



XIX CONGRESSO  
NAZIONALE  
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# TP53 nelle neoplasie linfoproliferative

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Università  
degli Studi  
di Ferrara



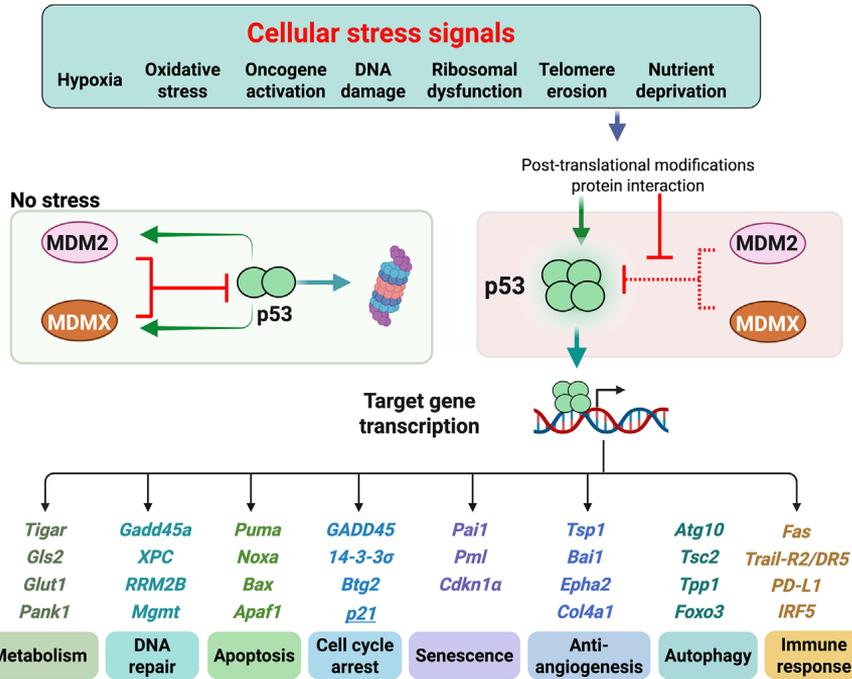
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Palazzo degli Affari

Dipartimento di scienze mediche

## Disclosures of Gian Matteo Rigolin

Company name	Research support	Employee	Consultant	Stockholder	Speakers bureau	Advisory board	Other
AstraZenca							x
Janssen-Cilag							x
BeOne							x
Pfizer							x
Roche							
Eli Lilly							
BMS							
AbbVie							
Sanofi							

# p53 signaling pathways



Signal  
 ↓  
 Mediators  
 ↓  
 Core regulation  
 ↓  
 Effectors  
 ↓  
 Response

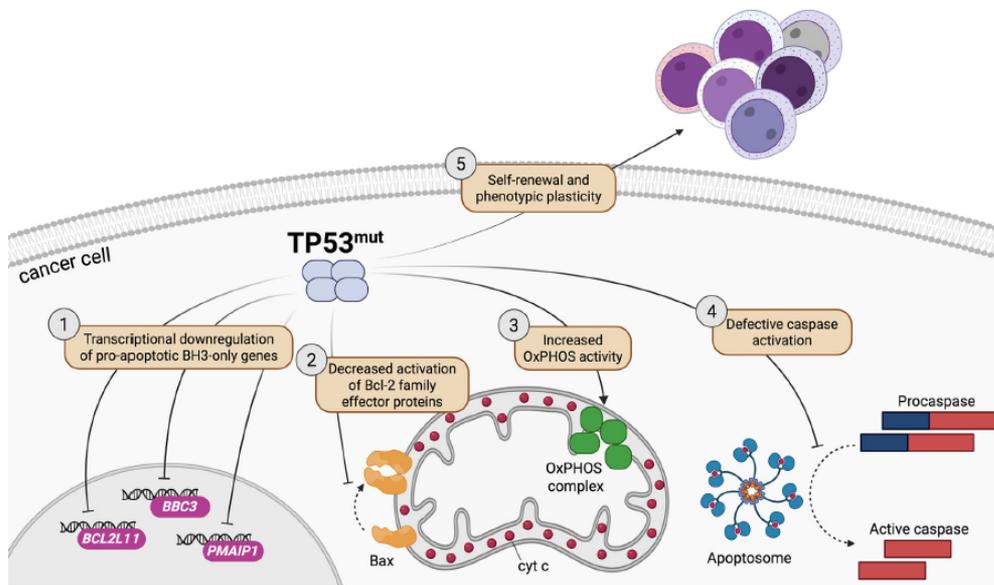
- Tumor-suppressor TP53 is crucial to**
- maintaining biological homeostasis
  - mediating in damaged cells
    - cell-cycle arrest
    - senescence
    - apoptosis

- TP53 disruptions impact on cancer trough:**
1. - mutant p53 **loss of function**
  2. - mutant p53 **dominant-negative**
  3. - mutant p53 **gain of function**

Baliakas P, Soussi T. J Intern Med. 2025 Aug;298(2):78-96.

# TP53 in lymphoproliferative disorders

## Effects of TP53 in cancer



Tamellini E, et al. Cell Death Dis. 2025 Nov 10;16(1):819.

