

# Biology of High-Risk Chronic Lymphocytic Leukemia

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*Unmet challenges in high risk hematological malignancies:  
from benchside to clinical practice (2<sup>nd</sup> edition)*

*Turin, September 13-14, 2021*



AO S. Croce e Carle  
Cuneo



2<sup>nd</sup> edition  
**Unmet challenges in high risk  
hematological malignancies:  
from benchside to clinical practice**

**Turin, September 13-14, 2021**  
Starhotels Majestic


*Scientific board:*  
**Marco Ladetto** (Alessandria)  
**Umberto Vitolo** (Candiolo-TO)



**Disclosures of NAME SURNAME**

Company name	Research support	Employee	Consultant	Stockholder	Speakers bureau	Advisory board	Other
AbbVie						x	
Roche						x	
Janssen						x	
Sanofi						x	

## high-risk cancer

 (hy-risk KAN-ser)

Cancer that is likely to recur (come back), or spread.

## Why does cancer recur or spread?

- standard treatments are inadequate in the short or long run;

## Why current treatments are inadequate?

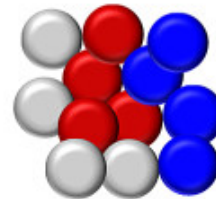
- unable to fully abrogate mechanisms of disease progression;
- CLL is heterogenous clonal disease under continuous evolution;



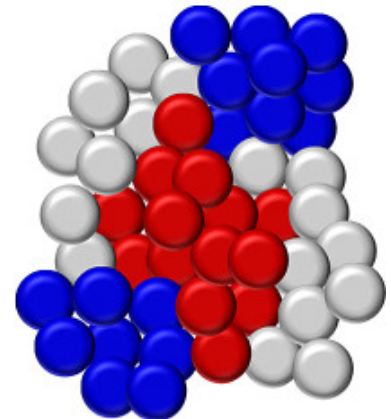
MBL



CLL



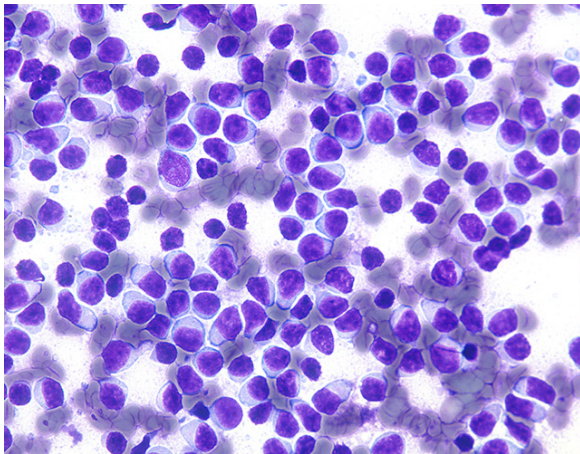
Progression



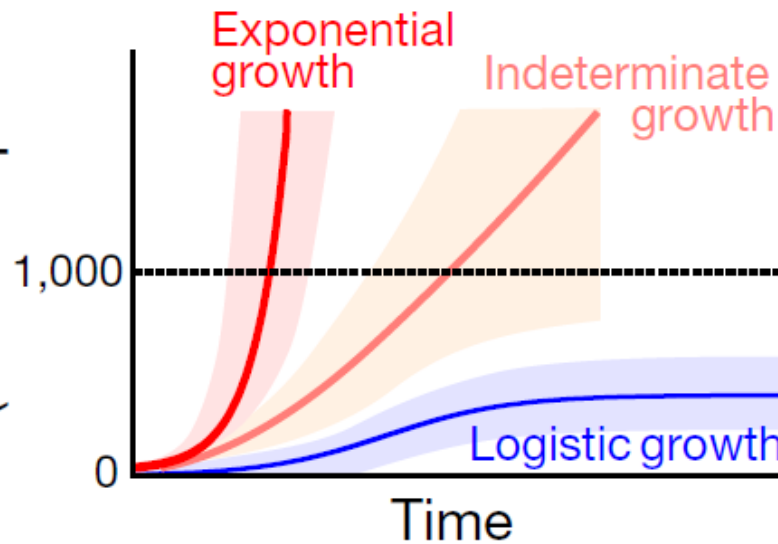


# CLL paradigm shift

progressive accumulation  
of functionally  
incompetent lymphocytes

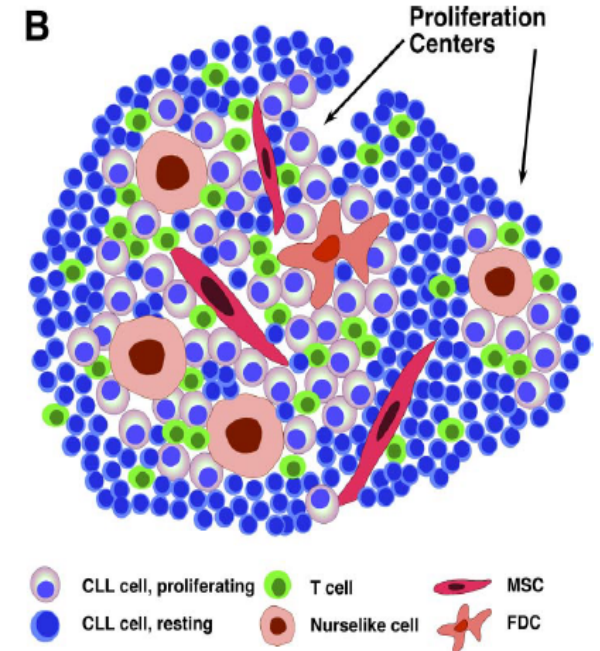


WBC ( $10^9$  cells per litre)



Gruber M et al., *Nature*. 2019 Jun;570(7762):474-479

CLL proliferation centers



Burger et al., *Blood*. 2009;114:3367-3375

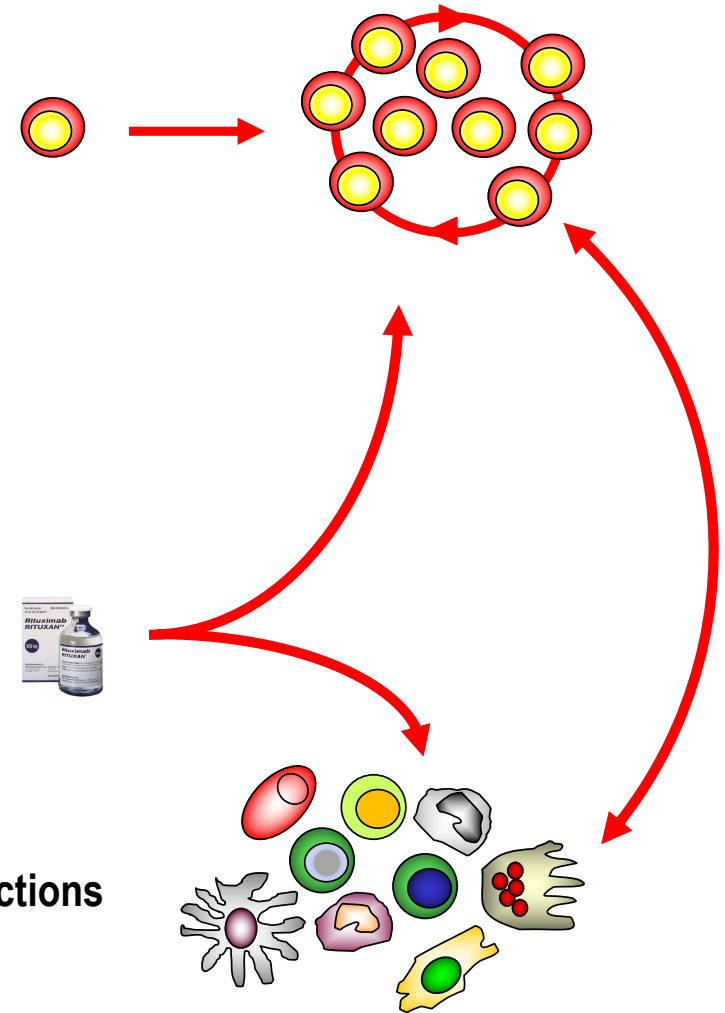
# Drivers of disease progression in CLL:

## Intrinsic drivers:

- genetic instability
- BCR (IGHV-M/UM)
- genetic abnormalities
- epigenetic alterations

## Extrinsic drivers:

- treatments 
- microenvironment interactions



## Innate high-risk (primary)

- genetic instability
- BCR (IGHV-M/UM)
- genetic abnormalities
- epigenetic alterations

- **treatments**



- 
- The diagram illustrates the mechanism of Rituximab (anti-CD20 antibody) in B-cell lymphoma treatment. It shows a cycle where B-cells proliferate, then are targeted by Rituximab, leading to cell death and the release of tumor cells. The text "ctions" is partially visible at the bottom left.

# Drivers of disease progression in CLL:

Innate high-risk (primary)

## Intrinsic drivers:

- genetic instability
- BCR (IGHV-M/UM)
- genetic abnormalities
- epigenetic alterations

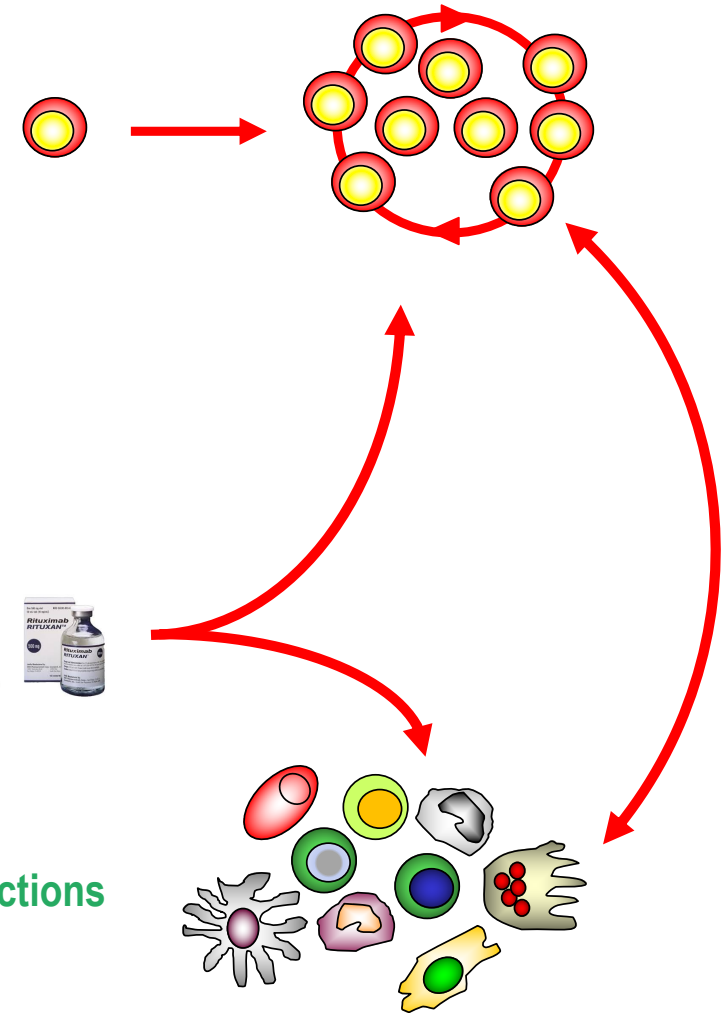
Adaptive high-risk (secondary)

