

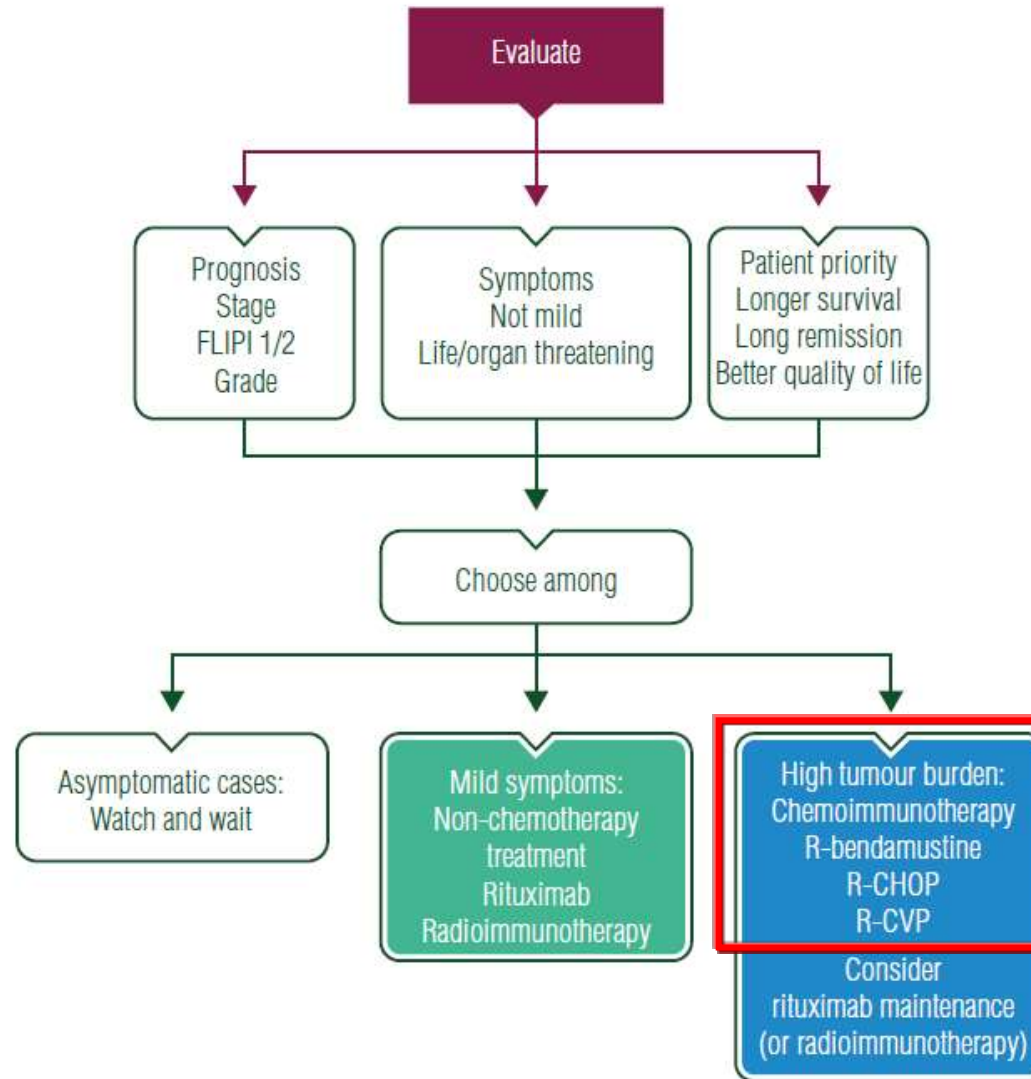
Incontro di aggiornamento sui disordini linfoproliferativi  
e sui protocolli della Fondazione Italiana Linfomi  
**Torino, 24 novembre 2017**

**Nuove prospettive  
nella terapia di prima linea**

Luca Arcaini

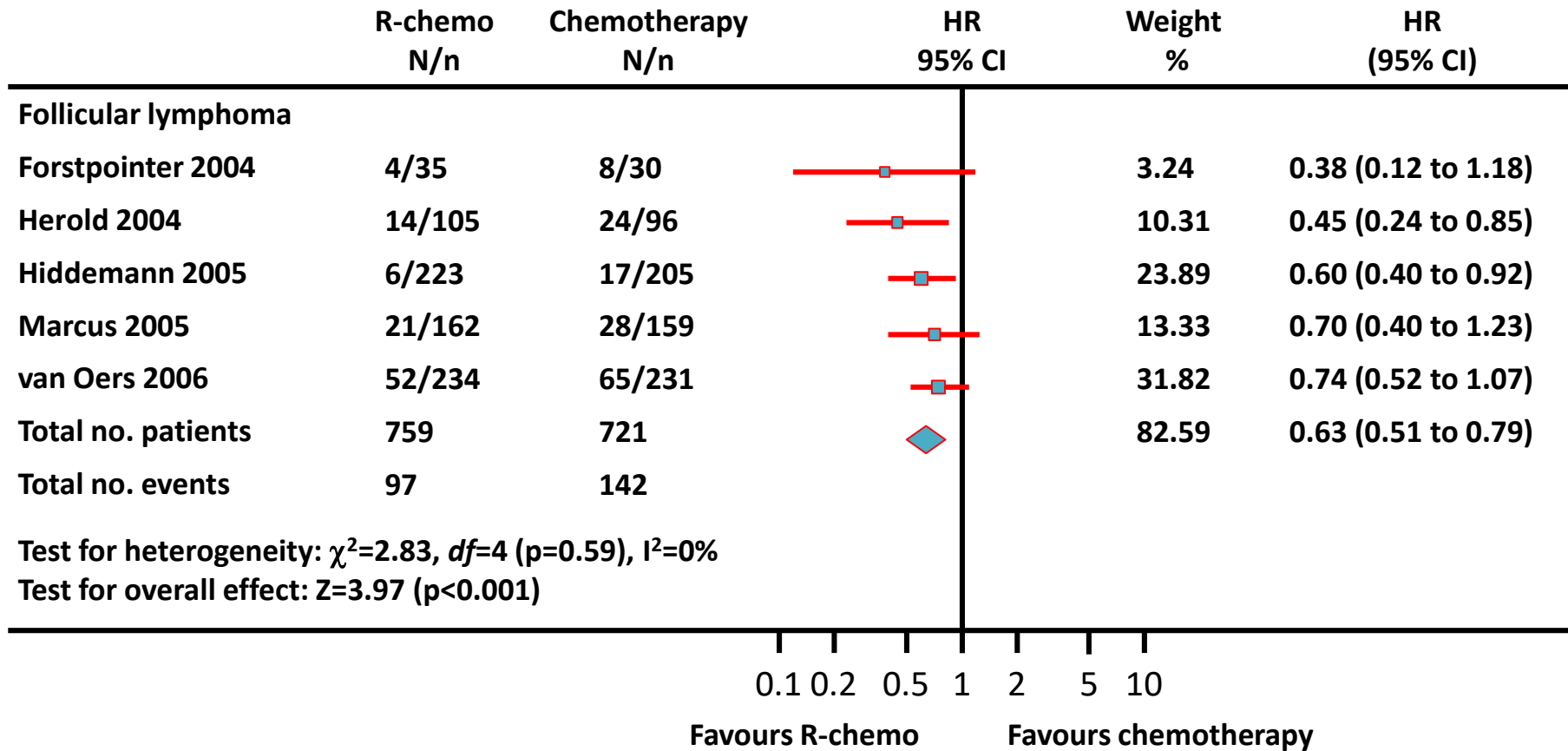
Department of Hematology Oncology, Fondazione IRCCS Policlinico San Matteo &  
Department of Molecular Medicine, University of Pavia, Italy