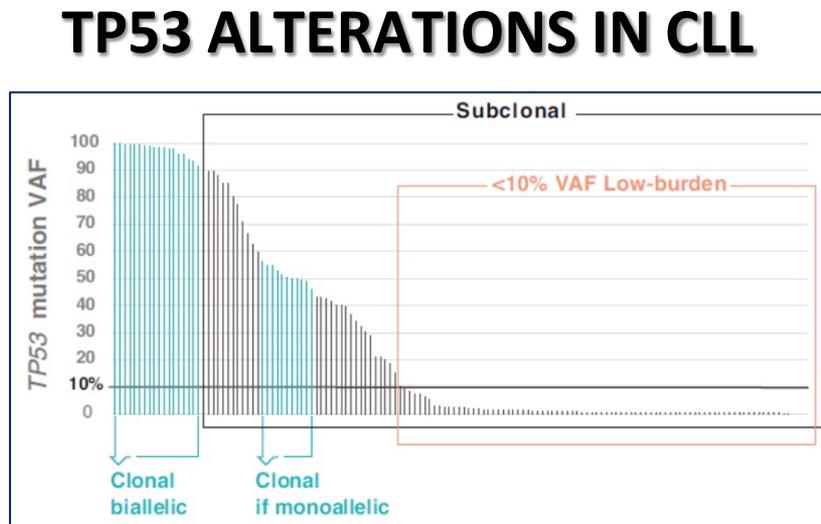
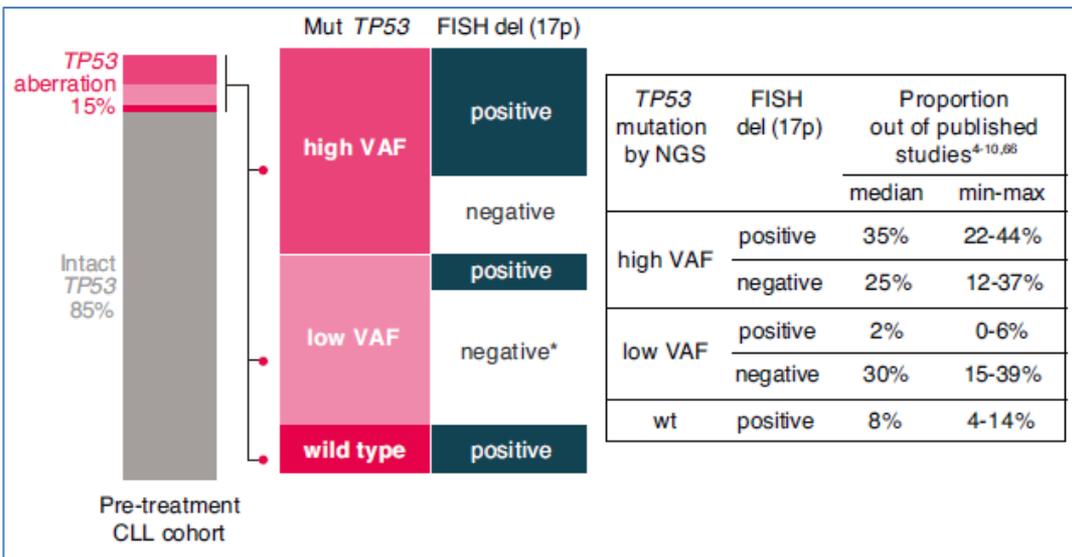
TP53 is working through a **2-hit mechanism**.

The **1<sup>st</sup> hit** is usually Single Nucleotide Variants  
 The **2<sup>nd</sup> hit** could be:

- Del 17p,
- Additional muts in the 2<sup>nd</sup> TP53 allele
- Copy neutral loss of heterozygosity (cnLOH)

## TP53 alterations in

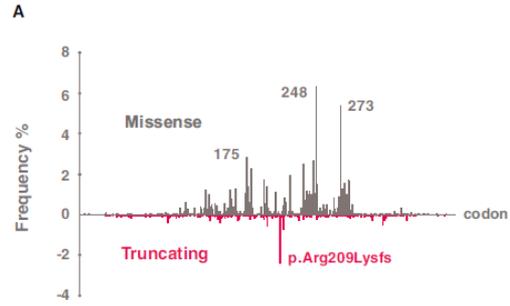
- Chronic Lymphocytic Leukemia
- Mantle Cell Lymphoma
- Diffuse Large B Cell Lymphoma
- Waldenstrom Macroglobulinemia



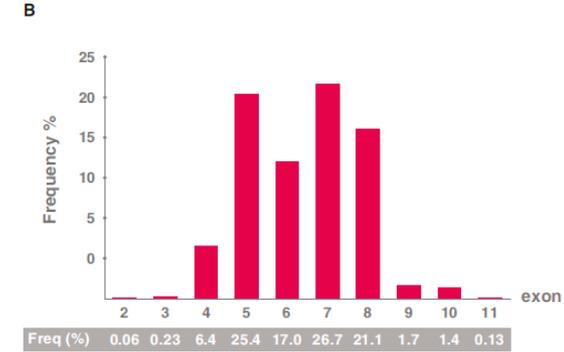
The frequency of TP53 abns is

- higher in those with **UM-IGHV genes**.
- **low at diagnosis (5-10%)**, higher at frontline treatment (10–20%), and in later disease stages, predominantly in CIT-treated pts and Richter transformation (**up to 50%**).

# Spectrum of TP53 defects in CLL



Variants are present along the gene and cluster in hot-spots  
 truncating frameshift variant in codon 209 is CLL-specific



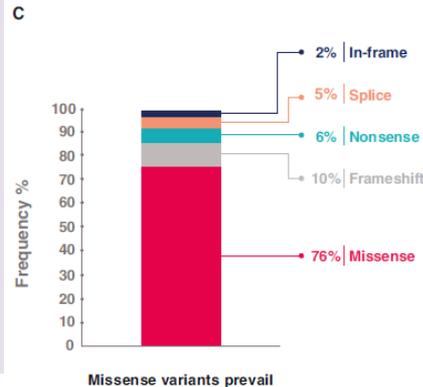
Majority of variants in exons 5-8

**A. Variants distribution**

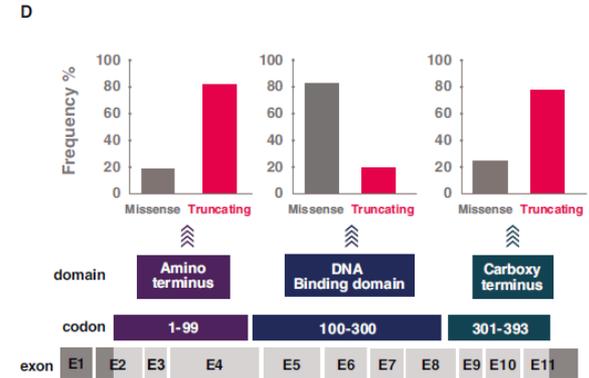
**B. Exon distribution of variants.**

**C. Proportion of variant types**

**D. Proportion of variant types**



Missense variants prevail



Truncating variants prevail in N- and C- terminus



# Classification of TP53 variants detected in CLL

A **classification algorithm** showing the basic principles of assigning variants into pathogenicity/oncogenicity classes.