## Extrinsic drivers:

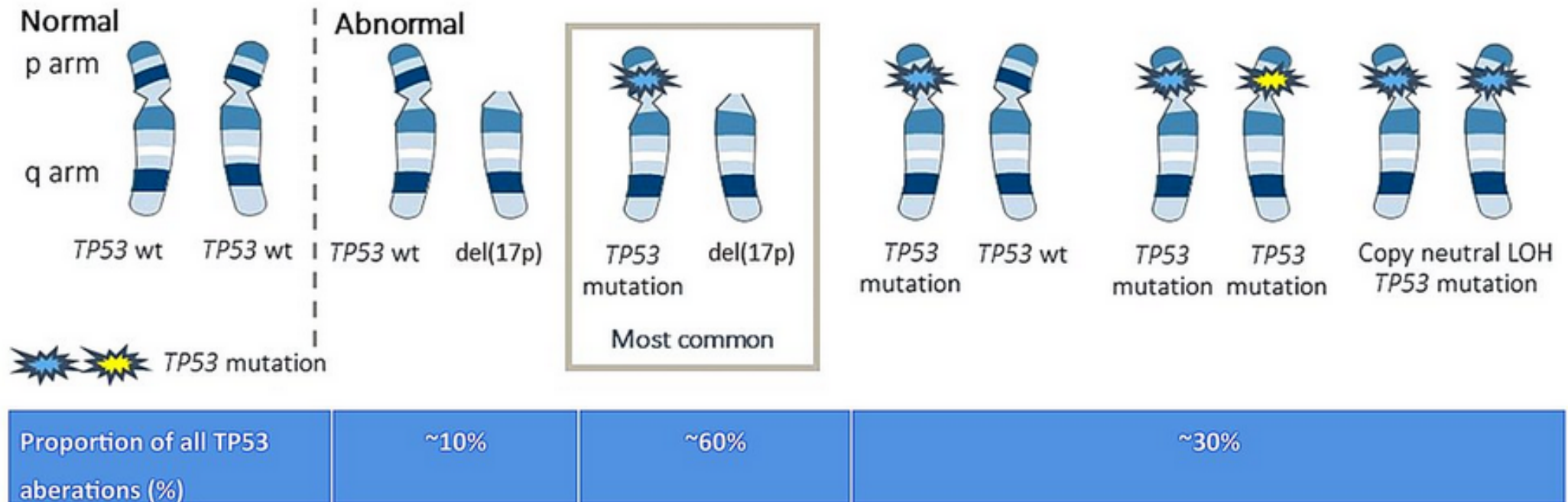
- treatments



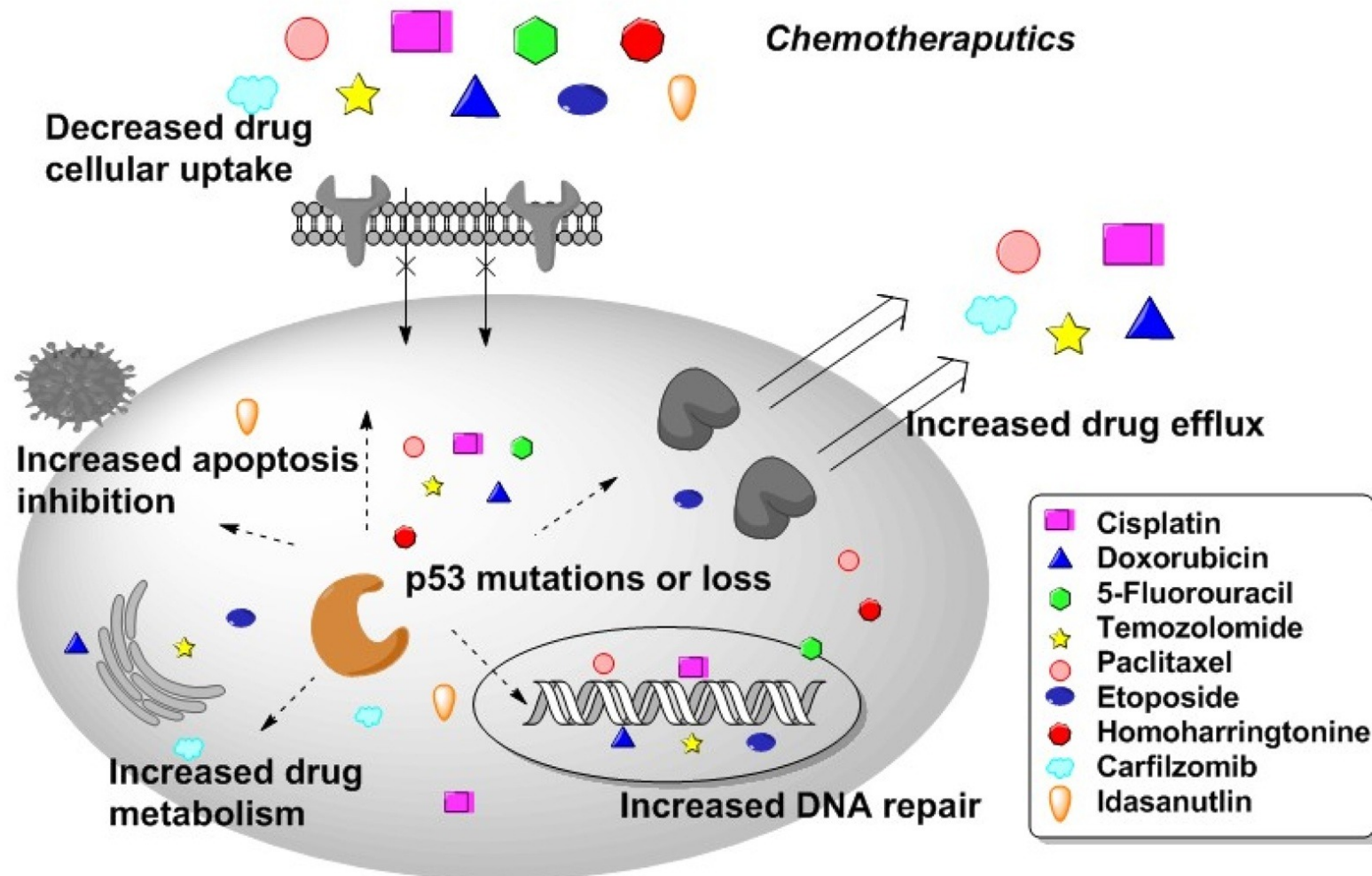
- microenvironment interactions



## TP53 aberrations in CLL

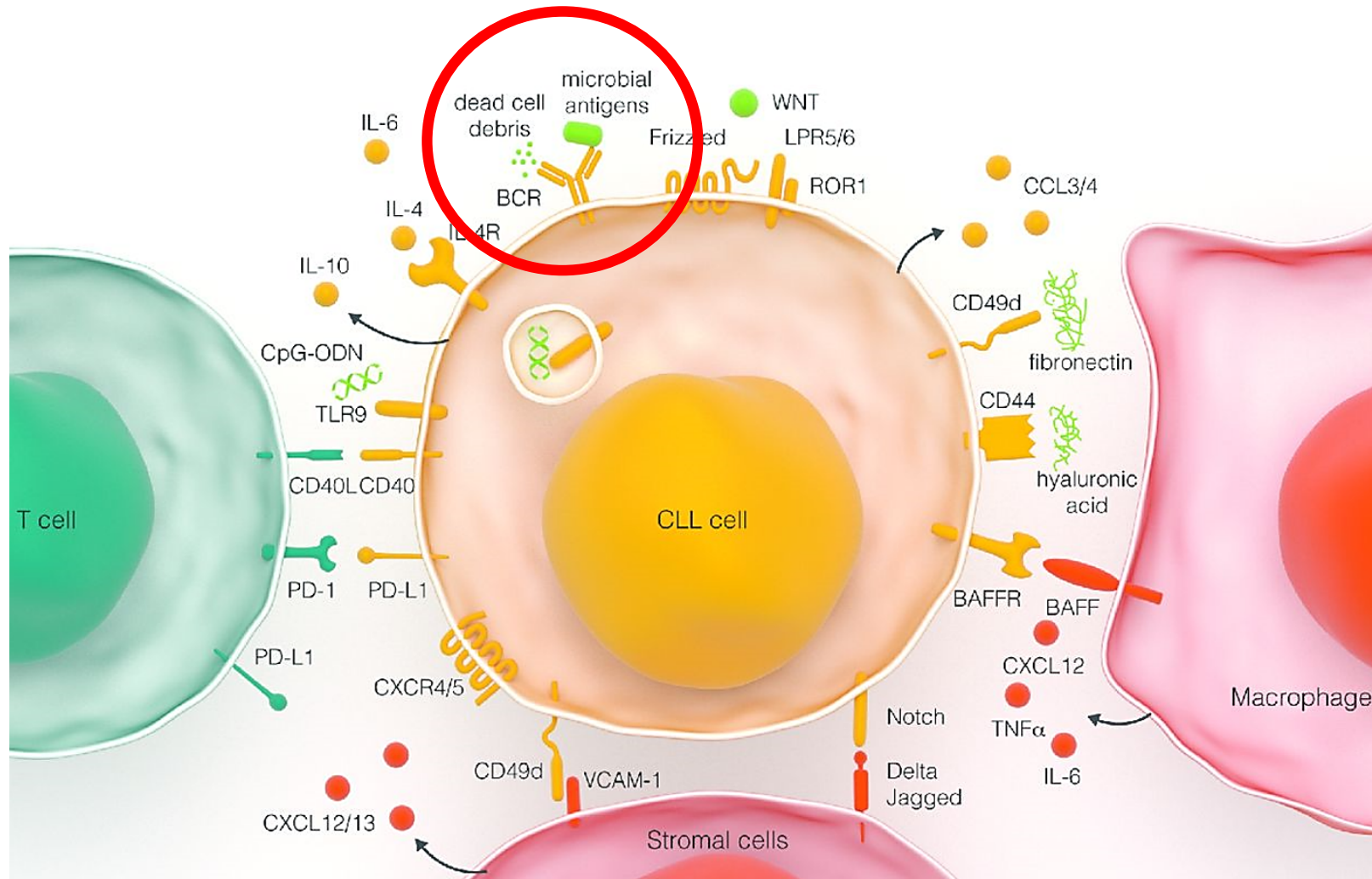


# Mechanisms for p53 mutation-associated chemotherapy resistance



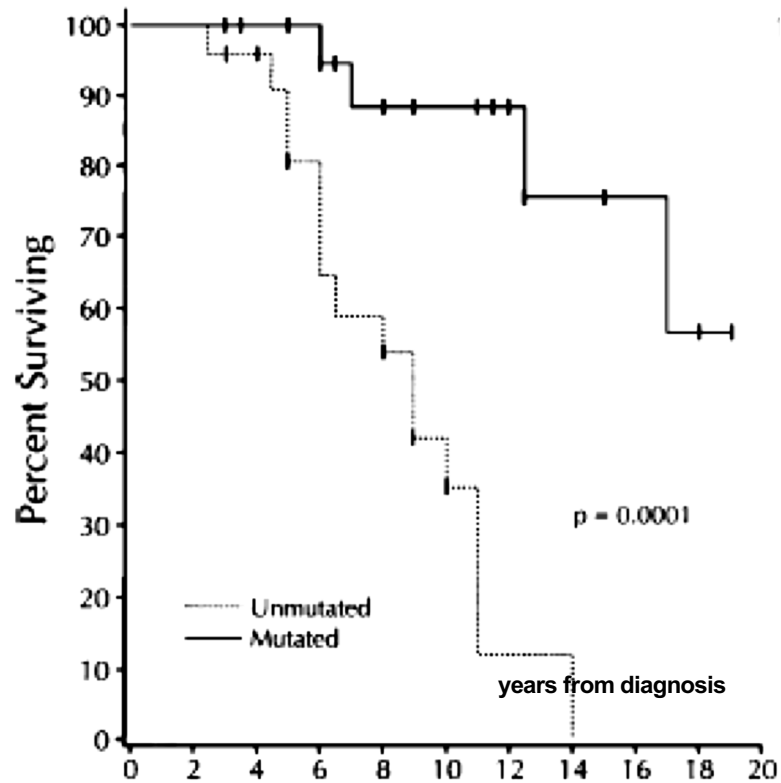


