

# Eppur si muove...

## La terapia nel MONDO LINFOMI

***Il razionale biologico delle  
combinazioni nei linfomi  
non Hodgkin***

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Staminali  
AO San Camillo  
Roma*



ROMA, 26 MAGGIO 2022

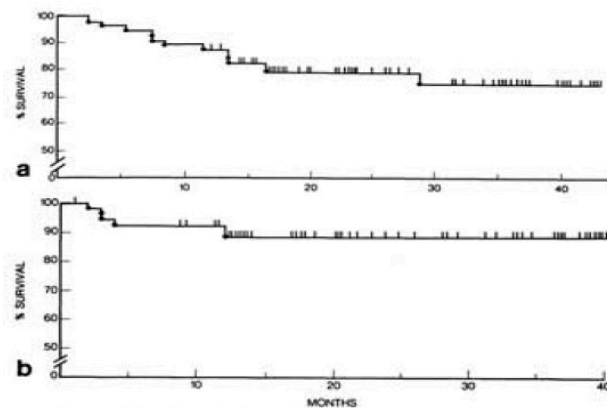
## Disclosures, Luigi Rigacci

Company name	Advisory board	Educational activities/Lecture fees
Astra Zeneca		X
Gilead-Sciences	X	X
Incyte		X
Janssen-Cilag	X	X
Novartis	X	
Roche	X	X
Gentili		X
Takeda	X	X
Abbvie	X	X
Sandoz		X

Once upon a time.....

# MACOP-B

*alternating weekly myelotoxic drugs with  
non-myelotoxic drugs!*

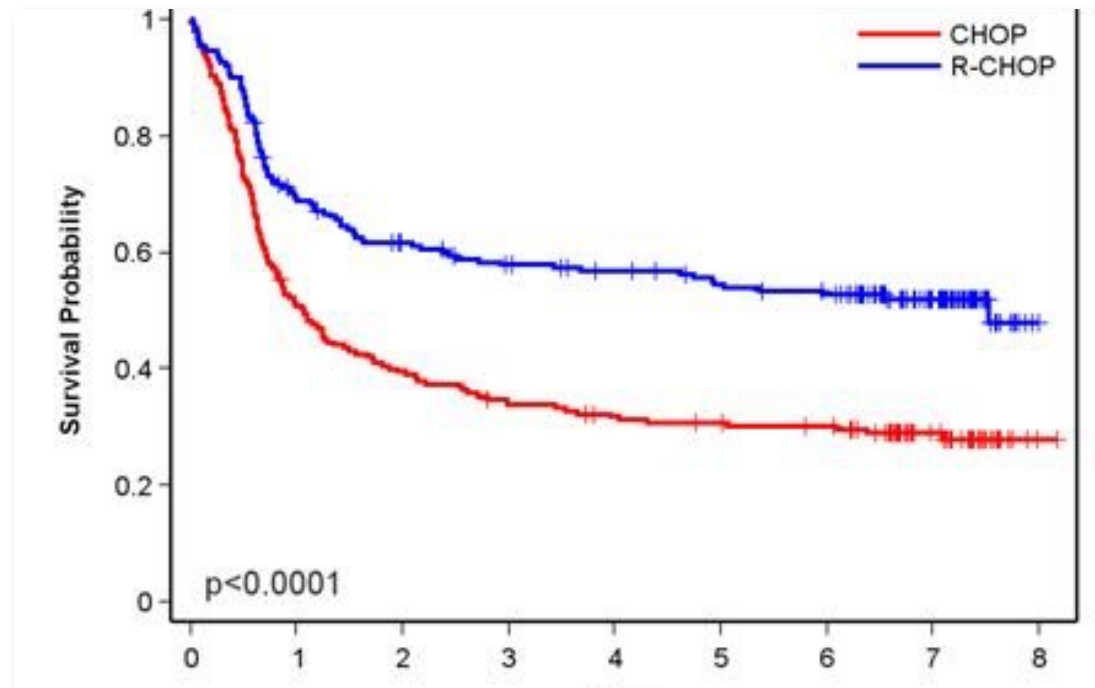


**MACOP-B Chemotherapy for the Treatment of Diffuse Large-Cell Lymphoma**

PAUL KLIMO, M.D.; and JOSEPH M. CONNORS, M.D.; Vancouver, British Columbia, Canada

**Annals of Internal Medicine. 1985;102:596-602.**

# Ours are coming!



Coiffier, NEJM 2002

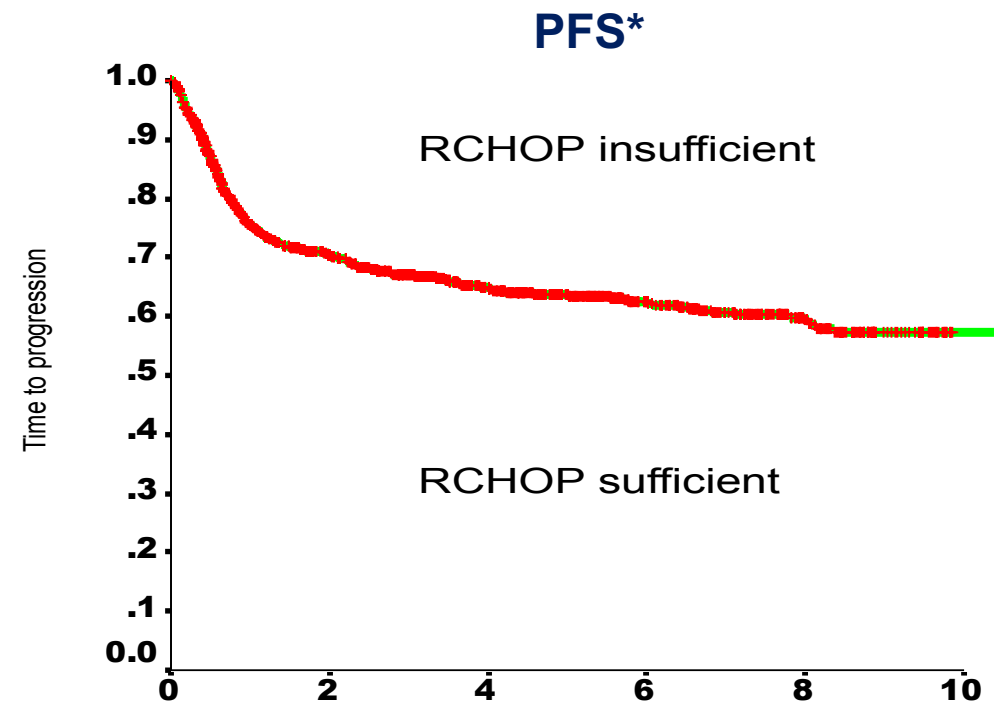
immunotherapy added a great benefit

## Heterogeneity of Outcomes in DLBCL

- Clinical factors
  - IPI (R-IPI)
- Interim PET scan
- GEP
  - ABC vs GCB
- Protein expression
  - MYC and BCL2
- Chromosomal alterations
  - MYC, BCL2, BCL6
- Deep sequencing mutation/combined expression analysis

Two broad strategies:

- Target both subgroups
  - Possibly overtreating RCHOP “sufficient group”
- Target RCHOP “insufficient” group provided
  - It can be identified
  - It can be targeted

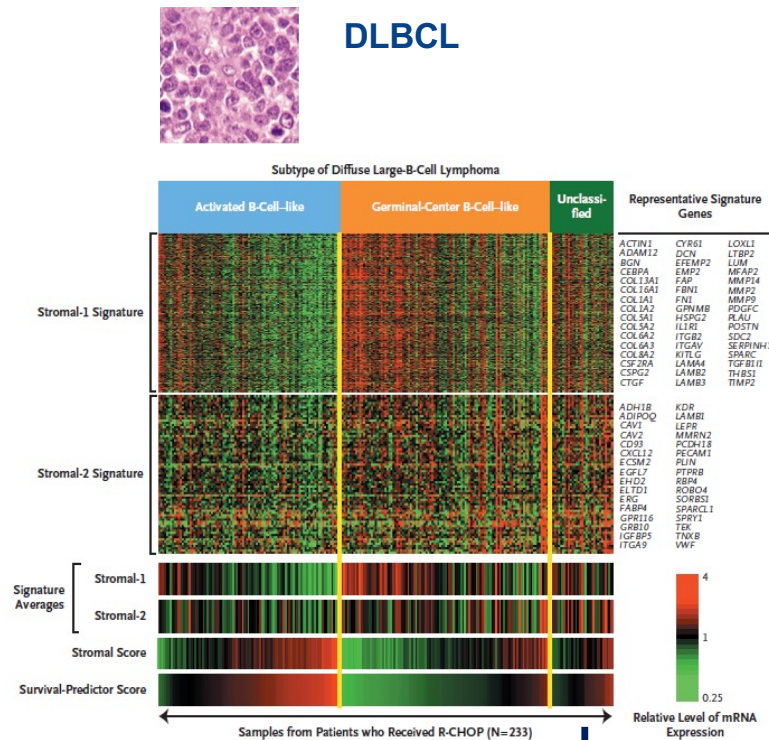


\*Patients with DLBCL treated with R-CHOP-21 at BCCA (n = 1,476).

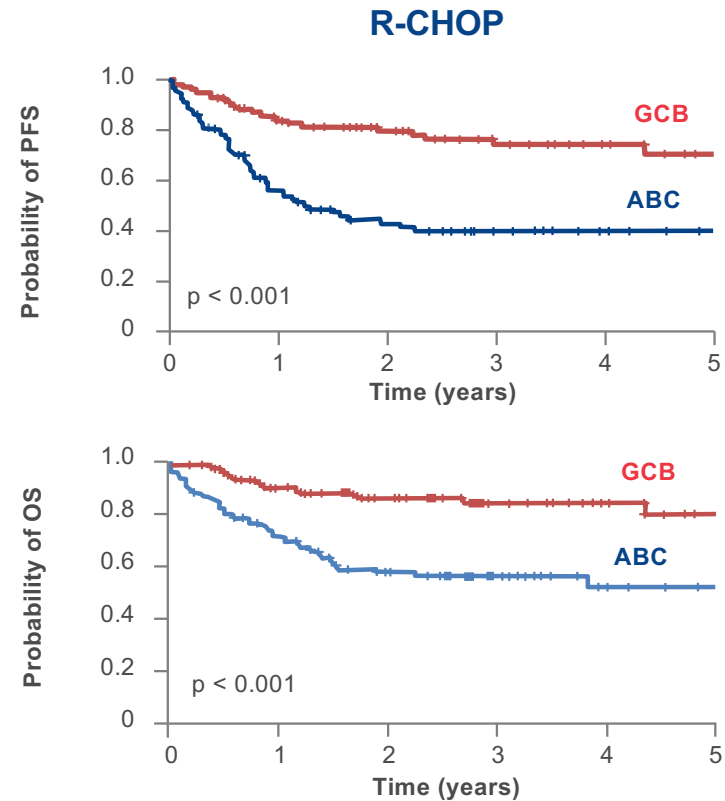
Sehn LH. ASH Education Book. 2012;1:402-9.