DNA event	CODING REGION						INTRONS	
	Deletion/insertion/duplication		Single nucleotide variant				Borderline exonic (2bp)	+/-2 bp Canonical splice site
Variant type	Frameshift	In-frame	Nonsense (stop-gain)	Missense	Synonymous	Possibly splice <sup>a</sup>	Splice	Possibly splice
Occurrence	10%	2%	6%	76%	<<1%	<1%	5%	<<1%
Consequence	Truncating	Deletion and/or insertion of aminoacids	Truncating	Amino acid change	No change	Various	Mainly truncating	Various
Classification <sup>a</sup>	Pathogenic		Pathogenic	Benign			Pathogenic	

In DNA-binding domain			Known population variant		Effect on splicing		Effect on splicing	
Yes	No		Yes	No	Confirmed	Predicted	Confirmed	Unknown
	Described to affect function							
	Yes	No/not sure						
	Likely pathogenic	Likely pathogenic	VUS	Benign	Pathogenic	VUS	Likely pathogenic	VUS

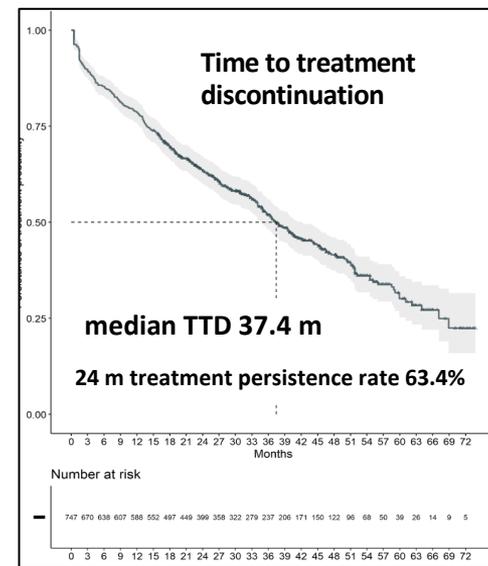
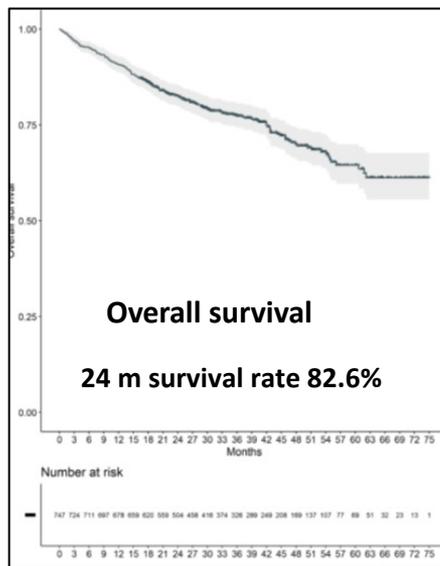
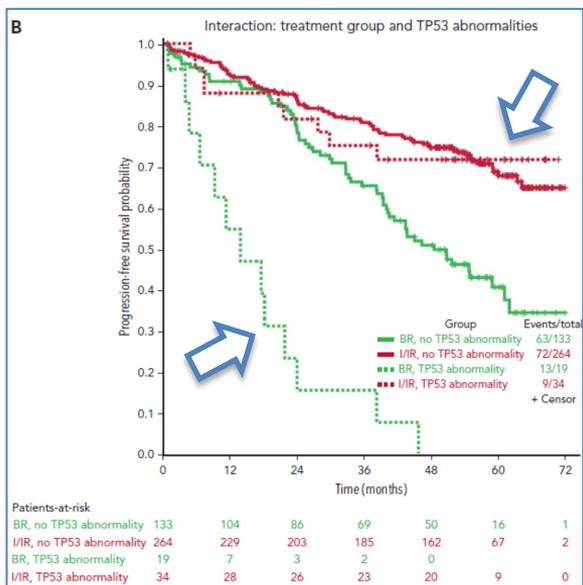
The TP53 database & Seshat:  
 Transactivation class (Kato et al., 2003, PMID: 12826609)  
 DNE\_LOF class (Giacomelli et al., 2018, PMID: 30224644)  
 Growth suppression (Kotler et al., 2018, PMID: 29979965)

Non-functional	Concordant		Discordant Partially functional No data
	Functional		
	Confirm		
	Confirmed	Not confirmed	Require further consideration and gathering information from multiple resources
Pathogenic	(Likely) Benign	False positive	

- Extent of functional loss
  - Frequency in tumors
  - Presence in healthy human populations
- »»»» ClinGen TP53 Variant Curation Expert Panel

# Prognostic and Predictive value of TP53 alterations in CLL

- TP53abs **predictive value** is clear for CIT regimens
- **Targeted agents outperformed CIT in the frontline and R/R settings.**



The A041202 study of ibr regimens for older untreated CLL  
Woyach JA et al. *Blood*. 2024;143(16):1616-1627.

CLL with TP53 abns and 1<sup>st</sup> line ibrutinib  
Rigolin GM, et al. *Blood Cancer J*. 2023;13(1):99.

