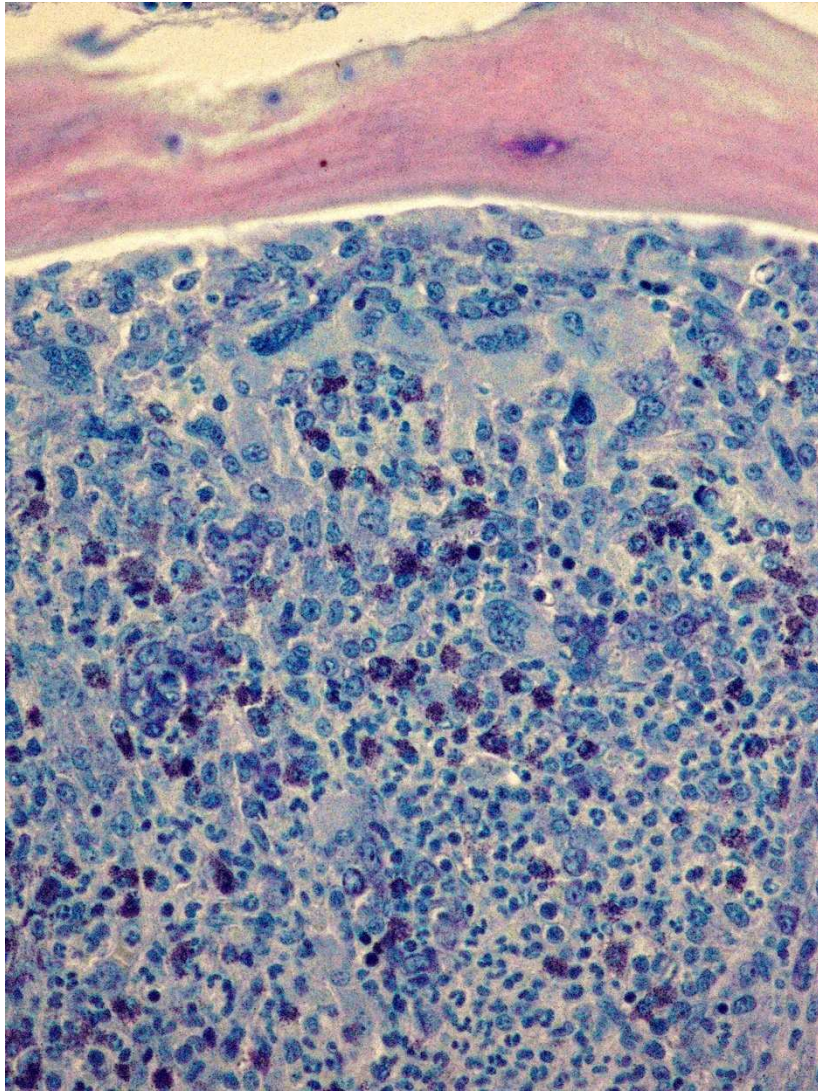
Cortesia Prof. Luigi Cavazzini, Ist. Anatomia Patologica, Ferrara

MECCANISMI DI PROGRESSIONE

STORIA NATURALE DELLA LMC



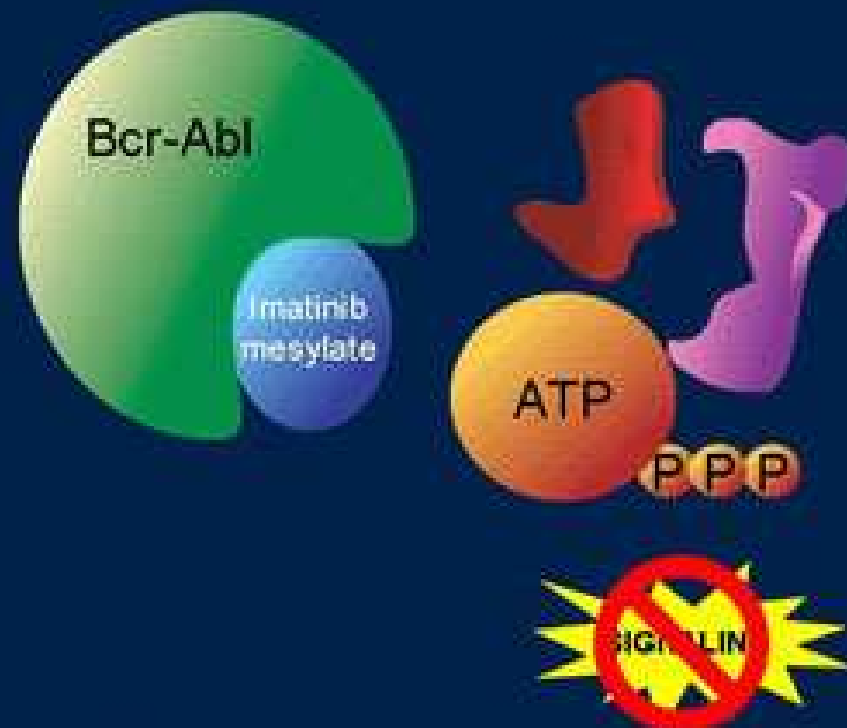
MECCANISMI DI PROGRESSIONE



Cortesia Prof. Luigi Cavazzini, Ist. Anatomia Patologica, Ferrara

Imatinib Mesylate: Mechanism of Action

- Imatinib mesylate occupies the ATP binding pocket of the Abl kinase domain
- This prevents substrate phosphorylation and signaling
- A lack of signaling inhibits proliferation and survival



**le lesioni citogenetiche
ricorrenti possono identificare
entità distinte di malattia**

LEUCEMIA ACUTA MIELOIDE (LAM)

Definizione:

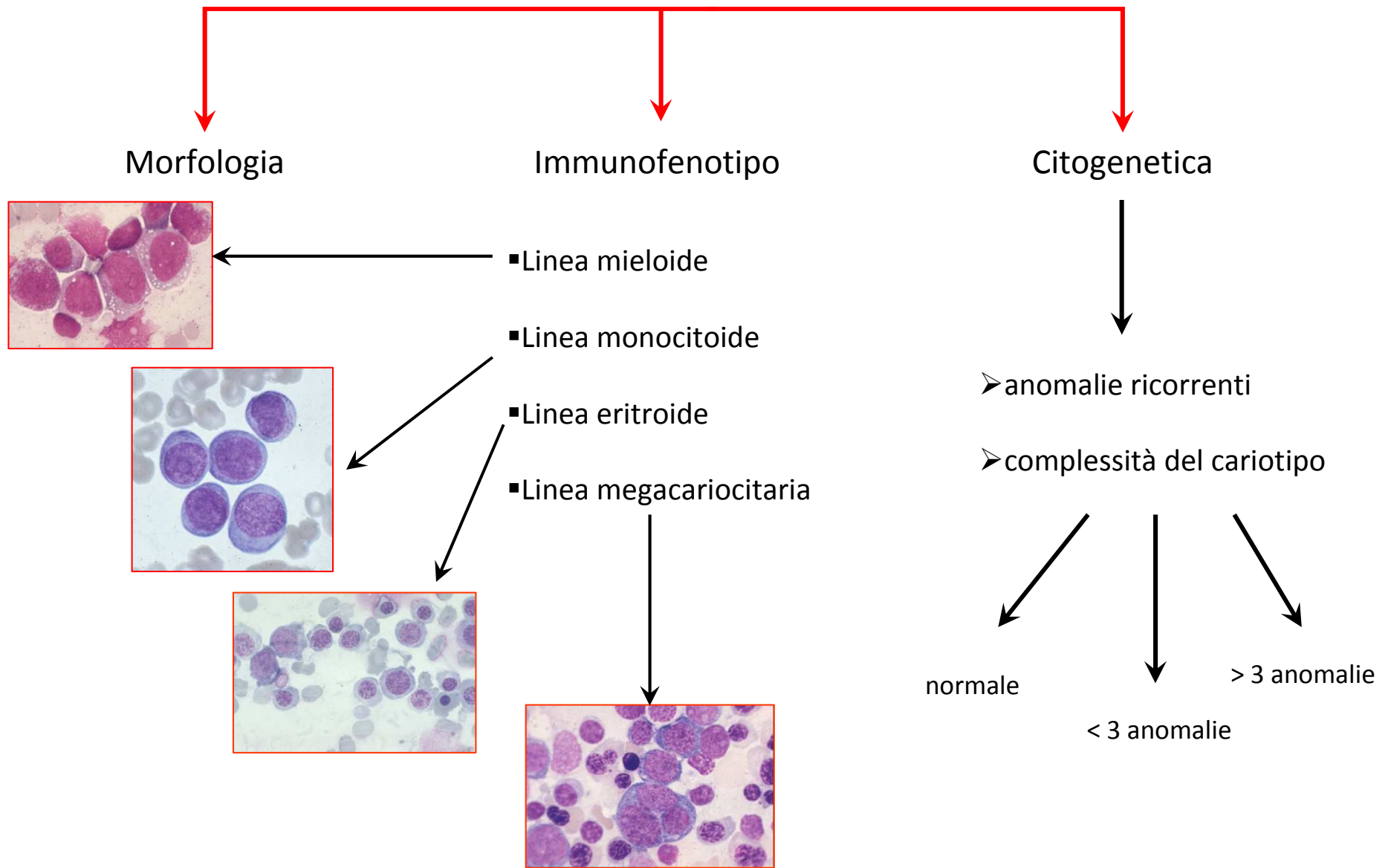
espansione clonale di elementi immaturi della serie mieloide che hanno subito un danno genetico tale da non permettere il completamento del processo differenziativo.

Diagnosi Clinica:

accumulo nel sangue periferico e nel midollo di precursori immaturi altamente indifferenziati:

- Leucocitosi/Leucopenia
- Anemia (da occupazione midollare)
- Piastrinopenia (idem)

CRITERI CLASSIFICATIVI PER LE LEUCEMIE ACUTE MIELOIDI



1) IDENTIFICAZIONE DI IDENTITA' DISTINTE DI MALATTIA

-Classificazione WHO delle Leucemie acute Mieloidi-

1. Leucemie mieloidi acute con traslocazioni citogenetiche ricorrenti

- LMA con t(8;21) (q22;q22), AML1(CBF α)/ETO
- Leucemia promielocitica acuta [LMA con t(15;17)(q22; q11-12) e varianti PML/RAR α]
- LMA con ipereosinofilia midollare [inv(16)(p13;q22) o t(16; 16)(p13;q11), CBF β /MYH11X]
- LMA con anomalie 11q23 (MLL)

2. Leucemia mieloide acuta con displasia multilineare

- Secondaria a sindrome mielodisplastica
- De novo

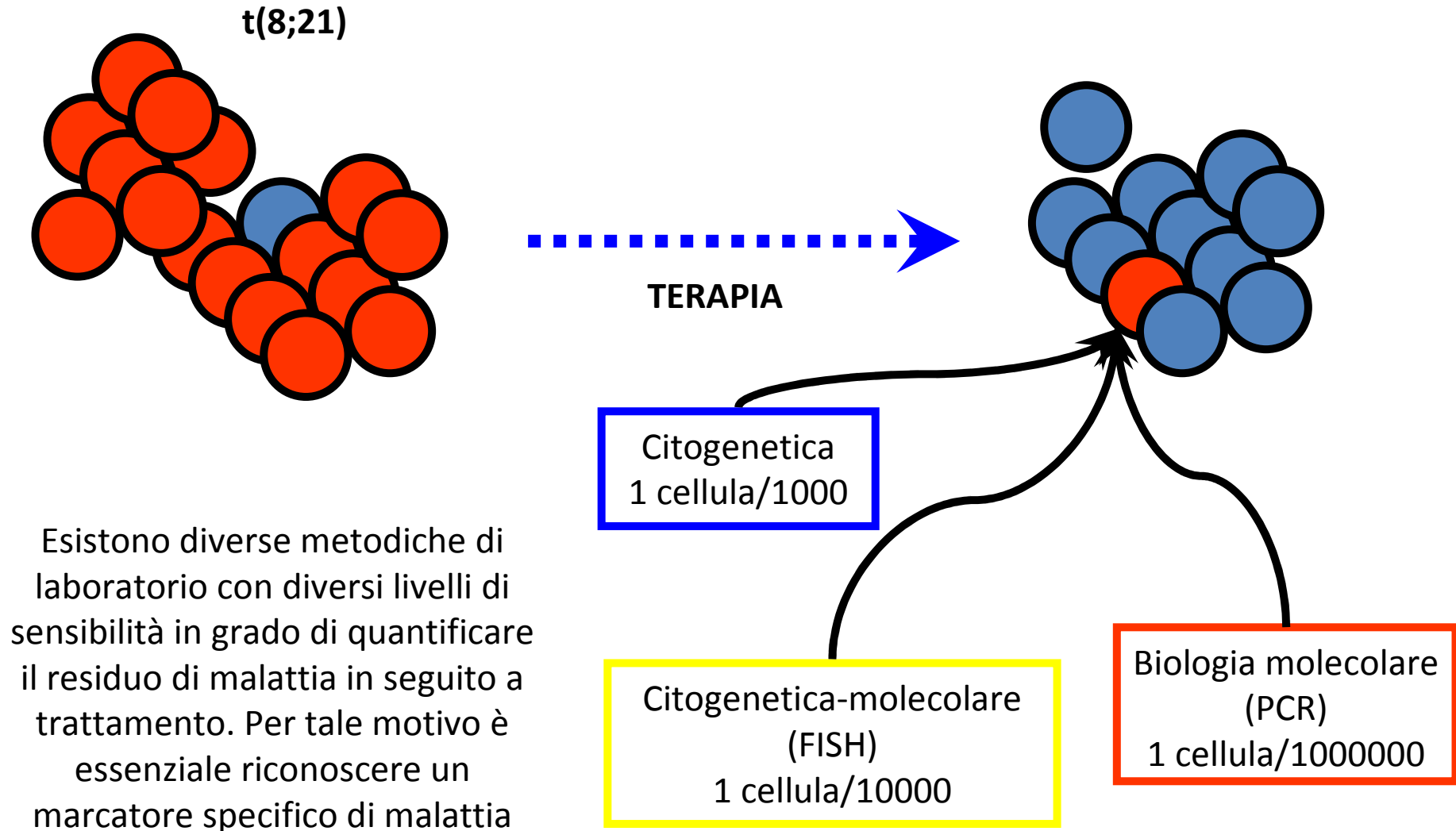
3. Leucemia mieloide acuta e sindromi mielodisplastiche secondarie a chemioterapia

- Secondaria ad agenti alchilanti
- Secondaria a epipodofillotossine
- Altri tipi

4. Leucemia mieloide acuta non altrimenti classificata

- LMA con differenziazione minima
- LMA senza maturazione
- LMA con maturazione
- Leucemia mielomonocitica acuta
- Leucemia monocitica acuta
- Eritroleucemia acuta
- Leucemia megacariocitica acuta
- Leucemia basofila acuta
- Panmielosi acuta con mielofibrosi

3) MONITORAGGIO DELLA MALATTIA RESIDUA



MRC/NCRI AML Trials: Overall Survival
Ages 16–59 excluding known prognostic abnormalities