# Targeting “Insufficient” R-CHOP Group

Evaluation of unfavourable DLBCL subsets: Cell of Origin profile subgroups by GEP



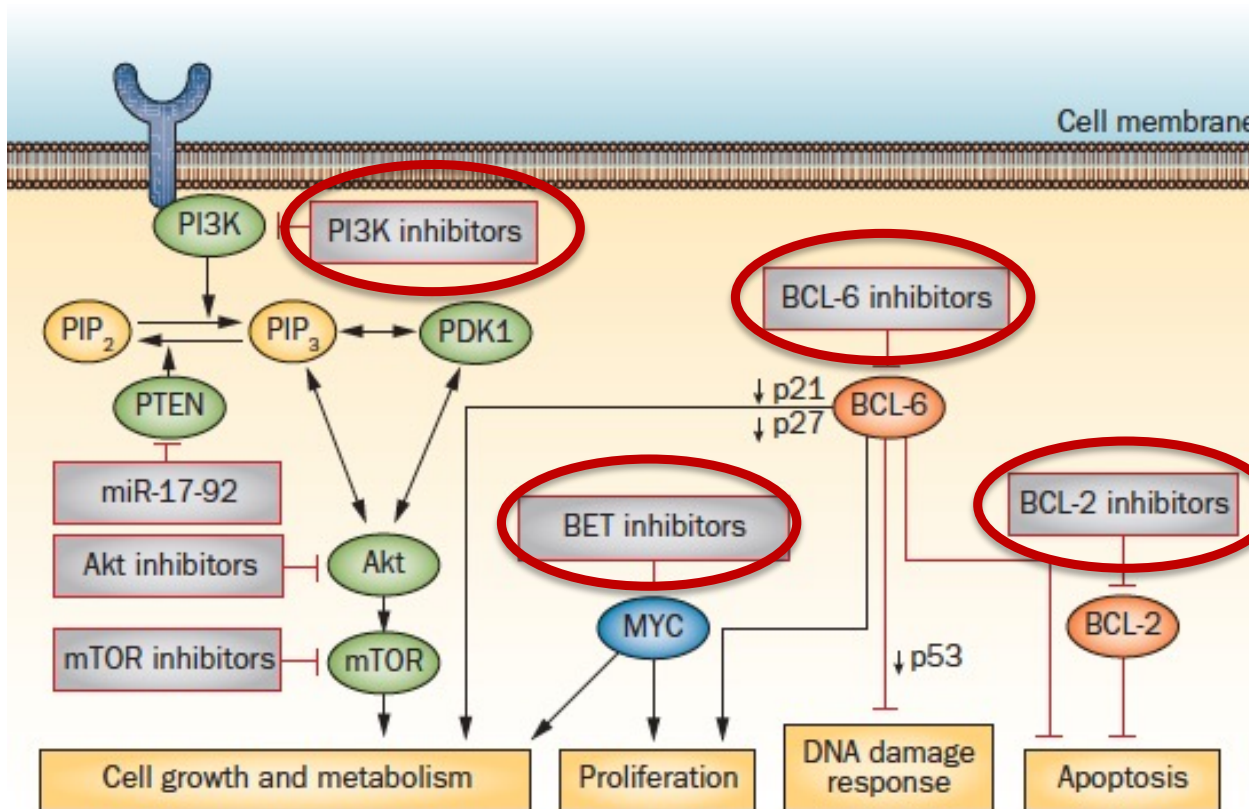
15% Unclassifiable



The GEP classification is not available in daily clinical practice

- GEP, gene expression profiling; OS, overall survival.
- Lenz G, et al. N Engl J Med. 2008;359:2313-23.

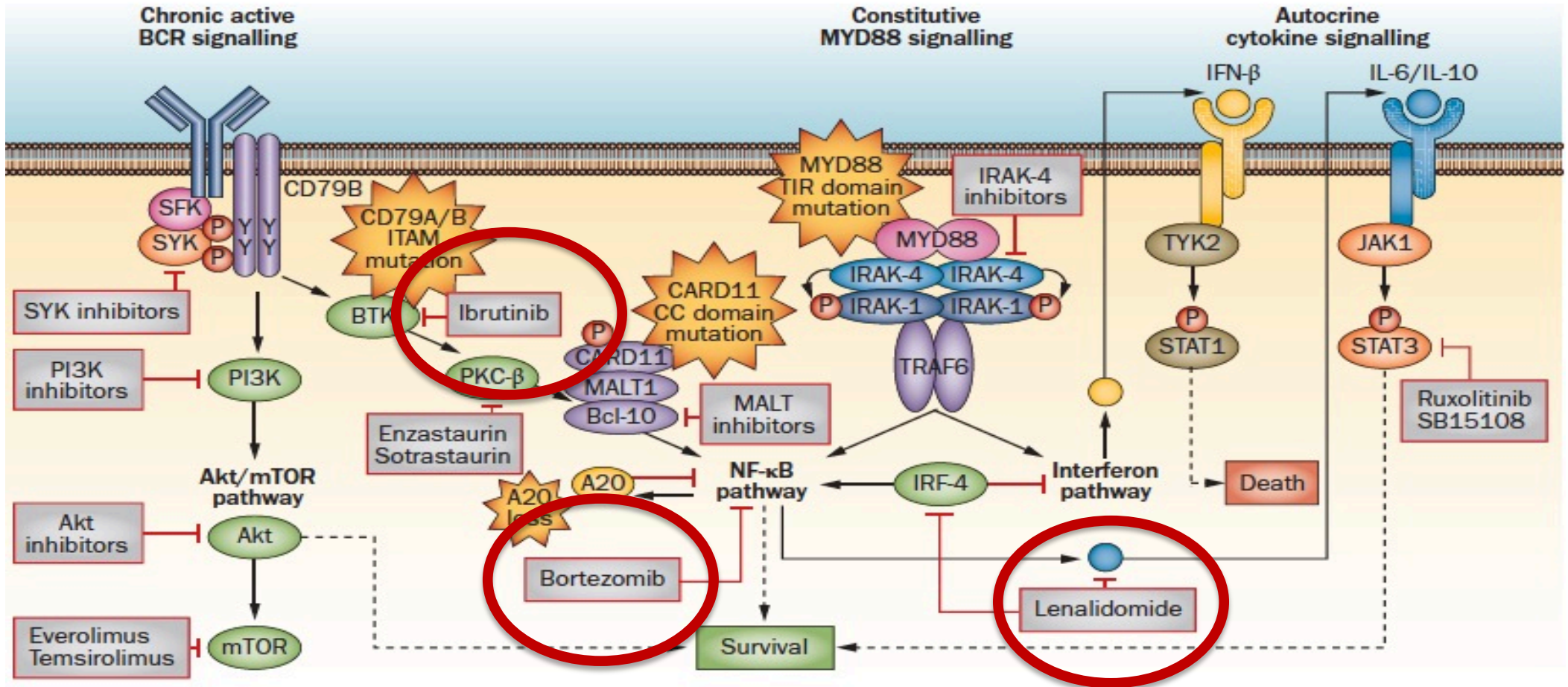
## The key signalling pathways implicated in GCB DLBCL



Roschewski M, Staudt LM, Wilson WH, Nat. Rev. Clin. Oncol. 2013.

- ✓ Loss of PTEN (Phosphatase and tensin homologue) expression in 55% of cases → activation of PI3K/Akt/mTOR signalling pathway; → **small-molecule inhibitors can be effective in GCB with decreased PTEN expression.**
- ✓ BCL-6 is frequently activated in GCB DLBCL; BCL6 deregulation results in enhanced tumour proliferation via decreased expression of the cell-cycle checkpoint proteins p21 and p27, impaired DNA damage response through decreased p53 expression, impaired cellular metabolism and resistance to apoptosis. → **inhibitors that target key co-repressor proteins of BCL-6.** In normal B cells, BCL-6 suppresses transcription of the MYC oncogene.
- ✓ **BET bromodomain inhibitors** represent a novel strategy of epigenetic regulation of MYC-driven tumours.
- ✓ BCL2 translocations are observed in up to 35% of GCB DLBCL cases, resulting in inhibition of apoptosis → **Inhibitors of BCL-2.**