# Prognostic and Predictive value of TP53 alterations in CLL

The role of TP53 abns in choosing between targeted agent regimens **is less well studied** (decision on risk of toxicities)

The prognostic impact of TP53 appears to be stronger with **time-limited regimens** than with continuous therapy (no definitive conclusions)

Stage	del(17p) or TP53mut	IGHV	Therapy
Inactive disease, Binet A-B, Rai 0–II	Irrelevant	Irrelevant	None
Active disease or Binet C or Rai III–IV	Yes	Irrelevant	Acalabrutinib <sup>1</sup> , zanubrutinib, ibrutinib, or venetoclax + ibrutinib, or venetoclax + obinutuzumab idelalisib-rituximab (only if contraindications for other options)
	No	mutated	Venetoclax + obinutuzumab, <sup>2</sup> or venetoclax + ibrutinib, <sup>2</sup> or acalabrutinib <sup>1</sup> , zanubrutinib, ibrutinib Chemoimmunotherapy <sup>3</sup> with FCR (BR above 65 years) or chlorambucil-obinutuzumab (unfit)
		unmutated	Acalabrutinib <sup>1</sup> , zanubrutinib, ibrutinib, or venetoclax + ibrutinib, <sup>2</sup> or venetoclax + obinutuzumab <sup>2</sup>

(1) The addition of obinutuzumab to acalabrutinib may be considered.

(2) Consider and discuss with the patient: Continuous vs. fixed-duration therapy, specific side effects of drug classes (myelosuppression, infections, secondary malignancies for CIT; cardiovascular toxicity and bleeding for BTKi (Acalabrutinib < Ibrutinib); TLS and infections for Ven-Obi; autoimmune disease and opportunistic infections for Idelalisib).

(3) Chemoimmunotherapy in regions where other options are not available or approved.

Hallek M. Am J Hematol. 2025 Mar;100(3):450-480.

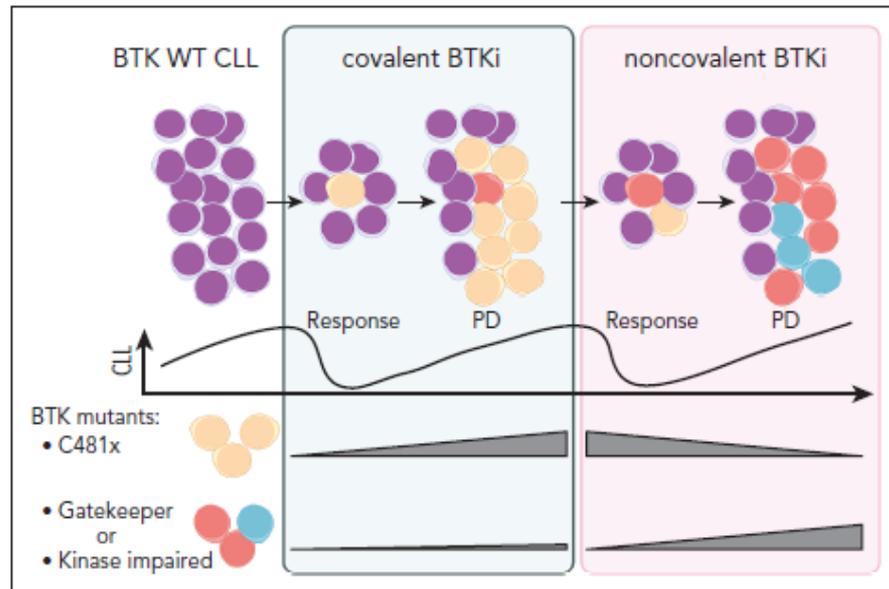
## Relevance of TP53 abns with novel agents

### Targeted agents

- **act independently of the p53 pathway**
- **no preferential pattern of evolution of TP53-aberrant clones**

### To be clarified

- how the TP53-aberrant clone **evolve**
- **Mechanisms of resistance** (1/3 of pts have no mutations at PD).
- the role of **genomic instability/complexity**



Brown JR, et al. *Blood*. 2026;147(1):24-34.

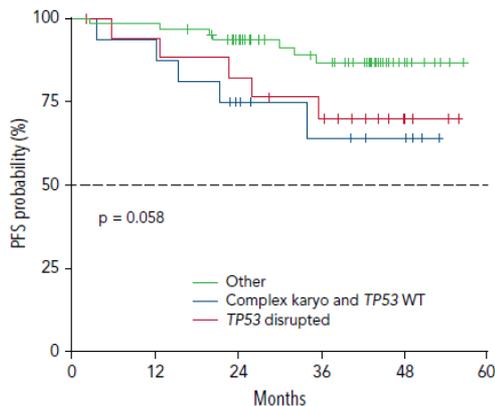
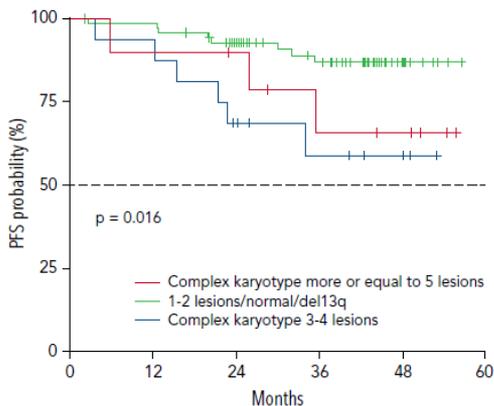
Jebaraj BMC, Stilgenbauer S. *Blood*. 2026 Jan 1;147(1):3-4.

