

# R-CHOP es insuficiente en LBCG de alto riesgo

Beatriz Wills, MD

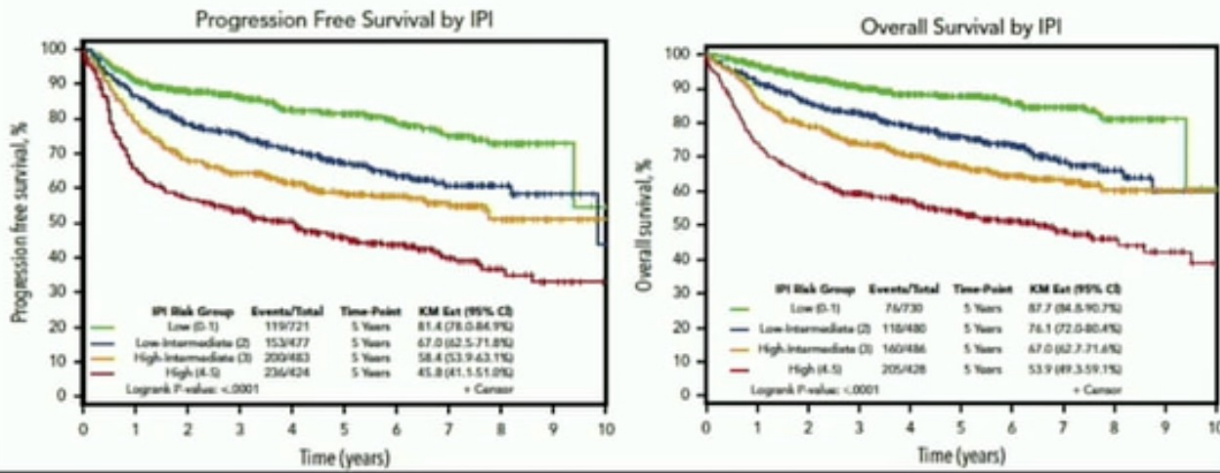
Fundación Santa Fe de Bogotá

# Características clínicas de nuestro paciente

## IPI risk factors, 1 point each: a, b

- Age > 60 years
- Serum LDH > normal
- Stage III–IV disease
- ECOG PS 2–4
- Two or more extranodal sites

L = 0 or 1  
 L-1 = 2  
 H-1 = 3  
 H = 4 or 5



**3** points

Poor prognosis (R-IPI)  
 High-intermediate risk  
 group (IPI)

**55** %

Overall survival (R-IPI)  
 49% overall survival (IPI)

**53** %

Progression-free survival (R-IPI)  
 57% progression-free  
 survival (IPI)

**IPI 3**

**NCCN IPI 4**

**Riesgo alto-intermedio**

Despite the ease in its implementation, IPI **does not fully represent disease heterogeneity.**

Therefore, efforts have **shifted to broad molecular profiles** that model and stratify risks of adverse outcomes.

# Índices clínicos pronósticos en LBCG tienen limitaciones

Análisis de 2124  
pacientes.

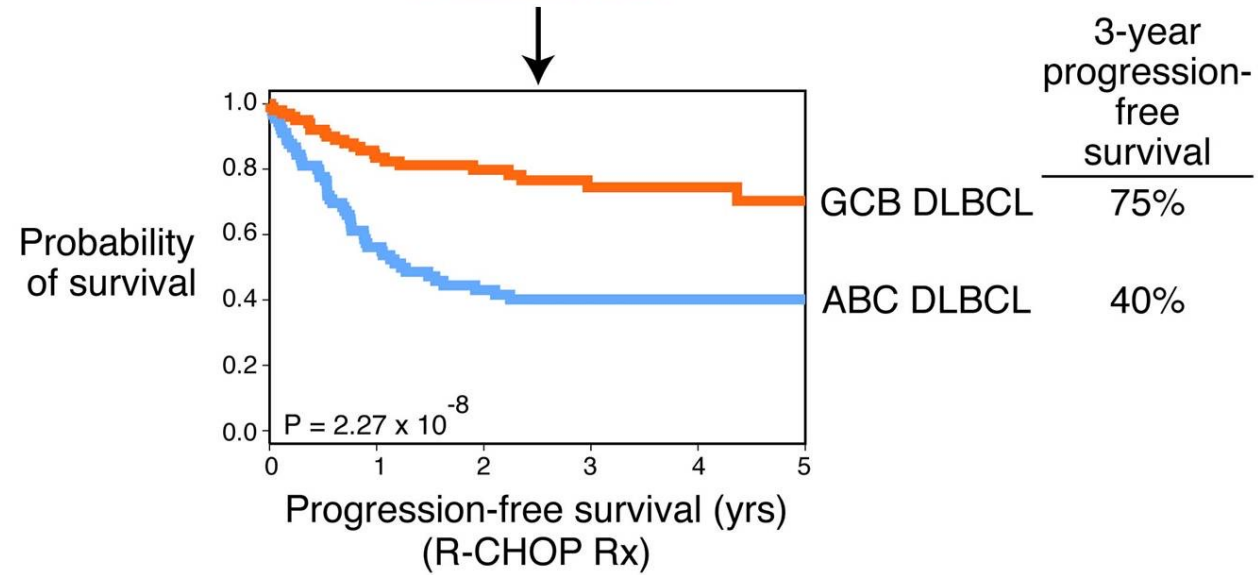
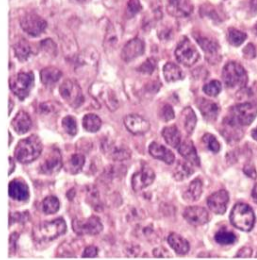
- Comparación de IPI, R-IPI, IPI NCCN

NCCN-IPI tuvo el mejor  
poder de  
discriminación para OS.

- No logró identificar un grupo de **pacientes de alto riesgo** con supervivencia global significativamente por debajo del 50%.

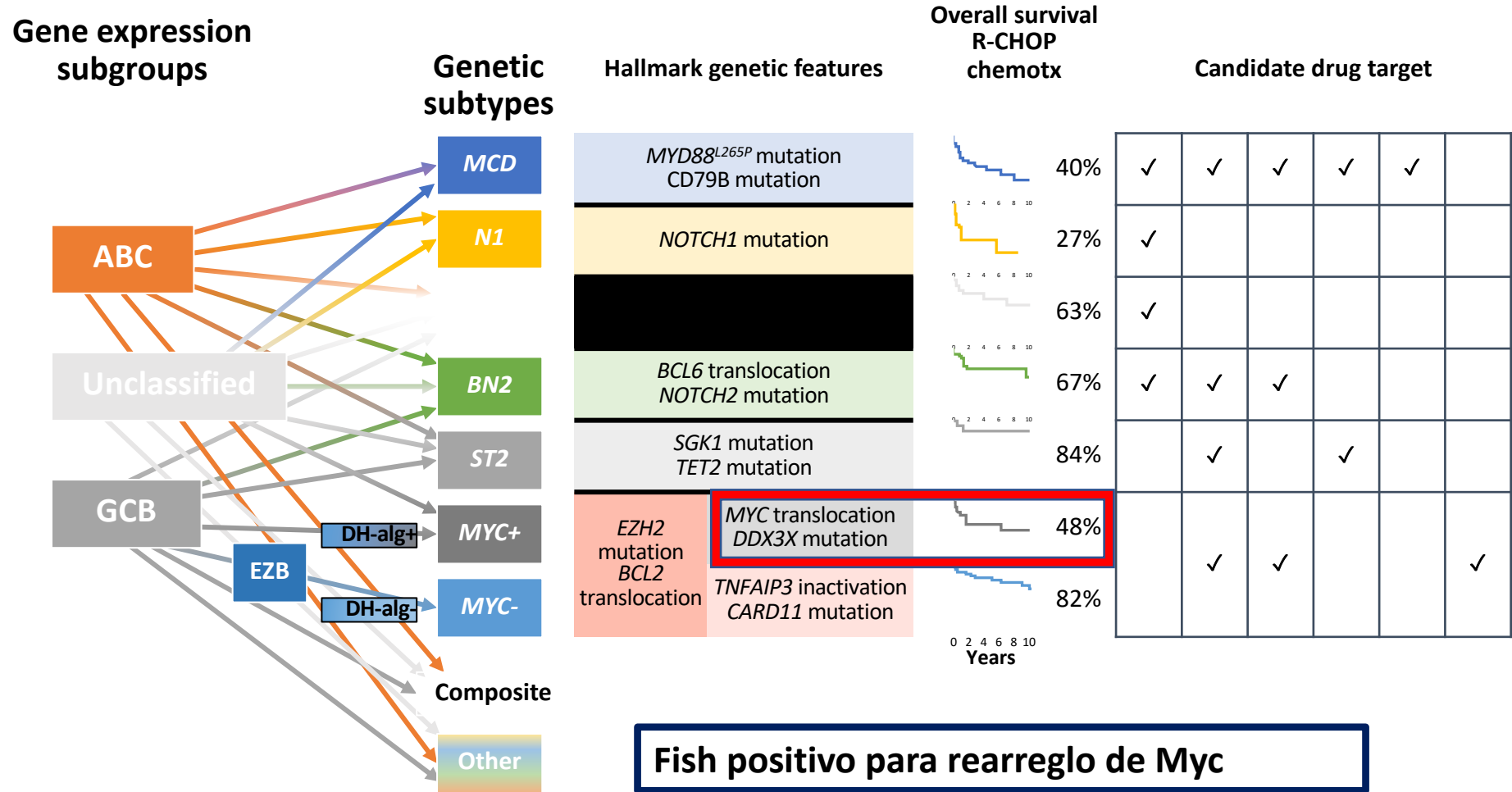
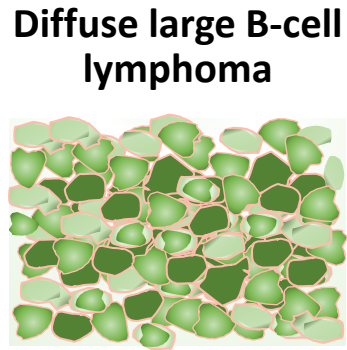
# Linfoma Difuso de Células Grandes B, centro germinal

Diffuse Large B Cell  
Lymphoma  
(DLBCL)



