

Recent Advances in the Frontline Treatment of DLBCL: Is there a new standard of care?

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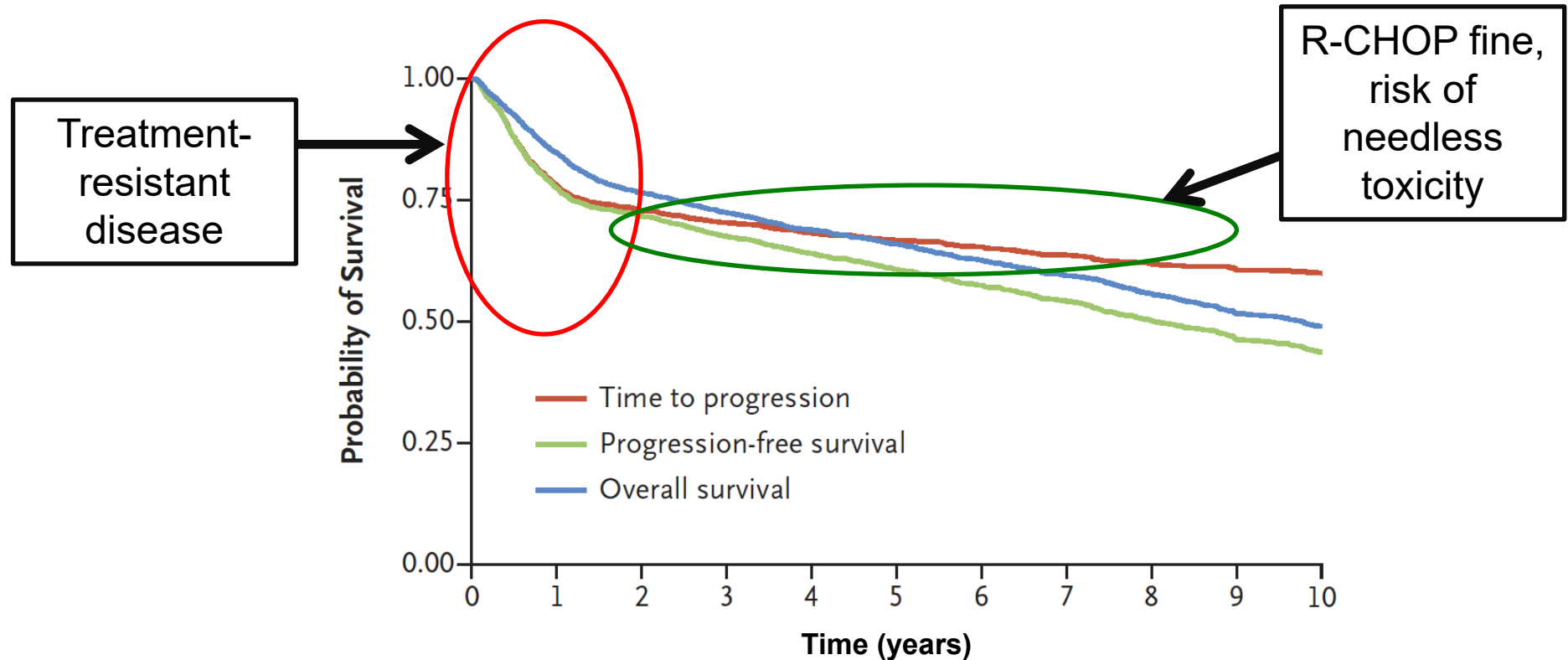
Disclosures

- **Consulting/Honoraria: Abbvie, Acerta, Amgen, Apobiologix, AstraZeneca, Celgene, Gilead, Incyte, Janssen, Kite, Karyopharm, Lundbeck, Merck, Morphosys, Roche/Genentech, Sandoz, Seattle Genetics, Servier, Teva, Takeda, TG Therapeutics, Verastem**
- **Research funding: Teva, Roche/Genentech**

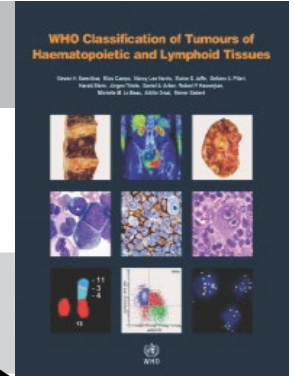
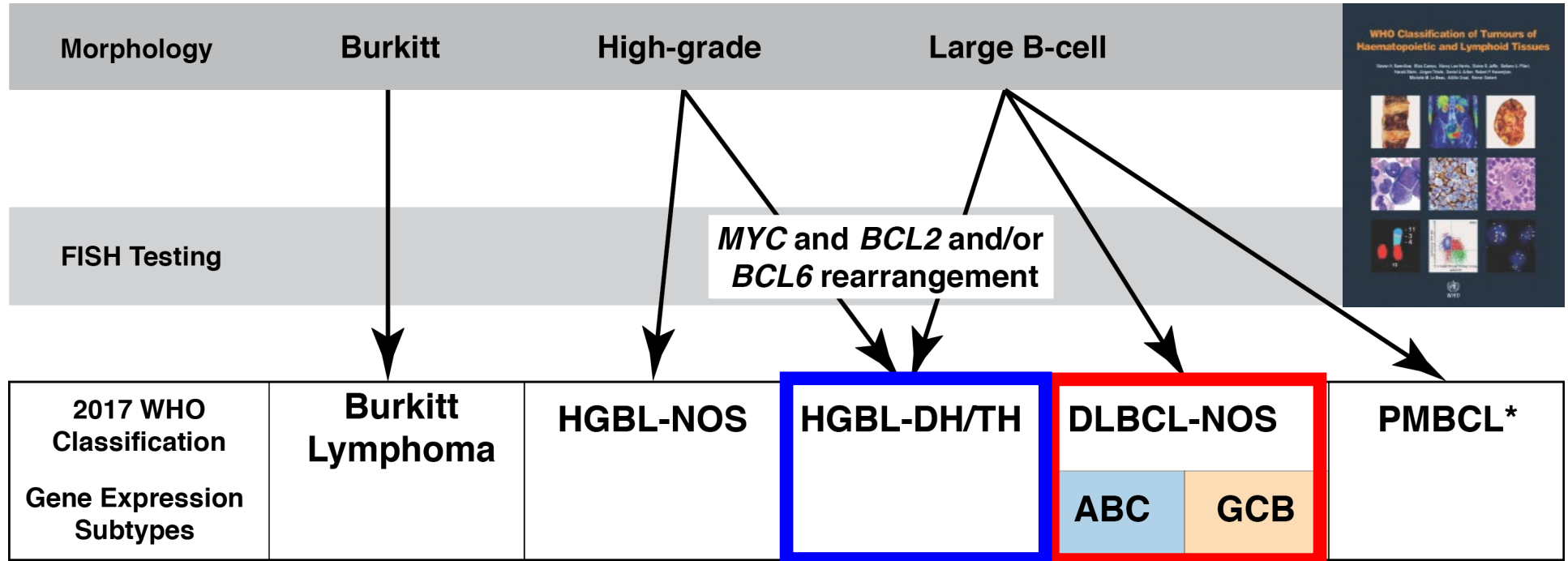
Key Considerations

- **What is standard of care and who do we treat differently now?**
- **Should we treat based on biology?**
- **Is the standard of care changing?**
- **What are the limitations to current trial design?**
- **How do we move the bar in the future?**

Outcomes with R-CHOP in Untreated DLBCL



WHO Classification – Aggressive B-cell Lymphoma



HGBL-NOS: high-grade B-cell lymphoma NOS

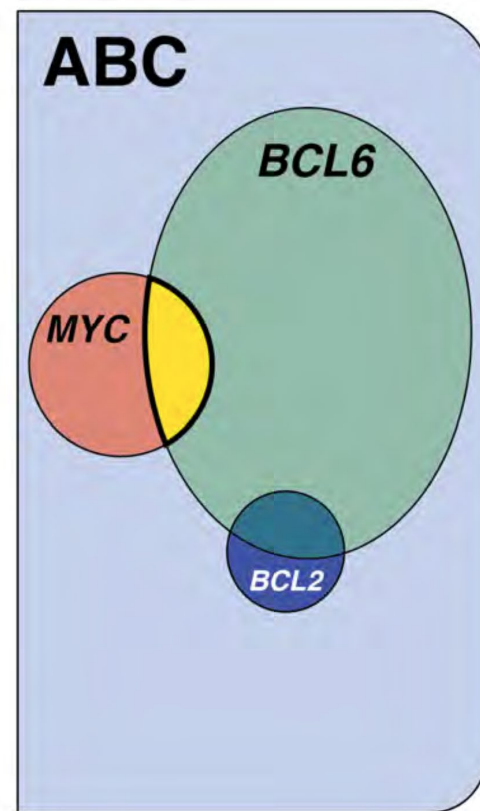
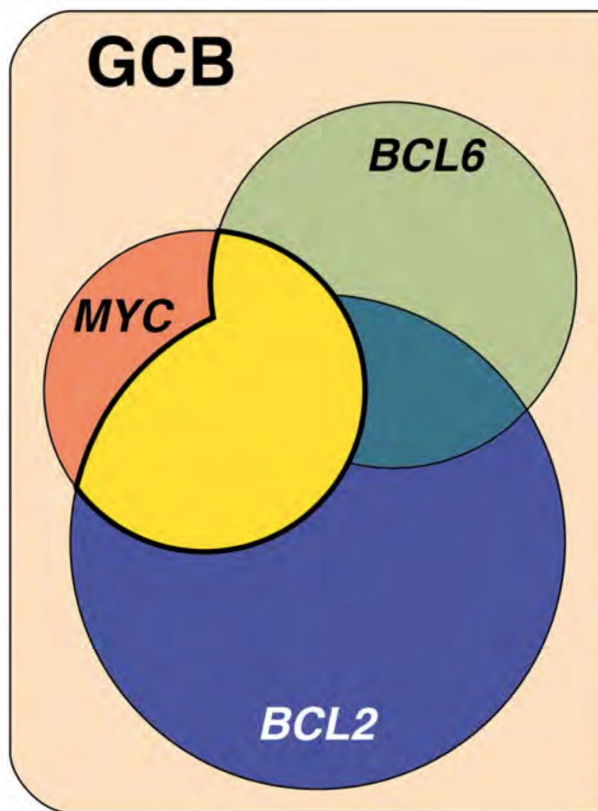
HGBL-DH/TH: high-grade B-cell lymphoma with *MYC* and *BCL2* and/or *BCL6* rearrangements

PMBCL: Primary mediastinal B-cell lymphoma

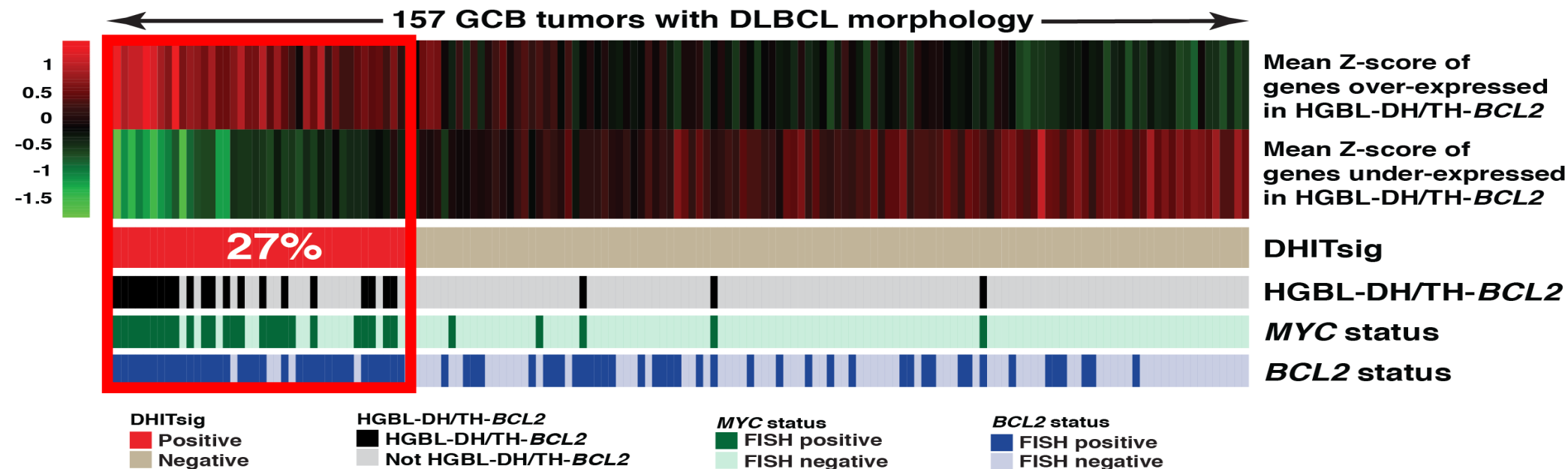
Swerdlow et al WHO revised 4th Edition 2017

Incidence of Double/Triple-Hit in DLBCL

- 12% harbour *MYC* rearrangements (> in GCB)
- ~7% are *MYC/BCL2* DHIT or *MYC/BCL2/BCL6* THIT
 - All cases are GCB
- ~1-2% are *MYC/BCL6*
 - GCB or ABC
- ~8% total incidence double/triple-hit

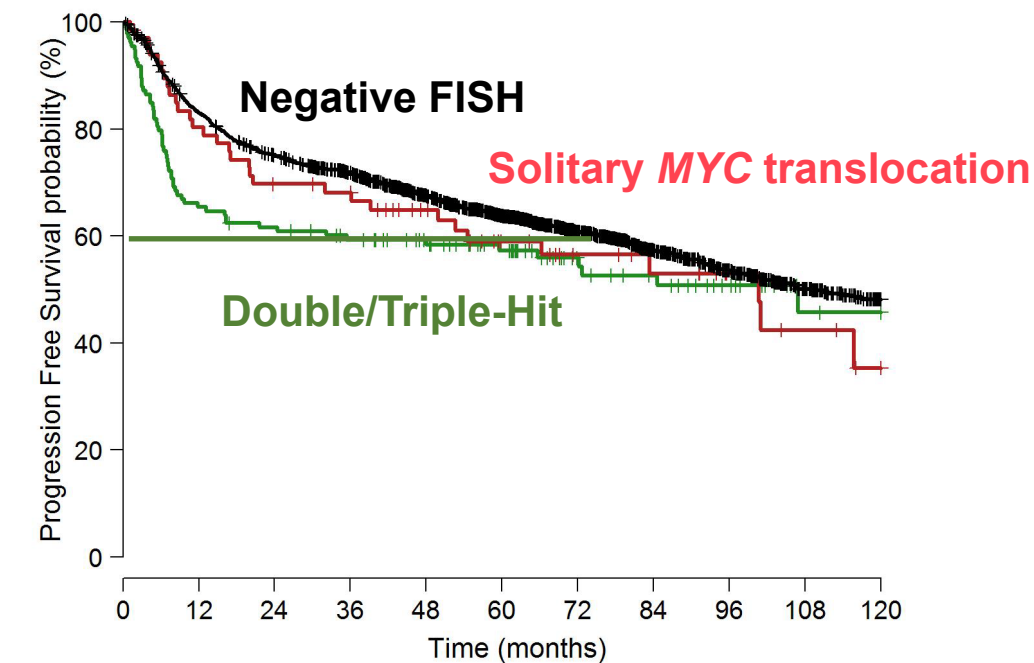


The “Double-Hit Gene Signature”



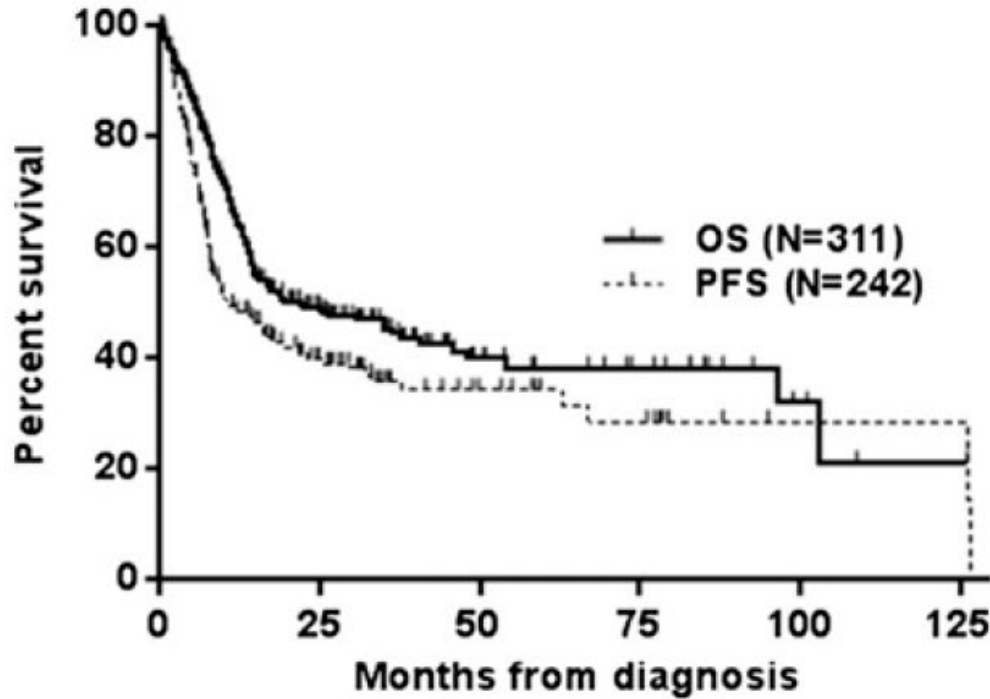
- Unique gene-expression signature identifies Double/Triple-Hit DLBCL
- Identifies an additional subset not detected by FISH

PFS in Patients with DLBCL Morphology Treated with R-CHOP According to FISH

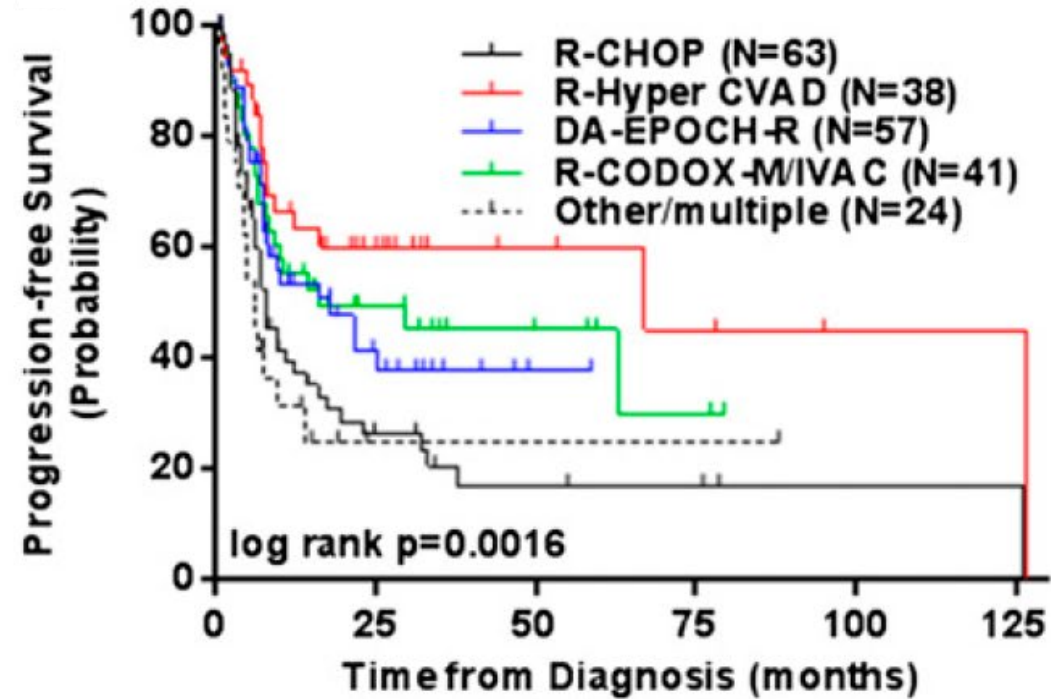


negative	2049	1687	1508	1371	1173	949	672	463	313	190	109
DH & TH	133	87	81	74	64	52	34	28	20	9	8
SH	67	53	45	43	34	25	18	15	10	7	4

