

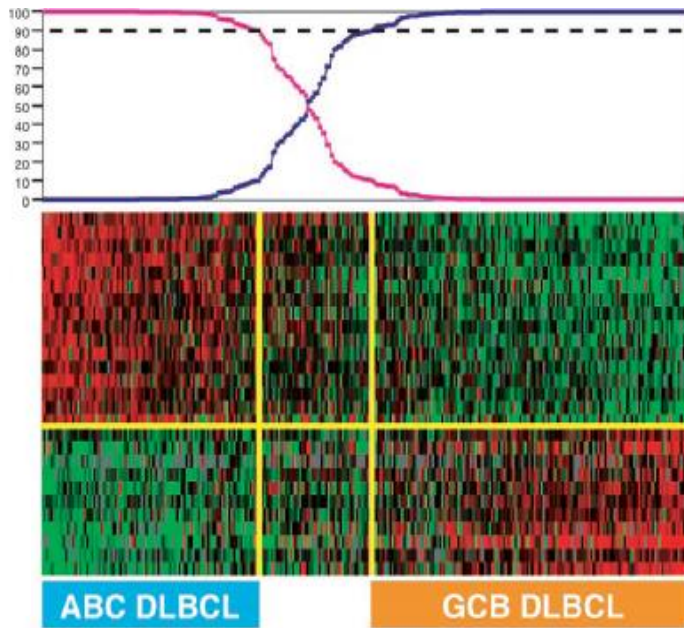
Biological Identification of High-Risk DLBCL

Georg Lenz

Norbert Schmitz

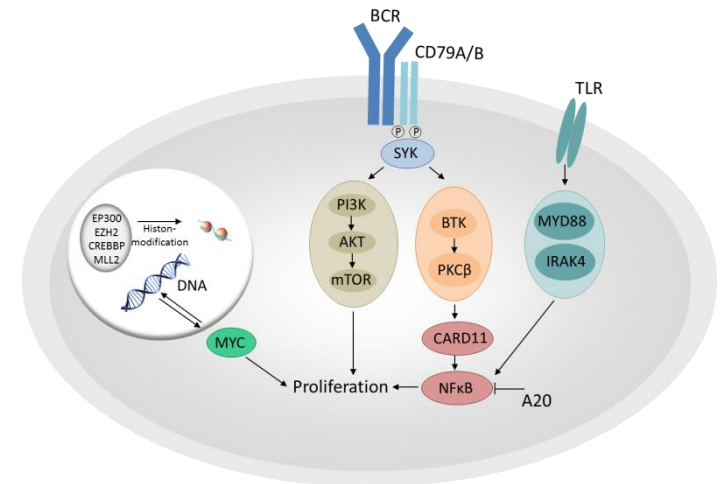
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Gene Expression Profiling Reveals Distinct Molecular Subtypes

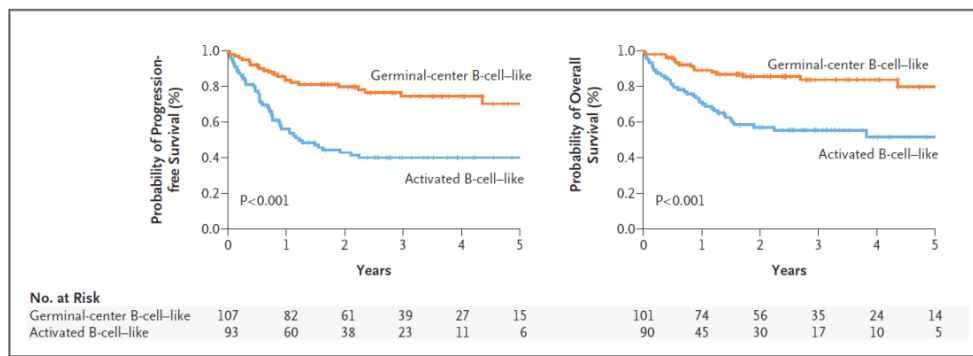


Key signaling pathways

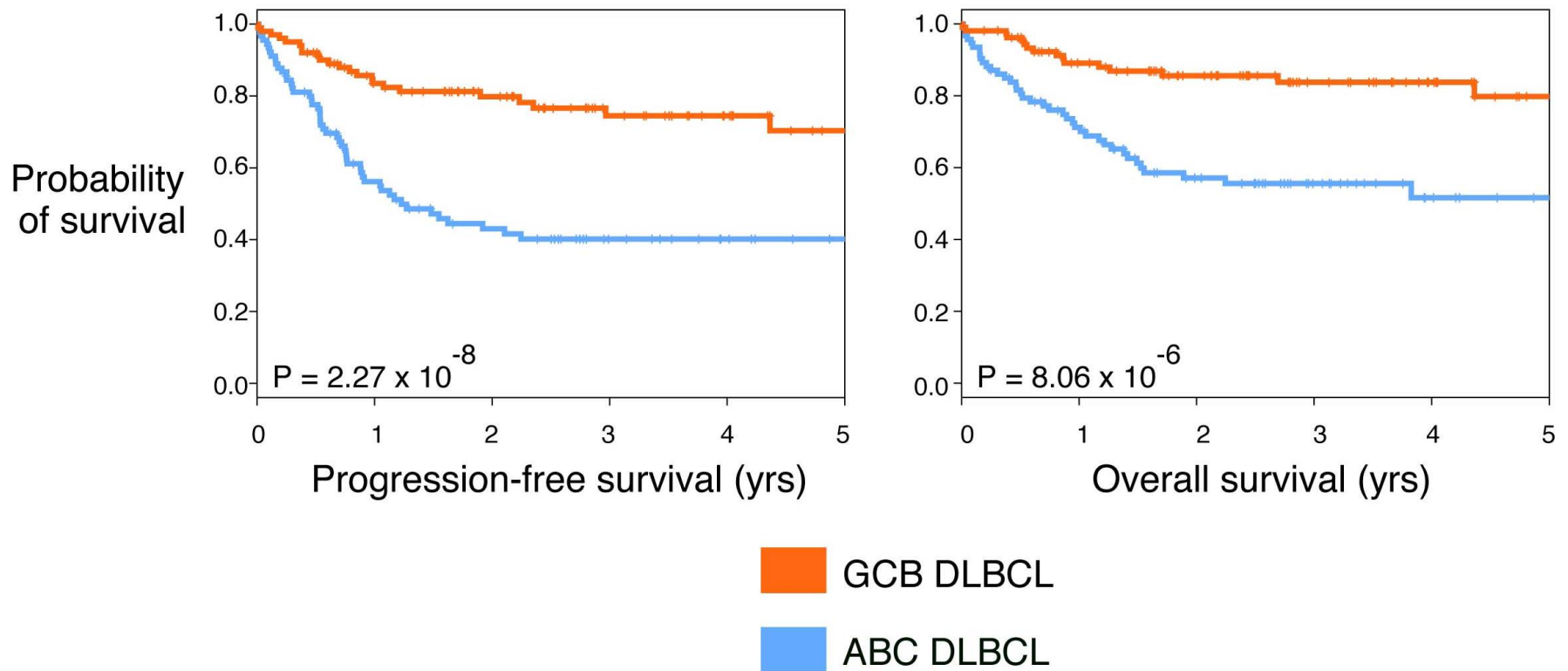
- ABC-DLBCL:
BCR signaling



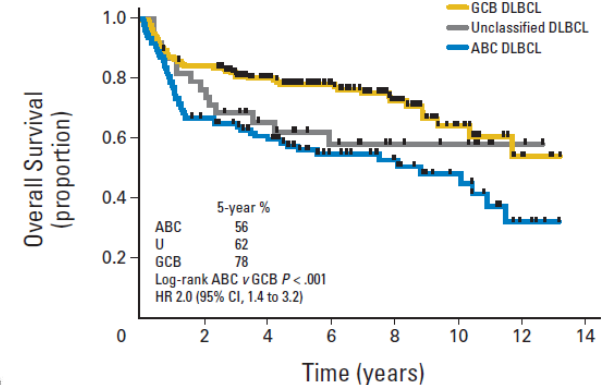
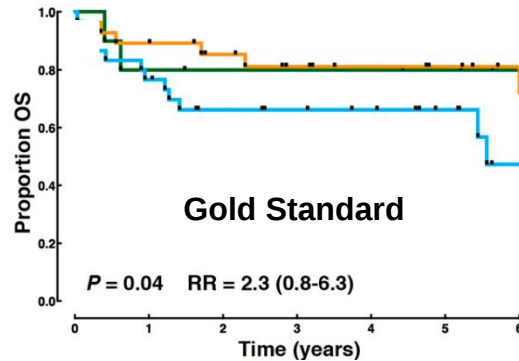
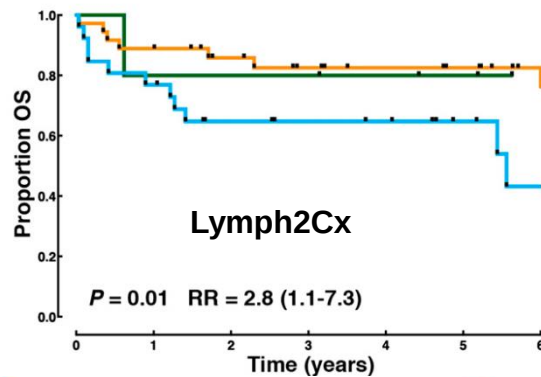
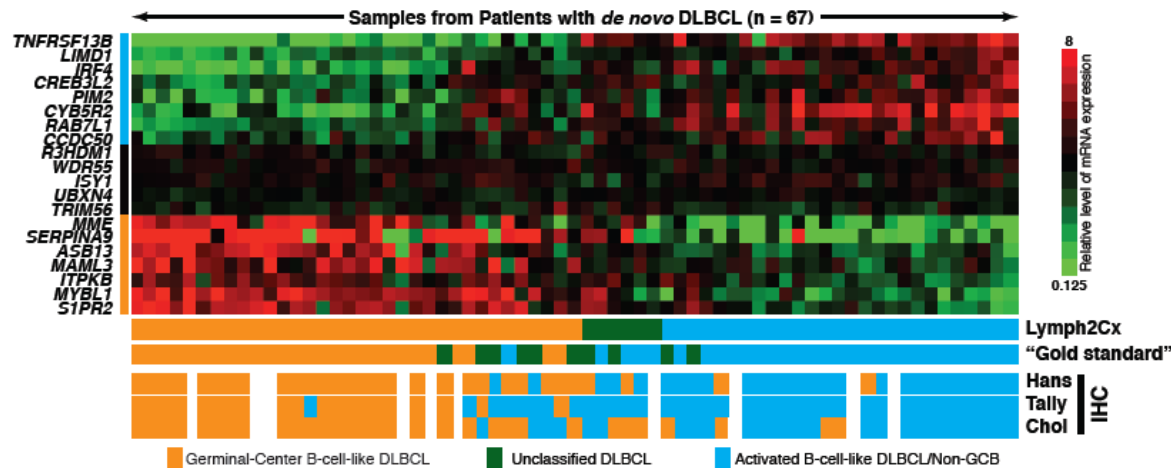
- GCB-DLBCL:
PI3K/Akt/mTOR signaling