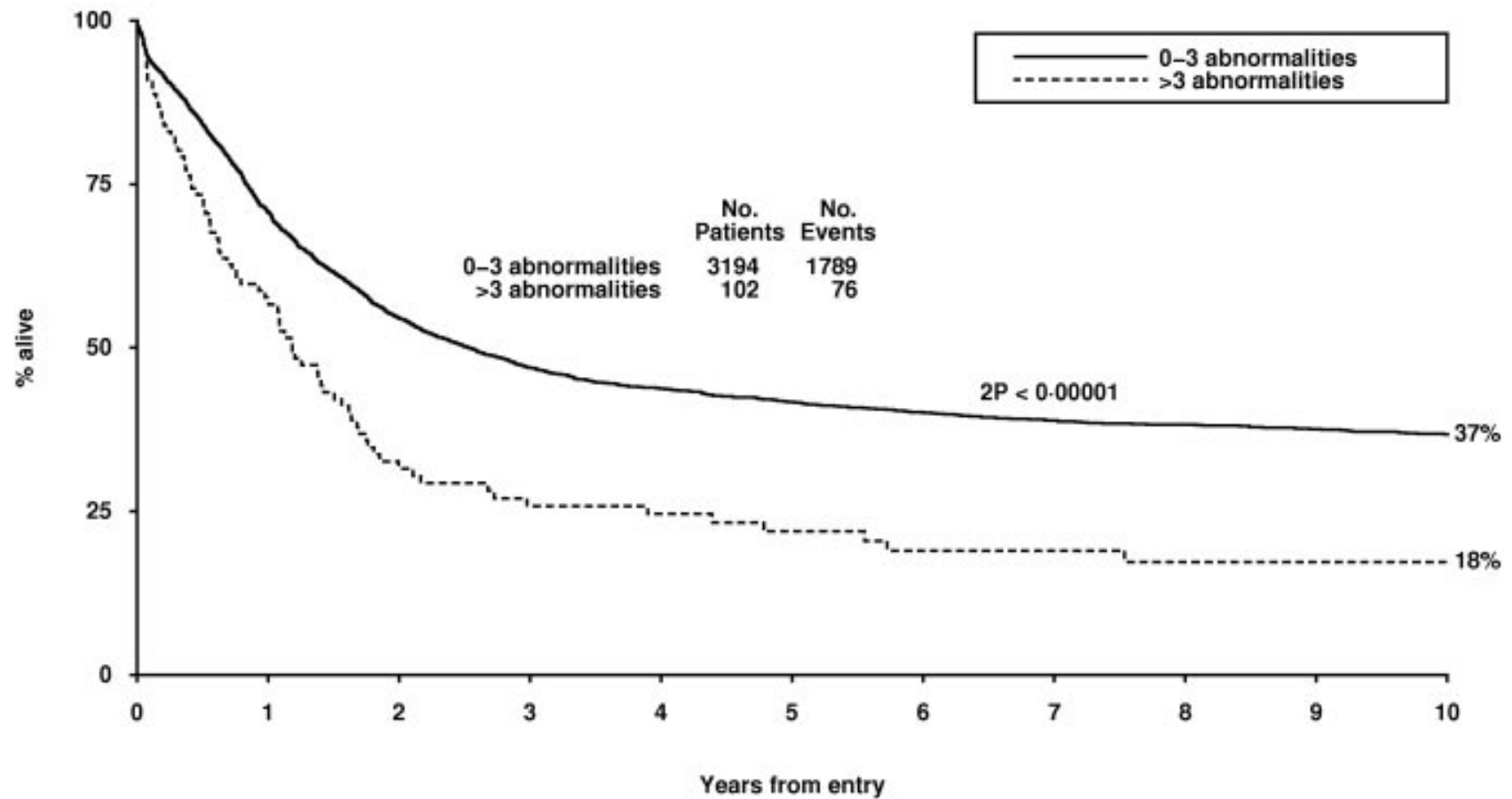


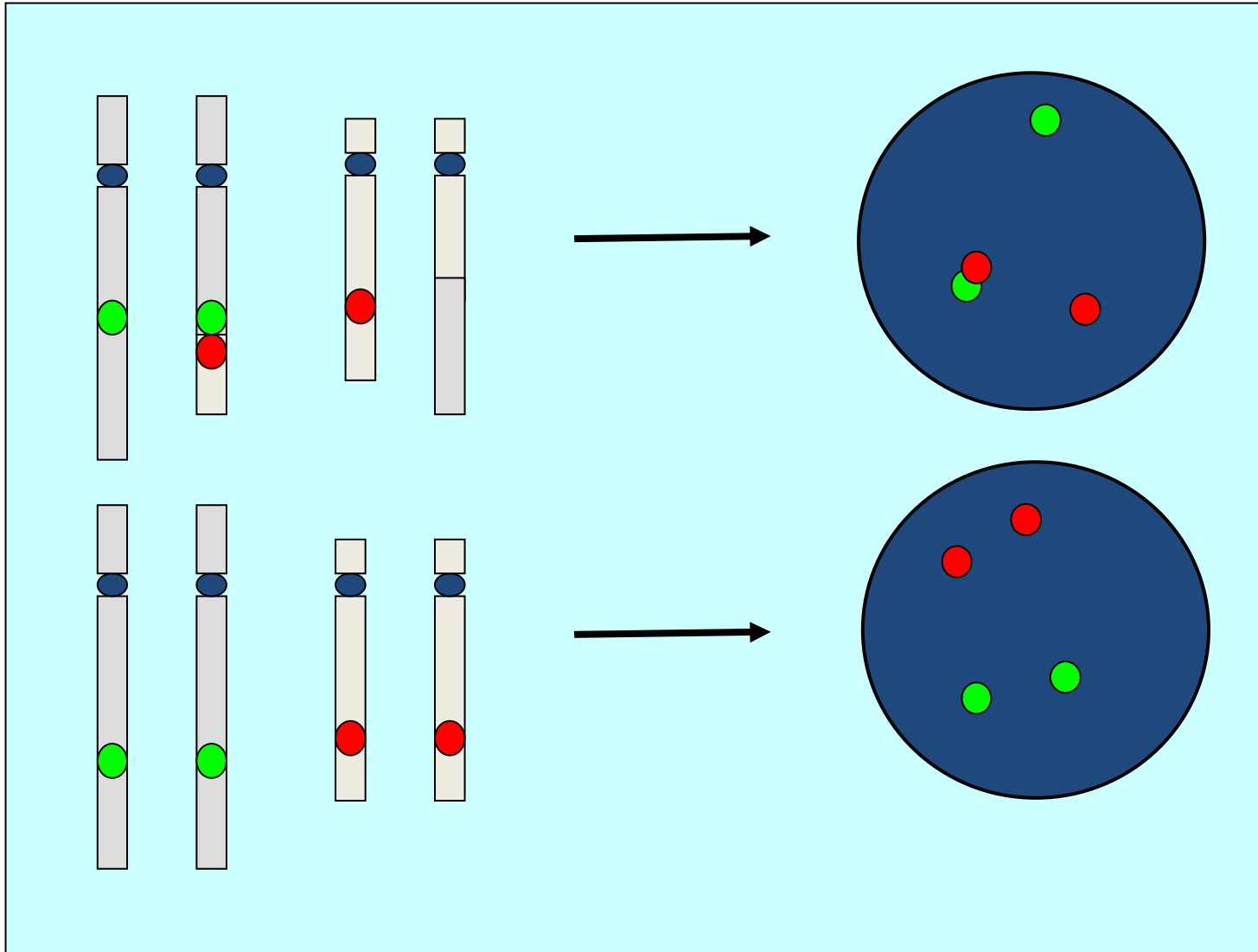
Table 4. Revised MRC prognostic classification based on multivariable analyses

Cytogenetic abnormality	Comments
Favorable	
t(15;17)(q22;q21)	
t(8;21)(q22;q22)	Irrespective of additional cytogenetic abnormalities*
inv(16)(p13q22)/t(16;16)(p13;q22)	
Intermediate	
Entities not classified as favorable or adverse	
Adverse	
abn(3q) [excluding t(3;5)(q21~25;q31~35), inv(3)(q21q26)/t(3;3)(q21;q26), add(5q), del(5q), -5, -7, add(7q)/del(7q),	Excluding cases with favorable karyotype†
t(6;11)(q27;q23),	
t(10;11)(p11~13;q23),	
t(11q23) [excluding t(9;11)(p21~22;q23) and t(11;19)(q23;p13)]	
t(9;22)(q34;q11),	
-17/abn(17p),	
Complex (≥ 4 unrelated abnormalities)	

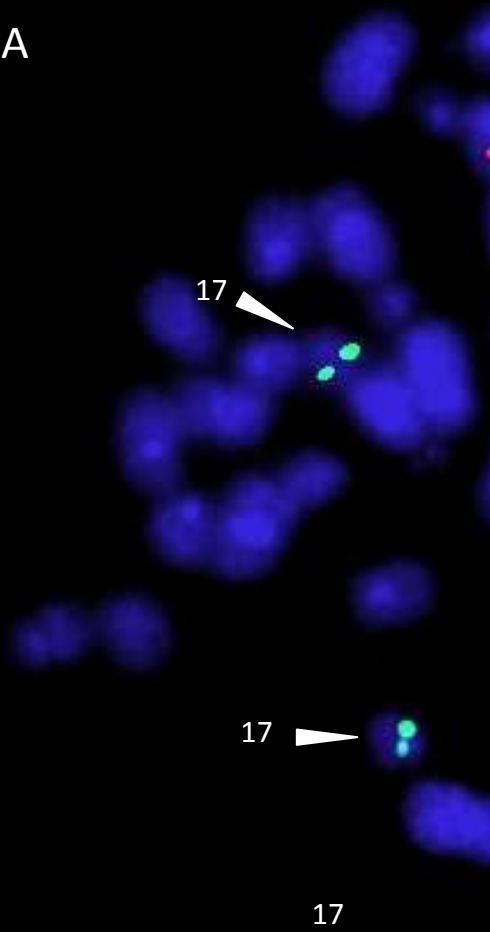
*All favorable-risk abnormalities.

†All adverse-risk abnormalities.

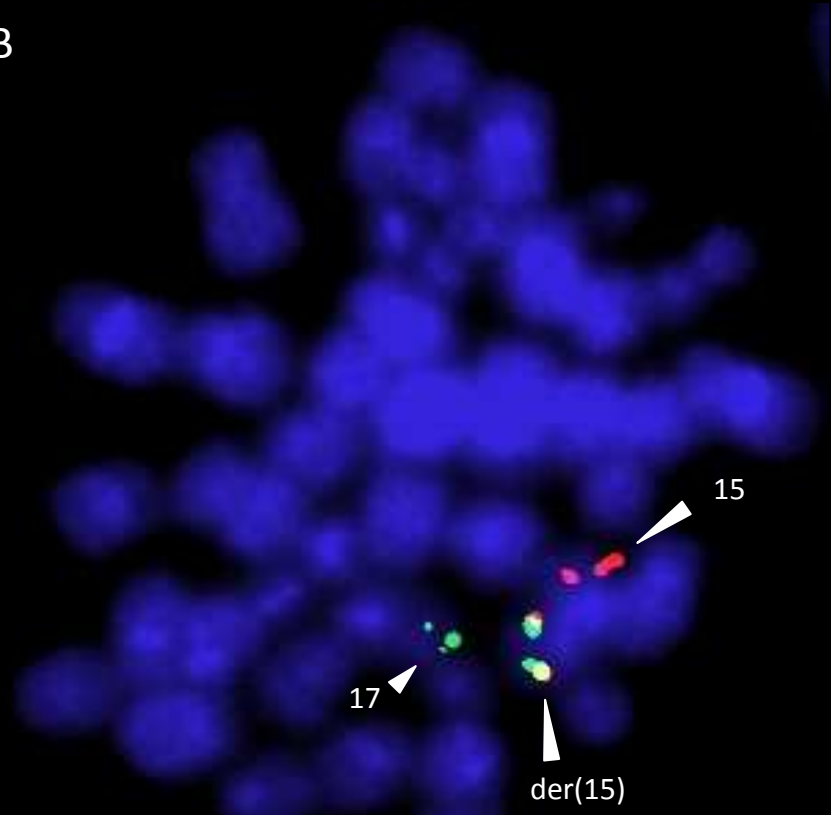
SISTEMA A COLOCALIZZAZIONE: SONDE SPECIFICHE PER GENI DIVERSI IDENTIFICANO UN SEGNALE DI FUSIONE IN PRESENZA DI TRASLOCAZIONE



