

# Minor-clone *TP53* mutations in CLL patients entering first-line treatment: clonal evolution and clinical impact

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# I have no disclosures



# Aims

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- To assess the risk of clonal expansion of minor-clone *TP53* mutations
- To evaluate the clinical impact of *TP53* mutations with different variant allele frequencies (VAFs)

**...in patients entering first-line treatment**

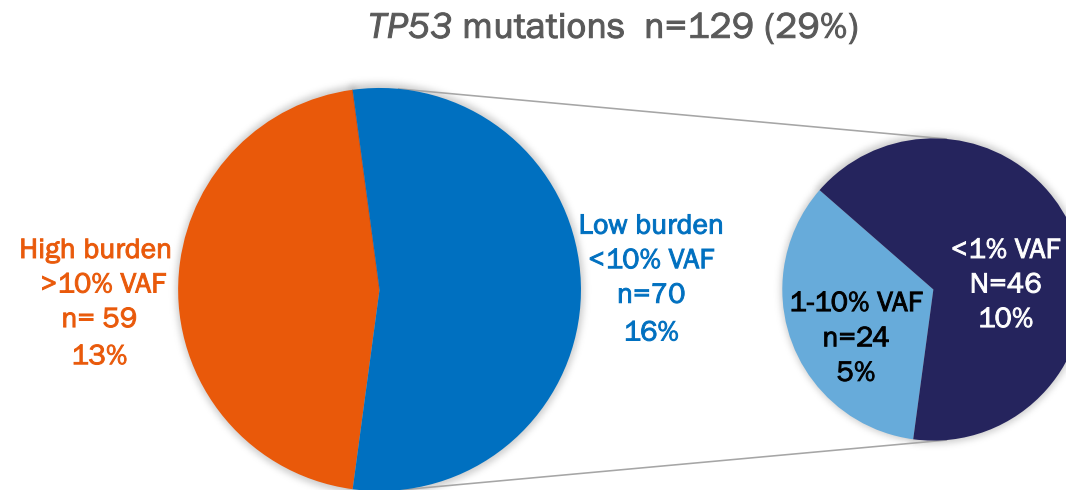
*“TP53 gene assessment should always be performed prior to initiation of the first and every subsequent line of treatment“*

*iwCLL guidelines: Hallek et al., 2018*

*TP53 recommendations: Malcikova et al., 2019*

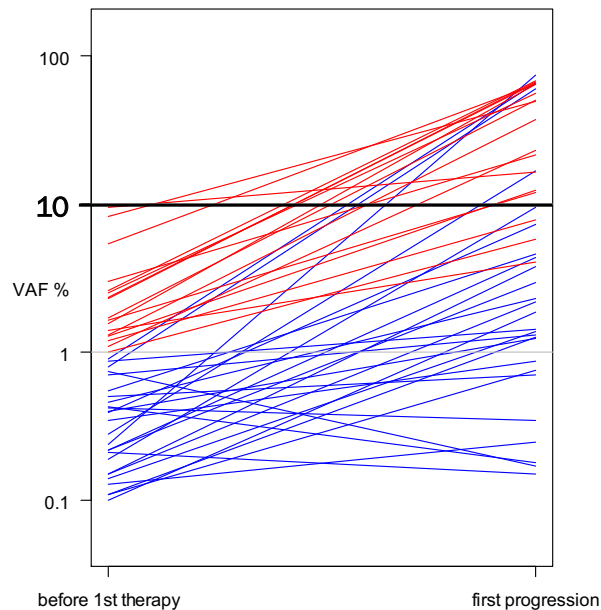
# Patient cohort

- 450 patients
  - 72% unmutated IGHV
  - 84% received chemoimmunotherapy
- Amplicon NGS *TP53* analysis with detection limit 0.1-0.2% VAF

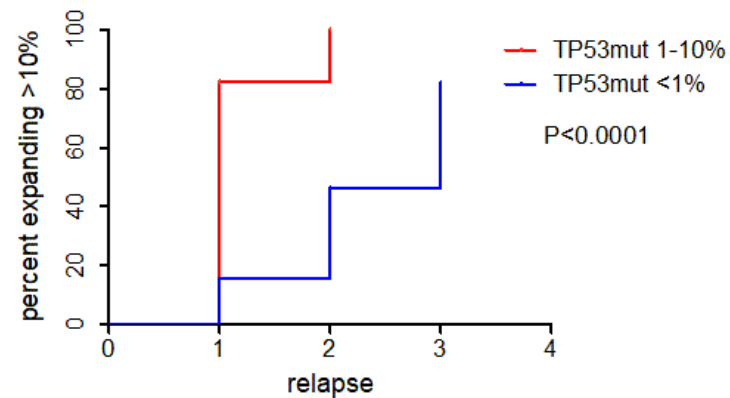


# Clonal evolution of minor-clone *TP53* mutations

- Samples in first relapse available in 43 patients with detected low-burden *TP53* mutation