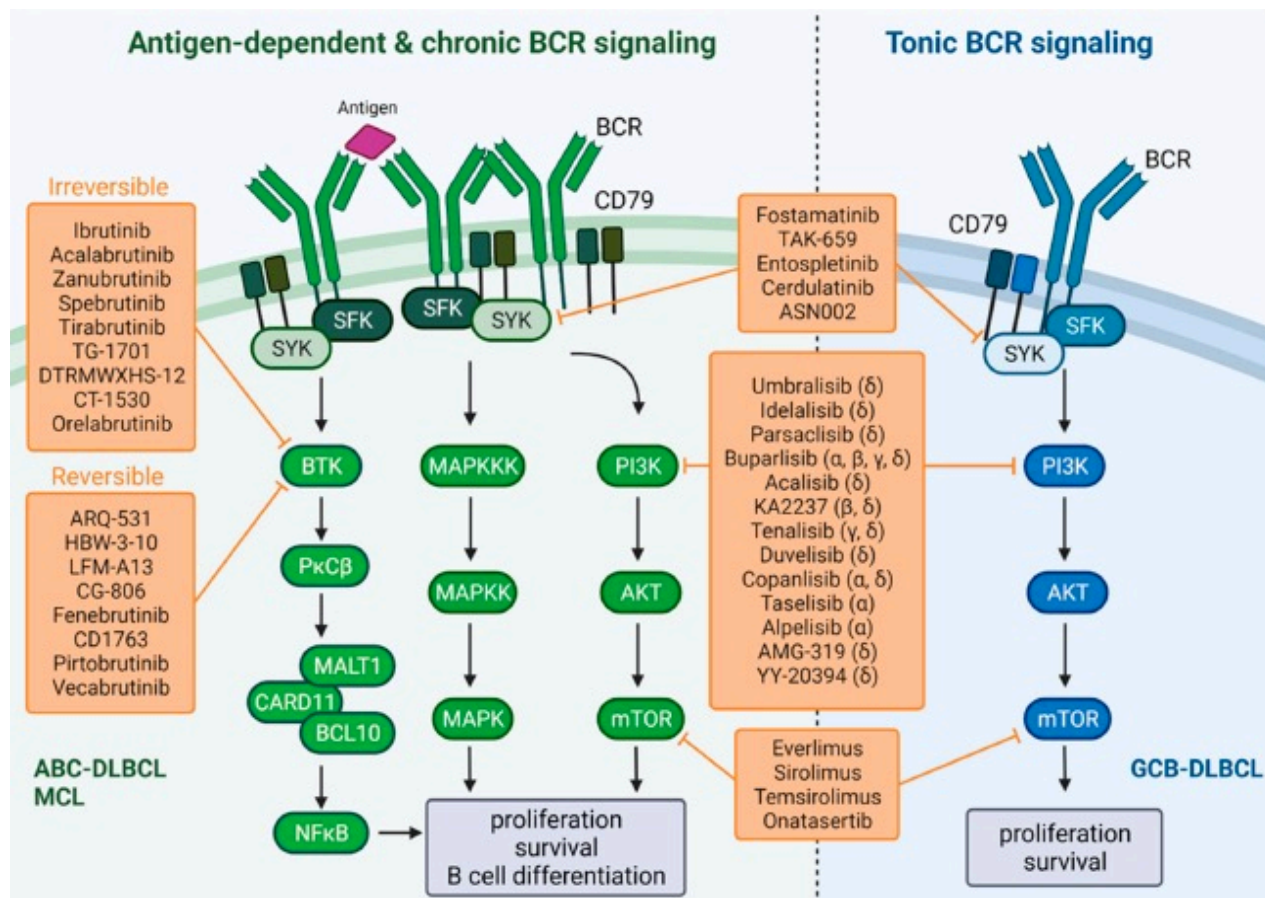
## The key signalling pathways implicated in ABC DLBCL



Roschewski M, Staudt LM, Wilson WH, Nat. Rev. Clin. Oncol. 2013.



## Regulation of BCR signaling and the therapeutic inhibition of BTK and PI3K in DLBCL



- ✓ ABC-DLBCL displays chronic active BCR signaling resulting in constitutive NF-kB activity
- ✓ In contrast to antigen and chronic active BCR signaling, the antigen-independent signal, termed 'tonic BCR signaling', is mediated by PI3K + PI3K /AKT/mTOR, but not the NF-kB pathway, to promote the proliferation and survival of malignant B cells. Genomic data have shown that GCB-DLBCL lines exclusively use tonic BCR signaling.

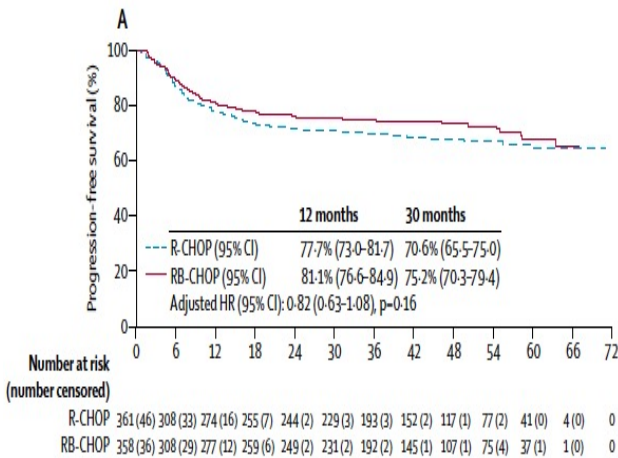
## Attempt to improve the outcome with the addition of novel drugs with or after R-CHOP: overall no significative advantage

Drug	Regimen	Subtype or not	Study	Results
<b>R-CHOP + X as induction</b>				
Bevacizumab <sup>1</sup>	RA-CHOP	DLBCL	Main	No advantage (PFS and OS)
Bortezomib <sup>2</sup>	BorR-CHOP	DLBCL	ReMoDL-B	No PFS advantage
Ibrutinib <sup>3</sup>	IR-CHOP	Non-GCB DLBCL	Phoenix	No EFS advantage
Lenalidomide <sup>4</sup>	R <sup>2</sup> -CHOP	ABC-DLBCL	Robust	No PFS advantage
Venetoclax <sup>5</sup>	VR-CHOP	DLBCL	Cavalli	Promising results
<b>R-CHOP + X as maintenance</b>				
Rituximab <sup>6</sup>	Rituximab	DLBCL	NHL-13	No EFS advantage 3-yr
Enzanstaurin <sup>7</sup>	Enzanstaurin	DLBCL	Prelude	No DFS advantage 4yr
Everolimus <sup>8</sup>	Everolimus	DLBCL	Pillar-2	No DFS advantage 2yr
Lenalidomide <sup>9</sup>	Lenalidomide	Elderly DLBCL	Remarc	PFS advantage, no OS

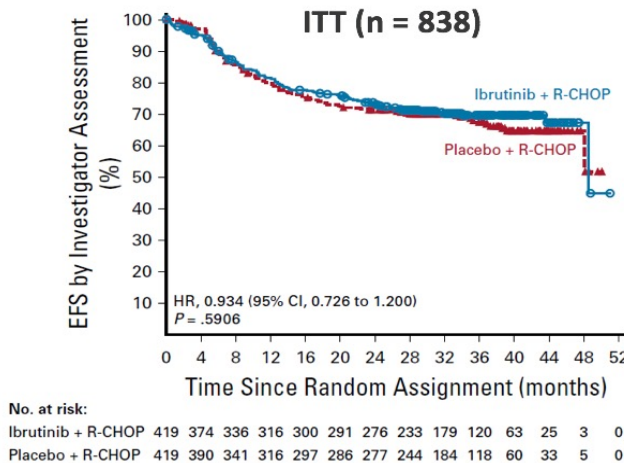
1. Seymour JF et al, Haematologica 2014; 2. Davies A et al, Lancet Oncol 2019; 3. Younes A et al, J Clin Oncol 2019; 4. Vitolo U et al, Hematol Oncol 2019; 5. Morschhauser F et al, Blood 2021; 6. Jagger U et al, Haematologica 2013; 7. Crump M et al, J Clin Oncol 2016; 8. Witzig T et al, Ann Oncol 2018; 9. Thieblemont C et al, J Clin Oncol 2019.

# Moving beyond R-CHOP... targeting ABC DLBCL

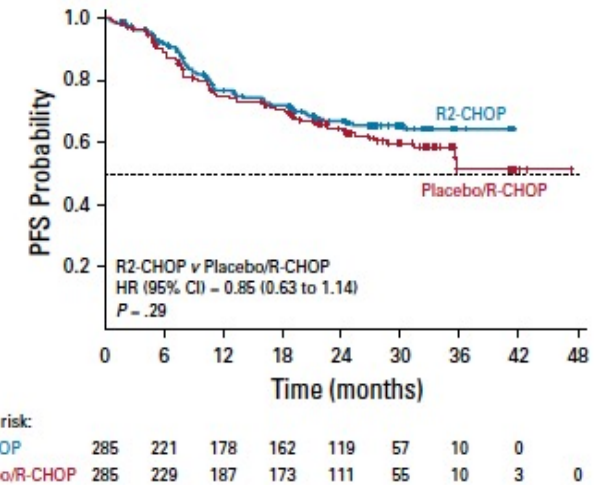
## R-CHOP + Bortezomib



## R-CHOP + iBTK



## R-CHOP + Lenalidomide



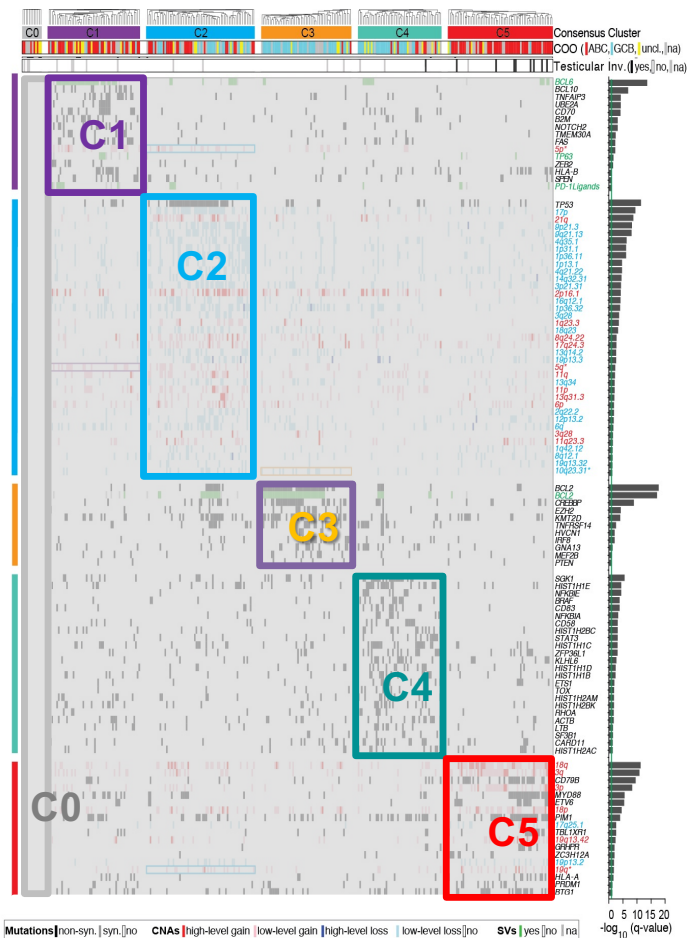
Davies A, et al. Lancet Oncol 2019; Younes A, et al. J Clin Oncol 2019; Nowakowski G, et al. J Clin Oncol 2021.