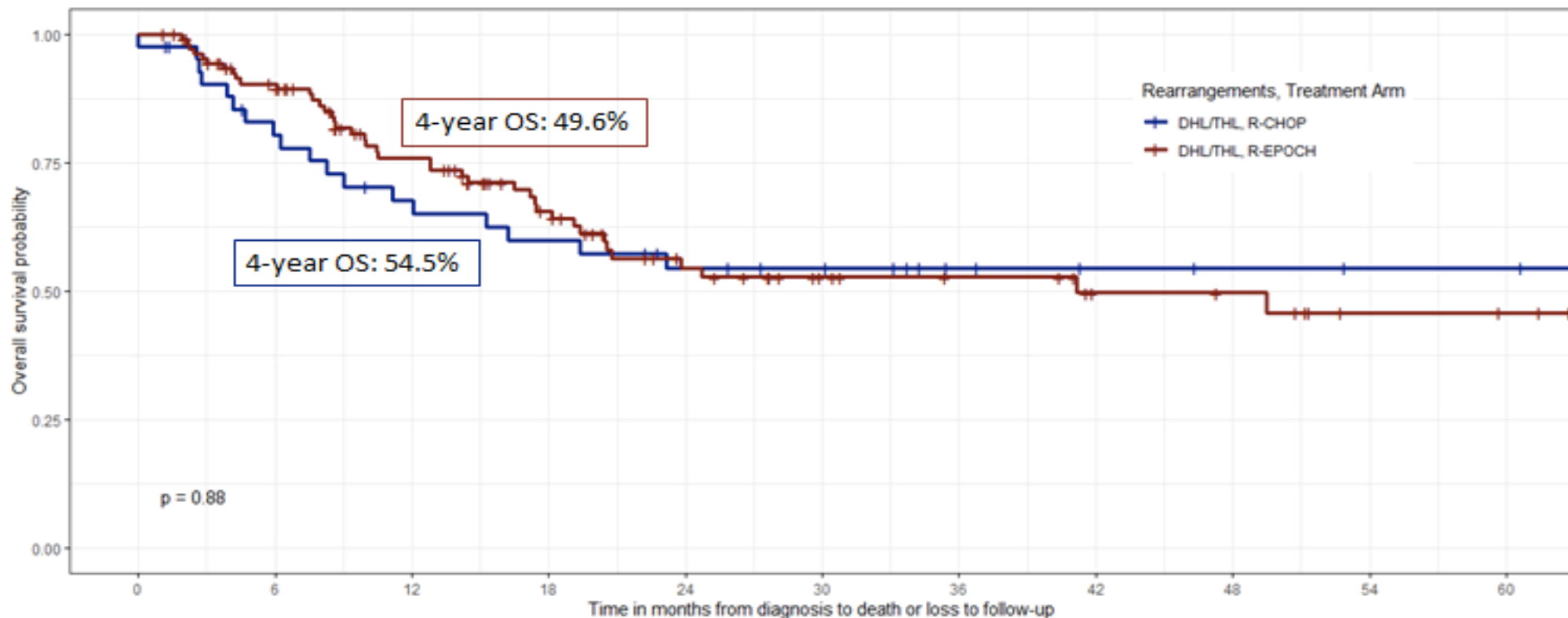
Outcome According to Induction Regimen in Double-Hit Lymphoma



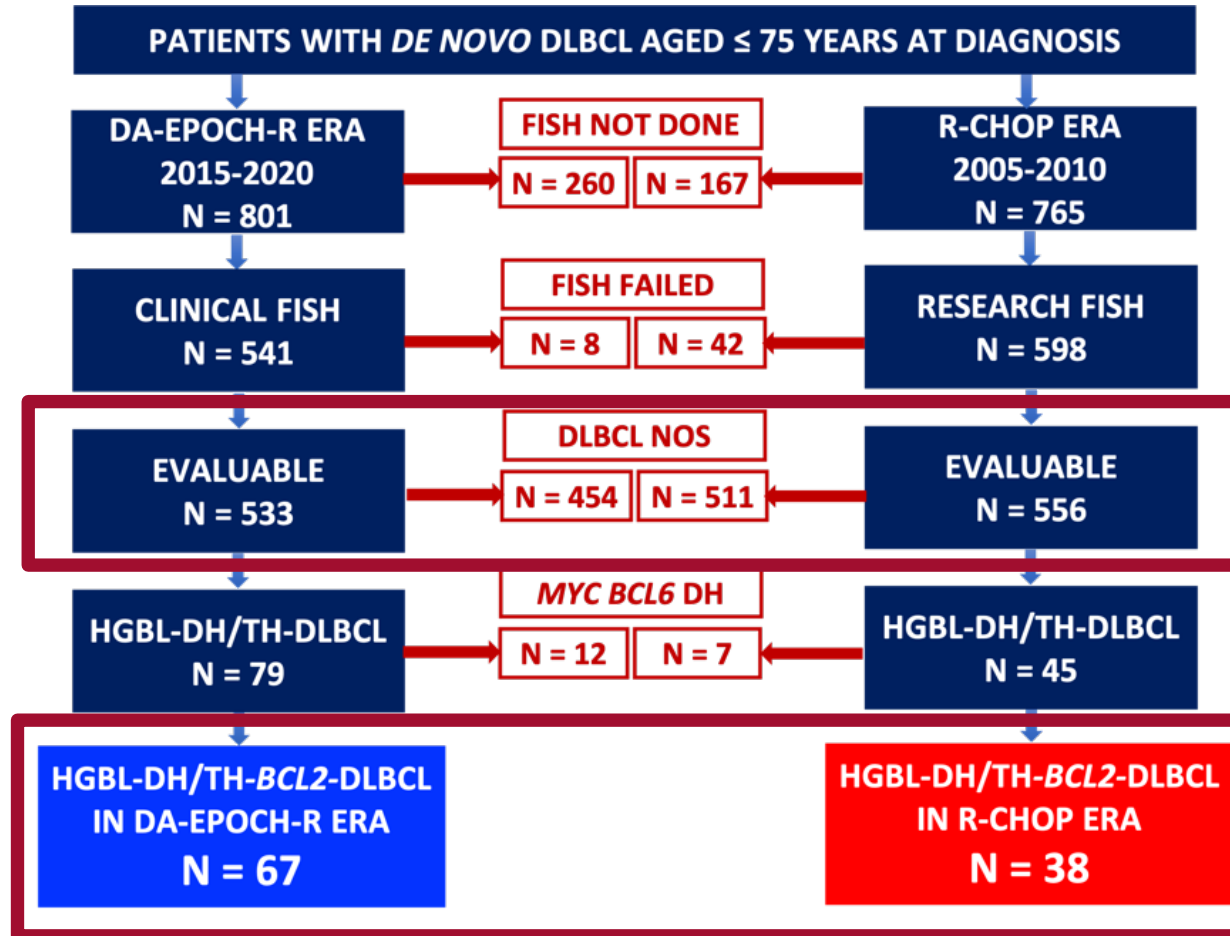
2y PFS 40%; 2y OS 49%



Retrospective Review Of R-EPOCH vs R-CHOP in Double/Triple-Hit DLBCL



Population Analysis: DA-EPOCH-R Era (Routine FISH) vs Historic Control



Baseline characteristics

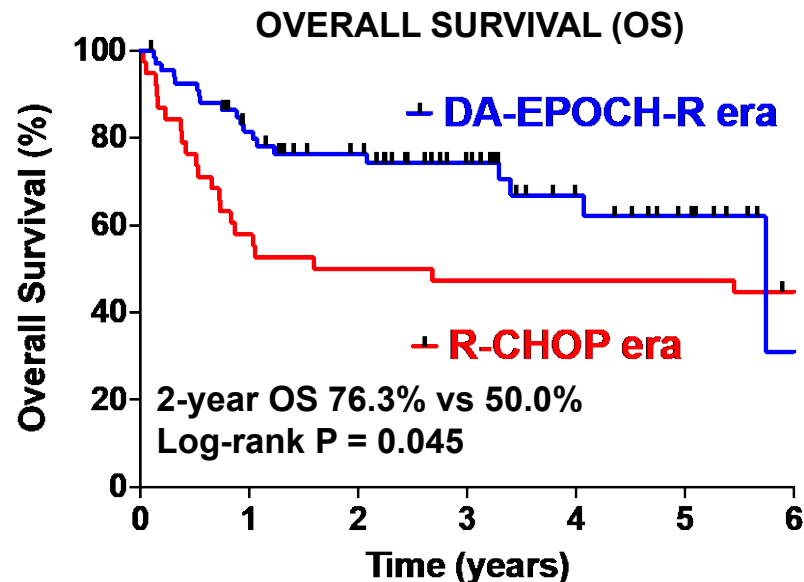
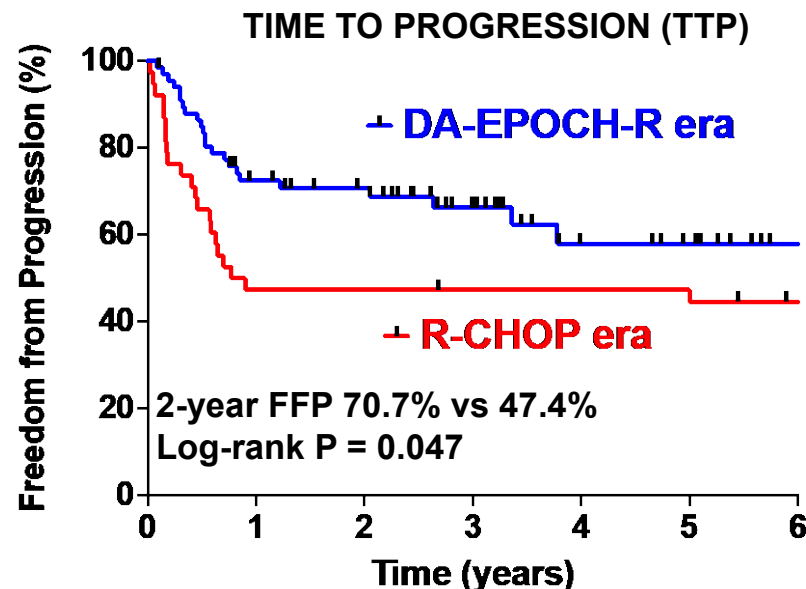
Characteristic	DA-EPOCH-R era 2015-2020 (n = 67)	R-CHOP era 2005-2010 (n = 38)	p
Age, Median (range, years)	64 (30-75)	63 (28-75)	0.89
Female (n, %)	24 (36)	14 (37)	1
Gene Rearrangements (n, %)			
MYC and BCL2 (double-hit)	52 (78)	29 (76)	1
MYC, BCL2 and BCL6 (triple-hit)	15 (22)	9 (24)	
Stage III/IV (n, %)	57 (85)	27 (71)	0.13
PS > 1 (n, %)	27 (40)	15 (39)	1
LDH > normal (n, %)	42 (63)	23 (61)	1
Extranodal sites >1 (n, %)	33 (49)	12 (32)	0.10
B symptoms (n, %)	31 (46)	16 (42)	0.84
Bulky disease ≥ 10 cm (n, %)	35 (52)	14 (37)	0.30
IPI risk group (n, %)			
Low (0-2)	22 (33)	18 (47)	0.15
High (3-5)	42 (63)	20 (53)	
Missing	3 (4)	0	
Treatment regimen (n, %)			
DA-EPOCH-R	47 (69)	0	
R-CHOP	16 (26)	32 (84)	
Highly Intensive*	3 (4)	4 (10)	
Palliative	1 (1.4)	2 (5)	

* CODOXMR/IVACR with consolidative autologous hematopoietic cell transplant.

PS: ECOG performance status, IPI: International Prognostic Index.



Era-on-era comparison: clinical outcomes

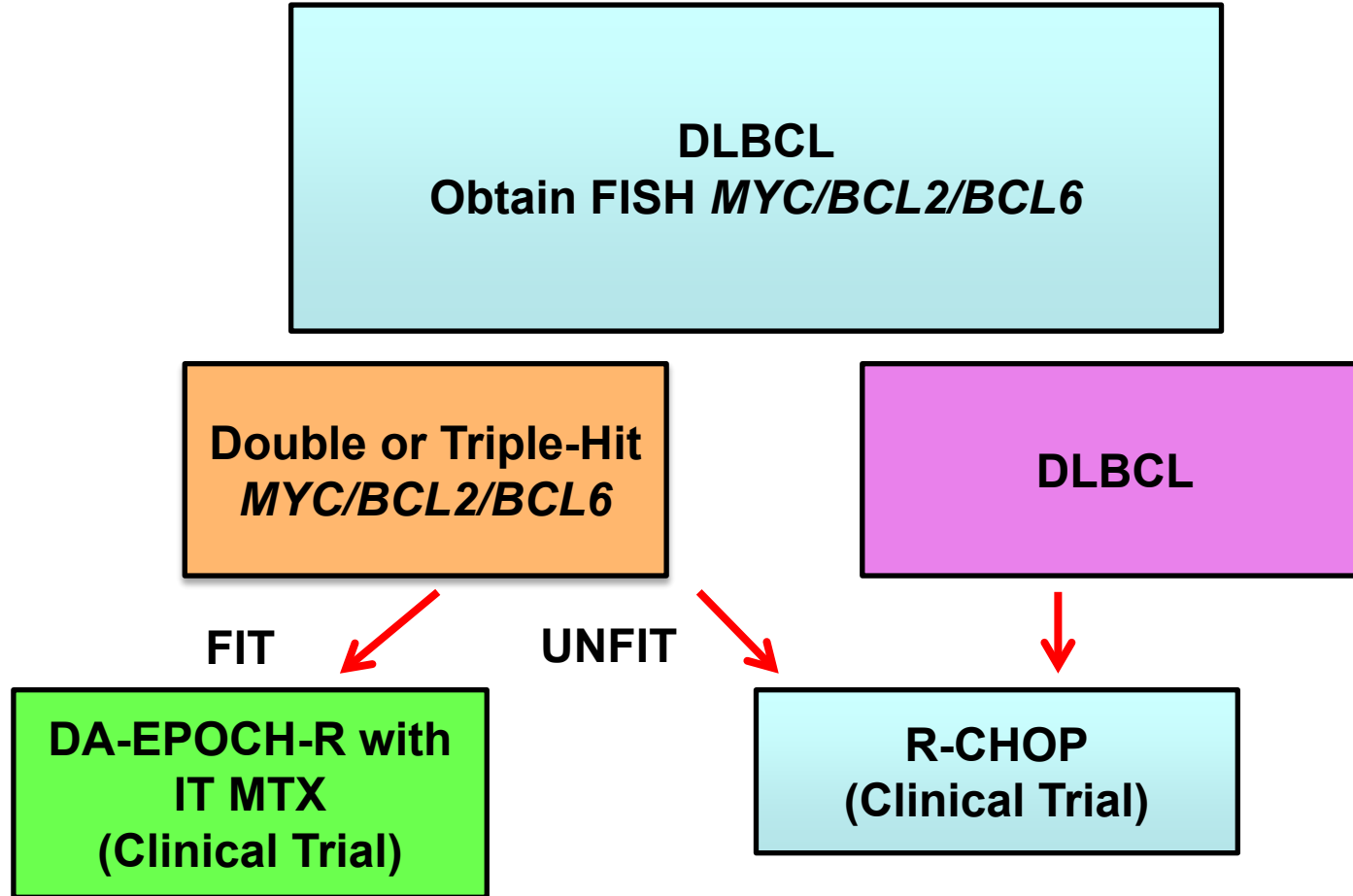


Cox model	DA-EPOCH-R era relative to R-CHOP era			
	TTP		OS	
	HR (95% CI)	p	HR (95% CI)	p
IPI low (0-2) vs high (3-5)	0.48 (0.26-0.89)	0.02	0.44 (0.23-0.83)	0.01
Individual IPI factors*	0.41 (0.21-0.82)	0.01	0.42 (0.21-0.85)	0.02

* Age >60 years, Stage III/IV, LDH>normal, PS>1 and extranodal sites>1. HR: adjusted Hazard Ratio, CI: Confidence interval

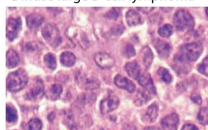


Treatment Algorithm for DLBCL

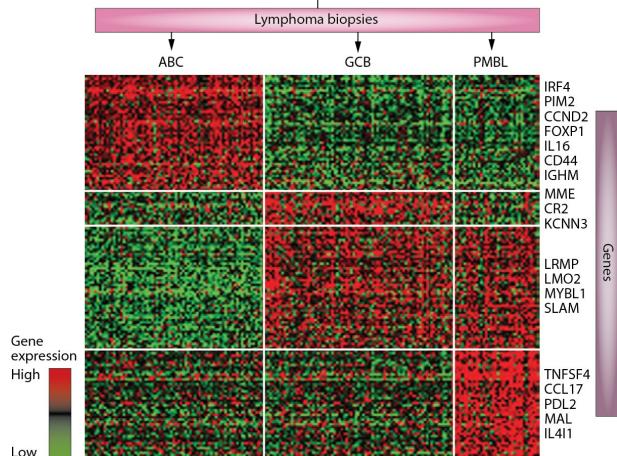


Prognosis According to Cell-of-Origin (ABC vs GCB) by GEP

Diffuse large B-cell lymphoma

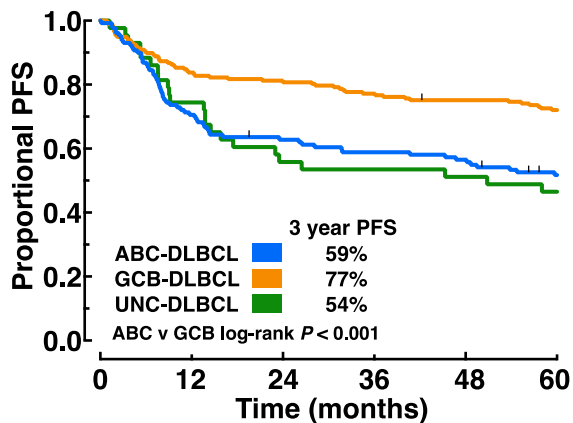


Lymphoma biopsies



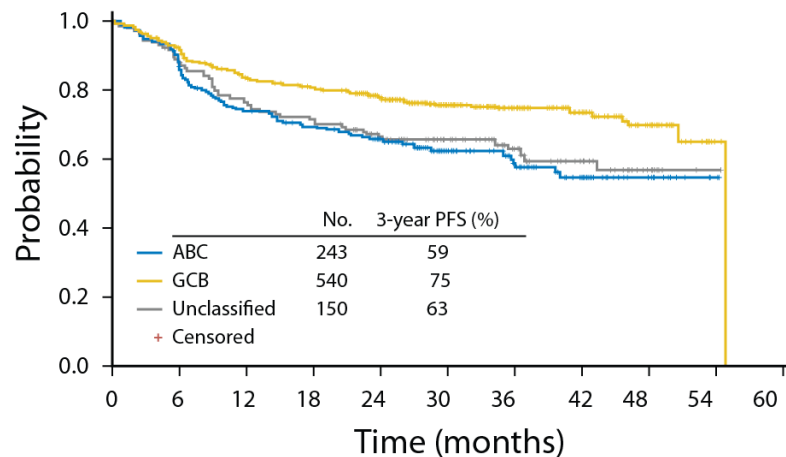
Alizadeh et al Nature 2000
Rosenwald et al, NEJM 2002
Lenz et al, NEJM 2008