ABC and GCB DLBCL have significantly different survival rates following R-CHOP

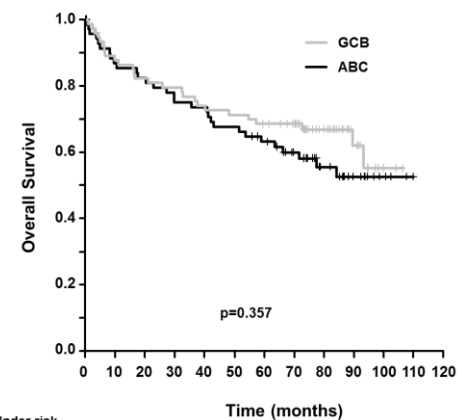
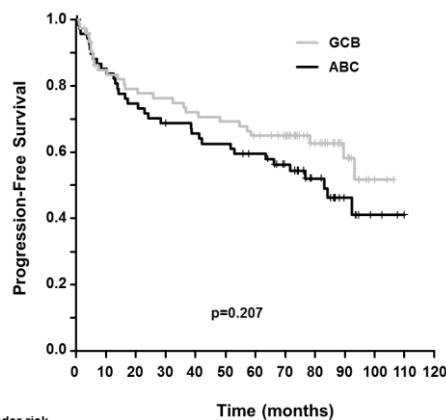
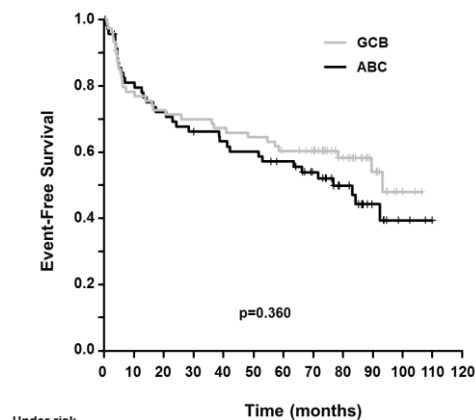


Nanostring nCounter-based GE profiling LLMPP Lymph2Cx

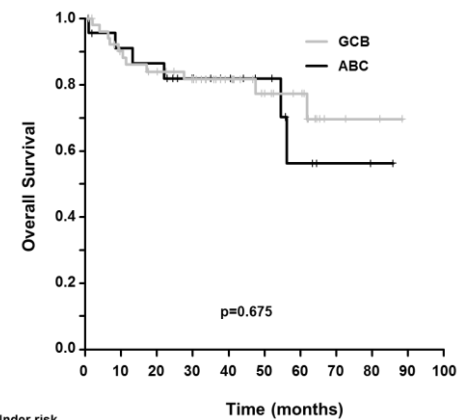
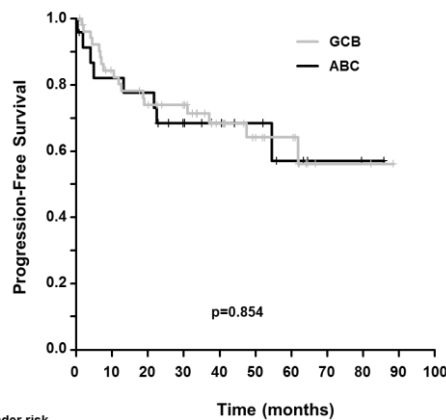
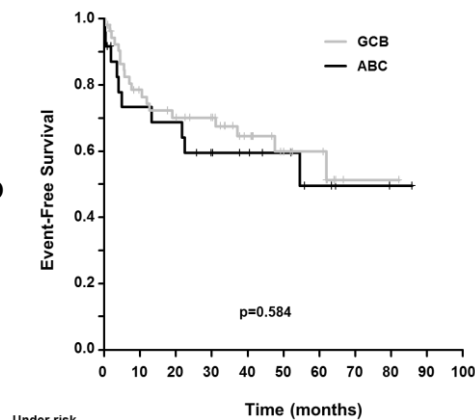


Lymph2Cx-based COO Classification and Survival Analysis

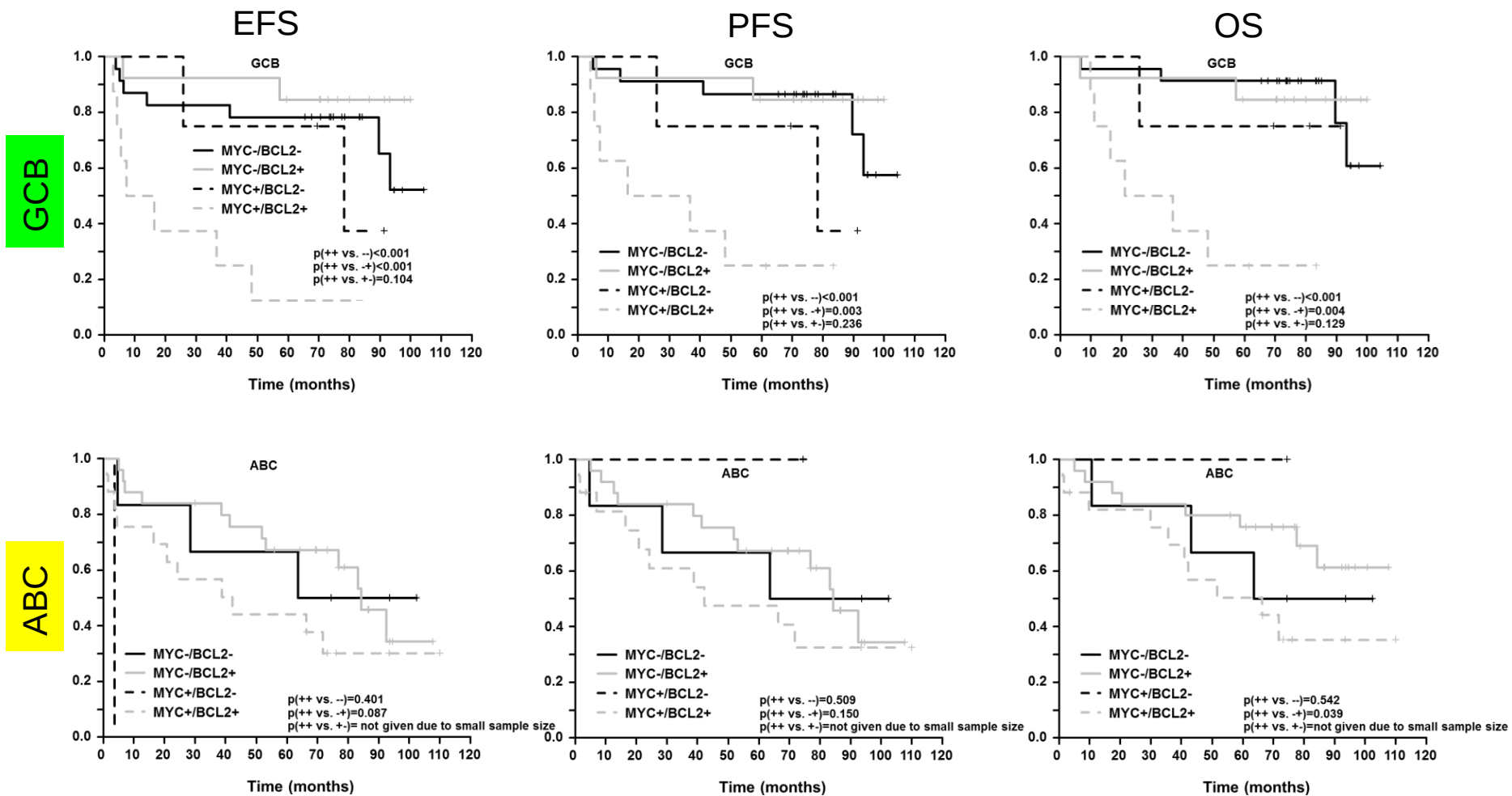
RICOVER-60
(R-CHOP-14)