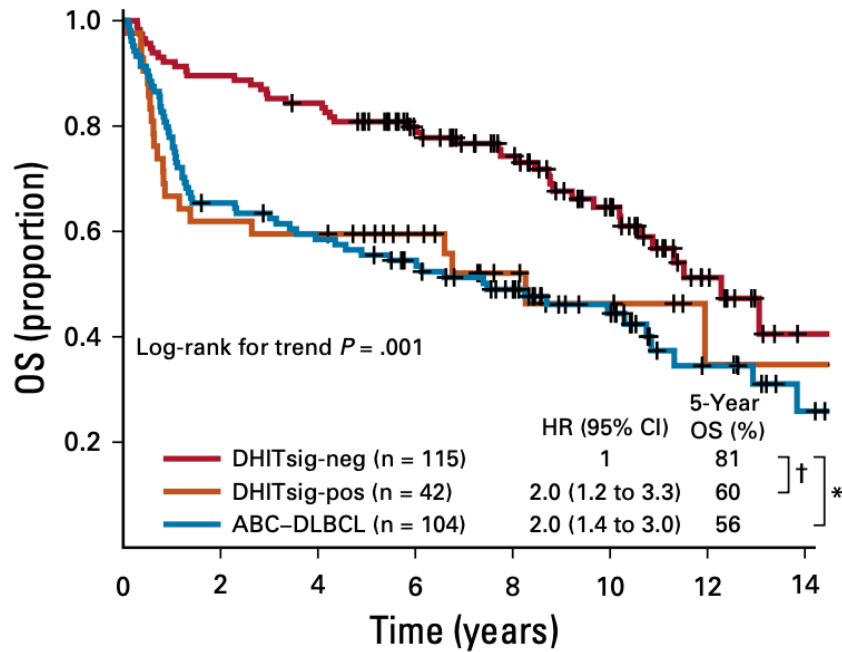
Subtype-specific response  
To chemotherapy

# Pero no todos los GCB se comportan igual... ¿Cuáles son las Características moleculares de nuestra paciente ?



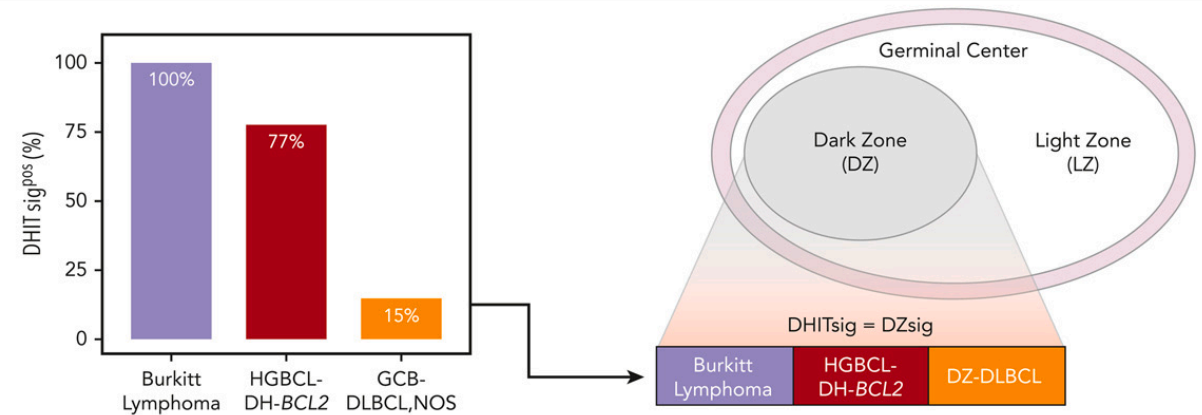
Heterogeneity observed within the COO subtypes

# Subtipo agresivo de linfoma CG caracterizado por perfil molecular Dark zone +

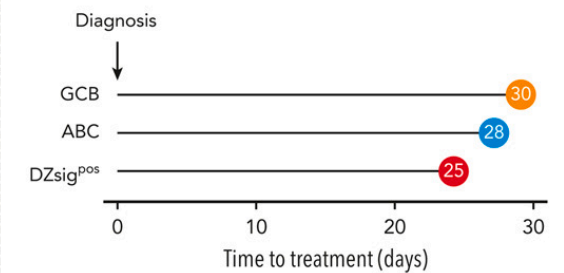
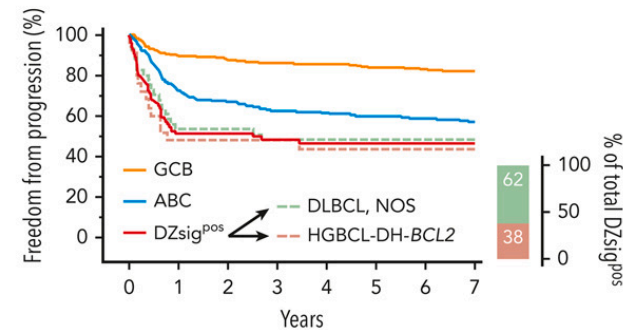


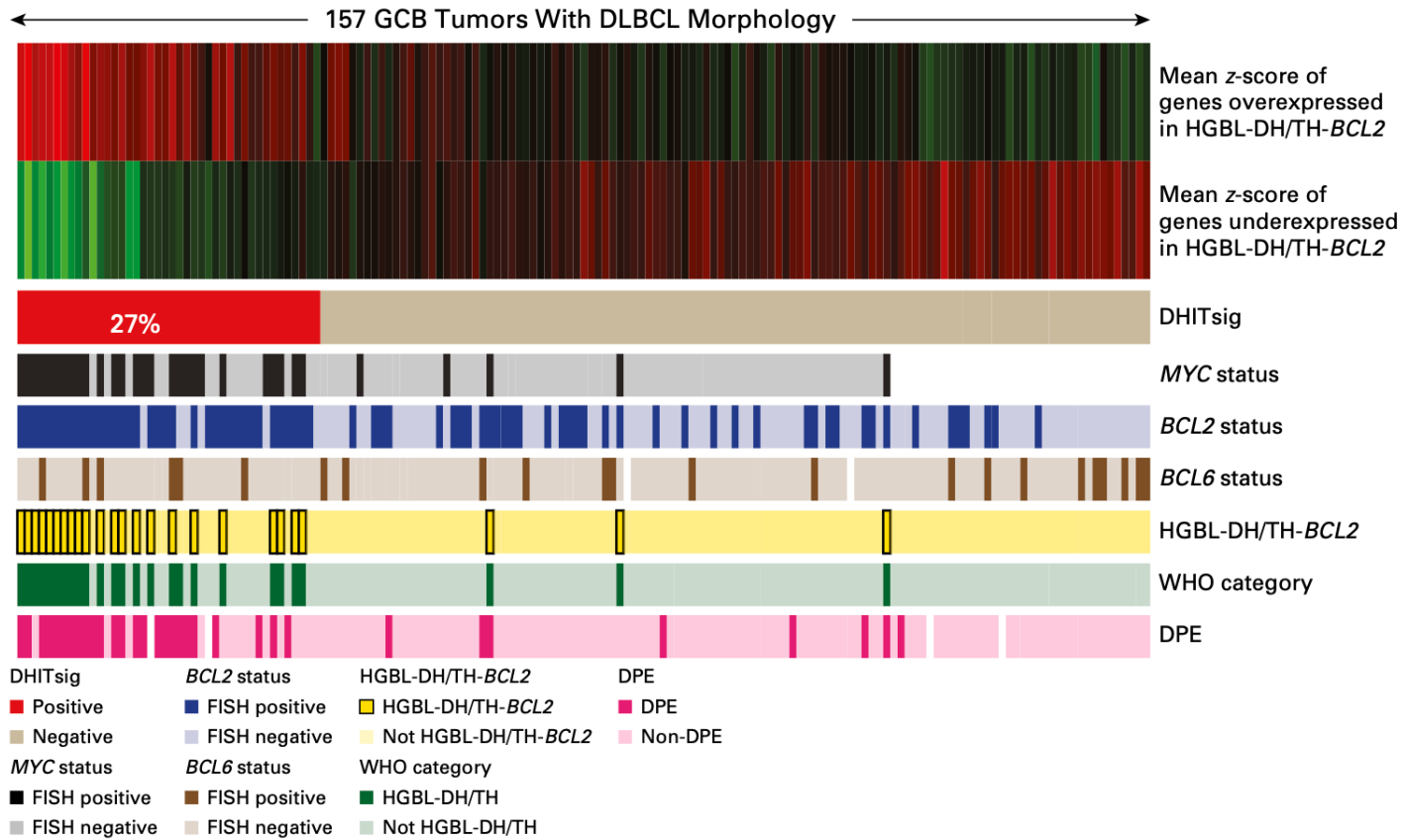
No. at risk	0	2	4	6	8	10	12	14
DHITsig-neg	115	103	96	79	62	41	17	4
DHITsig-pos	42	26	25	18	10	7	4	1
ABC-DLBCL	104	67	60	51	40	27	12	6

1. DHITsig expression extends beyond HGBCL-DH-BCL2 to identify dark zone lymphomas, and was thus renamed the "dark zone signature" (DZsig)

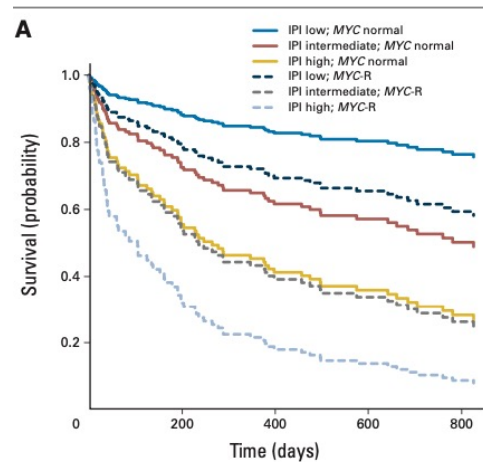
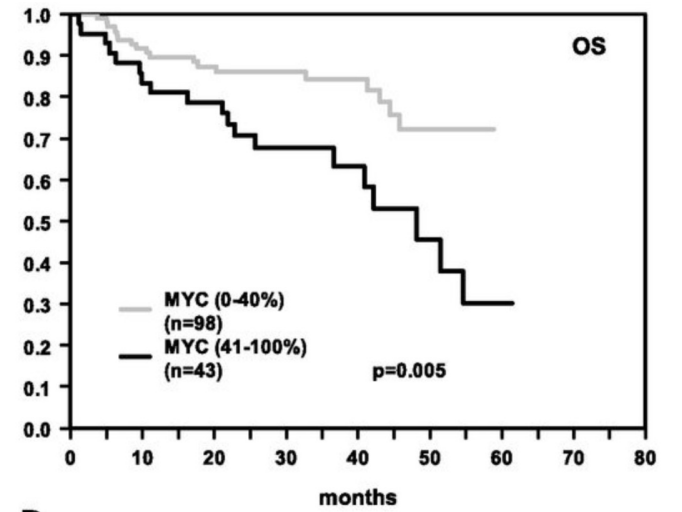
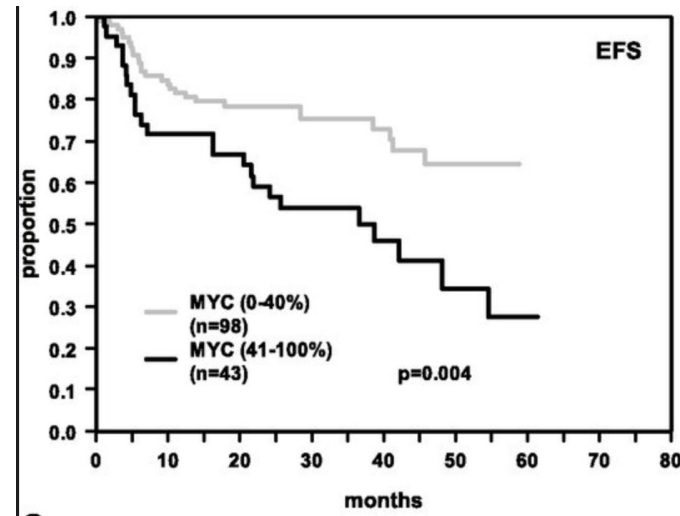
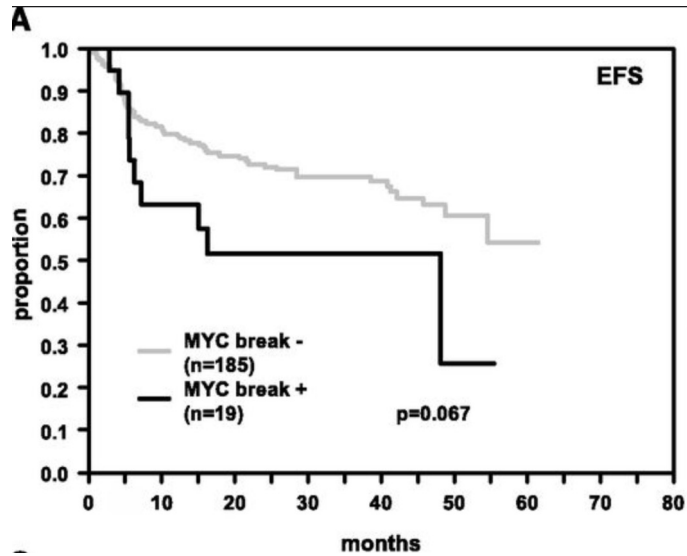