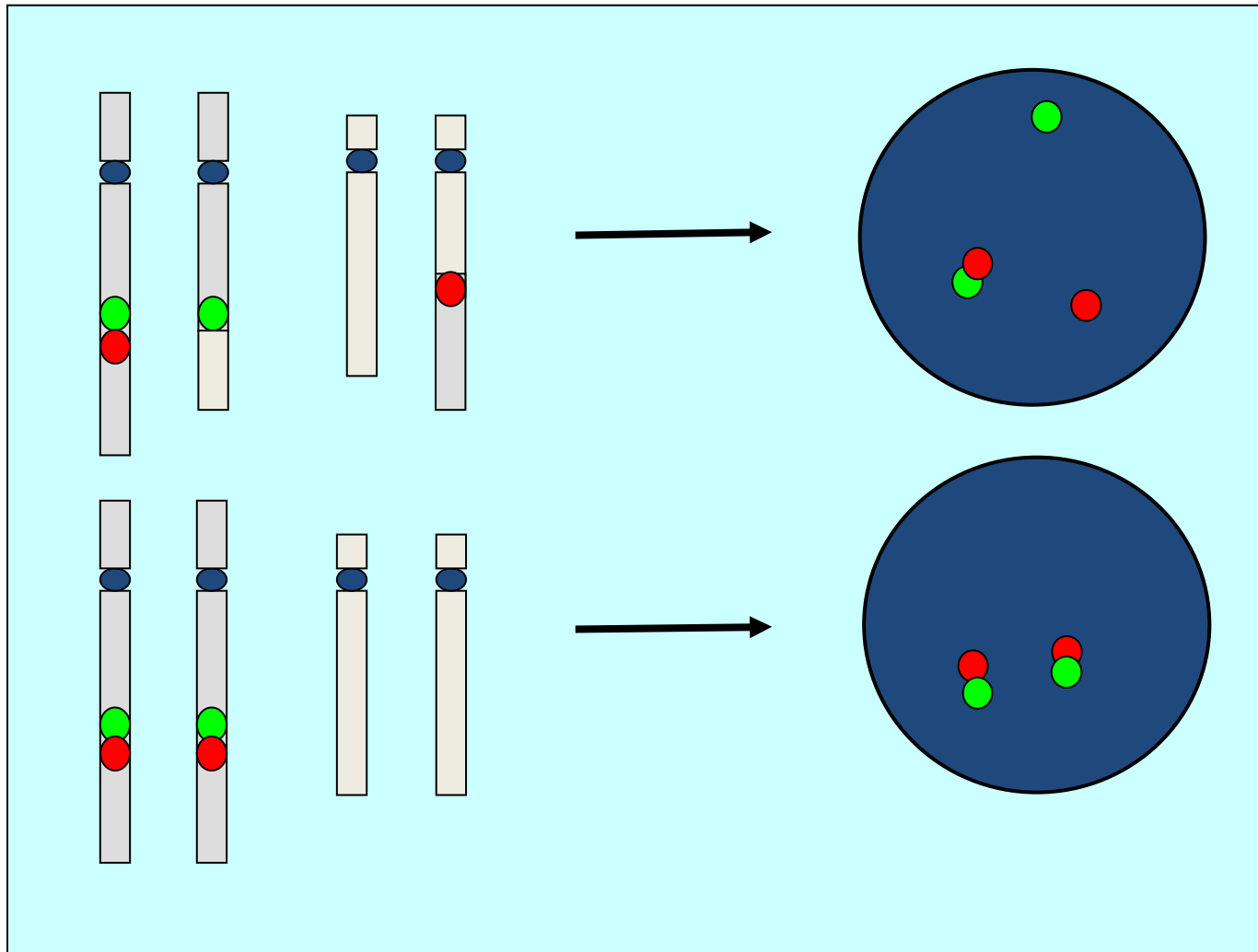
A



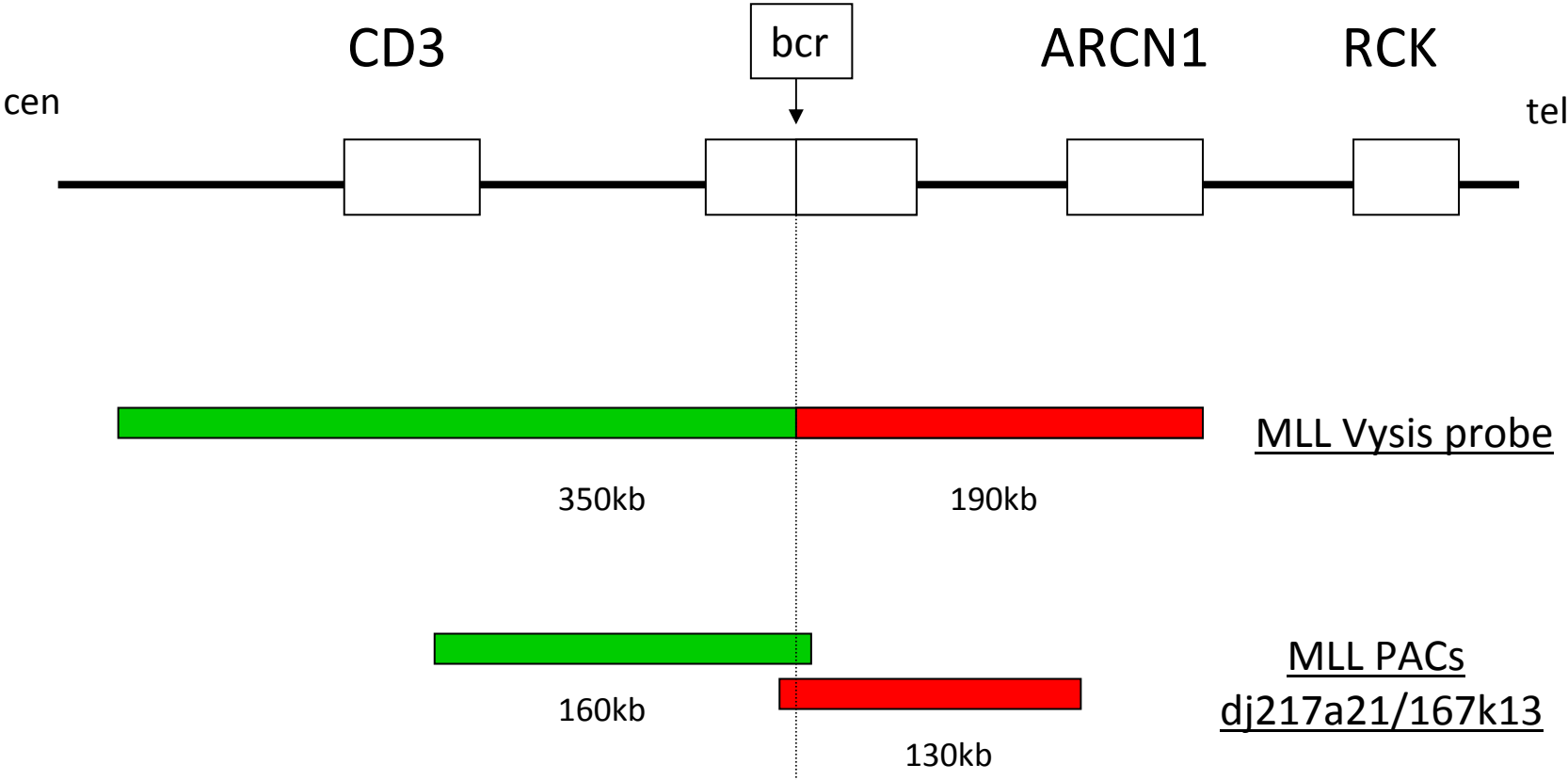
B



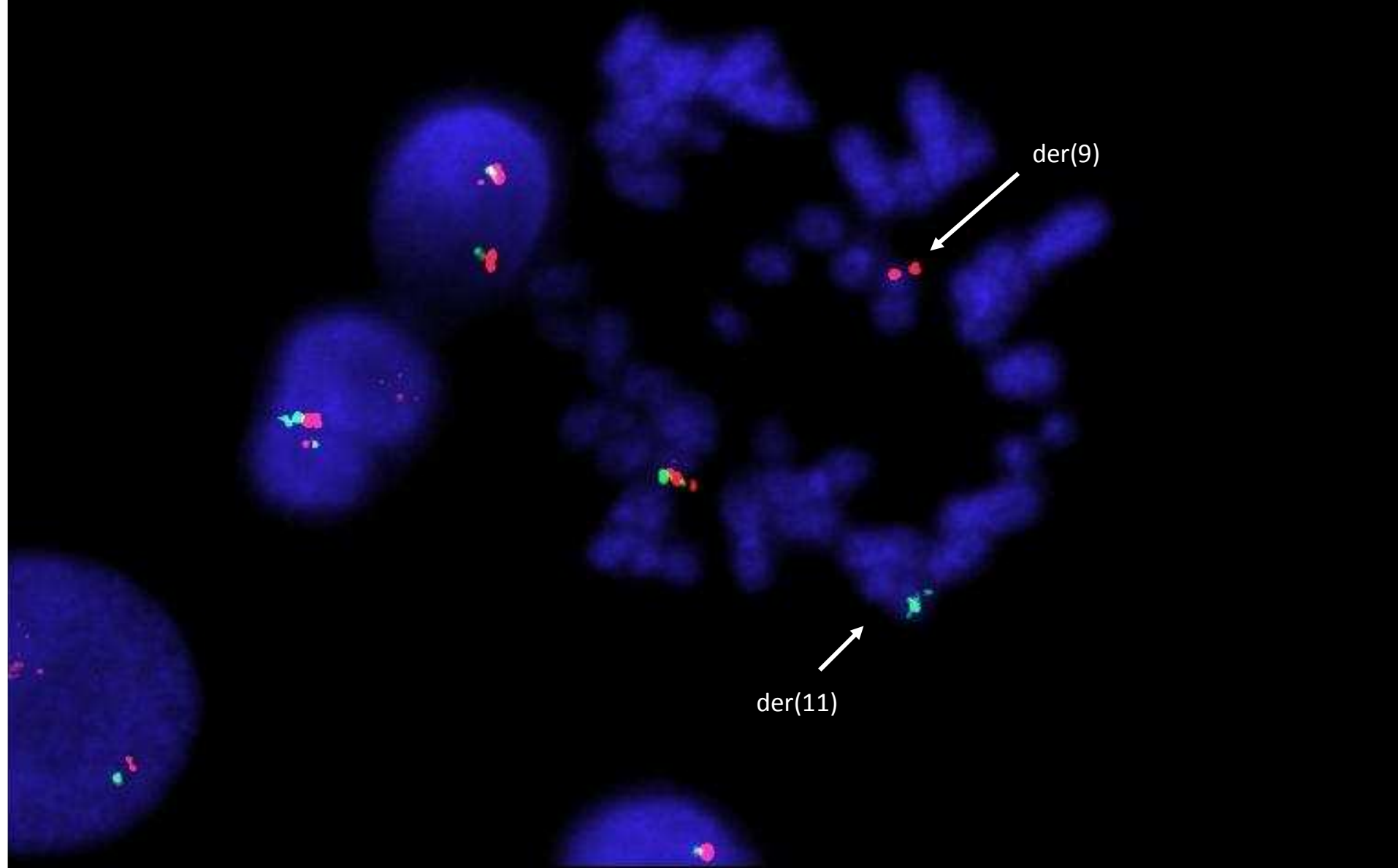
SISTEMA A SEGREGAZIONE: IL SEGNALE DI FUSIONE IDENTIFICA L'ALLELE NORMALE DEL GENE INTERESSATO NELLA TRASLOCAZIONE. QUANDO IL SEGNALE SEGREGA (LA FUSIONE SI DIVIDE IN DUE SEGNALI DI COLORE DIVERSO) È PRESENTE LA TRASLOCAZIONE