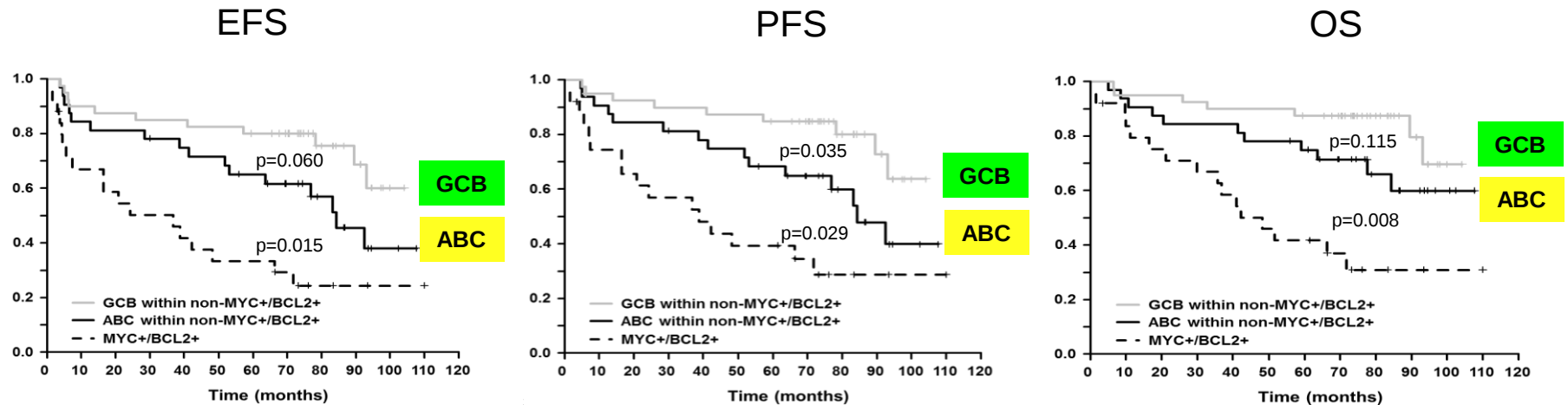
R-MegaCHOEP



Survival Analysis of COO-Classified MYC/BCL2 Dual Expressers



MYC/BCL2 Dual Expressers versus Non-Expressers

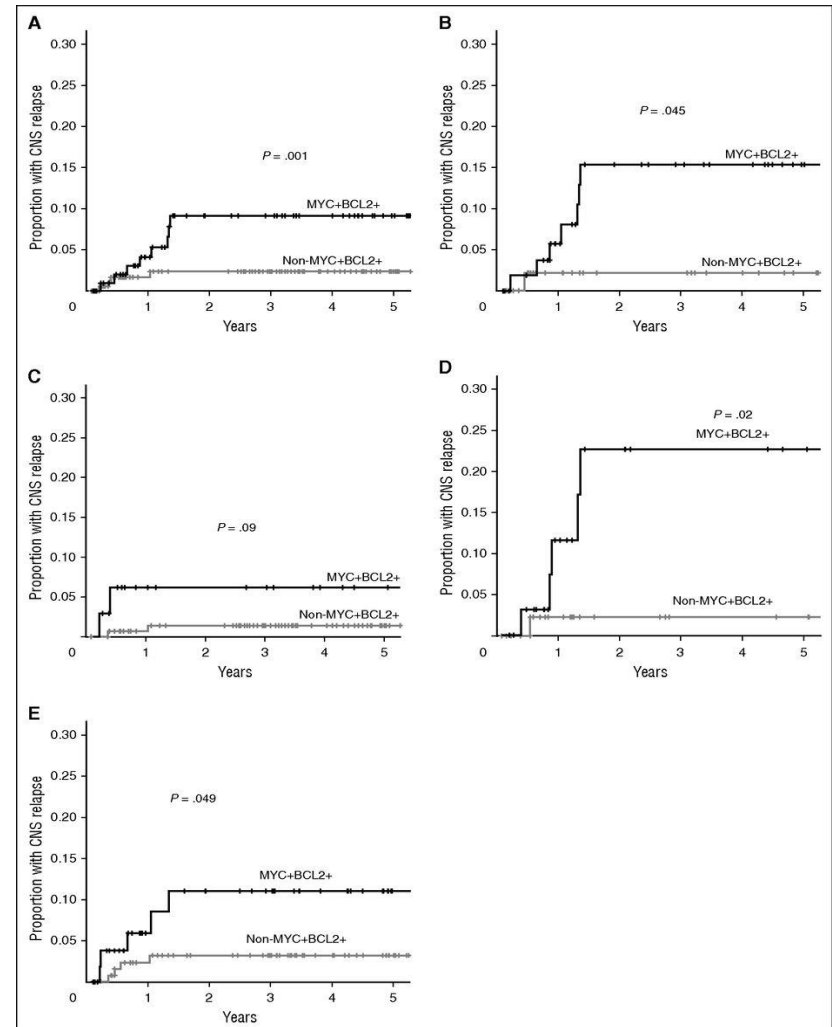


MYC/BCL2 DE status is a risk factor independent of COO classification

→ Corroborating data from Scott and colleagues 2015

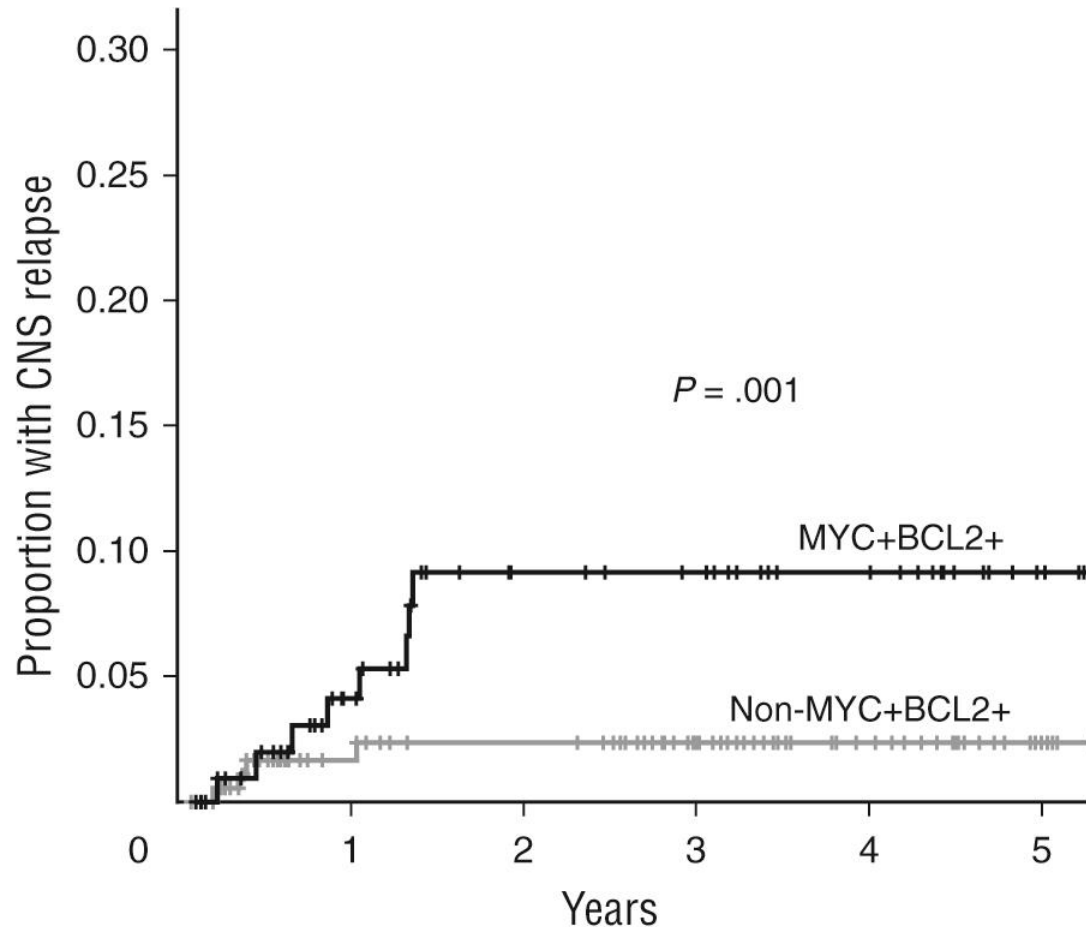
Risk of CNS relapse in dual expresser MYC+BCL2+.