# First line treatment in FL Therapeutic algorithm

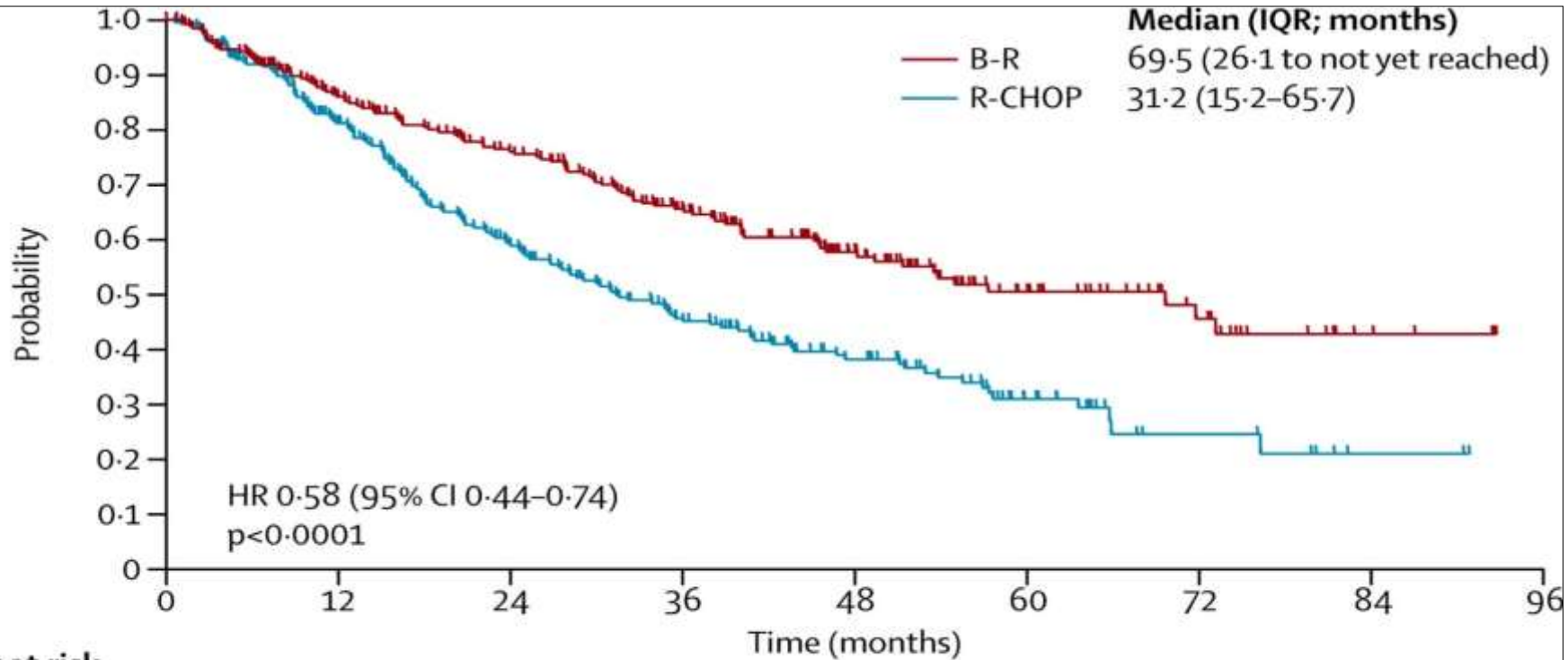


# Immunochemotherapy in high tumor burden patients

## Meta-analysis of chemotherapy vs R-chemotherapy Overall survival



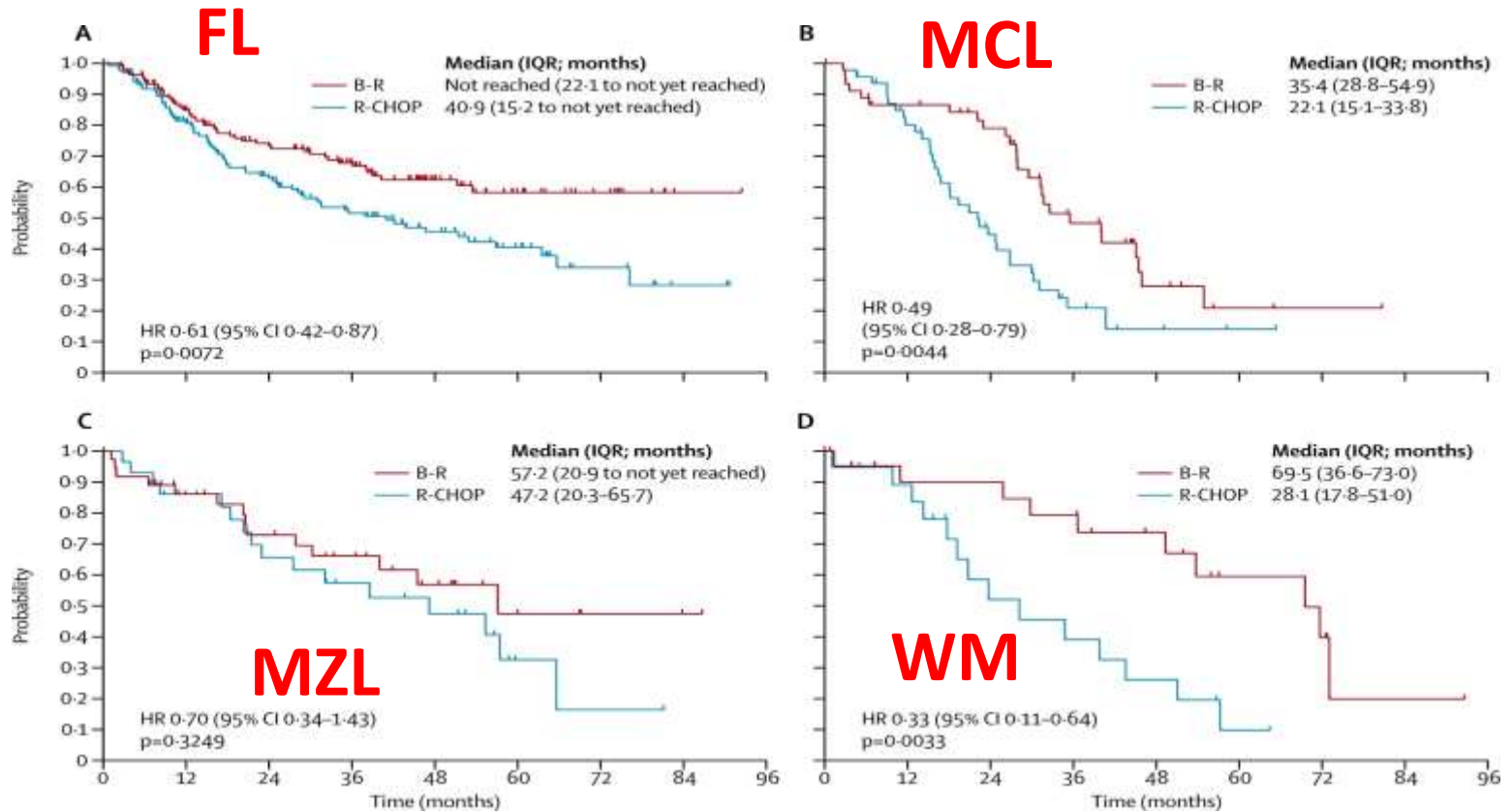
# Stil study: PFS



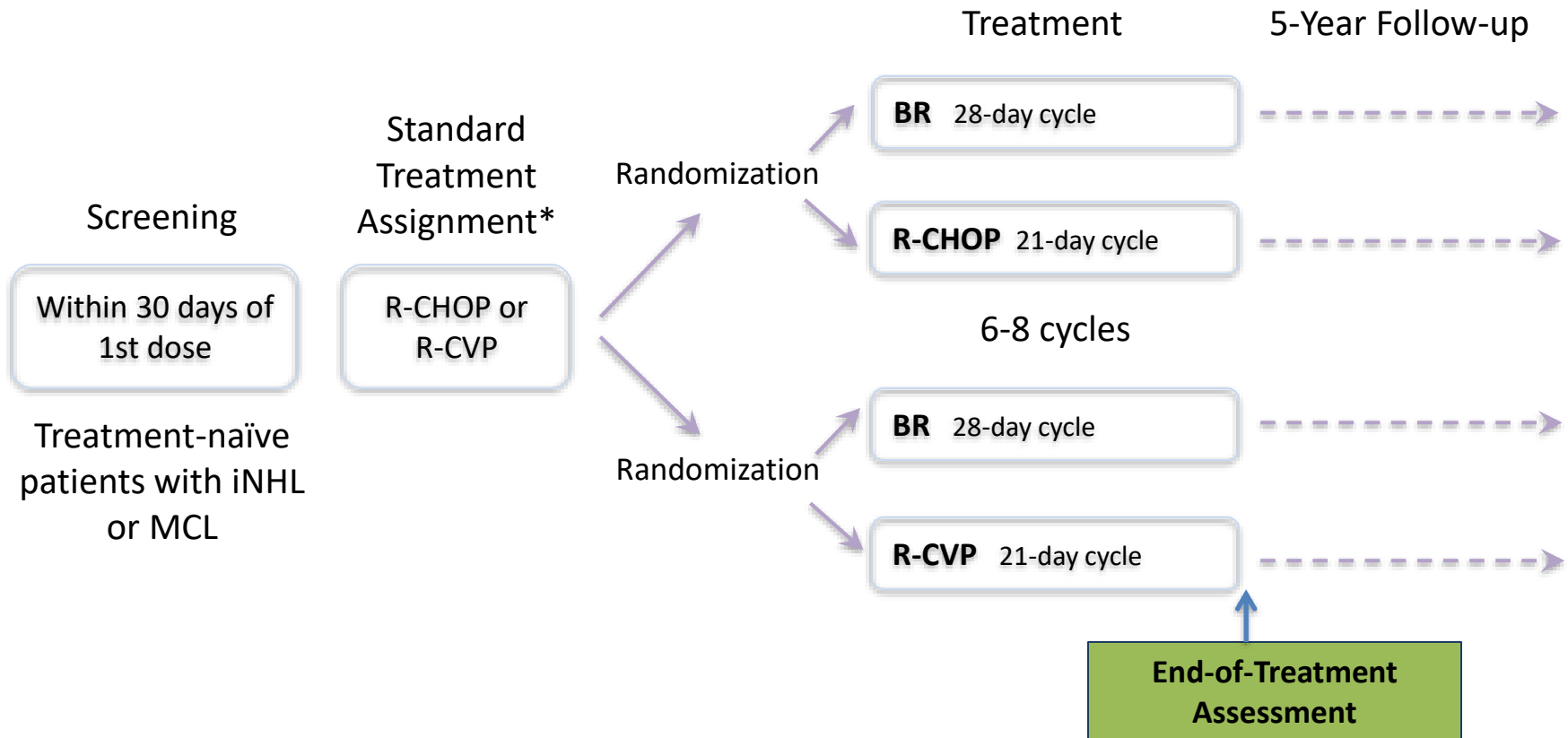
**Number at risk**

	0	12	24	36	48	60	72
B-R	207	169	125	71	35	19	
R-CHOP	185	123	83	54	24	9	

# PFS for histotype



# BRIGT Study Design



\*Based on investigator decision.