Bonfiglio S, et al. *Blood Adv*. 2023;7(12):2794-2806



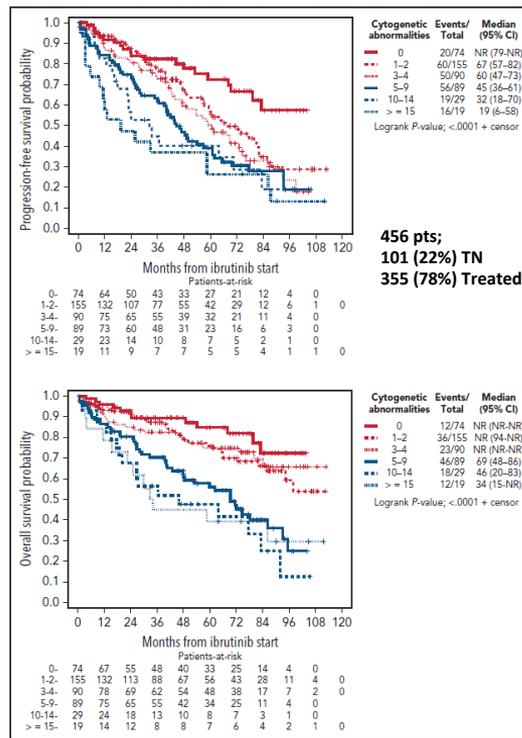
# Karyotype complexity and outcome in CLL treated with ibrutinib

## 1<sup>st</sup> line ibr-R: the GIMEMA LLC1114 phase 2 study



In MVA, the CK was significantly associated with a shorter PFS ( $P = .009$ ) along with a worse ECOG PS ( $P = .048$ ).

Rigolin GM, et al. Blood. 2021;138(25):2727-2730.



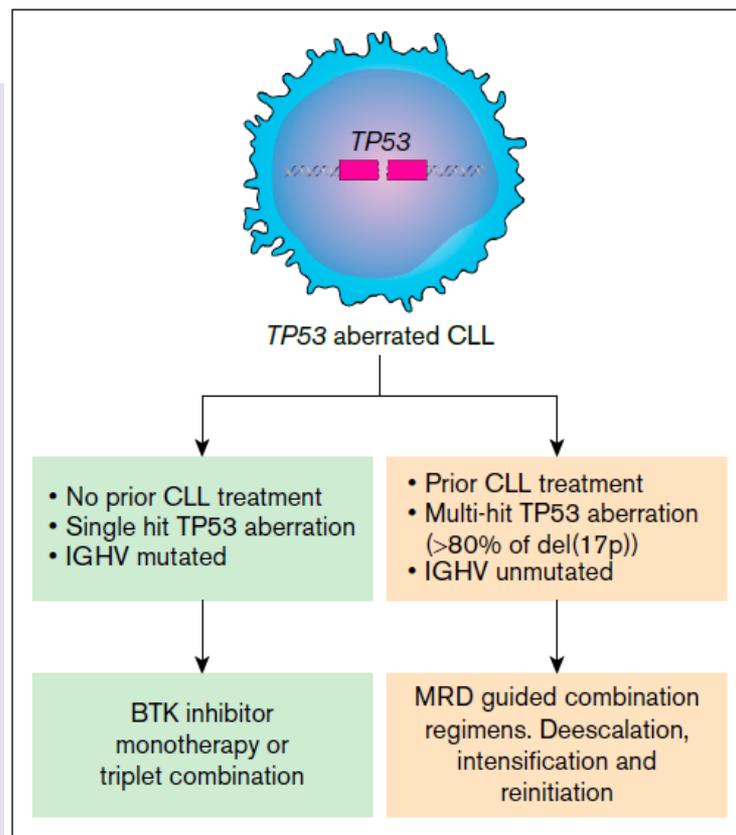
In MVA, karyotypic complexity independently associated with a worse PFS and OS

Kittai AS, et al. Blood 2021;138:2372-2382.

# Open questions in CLL

Given that pts with TP53 disruptions still perform worse

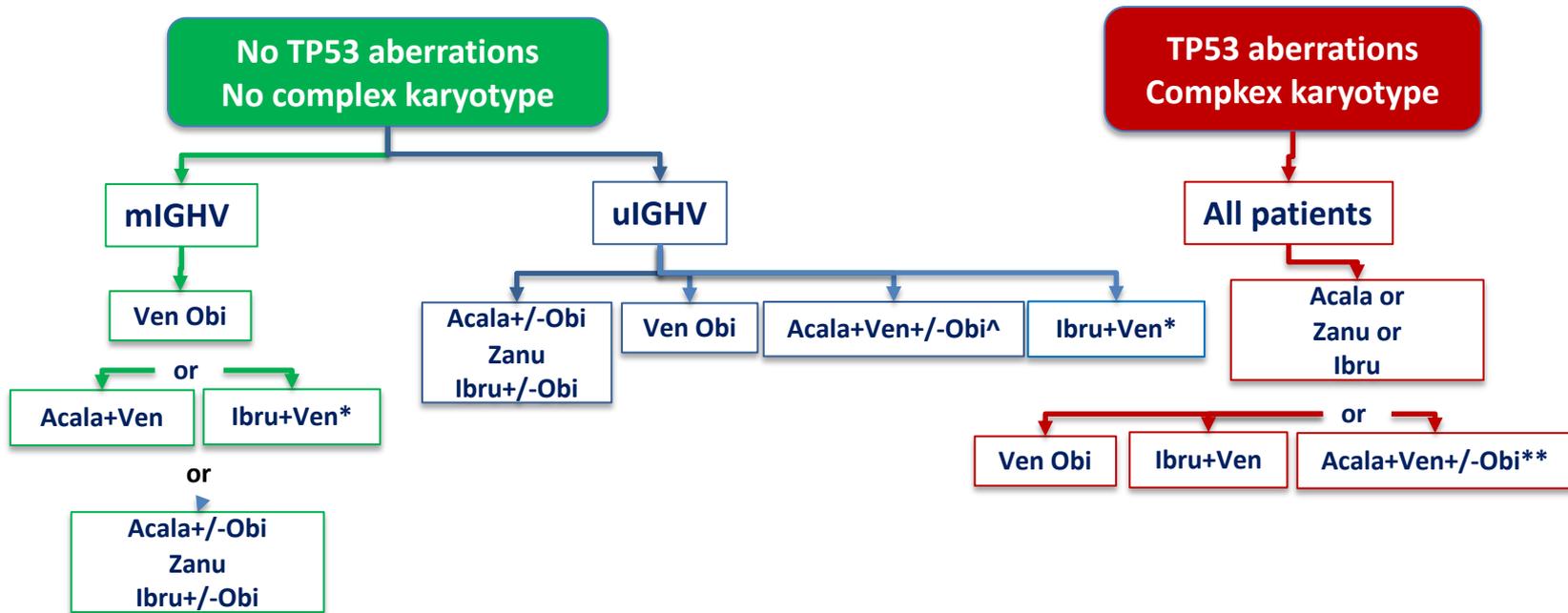
1. Which types and combinations of TP53 abns are predictive for tailoring treatment.
  - Mono, bi-allelic, multiple clones, micro clones, M/UM IGHV
2. The role of genomic complexity
  - karyotype, CGH arrays, chromothripsis, others
3. Is BTK inhibitor monotherapy sufficient?
4. What treatment alternative?