- A. All DLBCL
- B. ABC-subtype
- C. GCB-subtype
- D. CNS-IPI high
- E. CNS-IPI

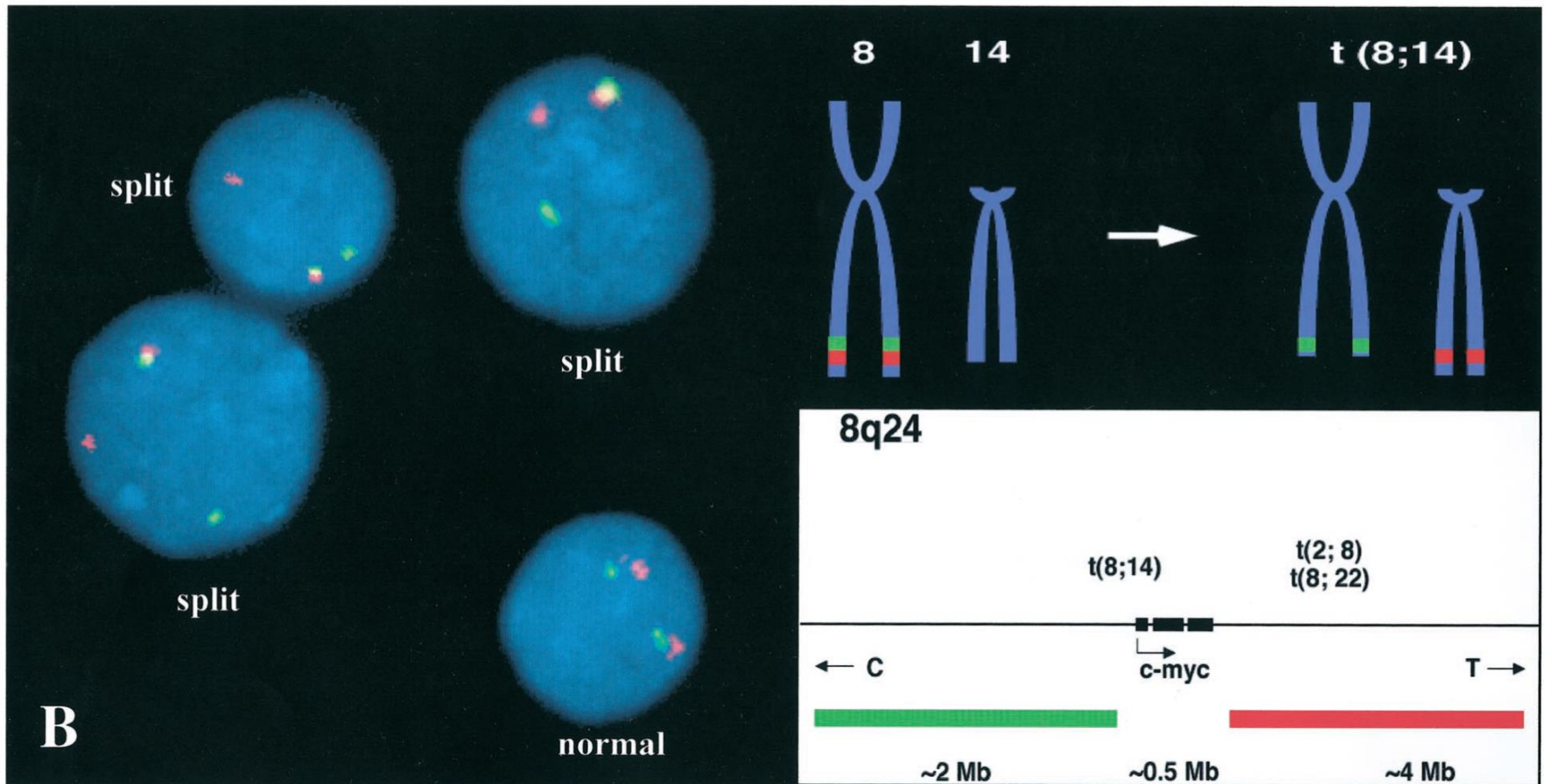


Savage et al. Blood 2016;127:2182-2188

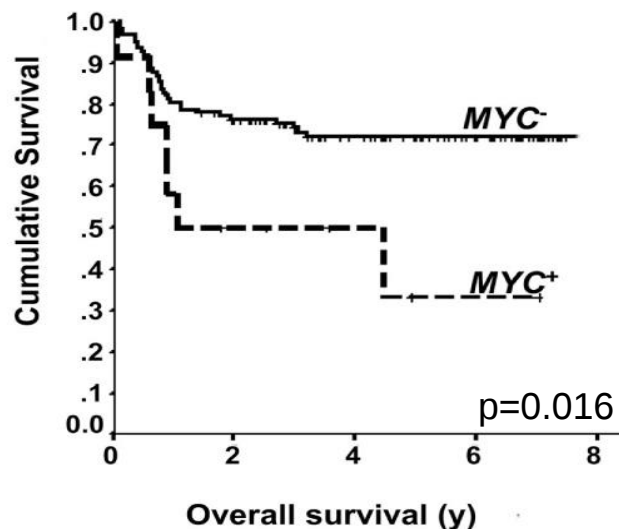
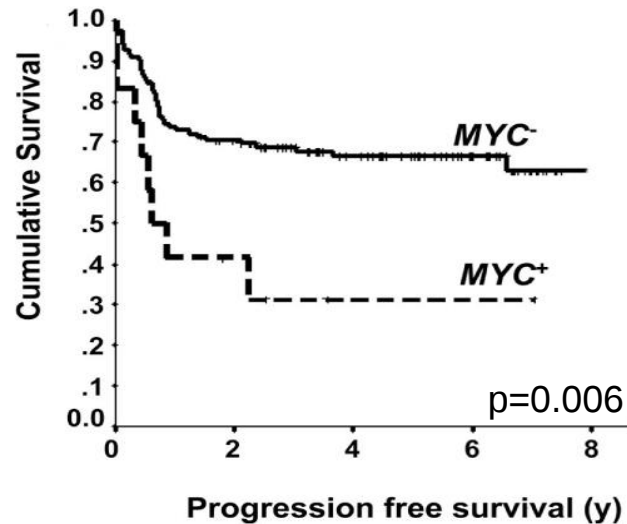
Double expressers are characterized by increased CNS relapse risk



8–10% of all DLBCLs are characterized by *MYC* translocations

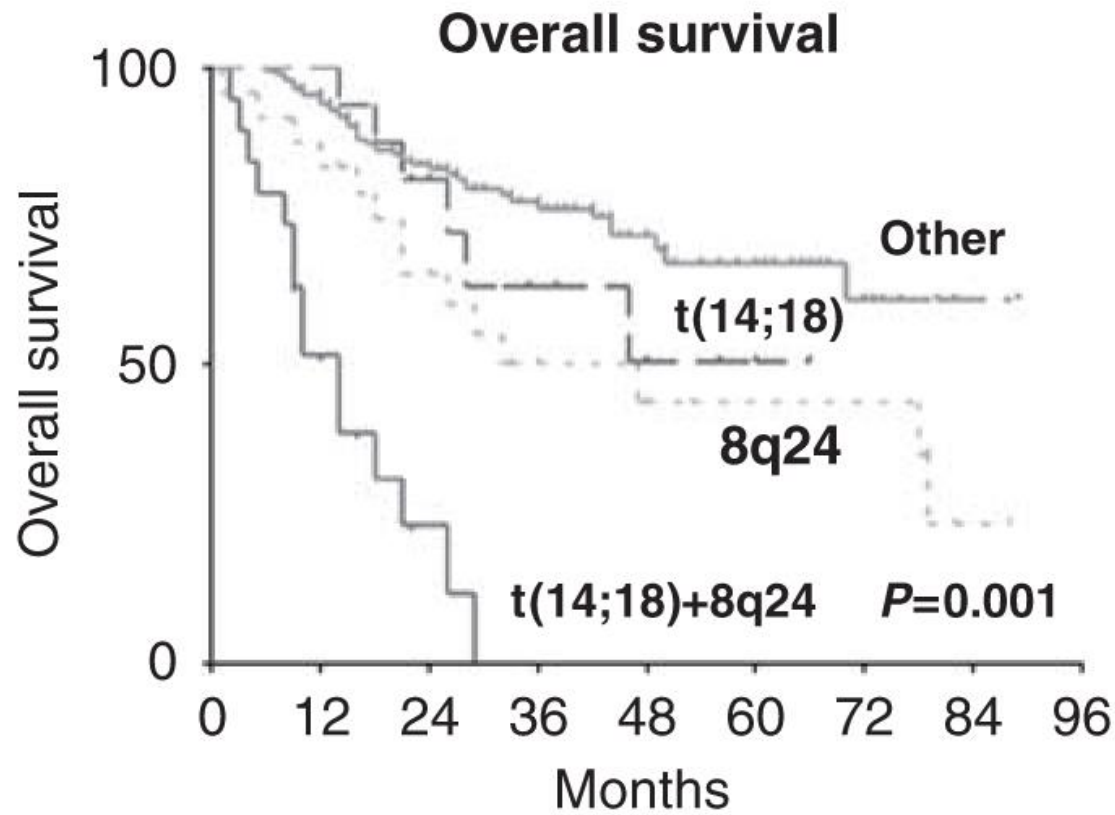


DLBCLs with *MYC* rearrangement are characterized by inferior survival after R-CHOP



- Incidence 8.8%
- Characterized by high proliferation
- Increased risk of CNS relapse
- 3/12 cases with *MYC* rearrangement had concurrent t(14;18)

DLBCLs with *MYC* and *BCL2* translocations are characterized by inferior survival



GCB?

ABC?

Double-hit?
lymphomas?

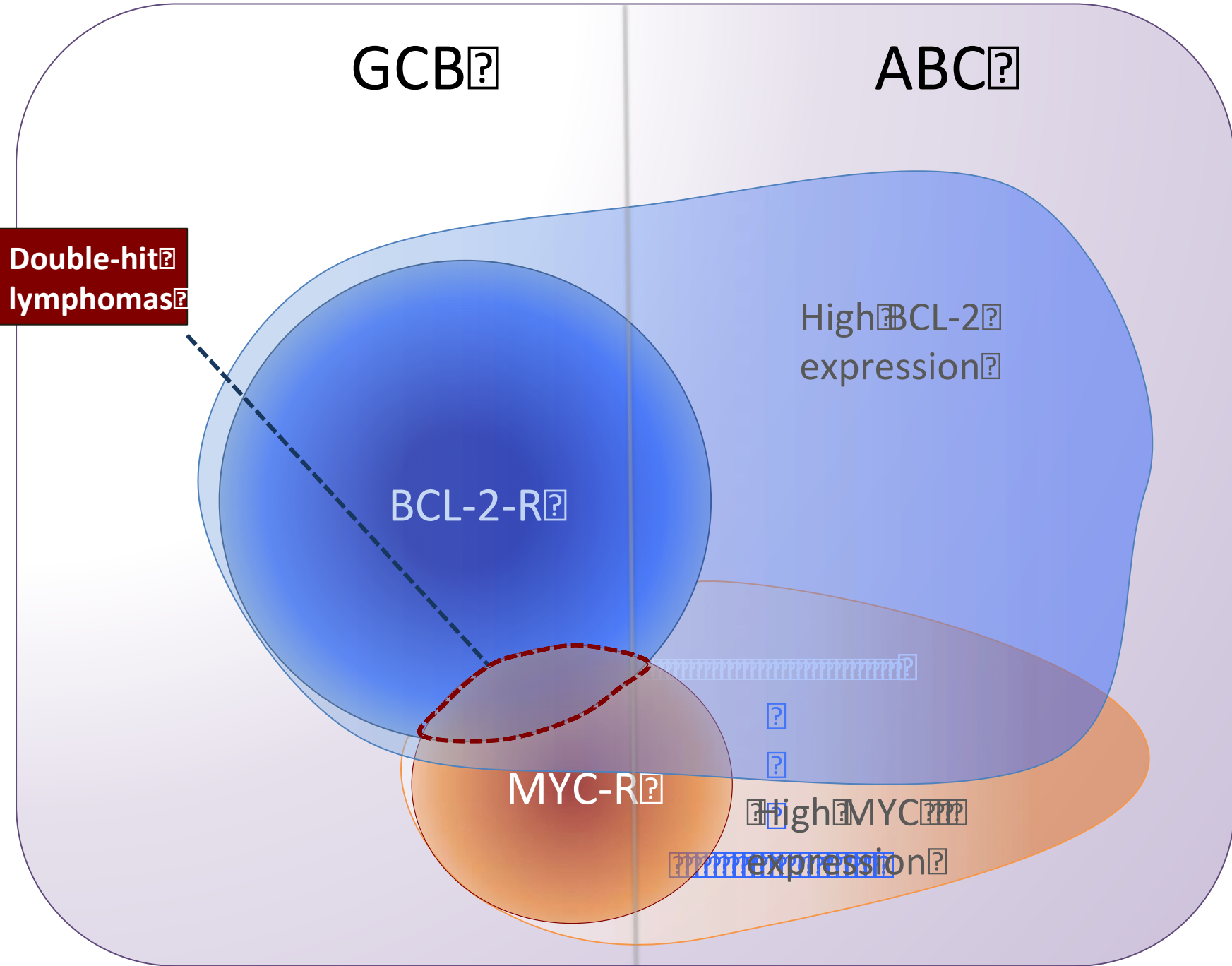
BCL-2-R?

High BCL-2?
expression?

MYC-R?

High MYC?

expression?