## Where are we wrong? Should we still care about COO?

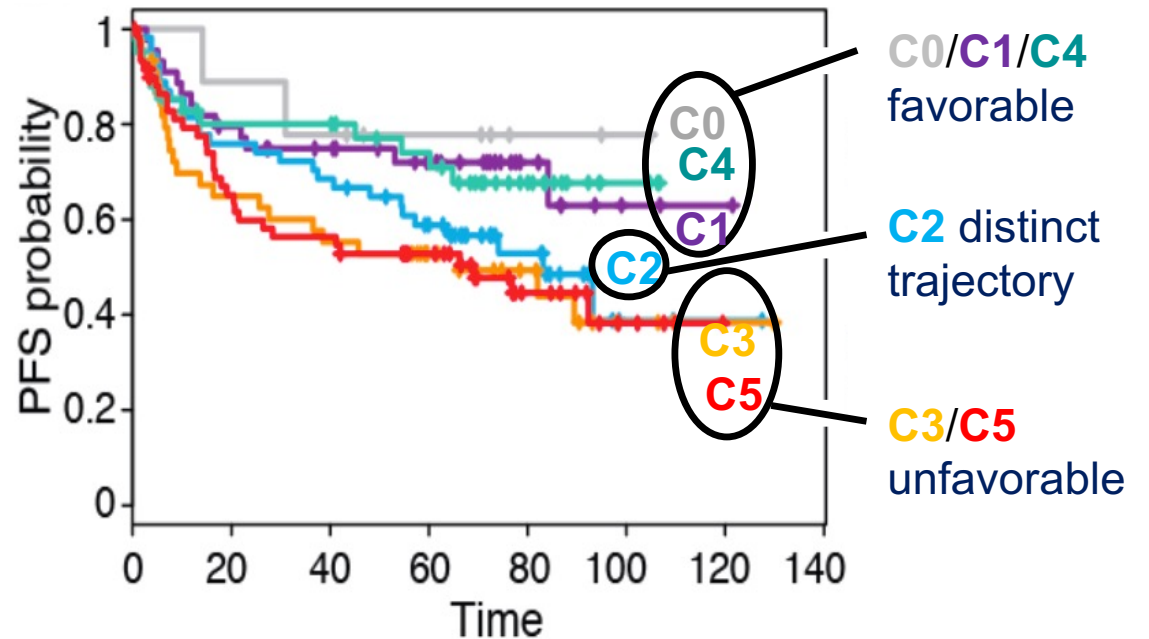
- ▶ Yes: the prognosis of ABC is still unsatisfactory
- ▶ Yes: subgroups of ABC patients benefit from the addition of specific drugs as ibrutinib in young and lenalidomide in high risk
- ▶ No: ABC *alone* is not the best target; DLBCLs are more heterogenous, mutational alterations, etc
- ▶ Maybe: ibrutinib or lenalidomide are not the best drugs, we need better drugs, novel-novel combinations
- ▶ .....to be continued.

# Genetically-distinct DLBCL Subsets are Predictive for Outcome

## Genetically-distinct DLBCLs

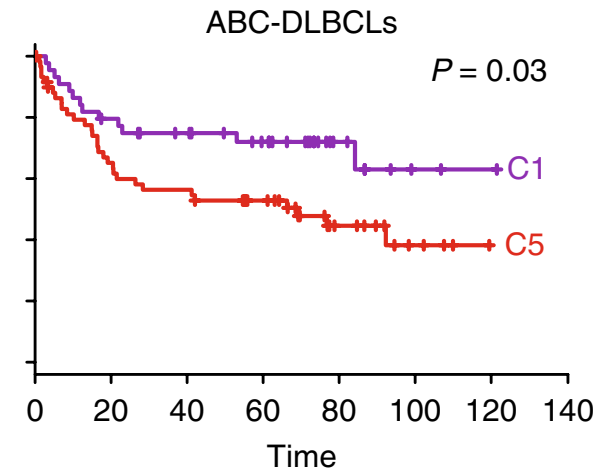
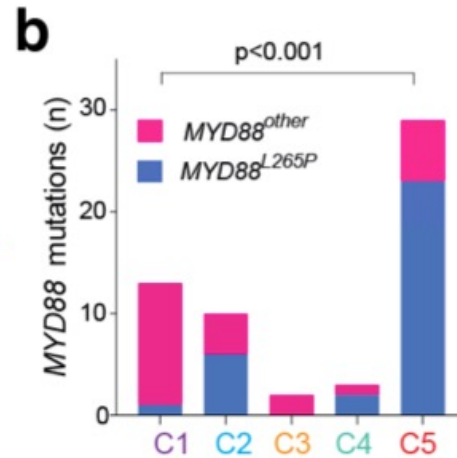
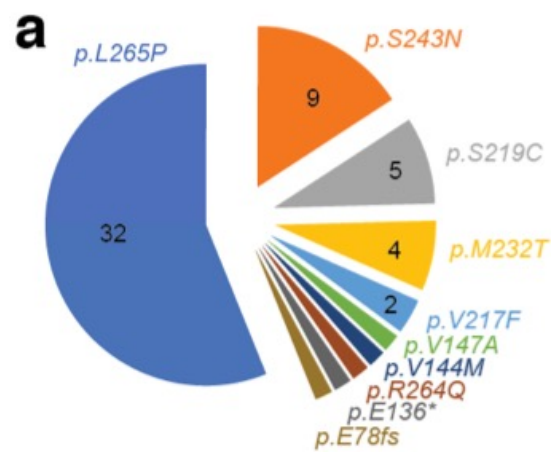


## Predictive for Outcome



- Genetic signatures comprised of
  - Mutations
  - Somatic copy number alterations (SCNAs)
  - Structural Variants (SVs)

# C1 vs. C5 DLBCLs – Two Genetically Distinct ABC-DLBCLs



	C1 DLBCLs	C5 DLBCLs
<b>c</b> MYD88 mutations	23% (13/56)	44% [28/64]
Type of MYD88 mutations	non-L265P	L265P
Concordant CD79B mutations	no	frequent

- Different types and incidences of MYD88 mutations

➔ C1 and C5 ABC-type DLBCLs arise by distinct pathogenetic mechanisms.

- C5 DLBCLs** - highest cAID activity
  - tumors passaged through the GC
- C1 DLBCLs** - low to absent cAID activity
  - suggestive of extrafollicular origin

## Evolving Strategies in the Treatment of DLBCL

Targeting all comers

Vs

Single gene/single drug model

Vs

Combination of genes/combination of drugs?

## Drugs by Molecular Classification Subgroups

Genetically defined category	Drugs
MCD/C5	ibrutinib, acalabrutinib, venetoclax
BN2/C1	ibrutinib, bortezomib, carfilzomib
EZB/C3	venetoclax, tazemetostat, idelalisib, copanlisib, duvelisib, umbralisib
C4	idelalisib, copanlisib, duvelisib, bortezomib, carfilzomib, ruxolitinib

**MCD:** MYD88<sup>L265P</sup> and CD79B mutations / **C5**

*predominantly ABC*

**BN2 :** BCL6 fusions and NOTCH2 mutations / **C1**

*both ABC and GCB*

**N1 :** NOTCH1 mutations

*predominantly ABC*

**EZB :** EZH2 and BCL2 mutations / **C3**

*predominantly GCB*



## Genetic Subtype Guided Rituximab-based Immunochemotherapy Improves Outcome in Newly Diagnosed Diffuse Large B-cell Lymphoma: First Report of a Randomized Phase 2 Study

<u>Subtype</u>	<u>Drug X</u>
MCD, BN2	ibrutinib
N1, NOS	lenalidomide
EZB	tucidinostat (HDAC) inhibitor
TP53	decitabine

- 128 patients:

