2nd 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)

BIOLOGY OF HIGH-RISK ACUTE LYMPHOBLASTIC LEUKAEMIA (IN CHILDREN AND YOUNG ADULTS)

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Jan Trka: No relevant conflict of interest

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ALL: genetics defines prognosis

- rarer in adults, the most frequent malignant disease in children: representation among malignancies
 Adults
- genetic aberrations: key role in pathogenesis
- heterogeneous biological nature → ALL subtypes
- excellent prognosis on modern therapy (children, AYA) AIEOP/BFM ALL 2000: CR=97.8% EFS=82.4% OS=90.4%
- significant improvement of treatment results: new schemes and adjustment of treatment by risk stratification plus new therapies

ALL risk stratification – AIEOP/BFM evolution

	BFM 70	BFM 76/79	BFM 81	BFM 83	BFM 86	BFM 90	BFM 95	ALL- IC	BFM 2000	BFM 2009	
Hepatosplenomegaly											
CNS involvement											
Mediastinum involvement											
Age											
WBC/blasts in PB											
Cytochemistry											
T-ALL											
BCR/ABL											
MLL/AF4											Genetics
Hypodiploidy											
Prednison response											
Morphology D+15											Early response
MRD (FCM D+15)											to treament
Morphology (D+33 CR)											(MRD)
MRD (qPCR D+33, D+78)											