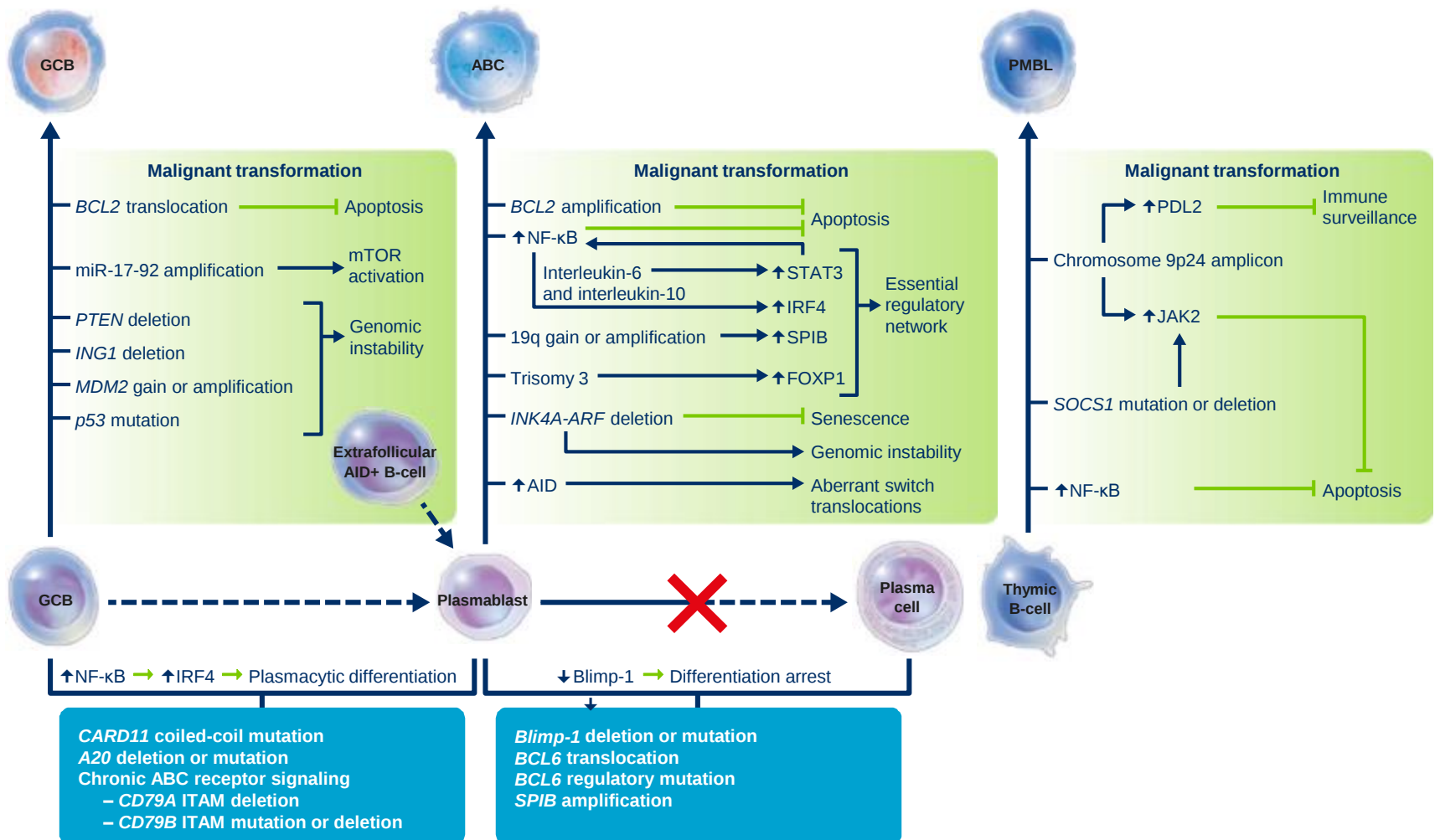
Prognostic implications of
biological markers
partly controversial

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BUT

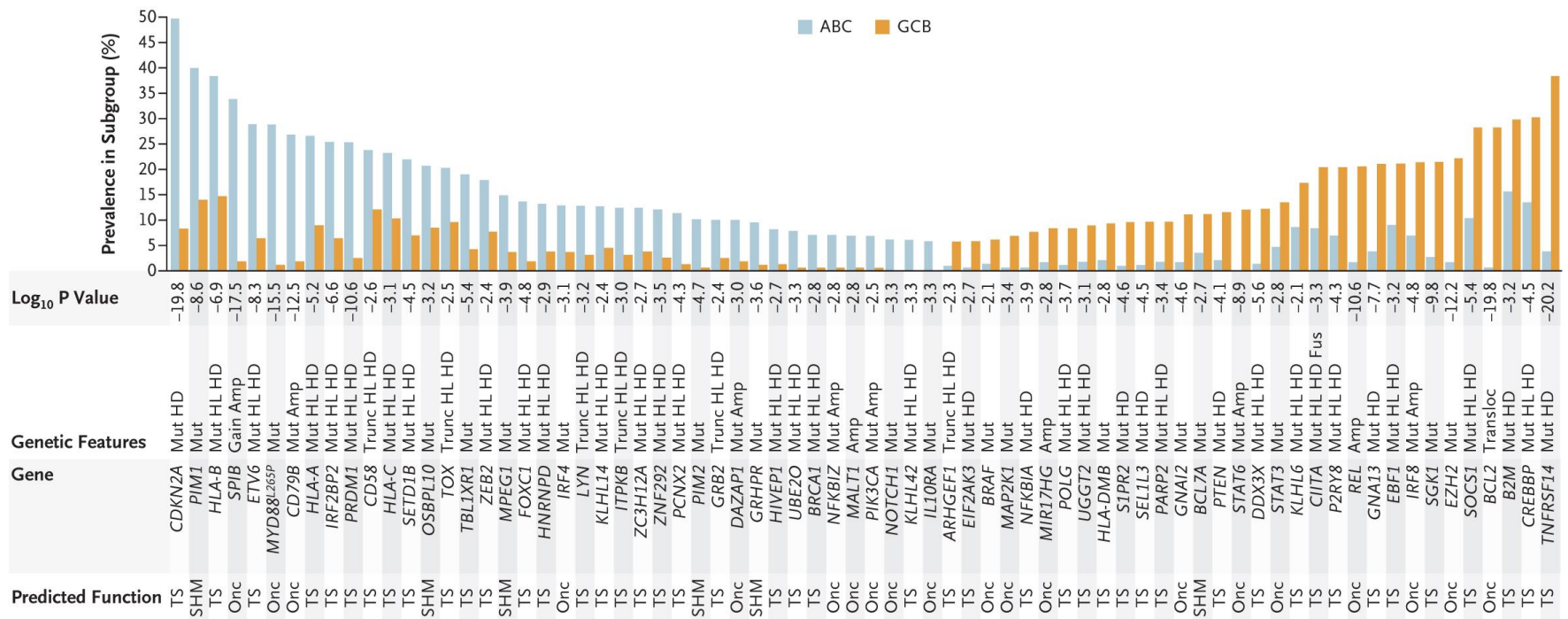
Molecular DLBCL subtypes
are addicted to different
oncogenic pathways