NOS – 50    TP-53 – 21

MCD – 26    EZB – 3

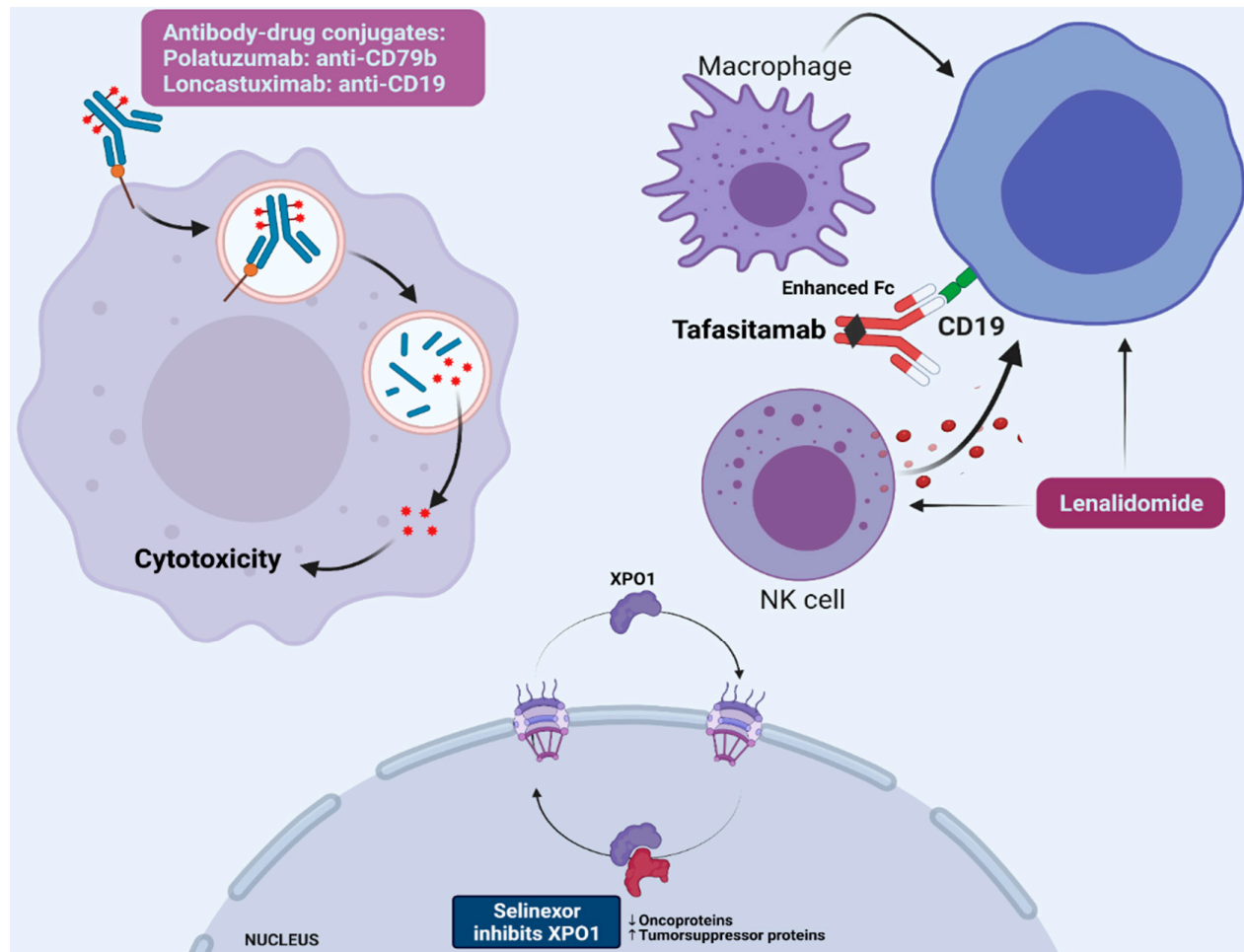
BN2 – 23    N1 – 3

- Outcome:  
CR  
1 yr. PFS<sub>12</sub>

CHOP-R  
65%  
79%

CHOP-R-X  
85%  
96%

## Mechanisms of Action for recent approved novel therapy in R/R DLBCL

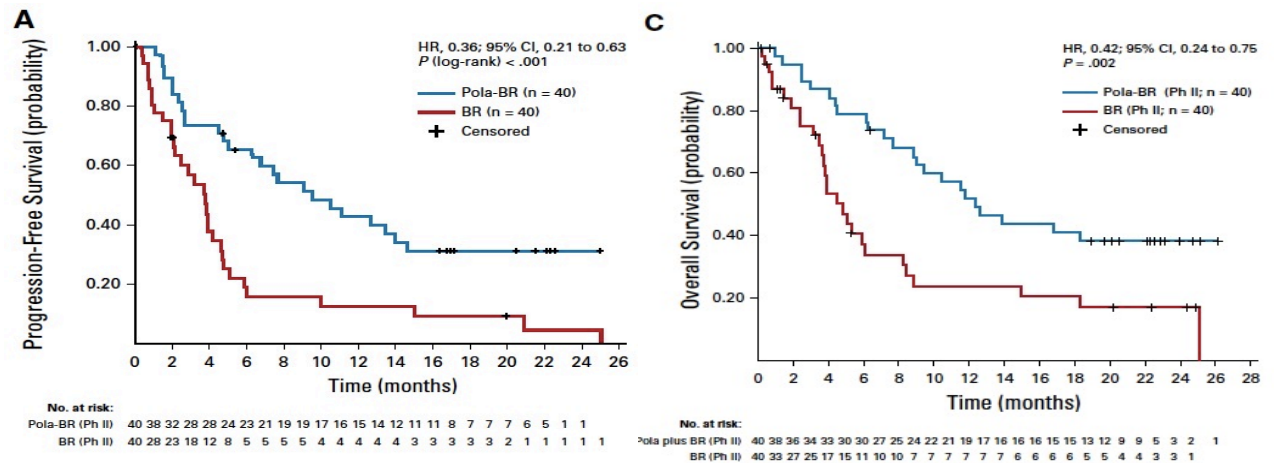


# ADC + R-chemotherapy

## Polatuzumab Vedotin in Relapsed or Refractory Diffuse Large B-Cell Lymphoma

Laurie H. Sehn, MD, MPH<sup>1</sup>; Alex F. Herrera, MD<sup>2</sup>; Christopher R. Flowers, MD, MSc<sup>1</sup>; Manali K. Kamdar, MD, MBBS<sup>4</sup>; Andrew McMillan, PhD<sup>5</sup>; Mark Hertzberg, MBBS, PhD<sup>6</sup>; Sarit Assouline, MDCM, MSc<sup>7</sup>; Tae Min Kim, MD<sup>8</sup>; Won Seog Kim, MD, PhD<sup>9</sup>; Muhit Ozcan, MD<sup>10</sup>; Jamie Hirata, PharmD<sup>11</sup>; Elicia Penuel, PhD<sup>11</sup>; Joseph N. Paulson, PhD<sup>11</sup>; Ji Cheng, PhD<sup>12</sup>; Grace Ku, MD<sup>11</sup>; and Matthew J. Matasar, MD<sup>13</sup>

J Clin Oncol 38:155-165. © 2019



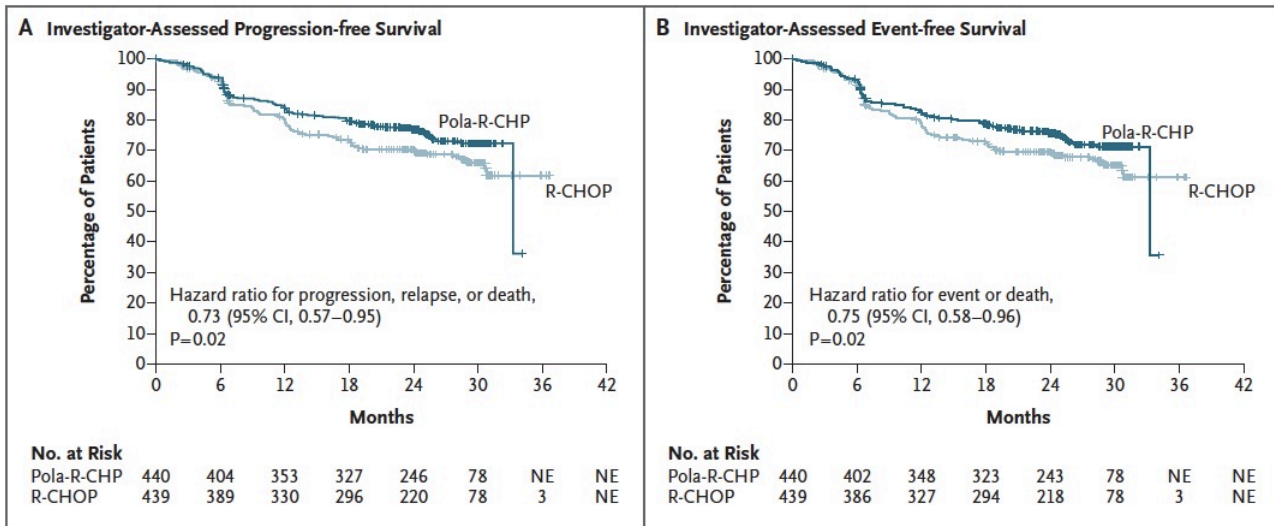
R/R

ORIGINAL ARTICLE

## Polatuzumab Vedotin in Previously Untreated Diffuse Large B-Cell Lymphoma

H. Tilly, F. Morschhauser, L.H. Sehn, J.W. Friedberg, M. Trněný, J.P. Sharman, C. Herbaux, J.M. Burke, M. Matasar, S. Raj, K. Izutsu, N. Mehta-Shah, L. Oberic, A. Chauchet, W. Jurczak, Y. Song, R. Greil, L. Mykhalska, J.M. Bergua-Burgués, M.C. Cheung, A. Pinto, H.-J. Shin, G. Hapgood, E. Munhoz, P. Abrisqueta, J.-P. Gau, J. Hirata, Y. Jiang, M. Yan, C. Lee, C.R. Flowers, and G. Salles

N Engl J Med 2022;386:351-63.

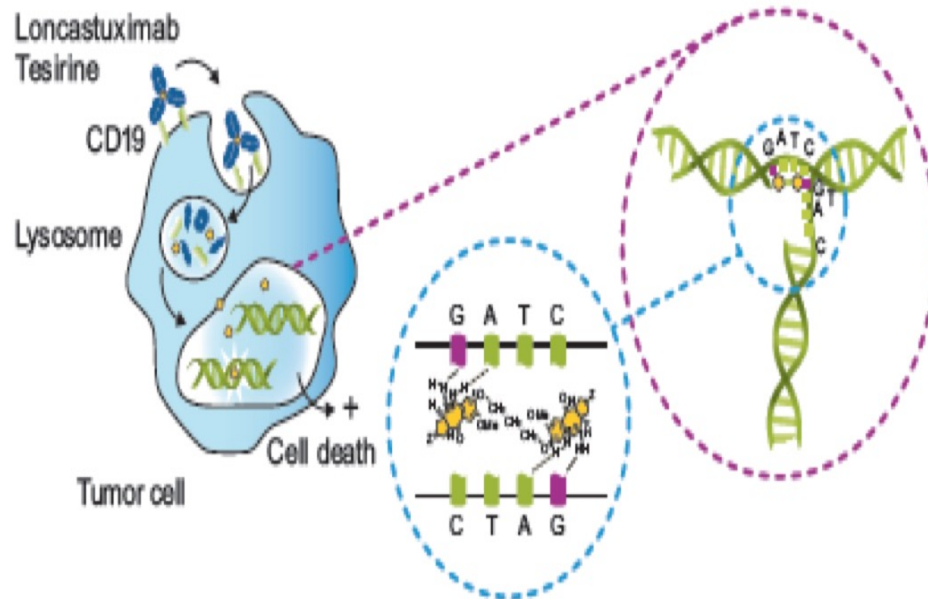


FIRST LINE

## ...adding mAb antiCD19?

Loncastuximab tesirine: humanized anti-CD19 antibody, stochastically conjugated through a cathepsin-cleavable valine-alanine linker to a pyrrolobenzodiazepine (PBD) dimer toxin causing DNA crosslinking.