Niemann CU. *Blood Adv.* 2026;10(3):707-709.

# First line therapy of CLL in 2025

Symptomatic early stage or advanced-stage CLL



\*When using I+V, cardiac toxicity must be considered, especially in elderly patients. ^When using AV+Obinutuzumab, infectious complications should be considered, especially in older patients

# TP53 in Mantle Cell Lymphoma

10- 40% of ND MCL, higher frequency at relapse.

- Associated with a del17p, high Ki-67%, CK, blastoid histology, and MYC rearrangement.

TP53 muts (w/wo del17p) had the **worst OS**.

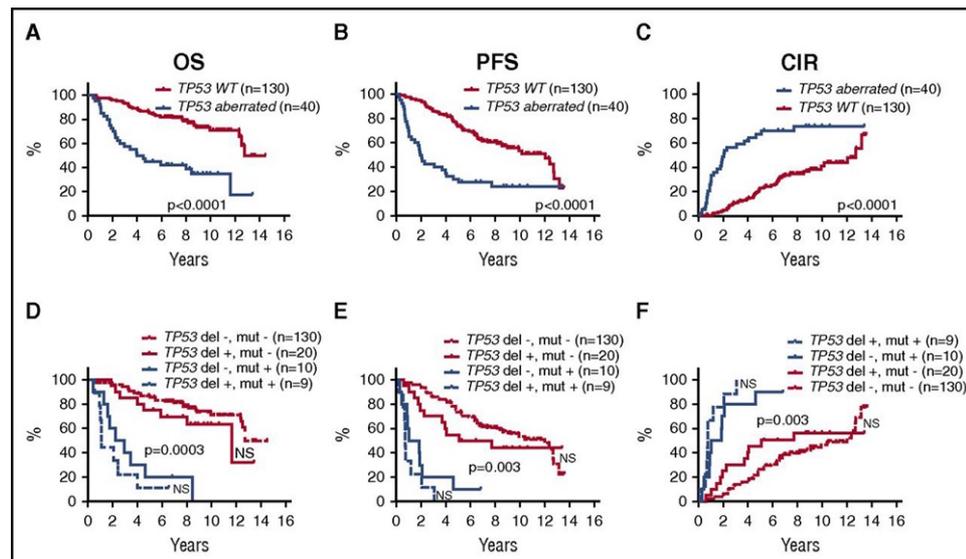
- Independent of genomic complexity
- Concomitant mutations in NOTCH1, CDKN2A, NSD2, and SMARCA4 associated with **ibr-ven resistance**

**High-risk MCL (unstable genome)**

Nadeu F, et al. Blood. 2020;136(12):1419-1432.

Jain P & Wang M. Blood. 2025;145(7):683-695.

## Prognostic impact of TP53 deletions and mutations in MCL



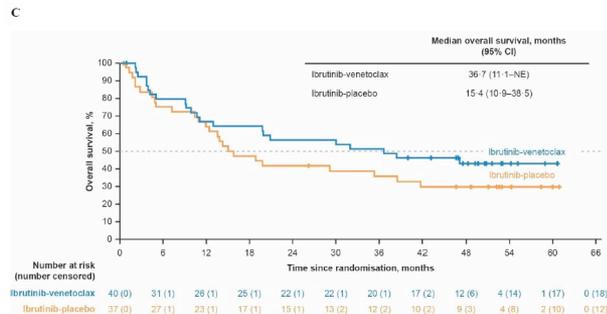
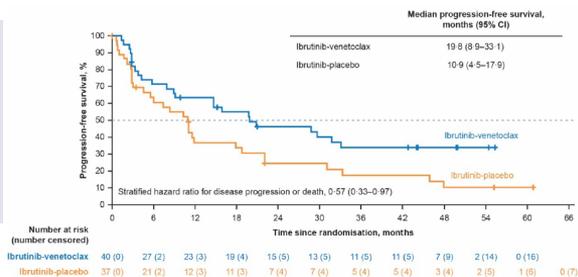
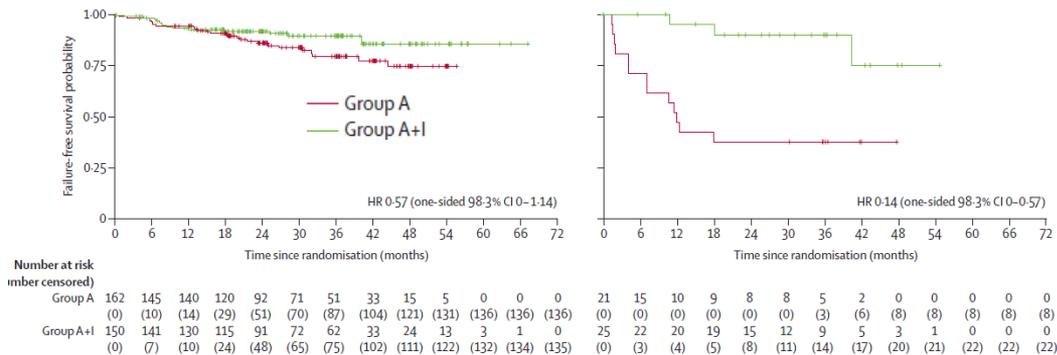
Eskelund CW, et al. Blood. 2017;130(17):1903-1910.