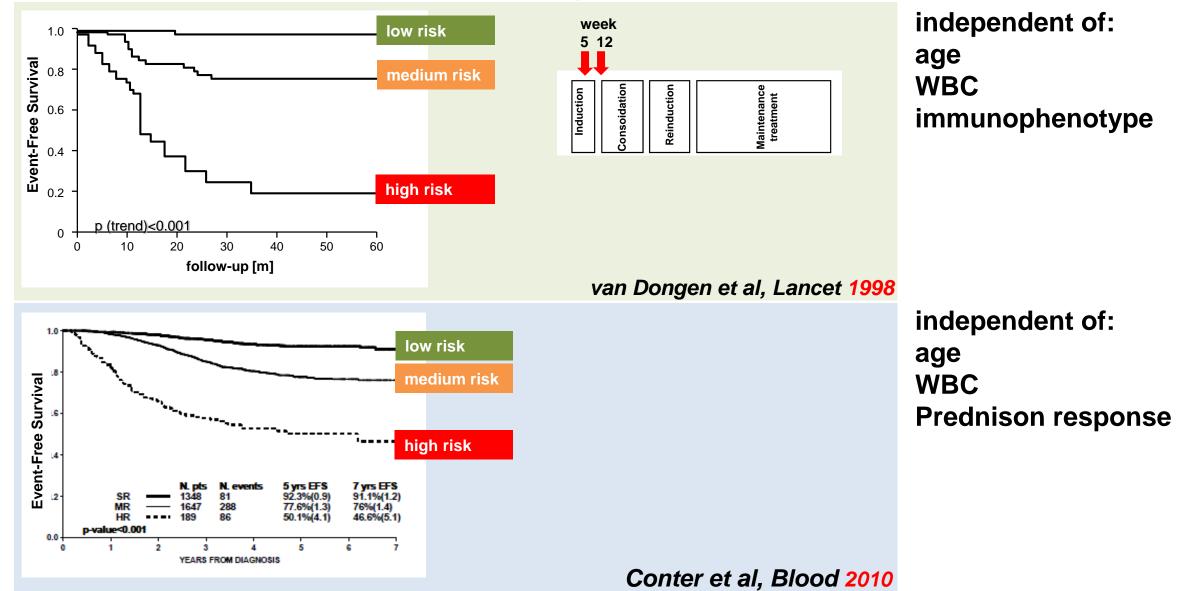
How to identify High-Risk ALL

Genetically-defined groups with empirically known poor overall outcome

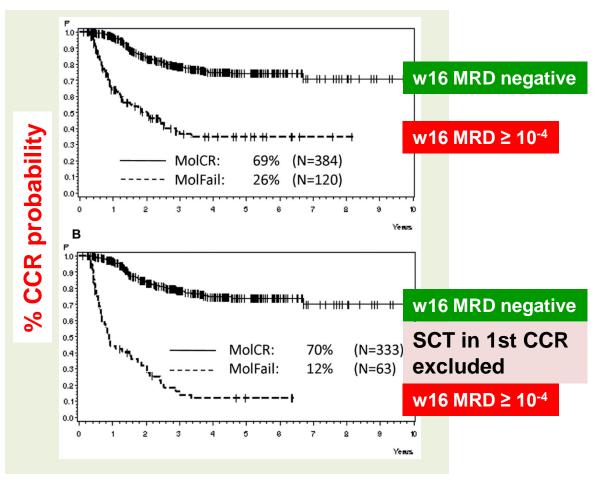
High Minimal Residual Disease during (early) phase of treatment predicts poor overall outcome