B: bendamustine; CHOP: cyclophosphamide, doxorubicin, vincristine, and prednisone;  
CVP: cyclophosphamide, vincristine, and prednisone; iNHL: indolent non-Hodgkin lymphoma;  
MCL: mantle cell lymphoma; R: rituximab.

# Background

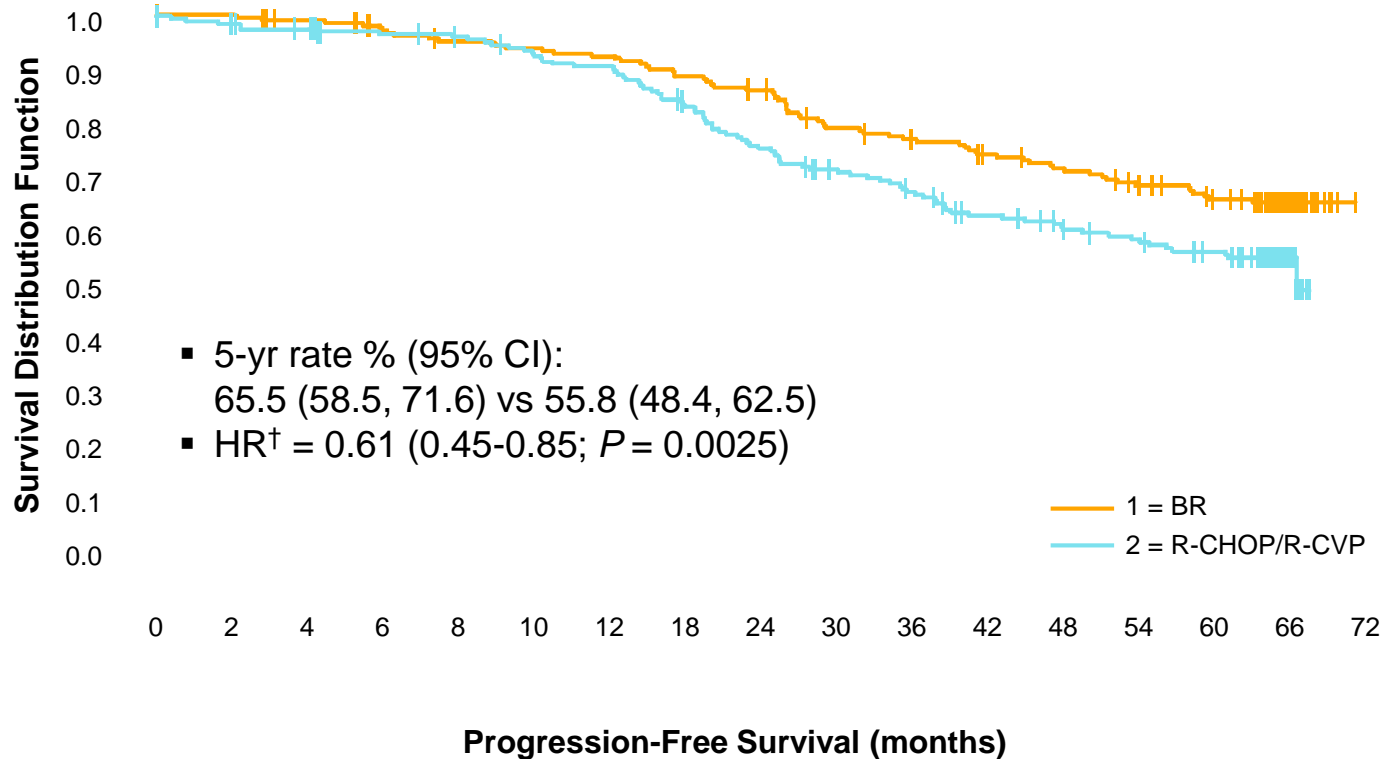
- Initial study results
  - Primary endpoint\*
    - CR: **31%** BR vs **25%** R-CHOP/R-CVP
    - CR rate ratio 1.26;  $P = 0.0225$  for non-inferiority
  - Safety\*
    - BR: ↑ hypersensitivity, vomiting and nausea, lymphocytopenia
    - R-CHOP/R-CVP: ↑ peripheral neuropathy, alopecia, neutropenia
  - Quality of life<sup>†</sup>
    - Improved for BR patients vs R-CHOP/R-CVP in the domains of QLQ-C30 which include Cognitive, Physical, Social, and Emotional Functioning, and GHS as well as reduction in dyspnea, constipation, and fatigue

\**Blood*. 2014;123(19):2944-2952; powered for non-inferiority of CR ratio.

<sup>†</sup>*Clin Lymphoma Myeloma Leuk*. 2016;16(4):182-190.

CR: complete response; GHS: global health status;  
QLQ-C30: 30-question quality-of-life questionnaire.

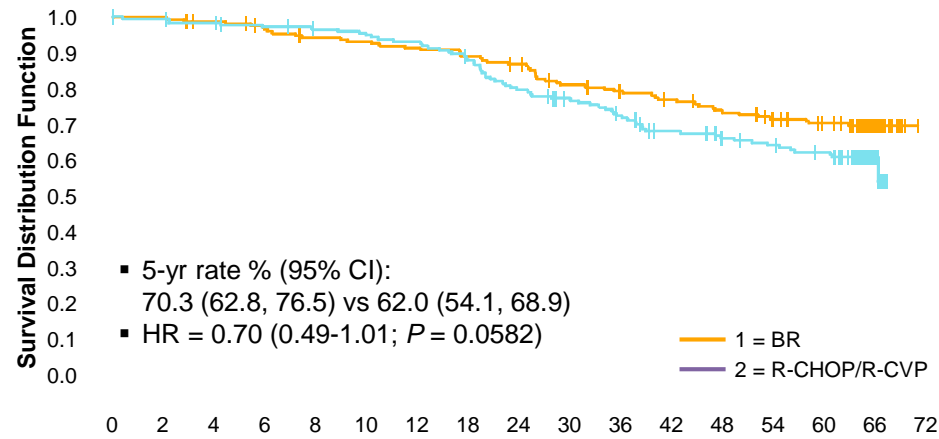
# Progression-Free Survival by Treatment Group\*