## CLL cell interactions in the TME

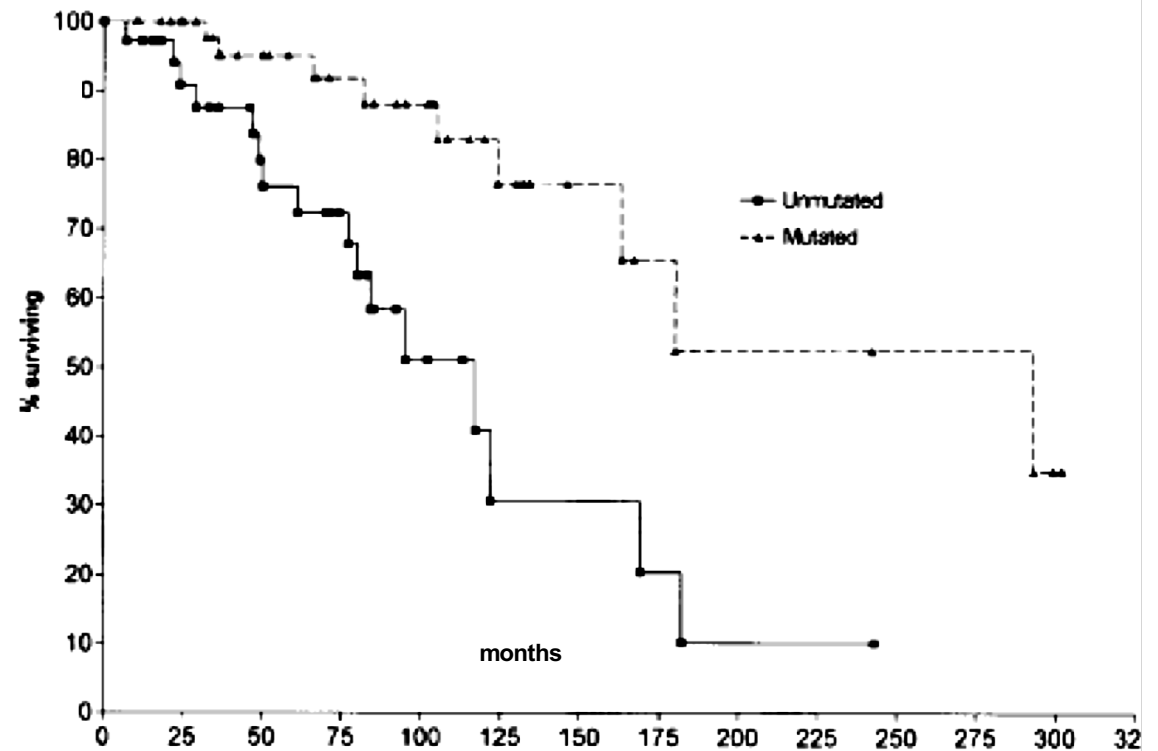




## Prognostic significance of IgVH mutational status in CLL

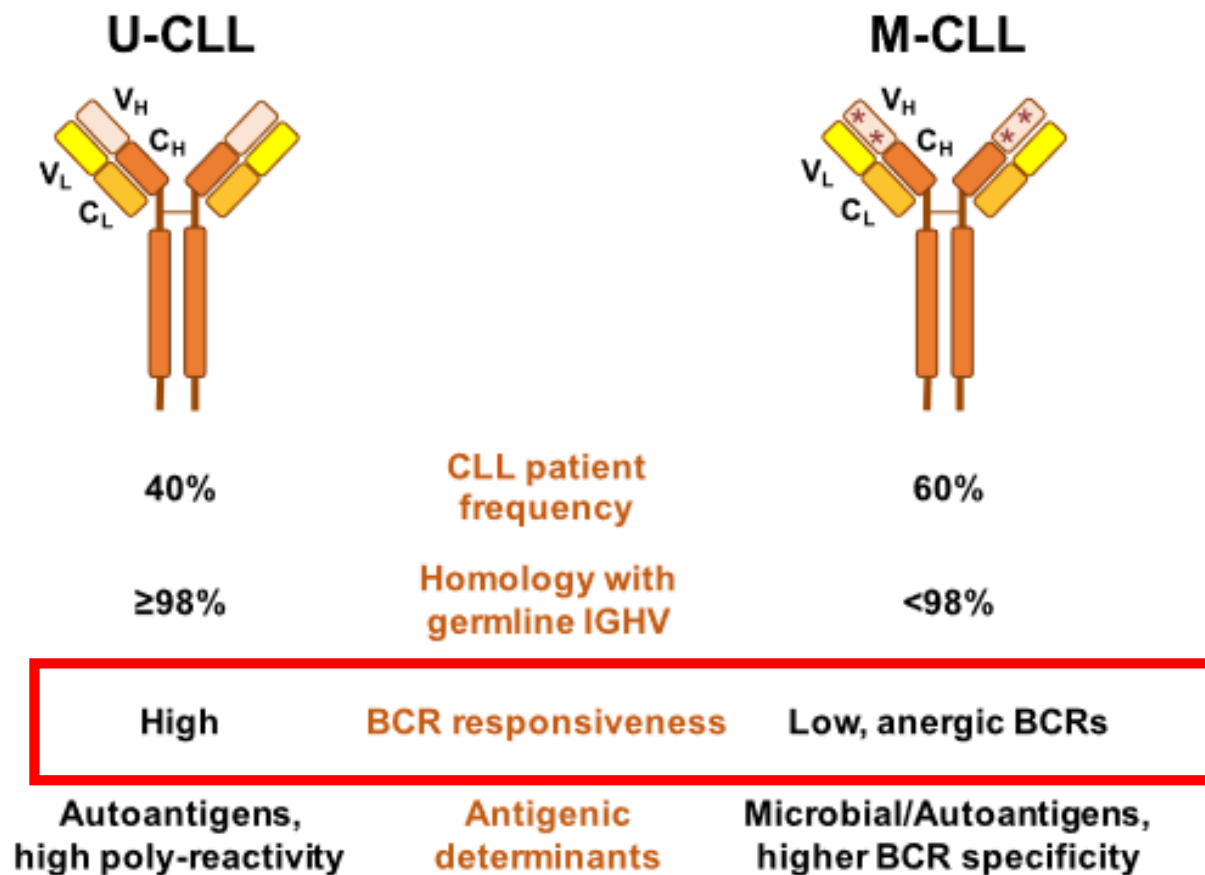


Damle RN et al., *Blood* 94: 1840-1847, 1999

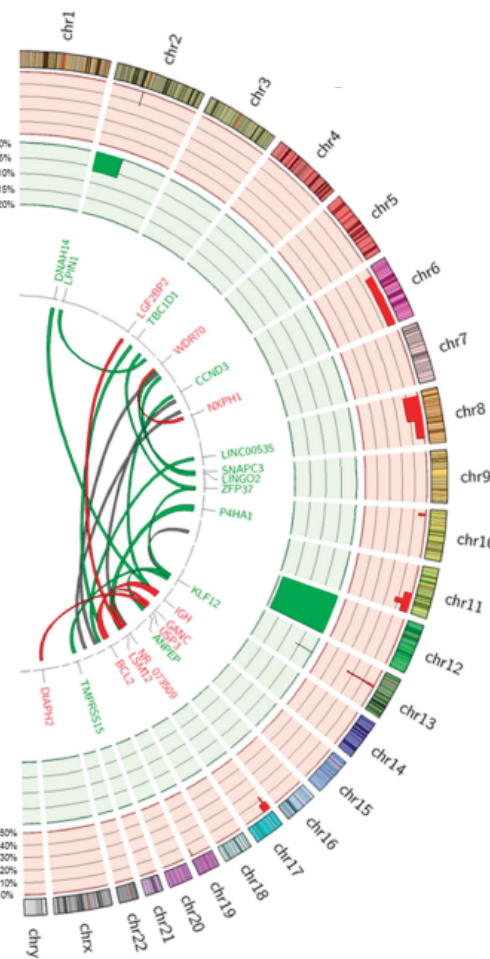
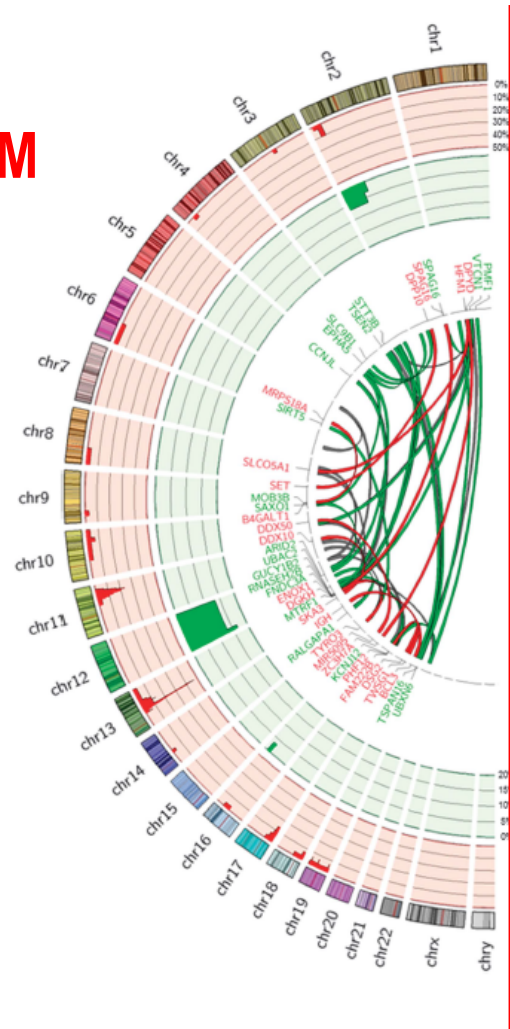


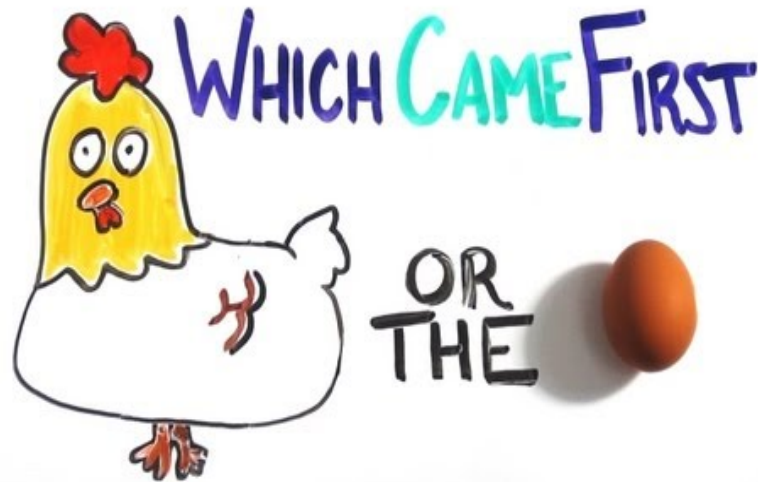
Hamblin TJ et al., *Blood* 94: 1848-1854, 1999