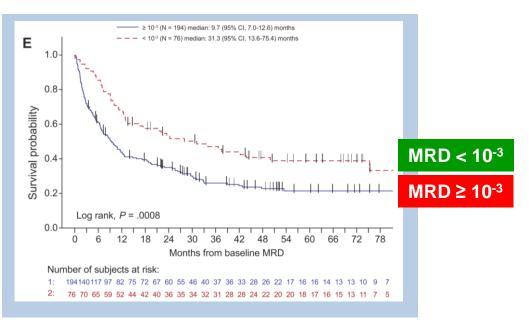
Prediction of relapse: MRD

based on MRD in two time-points in initial treatment



MRD confers poor prognosis also in adult ALL





Gokbuget et al, Hematology 2019

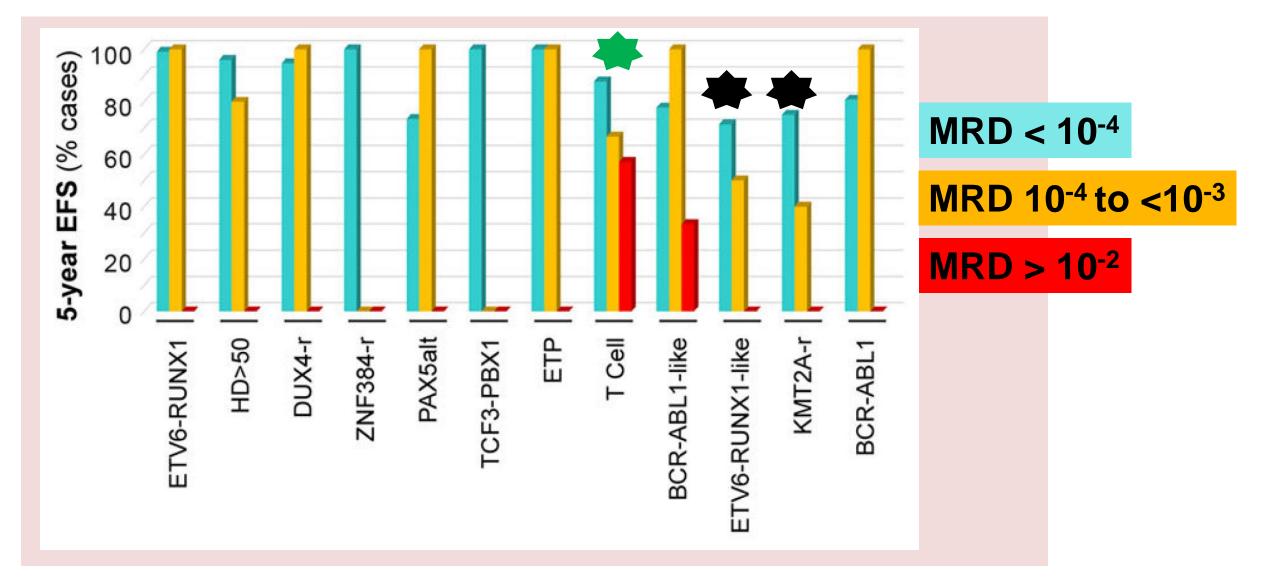
Gokbuget et al, Blood 2012

CHE

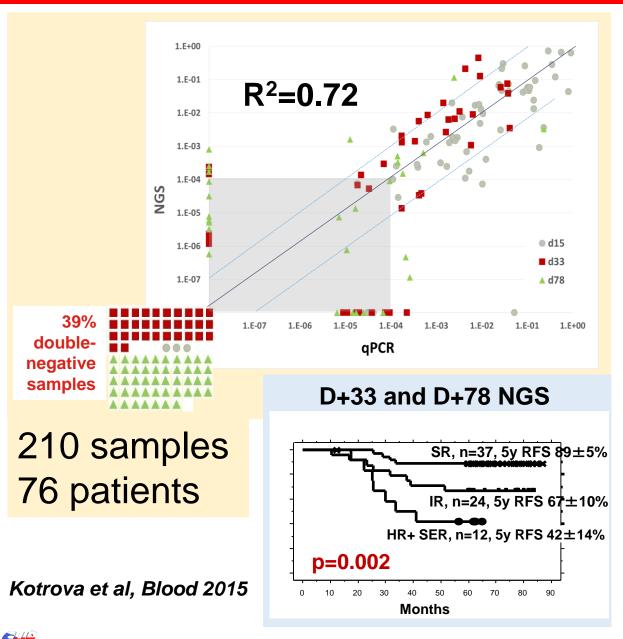
Systematic meta-analysis reveals MRD as the strongest predictor of outcome in B-cell adult ALL

Bassan et al, Haematologica 2019

Predictive value of MRD differs in subtypes of ALL



NGS MRD is equally predictive as qPCR MRD



CLIP study (unpublished data): 437 consecutive newly diagnoses ALL patients 780 IG/TR MRD markers

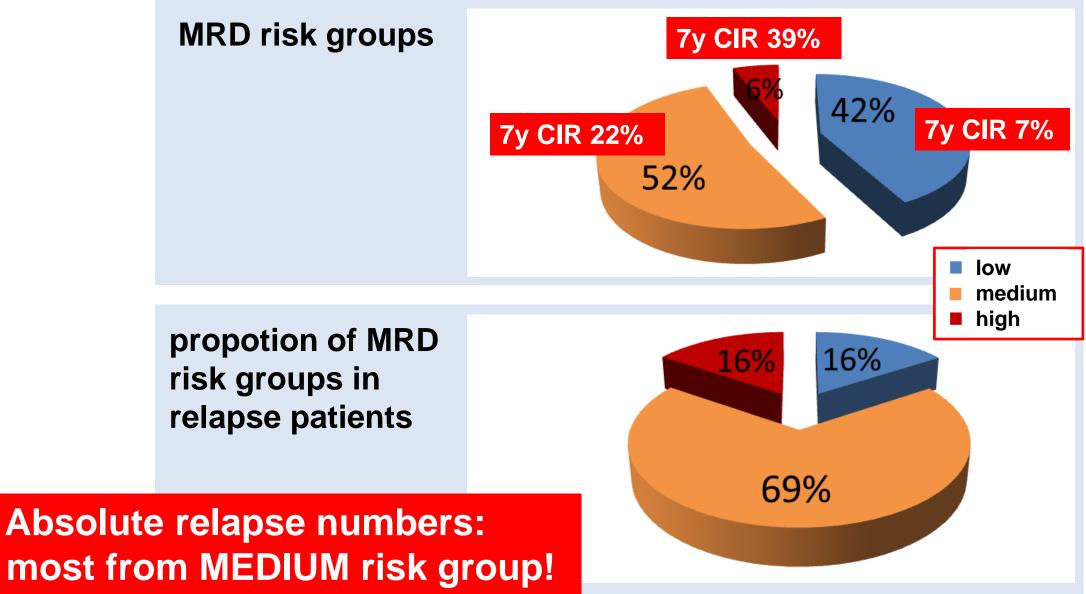
80.6% concordant MRD results7% (54) markers/results ruled out asfalse-positive by NGS