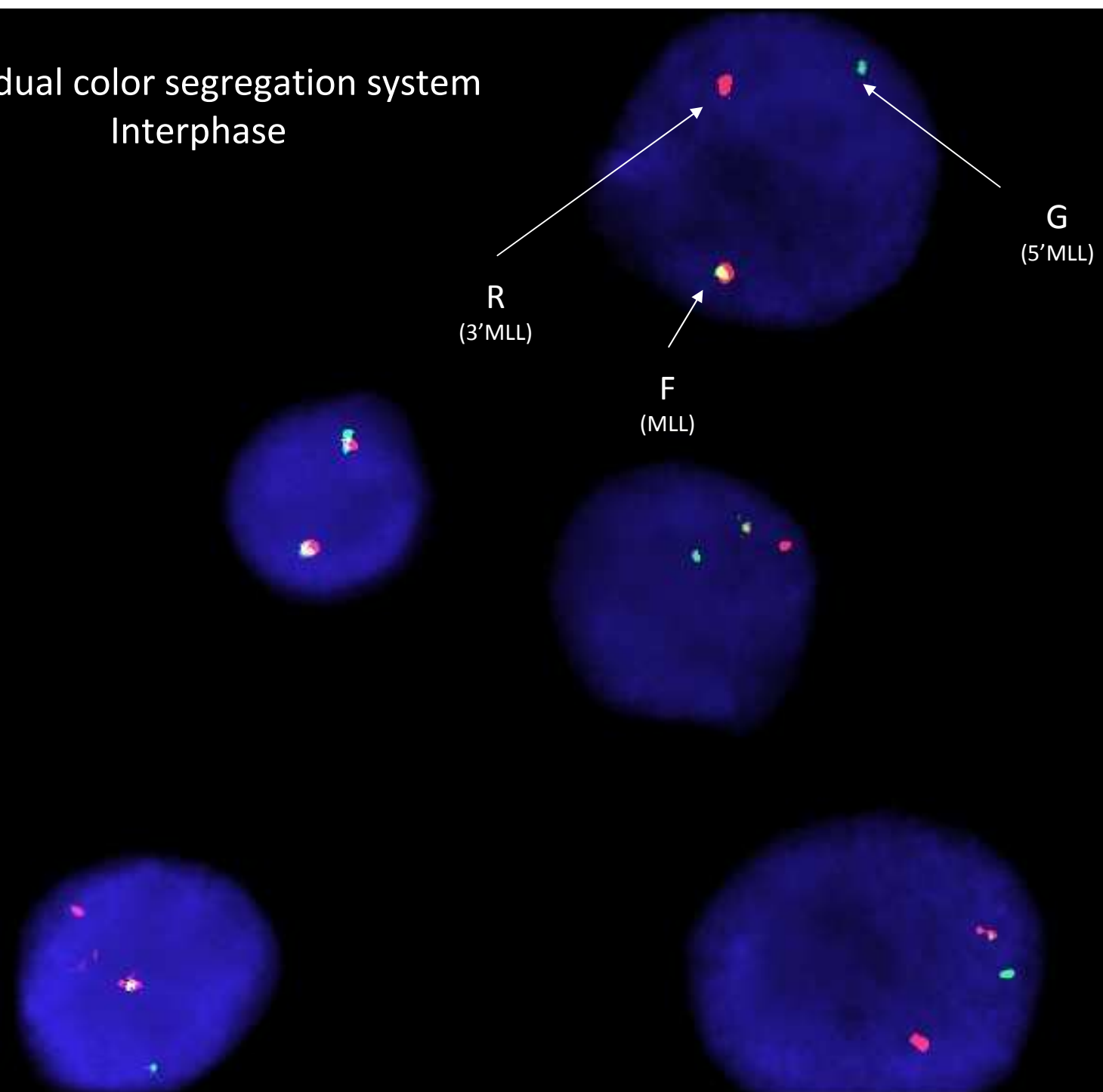
MLL probe systems

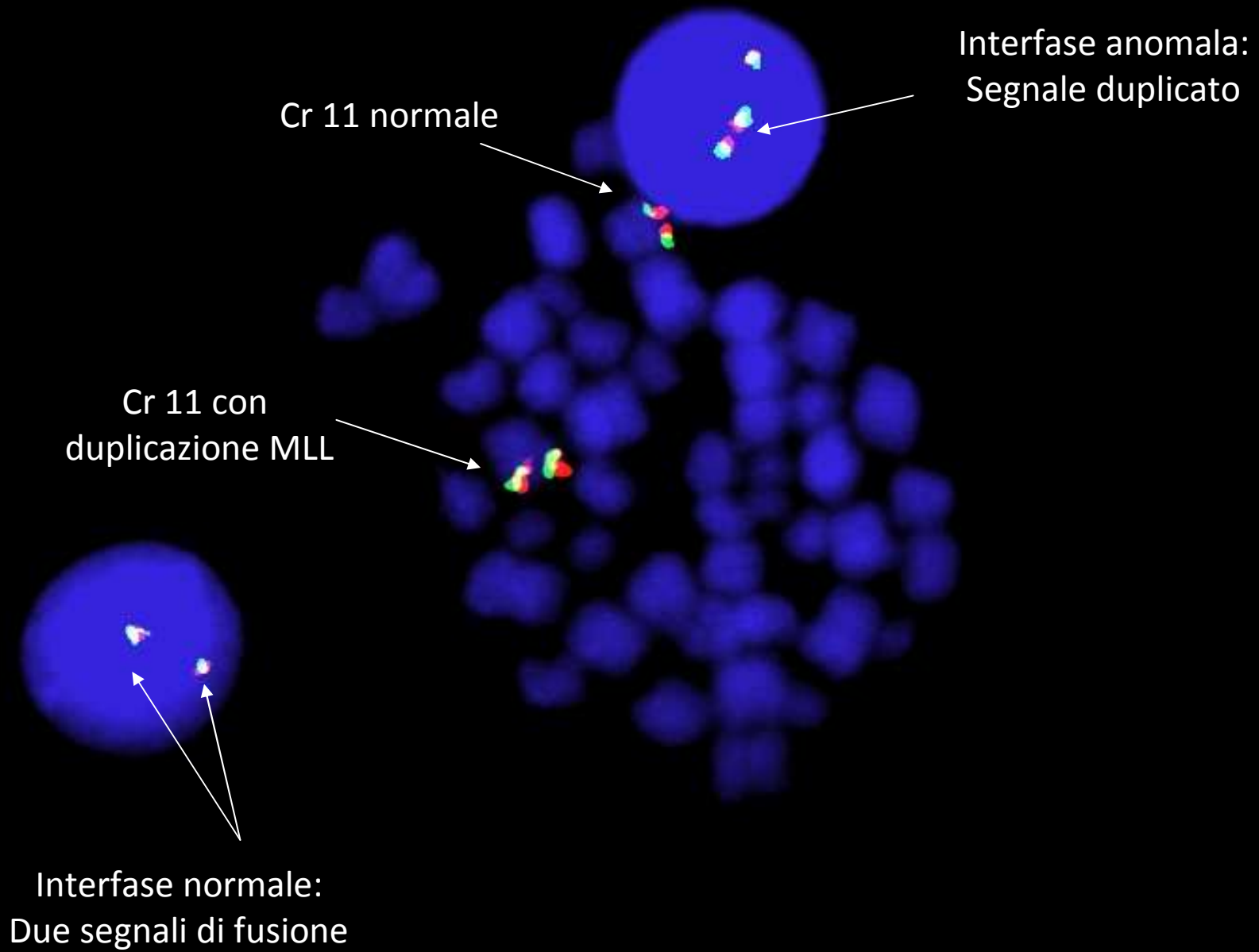


MLL dual color segregation system
Metaphase



MLL dual color segregation system
Interphase





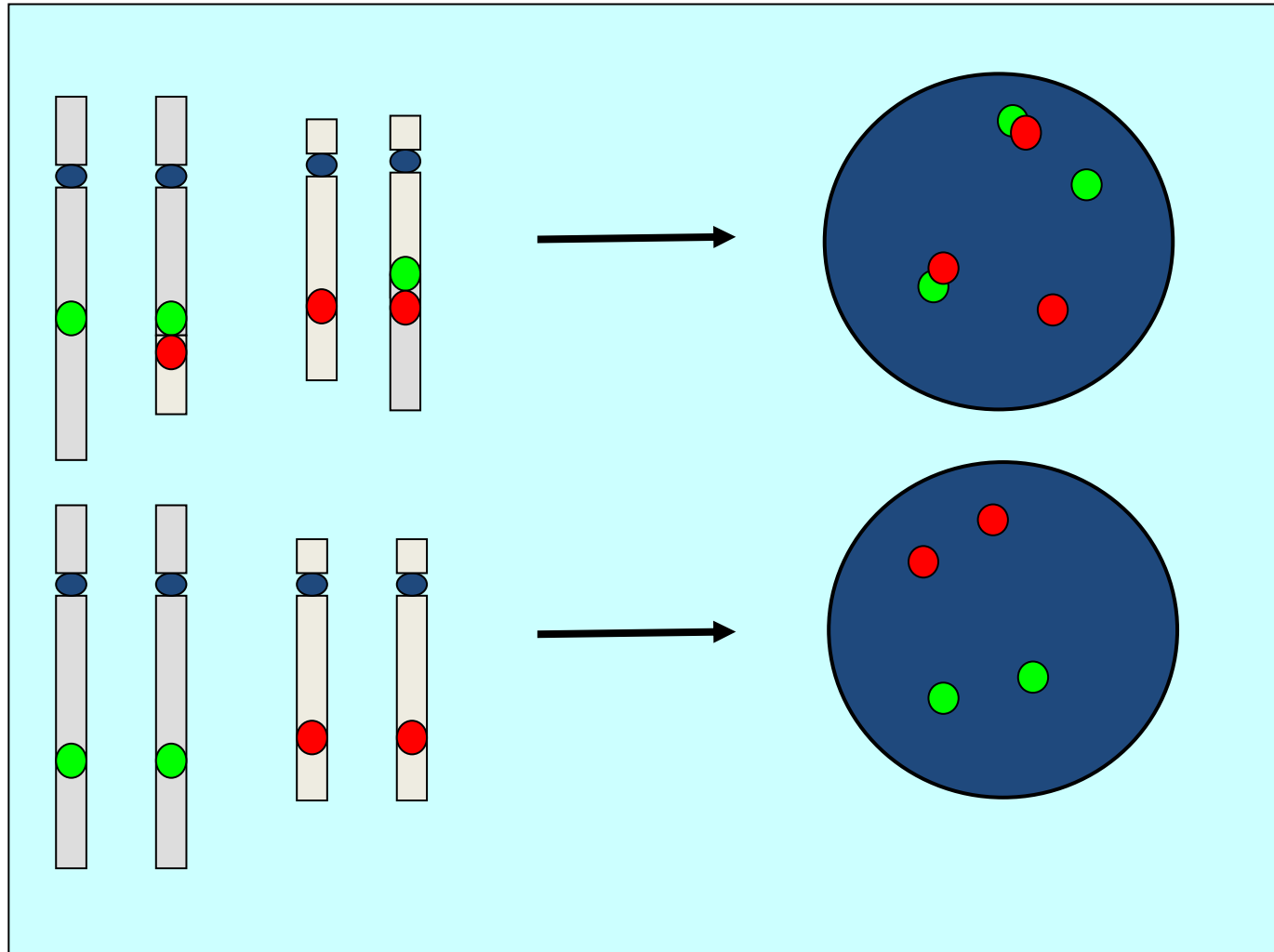
Cr 11 normale

Interfase anomala:
Segnale duplicato

Cr 11 con
duplicazione MLL

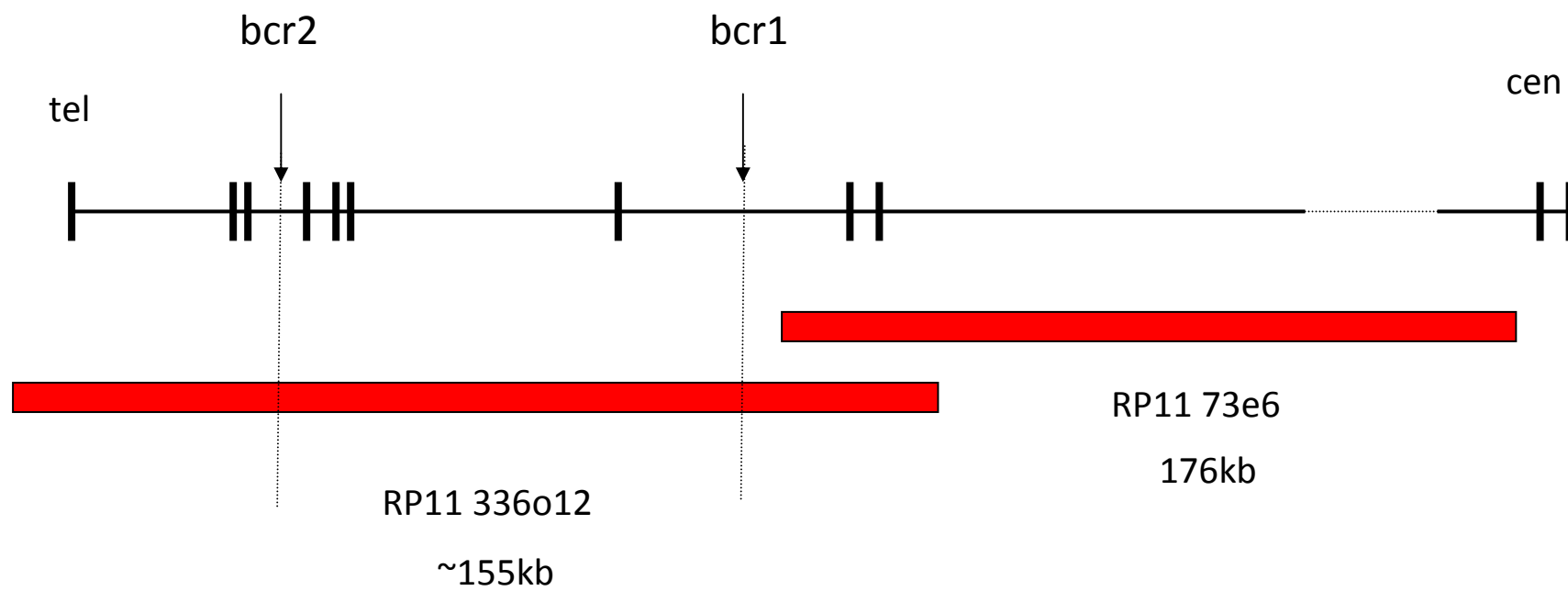
Interfase normale:
Due segnali di fusione

**SISTEMA COMBINATO I: I GENI COINVOLTI NELLA TRASLOCAZIONE
SONO MARCATI IN UNICO COLORE LA PRESENZA CONTEMPORANEA DI DOPPIA
FUSIONE E SEGNALI A SINGOLO COLORE RAPPRESENTA UNA DISTRIBUZIONE POSSIBILE,
IN UNA CELLULA 2N, SOLO IN CASO DI TRASLOCAZIONE**