### Mechanism of action of Lonca



### Tafasitamab (Fc-enhanced, anti-CD19 mAb)<sup>1-3</sup>

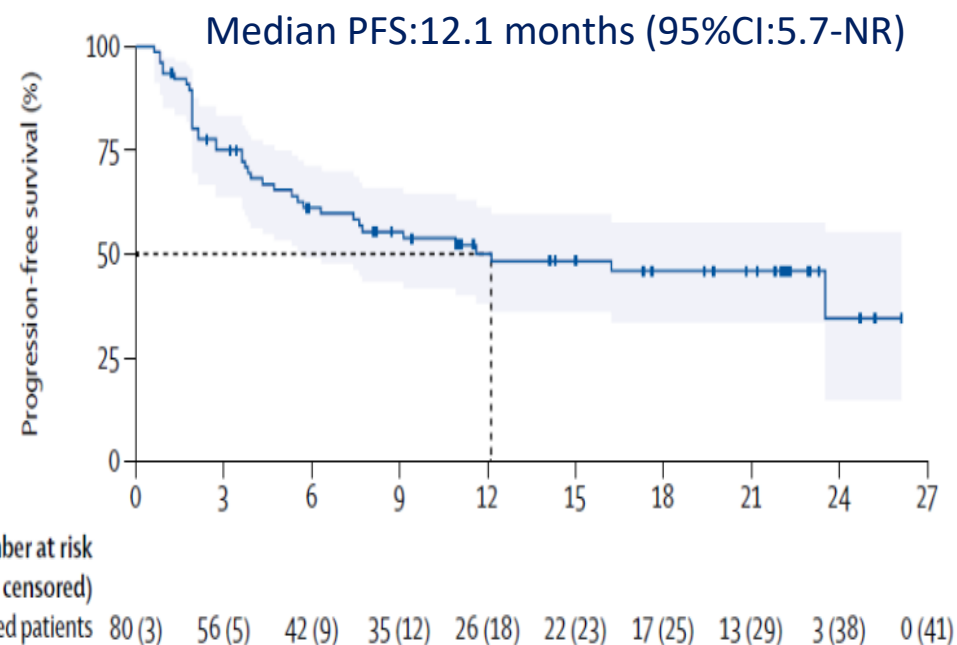
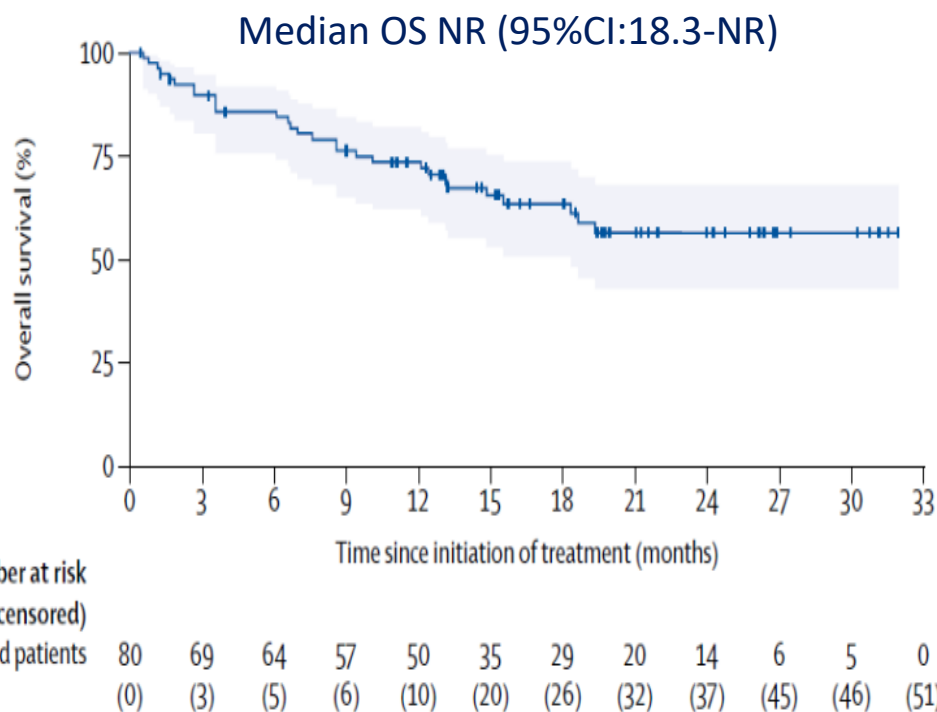
Affinity-matured  
CD19 binding site



- ADCC ↑
- ADCP ↑
- Direct cell death
- Encouraging single-agent activity in patients with R/R DLBCL and iNHL

# Tafasitamab + lenalidomide in R/R DLBCL

Single arm phase II study L-Mind



Median follow-up:19.6 months

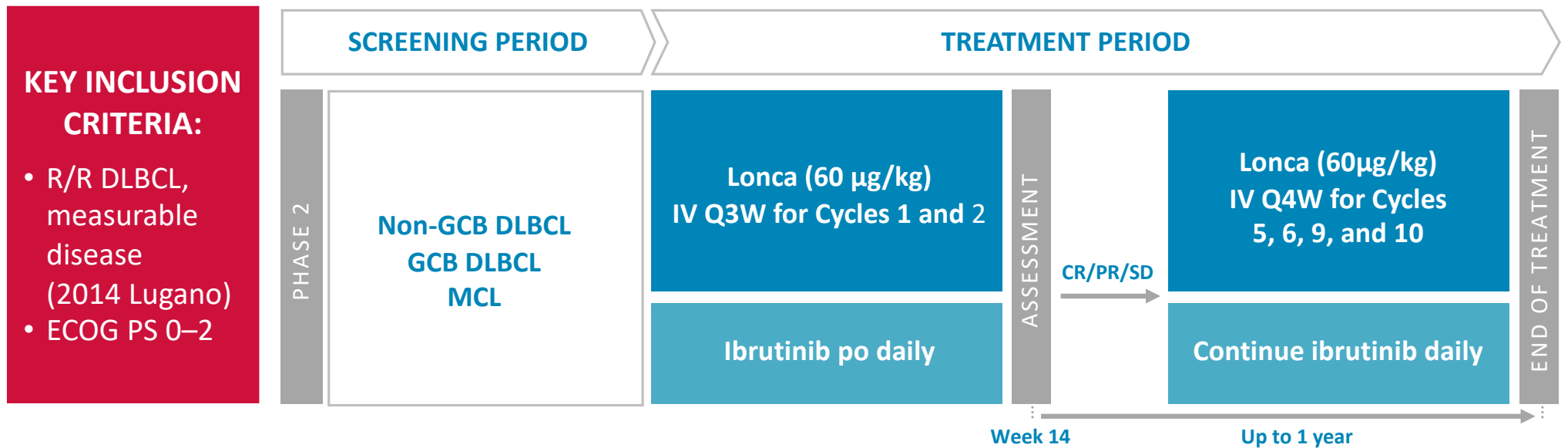
## Phase 2 Study of Loncastuximab Tesirine Plus Ibrutinib in RR-DLBCL (LOTIS-3)

### Primary phase 2 study objective:

- CRR assessed by central review in R/R non-GCB DLBCL (investigator-determined COO)

### Planned interim analysis objective:

- To determine if CRR in the non-GCB DLBCL cohort warranted the continuation of patient enrollment for study completion<sup>a</sup>



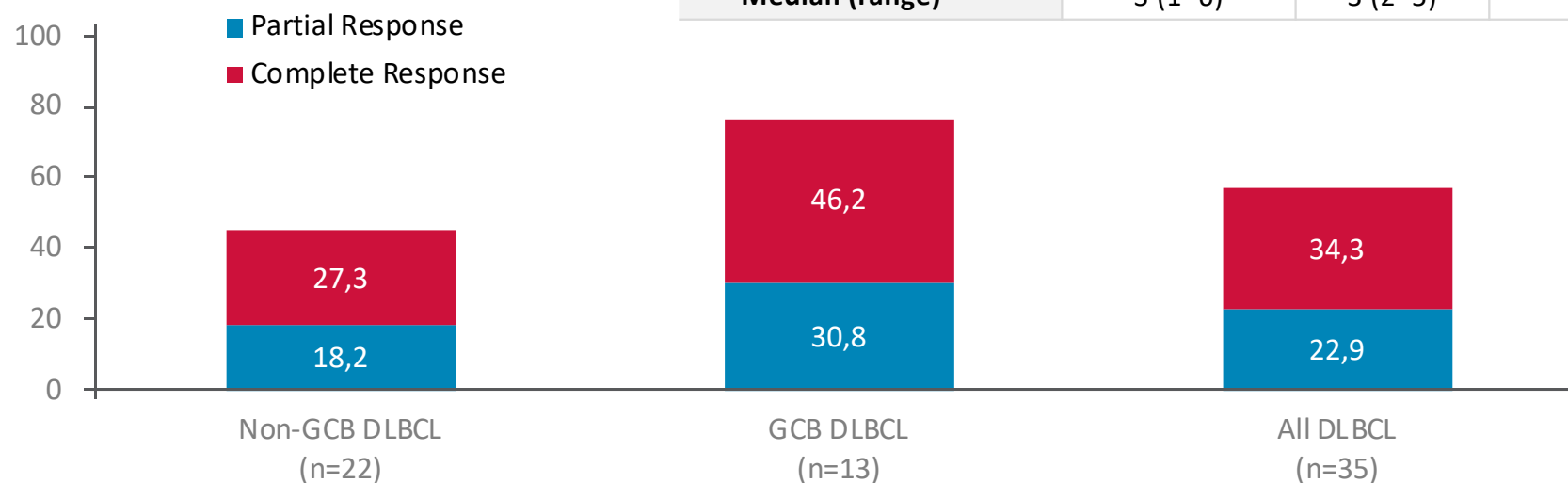
As of Aug 30, 2021, 35 patients with R/R DLBCL received Lonca 60 µg/kg plus ibrutinib 560 mg

## Phase 2 Study of Loncastuximab Tesirine Plus Ibrutinib in RR-DLBCL (LOTIS-3)

Median Lonca cycles: 2 (range: 1–6)

Median ibrutinib cycles: 3.5 (range: 1–15)

Characteristic	Non-GCB (n=22)	GCB (n=13)	All patients (n=35)
Age, yrs, median (range)	72 (19–82)	66 (53–82)	72 (19–82)
Prior systemic therapies, n Median (range)	3 (1–6)	3 (2–5)	3 (1–6)