10% patients assigned to lower-risk group (based on Eol MRD level)

Large datasets of identified and tested **IG/TR rearrangements**:

selecting markers of adequate specificity (10% of markers selected by traditional technique not enough specific!)

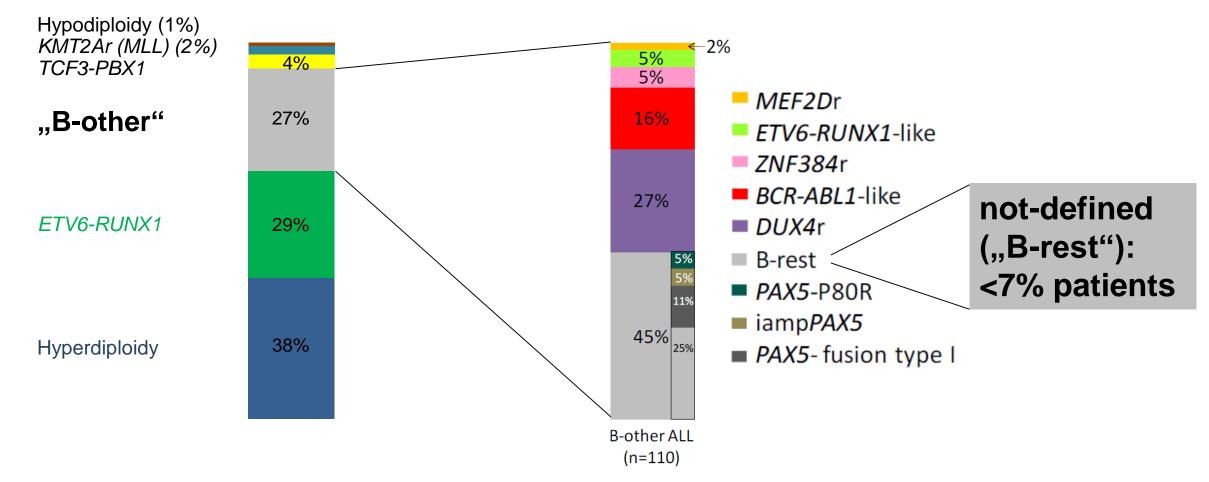
Relapses according to risk groups



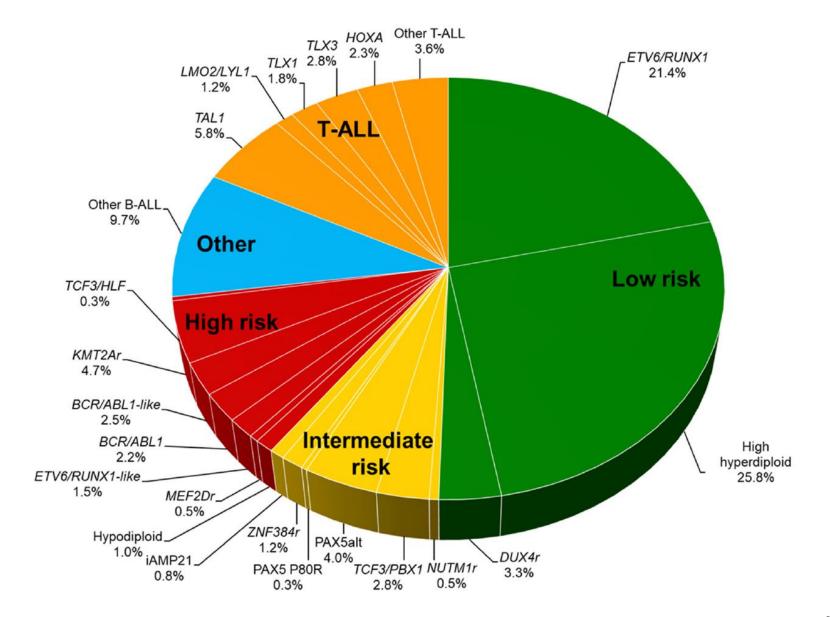
Conter et al, Blood 2010

Current genetic subtypes of ALL(Czech BFM ALL 2009)