15-20% difference in 3 year PFS



BC Cancer
2005 – 2010
R-CHOP treated

Unpublished Data



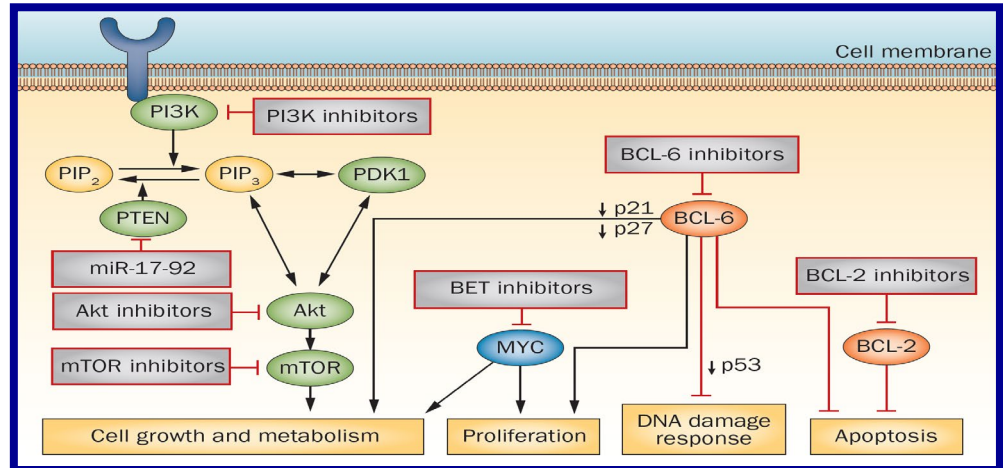
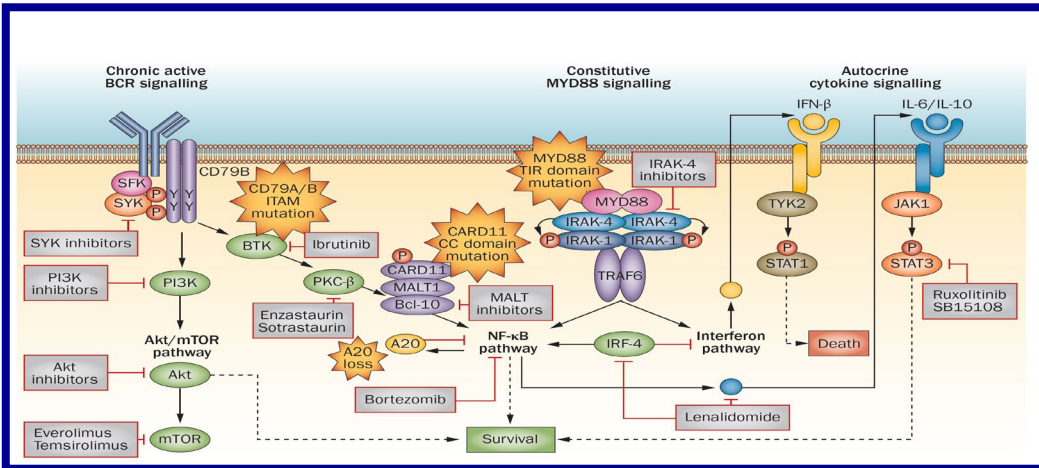
GOYA Trial
R-CHOP v G-CHOP
PFS including both arms

Vitolo et al J Clin Oncol 2017

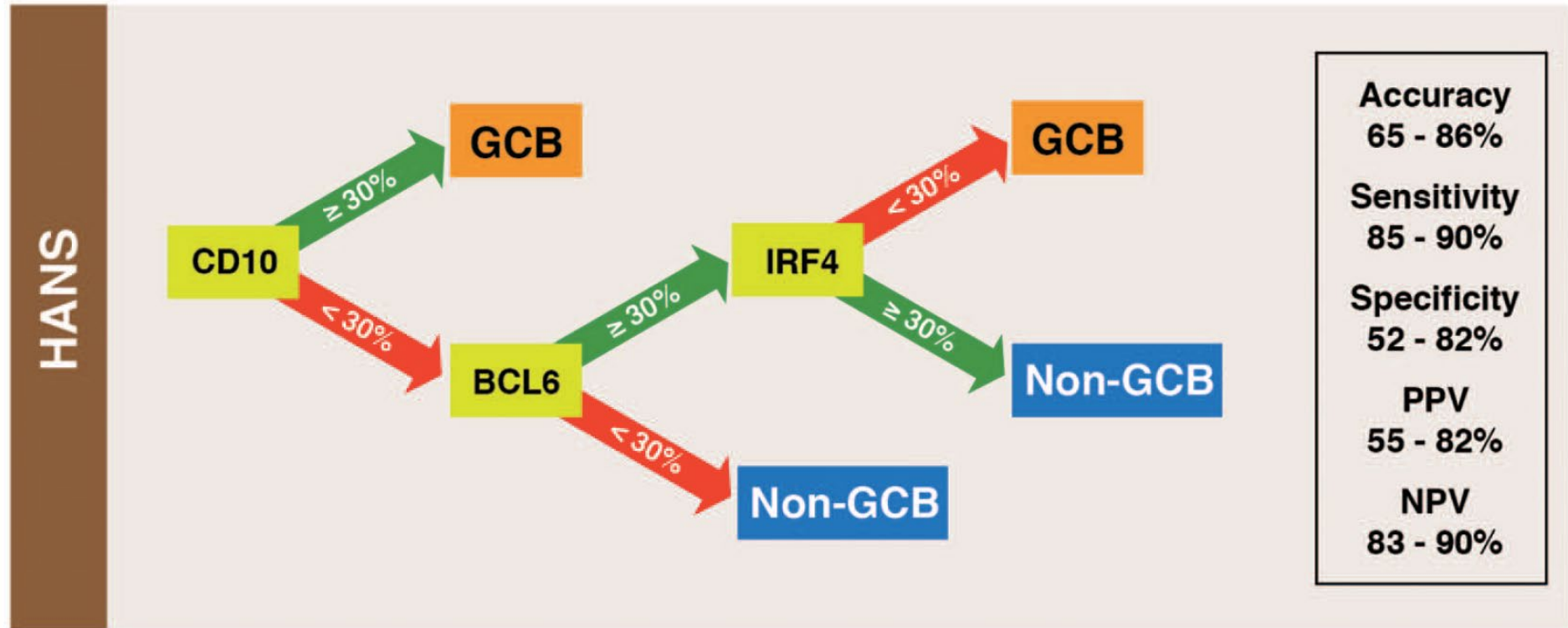
Distinct Signaling Pathways According to Cell-of-Origin & Potential Agents

ABC

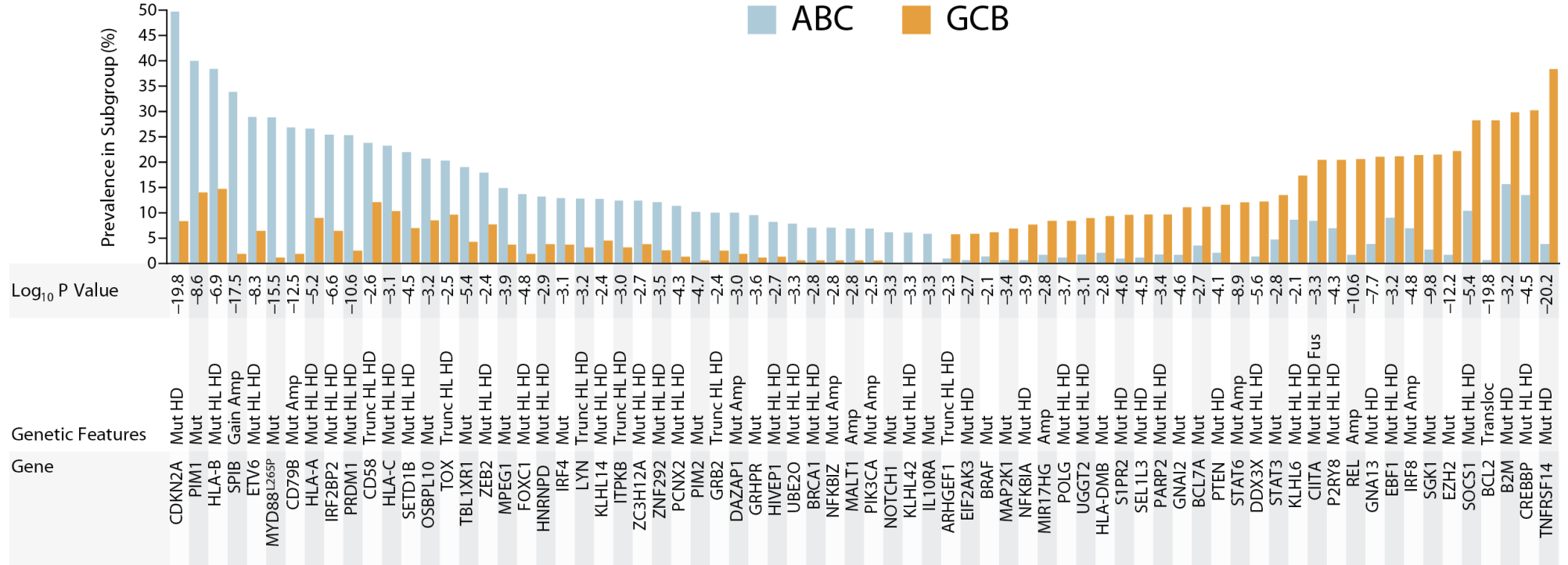
GCB



Using Immunohistochemistry to Assign COO



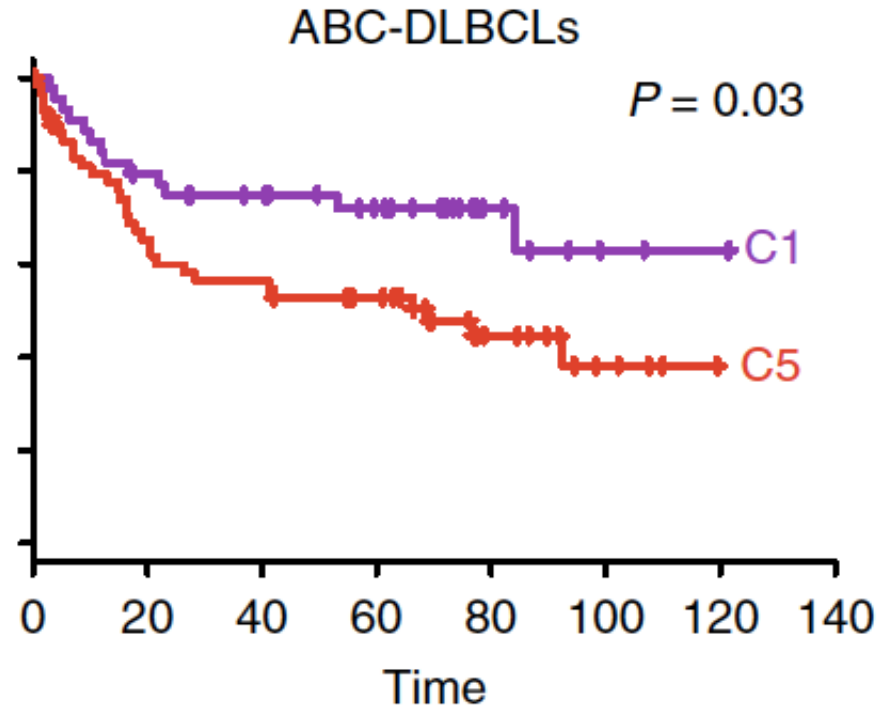
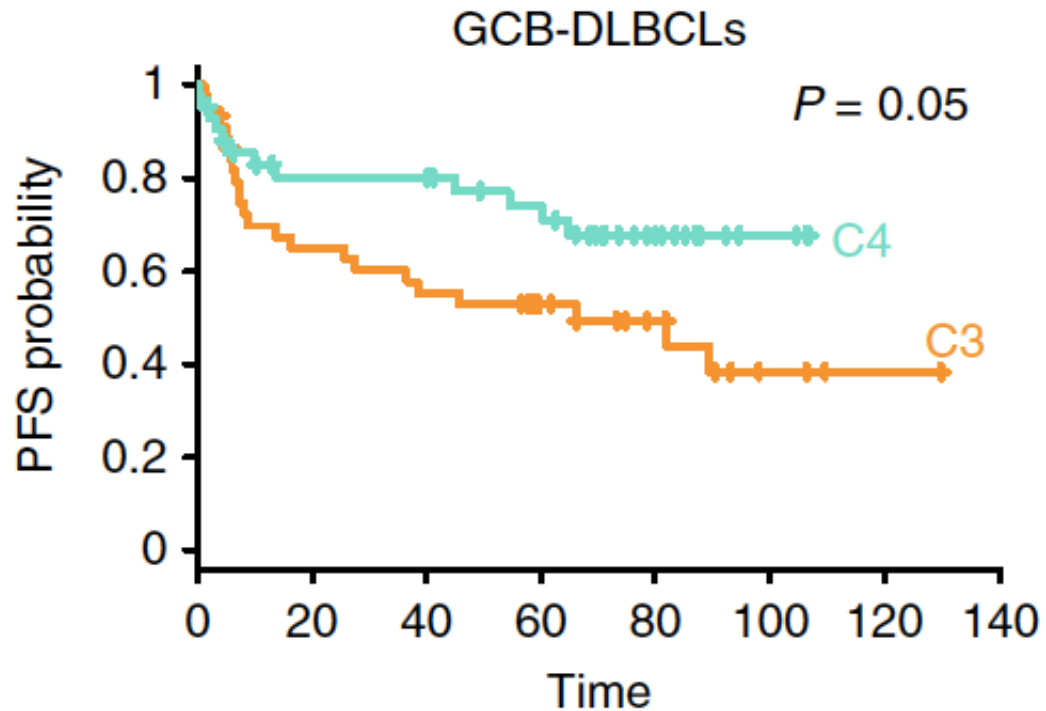
Cell-of-Origin – Distinct Mutational Landscapes



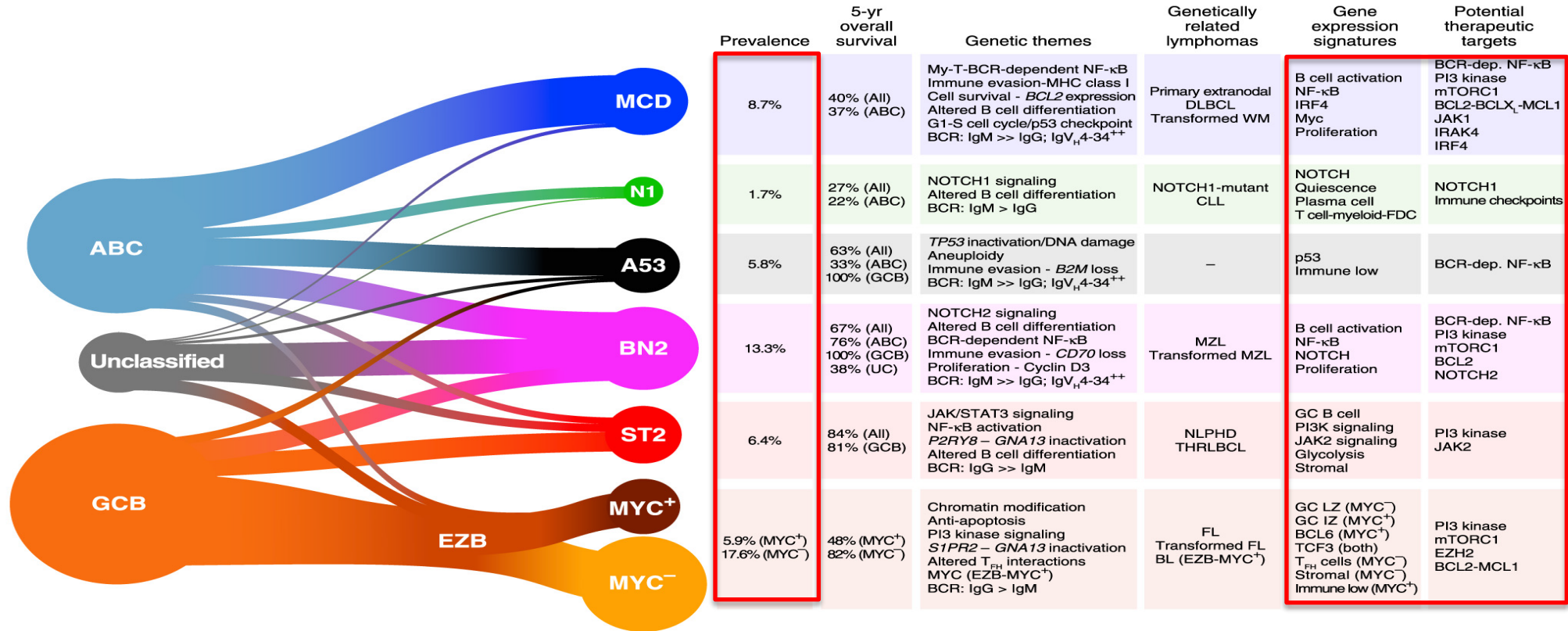
Heterogeneity within the COO subtypes is the likely explanation for variability in prognosis across populations

Schmitz et al N Eng J Med 2018

Novel Molecular Taxonomies of DLBCL Reveal Heterogeneity within COO Subtypes



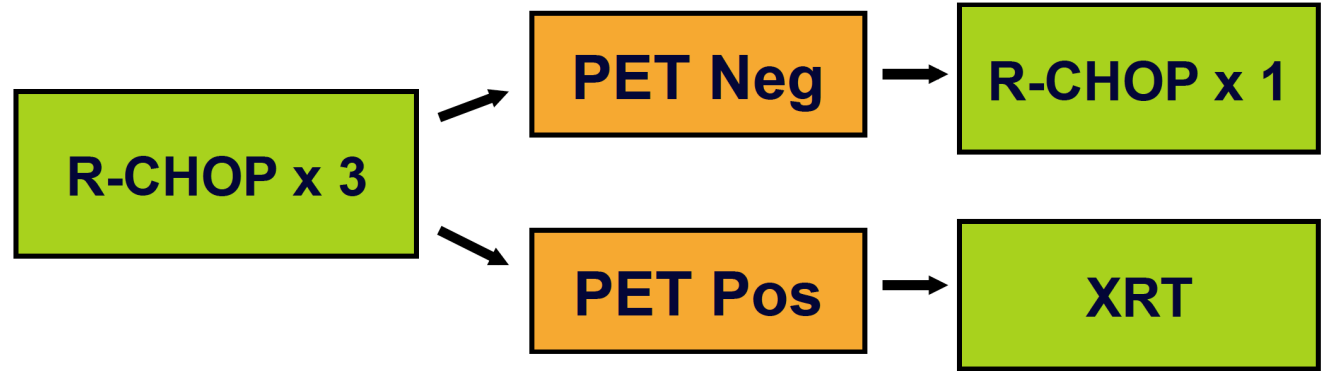
Novel subtypes within ABC and GCB DLBCL have been Identified



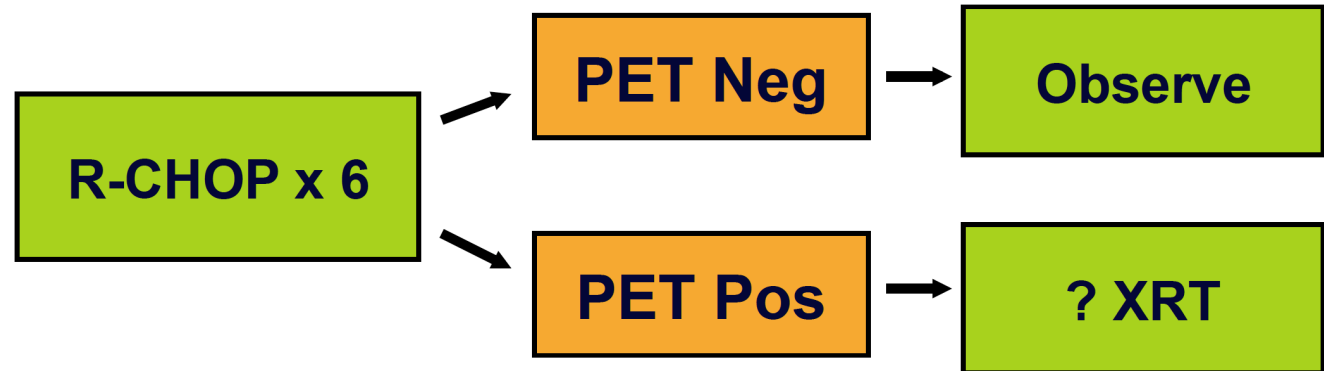
**Despite our growing understanding of
biology...we continue to treat patients with
DLBCL the same**

