

A functional prognostic model predicts progression free survival in patients with relapsed chronic lymphocytic leukemia treated with ibrutinib + venetoclax

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Let me guide you!

BACKGROUND

- Current prognostic models in CLL were developed to predict responses to chemoimmunotherapy
- These models have limited predictive value for newer combinations of targeted therapies

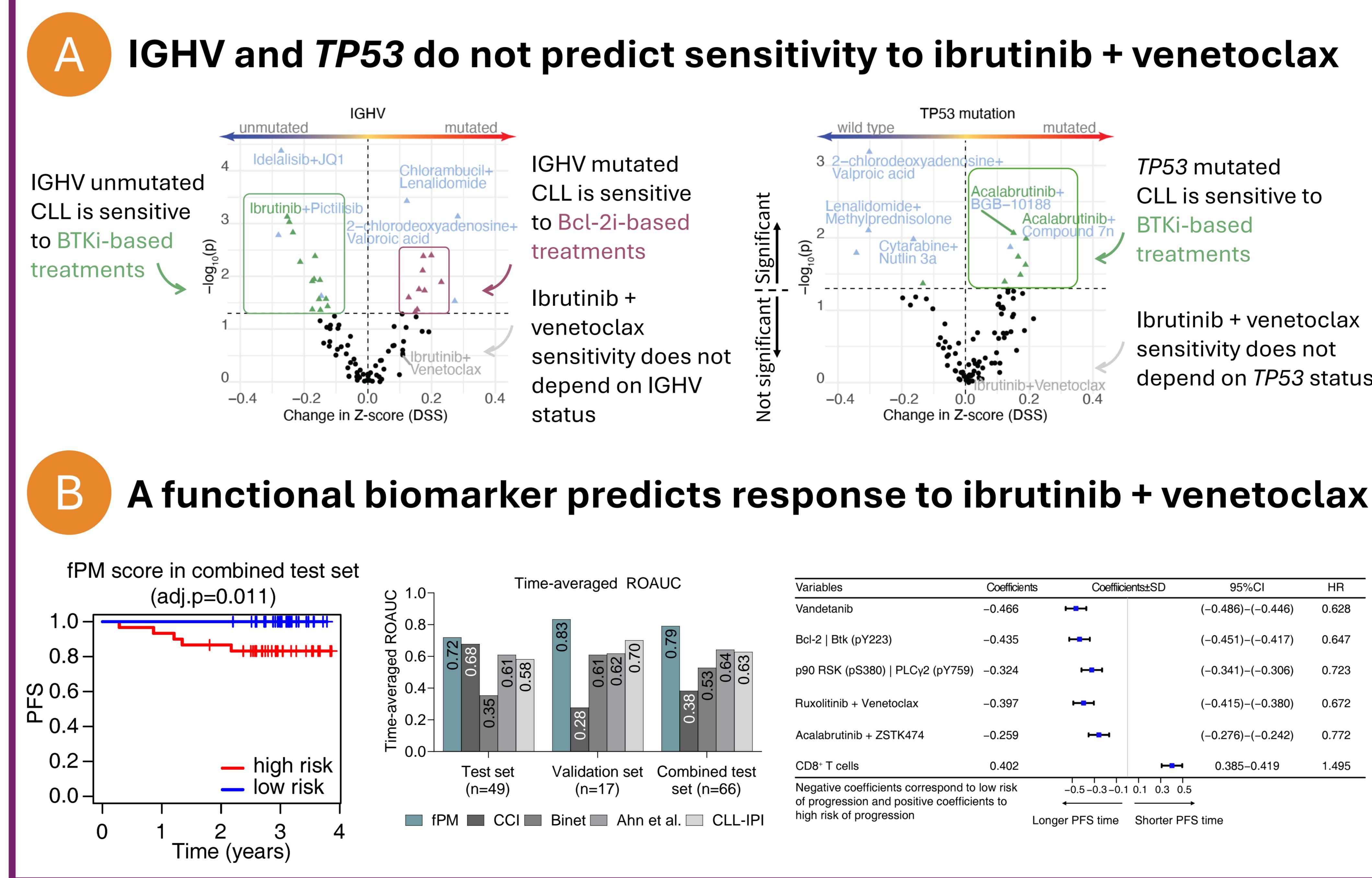
“Accurate prognostic models are needed to guide optimal treatment with targeted therapies”