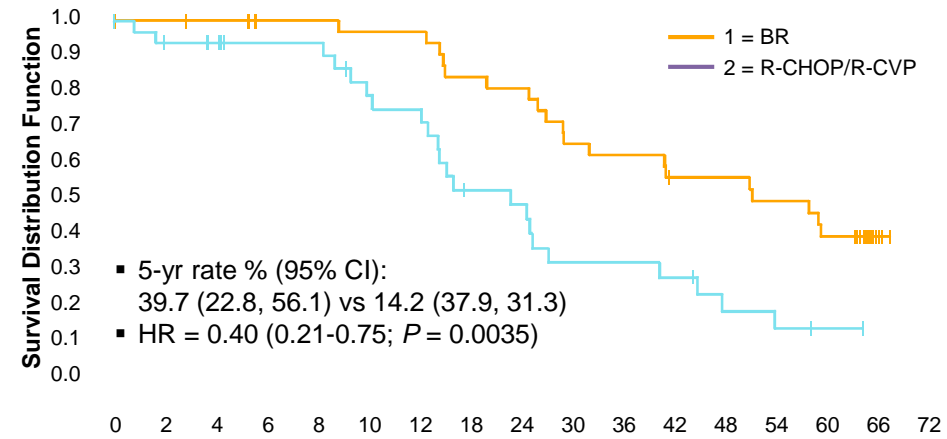


# Progression-Free Survival by Lymphoma Type\*

iNHL†



MCL

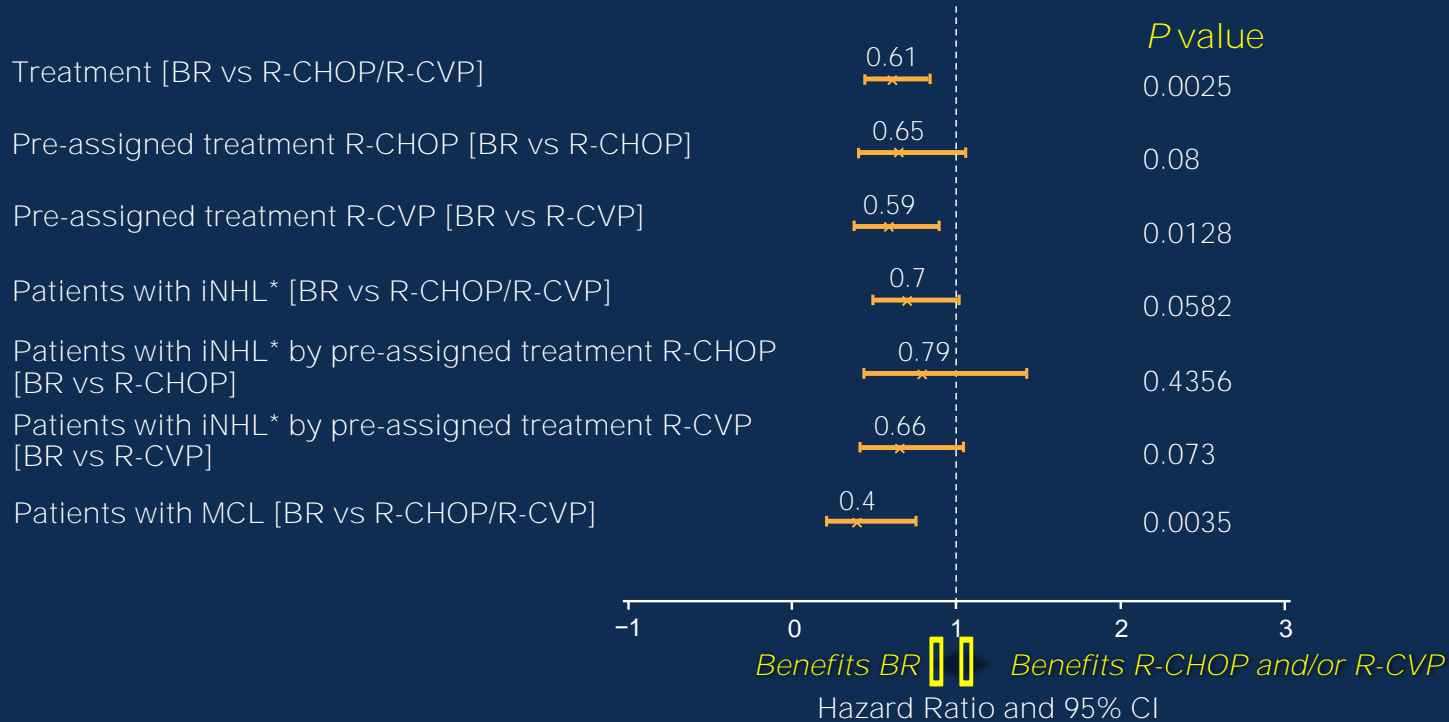


Progression-Free Survival (months)

\*BR vs R-CHOP/R-CVP.

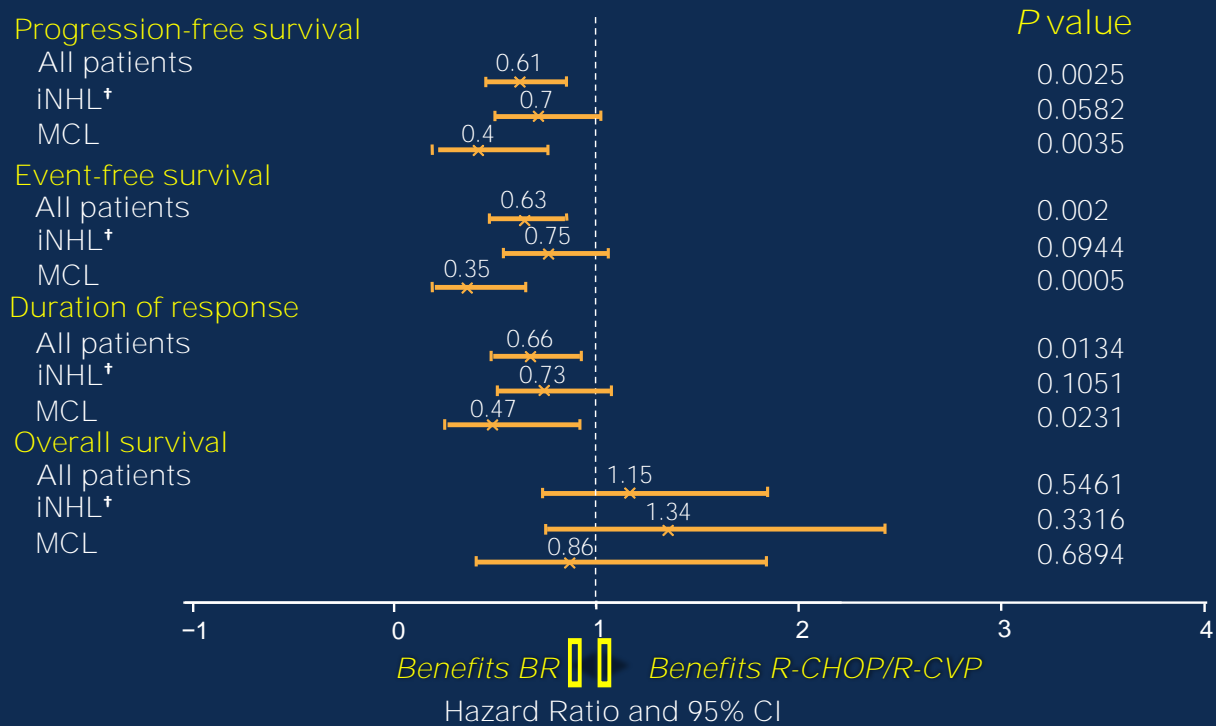
†Not including MCL.

# Forest Plot of PFS Subgroup Analyses



\*Not including MCL.

# Forest Plot of Time-to-Event Variables\*



\*BR vs R-CHOP/R-CVP.

†Not including MCL.

# Patient Mortality

Reported Causality	BR (n = 224)	R-CHOP/R-CVP (n = 223)
Disease progression	16	17
Other* and reason not reported	3	6
Cardiovascular	7	2
Respiratory	3	1
Infection	6	3
Secondary malignancy (excluding transformed NHL)	5	3
All causes <sup>†</sup>	40	32
Deaths up to 100 days of last dose	pneumonia (2) cardiac arrest (1) respiratory failure (1)	septic shock (1) disease progression (2)

\*Complications of stem cell transplant.

<sup>†</sup>Not statistically significant.

# Secondary Malignancy\*

	BR (n = 221)	R-CHOP/R-CVP (n = 215)	
Transformed NHL/DLBCL	5	7	
Basal cell carcinoma	9	4	
Squamous carcinoma of the skin	12	2	
Melanoma	2	1	
MDS	1	1	
Other solid malignancy	19	11	
Patients with secondary malignancy	42 (19%)	24 (11%)	<i>P</i> = 0.022
<b>Excluding NHL and non-melanoma skin cancer</b>	22 (10%)	13 (6%)	<i>P</i> = 0.133

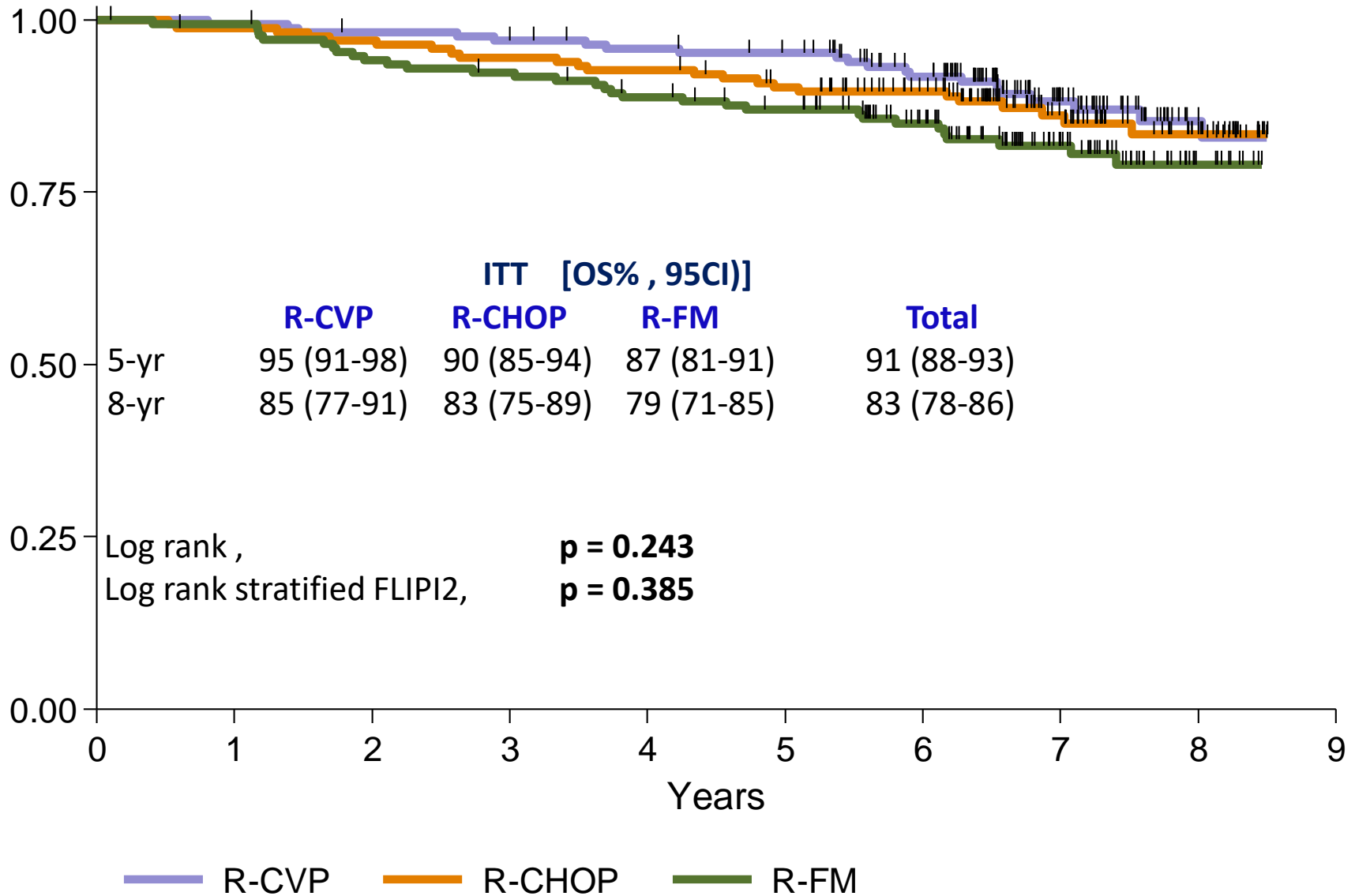
\*Exploratory analysis; histology not collected.