AF9 probe systems

AF9 ~300kb

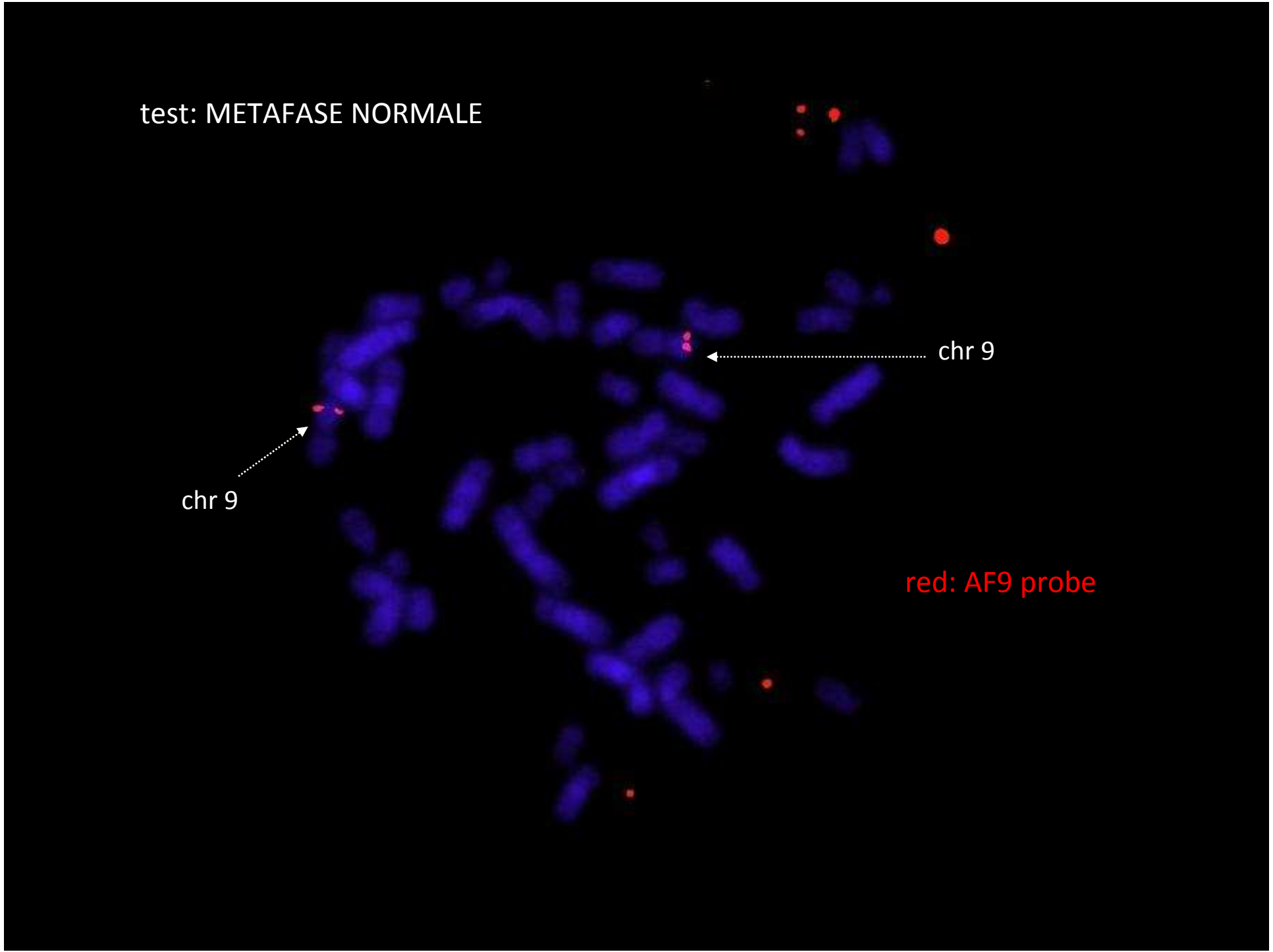


test: METAFASE NORMALE

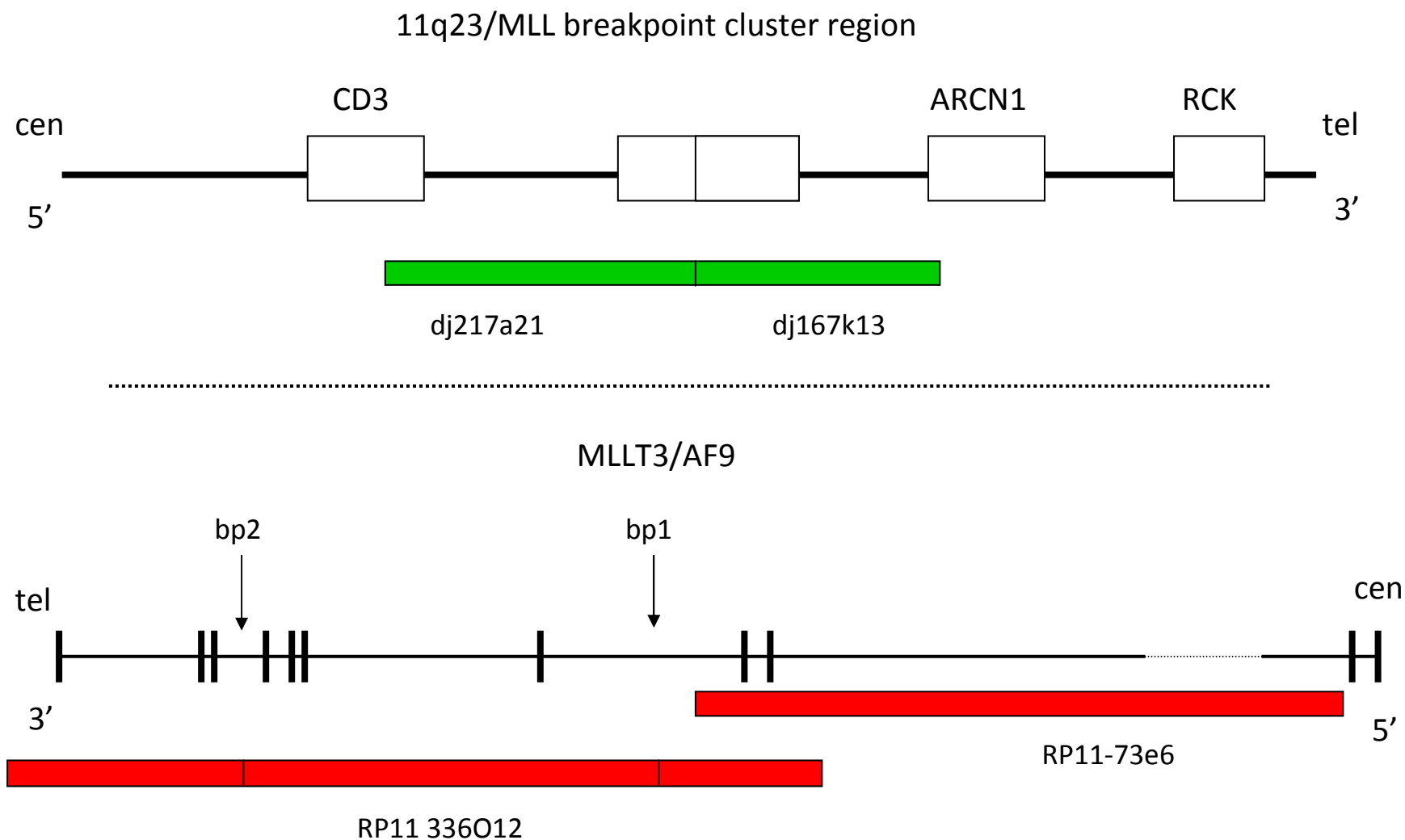
chr 9

chr 9

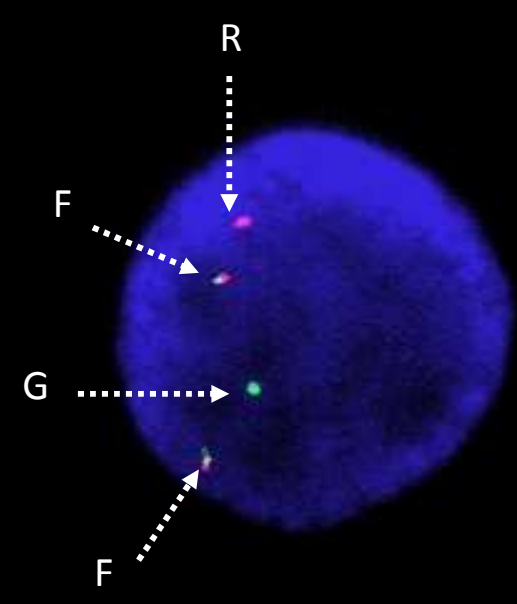
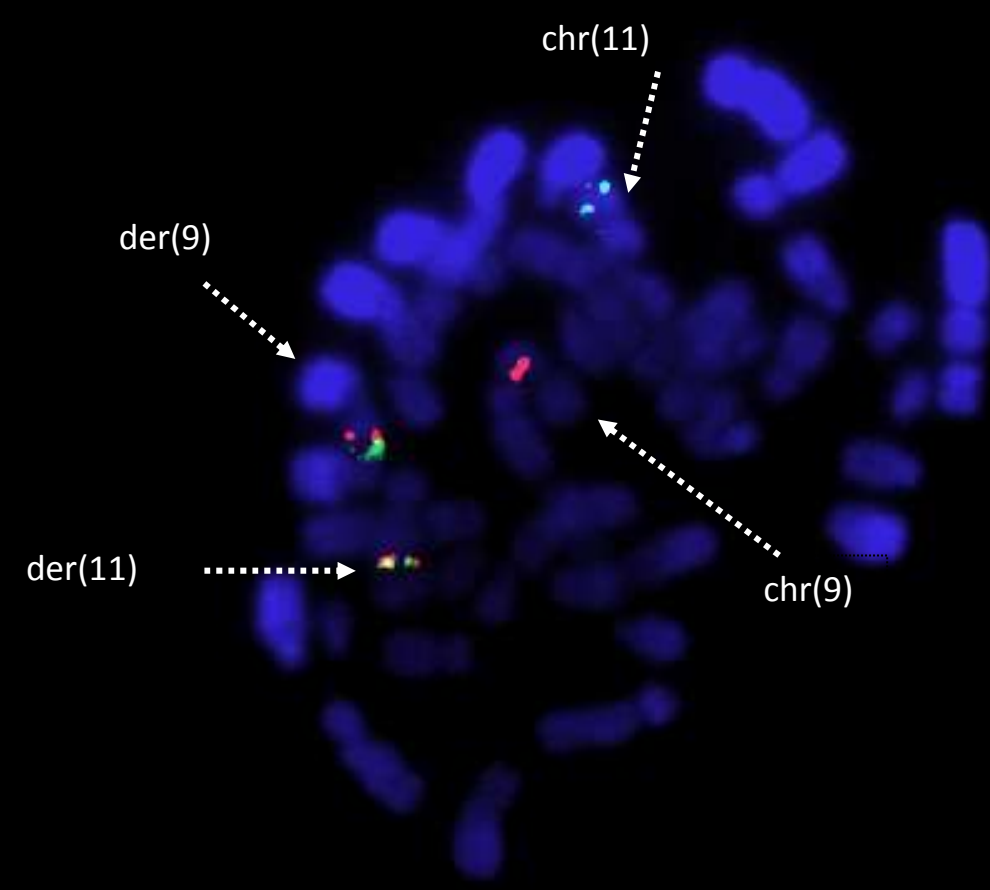
red: AF9 probe



t(9;11)(p22;q23) SISTEMA COMBIANTO SEGREGAZIONE: COLOCALIZZAZIONE, Dual Color Dual Fusion

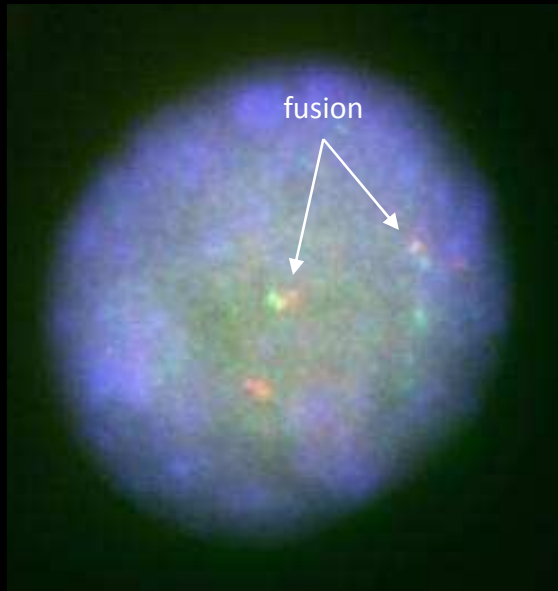


t(9;11)(p22;q23)
segregation/colocalization
combined system,
Dual Color Dual Fusion

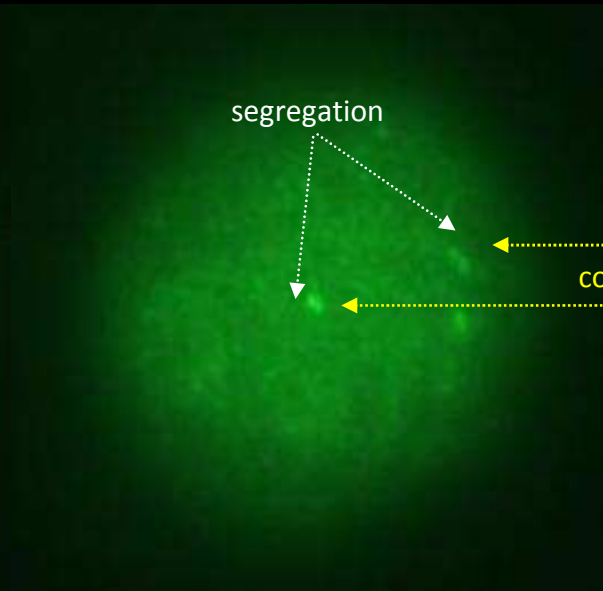


MLL/AF9 dual color dual fusion system: interphase

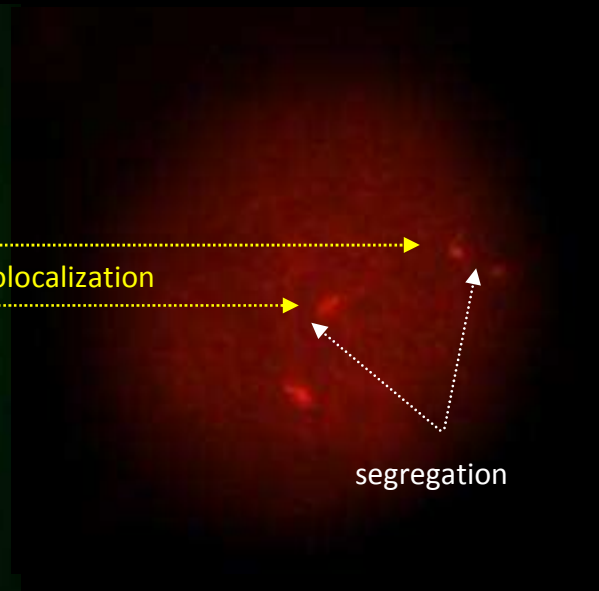
MLL/AF9
dual color dual fusion



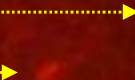
MLL probe set
Spectrum Green

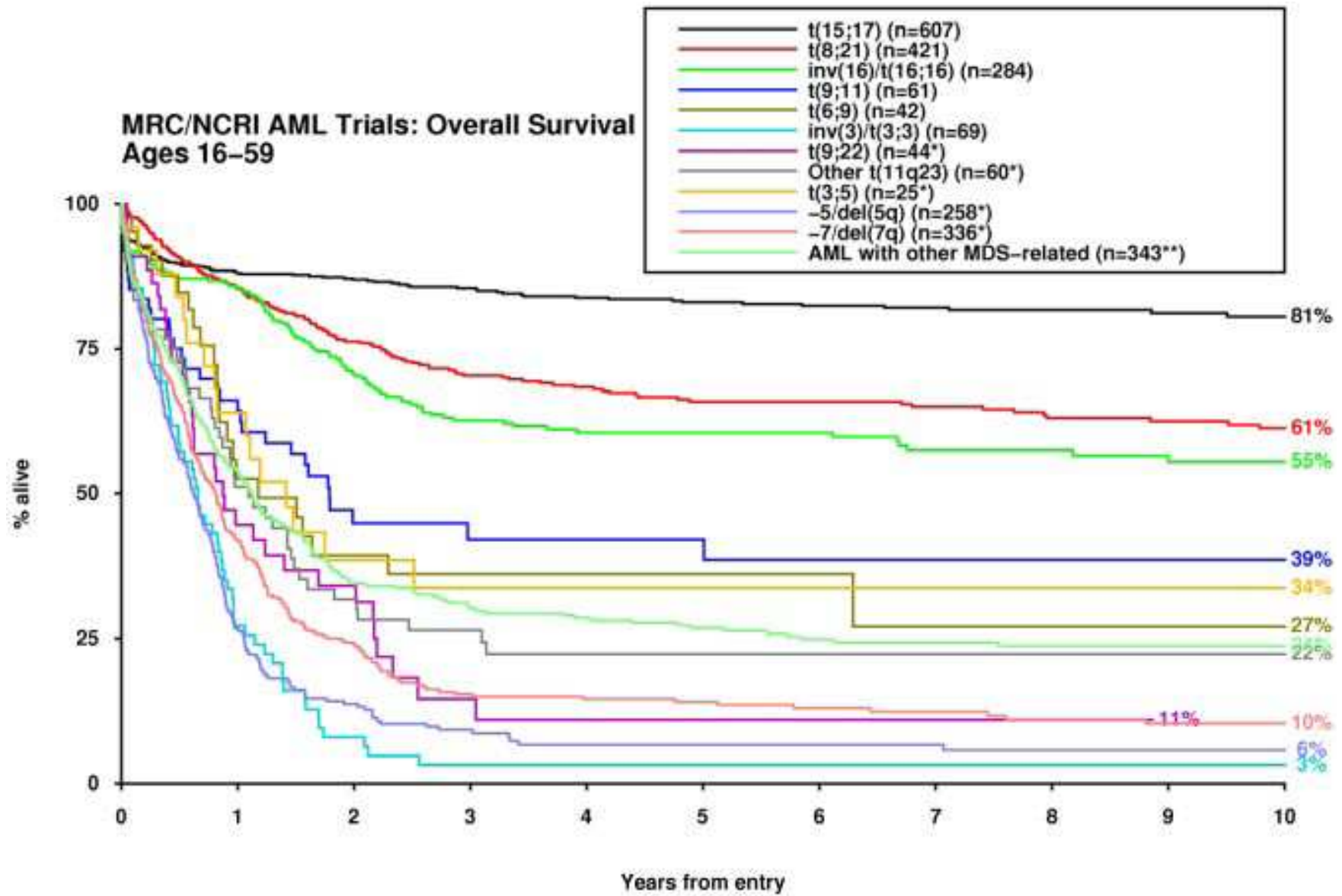


AF9 probe set
Spectrum Orange



colocalization





**Citogenetica e Citogenetica-Molecolare
come componenti integrativi
di sistemi diagnostico-prognostici**

L'analisi citogenetica nelle sindromi mielodisplastiche

DIFFERENTIAL DIAGNOSIS OF Myelodysplastic Syndromes

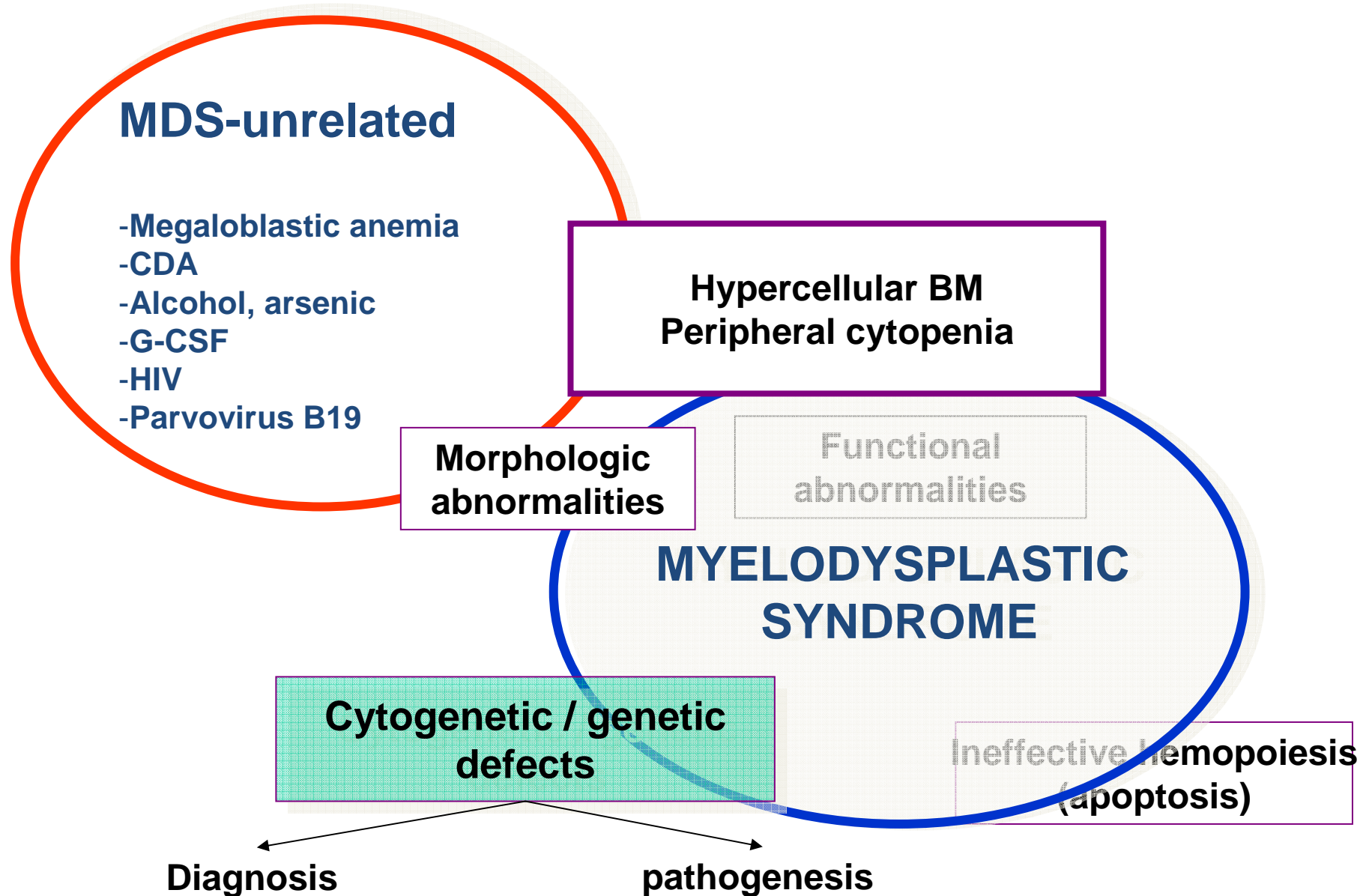


Table 2

IPSS Classification Criteria

Prognostic variable	Score				
	0	0.5	1.0	1.5	2.0
Bone marrow blasts	< 5%	5%–10%	—	11%–20%	21%–30%
Karyotype ^a	Good	Intermediate	Poor		
Cytopenias	0/1	2/3			