Molecular DLBCL subtypes are addicted to different oncogenic pathways

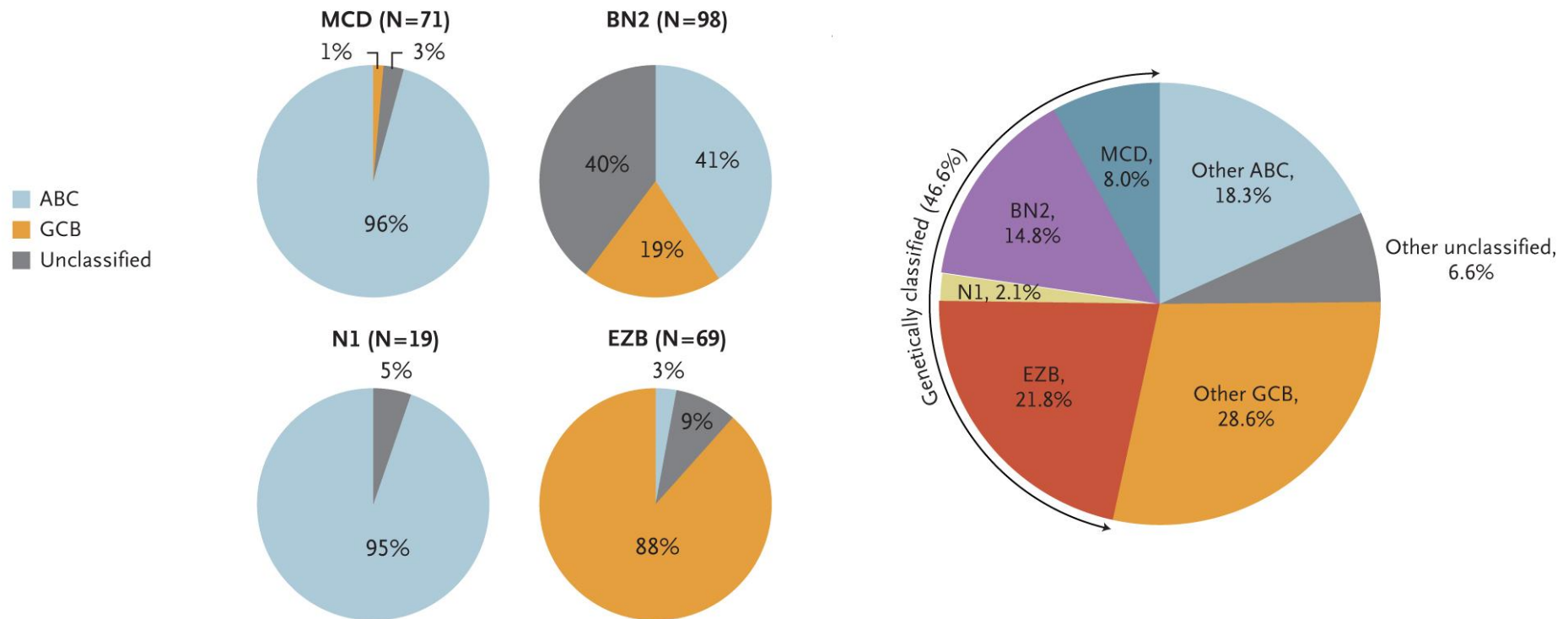


Future developments

Novel subtypes have been identified

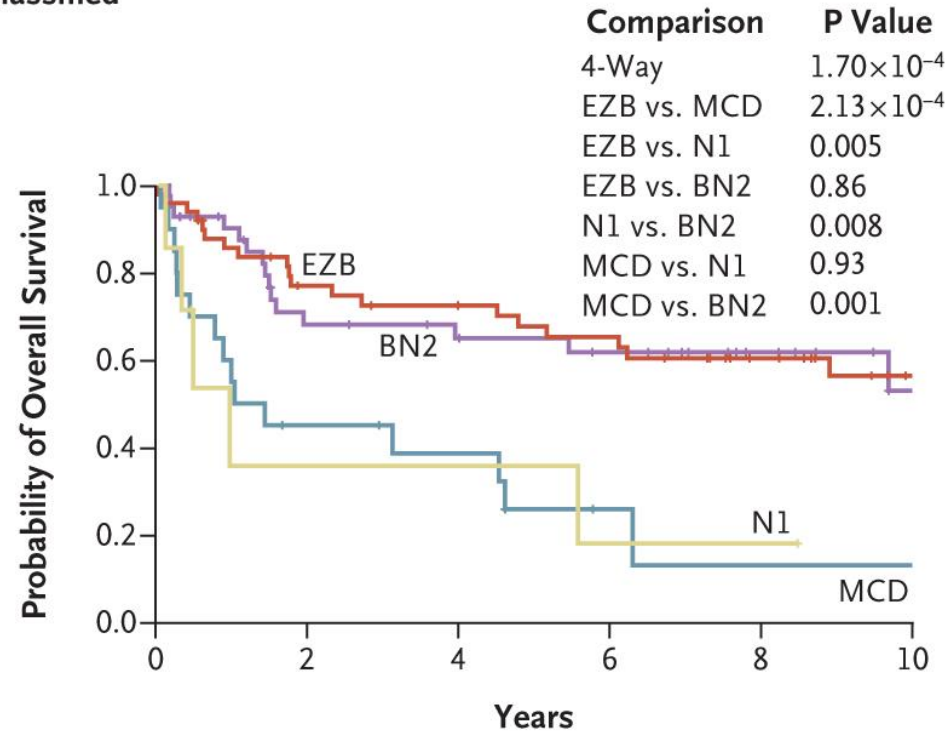


Novel subtypes have been identified



Novel subtypes have been identified

Overall Survival among Patients Whose Tumors Were Genetically Classified



No. at Risk

MCD	20	8	6	2	1	1
BN2	42	24	20	18	11	5
N1	7	2	2	1	1	0
EZB	50	34	30	27	19	11

Therapeutic implications

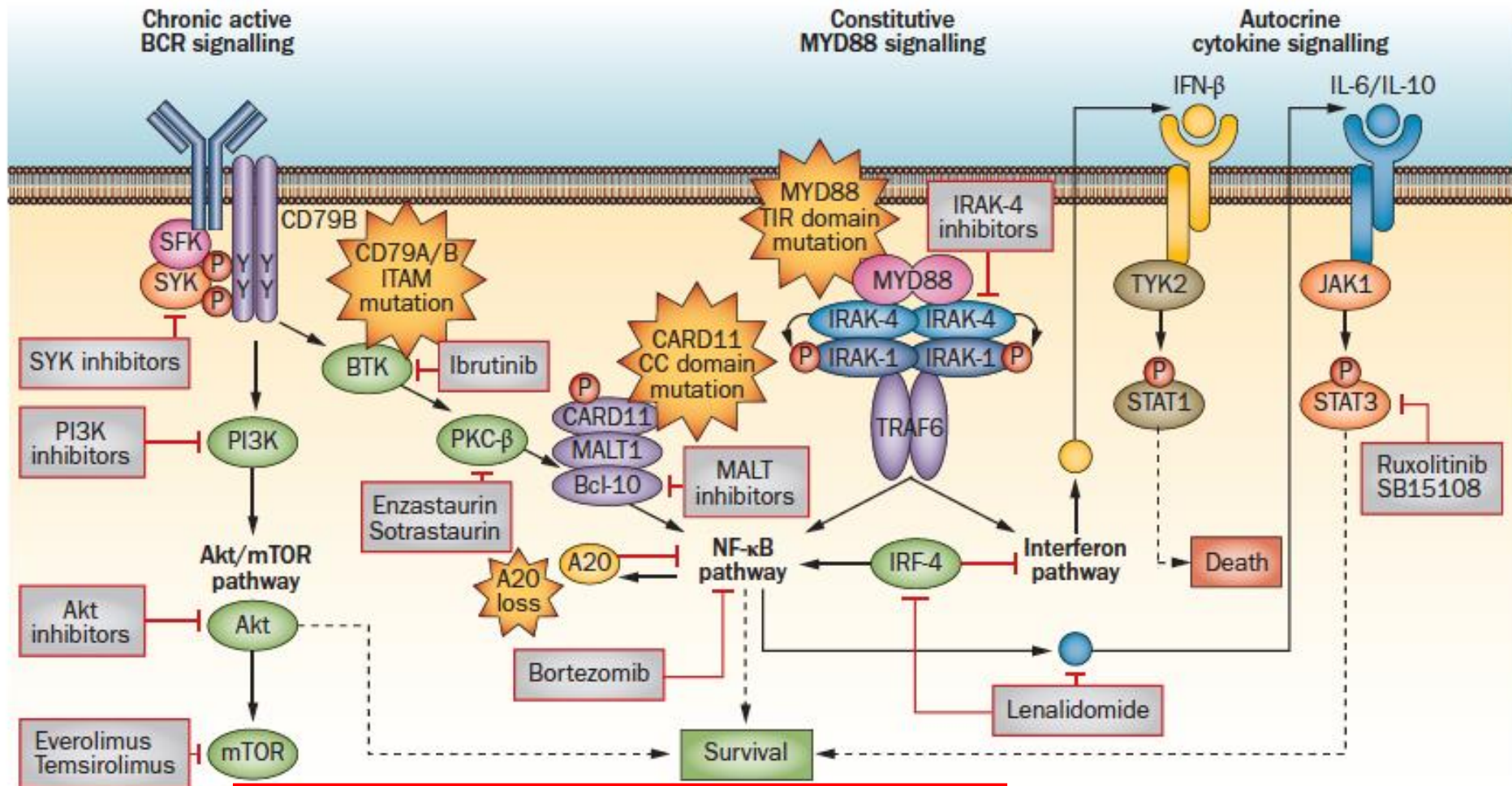
Molecular Characterisation: Implications for targeted therapy

20% of ABC DLBCL

Chronic active BCR signalling with intact NFkB genes *CD79B/A*

30% of ABC DLBCL

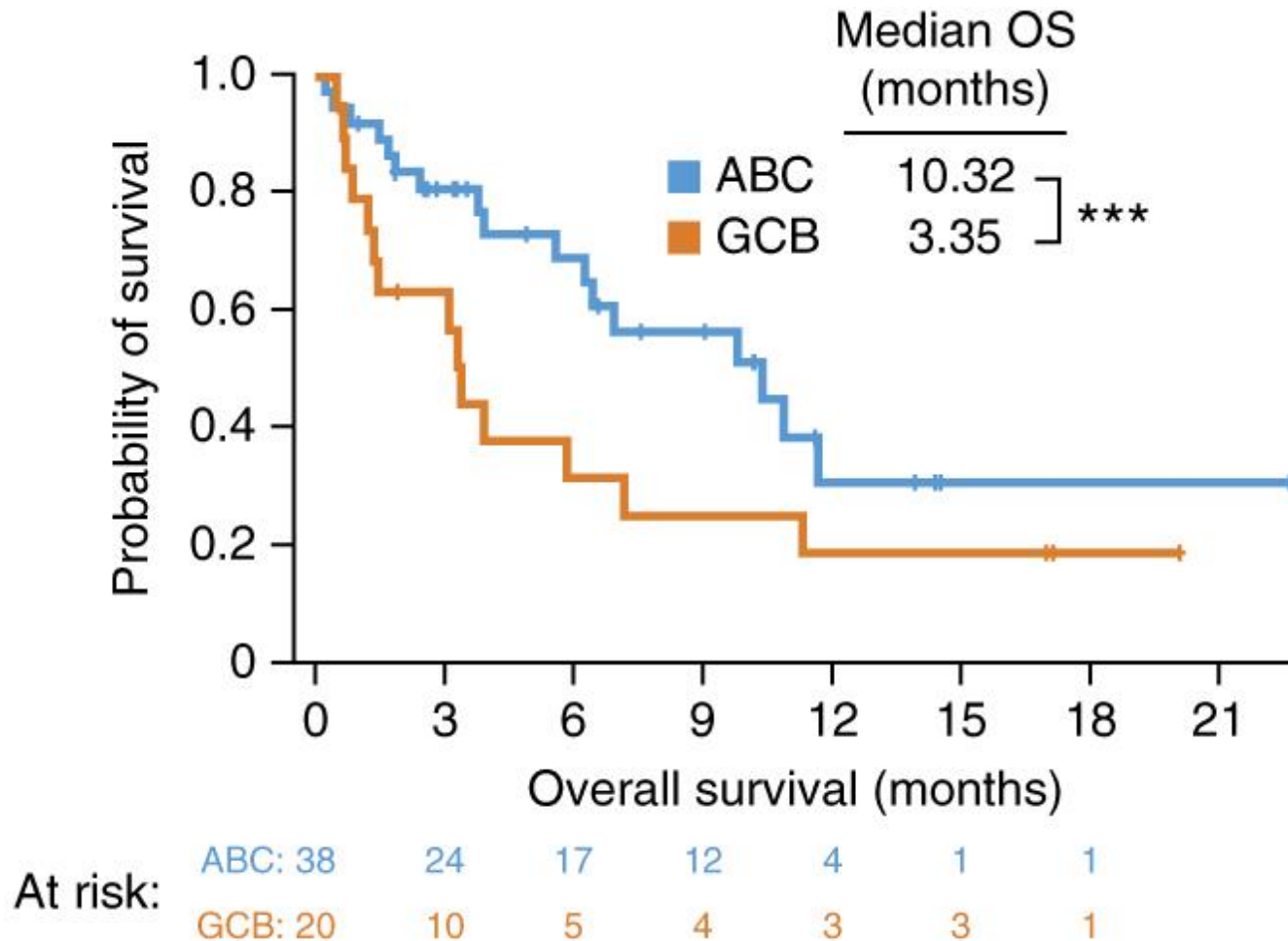
Activating mutations of *MyD88*
Association with *CD79* mutations