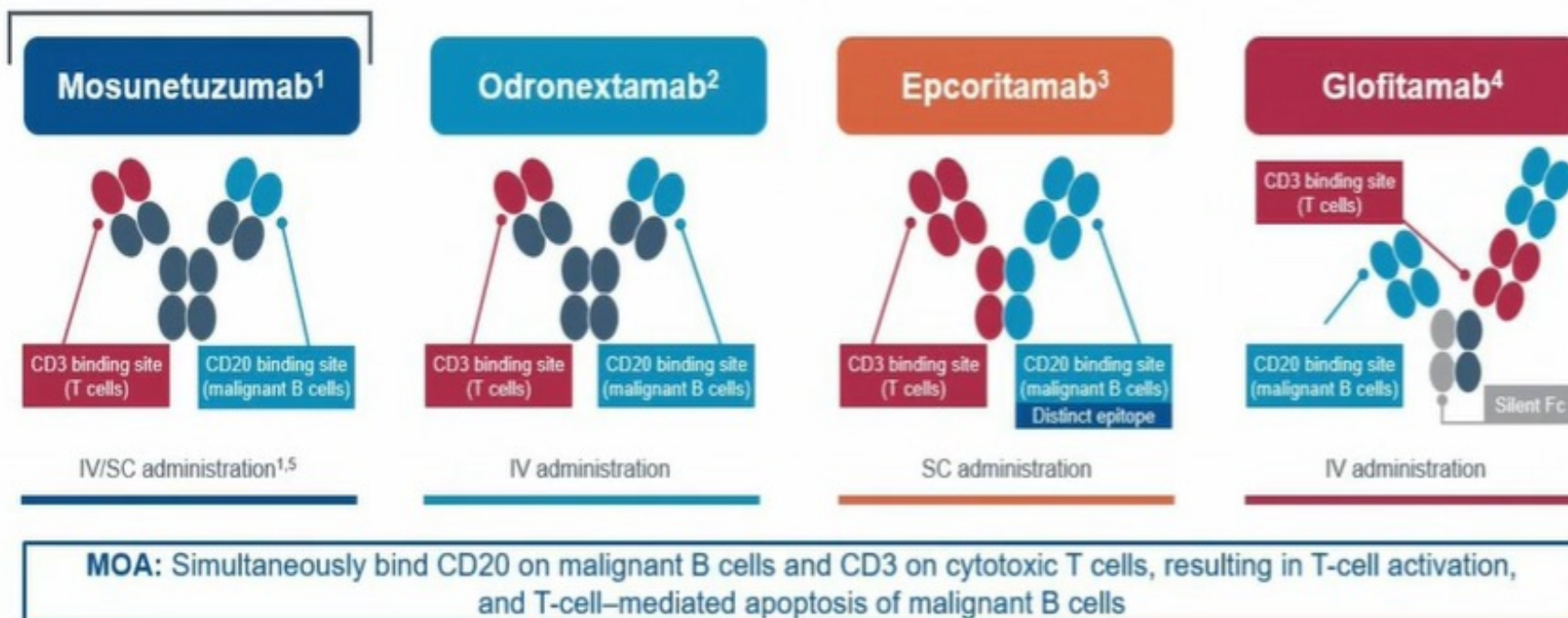


ORR (n/N) (95% CI) <sup>b</sup>	45.5% (10/22) (24.4, 67.8)	76.9% (10/13) (46.2, 95.0)	57.1% (20/35) (39.4, 73.7)
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## Emerging therapies: Bispecific Antibodies

Investigational CD20×CD3 bispecific antibodies for B-cell lymphomas:

FDA BTD for R/R FL (2020)





# CD20xCD3 bispecific antibodies + SoC

Rational combinations of targeted therapies

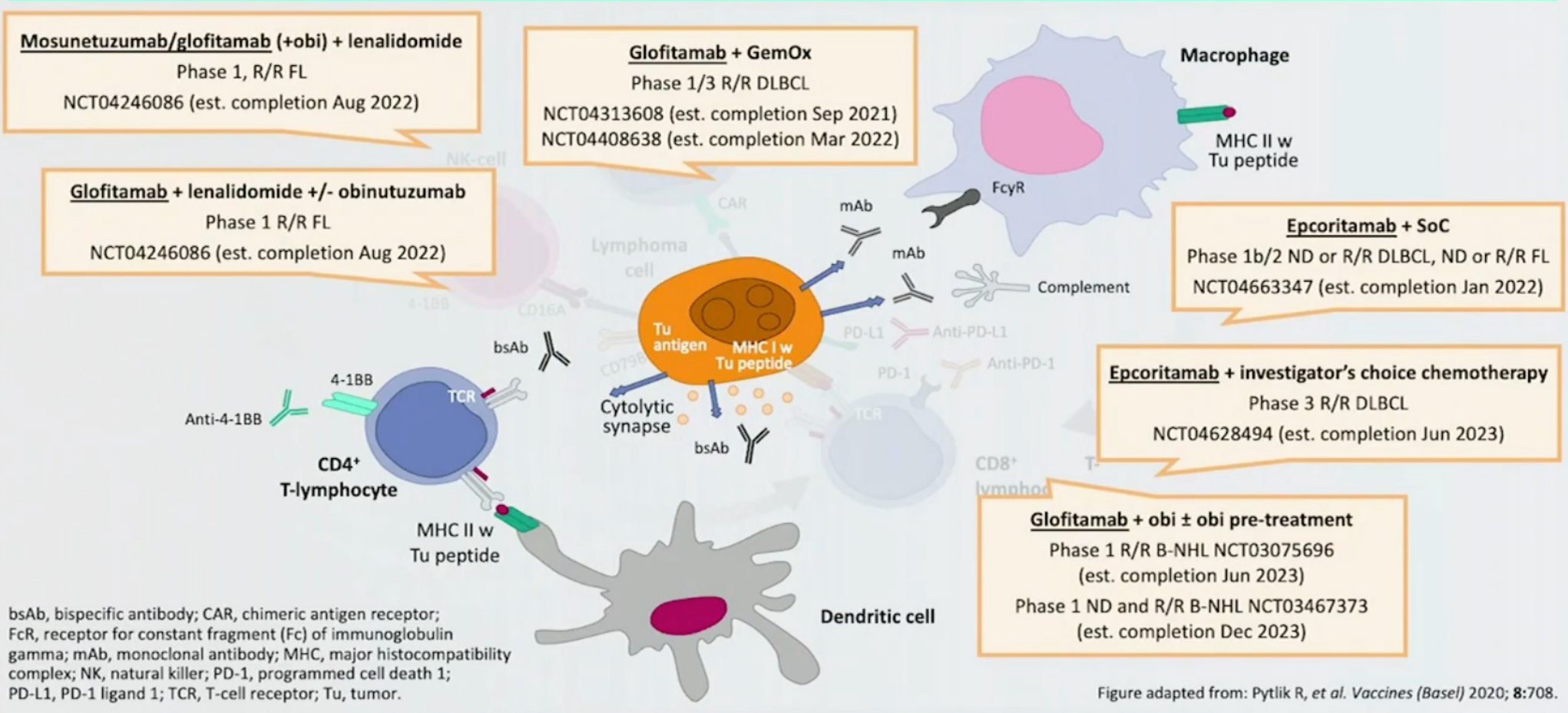
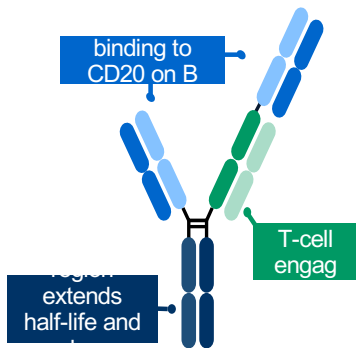


Figure adapted from: Pytlik R, et al. *Vaccines (Basel)* 2020; 8:708.

# Glofitamab in Combination with Polatuzumab Vedotin: Phase Ib/II in 59 pts with R/R Diffuse Large B-Cell Lymphoma (DLBCL)

Glofit + Pola arm: study design in R/R DLBCL



Key inclusion criteria (DLBCL arm)	Objectives	
<ul style="list-style-type: none"> <li>Age ≥18 years</li> <li>R/R DLBCL (including trFL and HGBCL)</li> <li>ECOG performance status 0–2</li> </ul>	<b>Primary:</b> <ul style="list-style-type: none"> <li>DLTs</li> <li>Determine MTD and/or RP2D for Glofit + Pola (including obinutuzumab pretreatment)</li> </ul>	<b>Secondary:</b> <ul style="list-style-type: none"> <li>Safety and tolerability</li> <li>Efficacy (CR rate and BORR per Lugano 2014<sup>1</sup>)</li> </ul>