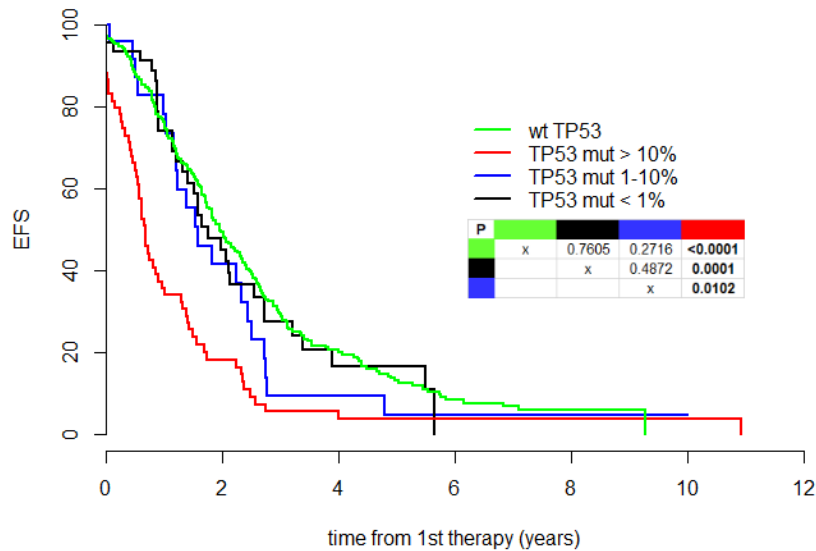


- 88% clonal expansion  $>1.5x$
- Median fold change = 10x (0.2-309x)
- Increase in mutation VAF above 10%
  - Patients with mutation 1-10% VAF before treatment – 82% (14/17)
  - Patients with mutations  $<1%$  VAF before treatment – 15% (4/26)
    - Gradual expansion in subsequent relapses

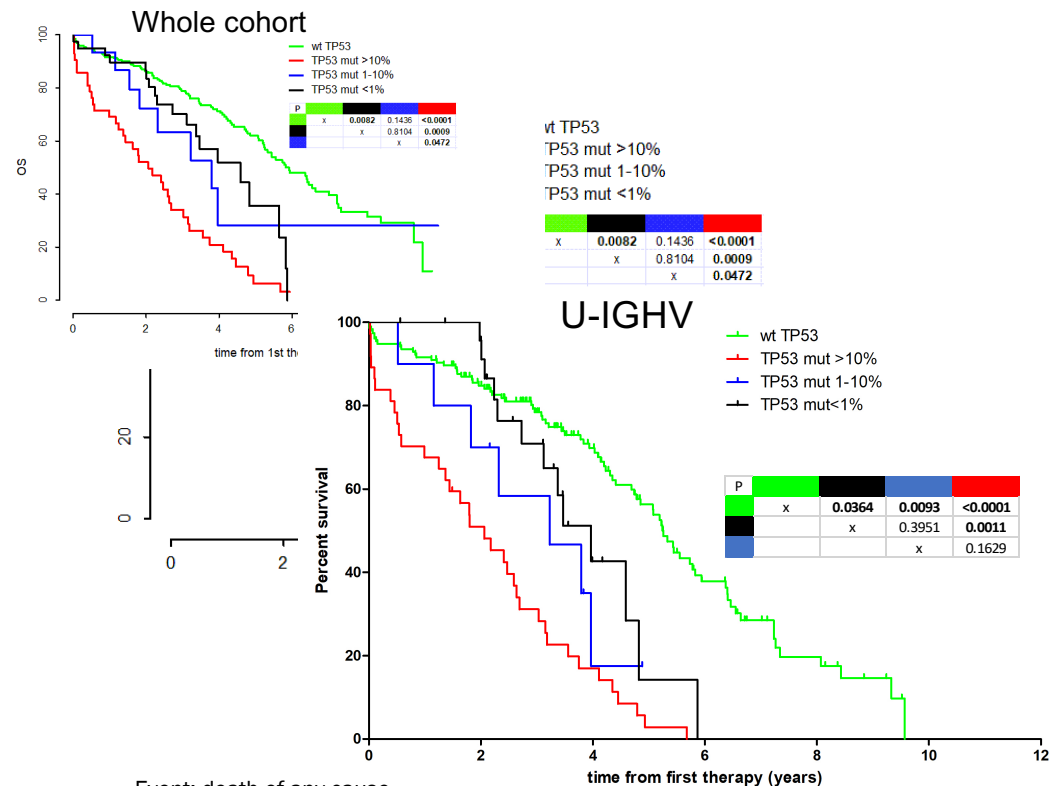


# Clinical impact

## Event-free survival



## Overall survival



Event: death of any cause  
 Excluded from OS analysis: switched to targeted therapy (88), allo-HSCT (14)

# Conclusions

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- Low-burden *TP53* mutations detected before first-line treatment
  - Increase in clonal proportion in first relapse in majority of patients with median fold change 10x
  - Did not significantly shorten response to first-line therapy
  - Affected OS at least in unmutated IGHV patients
- Unresolved issues remain → multicenter study is needed

Thank you for your attention