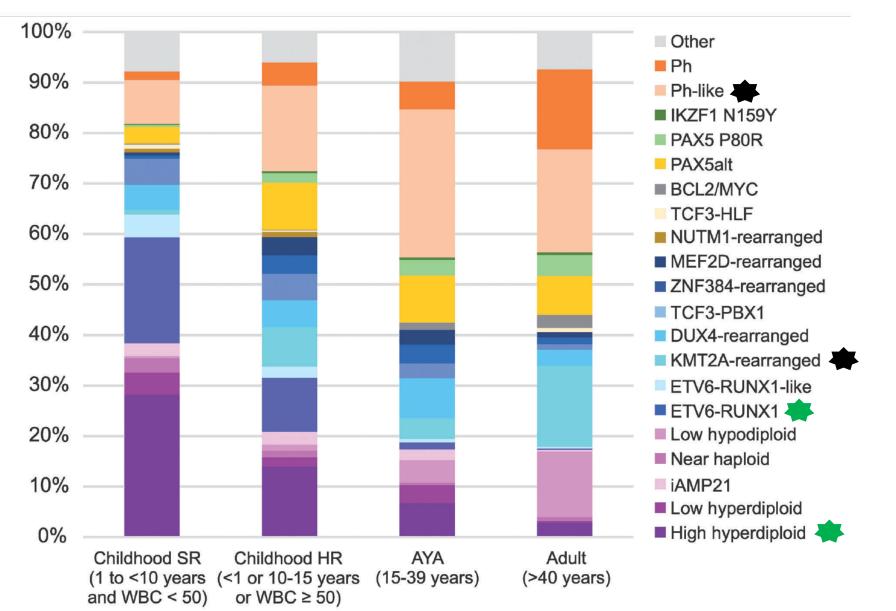
- heterogeneous group of diseases ('subtypes')
- genetic aberrations define biologic features of the cells
- various genotypes have different clinical outcomes



Separation of ALL subtypes into risk groups



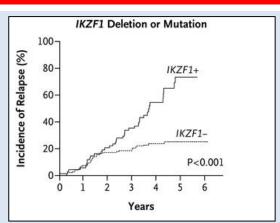
Representation of various genetic aberrations among age groups



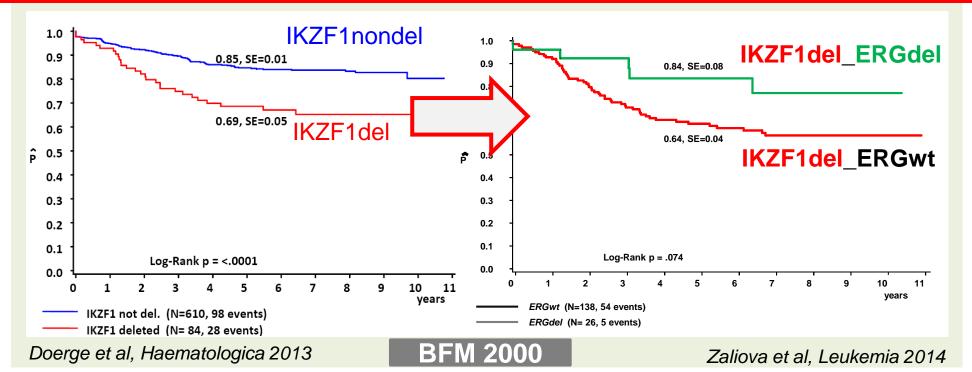
Inaba & Mullighan, Haematologica 2021

IKZF1 deletion (... and **ERG deletion**)