IPSS scores for risk groups are as follows:
 Low = 0; Int-1 = 0.5–2.0; Int-2 = 1.5–2.0; High = 2.5

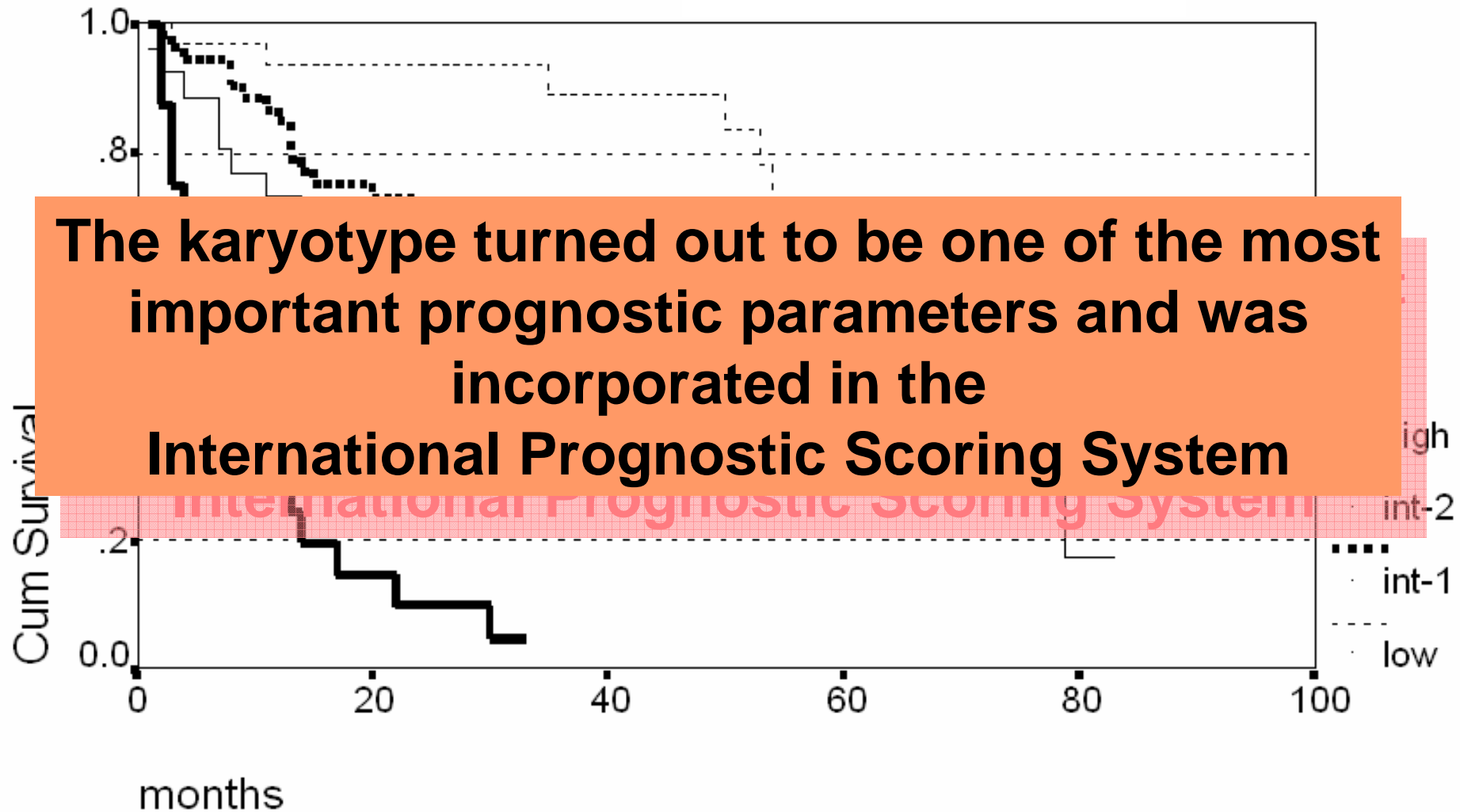
^aGood = normal, –Y, del(5q), del(20q); poor = complex (3 abnormalities) or chromosome 7 anomalies; intermediate = other abnormalities.

Int = intermediate; IPSS = International Prognostic Scoring System.

This research was originally published in *Blood*. Adapted, with permission, from Greenberg P et al.[6] Copyright © 1997 by the American Society of Hematology.

Risk Classification according to IPSS in 134 MDS patients studied at Section of Haematology, University of Ferrara (1998-2006)

% blasts, Karyotype, cytopenia



CYTOGENETIC ABNORMALITIES IN MDS

STRUCTURAL ABERRATIONS

BALANCED
(translocations and inversions)

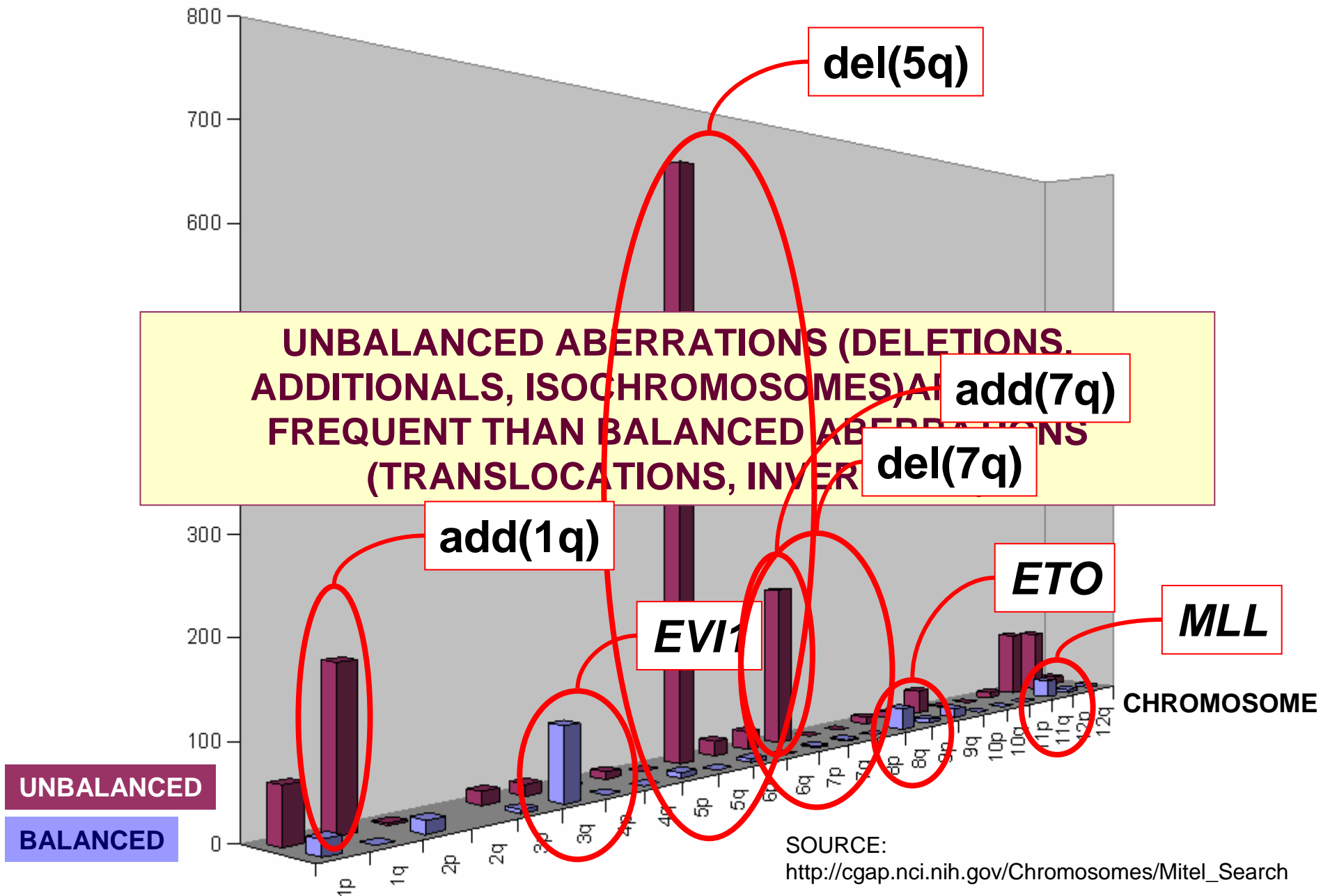
UNBALANCED
(deletions, additional, iso)