# Characteristics of U-CLL and M-CLL patient subsets

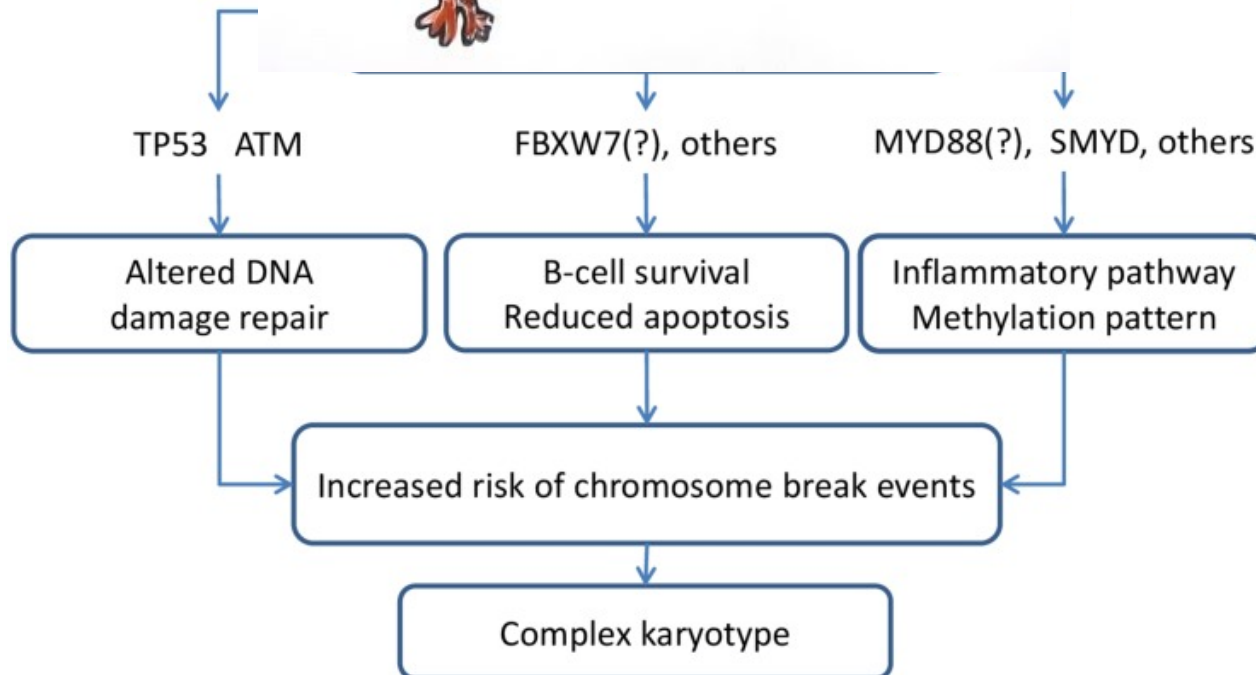


## Distinct mutational landscapes by WGS in IGHV-M and IGHV-UM CLL

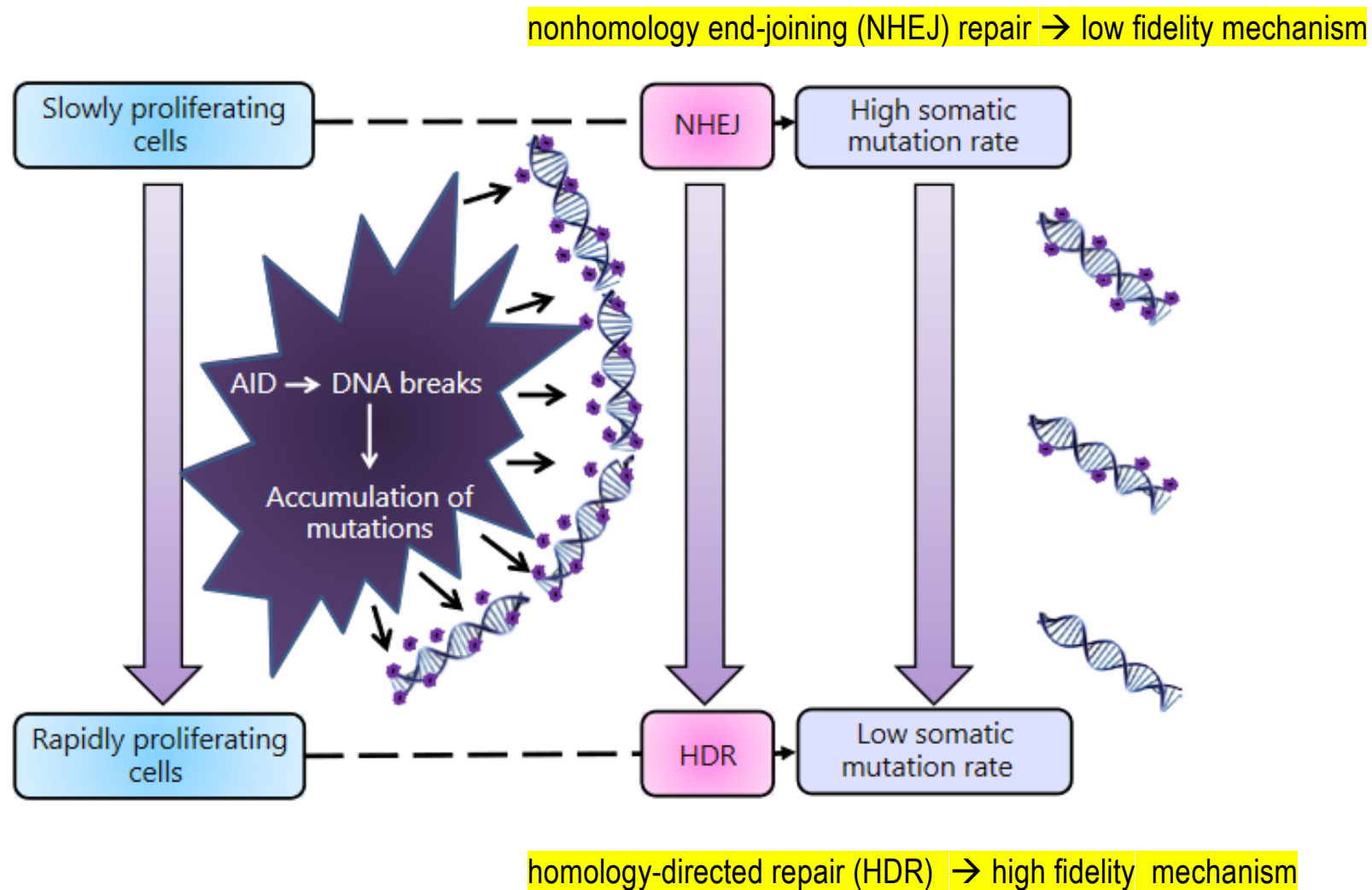




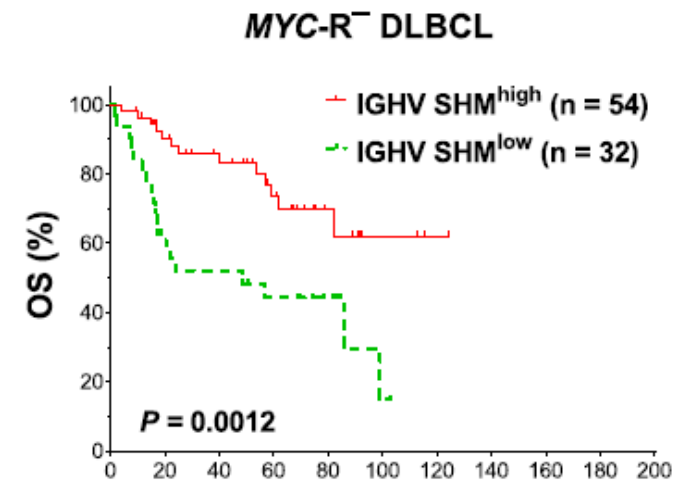
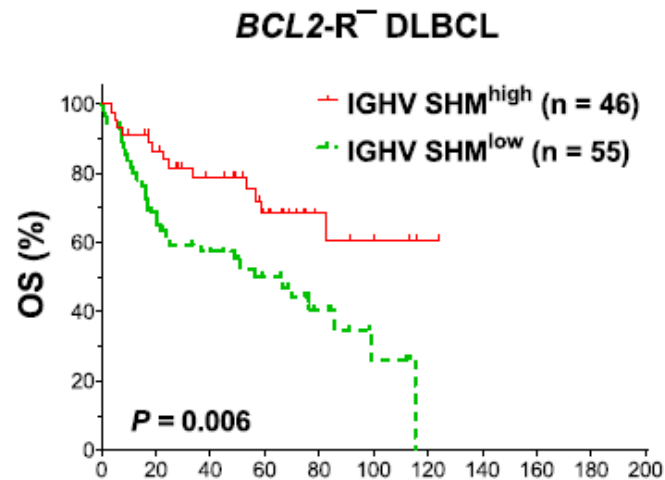
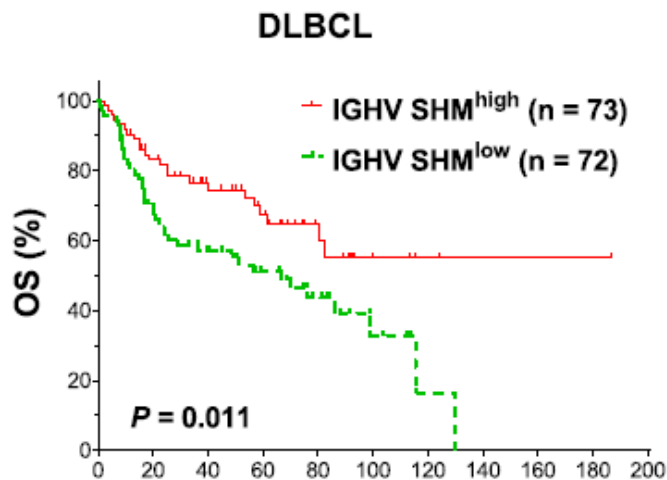
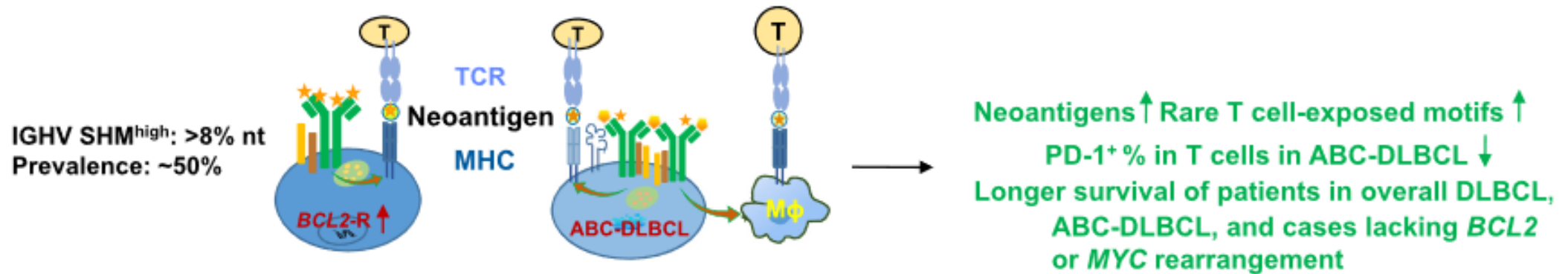
**Proliferation  
or  
IGHV mutational status ?**