AIM

- To identify predictors of PFS in response to BTK + Bcl-2 inhibitor combination therapy

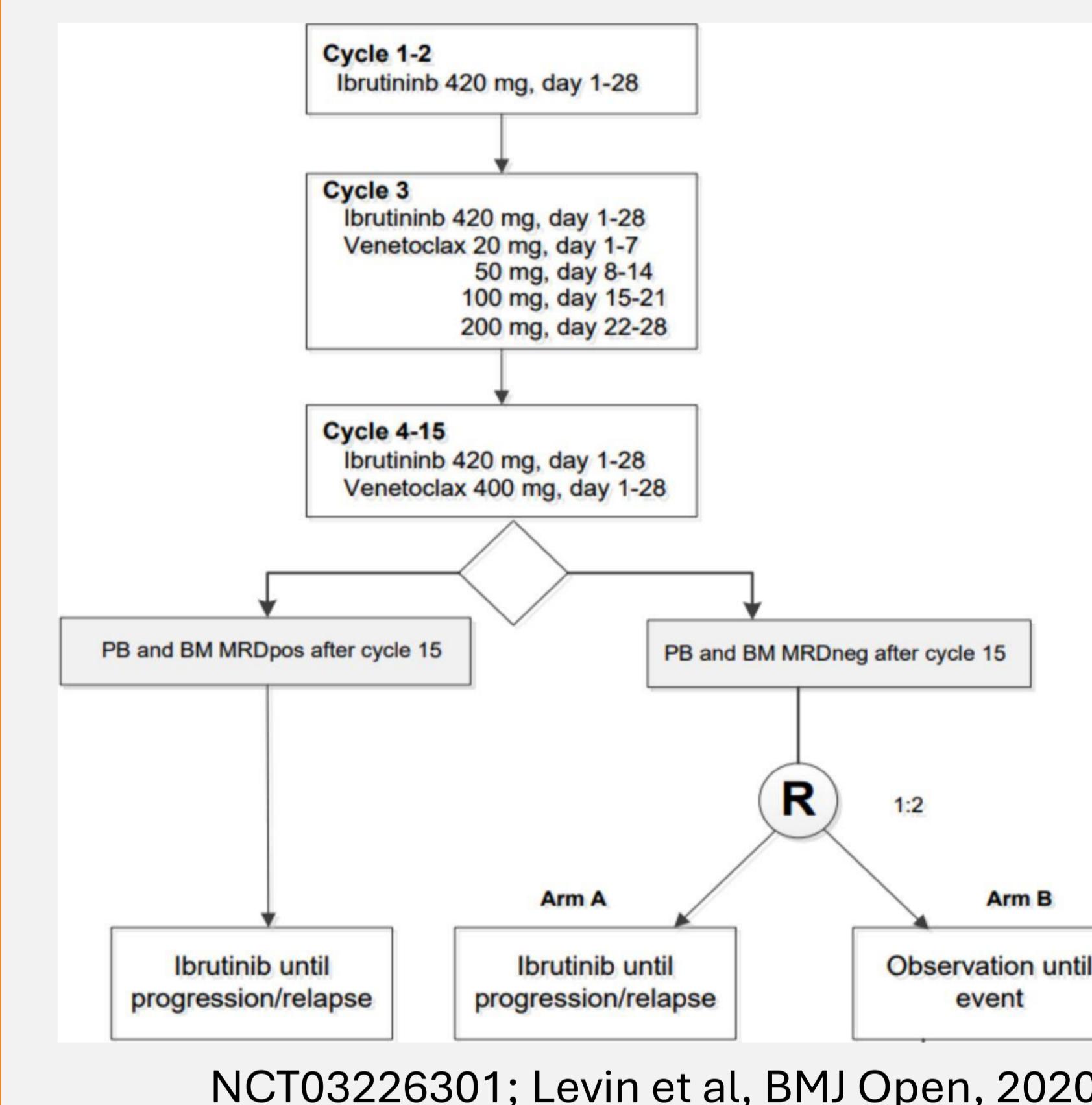
RESULTS

- IGHV and *TP53* do not predict ex vivo sensitivity to ibrutinib + venetoclax
- A functional biomarker predicts response to ibrutinib + venetoclax



“A functional biomarker score including signaling pathway phosphorylation, drug sensitivity, and T cells predicts PFS for R/R CLL treated with ibrutinib + venetoclax”