2. Among tumors of DLBCL morphology, gene expression profiling-defined molecular subgroups are associated with outcomes and diagnosis-to-treatment interval



**B**

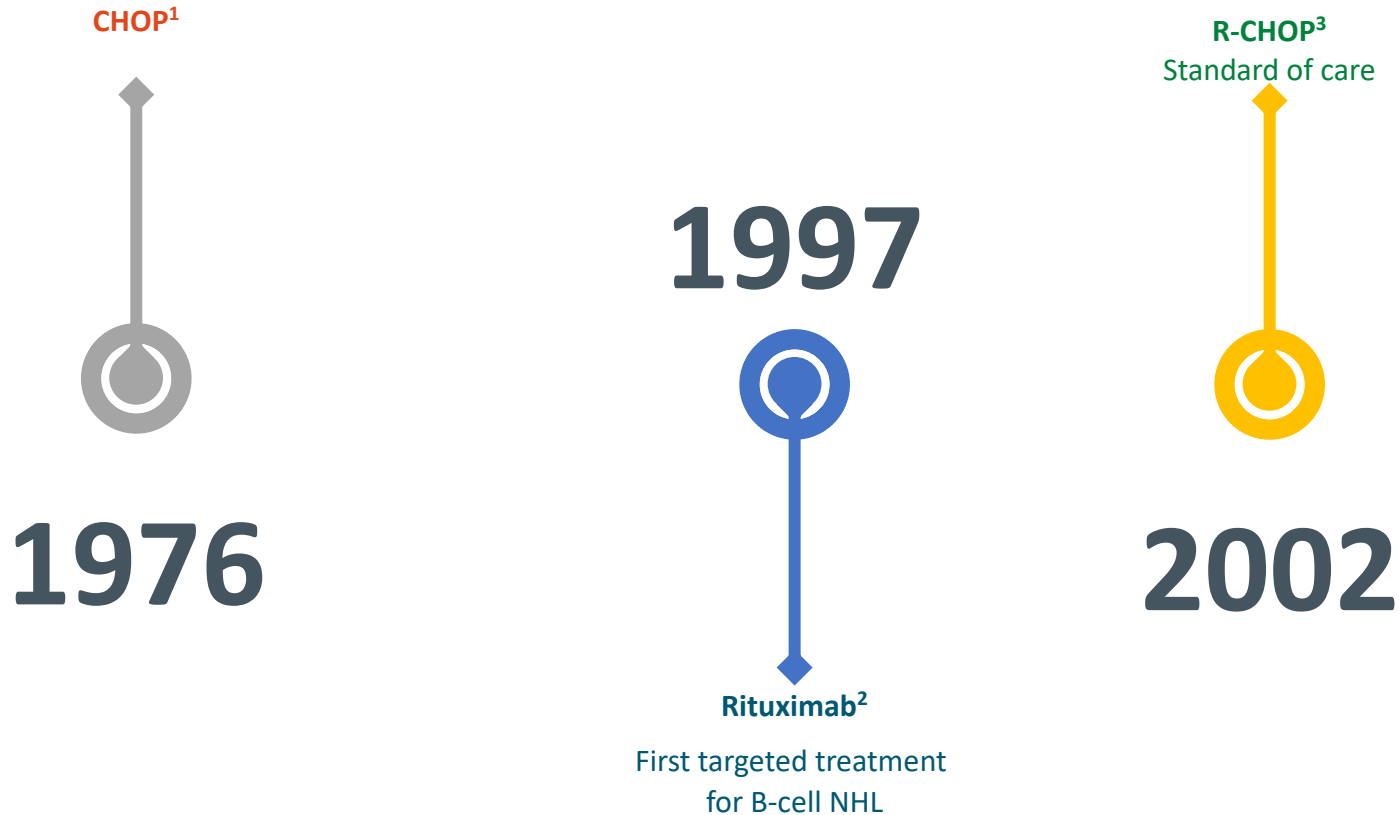
# Alteraciones en *MYC*--> fenotipo más agresivo y peores desenlaces



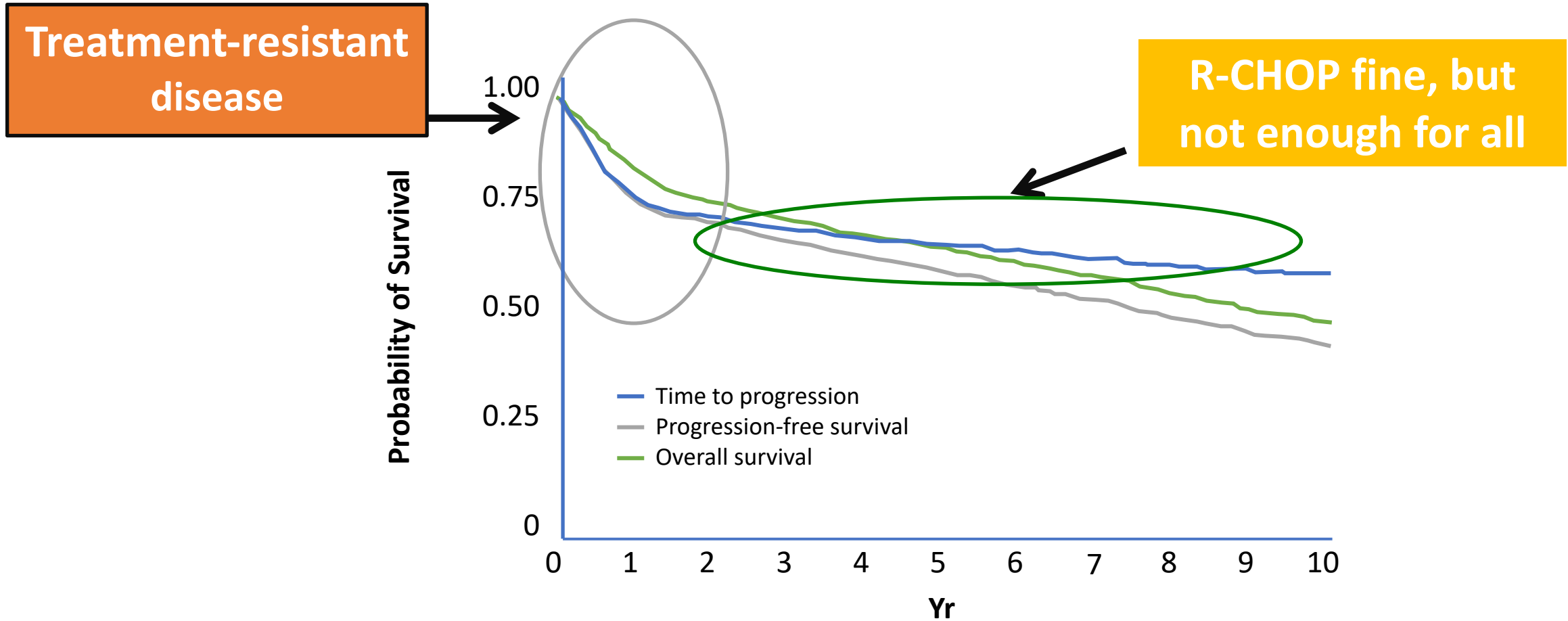
**Nuestra paciente:  
IHC MYC 40%  
FISH rearrreglo de Myc  
Pobre factor pronóstico**

rMYC +IPI → factor pronóstico desfavorable

# A pesar de entender mejor la biología de LBCG... Seguí(a)mos tratando a los pacientes con LDCBG igual por más de 20 años ...



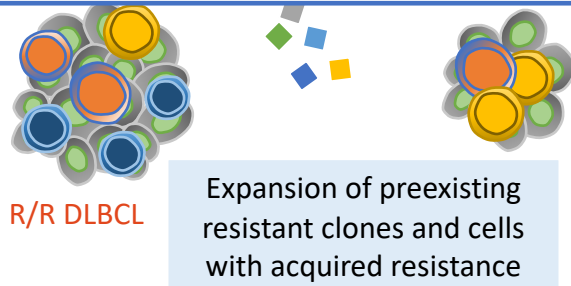
# Desenlaces con R-CHOP en primera línea para DLBCL



# Resistencia al tratamiento a primera línea es multifactorial

## Tumor Heterogeneity

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



Examples of genetic and/or epigenetic alterations in R/R DLBCL

### Genetic modifications

Cell cycle regulation

*CCND3, CDKN2A, CDKN2B*

DNA damage response

*TP53*

Epigenetic regulation

*EZH2, CREBBP, MEF2B, KMT2C, KMT2D*

Immune surveillance

*B2M, CD58, HLA-A, MS4A1*

Oncogenes

*MYC, PIM1, PRKCC, GATA3,*

*MLLT10, ABI1*

Signaling pathway activation

*STAT6, SOCS1, FOXO1, MYD88,*

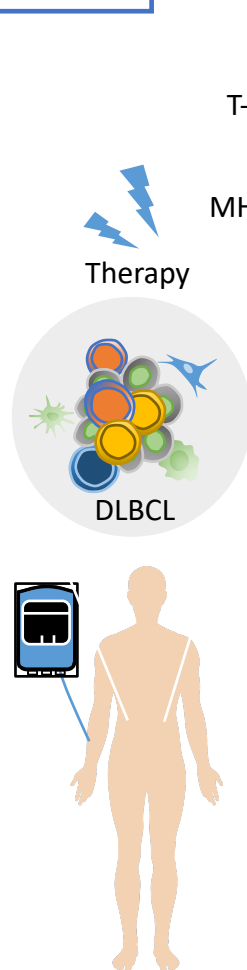
*CD79B, NFKBIE, NFKBIZ*

### Epigenetic modifications

DNA methylation

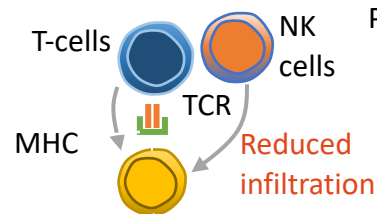
Histone

methylation/acetylation



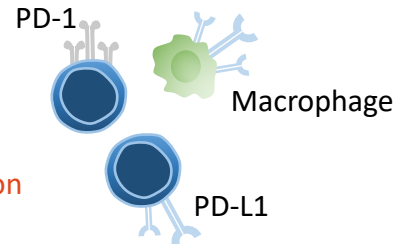
Immune dysfunction and supportive stromal cells promote a protumor environment for treatment resistance