BC Cancer DLBCL Treatment Algorithm

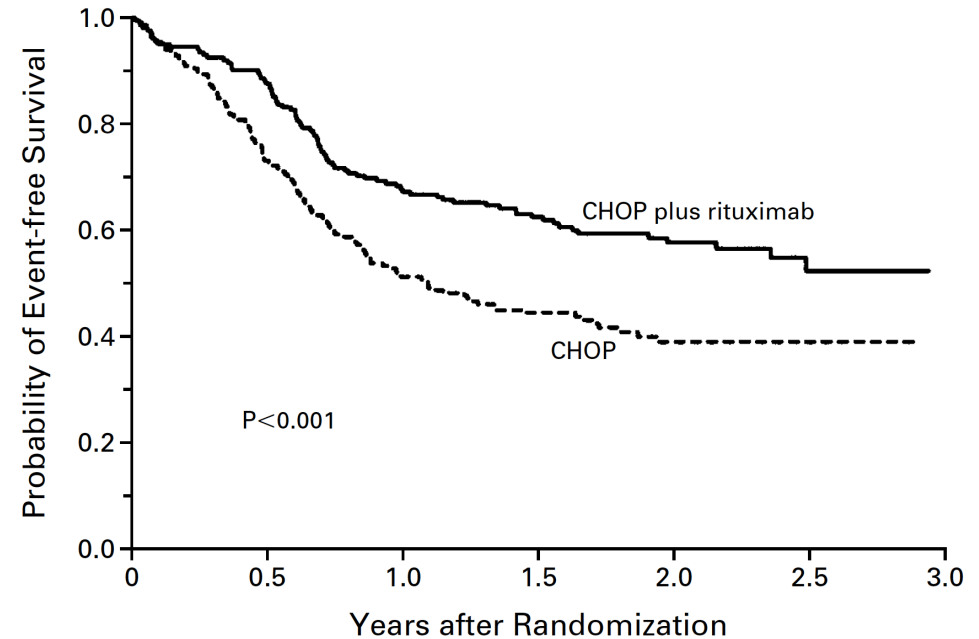
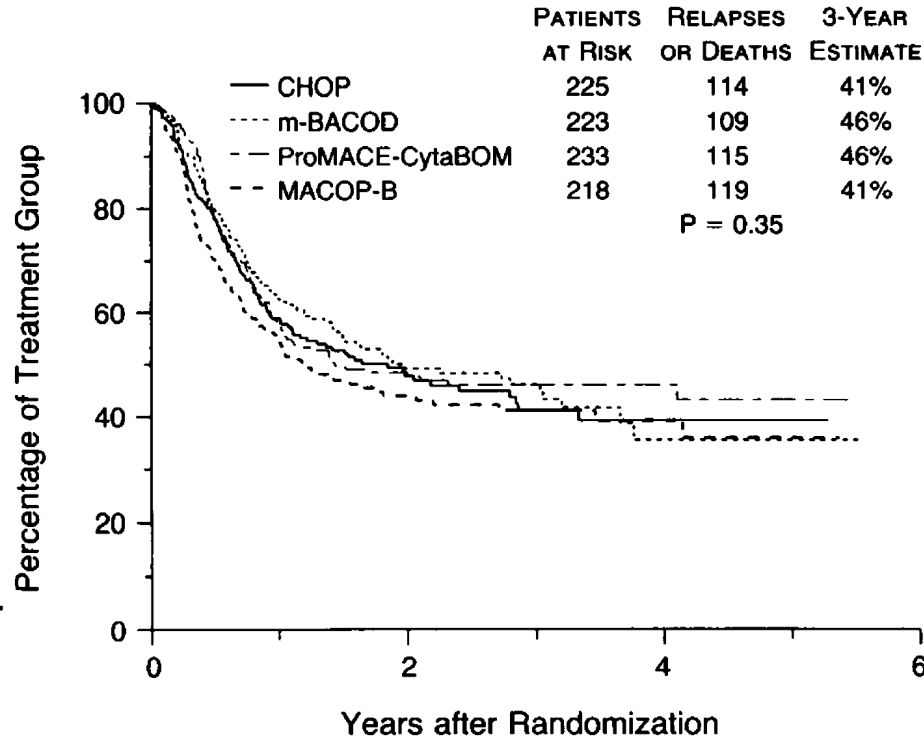
Limited Stage



Advanced Stage



R-CHOP Established as Standard of Care



Fisher et al, NEJM 1993

Coiffier, NEJM 2002

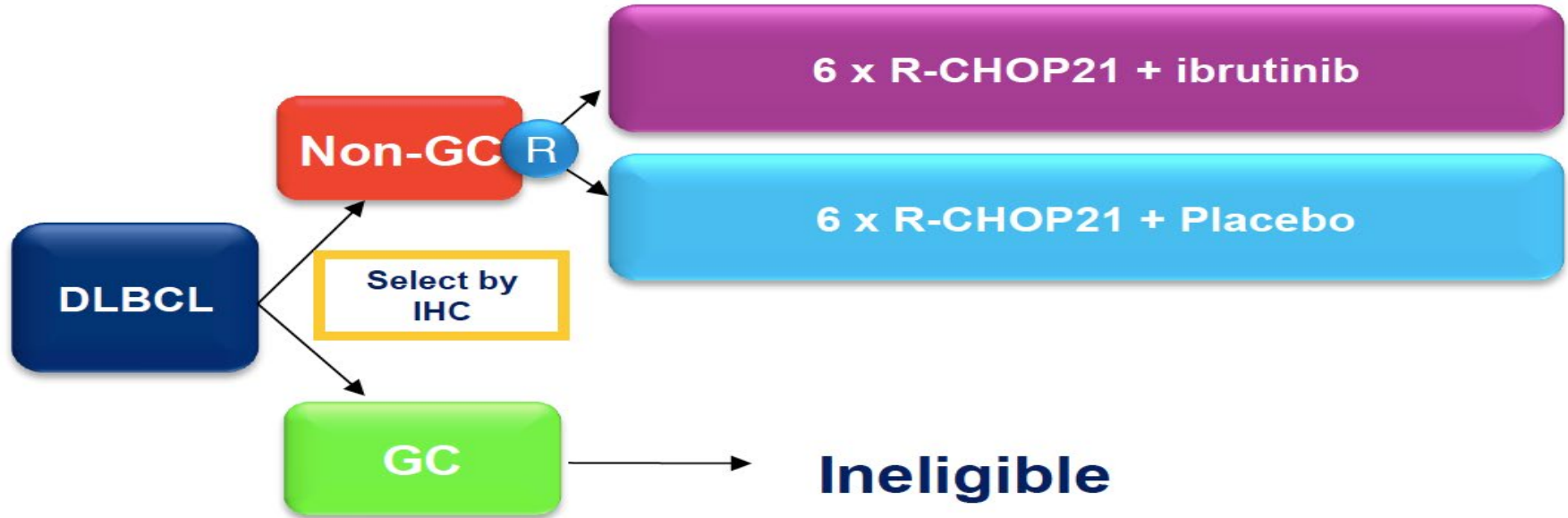
The Limit of Chemotherapy

Author	Therapy	Better than R-CHOP?
Cunningham, Lancet 2013	R-CHOP-14	No
Delarue, Lancet Oncol 2013	R-CHOP-14	No
Pfreundshuh, Lancet Oncol 2011	R-CHOEP	No
Recher, Lancet 2011	R-ACVBP	Yes (Age <60 y, IPI 1)
Wilson ASH 2016, Bartlett JCO 2019	DA-EPOCH-R	No
Le Gouill, ASCO 2011	ASCT v R-CHOP-14	No
Schmitz, Lancet Oncol 2012	R-Mega-CHOEP v R-CHOEP-14	No
Vitolo, ASH 2012 (#688)	ASCT v R-dose dense chemo	PFS Only
Stiff, NEJM 2013	ASCT v (R)-CHOP-21	PFS Only

Randomized Trials of Novel Agents

Author	Therapy	Better than R-CHOP
Leonard, JCO 2017	R-CHOP- Bortezomib	No
Davies, Lancet 2019	R-CHOP- Bortezomib	No
Vitolo, JCO 2017	Obinutuzumab-CHOP	No
Younes, JCO 2019	R-CHOP-Ibrutinib	? No
Nowakowski, JCO 2021	Lenalidomide-R-CHOP	? Yes (Phase II)
Nowakowski, JCO 2021	Lenalidomide-R-CHOP	No

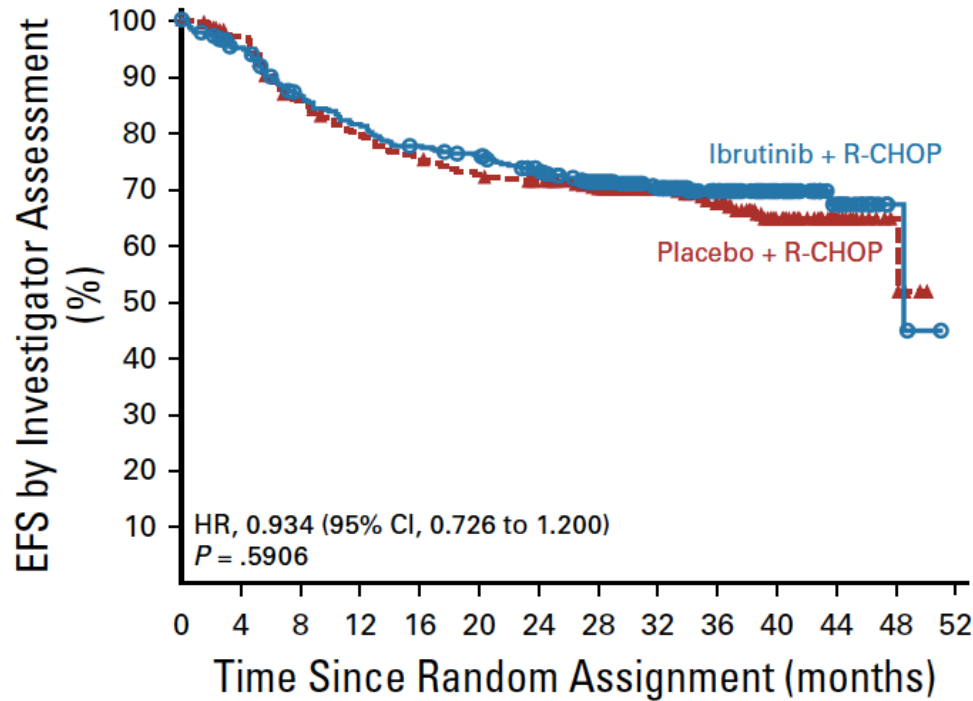
Phoenix Study: R-CHOP +/- Ibrutinib in Newly Diagnosed non-GCB DLBCL



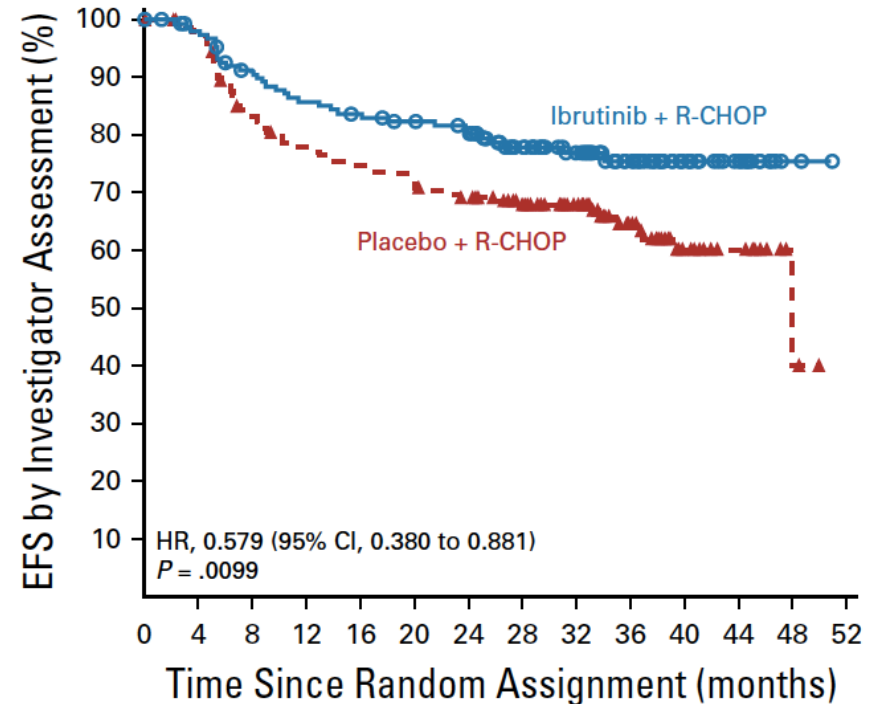
- Newly diagnosed DLBCL of non-GC
- ECOG PS ≤ 2 ; Age 18–80
- Primary Endpoint = EFS
- N = 800

* Ibrutinib 560 mg daily x 6 cycles
or placebo

Phoenix Study: R-CHOP +/- Ibrutinib in Newly Diagnosed non-GCB DLBCL



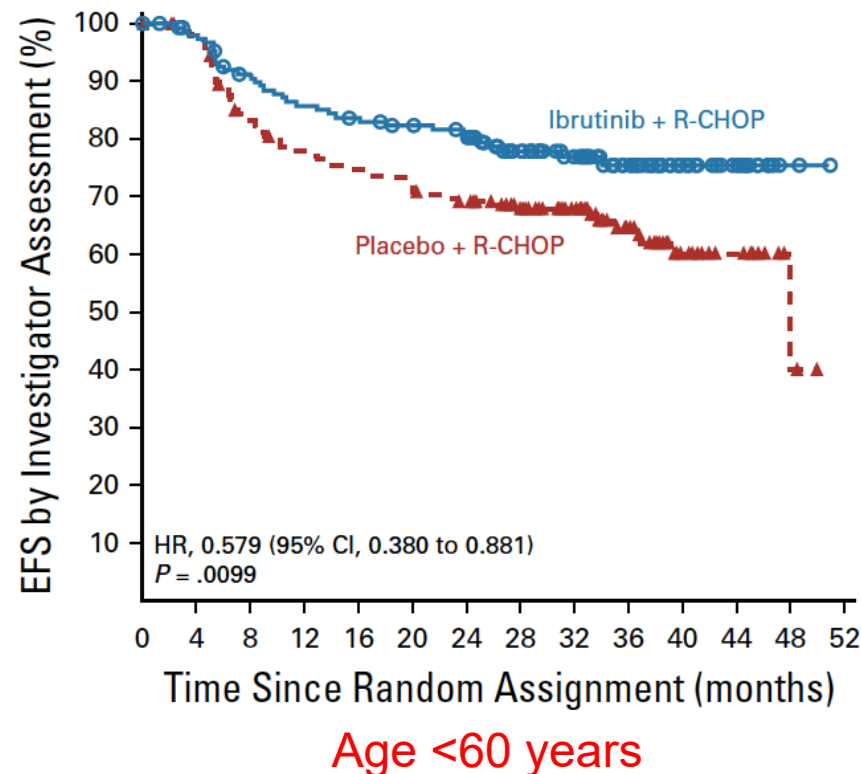
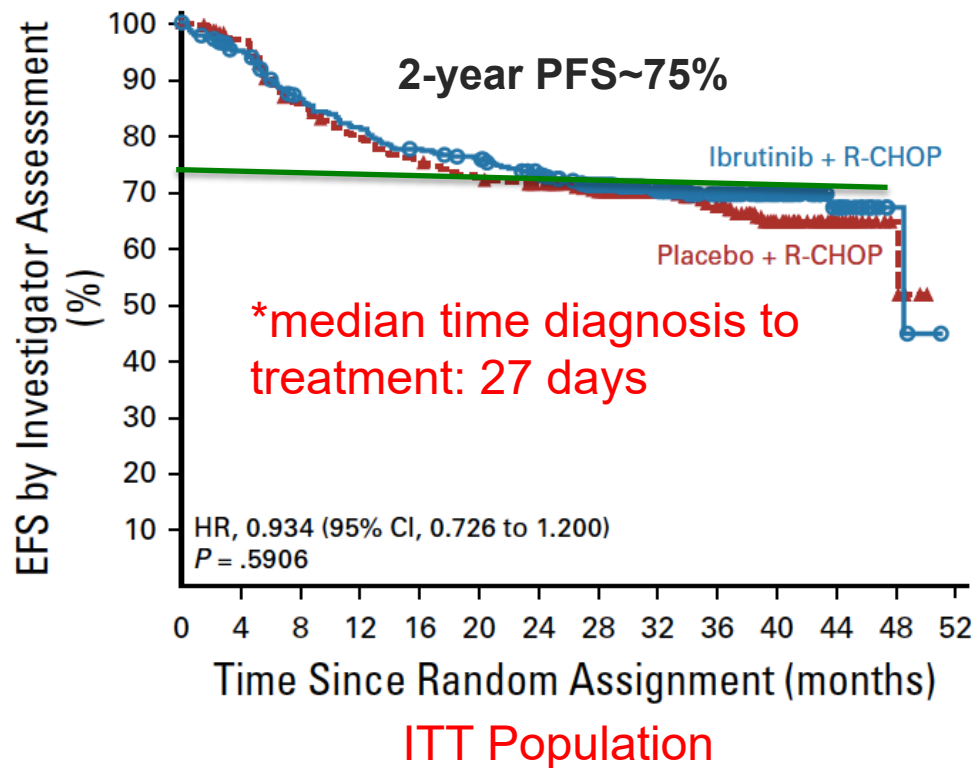
ITT Population



Age <60 years

Younes, A et al, JCO 2019

Phoenix Study: R-CHOP +/- Ibrutinib in Newly Diagnosed non-GCB DLBCL

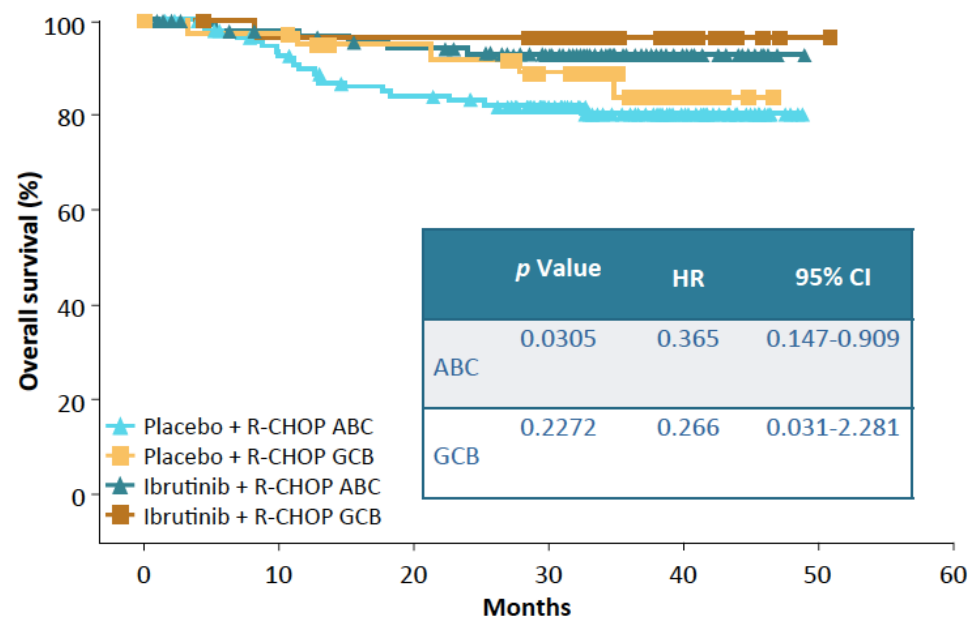


Concordance Between IHC and GEP

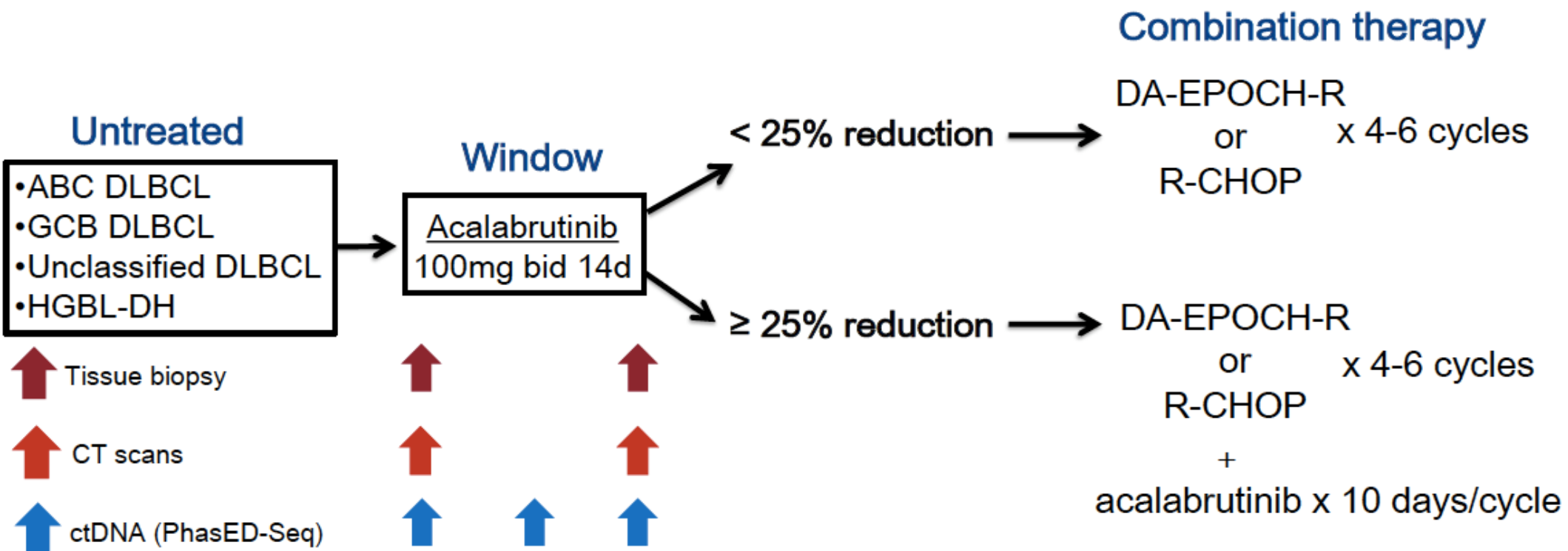
- 747 samples were evaluable from 838 enrolled patients
- 75.9% of enrolled (non-GCB by IHC) patients were ABC by GEP
- Non-GCB concordance = 82.7%