HO141/VISION TRIAL



GLOSSARY

Bcl-2	B-cell lymphoma-2
BTK	Bruton's tyrosine kinase
CCI	Charlson comorbidity index
CLL	Chronic lymphocytic leukemia
CLL-IPI	International prognostic index for CLL
fPM	Functional prognostic model
IGHV	Immunoglobulin heavy variable
MRD	Minimal residual disease
PBMC	Peripheral blood mononuclear cell
PFS	Progression free survival
R/R	Relapsed/refractory
ROAUC	Receiver operating area under the curve
TP53	Cellular tumor antigen p53



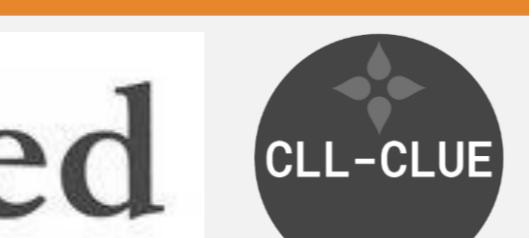
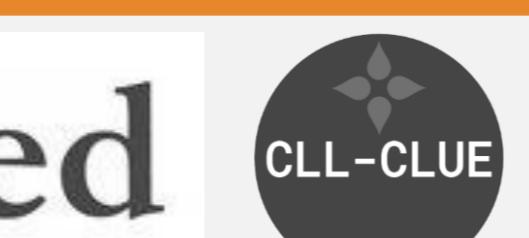
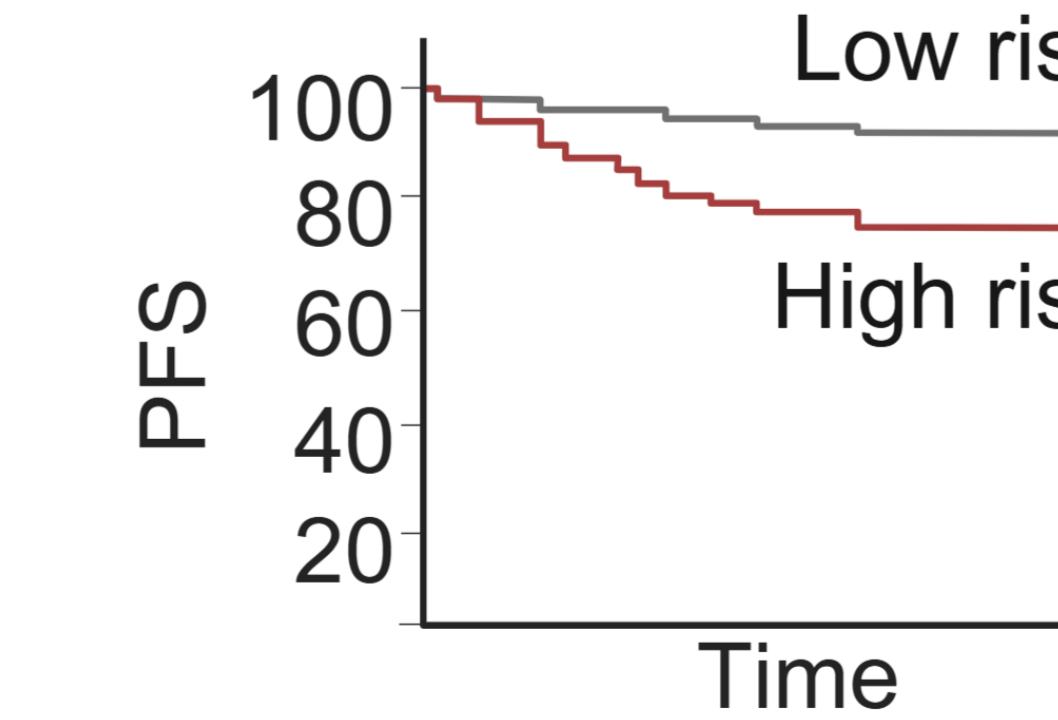
PBMCs from CLL patients (n=177) in the HO141/VISION trial (NCT03226301)



- Genetic
 - IGHV, *TP53*, karyotype
- Functional
 - Ex vivo drug sensitivity
 - Cell signaling analysis
 - Immunophenotyping
- Clinical
 - Binet stage, MRD, PFS



- Feature selection
- Machine learning



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