Deficiency in TIICs



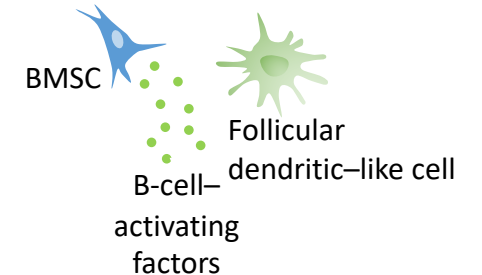
Inadequate apoptosis

Upregulation of inhibitory immune checkpoint molecules



Dysfunctional immune TME

Inhibition of apoptosis by cell adhesion-mediated resistance



Protection from apoptosis

## Host variabilities

Interpatient variabilities represented from multiple host-specific factors lead to highly variable responses to treatment

### Demographic and physical factors

Age

Body weight

Gender



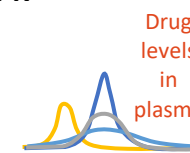
### PK

Absorption

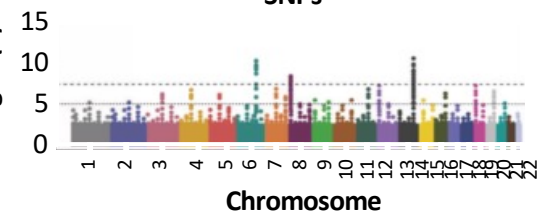
Distribution

Metabolism

Excretion

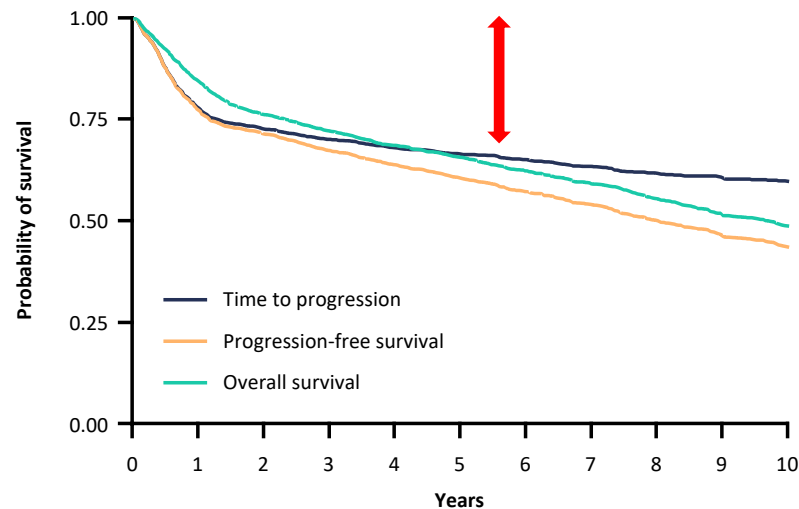


Log<sub>10</sub> (P)



# La mayoría de esfuerzos de mejorar R-CHOP han fallado

## R-CHOP-21



**Intensificar terapia:**  
*ASCT<sup>1</sup>, CHOP-14<sup>2</sup>, ACVBP<sup>3</sup>, DA-R-EPOCH<sup>4</sup>*

**Adicionar agente adicional:**  
*bortezomib<sup>5</sup>, ibrutinib<sup>6</sup>  
and lenalidomide<sup>7</sup>*

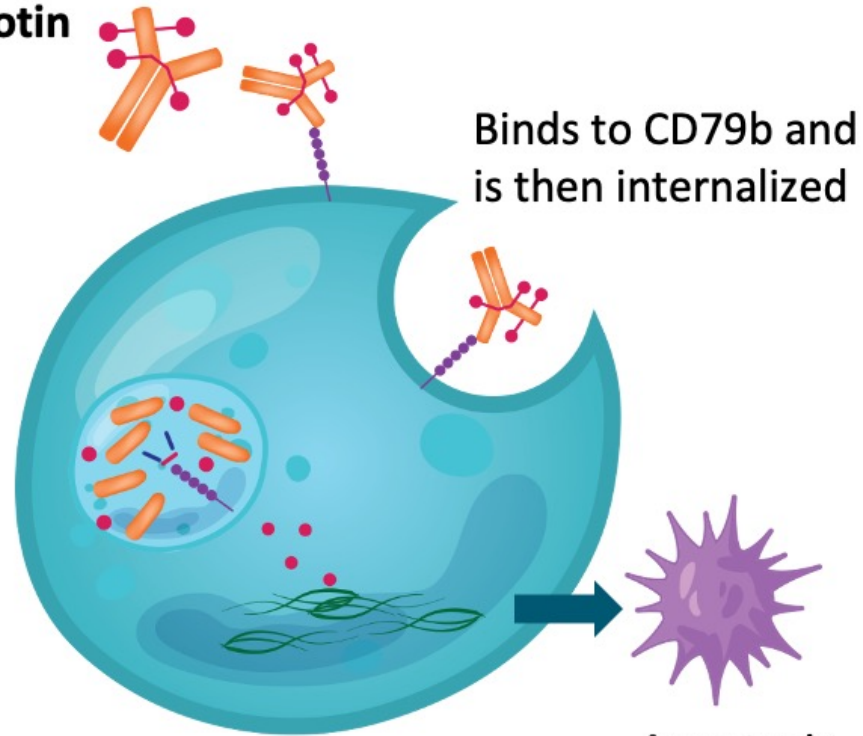
**Mantenimiento**  
*rituximab<sup>8</sup>, lenalidomide<sup>9</sup>,  
enzastaurin<sup>10</sup>, everolimus<sup>11</sup>*

**Reemplazar un componente**  
*bortezomib<sup>12</sup>, obinutuzumab<sup>13</sup>*

# Mecanismo de acción Polatuzuman Vedotin

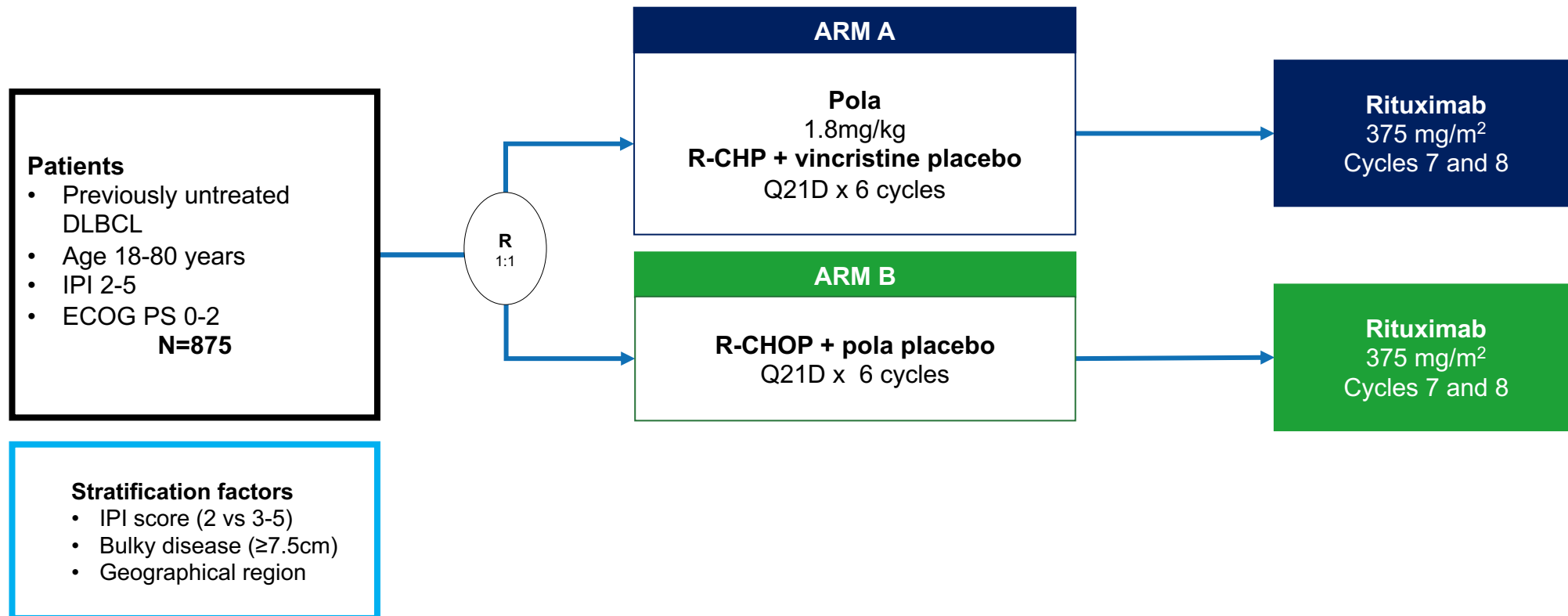
Anticuerpo monoclonal anti-CD79b  
conjugado a monometin auristatin E (MMAE)

