

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

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BACKGROUND

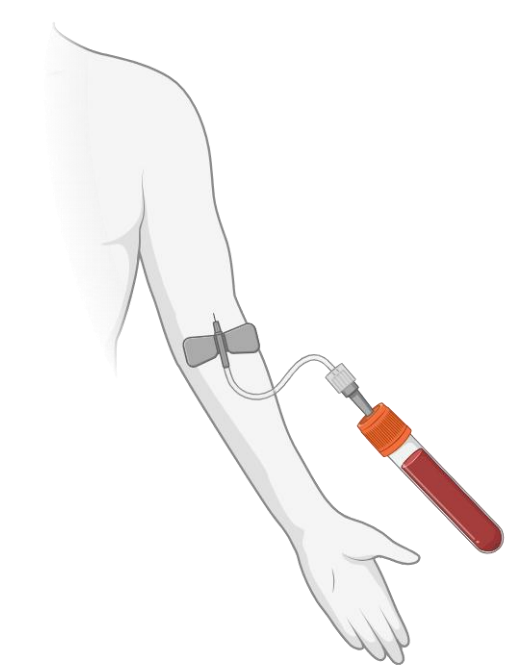
- Current prognostic models in CLL were developed to predict responses to chemoimmunotherapy
- These models have limited predictive value for newer combinations of targeted therapies

“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

1 Material

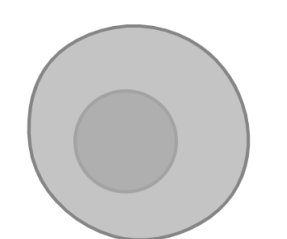


PBMCs from CLL patients (n=177) in the HO141/VISION trial (NCT03226301)

2 Measurements

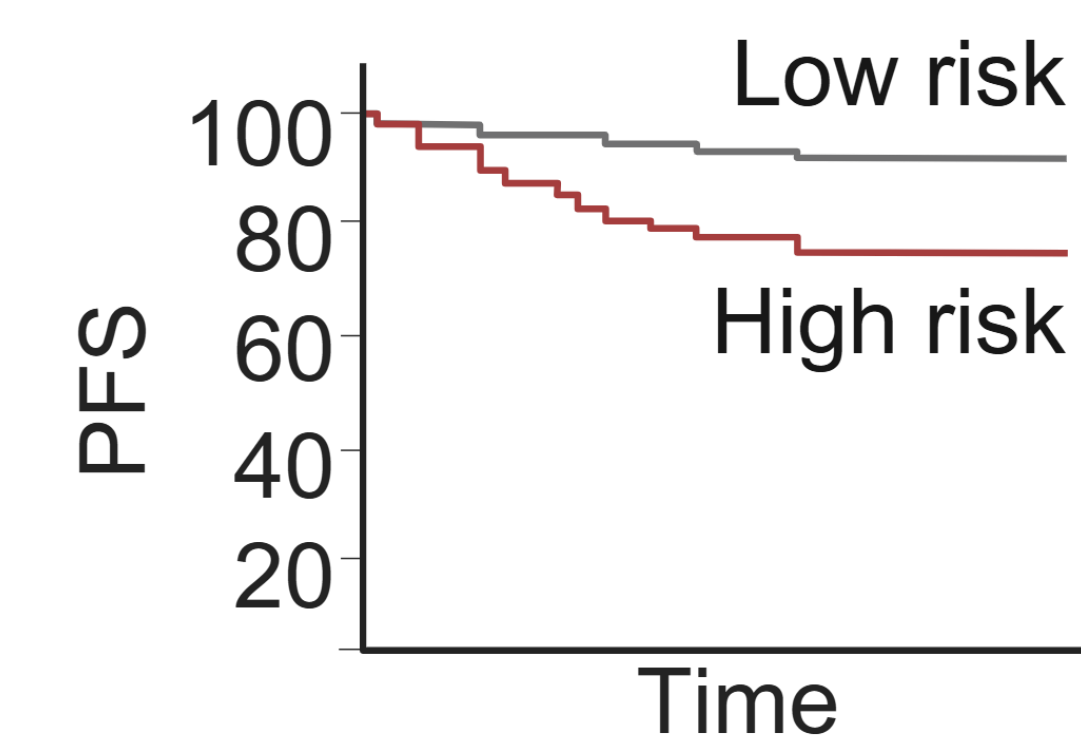


- Genetic
 - IGHV, TP53, karyotype
- Functional
 - Ex vivo drug sensitivity
 - Cell signaling analysis
 - Immunophenotyping
- Clinical
 - Binet stage, MRD, PFS