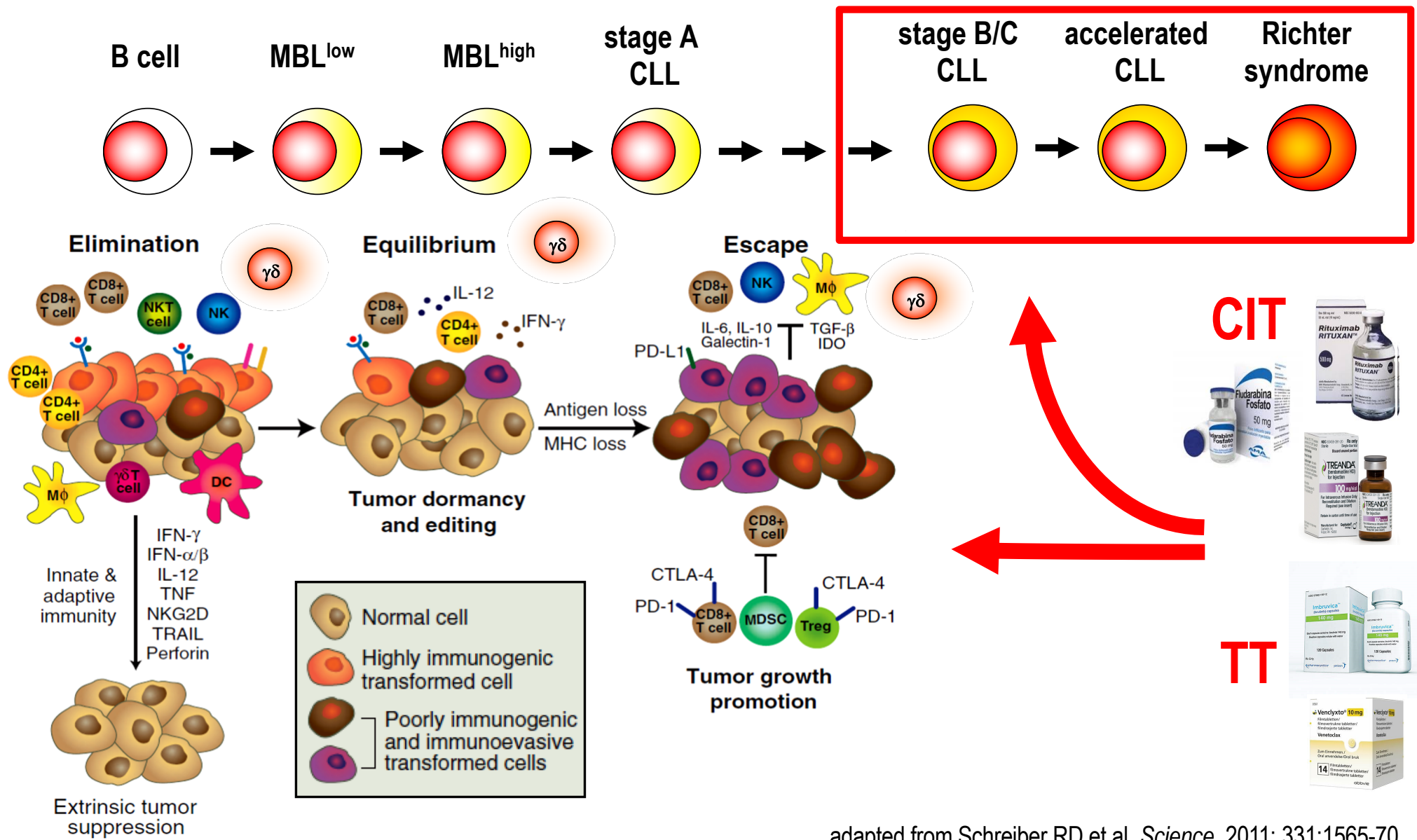
# The IGHV mutations status is determined by the proliferation rate



# Immunogenicity potential of IGHV-derived neoantigens generated by SHM



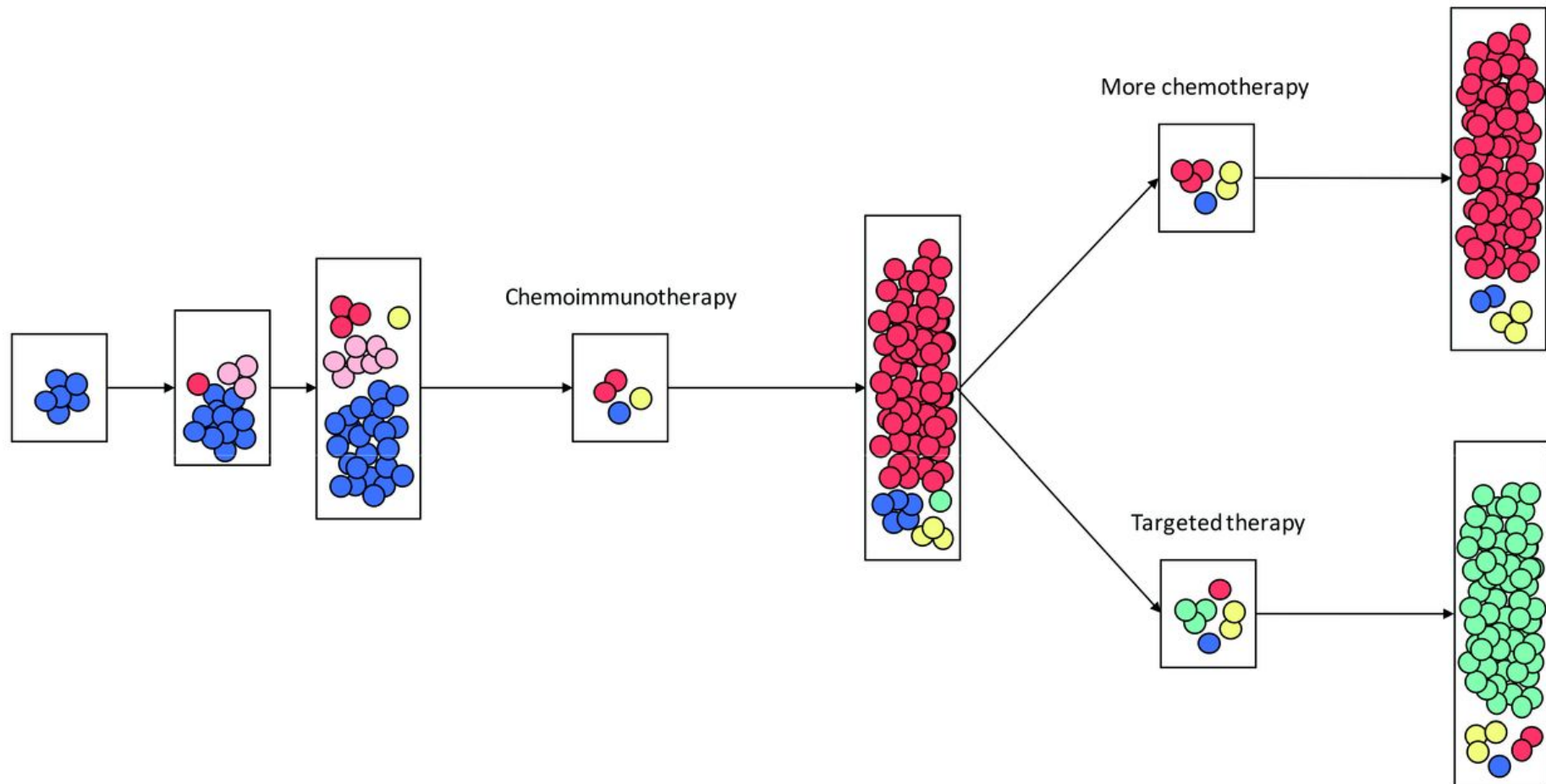
Xu-Monette ZY et al., *J Immunother Cancer*. 2019 Oct 22;7(1):272.



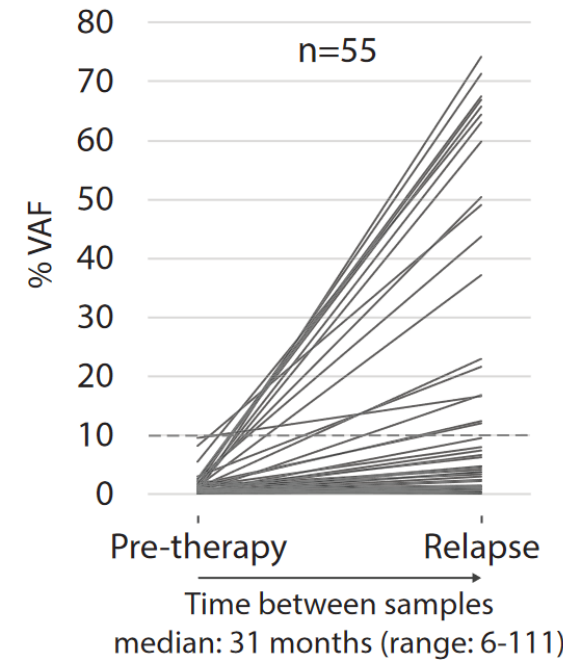
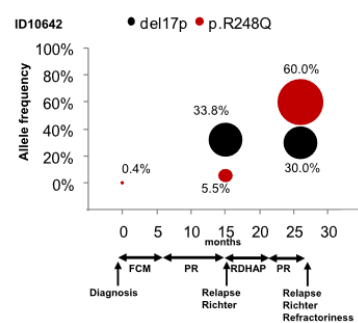
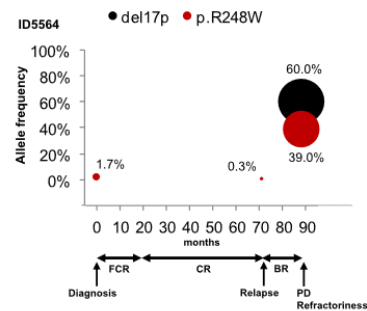
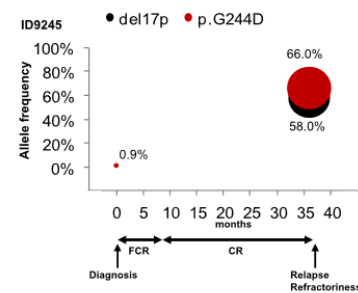
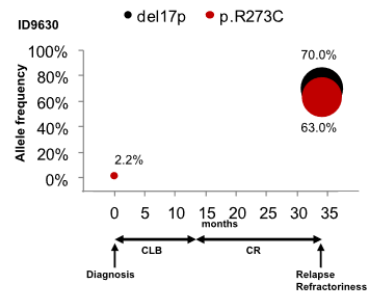
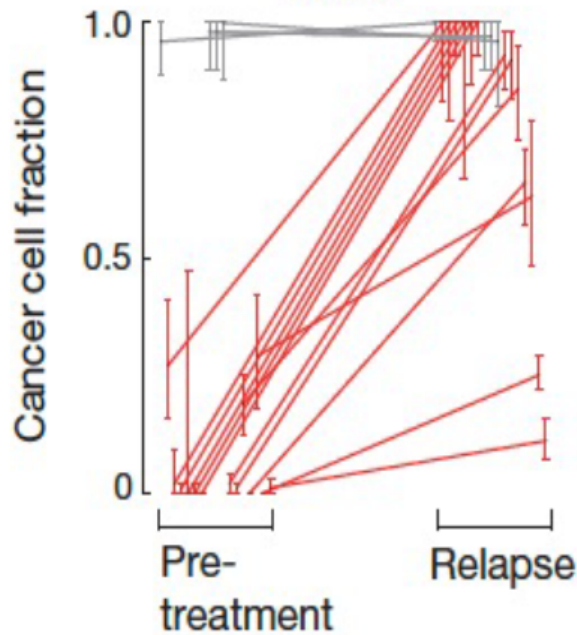
adapted from Schreiber RD et al, *Science*. 2011; 331:1565-70