		EdgeSeq COO GEP			
		GCB	Non-GCB		Total
			ABC	UNC	
Hans-based IHC	Non-GCB, n (%)	129 (17.2)	567 (75.9)	51 (6.8)	747 (100)

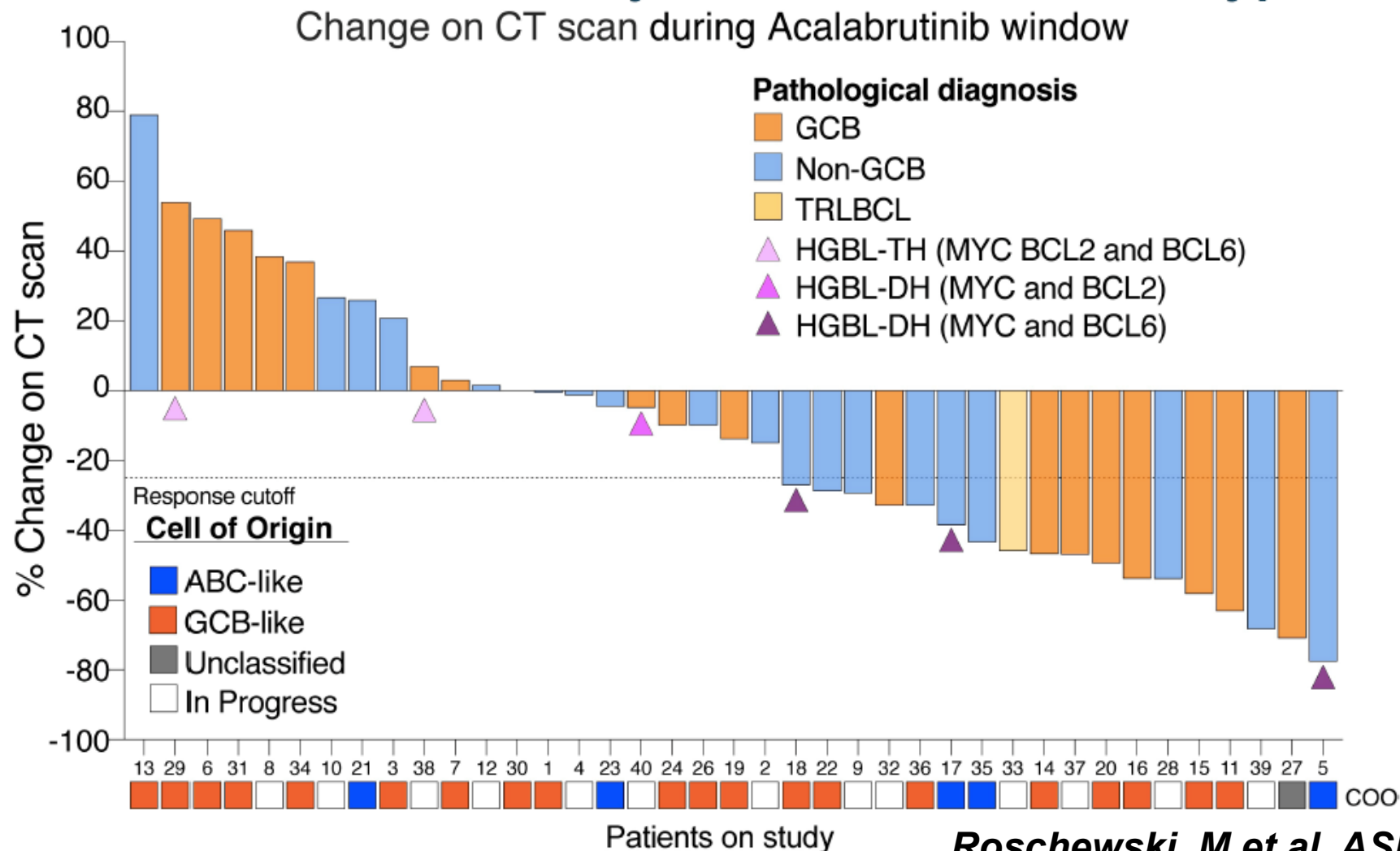
OS in Patients <60 years



Response-Adapted Acalabrutinib Window Study Design

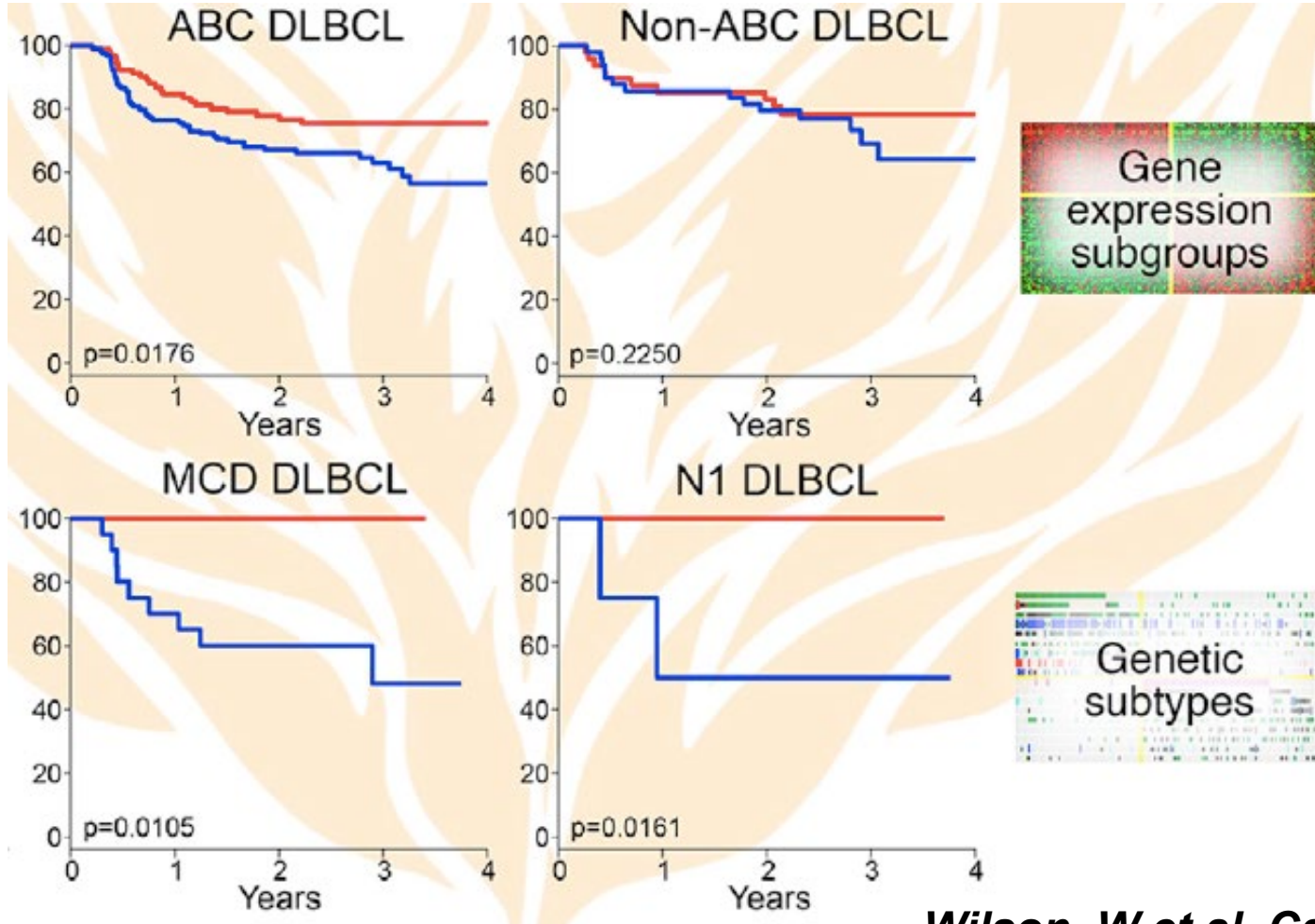


Acalabrutinib has Activity Across DLBCL Subtypes



Roschewski, M et al. ASH 2021

EFS According to GEP and Genetic Subtype

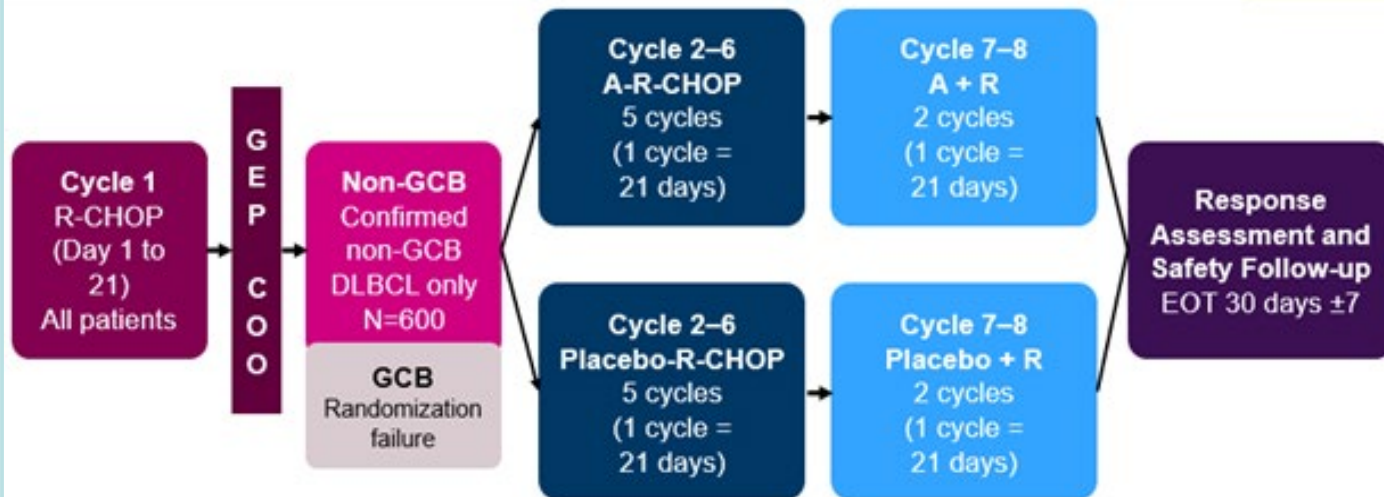


Learnings from PHOENIX informed the ESCALADE design

- Age ≤ 65 yo - instead of age-all comers
- COO by GEP – instead of IHC
- G-CSF – mandatory
- 1st R-CHOP cycle prior to randomization

Key Inclusion Criteria

- 18 - 65 years
- Histologically documented DLBCL
 - FFPE tumor tissue sample sent to the central laboratory prior to C1D1
 - Central laboratory confirmation by GEP of non-GCB subtype of DLBCL
- No prior treatment for DLBCL
- ECOG 0-2
- IPI 2-5
- Stage II-IV
- Measurable lesion by CT with contrast (or MRI)



Primary objective:

A+R-CHOP vs P+R-CHOP efficacy: INV assessed PFS

Key secondary objectives:

A+R-CHOP vs P+R-CHOP efficacy:

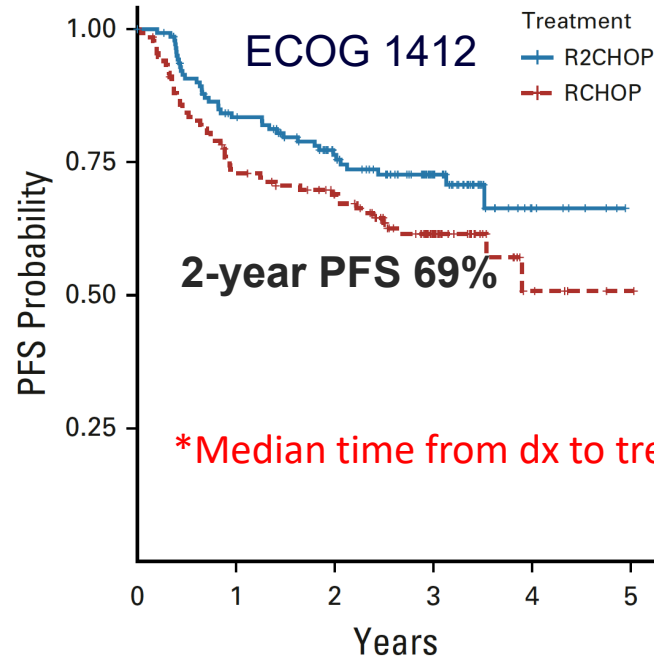
- INV assessed EFS
- BICR assessed CR rate at end of study treatment
- OS

Treatment and Duration:

- R-CHOP given every 21 days for 6 cycles starting on C1D1
- From Cycle 2 (randomized on Cycle 2 (C2D1)) to Cycle 6, pts with non-GCB DLBCL will receive either Acalabrutinib 100 mg twice daily plus R-CHOP or placebo plus R-CHOP
- Followed by 2 additional cycles of Rituximab + Acalabrutinib or placebo in Cycle 7 and Cycle 8, as 8 cycles of rituximab is recognized as a common standard per ESMO guidelines with 6 cycles of CHOP

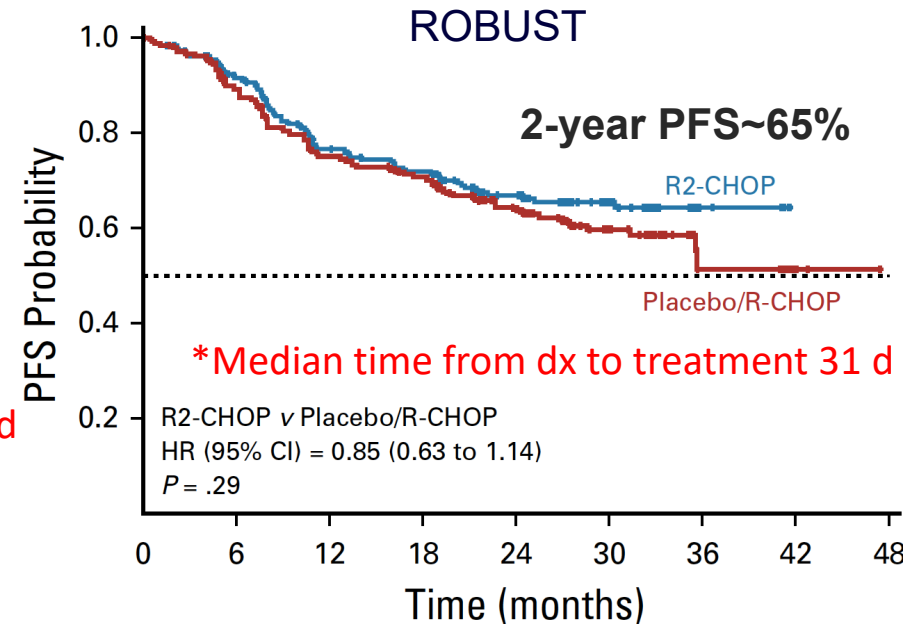
Interim and Final Analysis: One interim analysis for futility only and the final analysis will occur when **102** (45% of final) and **227**, respectively, INV-assessed PFS events combined in Arms A and B have been observed. IA is projected to occur 40 months after first subject randomized (FSR).

Lenalidomide-R-CHOP: ECOG 1412 Phase II and ROBUST Phase III



No. at risk:

R2CHOP	145	113	85	46	7	0
RCHOP	135	95	80	42	7	1

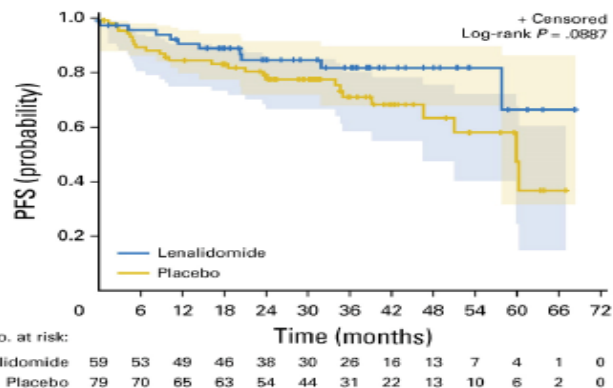


No. at risk:

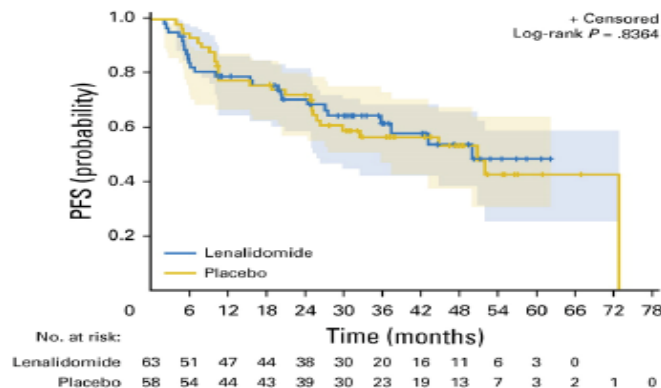
R2-CHOP	285	221	178	162	119	57	10	0
Placebo/R-CHOP	285	229	187	173	111	55	10	3