DLBCL: diffuse large B-cell lymphoma; MDS: myelodysplastic syndrome.

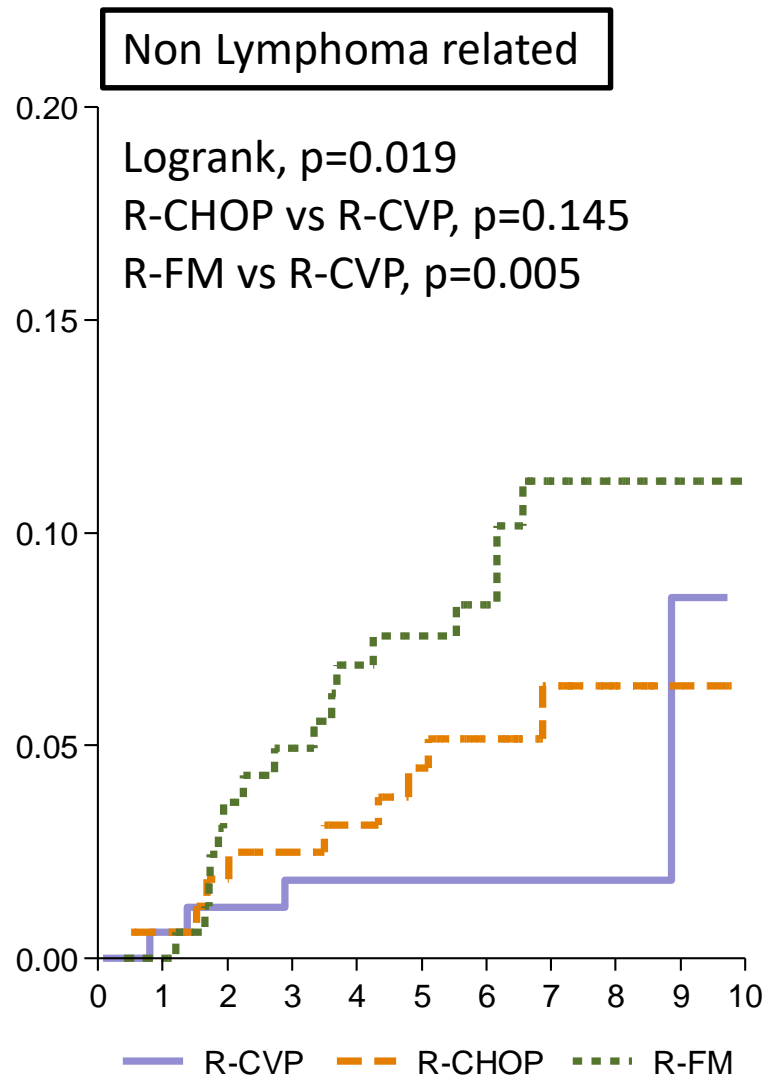
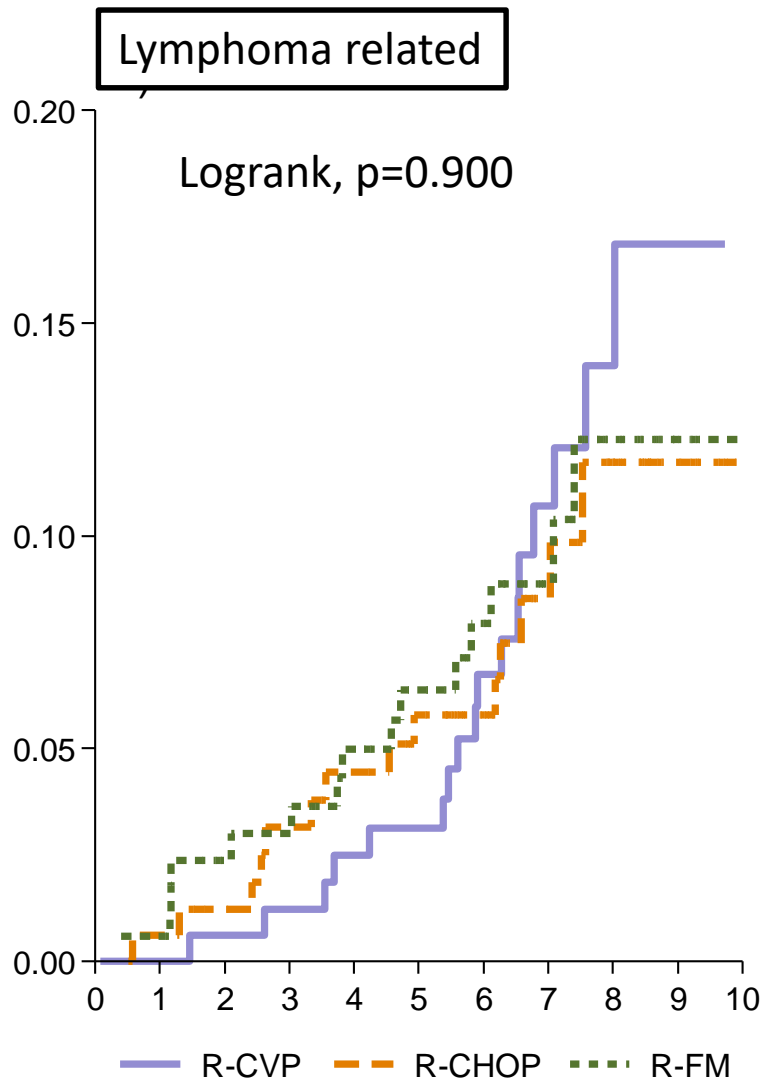
## Bright study

- PFS, EFS, and DOR were significantly in favor of BR vs R-CHOP/R-CVP regimen
  - Greater benefit vs R-CVP
  - Strongest benefit in MCL subgroup
- No difference in OS
- Overall safety profile was as previously reported, with the exception of a higher incidence of secondary malignancies in the BR treatment group

# FOLL05 Study Update: OS



# FOLL05 Study Update: Cause-specific mortality

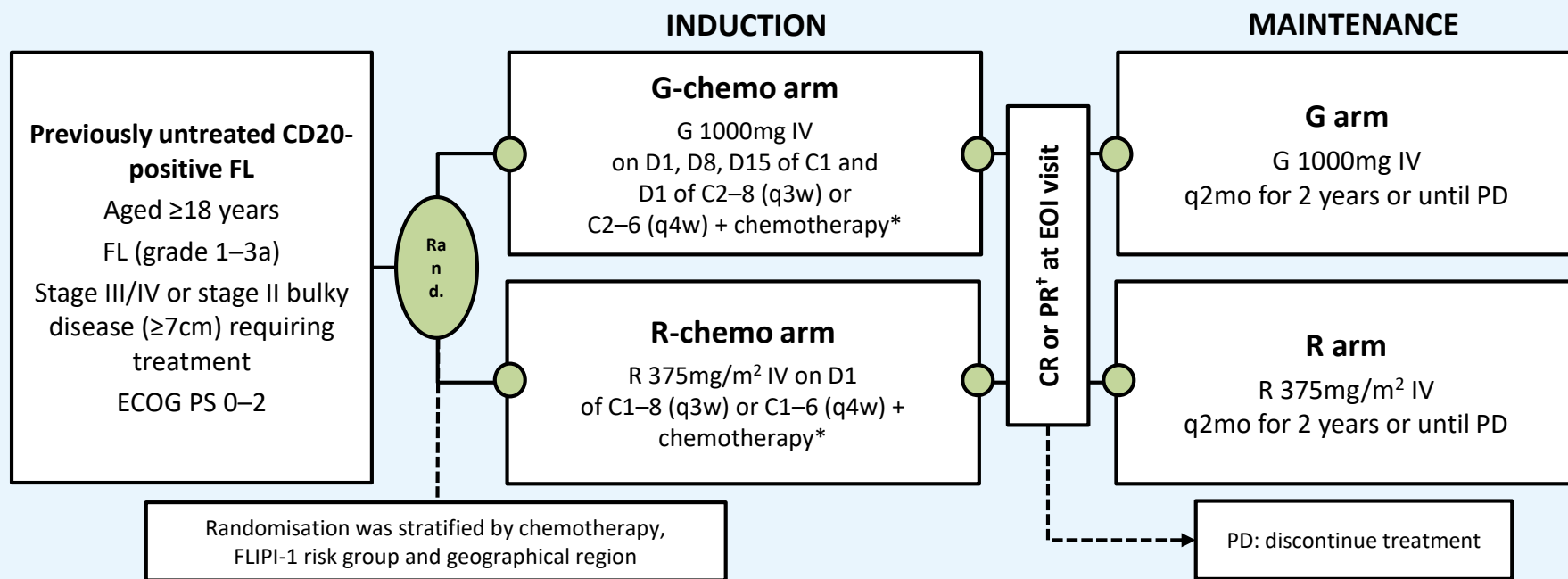


Follow-up, Years



# GALLIUM study design (FL)

International, open-label, randomised Phase III study in 1L pts (NCT01332968)