# TP53 and targeted agents

**In ND MCL**, the TRIANGLE trial demonstrated a failure-free survival benefit with the addition of ibrutinib to induction and maintenance therapy in pts with p53 overexpression

**In r/r classical MCL**, the SYMPATICO trial ibr + ven resulted in higher response rates and survival benefit in TP53-mut

Failure-free survival for group A+I vs group A in low ( $\leq 50\%$ ) p53 (C), and high ( $> 50\%$ ) p53 (D) (age  $< 65$  yrs and eligible for auto SCT)



Dreyling M, et al. Lancet. 2024;403(10441):2293-2306.

Wang M, J et al. Lancet Oncol 2025;26:200-213



# Zanubrutinib, obinutuzumab, venetoclax for 1st-line treatment of MCL with *TP53* mutation

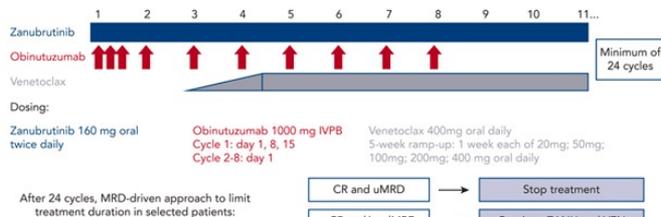
- **BOVen was safe and effective for untreated *TP53*-mutant MCL.**
- **Longer follow-up would determine the sustainability of such responses.**

## Context of Research

- *TP53*-mutant MCL is associated with poor survival outcomes with standard chemoimmunotherapy.
- We tested dual BTK and BCL2-inhibition with anti-CD20 monoclonal antibody therapy in *TP53*-mutant MCL

## Patients and Methods

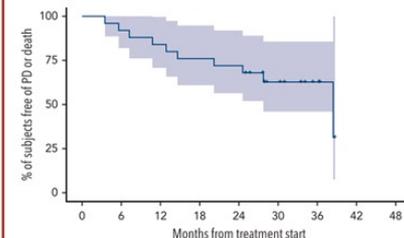
- Phase 2 clinical trial of zanubrutinib, obinutuzumab, and venetoclax (NCT03824483). Primary outcome measure: 2-year progression-free survival
- Enrolled 25 MCL patients with *TP53* mutation. Treatment schema (BOVen):



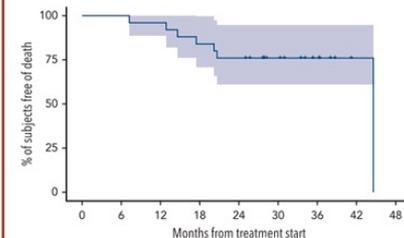
## Main Outcomes

- **Best Overall Response Rate 96% (24/25) and Complete Response Rate 88% (22/25).**
- **Toxicity was manageable. 32% (8/25) w/neutropenia, no febrile neutropenia, 20% (5/25) received growth factor support.**