NUMERICAL ABERRATIONS

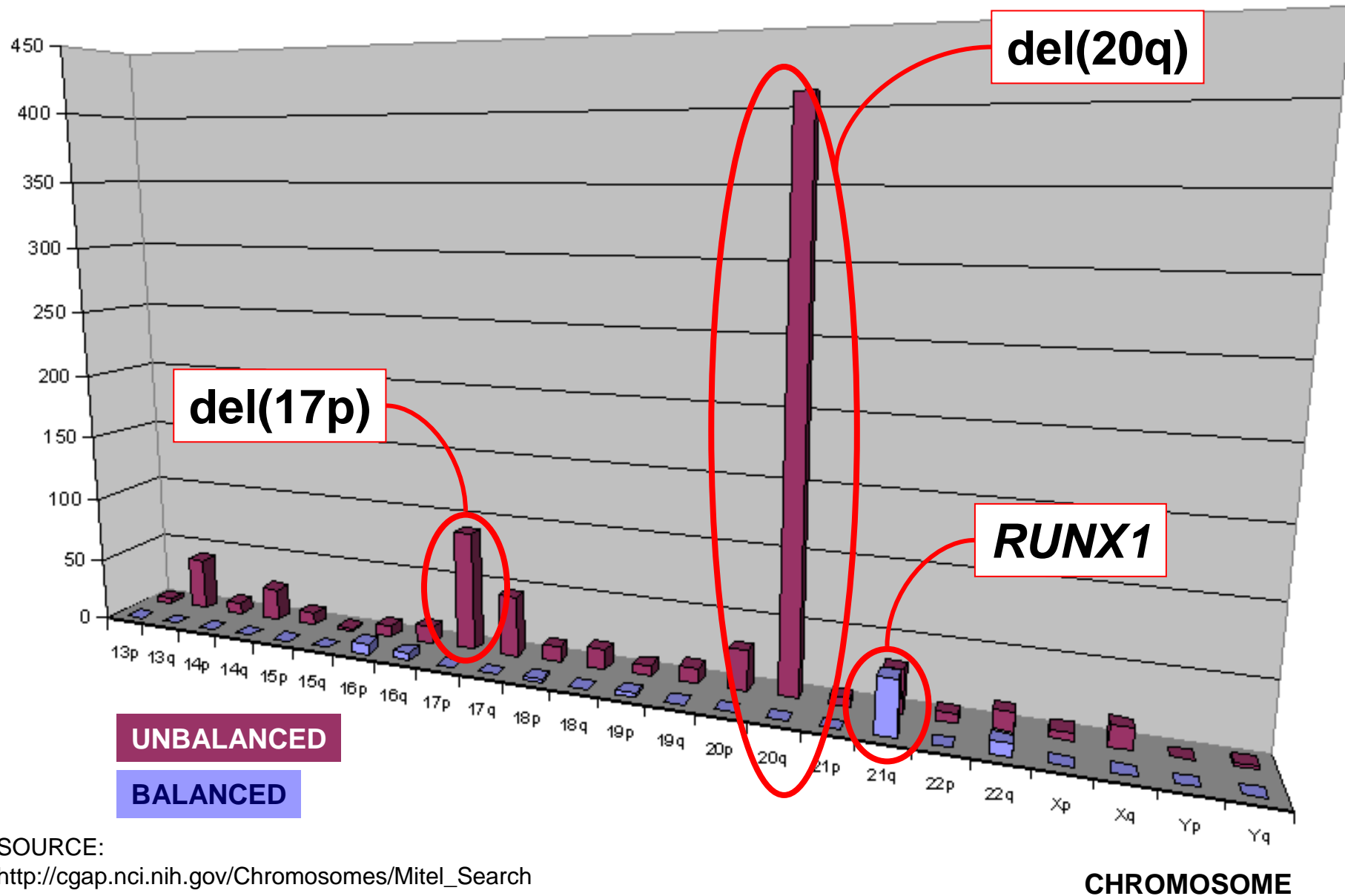
TRISOMIES

MONOSOMIES

STRUCTURAL ABERRATIONS (balanced and unbalanced) IN MDS



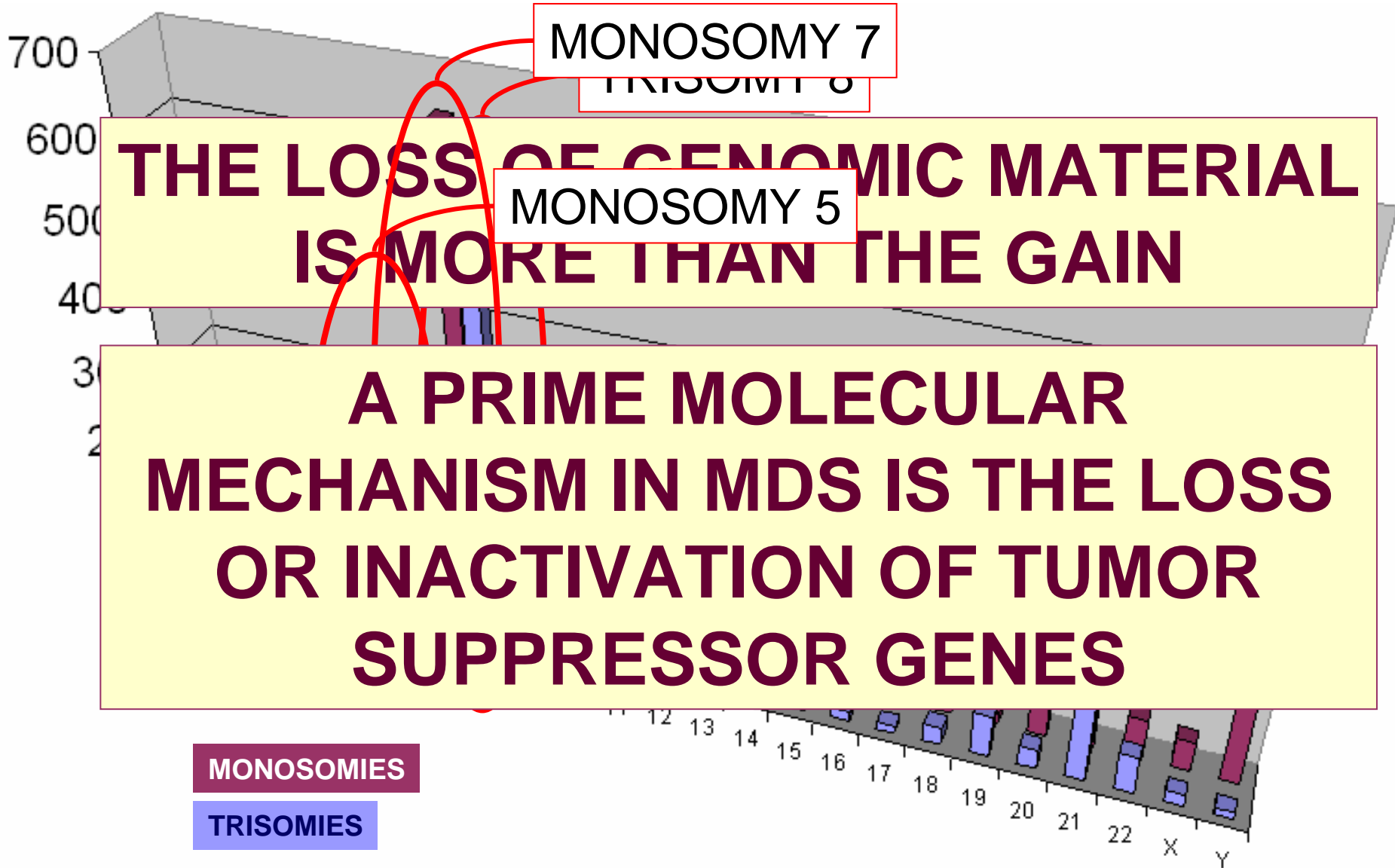
STRUCTURAL ABERRATIONS (balanced and unbalanced) IN MDS



SOURCE:
http://cgap.nci.nih.gov/Chromosomes/Mitel_Search

CHROMOSOME

NUMERICAL ABERRATIONS IN MDS

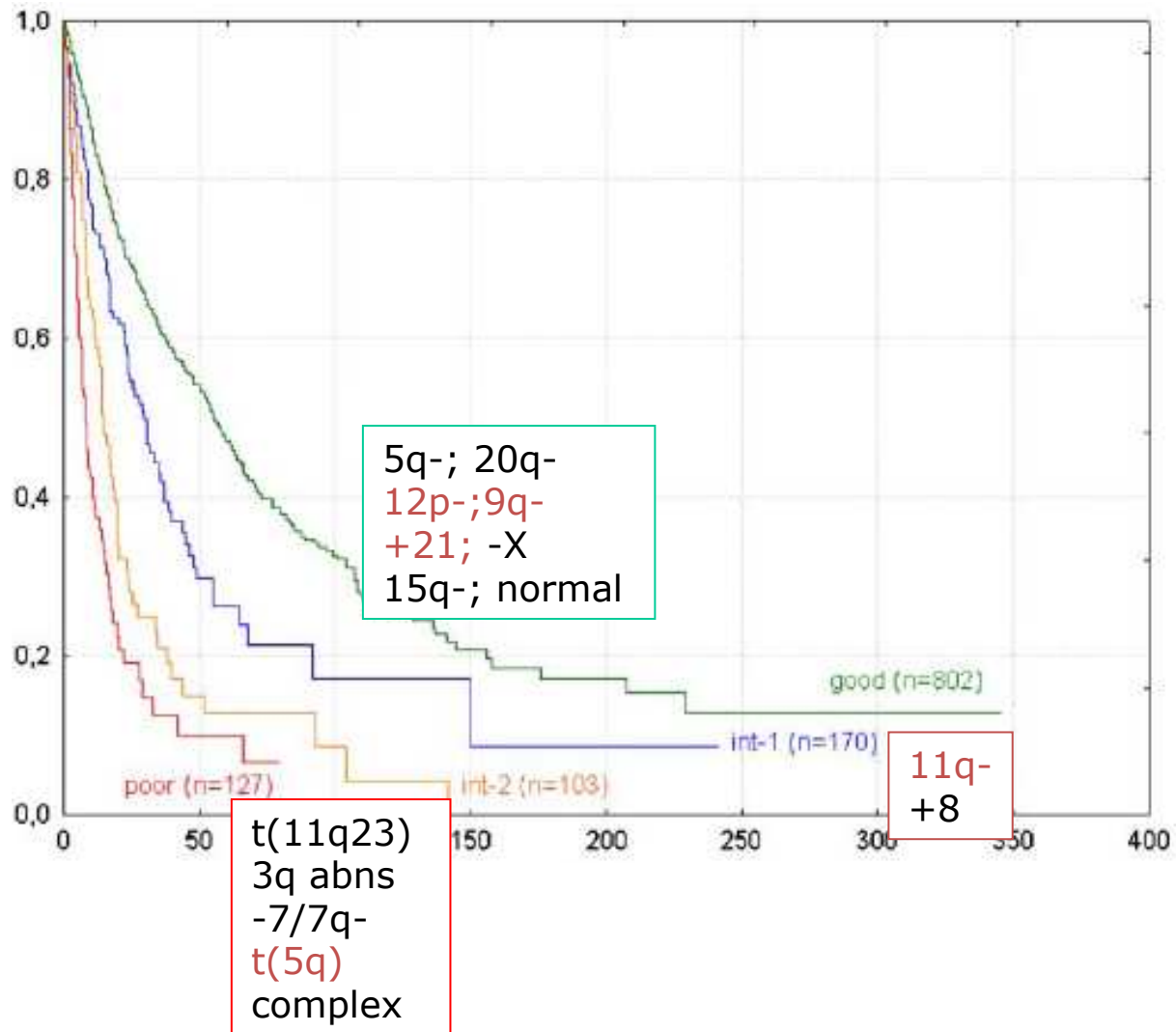


SOURCE:

http://cgap.nci.nih.gov/Chromosomes/Mitel_Search

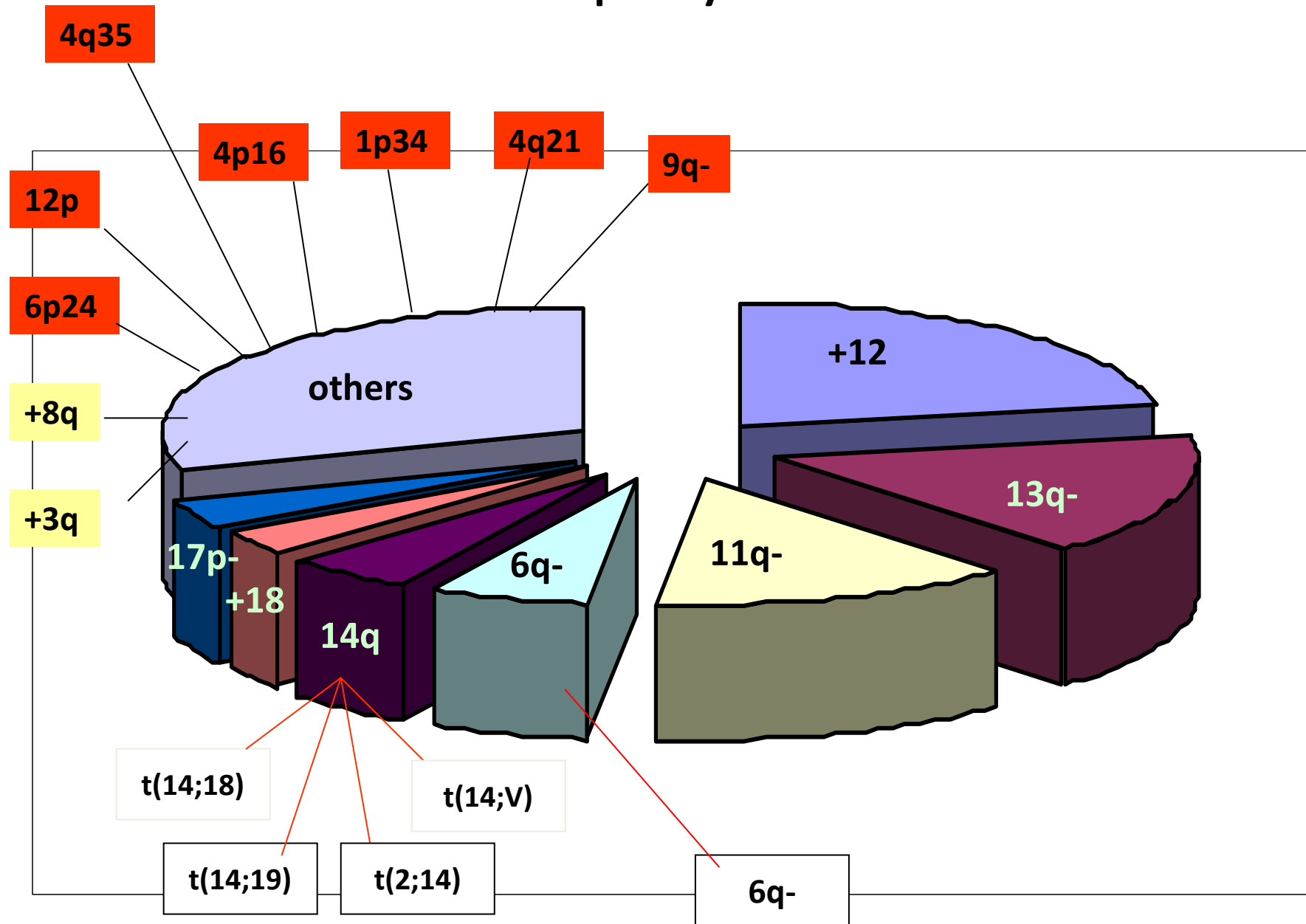
PROGNOSTIC SIGNIFICANCE OF CHROMOSOME ABERRATIONS IN MDS

Haase D. Ann Hematol, 87: 515, 2008

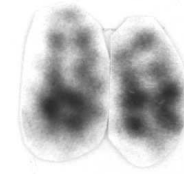
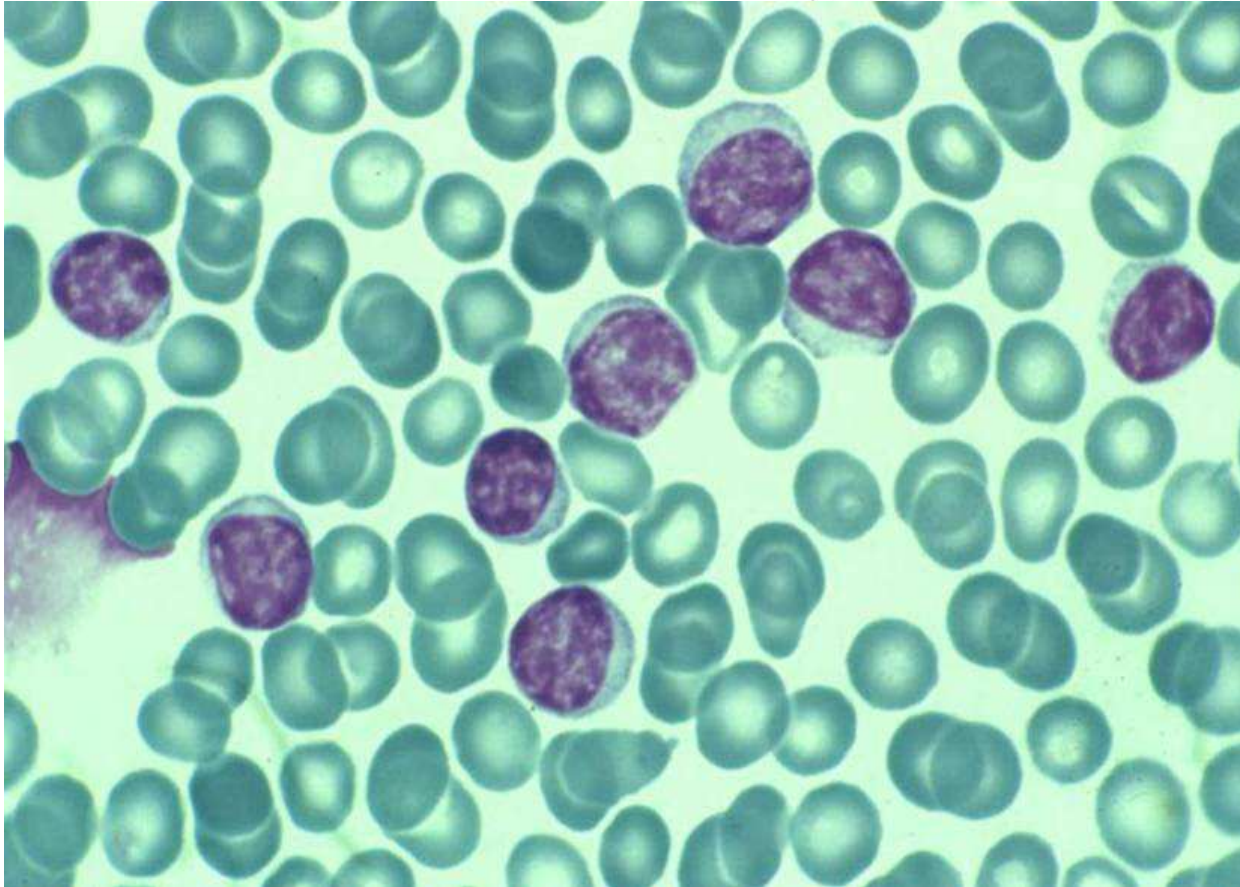