## Primary endpoint

- PFS (INV-assessed)

## Secondary endpoints

- PFS (IRC-assessed)
- OS, EFS, DFS, DoR, TTNALT
- ORR/CR at EOI (+/- FDG-PET)
- Safety
- PROs

\*CHOP q3w × 6 cycles, CVP q3w × 8 cycles, bendamustine q4w × 6 cycles; chemo regimen chosen by site prior to initiation and received by all FL pts at site; †patients with SD at EOI entered observation for up to 2 years or until PD if earlier

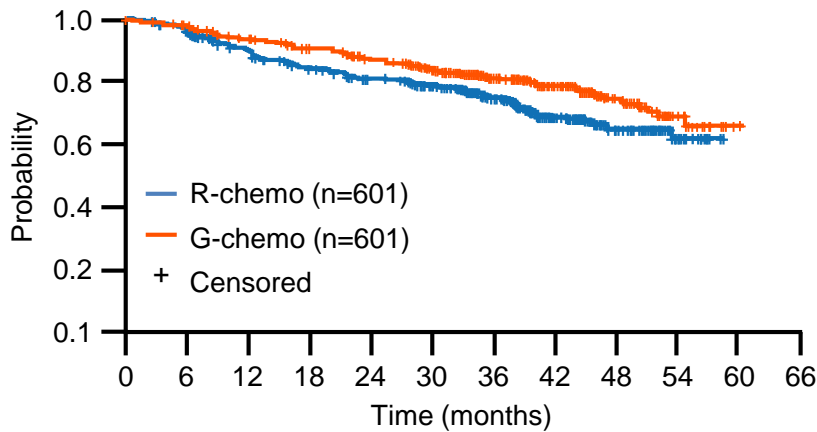
# Baseline characteristics\*

<i>n (%)</i>	<i>R-chemo, n=601</i>	<i>G-chemo, n=601</i>
Median age, years (range)	58.0 (23–85)	60.0 (26–88)
Male	280 (46.6)	283 (47.1)
Ann Arbor stage at diagnosis		
I	8 (1.3) <sup>†</sup>	10 (1.7) <sup>‡</sup>
II	44 (7.4) <sup>†</sup>	41 (6.9) <sup>‡</sup>
III	208 (34.8) <sup>†</sup>	209 (34.9) <sup>‡</sup>
IV	337 (56.4) <sup>†</sup>	338 (56.5) <sup>‡</sup>
FLIPI risk group		
Low (0–1)	125 (20.8)	127 (21.1)
Intermediate (2)	223 (37.1)	225 (37.4)
High (≥3)	<b>253 (42.1)</b>	<b>249 (41.4)</b>
Bone marrow involvement	295 (49.3) <sup>‡</sup>	318 (53.7) <sup>§</sup>
Extranodal involvement	396 (65.9)	392 (65.2)
Bulky disease (≥7cm)	<b>271 (45.2)<sup>¶</sup></b>	<b>255 (42.5)<sup>¶</sup></b>
Median t from dg to random, mo (range)	<b>1.4 (0–168.1)<sup>‡</sup></b>	<b>1.5 (0.1–121.6)<sup>‡</sup></b>

\*ITT population; <sup>†</sup>n=597; <sup>‡</sup>n= 598; <sup>§</sup>n=592; <sup>¶</sup>n=600

# PFS after 41.1 mo median f-up\*

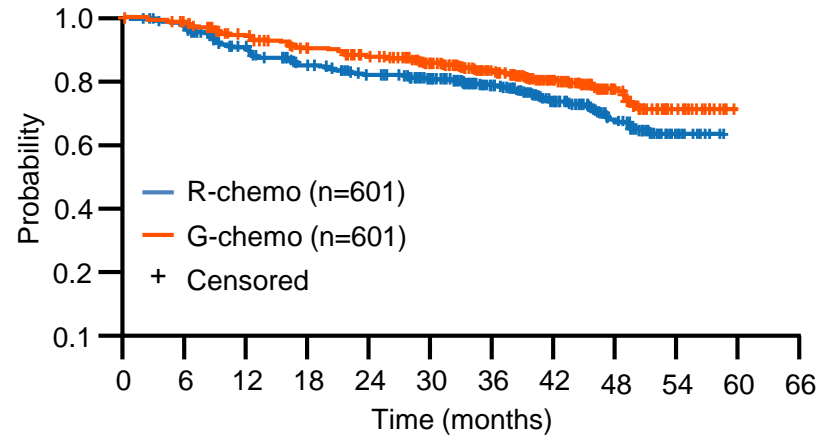
## INV-assessed PFS



No. of patients at risk

G-chemo	601	561	505	464	438	396	267	149	77	18
R-chemo	601	569	535	505	478	420	291	176	85	25

## IRC-assessed PFS



No. of patients at risk

G-chemo	601	563	502	463	438	394	271	151	73	16
R-chemo	601	571	532	497	476	414	287	179	79	22

**R-chemo, n=601**      **G-chemo, n=601**

3-yr PFS, % (95% CI)	75.0 (71.0, 78.5)	81.5 (77.9, 84.6)
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HR (95% CI), p-value<sup>†</sup>      0.68 (0.54, 0.87), **p=0.0016**

**R-chemo, n=601**      **G-chemo, n=601**

3-yr PFS, % (95% CI)	78.9 (75.2, 82.1)	83.4 (79.9, 86.3)
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HR (95% CI), p-value<sup>†</sup>      0.72 (0.56, 0.93), **p=0.0118**

\*ITT population; <sup>†</sup>stratified analysis; stratification factors = FLIPI, chemotherapy regimen

# Adverse events\*

<i>n (%) of pts reporting <math>\geq 1</math> one event</i>	<i>R-chemo, n=597</i>	<i>G-chemo, n=595</i>
Any AE	585 (98.0)	593 (99.7)
Grade 3–5 AEs	409 (68.5)	<b>449 (75.5)</b>
SAE	246 (41.2)	<b>281 (47.2)</b>
Grade 5 (fatal) AE	21 (3.5)	24 (4.0)
AE leading to treatment discontinuation	88 (14.7)	98 (16.5)

- All-cause deaths in each arm: R-chemo, 52 (8.7%); G-chemo, 42 (7.1%)

\*Safety population

# Selected grade 3–5 AEs of particular interest (frequency >2%)\*

<i>n (%) of pts reporting <math>\geq 1</math> one event</i>	<i>R-chemo, n=597</i>	<i>G-chemo, n=595</i>
Neutropenia	236 (39.5)	<b>278 (46.7)</b>
Infections <sup>†</sup>	98 (16.4)	<b>121 (20.3)</b>
Infusion-related reactions <sup>‡</sup>	40 (6.7)	<b>74 (12.4)</b>
Thrombocytopenia	16 (2.7)	<b>36 (6.1)</b>
Second malignancies (SMQ) <sup>§</sup>	21 (3.5)	29 (4.9)
Cardiac events	17 (2.8)	23 (3.9)

\*AEPIs occurring in >2% of patients in safety population, in either treatment arm

<sup>†</sup>System Organ Class 'Infections and Infestations'

<sup>‡</sup>Related to study treatment and occurring during or in the 24 hours after infusion

<sup>§</sup>Standardised MedDRA query, i.e. malignant or unspecified tumours occurring >6 mo after study drug intake

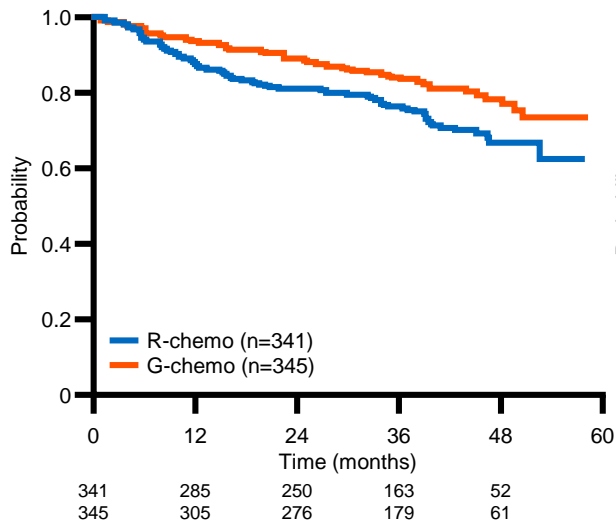
# Baseline characteristics by chemo\*

<i>n (%)</i>	<i>Benda, n=686</i>	<i>CHOP, n=399</i>	<i>CVP, n=117</i>
Median age, years (range)	59 (23–88)	58 (31–85)	59 (32–85)
Age ≥80 years	<b>23 (3.4)</b>	3 (0.8)	<b>4 (3.4)</b>
Male	332 (48.4)	177 (44.4)	54 (46.2)
Charlson Comorbidity Index score ≥1 <sup>†</sup>	<b>163 (23.8)</b>	69 (17.3)	22 (18.8)
ECOG PS 2	24 (3.5)	8 (2.0)	6 (5.1)
FLIPI high risk (≥3)	274 (39.9)	<b>187 (46.9)</b>	41 (35.0)
Bulky disease (≥7cm)	274 (39.9)	<b>206 (51.6)</b>	46 (39.3)