**Polatuzumab  
vedotin**



# POLARIX: Diseño del estudio

Ensayo doble ciego, fase 3, controlado con placebo

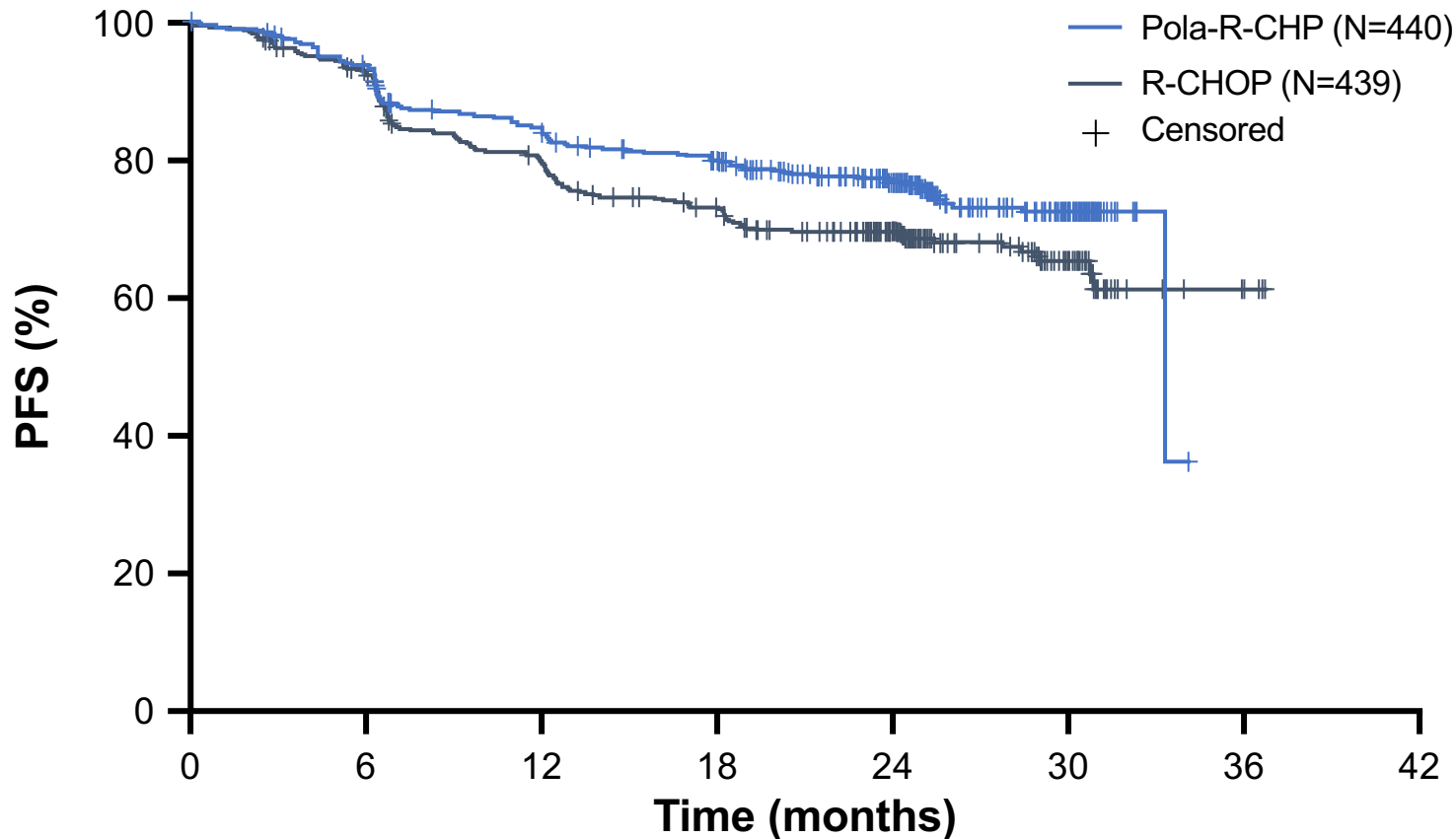


ECOG PS, Eastern Cooperative Oncology Group Performance Status; IPI, international prognostic index; LYSA, Lymphoma Study Association; LYSARC, Lymphoma Academic Research Organisation; Q21D, every 21 days; R-CHOP, rituximab, cyclophosphamide, doxorubicin, vincristine, and prednisone; R-CHP, rituximab, cyclophosphamide, doxorubicin, and prednisone.



Collaboration with LYSA and LYSARC

# POLARIX: Pola-R-CHP mejoró significativamente la supervivencia libre de enfermedad frente a R-CHOP.



**HR 0.73** 95% CI, 0.57, 0.95

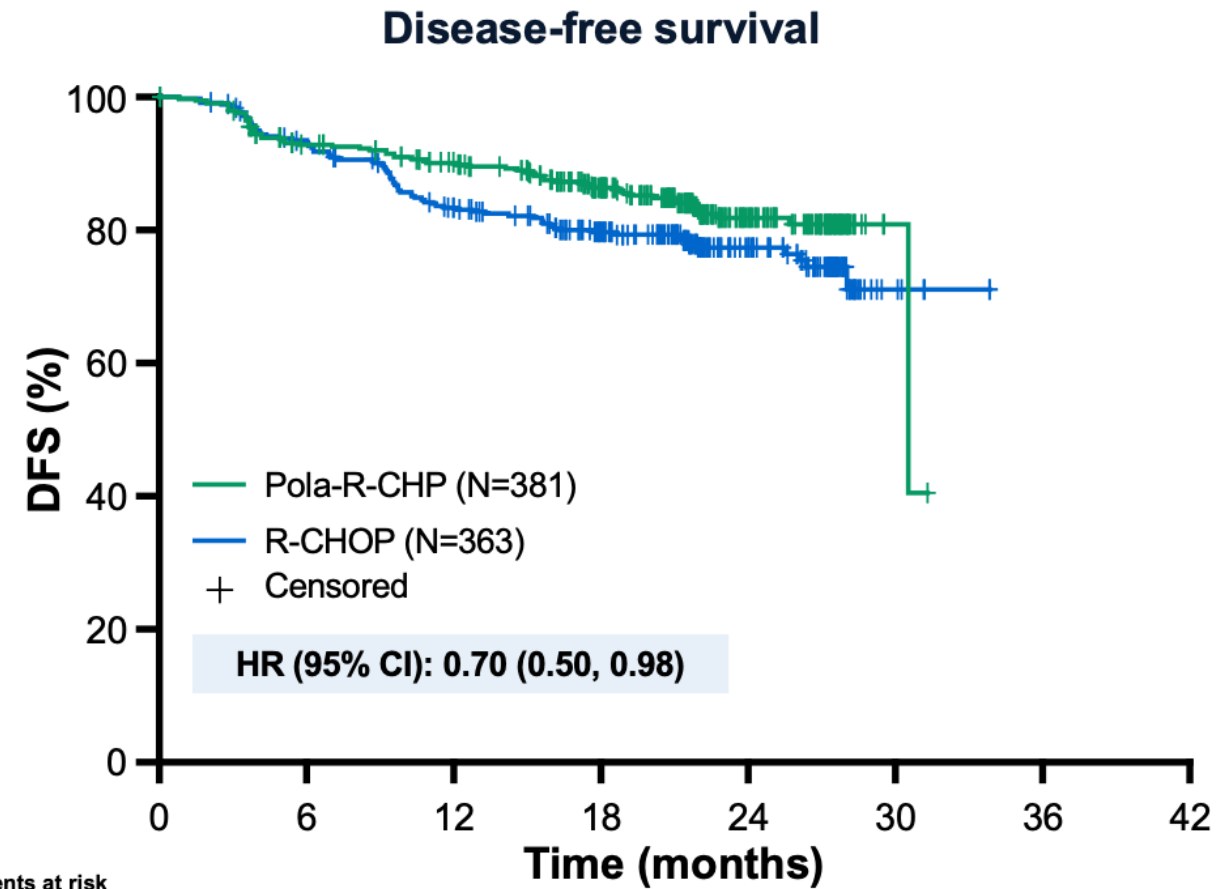
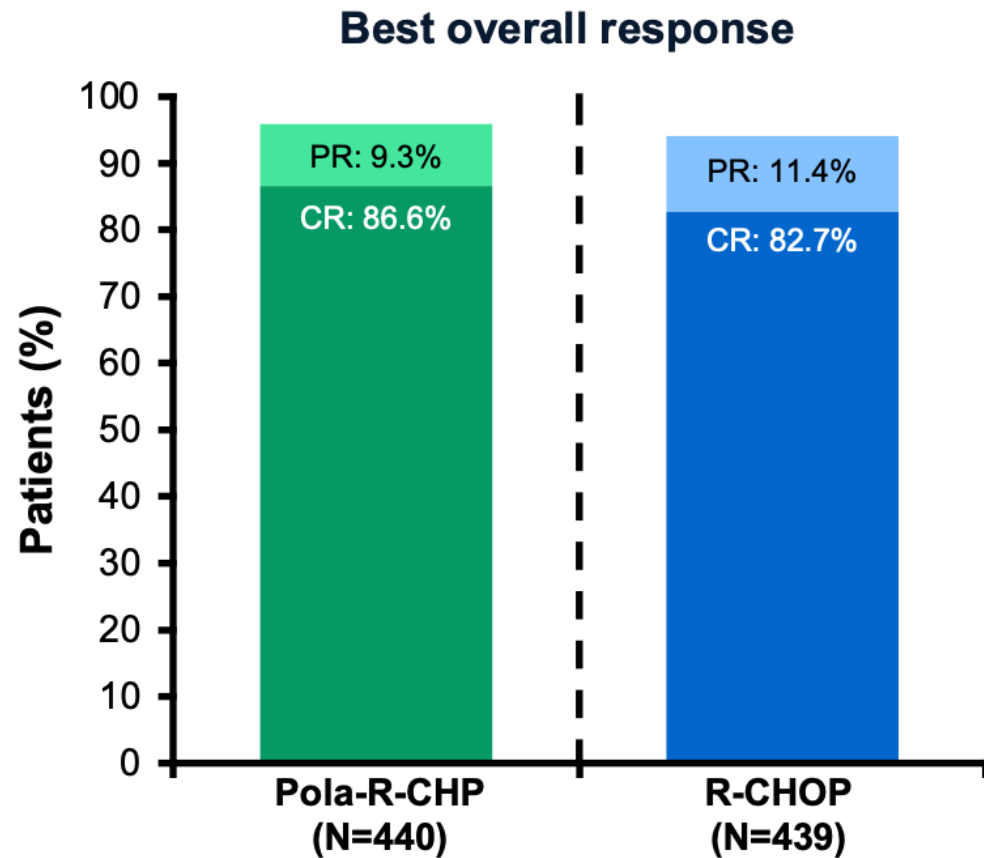
- Pola-R-CHP demonstrated a **27% reduction in the relative risk of disease progression, relapse, or death** versus R-CHOP
- **24-month PFS:** 76.7% with Pola-R-CHP versus 70.2% with R-CHOP ( $\Delta=6.5\%$ )