REMARC Trial: R-CHOP followed by Lenalidomide Maintenance in DLBCL

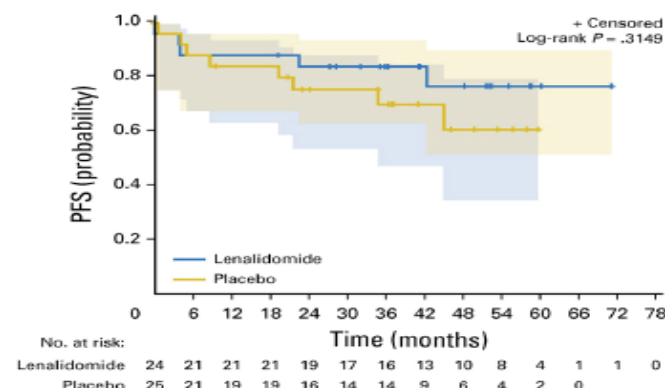
GCB



ABC

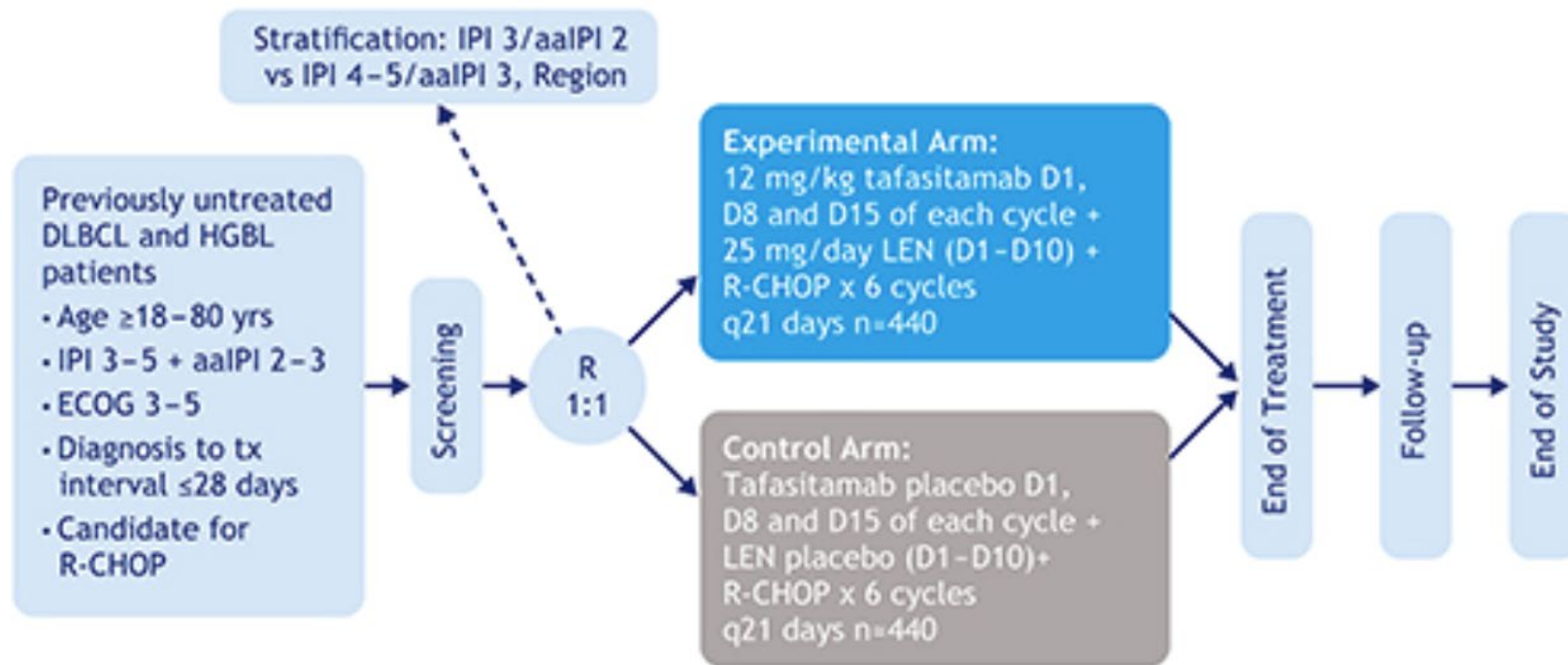


Unclassified



Front-MIND Trial: Tafa-Len-R-CHOP vs R-CHOP in DLBCL

Study Design



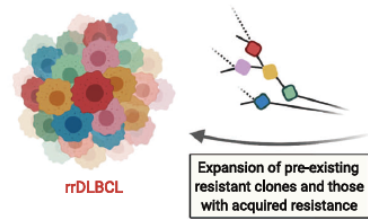
Challenges with Recent Phase 3 Trials

- **High bar to beat with R-CHOP**
- **Large patient numbers required**
- **Biomarker requirement caused delay in treatment leading to patient selection**
- **Highest risk patients excluded**
- **Biological heterogeneity despite patient enrichment**

Treatment Resistance in DLBCL

Tumor Heterogeneity

Genetic and/or epigenetic alterations in cancer cells generate spatial and temporal diversity to confer treatment resistance



Examples of genetic and/or epigenetic modifications in rDLBCL

Genetic modifications

Immune surveillance

B2M, CD58, HLA-A, MS4A1

Epigenetic regulation

EZH2, CREBBP, MEF2B, KMT2C, KMT2D

DNA damage response

TP53

Cell cycle regulation

CCND3, CDKN2A, CDKN2B

Signaling pathway activation

STAT6, SOCS1, FOXO1, MYD88, CD79B, NFKBIE, NFKBIZ

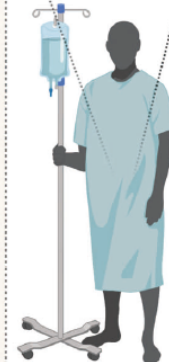
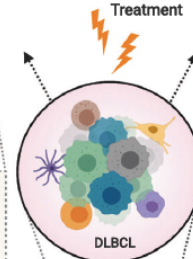
Oncogenes

MYC, PIM1, PRKCQ, GATA3, MLLT10, ABI1

Epigenetic modifications

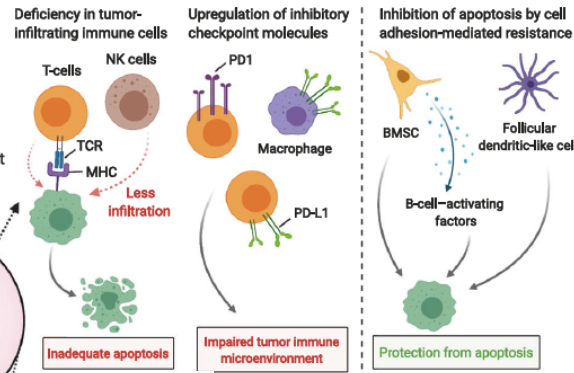
DNA methylation

Histone methylation/acetylation



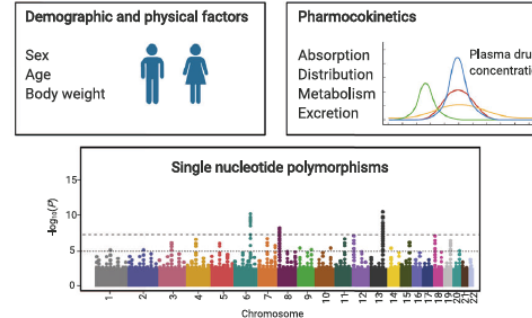
Tumor Microenvironment

Impaired immune function and supportive stromal cells promote a pro-tumor environment to mediate treatment resistance



Host Variabilities

Interpatient variabilities represented from several host-specific factors result in highly variable treatment responses



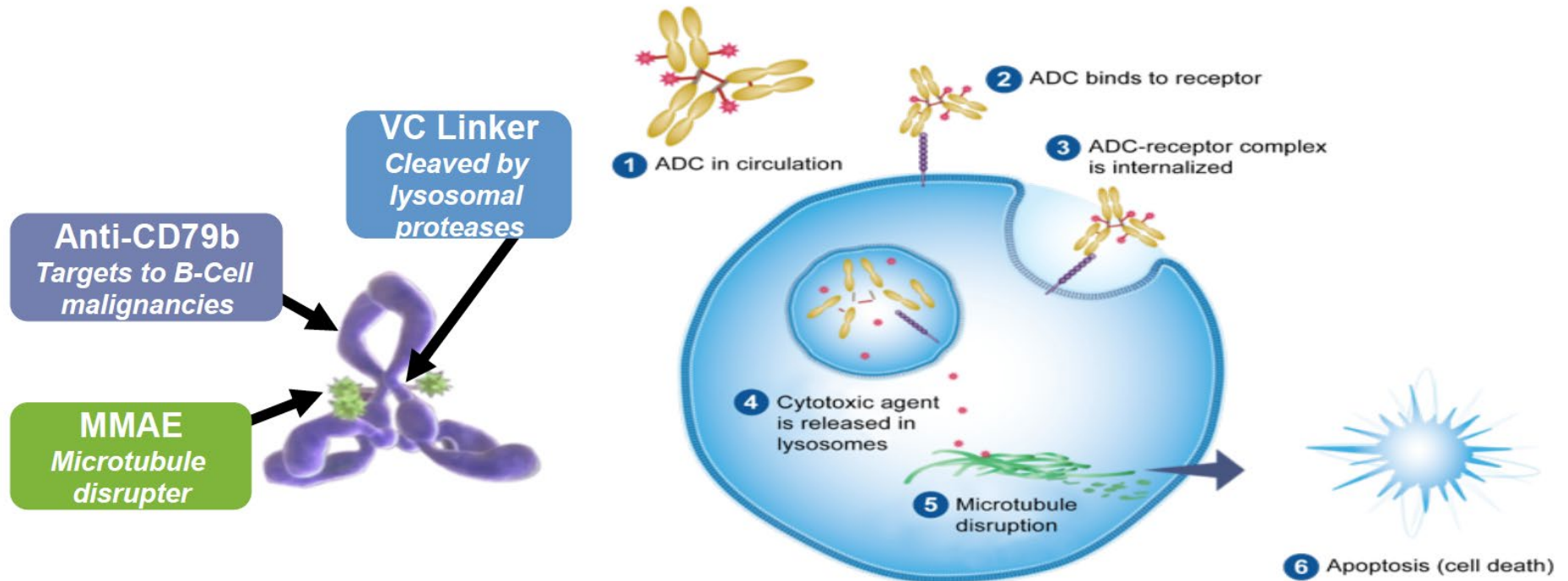
**Novel Agents have Emerged Allowing
Durable Disease Control**

Novel Agents Recently Approved in R/R DLBCL

	Pola-BR	Selinexor	Tafasitamab/Lenali domide	Loncastuximab Tesirine
MOA	Anti-CD79b ADC	XPO-1 inhibitor	Anti-CD19 MAb/Immunomodulat or	Anti-CD19 ADC
ORR	45%	28%	58%	48%
CR rate	40%	10%	40%	24%
PFS	9.2m	2.6m	11.6m	4.9m
DOR	12.6m	9.3m	43.9m	10.3m
OS	12.4m	NR	33.5m	9.9m

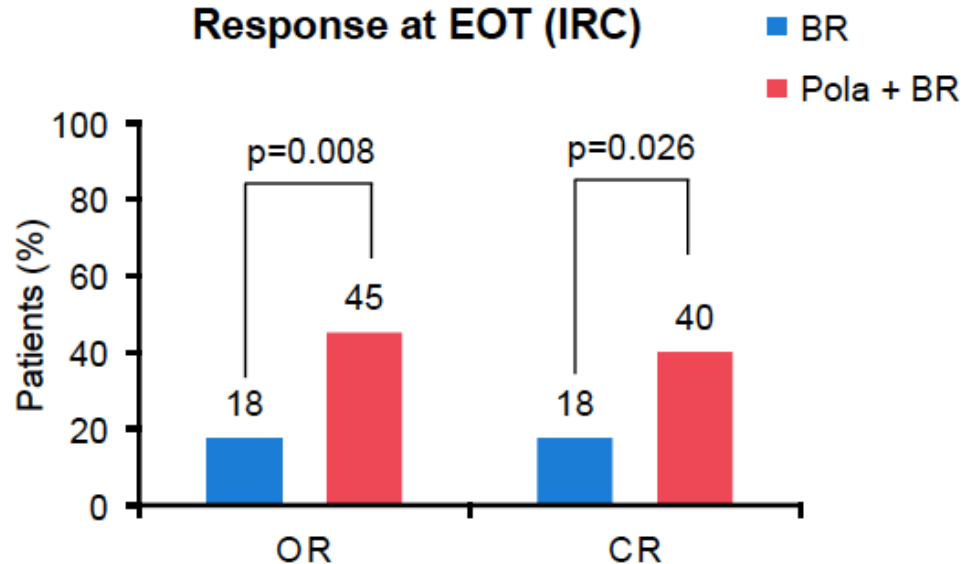
Polatuzumab Vedotin: Anti-CD79b Drug Conjugate

- Microtubule inhibitor MMAE conjugated to CD79b monoclonal antibody via a protease-cleavable peptide linker

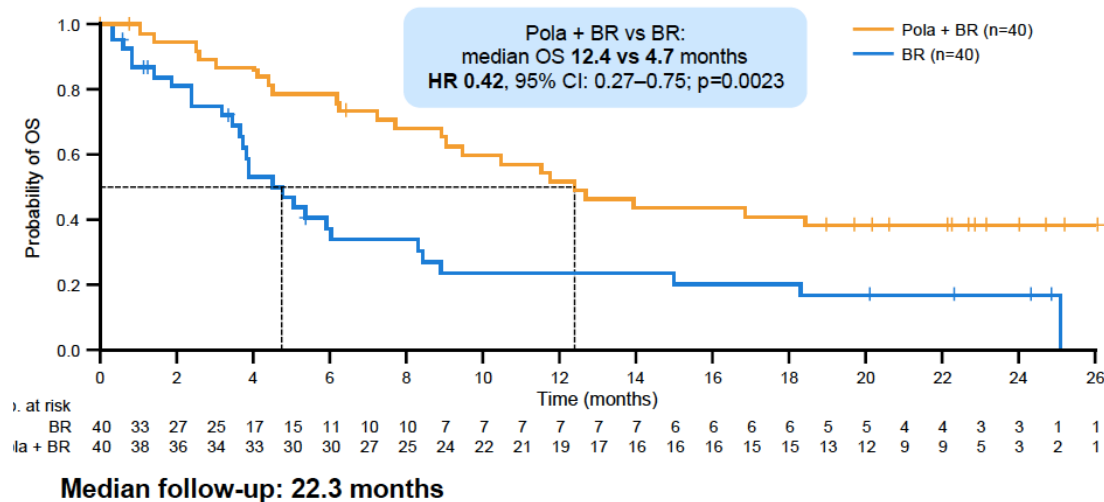


Randomized Phase II: Pola-BR vs BR

Response at EOT (IRC)



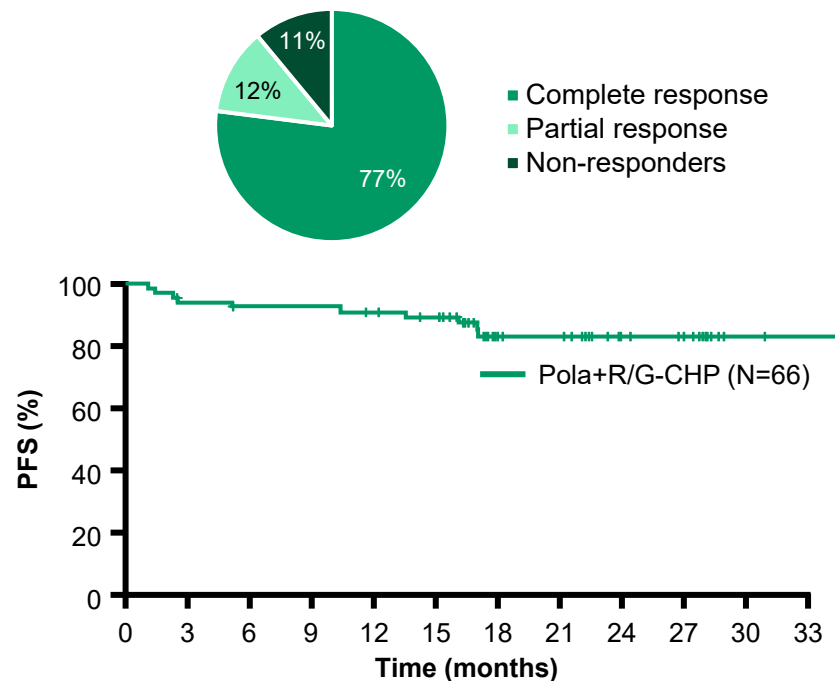
Overall Survival



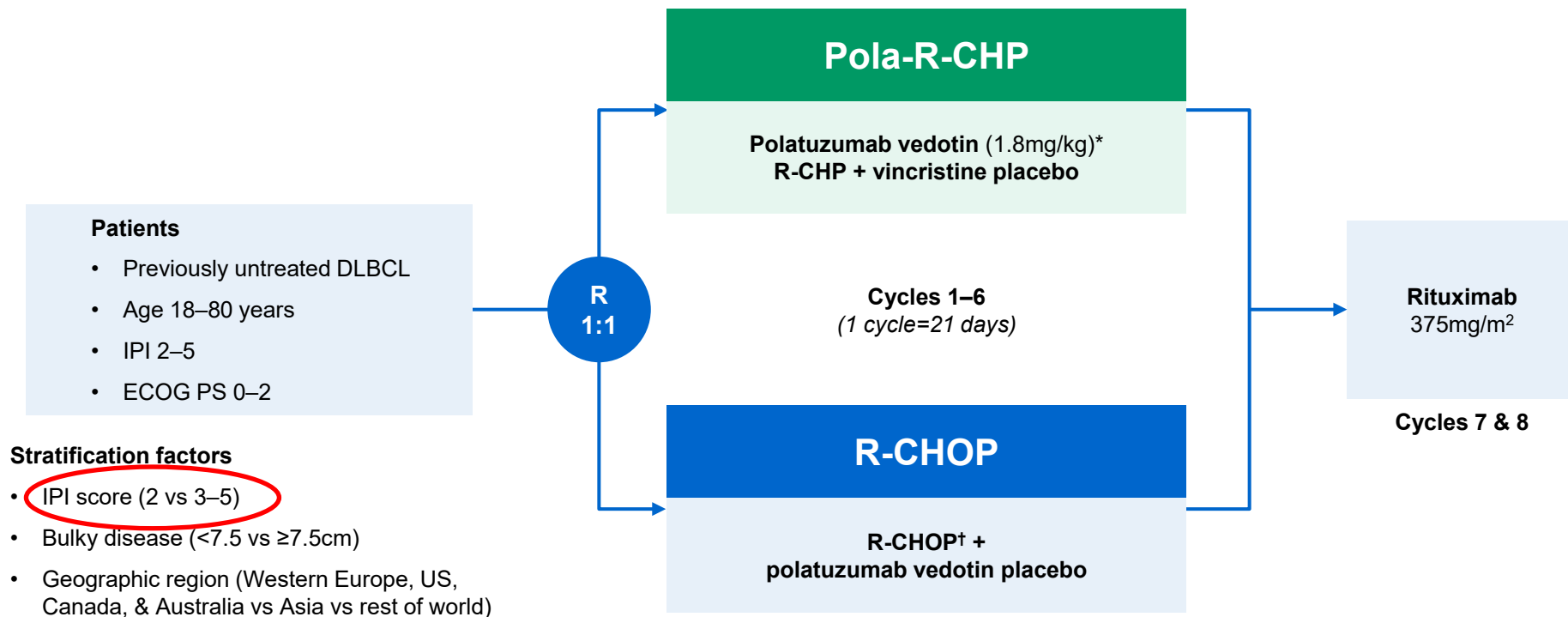
Pola + R/G-CHP in First-line DLBCL

- Open-label phase 1b/2 study
- Phase 2 population: DLBCL, IPI ≥ 2
- ORR: 89%; CR 77%
- Median f/up: 21.5 months
- 2-yr PFS: 83%

Pola+R/G-CHP demonstrated activity in first-line DLBCL



POLARIX: A randomized double-blinded study



*IV on Day 1; †R-CHOP: IV rituximab 375mg/m², cyclophosphamide 750mg/m², doxorubicin 50mg/m², and vincristine 1.4mg/m² (max. 2mg) on Day 1, plus oral prednisone 100mg once daily on Days 1–5.

IPI, International prognostic index; ECOG PS, Eastern Cooperative Oncology Group performance status; R, randomized.