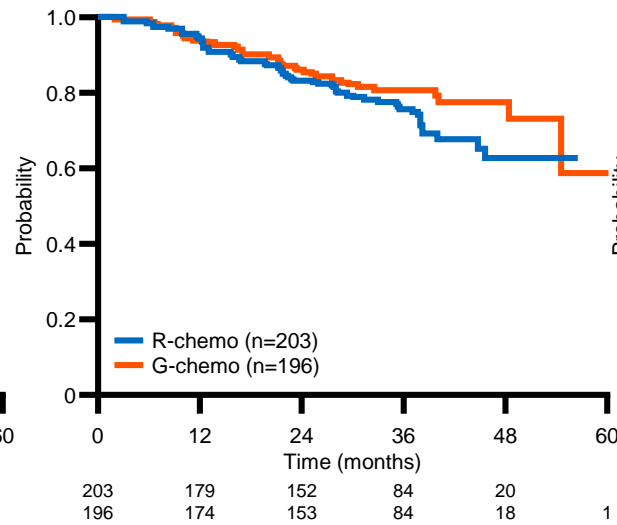
\*ITT population; <sup>†</sup>scored retrospectively based on conditions reported on medical history page of CRF

# INV-assessed PFS by chemo\*

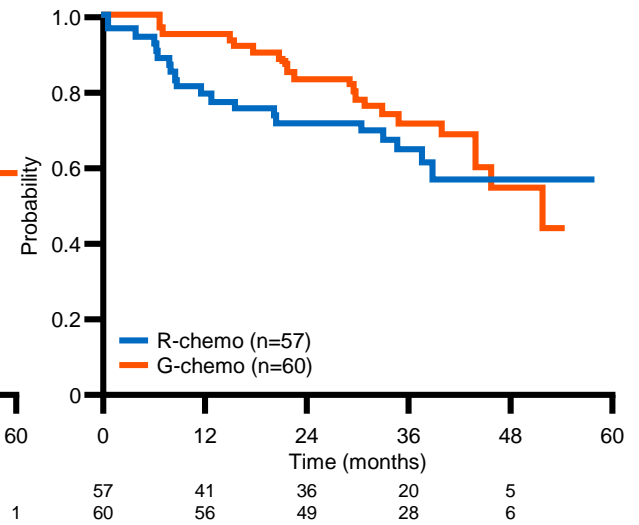
**Benda**



**CHOP**



**CVP**



HR (95% CI)<sup>†</sup> 0.63 (0.46, 0.88)

3-yr PFS 84.1% G-B vs 76.4% R-B

HR (95% CI)<sup>†</sup> 0.72 (0.48, 1.10)

3-yr PFS 80.6% G-CHOP vs 75.6% R-CHOP

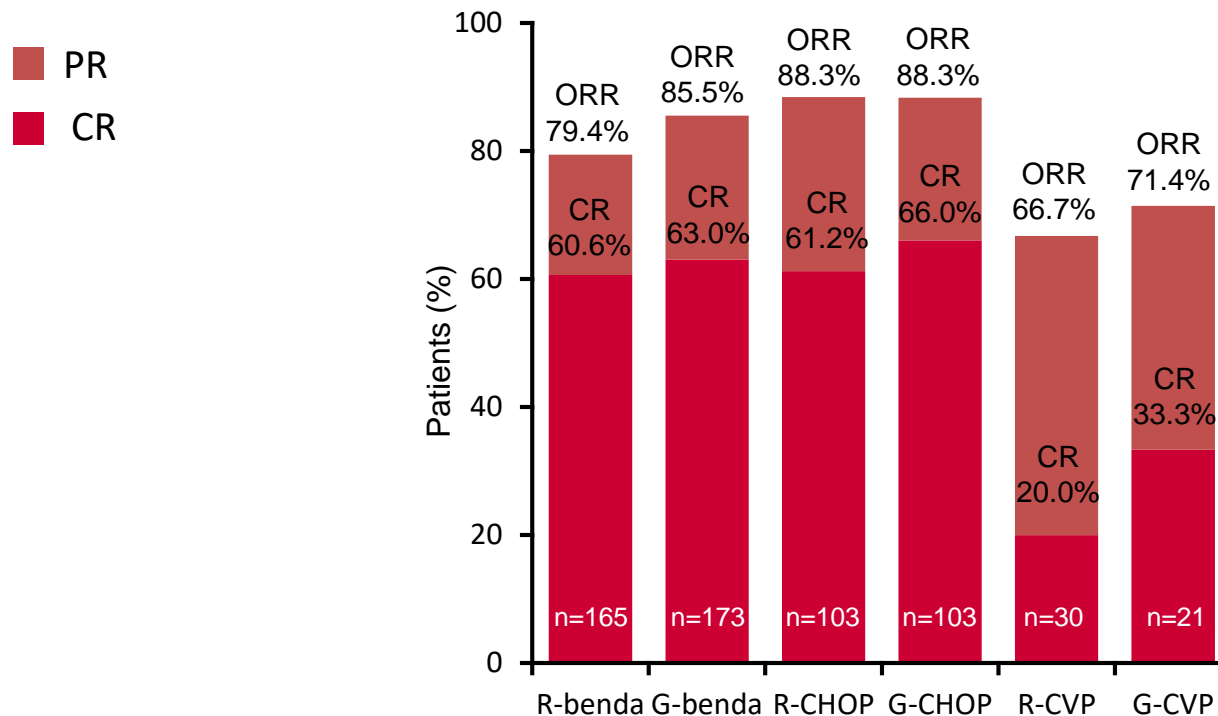
HR (95% CI)<sup>†</sup> 0.79 (0.42, 1.47)

3-yr PFS 71.3% G-CVP vs 64.2% R-CVP

- By chemo analysis not powered to demonstrate statistically significant differences between treatment arms
- \*ITT population; <sup>†</sup>analysis stratified by FLIPI (as well as chemotherapy regimen)

# INV-assessed response at end of induction

*CT- and PET-based\*†*



\*PET ITT population i.e. all randomised FL pts who had PET-avid lesions representing lymphoma at baseline

†assessed according to Cheson 2007 criteria



# AEs by treatment arm\*

<i>n (%) of pts reporting ≥1 event</i>	<i>R-benda, n=338</i>	<i>G-benda, n=338</i>	<i>R-CHOP, n=203</i>	<i>G-CHOP, n=193</i>	<i>R-CVP, n=56</i>	<i>G-CVP, n=61</i>
Any AE	331 (97.9)	338 (100)	201 (99.0)	191 (99.0)	56 (100)	61 (100)
Grade 3–5 AE	228 (67.5)	233 (68.9)	<b>151 (74.4)</b>	<b>171 (88.6)</b>	<b>30 (53.6)</b>	<b>42 (68.9)</b>
SAE	160 (47.3)	176 (52.1)	<b>67 (33.0)</b>	<b>76 (39.4)</b>	<b>19 (33.9)</b>	<b>26 (42.6)</b>
Grade 5 (fatal) AE	16 (4.7)	20 (5.9)	4 (2.0)	3 (1.6)	1 (1.8)	1 (1.6)
AE leading to treatment discontinuation	48 (14.2)	52 (15.4)	31 (15.3)	32 (16.6)	9 (16.1)	11 (18.0)

- Difference between treatment arms in grade 3–5 AEs and SAEs was more pronounced in patients treated with CHOP and CVP
- Comparisons confounded by imbalances in baseline patient and disease characteristics between chemo groups

\*Safety population, i.e. all randomised FL pts who received at least one dose of study drug

# AEs by chemo\*

<i>n (%) of pts reporting ≥1 event</i>	<i>R-benda, n=338</i>	<i>G-benda, n=338</i>	<i>R-CHOP, n=203</i>	<i>G-CHOP, n=193</i>	<i>R-CVP, n=56</i>	<i>G-CVP, n=61</i>
Any AE	331 (97.9)	338 (100)	201 (99.0)	191 (99.0)	56 (100)	61 (100)
Grade 3–5 AE	228 (67.5)	233 (68.9)	<b>151 (74.4)</b>	<b>171 (88.6)</b>	30 (53.6)	42 (68.9)
SAE	<b>160 (47.3)</b>	<b>176 (52.1)</b>	67 (33.0)	76 (39.4)	19 (33.9)	26 (42.6)
Grade 5 (fatal) AE <sup>†</sup>	<b>16 (4.7)</b>	<b>20 (5.9)</b>	4 (2.0)	3 (1.6)	1 (1.8)	1 (1.6)
AE leading to treatment discontinuation	48 (14.2)	52 (15.4)	31 (15.3)	32 (16.6)	9 (16.1)	11 (18.0)

- Grade 3–5 AEs most frequent with CHOP (neutropenia, leukopenia, febrile neutropenia, IRRs); SAEs and fatal AEs most frequent with benda
  - Frequency of grade 5 AEs similar to R-CHOP arms in SABRINA (5.7%, i.v.; 3.6%, s.c.)