**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.

# Tasas de respuesta y supervivencia libre de enfermedad

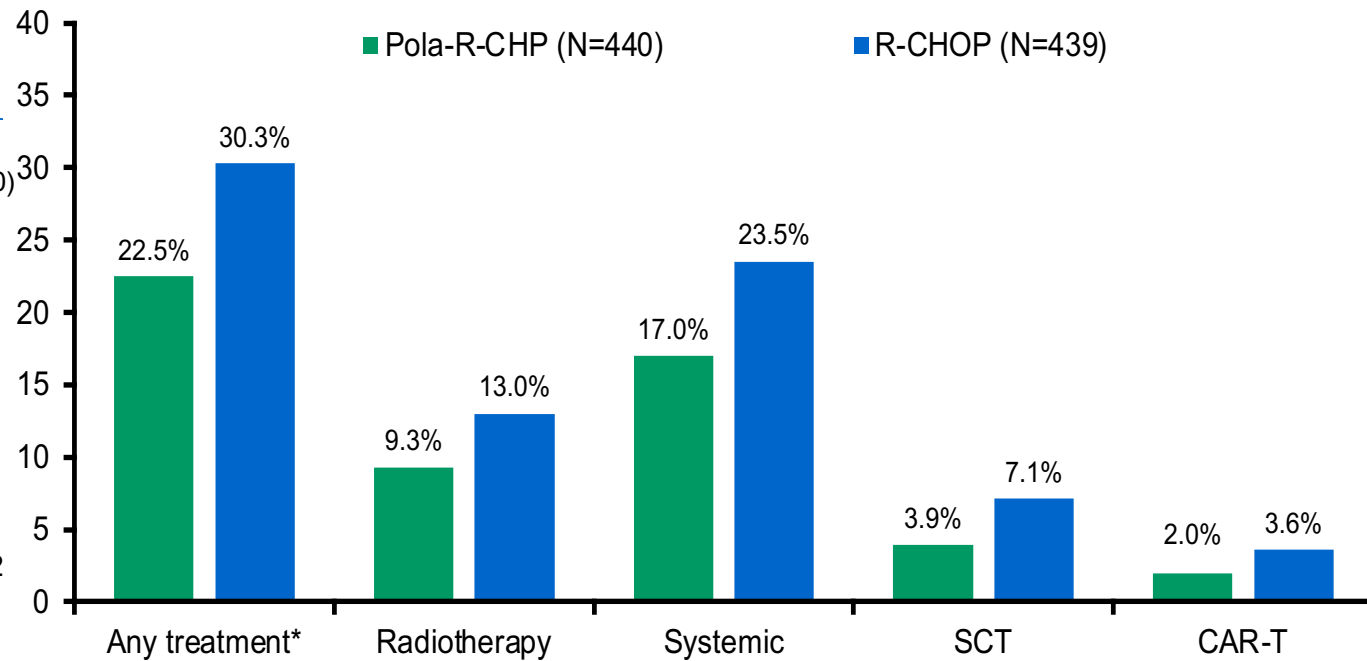
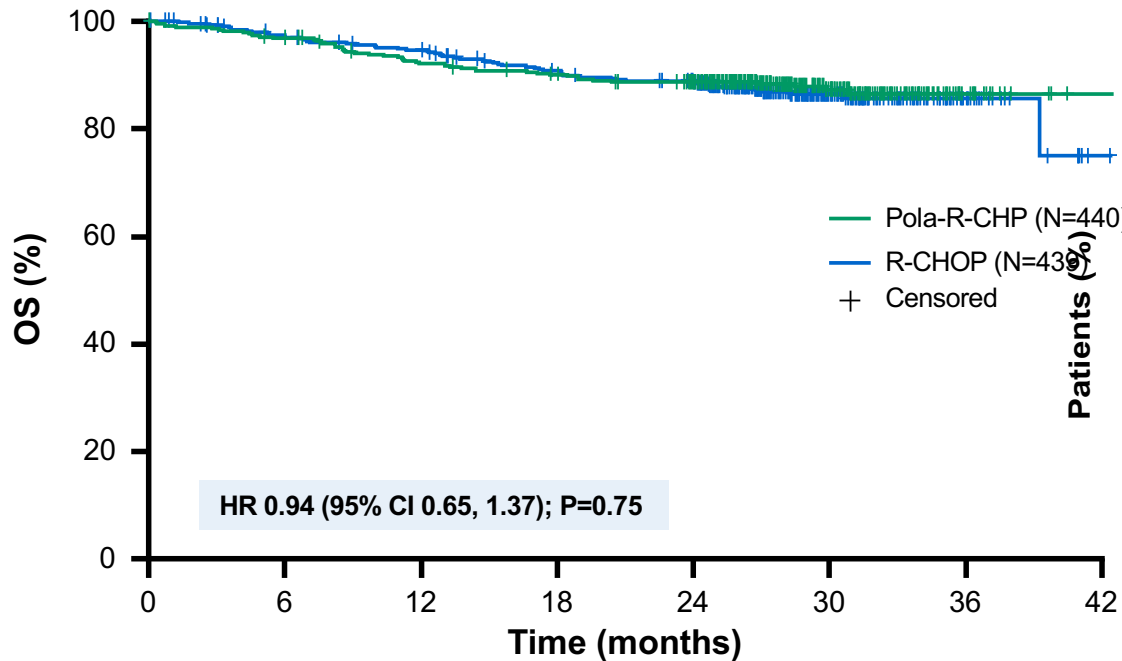


No. of patients at risk

Pola-R-CHP	381	342	322	266	106	2	NE	NE
R-CHOP	363	326	282	238	96	5	NE	NE

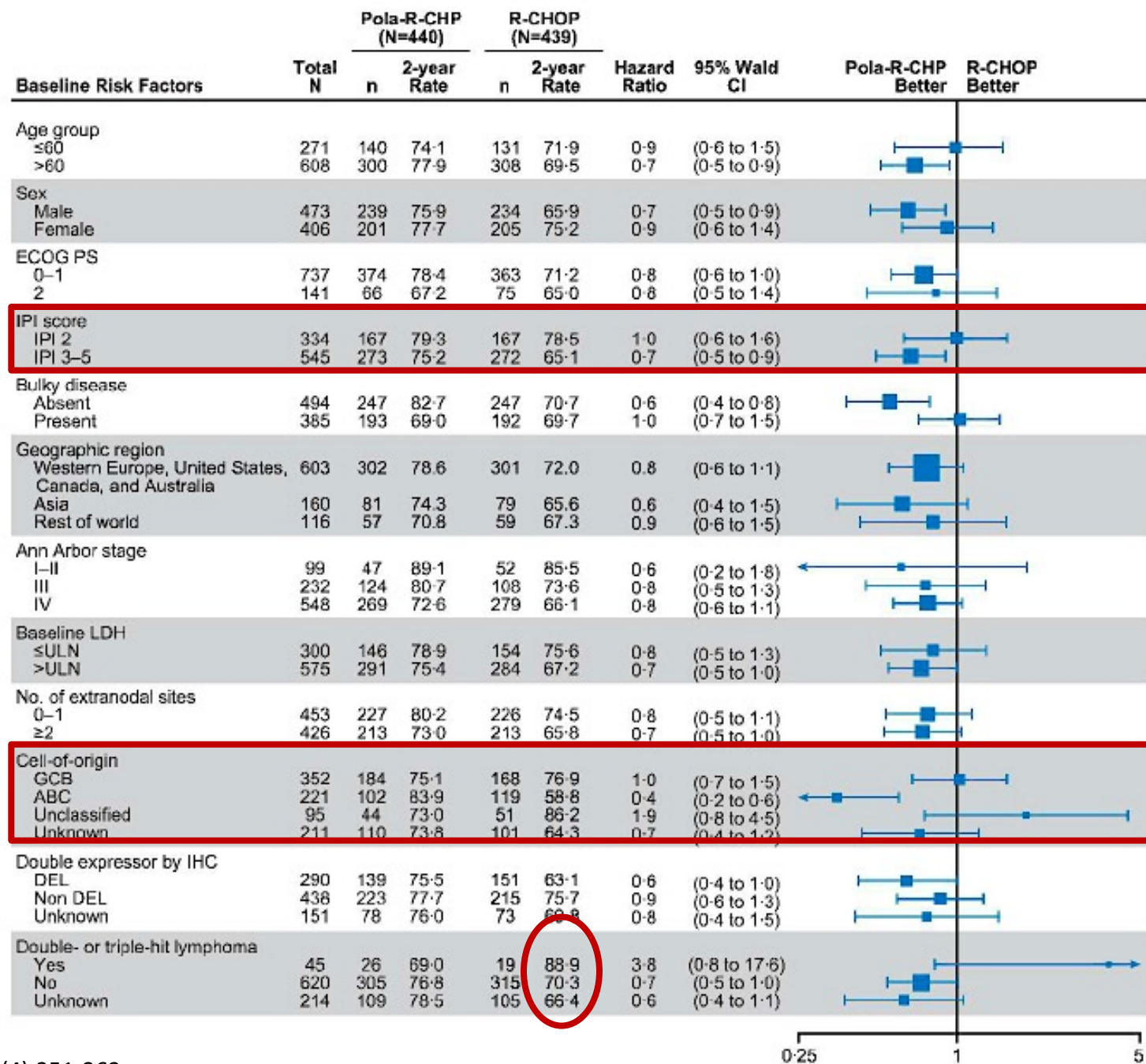
# POLARIX: Supervivencia global y necesidad de tratamientos adicionales