3 Modeling

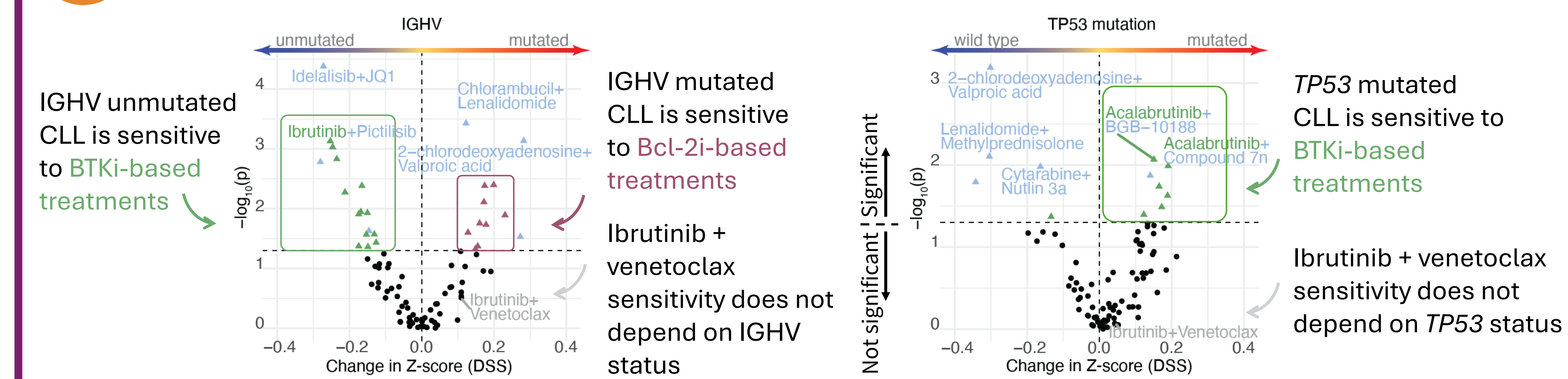
- Feature selection
- Machine learning



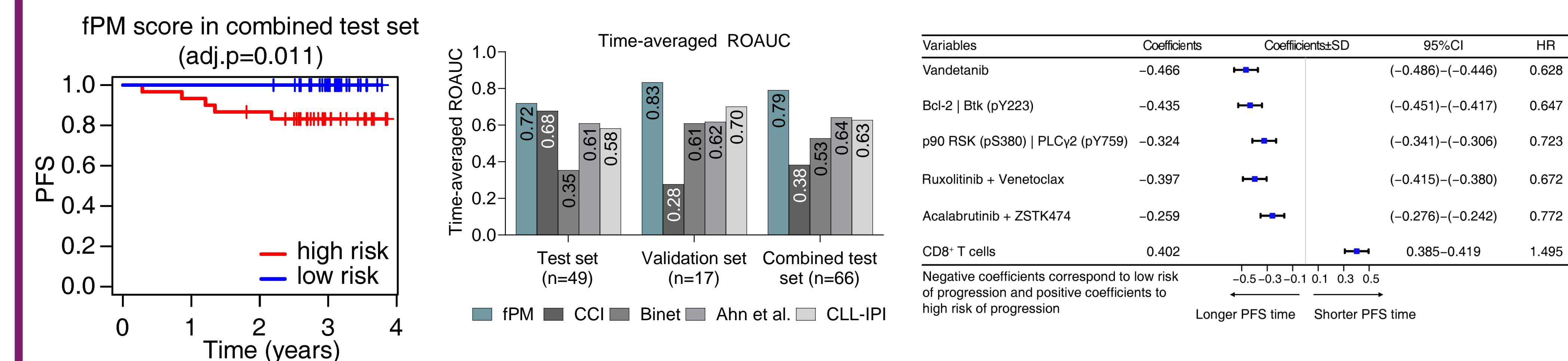
RESULTS

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

A IGHV and TP53 do not predict sensitivity to ibrutinib + venetoclax