# Clonal selection is driven by different treatment regimens.



# CIT-induced clonal expansion of *TP53* mutants



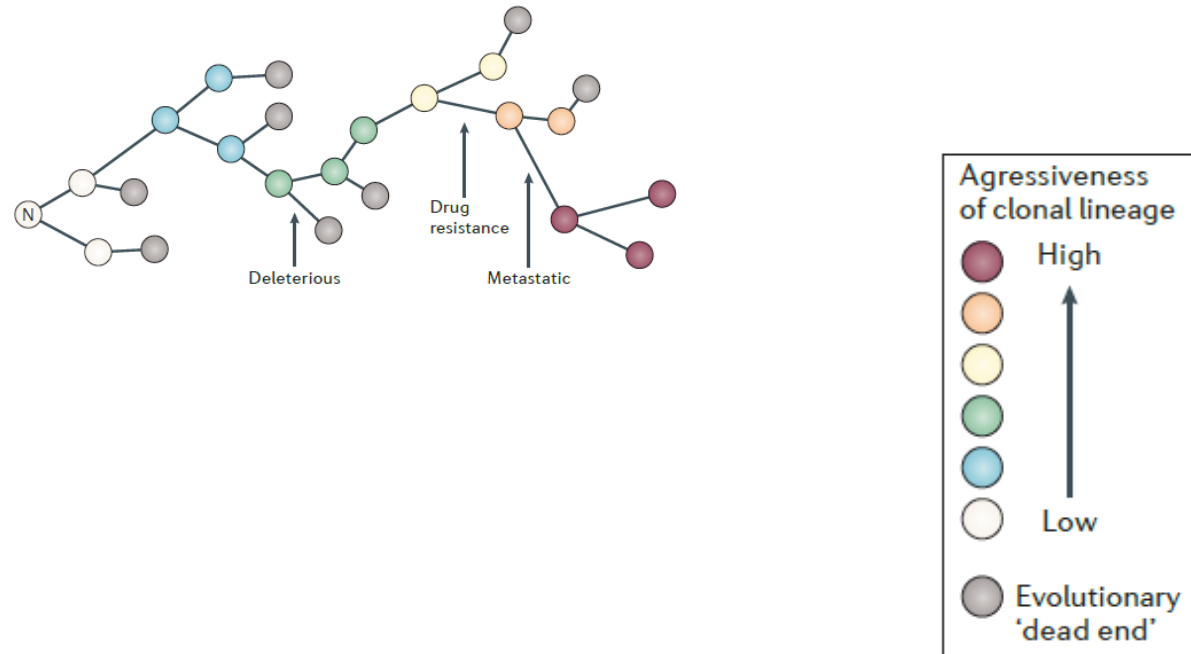
Landau DA et al., *Nature*. 2015 Oct 22;526(7574):525-30.

Rossi D et al., *Blood*. 2014 Apr 3;123(14):2139-47

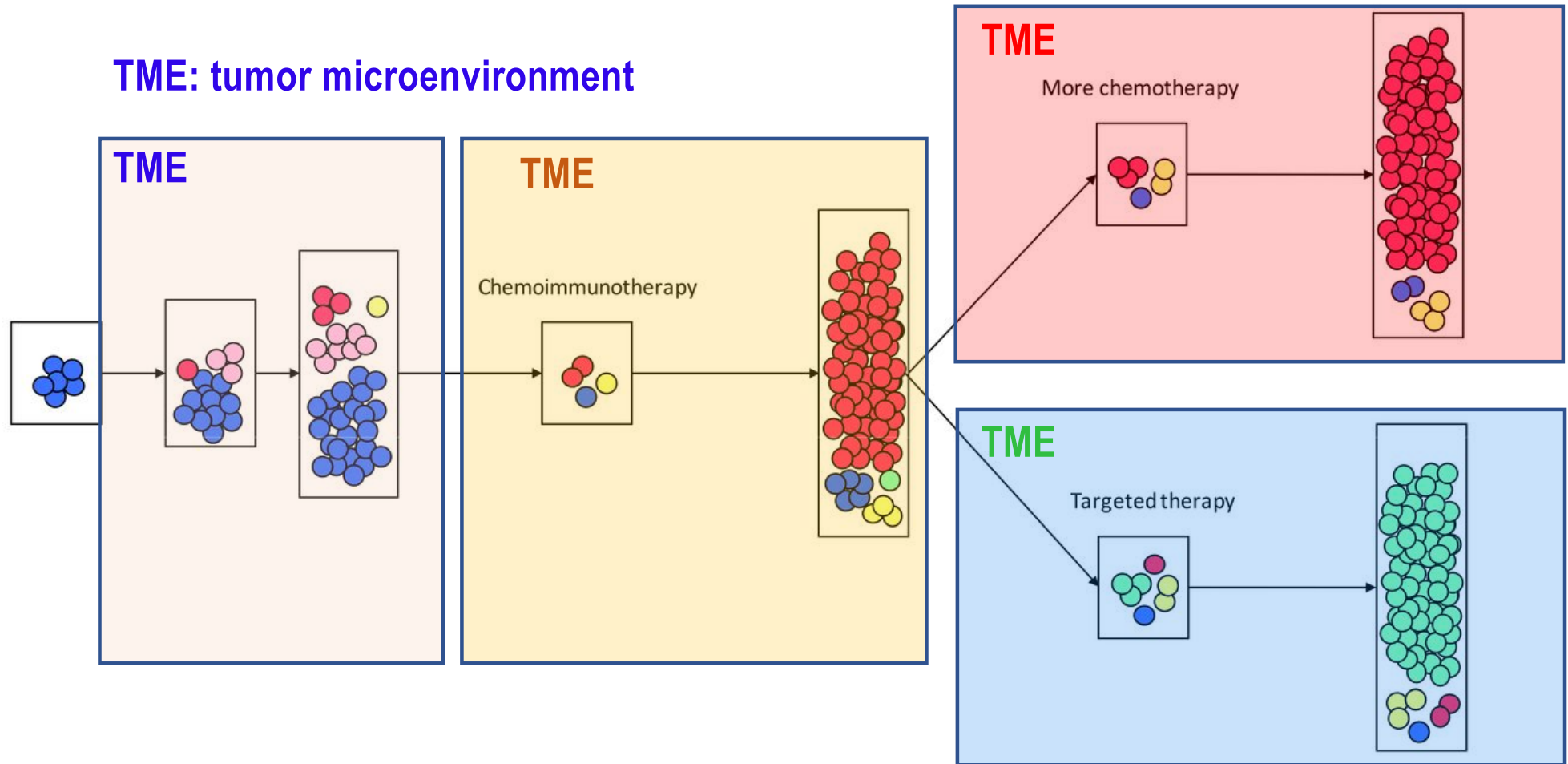
Malcikova J et al., *Blood*. 2021 May 4; pub ahead of print

# Alternative clonal evolution models to survive in the TME

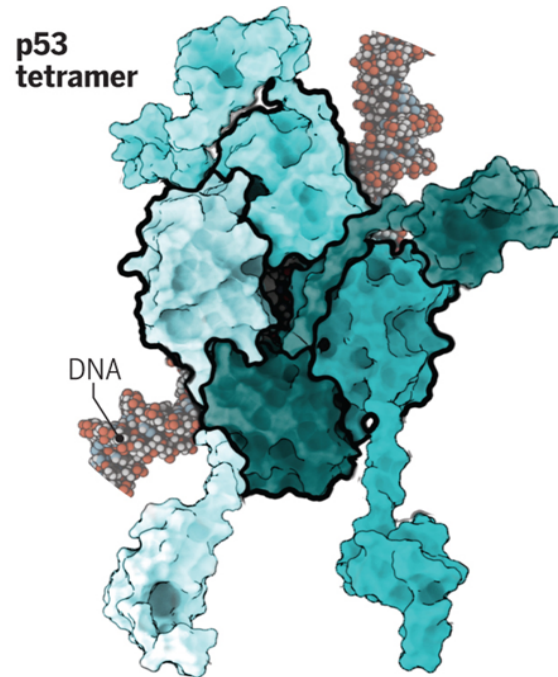
## Clonal evolution model



## TME: tumor microenvironment



# Effects of TP53 mutations



**Wild-type p53**  
Wild-type (WT) p53 is a tetrameric transcription factor.



**DNE mutant/  
wild type**  
Mutation in one allele can cause DNEs in which the mutant (MT) dimer poisons the wild-type dimer.