50% of ABC DLBCL

Mutations in NFkB regulators : *A20*, *CARD11*, others

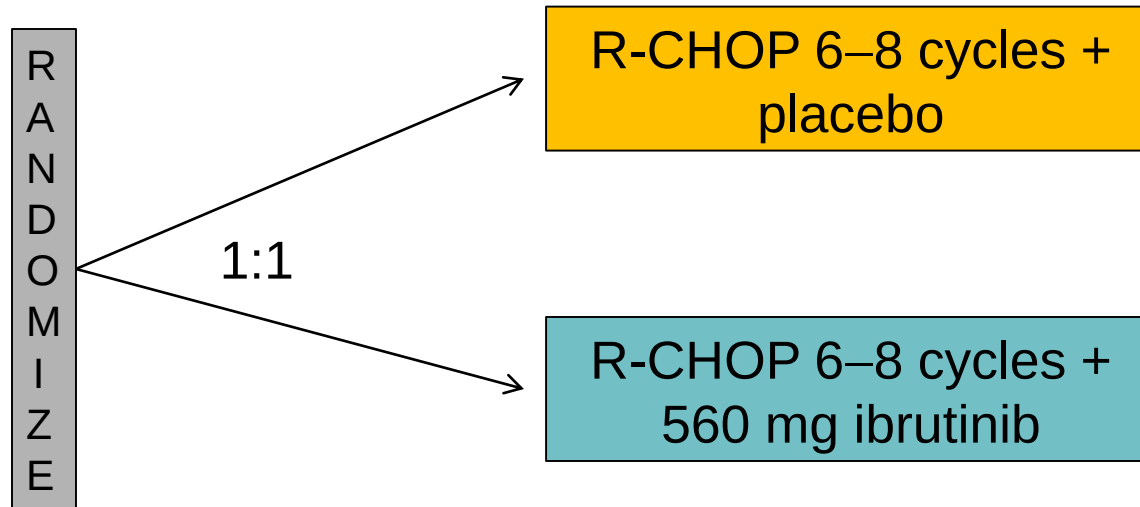
Ibrutinib is preferentially active in ABC DLBCL



Ibrutinib + R-CHOP in newly diagnosed non-GCB DLBCL

PHOENIX (NCT01855750): study design

Phase III trial



Stratified by R-IPI 1-2 vs 3-5, region
and number of prespecified cycles

Primary endpoint: EFS

Secondary endpoint: PFS, OS, CR, time to worsening (LYM subscale), pharmacokinetics, safety

Exploratory endpoint: quality of life, biomarkers related to resistance

Key Inclusion Criteria

Newly diagnosed histologically confirmed non-GCB DLBCL

Stage II-IV disease

At least one measurable site of disease

Age ≥ 18 years

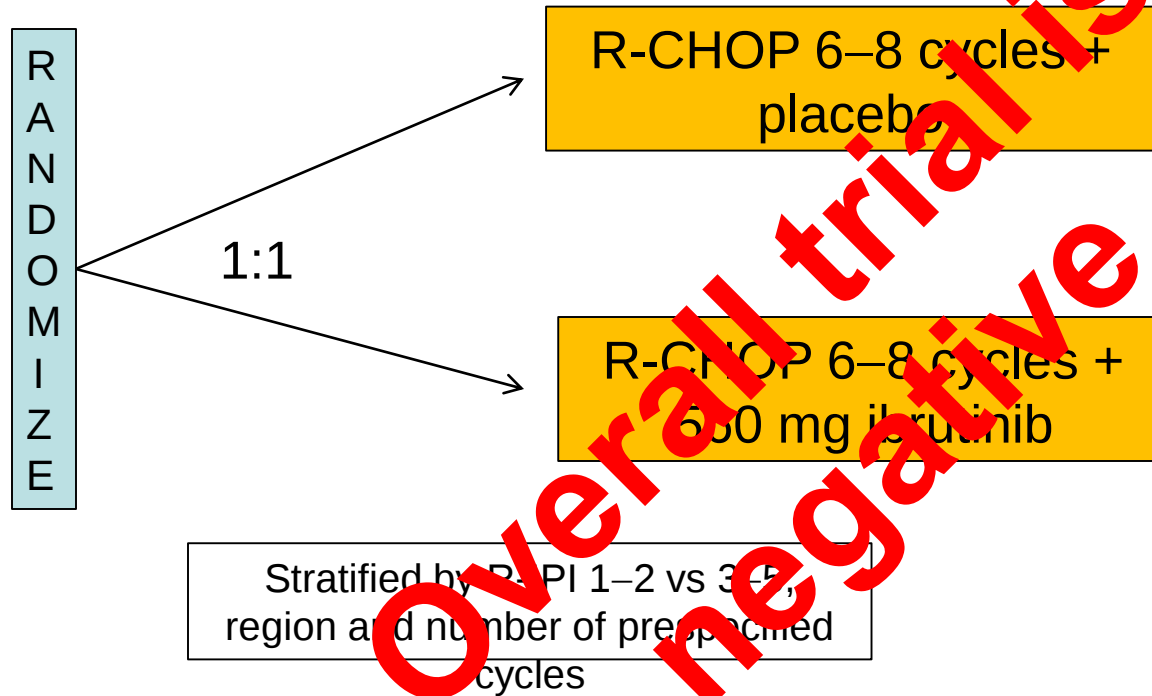
R-IPI ≥ 1

ECOG PS 0-2

Ibrutinib + R-CHOP in newly diagnosed non-GCB DLBCL

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R-CHOEP-brut

Indication

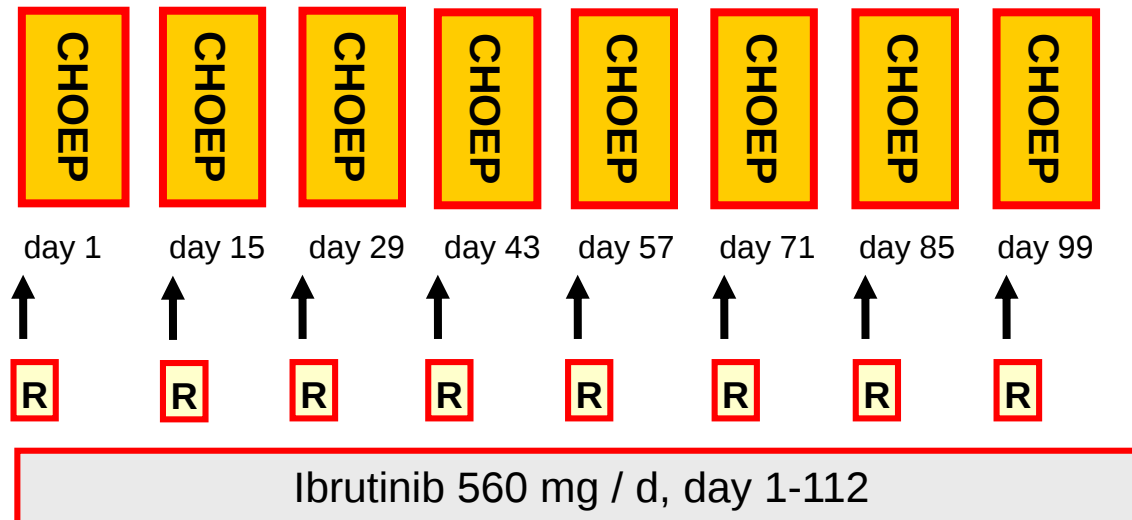
First-line therapy of diffuse large B-cell lymphoma
Younger patients (18-60 years)
Age-adjusted International Prognostic Index 2-3

Design

Prospective, multicenter, phase II study

No. of patients

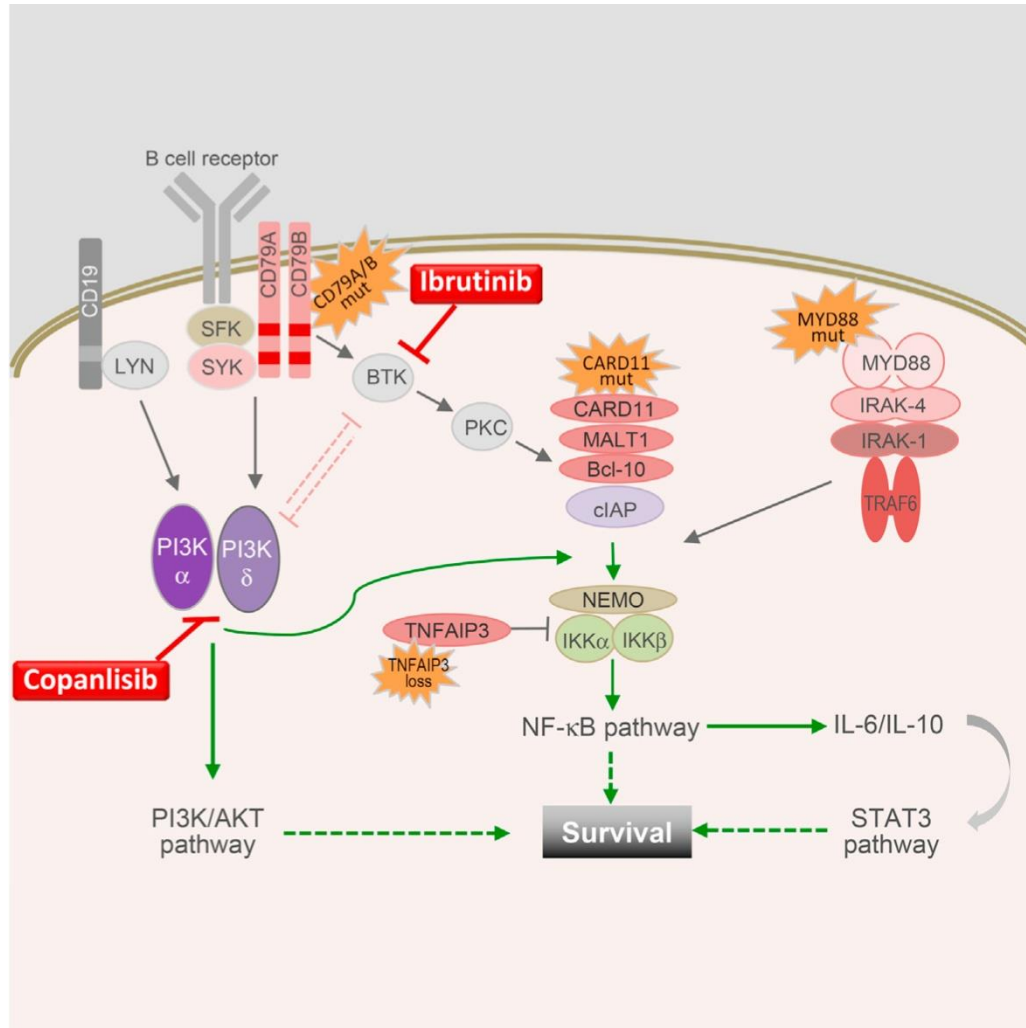
75



Patients with bulky disease/extranodal involvement:
radiotherapy (dose 39.6 Gy)