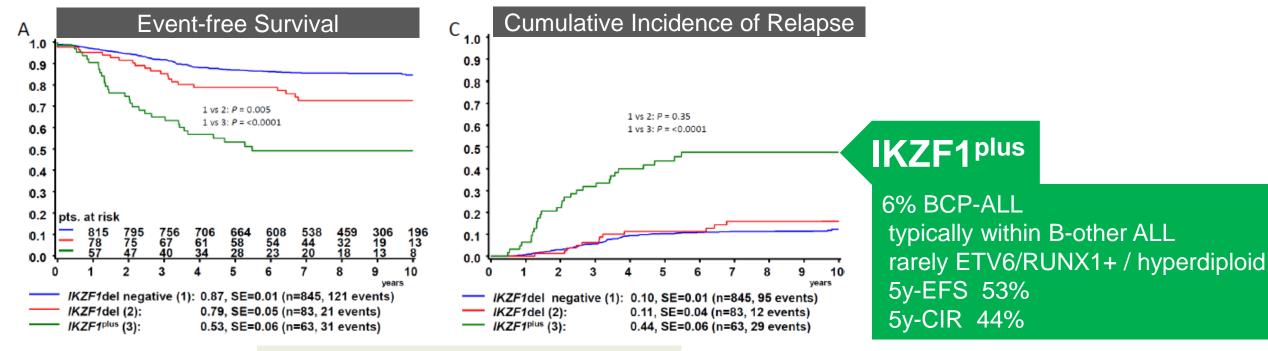
- identified with SNParray Mullighan et al, N Engl J Med 2009
- poor initial response (MRD)
- secondary aberration 29% HR ALL
- dismal outcome
- also in Ph+ ALL and BCR/ABL-like ALL Roberts et al, N Engl J Med 2014 van der Veer, Zaliova, Mottadelli et al, Blood 2014



in presence of ERGdel \rightarrow IKZF1del looses negative prognostic significance



Negative prognostic impact of IKZF1^{plus}



IKZF1^{plus}

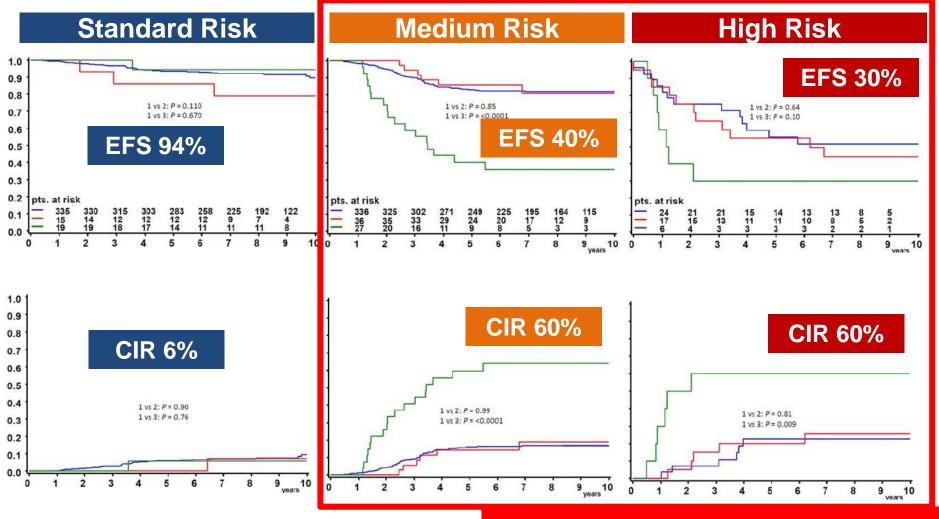
deletion IKZF1 and

- deletion PAX5 and/or
- deletion CDKN2A and/or
- deletion CDKN2B and/or
- deletion PAR1 (P2RY8-CRLF2) and
- (lack of ERG deletion)