## Necesidad de tratamiento adicionales



No. of patients at risk

Pola-R-CHP	440	423	397	384	362	140	15	1
R-CHOP	439	414	401	376	355	132	20	1

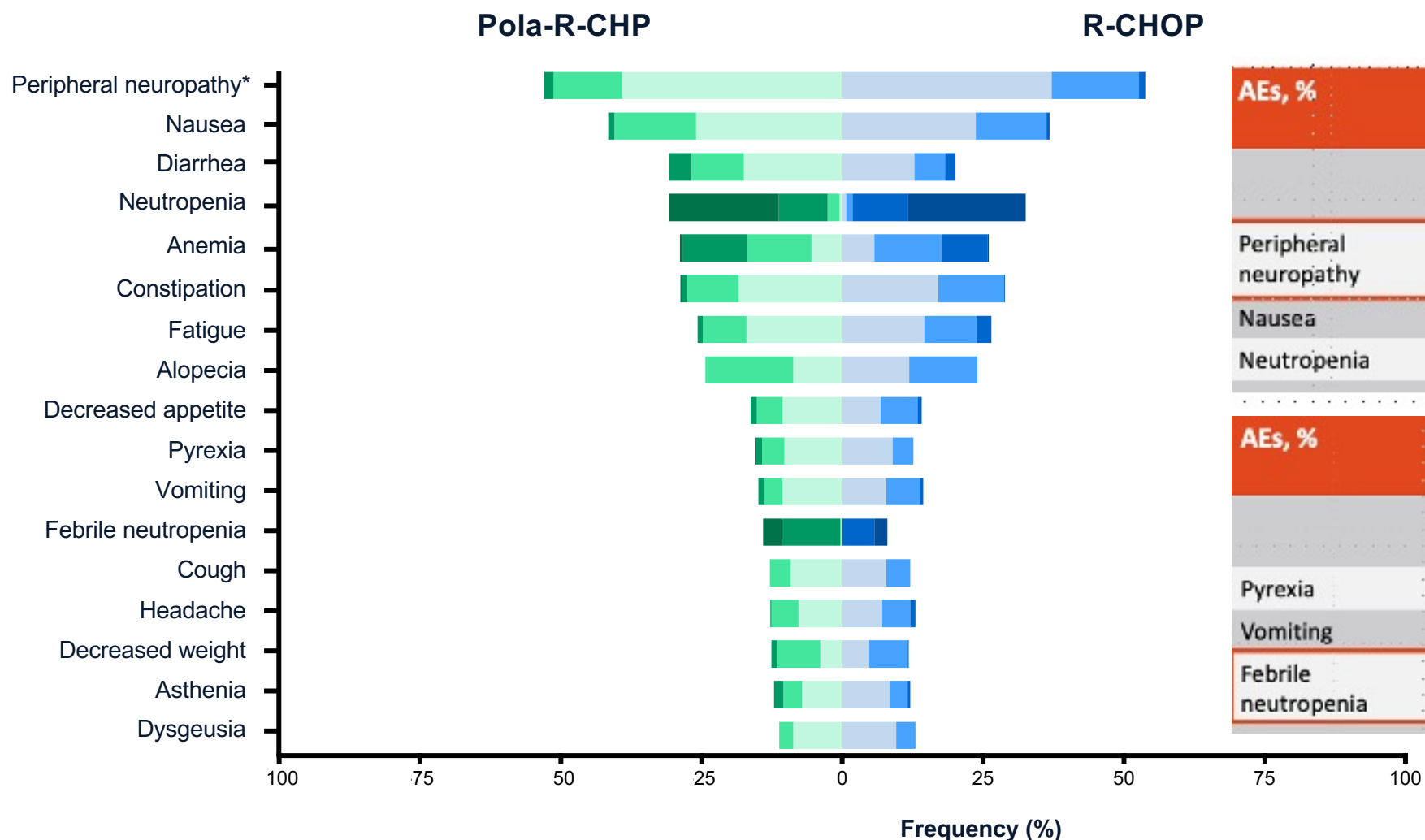


10% PFS difference in pts with IPI 3-5

15% PFS difference in pts with ABC DLBC

Unusual outcome for control arm pts with DH/TH

# POLARIX: Efectos adversos son similares



AEs, %	Pola + R-CHP (n = 435)		R-CHOP (n = 438)	
	Any Grade	Grade 3/4	Any Grade	Grade 3/4
Peripheral neuropathy	52.9	1.6	53.9	1.1
Nausea	41.6	1.1	36.8	0.5
Neutropenia	30.8	28.3	32.6	30.8

AEs, %	Pola + R-CHP (n = 435)		R-CHOP (n = 438)	
	Any Grade	Grade 3/4	Any Grade	Grade 3/4
Pyrexia	15.6	1.4	12.6	0
Vomiting	14.9	1.1	14.4	0.7
Febrile neutropenia	14.3	13.8	8.0	8.0

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.

# Deciphering the Clinical Benefit of Pola-R-CHP versus R-CHOP in Different Genetic Subtypes Beyond Cell of Origin in the POLARIX Study

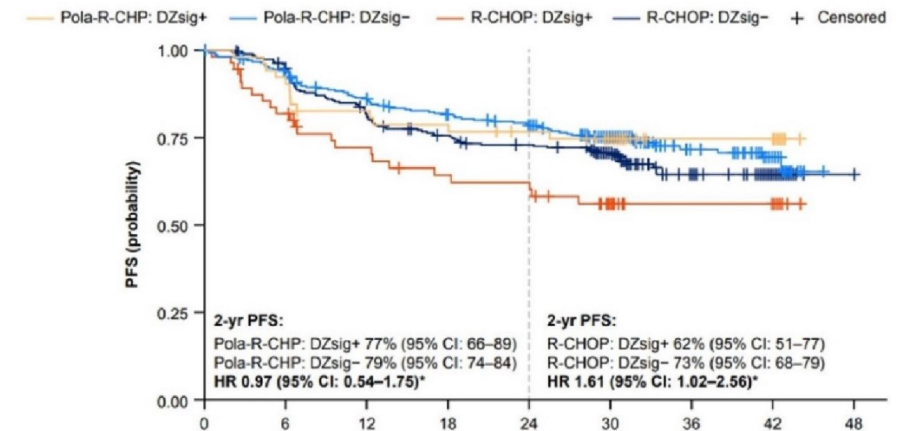
Franck Morschhauser, Wilfred Leung, Vibha Raghavan, Georg Lenz, Fabrice Jardin, Alex F. Herrera, Laurie H. Sehn, Jeff P. Sharman, Christopher R. Flowers, Jonathan W. Friedberg, Marek Trněný, Hervé Tilly, Charles Herbaux, Samuel Tracy, Christopher R. Bolen, Will Harris, Jamie H. Hirata, Calvin Lee, Yanwen Jiang, Gilles Salles



Blood (2023) 142 (Supplement 1): 3000.

<https://doi.org/10.1182/blood-2023-178623>

- Ciertos subtipos moleculares ( independiente del COO) se han asociado con mal pronóstico con R-CHOP
  - *EZH2* mutations/*BCL6* translocations (EZB), *MYD88/CD79B*, dark zone gene expression (DZ)
- Análisis post hoc POLARIX.
  - Pts con GCB DLBCL DZsig+ beneficio significativo Pola-R-CHP vs R-CHOP.
- DZsig refines cell-oforigin classification by identifying patients within GCB-DLBCL with inferior outcomes and shorter time to treatment.

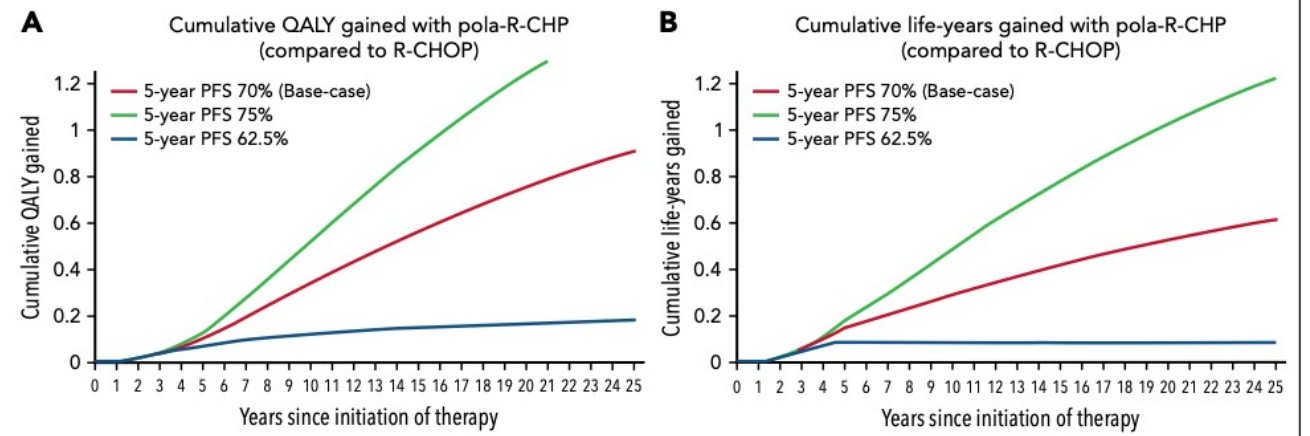


\*HRs compare DZsig+ versus DZsig- in each treatment arm.  
 CI, confidence interval; DZsig, dark zone gene expression signature; HR, hazard ratio; PFS, progression-free survival;  
 Pola-R-CHP, polatuzumab vedotin in combination with rituximab plus cyclophosphamide, doxorubicin, and prednisone;  
 R-CHOP, rituximab, cyclophosphamide, doxorubicin, vincristine, and prednisone.

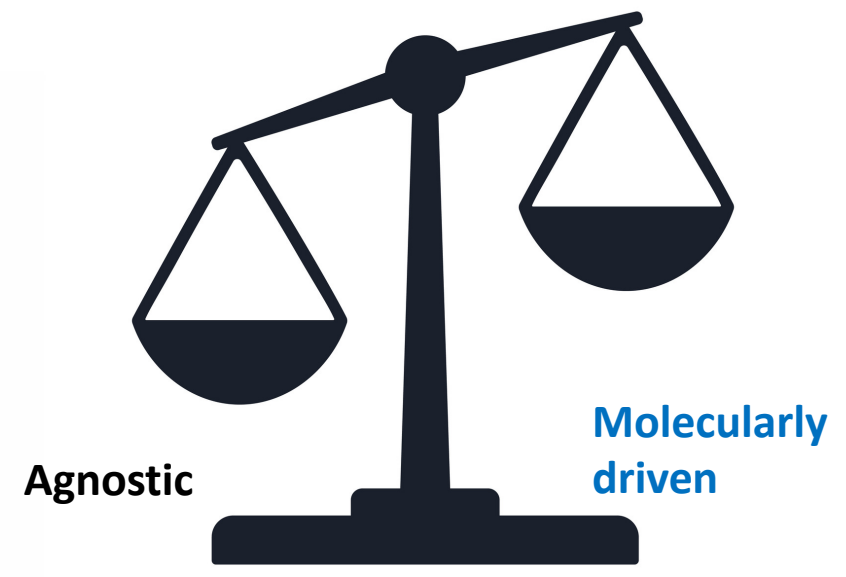
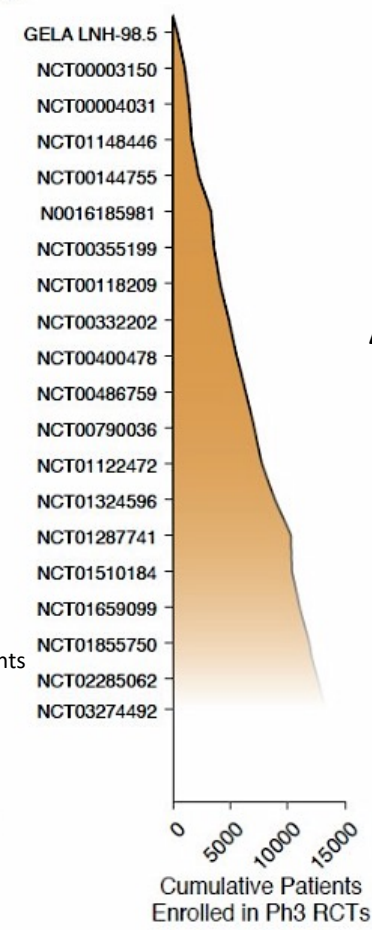
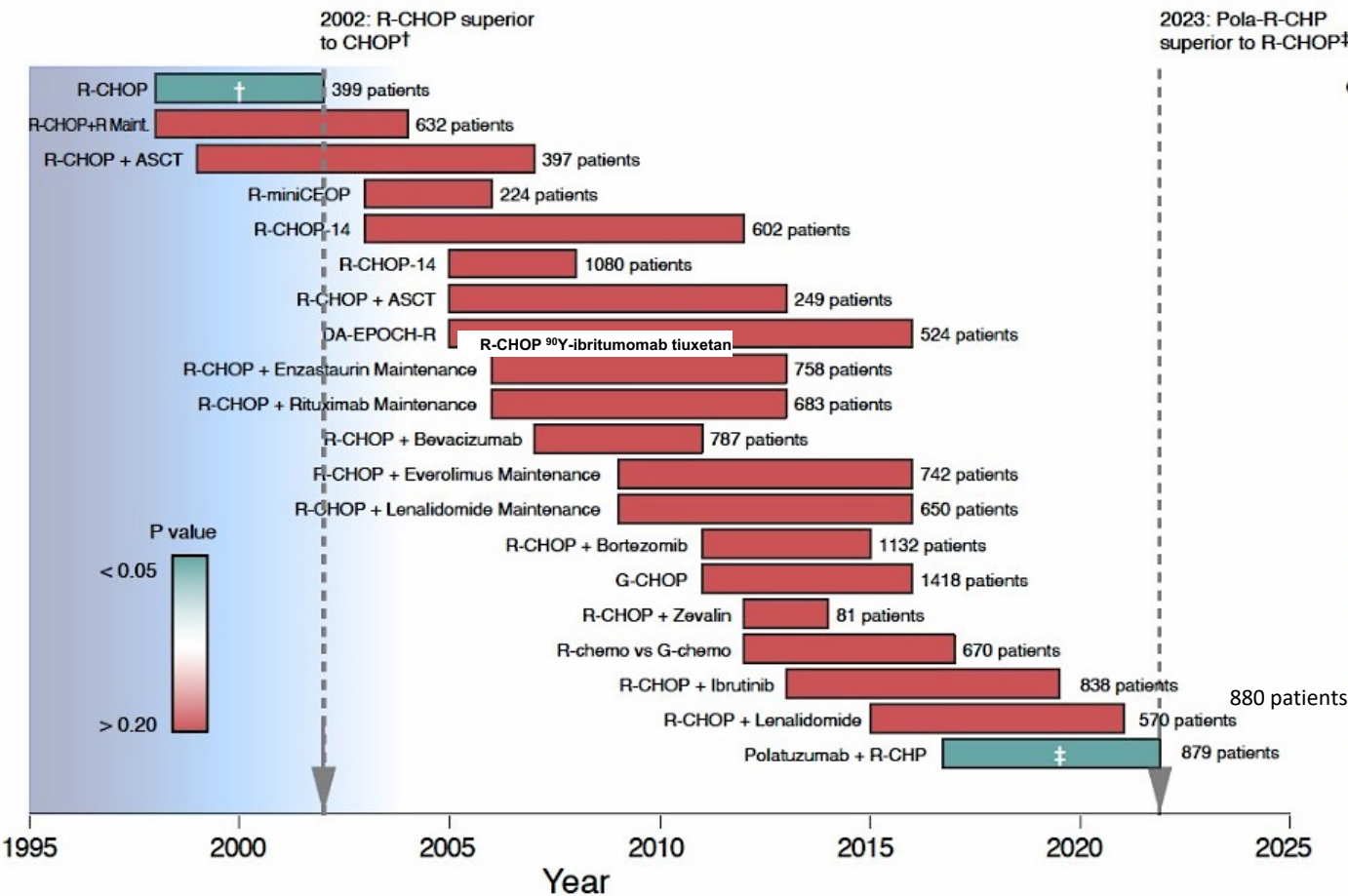
# POLARIX: costo-efectividad de Polatuzumab Vedotin + R-CHP comparado con R-CHOP