Tilly H et al, NEJM 2021

POLARIX: Key endpoints and analysis timing

Key endpoints

Primary endpoint	Progression-free survival (Investigator-assessed)
Secondary endpoints	Event-free survival Complete response rate at end of treatment (PET/CT, IRC-assessed) Disease-free survival Overall survival
Safety endpoints	Incidence, nature, and severity of adverse events

Statistical design and timing of primary analysis:

- 875 patients, **all on study for ≥ 24 months** with approximately 228 PFS events, were required for the primary analysis. This occurred on June 28, 2021 (clinical cut-off date)
- Median follow up at the primary analysis was **28.2 months**

Baseline characteristics

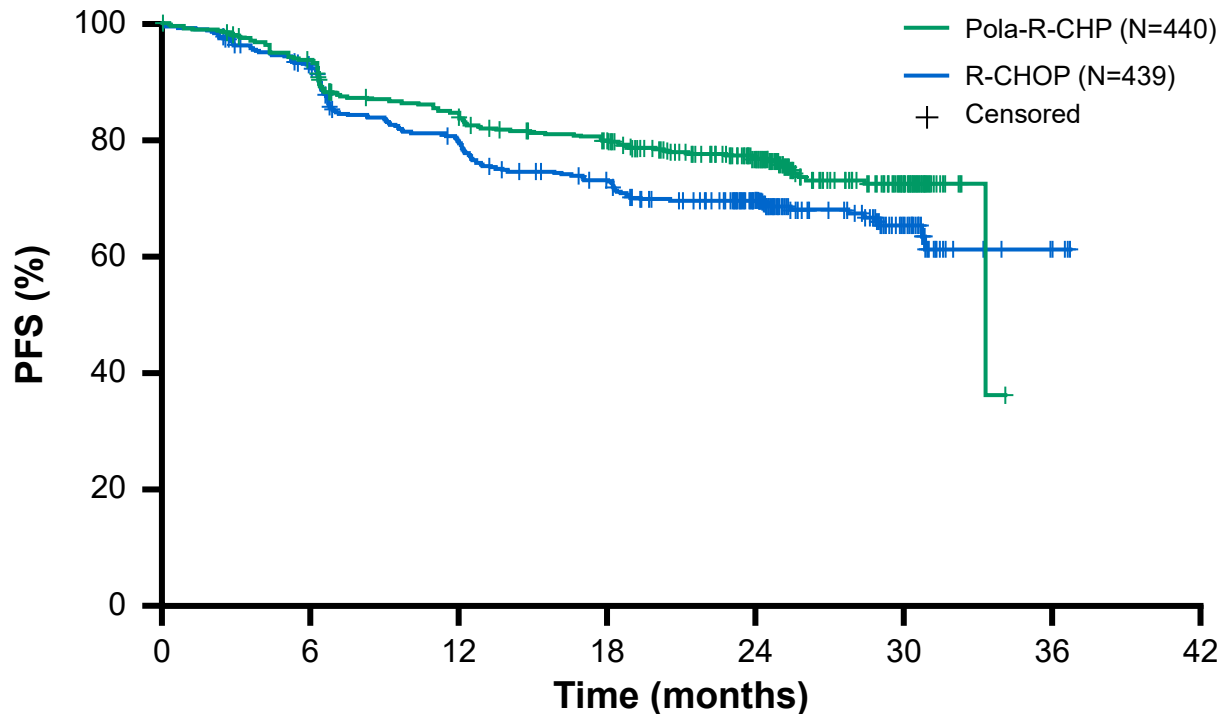
ITT population		Pola-R-CHP (N=440)	R-CHOP (N=439)
Age	Median (range), years	65.0 (19–80)	66.0 (19–80)
Sex, n (%)	Male	239 (54)	234 (53)
ECOG PS, n (%)	0–1	374 (85)	363 (83)
	2	66 (15)	75 (17)
Bulky disease (≥ 7.5 cm), n (%)	Present	193 (44)	192 (44)
Elevated LDH, n (%)	Yes	291 (66)	284 (65)
Time from diagnosis to treatment initiation	Median, days	26	27
Ann Arbor Stage, n (%)	III–IV	393 (89)	387 (88)
Extranodal sites, n (%)	≥ 2	213 (48)	213 (49)
IPI score, n (%)	2	167 (38)	167 (38)
	3–5	273 (62)	272 (62)
Cell-of-origin, (%)*	ABC	102 (31)	119 (35)
	GCB	184 (56)	168 (50)
	Unclassified	44 (13)	51 (15)
MYC/BCL2 expression, n (%)*	Double expression	139 (38)	151 (41)
MYC/BCL2/BCL6 rearrangement, n (%)*	Double-/triple-hit	26 (8)	19 (6)

*In the Pola-R-CHP and R-CHOP groups, respectively, the numbers of patients evaluable for cell-of-origin were 330 and 338, with IHC for MYC/BCL2 expression were 362 and 366, and with FISH for MYC/BCL2/BCL6 rearrangements were 331 and 334.

ABC, activated B-cell; FISH, fluorescence in situ hybridization; GCB, germinal center B-cell; LDH, lactate dehydrogenase.

Primary endpoint: Progression-free survival

Pola-R-CHP significantly improved PFS versus R-CHOP



HR 0.73 (P<0.02)

95% CI: 0.57, 0.95

- Pola-R-CHP demonstrated a **27% reduction in the relative risk of disease progression, relapse, or death** versus R-CHOP
- **24-month PFS:**
76.7% with Pola-R-CHP versus
70.2% with R-CHOP ($\Delta=6.5\%$)

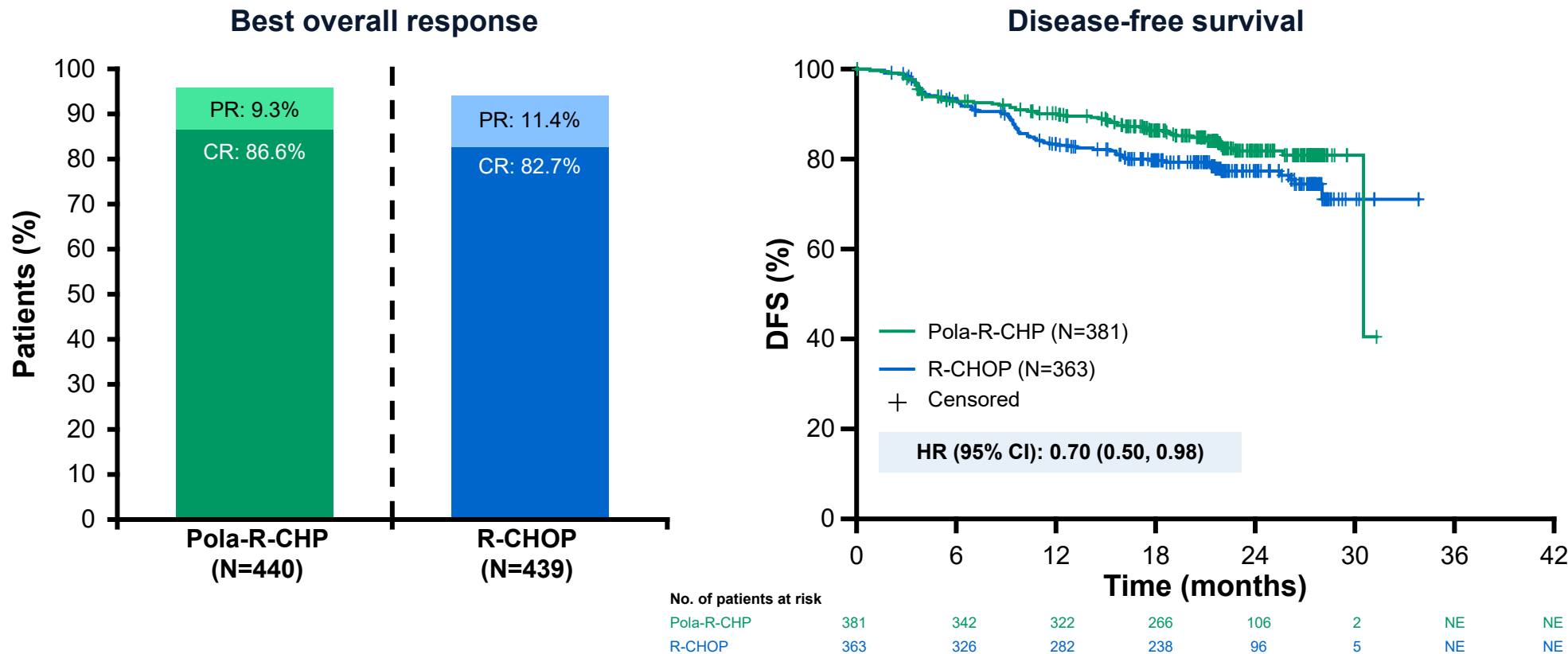
No. of patients at risk

Pola-R-CHP	440	404	353	327	246	78	NE	NE
R-CHOP	439	389	330	296	220	78	3	NE

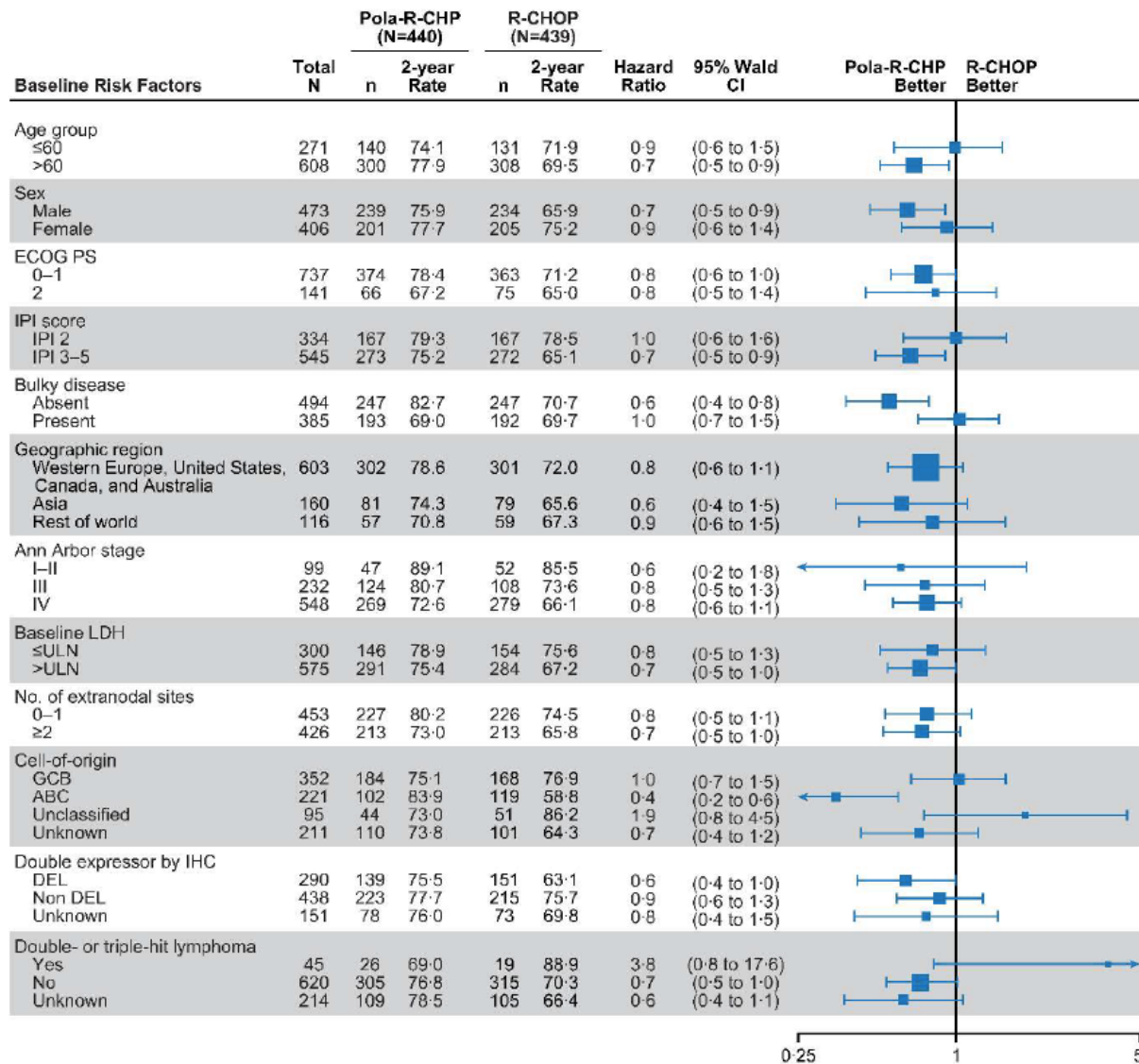
ITT population. Data cut-off: June 28, 2021; median 28.2 months' follow-up.

NE, not evaluable.

Response rates and disease-free survival



ITT population. Data cut-off: June 28, 2021; median 28.2 months' follow-up. Disease-free survival (DFS) defined as the time from the date of the first occurrence of a documented complete response to the date of progression, relapse, or death from any cause for the subgroup of patients with a best overall response of CR.



? Benefit

Younger ≤ 60y

Females

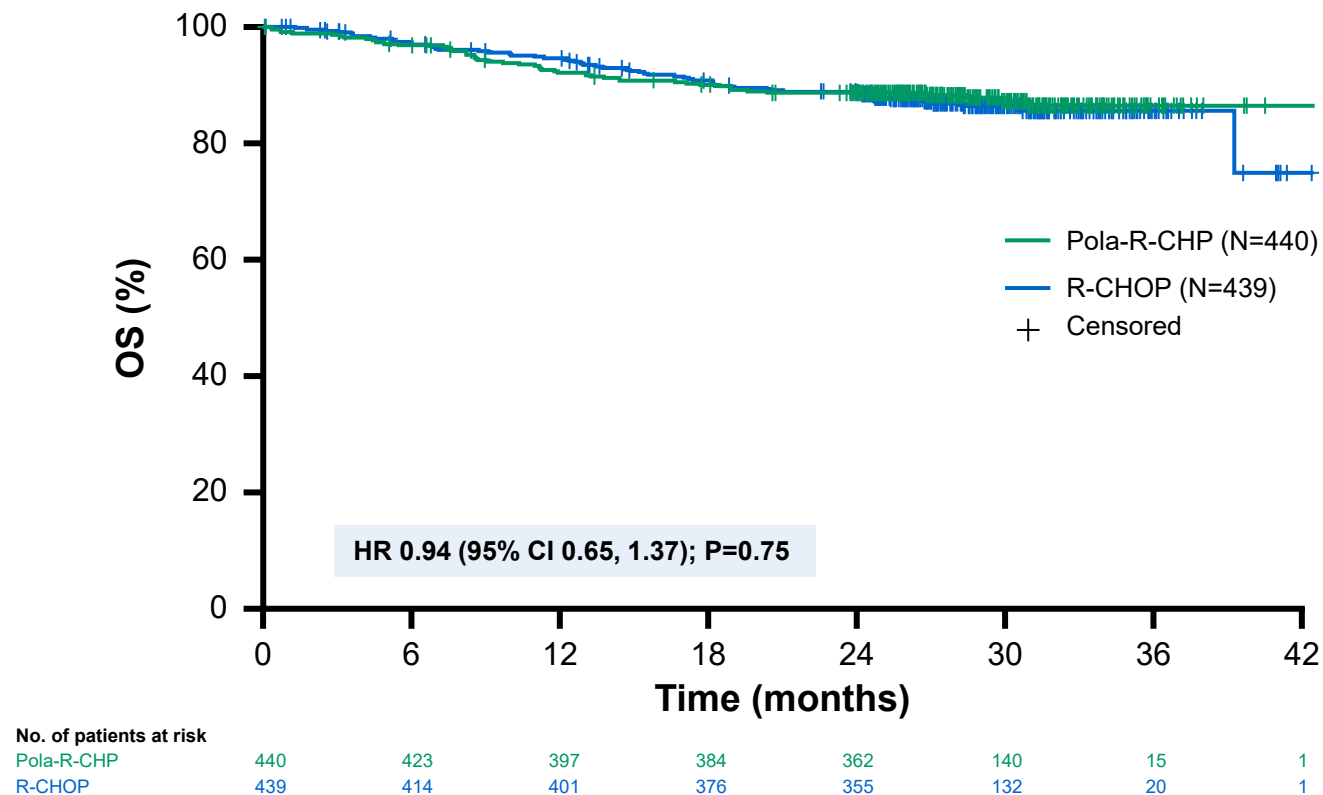
IPI = 2

Bulk ≥ 7.5 cm

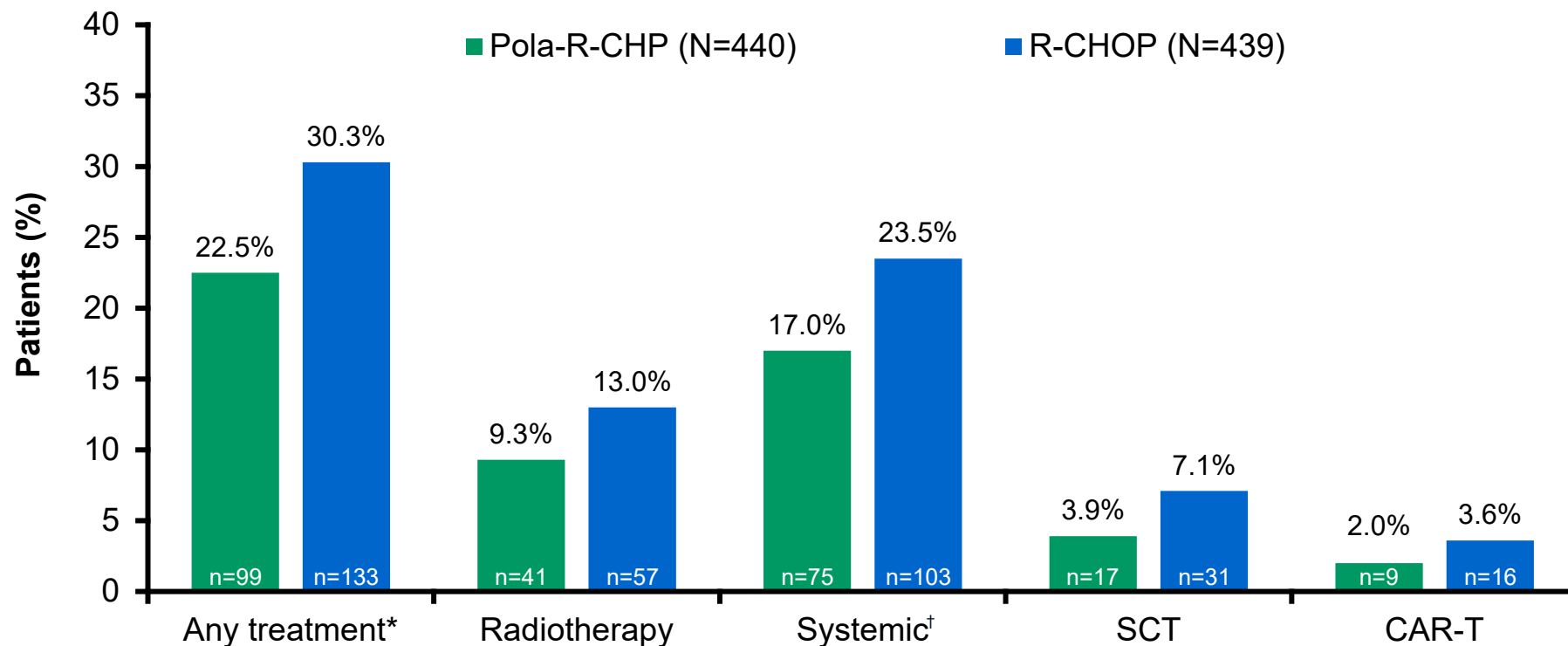
GCB Subtype

DH/TH lymphoma

Overall survival



Patients receiving subsequent treatments



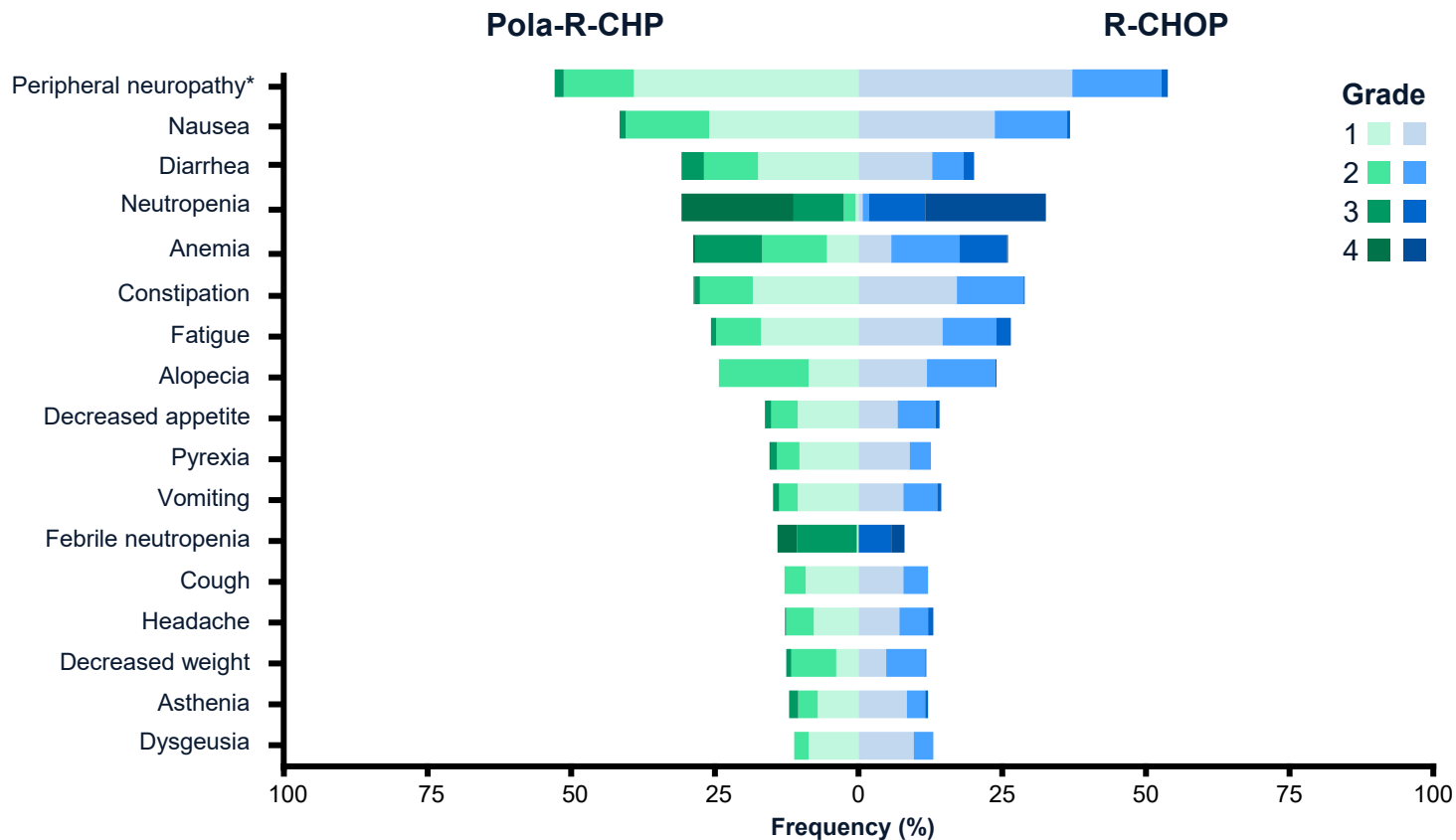
Data cut-off: June 28, 2021. *Subsequent lymphoma treatment was defined as non-protocol anti-lymphoma therapy; †Includes any monotherapy, multi-drug, or cell-based regimen. CAR-T, chimeric antigen receptor T-cell therapy; SCT, stem cell transplant.