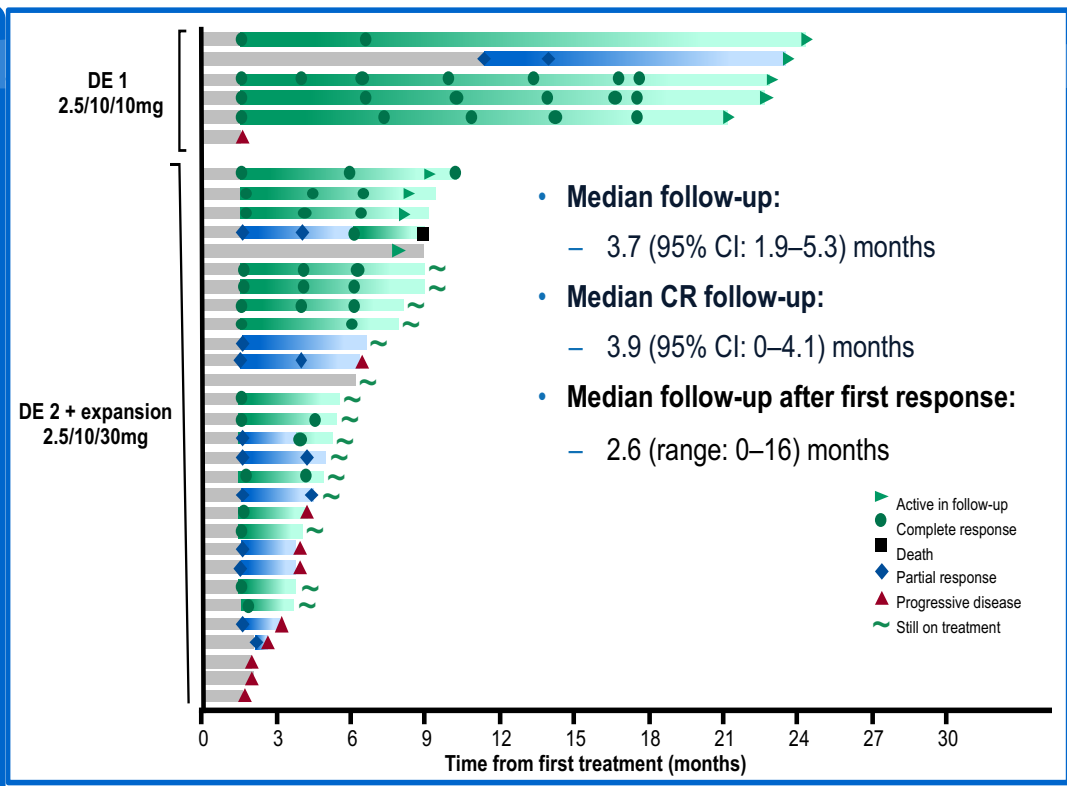
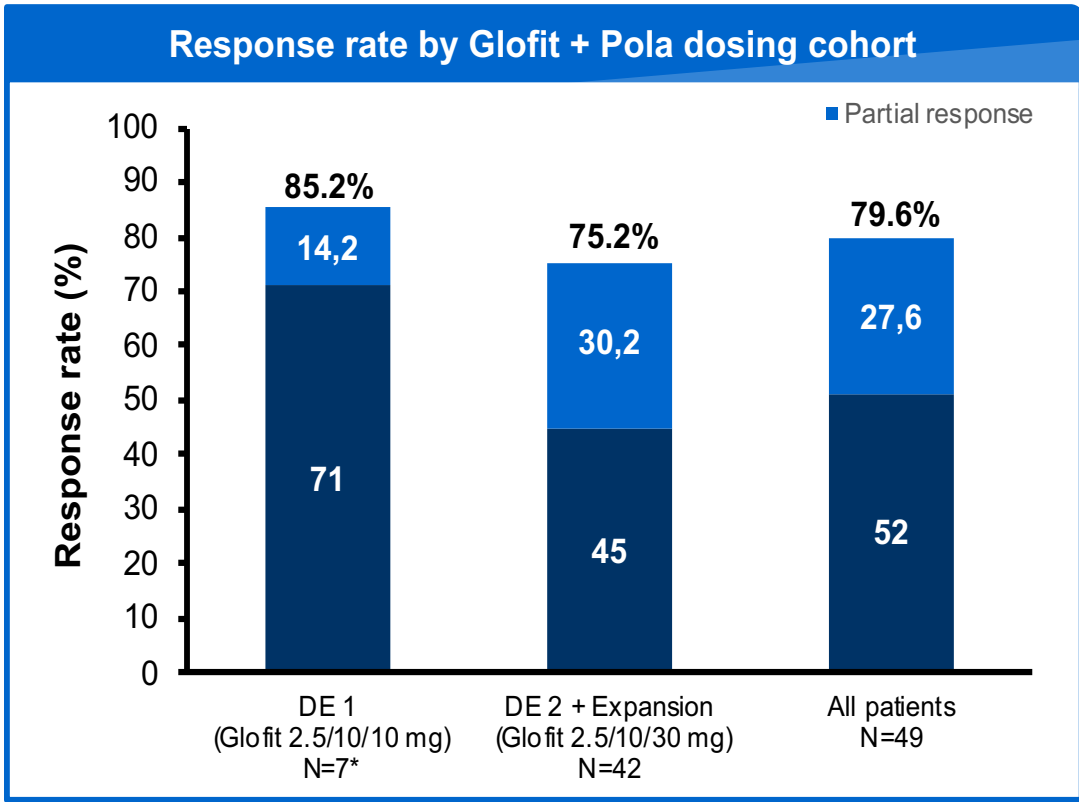
### Glofit + Pola administration in R/R DLBCL

- Target enrollment ~90 patients
- CRS mitigation: obinutuzumab IV 1000 mg 7 days prior to glofitamab administration (step-up dosing)
- Efficacy assessments with PET-CT C3D1, C6D1 C8D15, EOT and Q3M

C1	C2–6	C7–12
D1: Obinutuzumab pretreatment 1000 mg D2: Polatuzumab vedotin 1.8 mg/kg D8: Glofitamab 2.5 mg D15: Glofitamab 10 mg	D1: Polatuzumab vedotin 1.8 mg/kg D1: Glofitamab 10 or 30 mg	D1: Glofitamab 10 or 30 mg
Glofitamab step-up dosing	Glofitamab target dose (Q3W)	Glofitamab target dose (Q3W)

*Cycle = 21 days; glofitamab (C2–C12) and polatuzumab vedotin (C2–C6) administered Q3W*

# Glofitamab in Combination with Polatuzumab Vedotin: response rate and adverse events



Hutchings M et al. Abs#525, ASH 2021.



## topMIND: PHASE 1B/2A BASKET STUDY TO EVALUATE TAFASITAMAB<sup>a</sup> AND THE PI3K $\delta$ INHIBITOR PARSACLISIB IN RELAPSED/REFRACTORY NON-HODGKIN LYMPHOMA OR CHRONIC LYMPHOCYTIC LEUKAEMIA<sup>1</sup>

Adult patients with R/R B-cell malignancies, including DLBCL, MCL, FL, MZL and CLL/SLL, with ECOG PS 0–2 and  $\geq 2$  prior systemic antilymphoma/antileukemia therapies (N=100)

Tafasitamab 12 mg/kg IV QW (Cycles 1–3) then Q2W (Cycle 4 onward), plus parsaclisib 20 mg QD (Cycles 1–2) then 2.5 mg QD (Cycle 3 onward)

Phase 1b	R/R DLBCL n=10	R/R MCL n=10	R/R FL n=10	R/R MZL n=10	R/R CLL/SLL n=10
Phase 2a	n=10	n=10	n=10	n=10	n=10

### Primary Endpoint:<sup>b</sup>

- Phase 1b: incidence and severity of TEAEs and incidence of DLTs
- Phase 2a: ORR

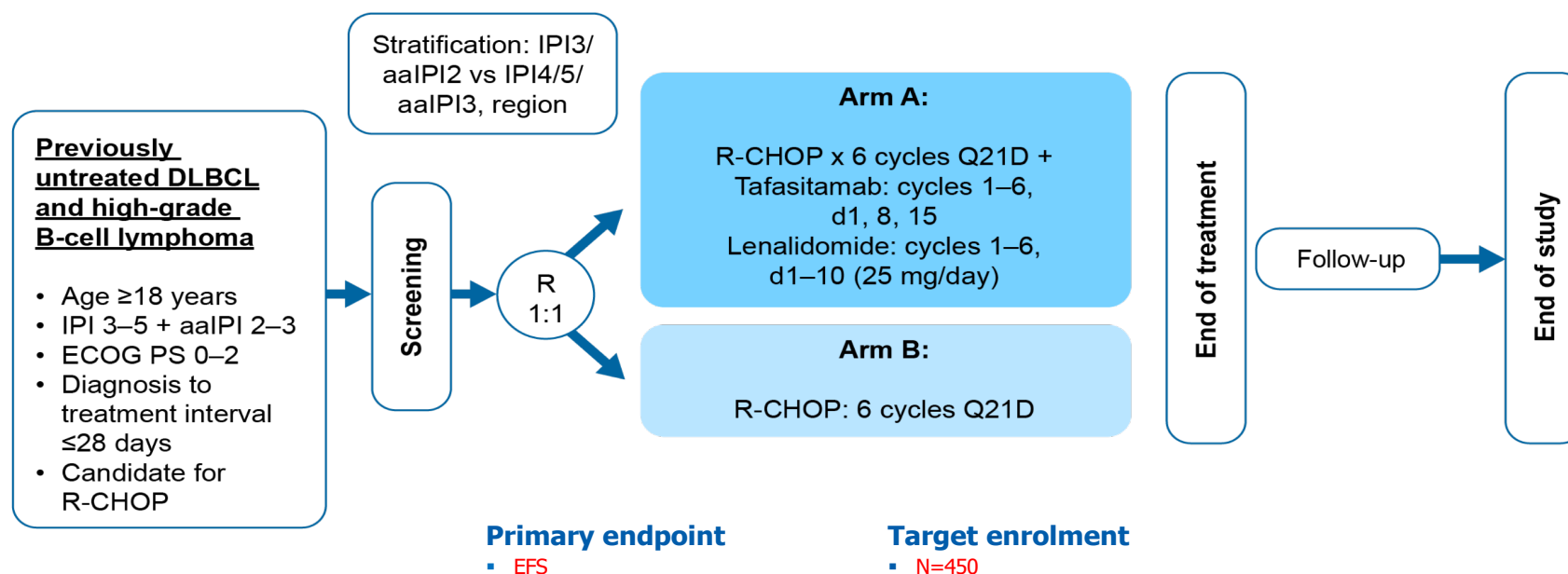
### Key Secondary/Exploratory Endpoints:<sup>b</sup>

- PK parameters of tafasitamab in combination with parsaclisib
- PK parameters of parsaclisib in combination with tafasitamab
- CRR, DOR, PFS, OS, MRD

- Immunogenicity of tafasitamab
- Cytokine, immune cell and tumour microenvironment response to tafasitamab plus parsaclisib
- Molecular markers for response or resistance

## frontMIND: STUDY DESIGN (MORPHOSYS TRIAL)

INTERNATIONAL, PROSPECTIVE, OPEN-LABEL PHASE 3 STUDY IN 1L DLBCL AND HIGH-GRADE B-CELL LYMPHOMA



1L, first-line; aaIPI, age-adjusted International Prognostic Index; d, day(s); DLBCL, diffuse large B-cell lymphoma; ECOG PS, Eastern Cooperative Oncology Group performance status; EFS, event-free survival; IPI, International Prognostic Index; Q21D, every 21 days; R, randomisation; R-CHOP, rituximab, cyclophosphamide, doxorubicin hydrochloride, vincristine and prednisolone.

## Conclusions

- ▶ R-CHOP is still the standard of care in DLBCL but we need to move forward to improve the outcome of our patients.
- ▶ COO is predictive of the outcome with ABC subtype having a worst prognosis in terms of systemic and also CNS progression but we cannot based anymore on this simple subgrouping
- ▶ A single target approach have failed underlining the molecular complexity of DLBCL
- ▶ A more accurate recognition of unfavourable DLBCL subsets is recommended to better tailor the treatment
- ▶ New study designs potentially focused on mutational alterations with combination of multiple novel drugs may have a greater chance of success.
- ▶ Novel-novel combinations as anti-CD19 and immunomodulators, or bispecific antibodies + different novel biological drugs or chemoimmunotherapy represent a step forward the cure of all DLBCL



**GRAZIE**