Stanulla et al, J Clin Oncol 2018



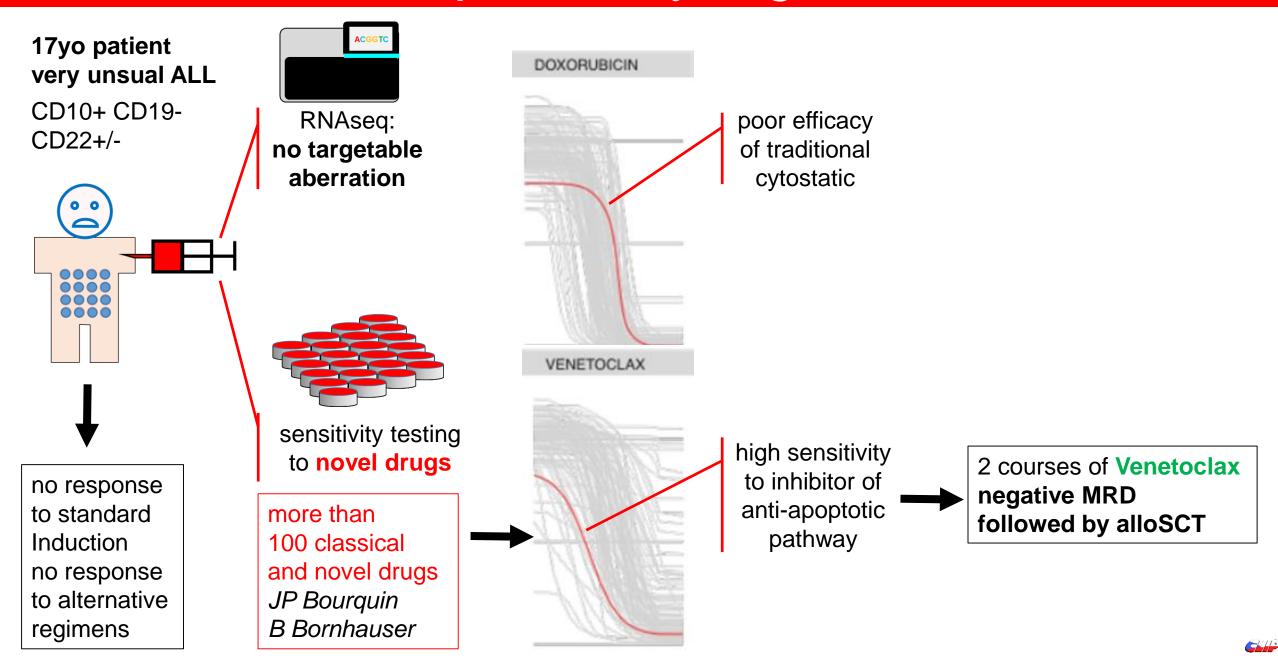
Prognostic impact of IKZF1^{plus} depends on MRD



stratification into High-Risk Group with **Blinatumomab** randomisation

Stanulla et al, J Clin Oncol 2018

Case Report: very-high risk ALL





- new subtypes of ALL recently described (incl. High-Risk) less clear definitions, overlaps, different age- and population-based frequencies
- MRD remains crucial in identification of High-Risk patients time-points specific for particular protocols, most relapses in MRD Medium-Risk group
- progressive integration of new prognostically relevant genetic subtypes into stratification algorithms

specific for particular protocols, MRD as important co-factor, non-randomizable small subgroups

- new methods in diagnostic approaches (NGS- and multiFCM-based) change in diagnostic paradigm: increased analytical and interpretational demand
- identification of new drugable aberrations? very rare in children x wider use of immunotherapy and/or new therapeutics (Venetoclax)