Safety summary

Safety profiles were similar with Pola-R-CHP and R-CHOP

n (%)	Pola-R-CHP (N=435)	R-CHOP (N=438)
Any-grade adverse events	426 (97.9)	431 (98.4)
Grade 3–4	251 (57.7)	252 (57.5)
Grade 5	13 (3.0)	10 (2.3)
Serious adverse events	148 (34.0)	134 (30.6)
Adverse events leading to:		
Discontinuation of any study drug	27 (6.2)	29 (6.6)
Polatuzumab vedotin / vincristine	19 (4.4)	22 (5.0)
Dose reduction of any study drug	40 (9.2)	57 (13.0)

Common adverse events



Data cut-off: June 28, 2021. Adverse events are Medical Dictionary for Regulatory Activities version 24.0 preferred terms; shown are all-grade adverse events occurring in $\geq 12\%$ of patients in any treatment arm. *Peripheral neuropathy is defined by standard organ class group of preferred terms.

What will be Required to Replace R-CHOP

- **Phase III trial confirming better efficacy or lower toxicity**
- **Must be tolerable in the majority of patients**
 - **Increase in toxicity must be offset by greater increase in benefit**
- **Must be broadly deliverable and affordable**
- **If targeted to a molecular subgroup, require a validated biomarker to identify appropriate patients**

Questions

- Is pola-R-CHP the new standard of care?
- Should it be used in all patients?
 - Regardless of IPI, COO?
 - What about limited stage protocols?
- Should pola-R-CHP be the comparator in all clinical trials?

Future Trial Design

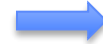
- Require adaptive designs to capture higher risk population
 - Limit exclusion criteria (lab restrictions, ECOG PS)
 - Decentralize biomarker testing
 - Allow initial cycle of therapy prior to randomization
 - Allow initial cycle prior to enrollment (retrospective screening)
 - Statistical power for realistic expectation of outcomes

Alternative Approaches

R-CHOP + X

R-CHOP → X

**Response Adapted
R-CHOP**



Negative trials:

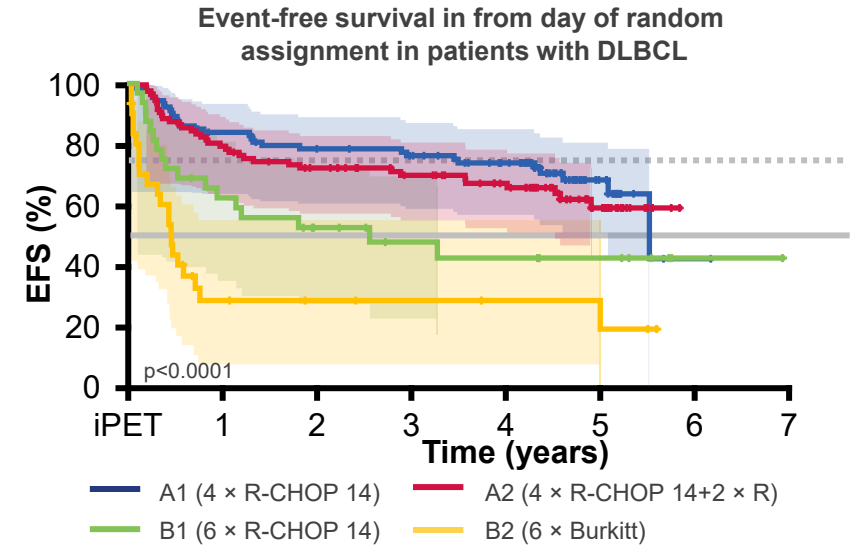
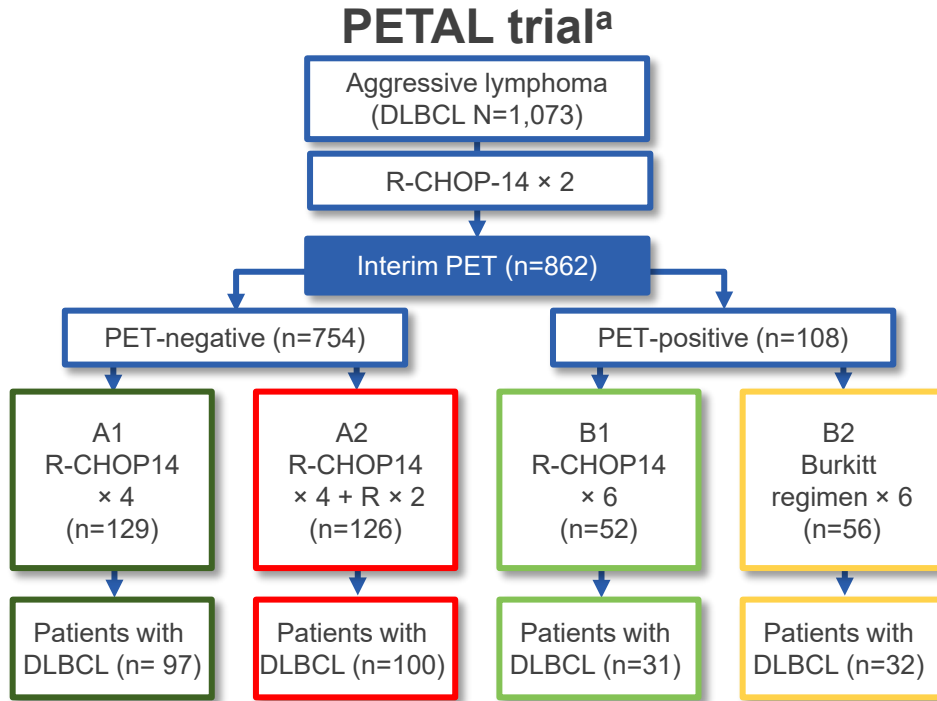
rituximab, enzastaurin,
lenalidomide, everolimus



Negative trials:

PET-adapted intensification
? PET-adapted CAR T-cell
? ctDNA response adapted

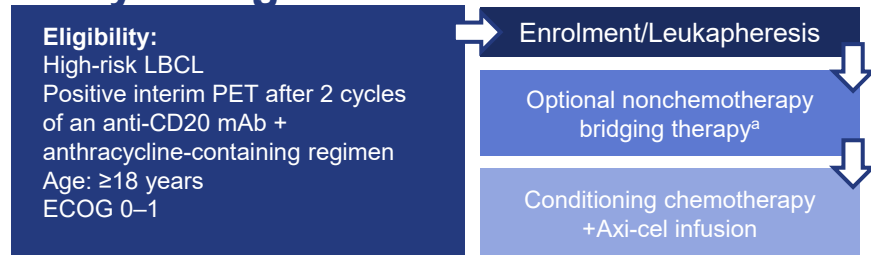
PETAL trial: Intensification of therapy based on interim PET status does not improve outcomes



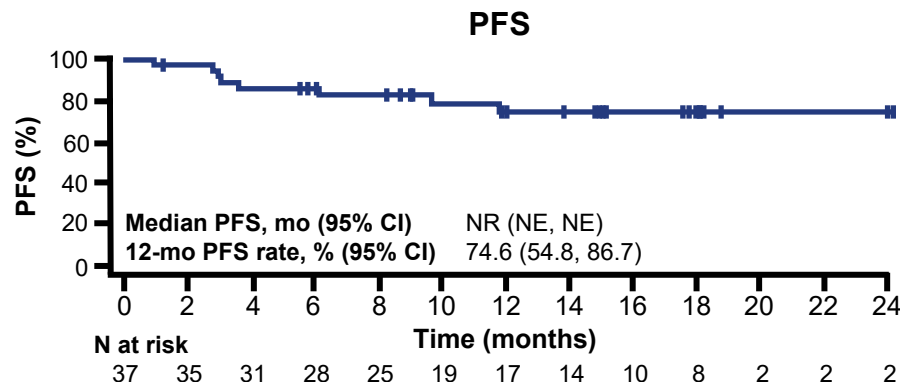
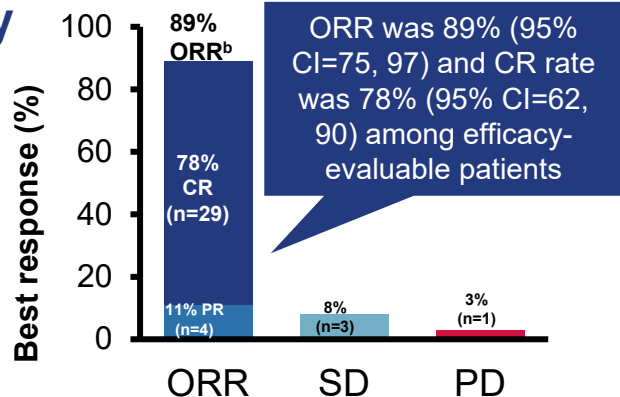
A1 (4xR-CHOP 14)	97	79	71	66	56	18	1
A2 (4xR-CHOP 14+2xR)	100	78	68	57	46	19	0
B1 (6xR-CHOP 14)	32	19	13	9	8	6	1
B2 (6xBurkitt)	31	7	5	4	3	3	0

ZUMA-12: Phase II study using axi-cel as 1L therapy in patients positive PET after 2 cycles

Study Design



Efficacy



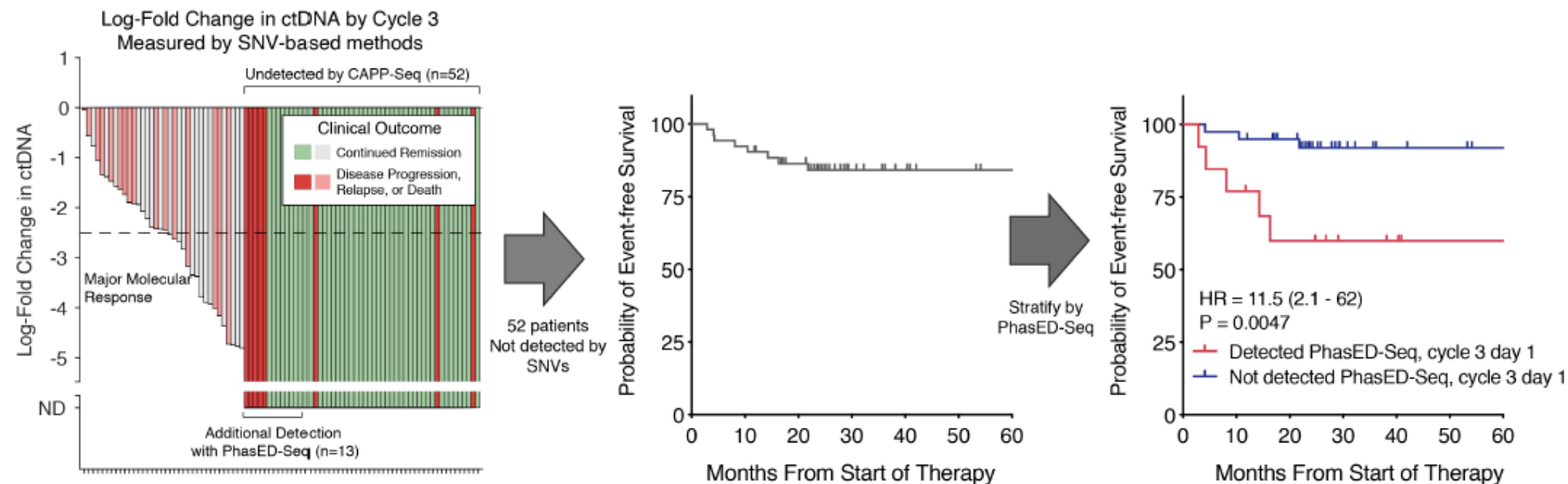
Safety

Toxicities	N=40
Any grade CRS, n (%) ^c	40 (100)
Grade 3	3 (8)
Any grade NE, n (%) ^d	29 (73)
Grade ≥3	9 (23)

Neelapu SS, *et al.* ASH 2021 (Abstract 739; oral).

ctDNA detection by PhasED-Seq improves outcome prediction at interim time-points

- 88 patients with DLBCL undergoing first-line treatment with cycle 3, day 1 samples available
 - ctDNA by CAPP-Seq stratifies patients based on Major Molecular Response (Kurtz et al, JCO 2018)
 - 52 patients undetectable at cycle 3, day 1 by SNV-based CAPP-Seq
- PhasED-Seq further stratifies patients who have undetectable ctDNA by CAPP-Seq



Alternative Approaches

R-CHOP + X

R-CHOP → X

**Response Adapted
R-CHOP**

Replace R-CHOP

Negative trials:

rituximab, enzastaurin,
lenalidomide, everolimus

Negative trials:

PET-adapted intensification
? PET-adapted CAR T-cell
? ctDNA response adapted

? Novel agents (Bispecific Abs)
? CAR T-cell therapy

Guidance-01: Randomized Phase 2 Trial of Genetic Subtype Guided Immunochemotherapy

Study Design (NCT04025593)

- The study started from **July, 2019**.
- All patients were treated with ONE cycle of standard R-CHOP immediately at diagnosis.
- Patients were randomly assigned 1:1 and stratified by genetic subtype.
- Using targeted sequencing and FISH for BCL2, MYC translocation and BCL6 fusion to classify patients into six genetic subtypes MCD like, BN2 like, N1 like, EZB like, according to **NEJM classification (2018)**, TP53 mutation, and others.

Untreated DLBCL

- Age 18-80
- IPI ≥ 2

Stratified by K-medoids algorithm (PAM) simulated genetic subtyping using targeted sequencing panel of 18 genes: *BTG1, CD70, CD79B, CREBBP, DTX1, EP300, EZH2, MPEP1, MTOR, MYD88, NOTCH1, NOTCH2, PIM1, STAT6, TBL1XR1, TNFAIP3, TNFRSF14, and TP53*

R-CHOP $\times 1$

R

1:1

MCD like: Ibrutinib+R-CHOP $\times 5$

BN2 like: Ibrutinib+R-CHOP $\times 5$

N1 like: Lenalidomide+R-CHOP $\times 5$

EZB like: Tucidinostat+R-CHOP $\times 5$

TP53 mutated: Decitabine+R-CHOP $\times 5$

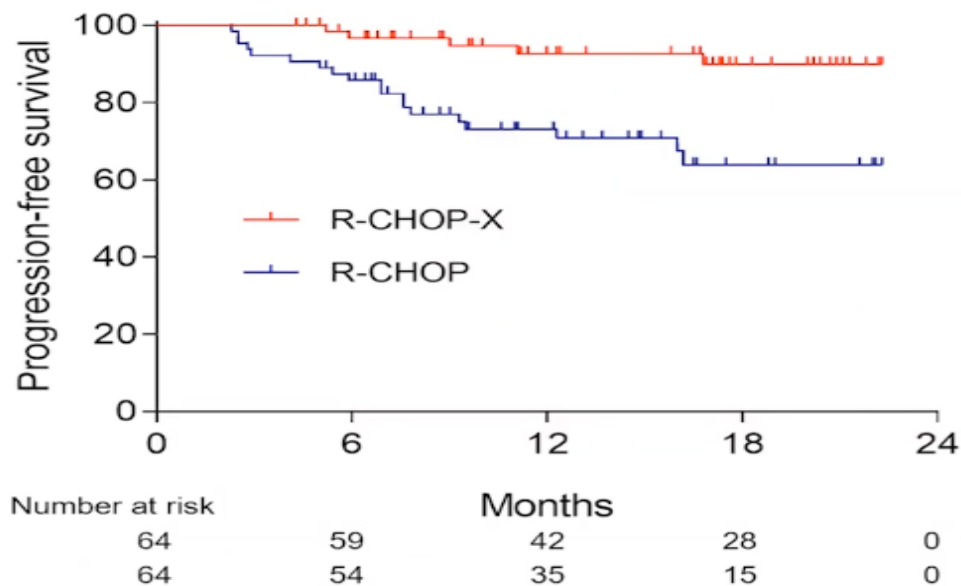
Others: Lenalidomide+R-CHOP $\times 5$

Ibrutinib ¹	420mg po qd
Lenalidomide ²	25mg d1-10 po
Tucidinostat ³	20mg d1, 4, 8, 11 po
Decitabine ⁴	10 mg/m ² d1-5
R-CHOP	Standard dose

G-CSF prophylaxis was given from the second cycle of chemotherapy if grade ≥ 3 neutropenia was present in the first cycle.

1. Younes et al., J Clin Oncol 2019. 2. Nowakowski et al., J Clin Oncol 2021. 3. Zhang et al., Clin Epigenet 2020. 4. Zhang et al., ICML 2019 abstract (NCT02951728)

Secondary Endpoint: PFS



Median follow-up 16.1 months

	R-CHOP-X	R-CHOP
1-year PFS (95%CI)	93% (81%-97%)	73% (60%-83%)

Ongoing/Planned Trials in Upfront DLBCL

- **BTK-inhibitor R-CHOP trials**
 - Escalade (acala); UK trial; zanabrutinib
- **First-Mind Trial**
 - Tafasitamab/Lenalidomide + R-CHOP
- **Bispecific antibodies + R-CHOP**
- **Biology-driven trials**
- **Response-adapted trials (ctDNA, quantitative PET/CT)**

Summary

- Moving beyond R-CHOP has been a challenge
- Pola-R-CHP results in improved PFS with similar toxicity
- Improving the cure rate in frontline setting is important as secondary therapies associated with higher toxicity, cost and poor outcomes
- Further improvement needed and trials of novel therapies remain important
- Identification of predictive biomarkers (using validated tools) will be essential to optimize outcome in individual patients