Treatment	Life, yr	Effectiveness, QALYs	Incremental effectiveness	Cost Effectiveness		
				Cost in 2021, \$	Increase in cost, \$	Incremental cost effectiveness ratio, \$/QALY
R-CHOP	11.4	10.89	--	342,833	--	--
Pola-R-CHP	11.9	11.80	0.91	419,769	76,936	84,308

- Compared with R-CHP, Pola-R-CHP is **cost-effective (\$84,308)** at a willingness-to-pay (WTP) level of \$150,000/QALY, **if a PFS improvement is maintained with Pola-R-CHP in the long term (5-yr PFS: 70%)**
- Pola-R-CHP has an **incremental effectiveness** of 1.23 life-years and 1.16 QALYs
- Cost effectiveness of Pola-R-CHP depends largely on the **high cost of CAR T-cell therapy**
- Further efforts are needed **to reduce the cost of pola** and other novel therapies in DLBCL.

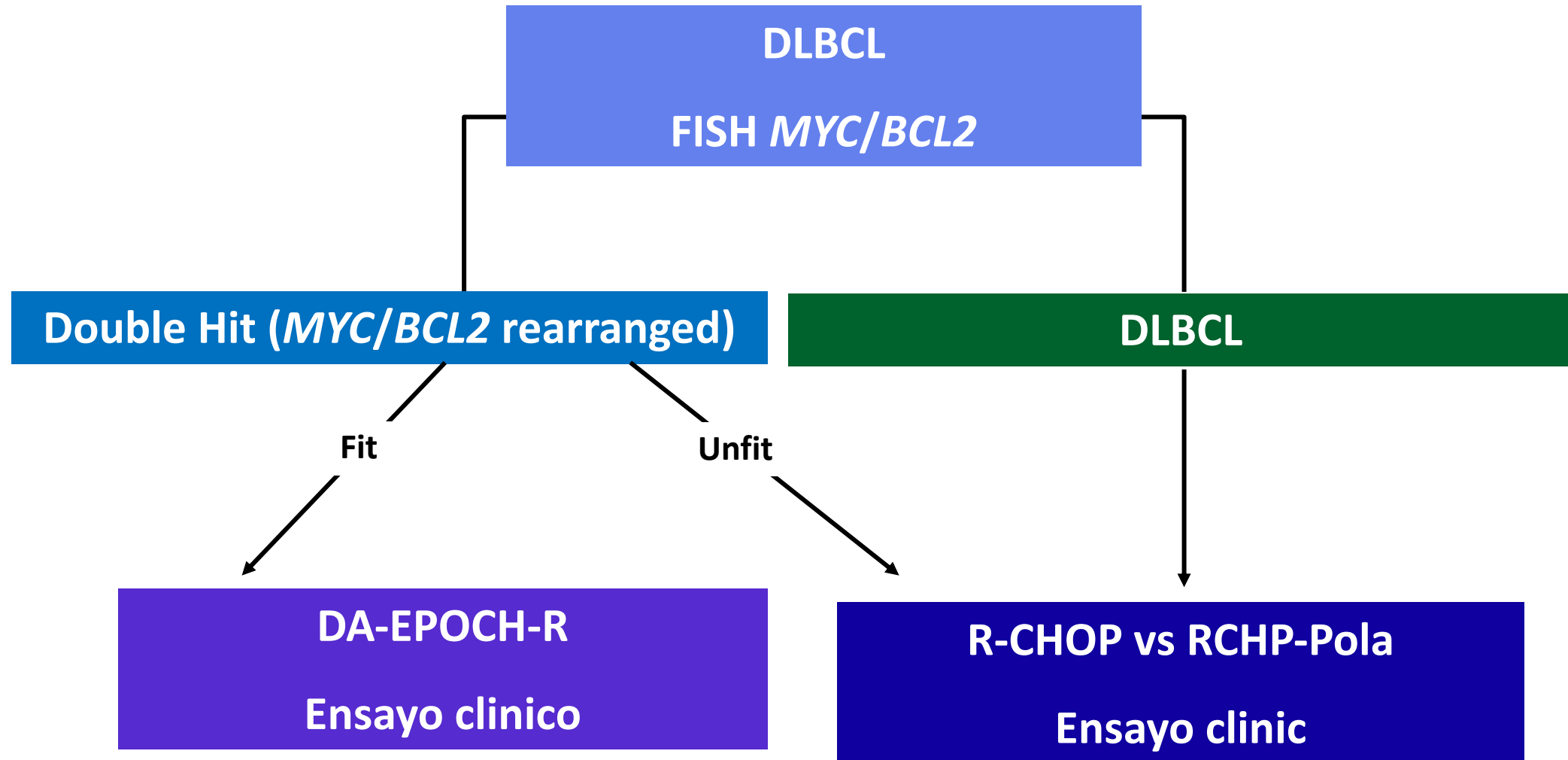


# ¿Más allá de RCHOP – Terapia guiada por perfil Molecular o agnóstico?



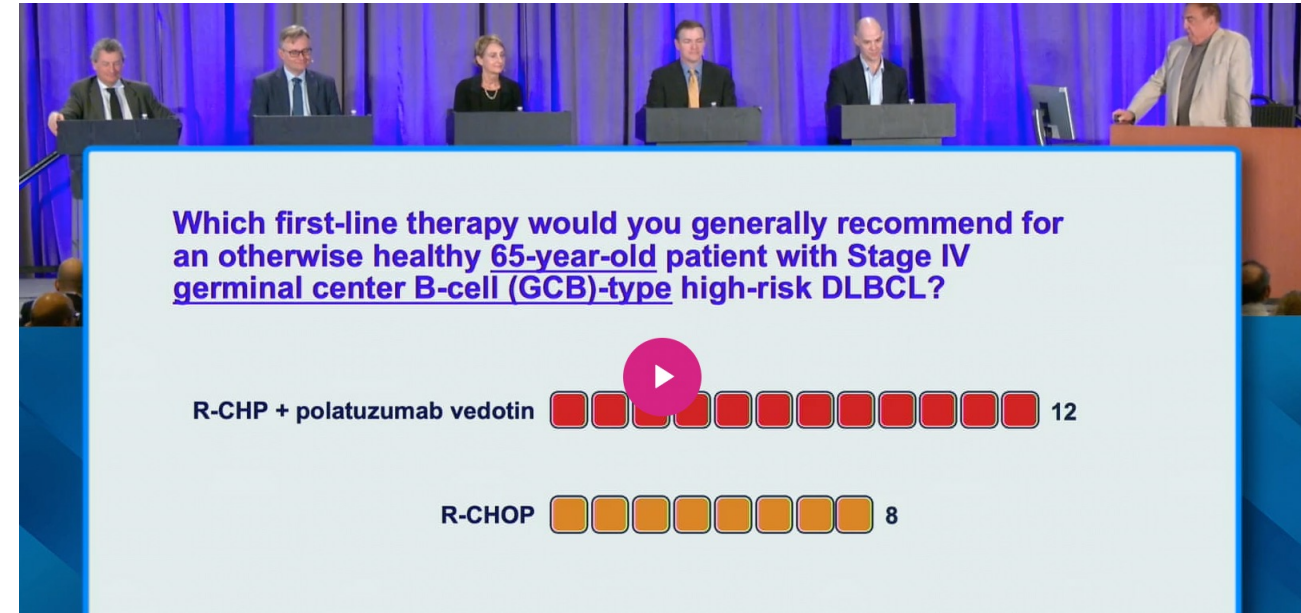
- | Molecularly Agnostic  | Molecularly selected   |
|---|--|
| <ul style="list-style-type: none"> <li>CAR T-cell therapy</li> <li>Tafasitamab/lenalidomide</li> <li>Bispecifics</li> <li>COO agnostic small molecules</li> </ul> | <ul style="list-style-type: none"> <li>BTK inhibitors</li> <li>PI3K inhibitors</li> <li>BCL2 inhibitors</li> <li>Others</li> </ul> |

# Mi enfoque en este caso



# Conclusion

- La terapia de primera línea para (DLBCL) **con Pola R-CHP es prometedora**
  - Hay margen para mejorar
  - No hay suficiente evidencia para excluir GCB del beneficio de Pola RCHP
- El uso de terapias **guiadas por el perfil molecular/ctDNA/MTV** es importante para optimizar y personalizar la terapia
- Debemos analizar la costo efectividad en nuestro contexto



C), genetic classification is yet to be optimized for routine clinical application.<sup>12</sup>