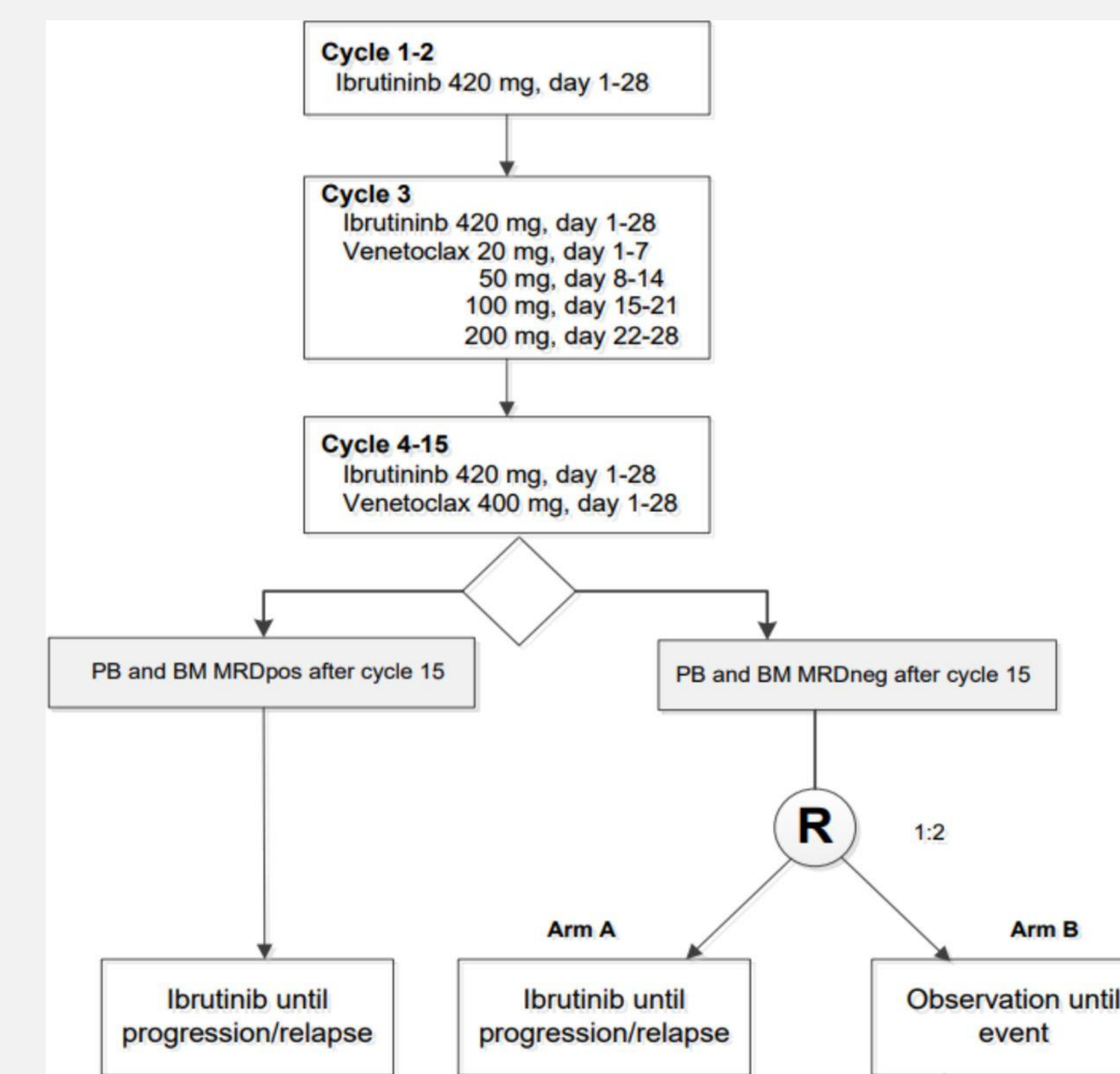
B A functional biomarker predicts response to ibrutinib + venetoclax



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

Let me guide you!

HO141/VISION TRIAL



NCT03226301; Levin et al, BMJ Open, 2020

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