L'analisi citogenetica nella leucemia linfatica cronica

B-CLL: Frequency of chromosome aberrations



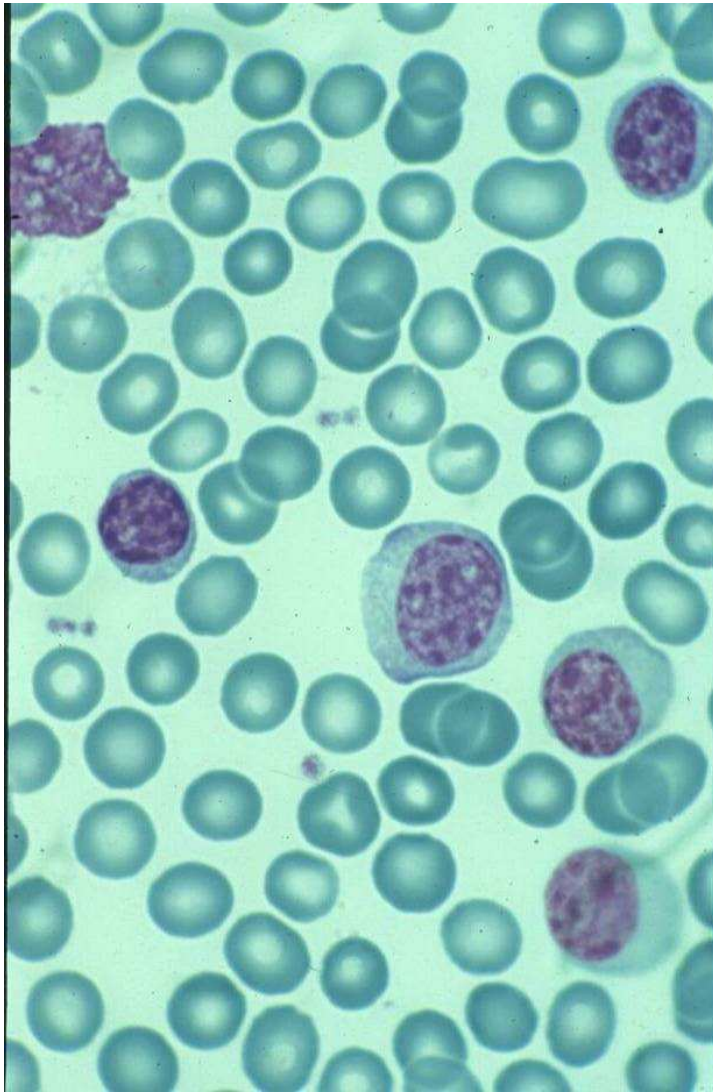
13q- and typical CLL



13q-

- “mutated” IgVh
- CD38- ZAP70-
- Typical CLL

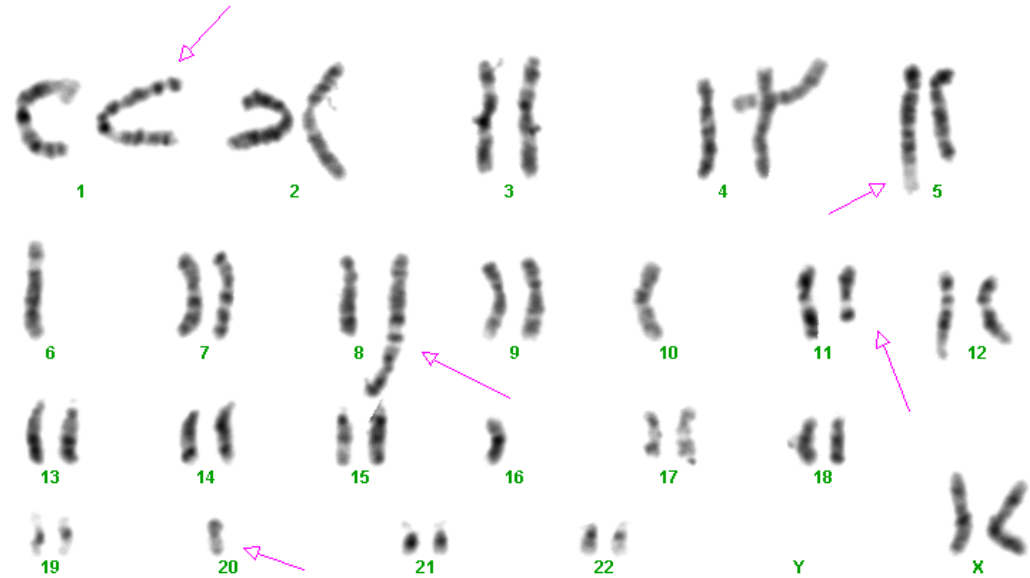
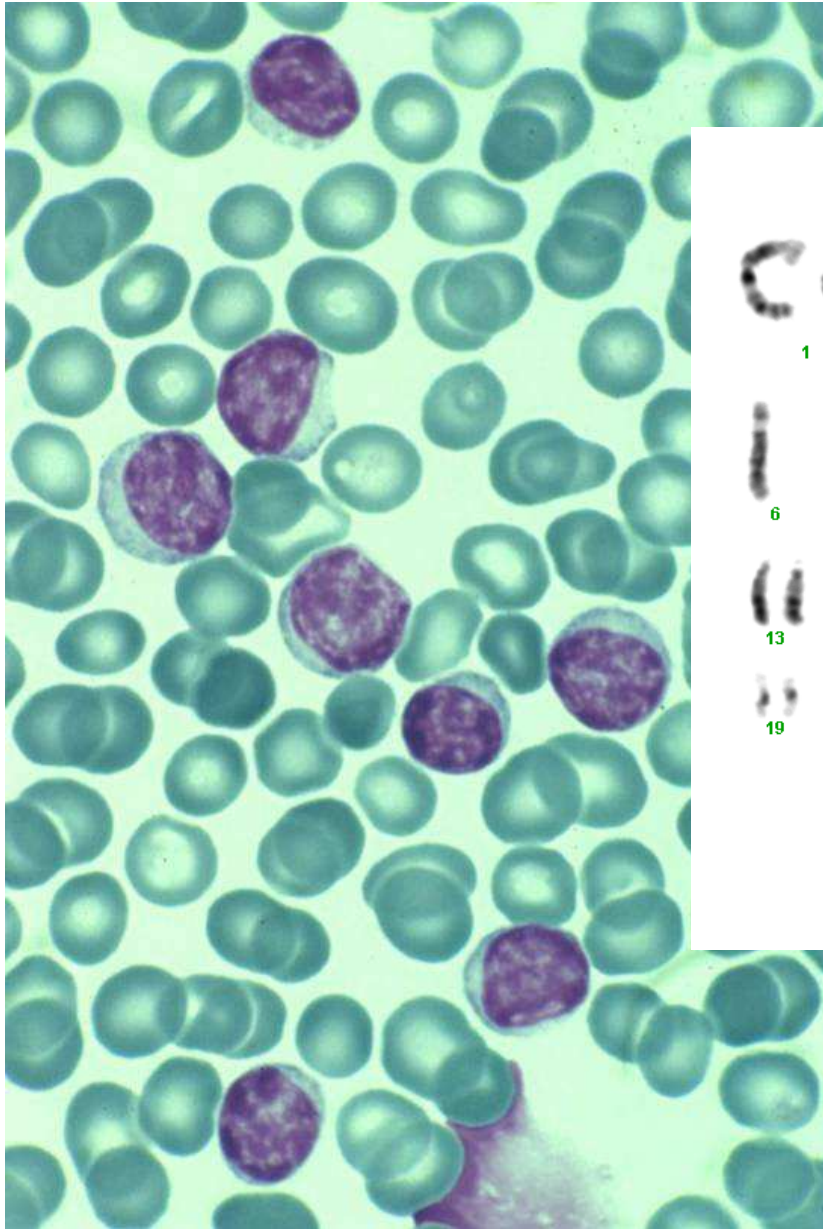
Trisomy 12 and atypical CLL



- “approx 50% mutated” IgVh

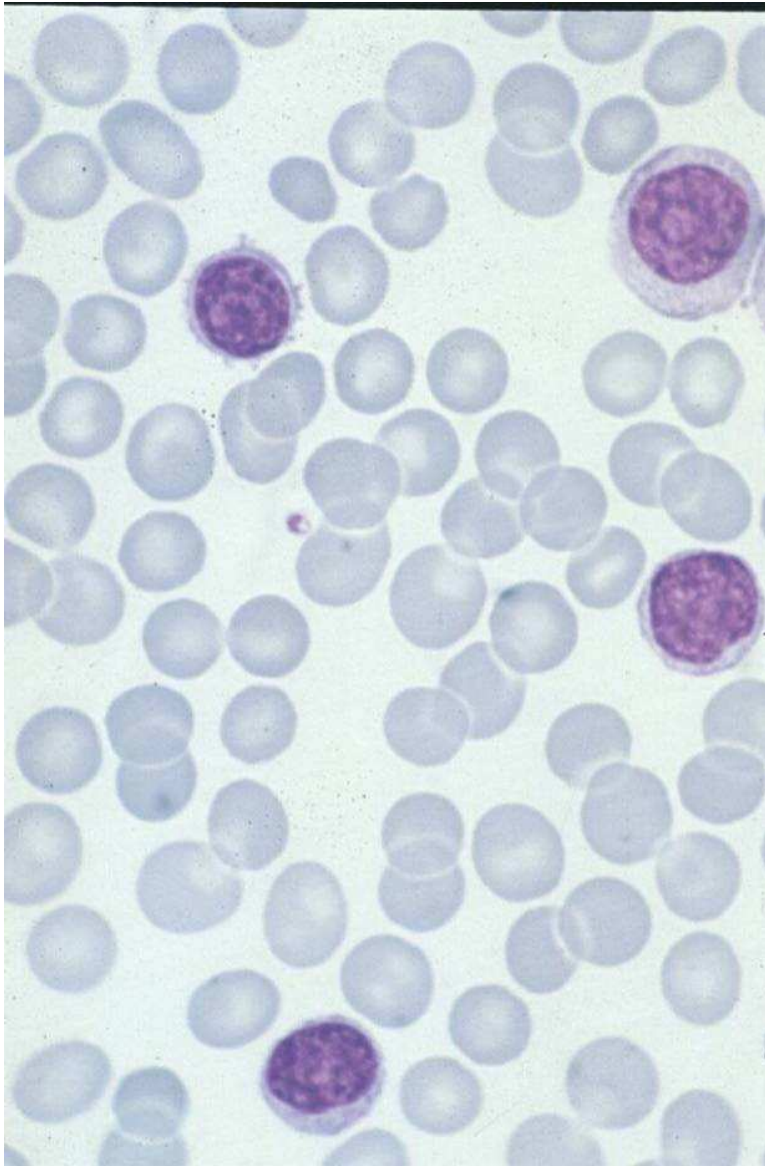
- CD38-/+ ZAP70+/-

11q- and typical CLL

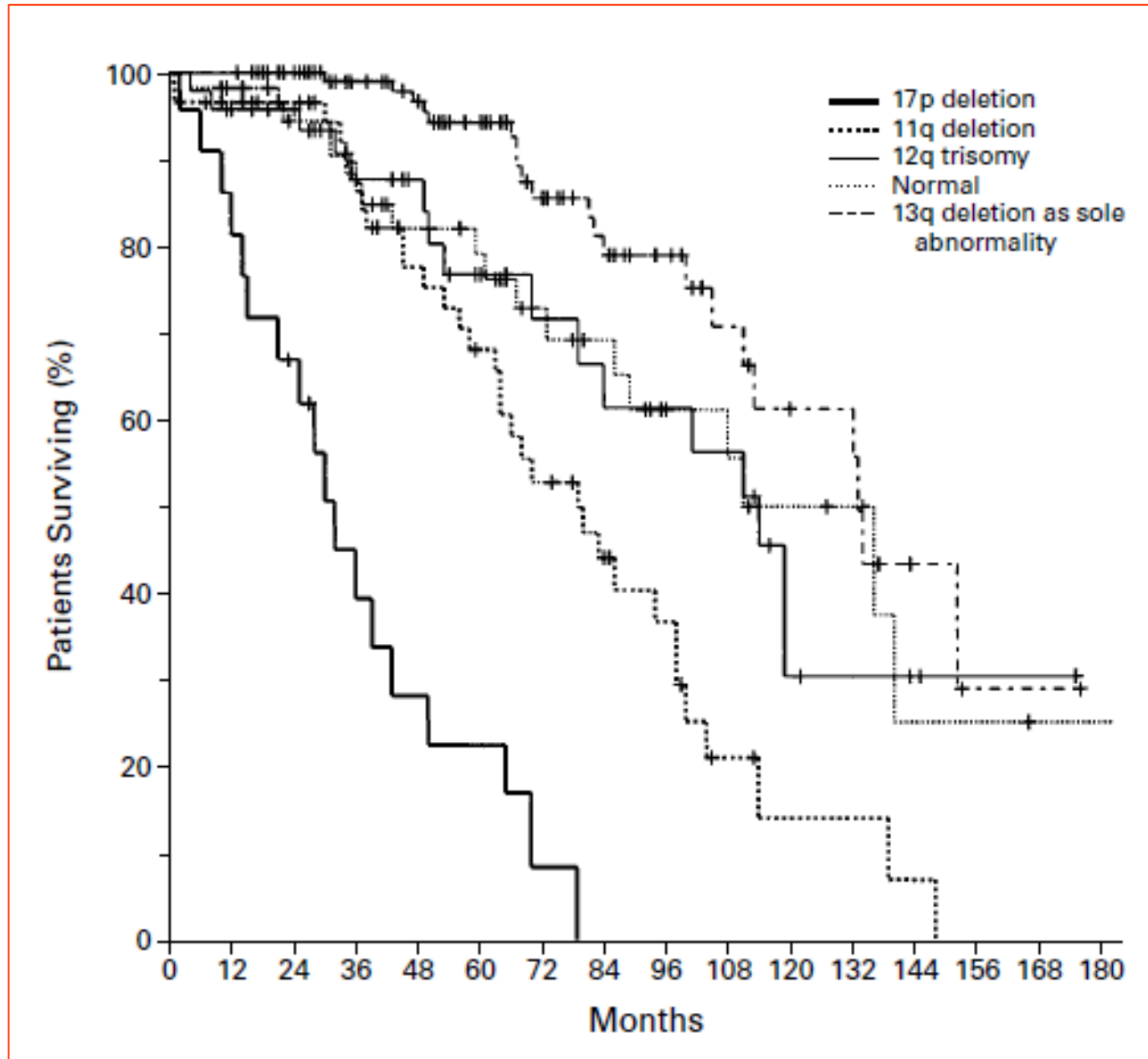


- “unmutated” IgVh
- CD38+ ZAP70+/-
- ATM deletion
- COMPLEX KARYOTYPE

17p- and CLL/PL



- “unmutated” IgVh
- CD38+/ZAP70+
- TP53 deletion
- refractory to purine analogs



Conclusioni

L'analisi citogenetica è necessaria per la diagnosi di alcune malattie oncoematologiche (LMC, APL, CEL, etc.) in cui il trattamento si basa sulla anatomia genetica-molecolare della lesione.

L'analisi citogenetica è necessaria per un completo inquadramento prognostico

L'analisi citogenetico-molecolare rappresenta un metodo rapido ed integra lo studio citogenetico convenzionale in casi a presentazione atipica

L'analisi citogenetico-molecolare offre la possibilità di un rapido screening di lesioni note alla diagnosi su nuclei in interfase