- **2-year PFS: 72% (56, 92)**

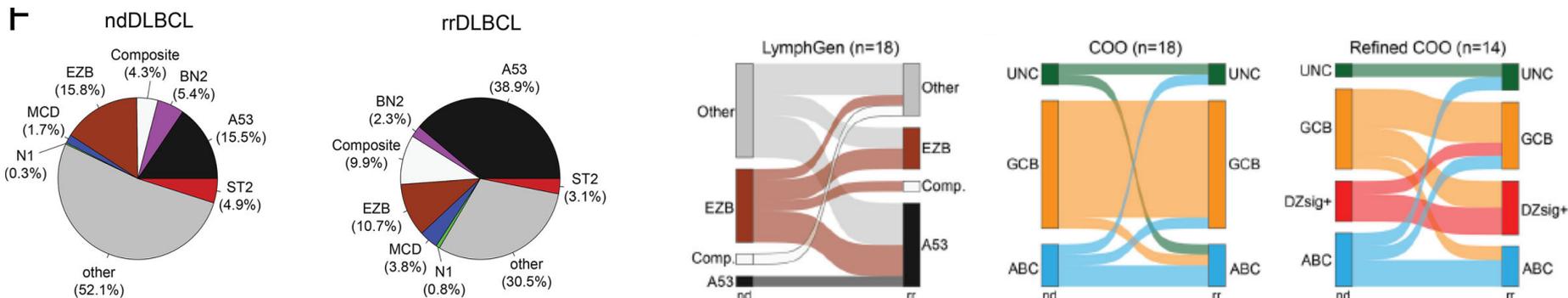


- **2-year OS: 76% (61, 95)**





# Mechanisms of treatment resistance and relapse in DLBCL



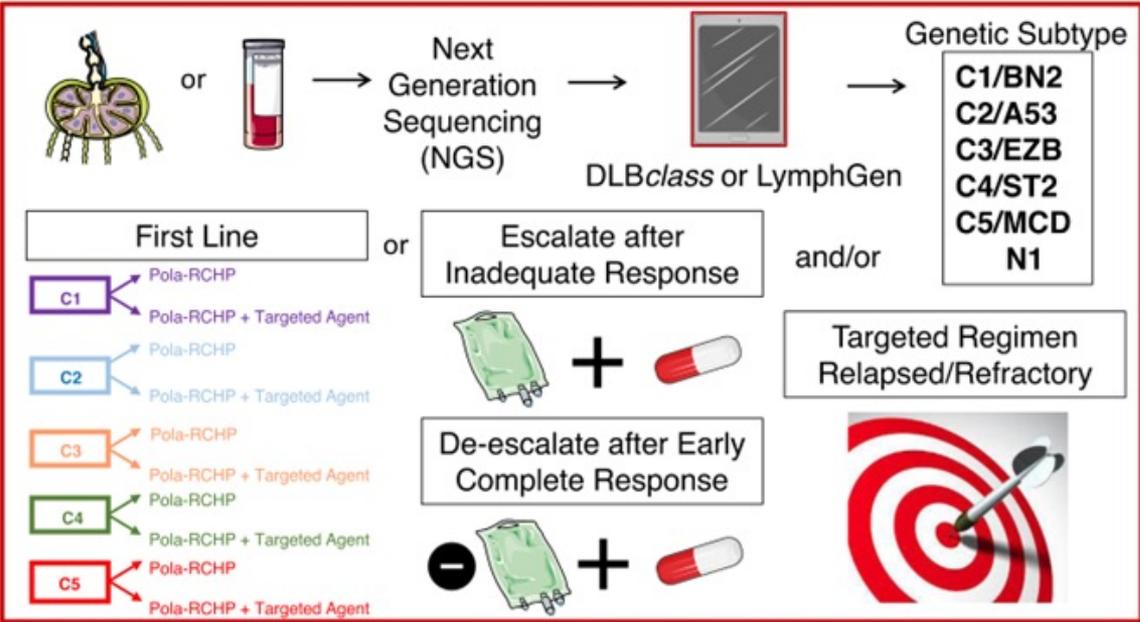
## rrDLBCL pts have

- More DNA gains, losses, and rearrangements including TP53 depletion
- More TP53 somatic mutations, 17p13 copy number variations and higher TP53 VAF
- Clonal expansion of tumor populations harboring TP53 mutations

Alterations in TP53 represent **one the different subtypes of specific mechanisms of resistance**



# DLBCL Molecular Subtypes to Tailor Treatment



**Conclusions:** Genetic subtypes can be determined using NGS and DLBclass or LymphGen classifiers. Treatment could be tailored in first line with the proposed LymphoMatch clinical trial algorithm, or in patients with inadequate or early complete responses, and/or in relapsed/refractory DLBCL. **Rutherford SC. DOI**

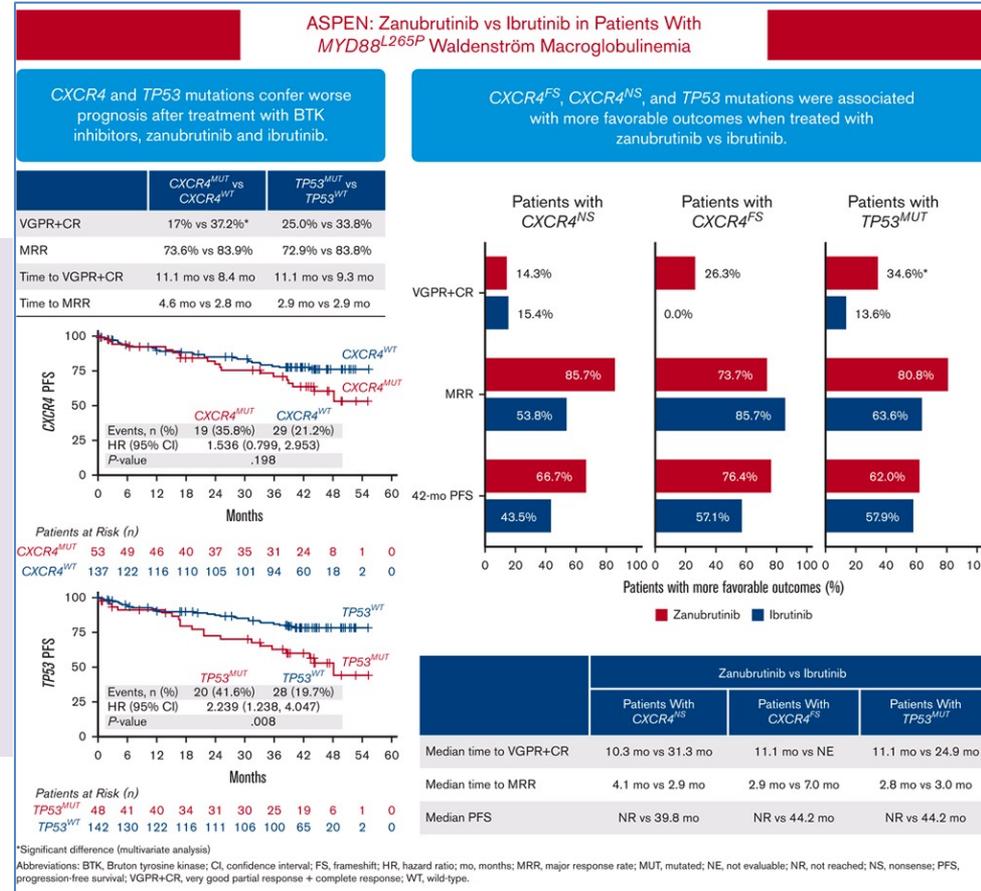
# TP53 and Waldenström macroglobulinemia

## TP53 abns

- In 5-10% treatment-naïve pts
- In 25%-30% of previously treated pts
- **shorter OS and PFS with BTKi**
- **May guide treatment considerations**
  - Pts with CXCR4 or TP53 muts had more favorable outcomes with **zanu vs ibr**

Tam CS, et al. Blood Adv. 2024 Apr 9;8(7):1639-1650.

Treon SP, et al. Blood. 2024 Apr 25;143(17):1702-1712.



# Conclusions

- Treatment with **targeted agents might prevent the expansion** of TP53-mut clones.
- Data on TP53 mutations **is still evolving** in the targeted agent setting.
- The importance of **precise classification** of TP53 abns in the design of clinical trials in order to obtain robust evidence for improving the treatment tailoring.

TP53 status	Strategies	Drugs
No p53	p53 gene delivery	Gendicine
wt p53	Disruption of wtp53-MDM2/MDM4 interaction	Nutlin; RG7112; SAR405838; KT253
p53 missense variants	Restoration of p53 functions	APR-46; ATO; PC14586; CP-31398; COTI-2; RETRA
	Eradication of p53 variants	
	Synthetic lethality	Adavosertib
p53 nonsense variants	boost anti-tumor immunity	SGT-53
	Restoration of p53 functions	Aminoglycosides 5-Fluorouridine

Baliakas P, Soussi T. J Intern Med. 2025 Aug;298(2):78-96.