**p53 LOF mutant**  
An all-mutant tetramer may have LOF of wild-type activity.

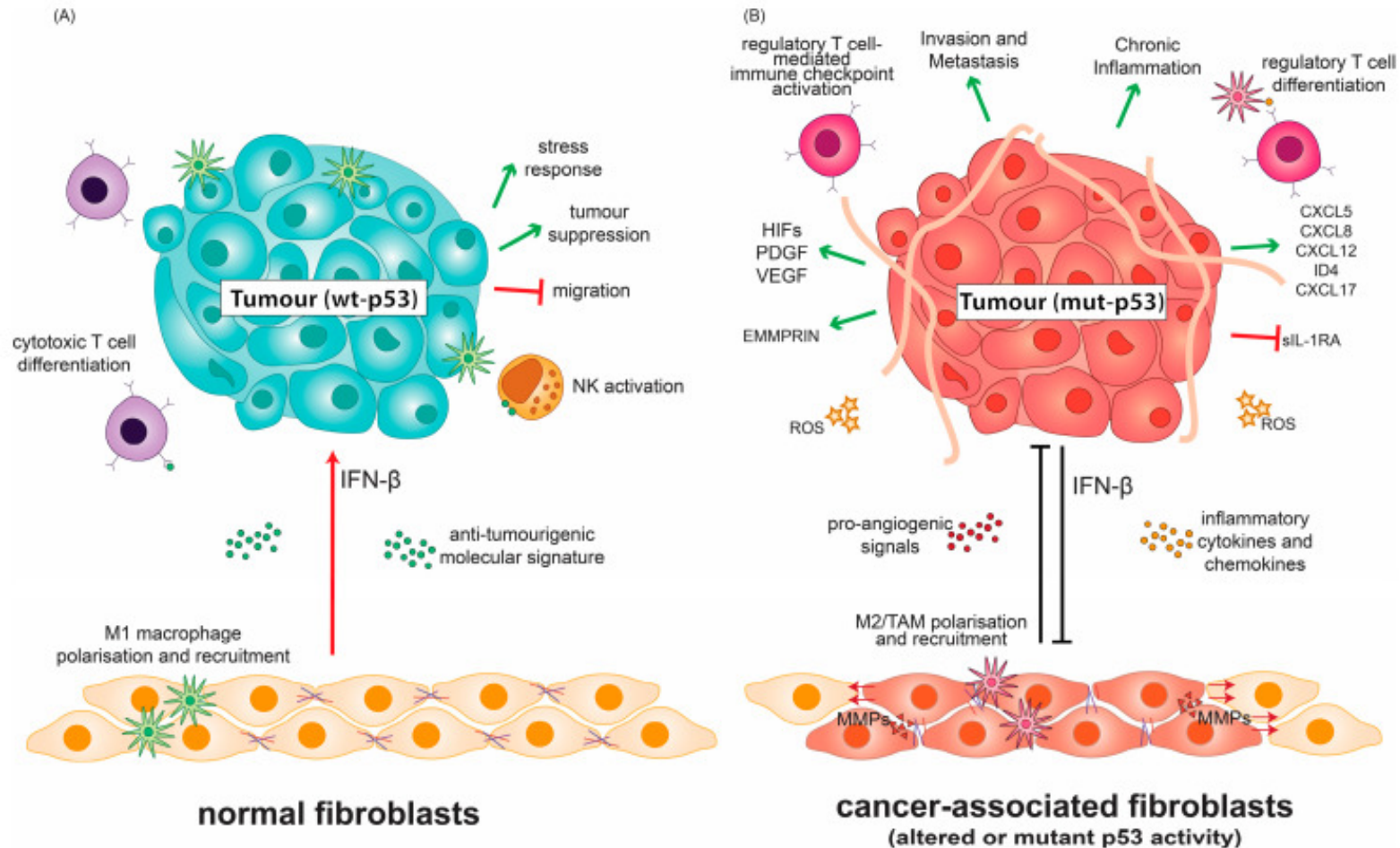


**p53 GOF mutant**  
All-mutant tetramers may show GOF through new protein interactions with protein X.



**DNE: dominant negative effects;  
LOF: loss of function;  
GOF: gain of function;**

# Mutant p53 modifies the TME to suppress immune responses

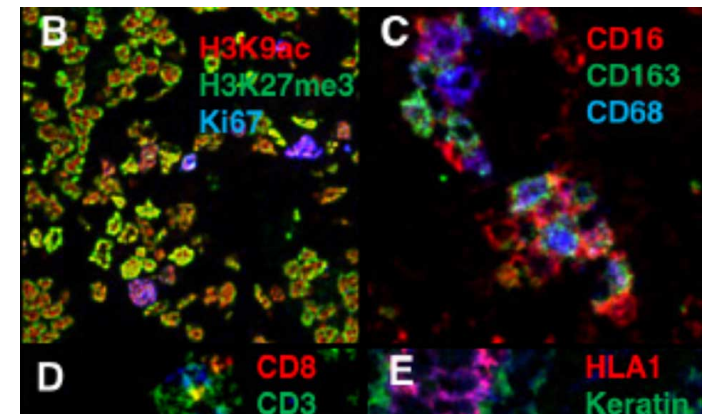
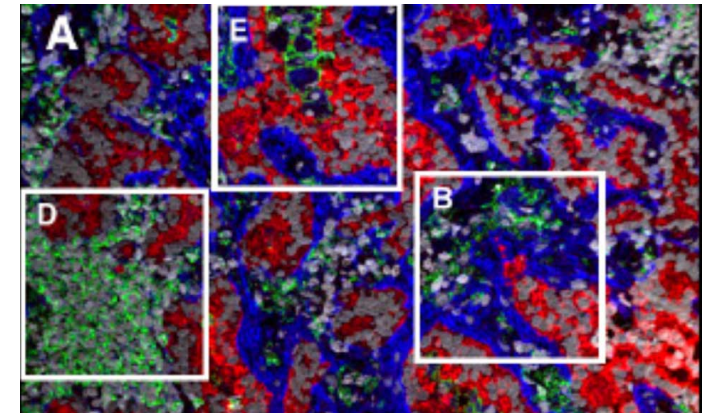




Precision  
Medicine

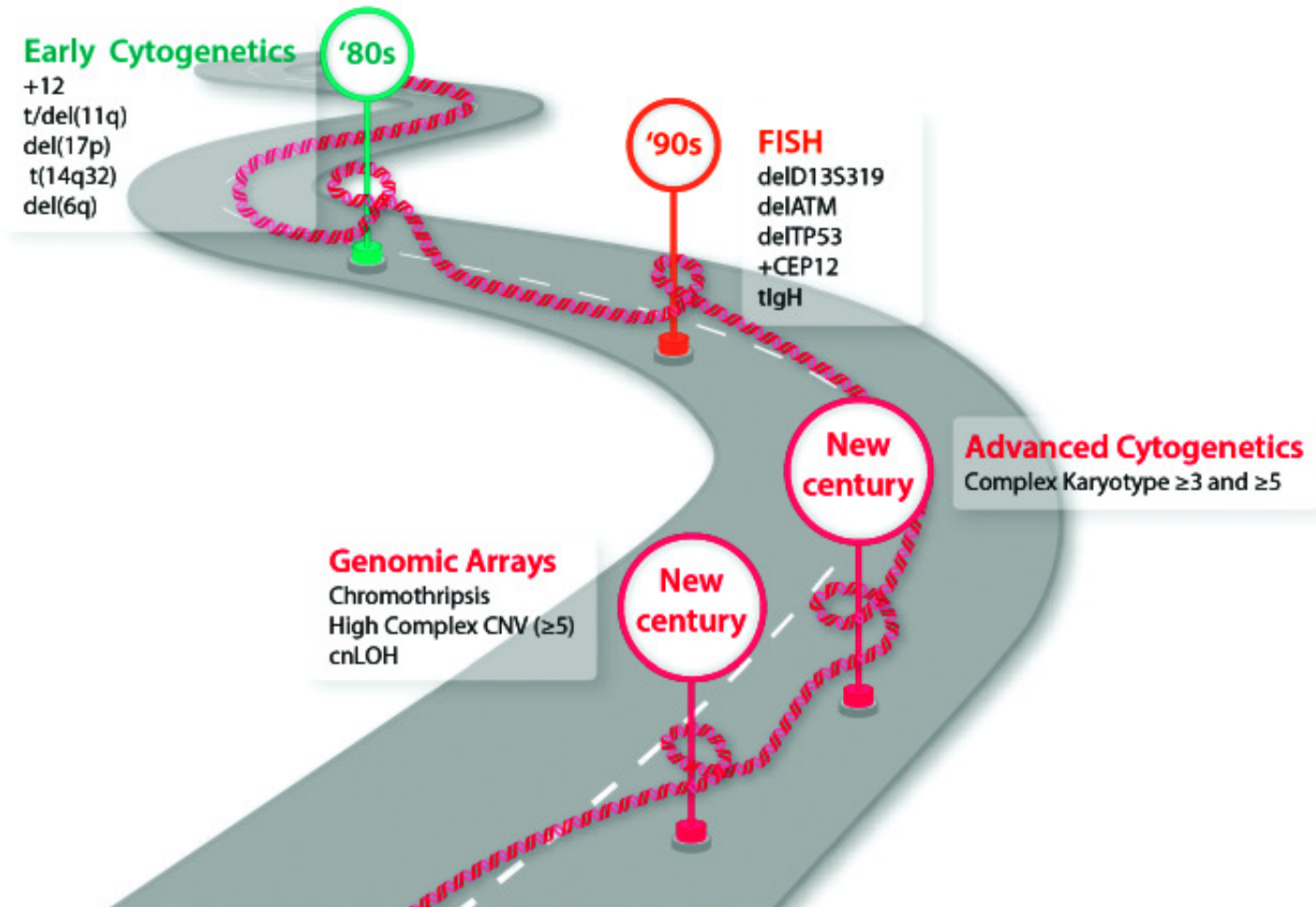


Precision  
Immunology

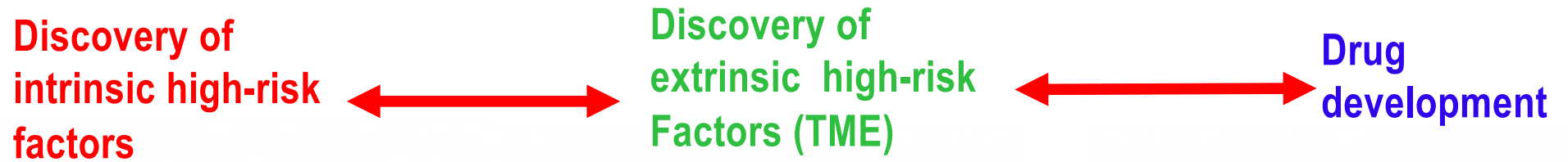




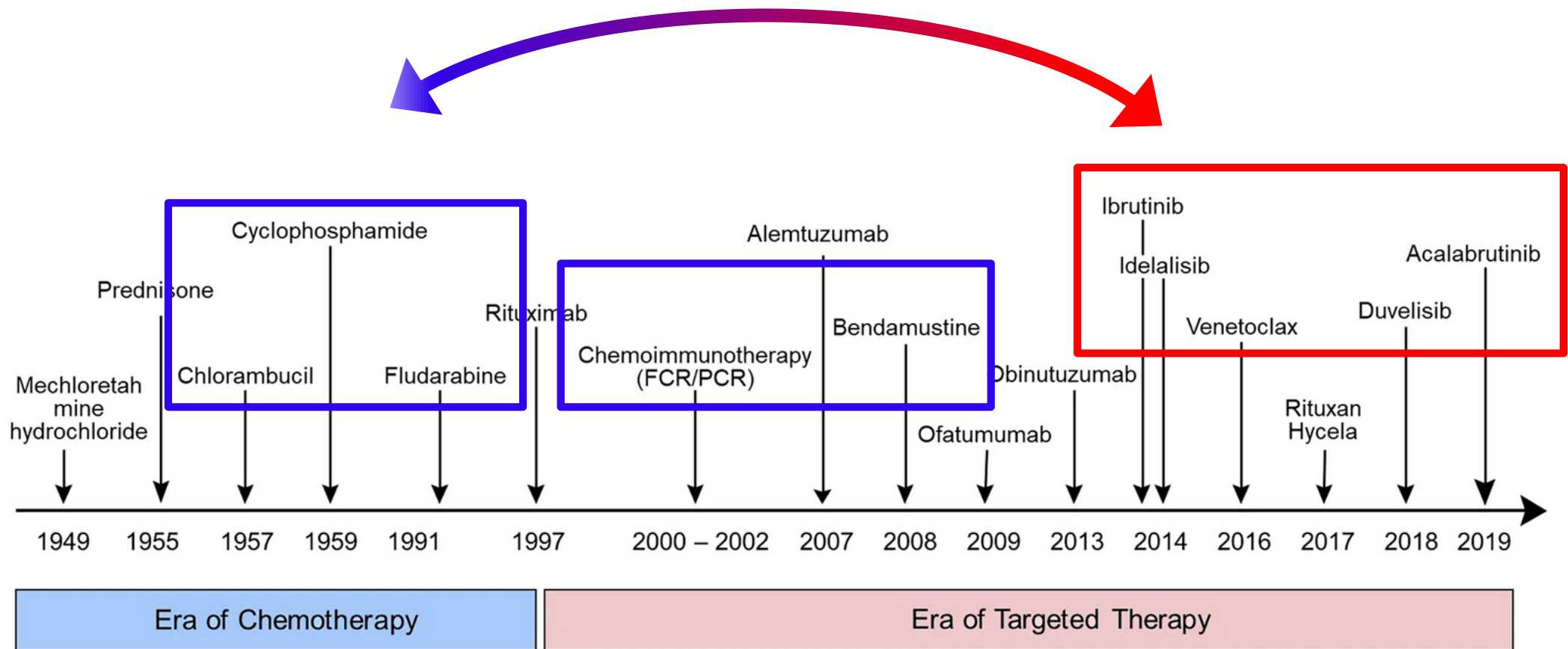
# The long road of genetic advances with a clinical impact on CLL