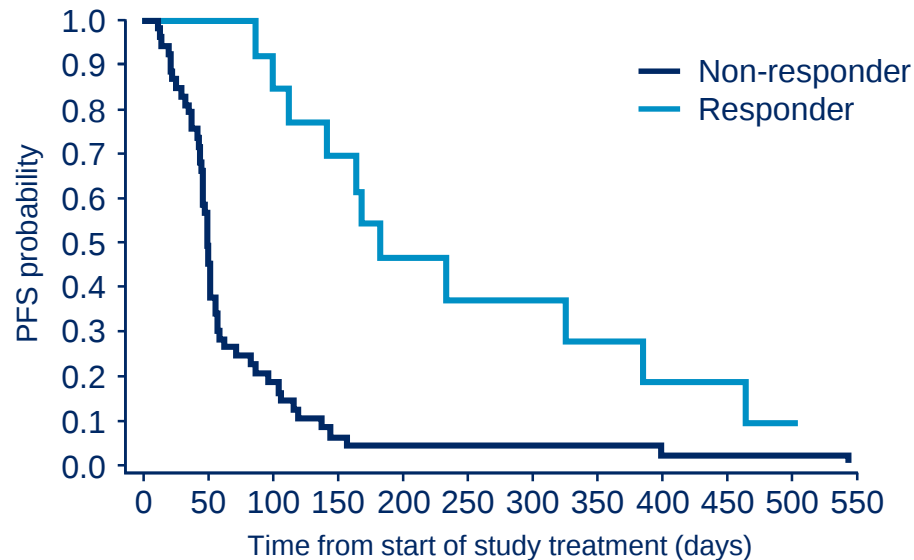
Molecular DLBCL subtypes
are addicted to different
components of the
PI3K/AKT pathway

Copanlisib is active in preclinical models of ABC DLBCL



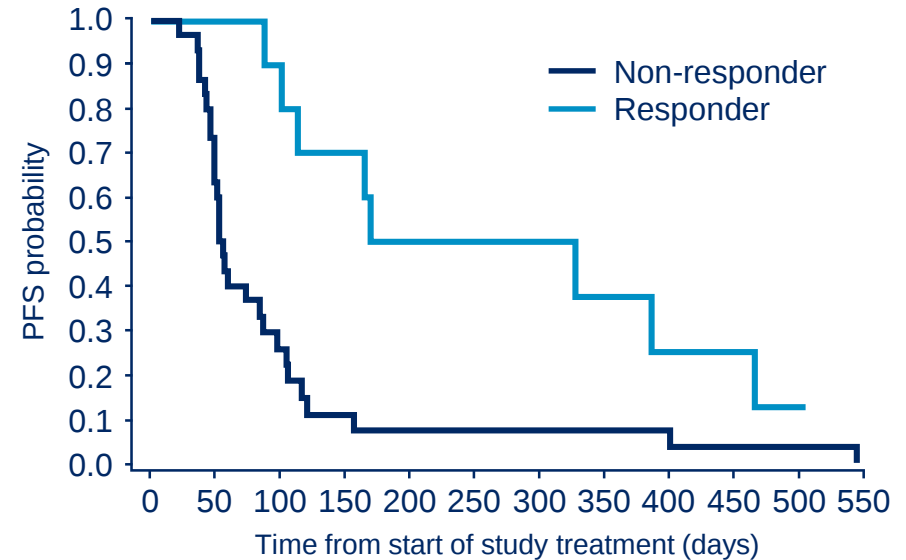
COPANLISIB: Progression-free survival for patients with objective responses

FAS



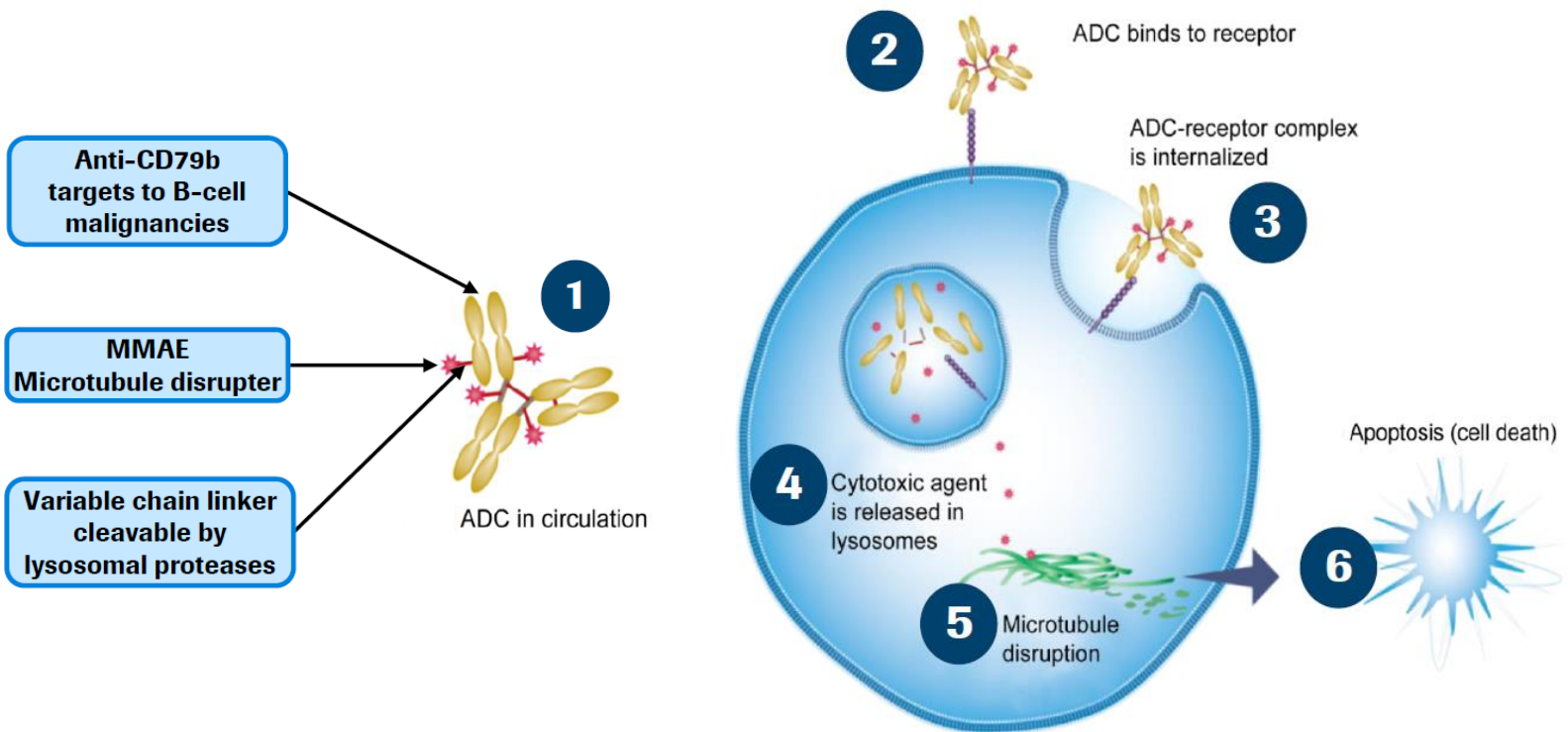
	Overall FAS (n=67)	Responder (n=13)	Non-responder (n=54)
Median PFS, days (95% CI)	54 (50-84)	183 (113-385)	50 (46-56)

PPS



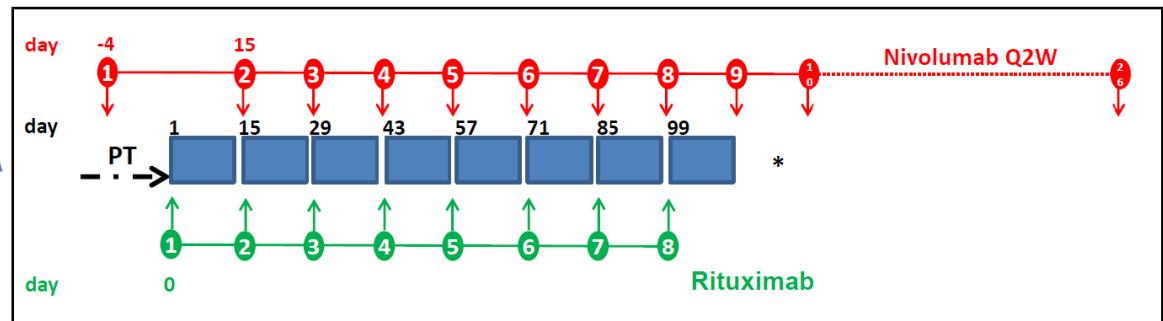
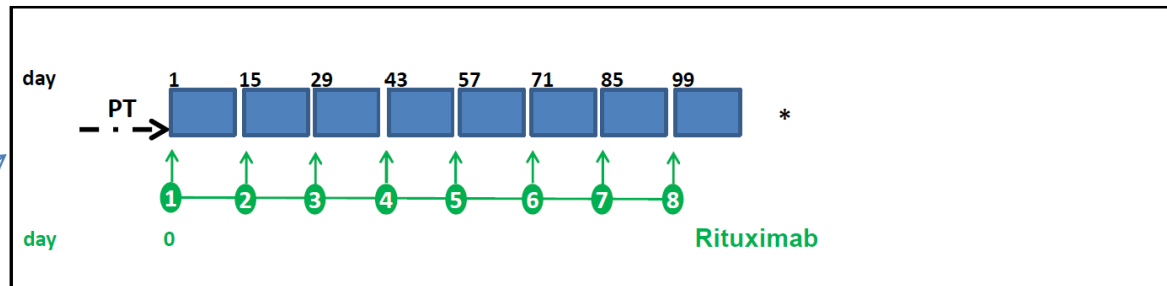
	Overall PPS (n=40)	Responder (n=10)	Non-responder (n=30)
Median PFS, days (95% CI)	84 (52-106)	248 (88-465)	54 (50-84)

Polatuzumab Vedotin: anti-CD79b ADC (Antibody drug conjugate)



Design

R



Gemcitabine 1000 mg/m², Oxaliplatin 100 mg/m², recycle on d15