## The evolutionary concept of high-risk CLL



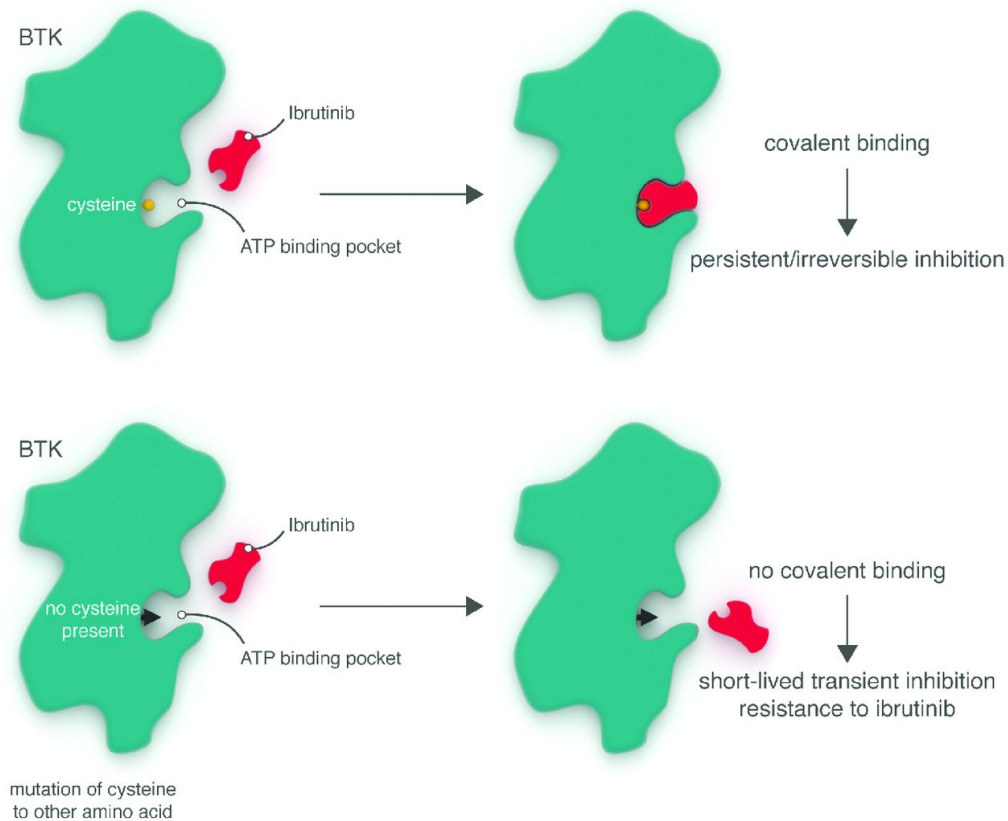
# Paradigm shift of high-risk concept in CLL



adapted from Parikh, S.A. et al., *Leukemia* **34**, 1979–1983 (2020)

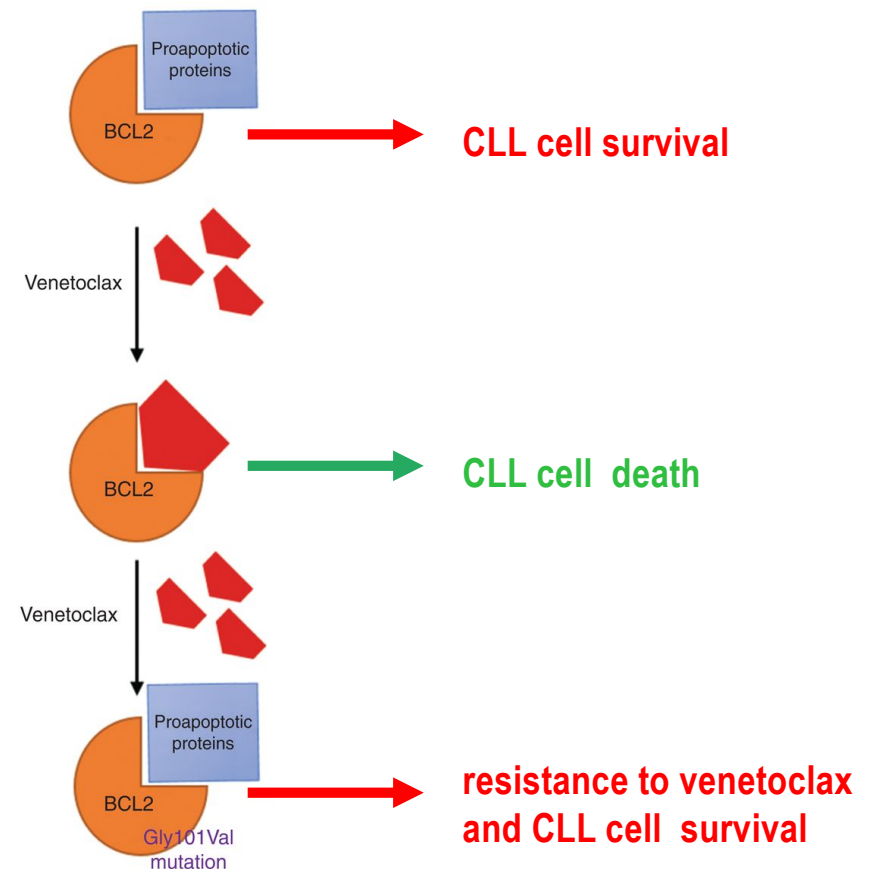
# Targeted therapy is not exempted from acquired resistance

## ibrutinib



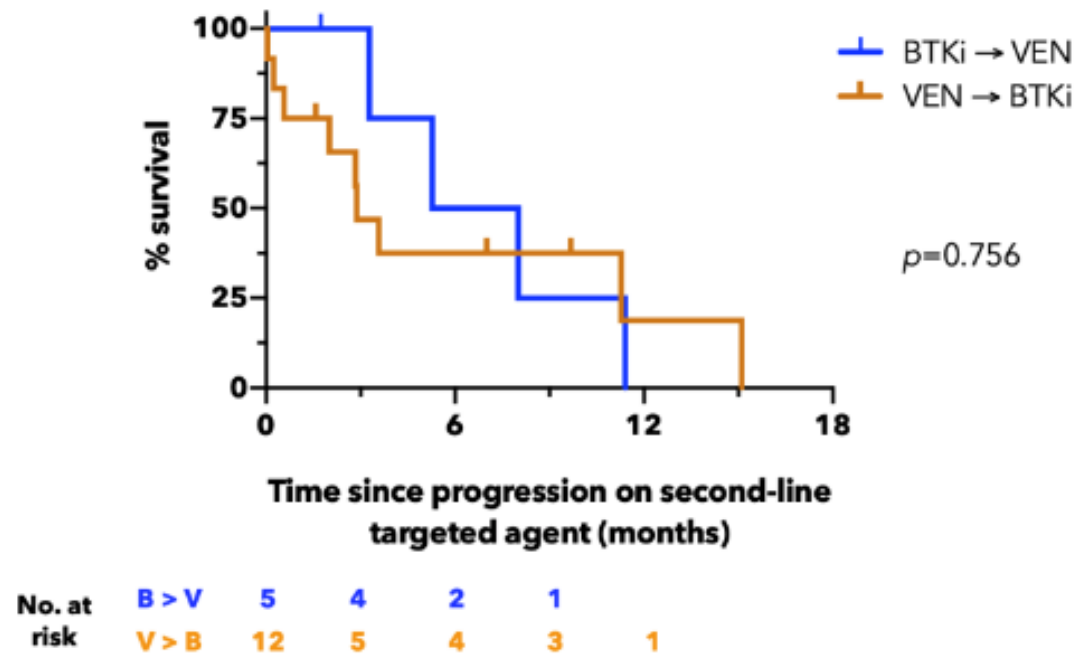
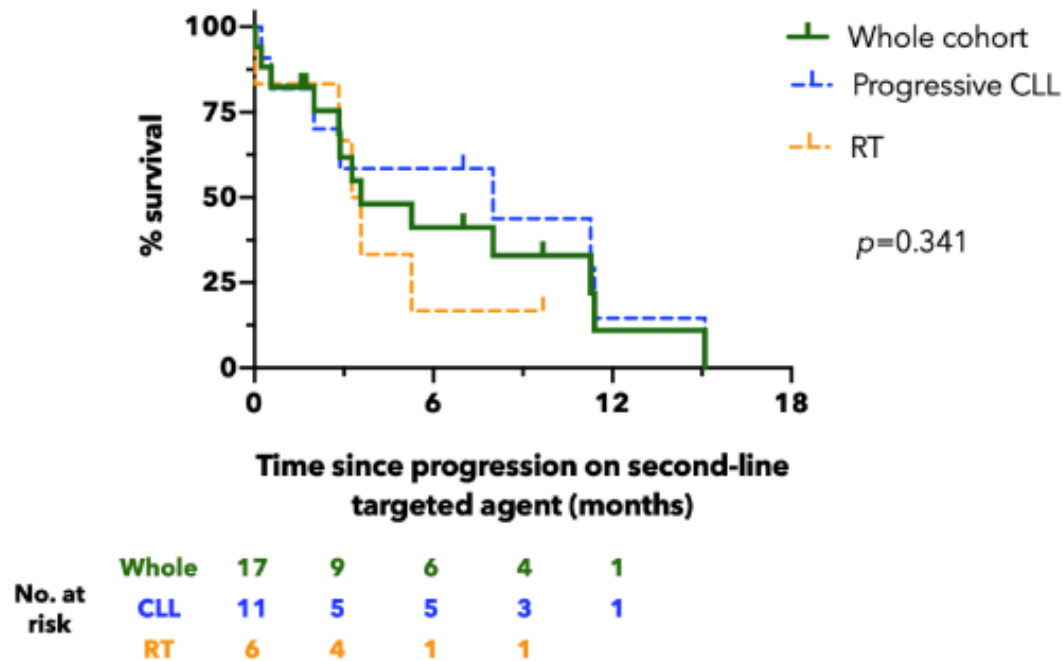
Zhang SQ et al., *Br J Haematol.* 2015 Aug;170(4):445-56.

## venetoclax



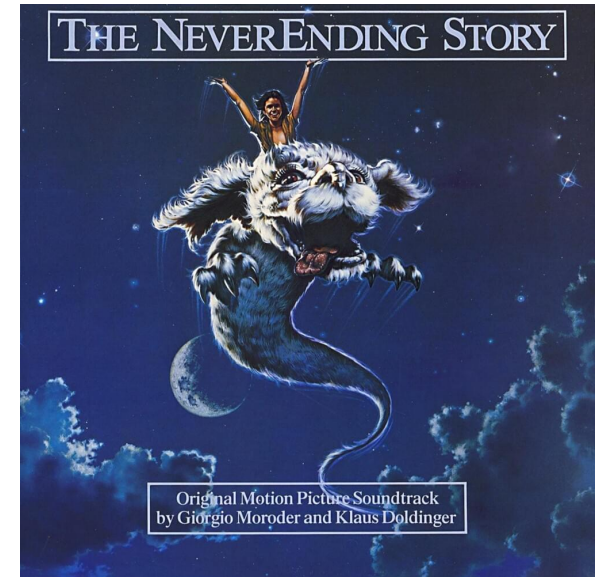
modified from Thangavadi S et al., *Cancer Discov.* 2019 Mar;9(3):320-322.

# Dismal prognosis of CLL pts resistant to BTKi and venetoclax



## Final Thoughts:

- High-risk CLL cannot be cured by CIT
- Therapeutic switch from CIT to TT to (C)IT to IT/TT
- High-risk is an evolving concept
- Decoding mechanisms of disease progression
- Decoding mechanisms of primary and acquired drug resistance



?

2<sup>nd</sup> edition

Unmet challenges in high risk  
hematological malignancies:  
from benchside to clinical practice



3<sup>rd</sup> edition

Towards meeting  
unmet challenges in high-risk CLL



2<sup>nd</sup> Cuneo City ImmunoTherapy Conference (CCITC)

# Immunotherapy in Hematological Malignancies 2021

September 29-30,  
October 1-2, 2021  
VIRTUAL

Organized by Prof. Massimo Massaia, SC Ematologia AO S.Croce e Carle, Cuneo, Italy and  
Centro Interdipartimentale di Ricerca in Biologia Molecolare (CIRBM), Torino, Italy

## Main Topics

- Targeting tumor cells
- Targeting immune cells and bystander cells
- Antibody-drug conjugates
- Bispecific antibodies
- Abscopal effect of radiotherapy
- CAR-T cells, NK, and  $\gamma\delta$  T cells
- Targeting the microbioma
- Cancer immunometabolism
- 3D cell culture models
- Mechanisms of immune resistance
- Immune lessons from allogeneic transplantation
- Tumor vaccination: back to the future

ER Congressi <https://ccitc2021.it>