\*Safety population, i.e. all randomised FL pts who received at least one dose of study drug; <sup>†</sup>includes 6 pts with fatal AEs that occurred after start of new anti-cancer therapy (G-benda, 4; R-benda, 2)

# Grade 5 (fatal) AEs excluding six pts who started new anti-cancer treatment\*†

<i>SOC</i>	<i>R-benda, n=338</i>	<i>G-benda, n=338</i>	<i>R-CHOP, n=203</i>	<i>G-CHOP, n=193</i>	<i>R-CVP, n=56</i>	<i>G-CVP, n=61</i>
Infections and infestations	1 (0.3)	4 (1.2)	0	1 (0.5)	0	0
Cardiac disorders	2 (0.6)	2 (0.6)	0	0	0	0
Gastrointestinal disorders	0	1 (0.3)	0	0	0	1 (1.6)
General disorders and administration site conditions	2 (0.6)	1 (0.3)	0	1 (0.5)	1 (1.8)	0
Metabolism and nutrition disorders	1 (0.3)	1 (0.3)	0	0	0	0
Neoplasms benign, malignant and unspecified	3 (0.9)	4 (1.1)	2 (1.0)	1 (0.5)	0	0
Nervous system disorders	4 (1.2)	0	2 (1.0)	0	0	0
Respiratory, thoracic and mediastinal disorders	1 (0.3)	3 (0.9)	0	0	0	0
Total (all SOCs)	<b>14 (4.1%)</b>	<b>16 (4.7%)</b>	4 (2.0%)	3 (1.6%)	1 (1.8%)	1(1.6%)

- Many pts with fatal AEs had adverse risk factors, e.g. aged ≥80 years (benda, 6; CVP, 1), ECOG PS grade 2 (benda, 4; CVP, 1), and CCI ≥1 (benda, 11; CHOP, 1; CVP, 1)

\*Safety population; †fatal AEs in 5 G-benda and 1 R-benda pts that occurred after new systemic anti-cancer treatment had started were not included (PD, 4 pts; new malignancies, 2 pts); CCI, Charlson Comorbidity Index

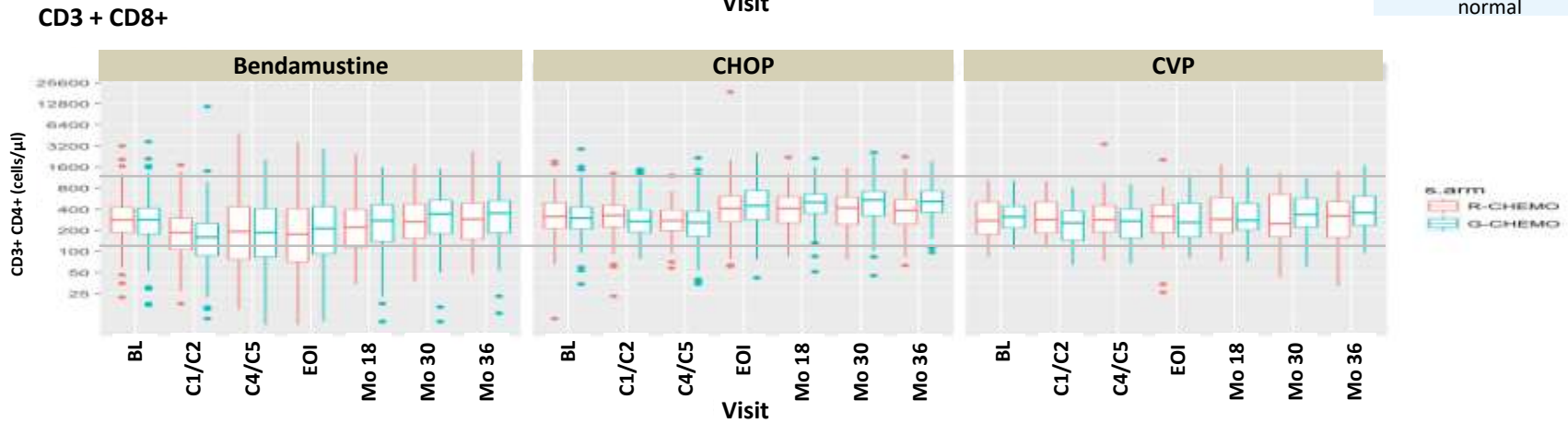
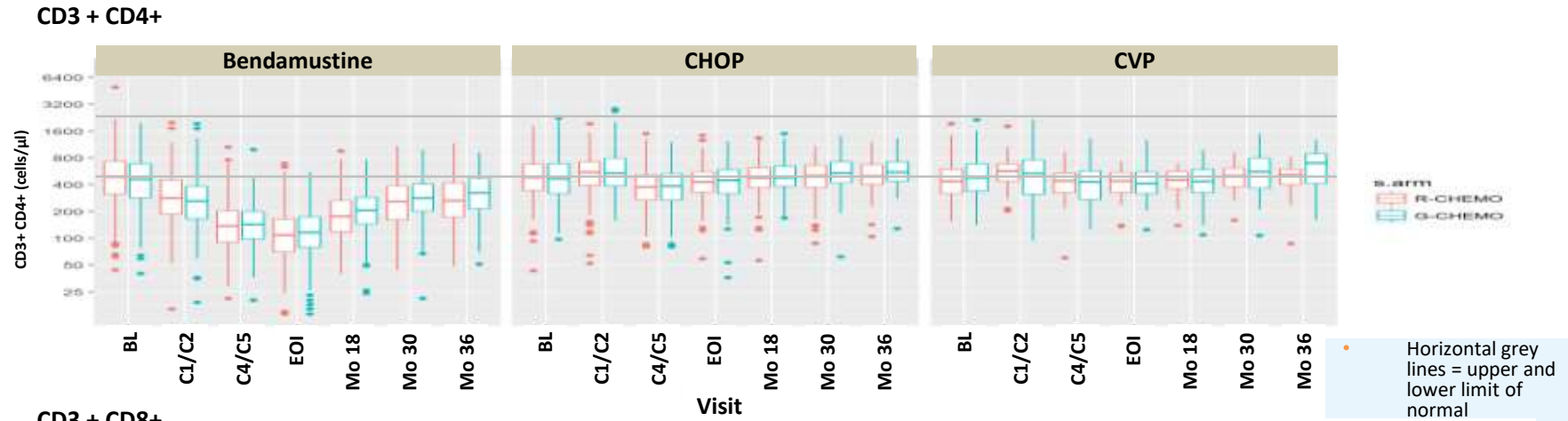
# Selected grade 3–5 AEs of particular interest by chemo\*

<i>n (%) of pts reporting ≥1 event</i>	<i>R-benda, n=338</i>	<i>G-benda, n=338</i>	<i>R-CHOP, n=203</i>	<i>G-CHOP, n=193</i>	<i>R-CVP, n=56</i>	<i>G-CVP, n=61</i>
Cardiac events	12 (3.6)	13 (3.8)	5 (2.5)	6 (3.1)	0 (0.0)	4 (6.6)
Neutropenia	107 (31.7)	107 (31.7)	<b>115 (56.7)</b>	<b>142 (73.6)</b>	14 (25.0)	29 (47.5)
Febrile neutropenia	13 (3.8)	18 (5.3)	<b>14 (6.9)</b>	<b>22 (11.4)</b>	2 (3.6)	2 (3.3)
Second malignancies <sup>†</sup>	<b>12 (3.6)</b>	<b>21 (6.2)</b>	7 (3.4)	7 (3.6)	2 (3.6)	1 (1.6)
Other solid tumours	9 (2.7)	11 (3.3)	7 (3.4)	4 (2.1)	2 (3.6)	0
Hematological tumours <sup>‡</sup>	0	3 (0.9)	0	3 (1.6)	0	0
Non-melanoma skin cancer	<b>3 (0.9)</b>	<b>7 (2.1)</b>	0	0	0	1 (1.6)
Infections	<b>66 (19.5)</b>	<b>89 (26.3)</b>	25 (12.3)	23 (11.9)	7 (12.5)	8 (13.1)
Opportunistic infections <sup>§</sup>	6 (1.8)	10 (3.0)	2 (1.0)	5 (2.6)	0	0

- Frequency of grade 3–5 second malignancy and infections similar in G and R arms for CHOP and CVP groups but not for benda
- Comparisons confounded by imbalances in baseline patient and disease characteristics between chemo groups

\*Safety population; <sup>†</sup>standardised MedDRA query = malignant or unspecified tumours occurring >6 months after first study drug intake; <sup>‡</sup>Hodgkin disease (n=3), AML (n=2), and ALL (n=1); <sup>§</sup>including fungal infections, cytomegalovirus, herpes zoster and pneumocystis jirovecii pneumonia

# T-cell counts over time



*Low T-cell count at baseline*

*R-benda,  
n=341*

*G-benda,  
n=345*

*R-CHOP,  
n=203*

*G-CHOP,  
n=196*

*R-CVP,  
n=57*

*G-CVP,  
n=60*

CD3+/CD4+ cell count of  $\leq 200/\text{mm}^3$

36 (12.5%)

36 (11.4%)

12 (7.2%)

9 (5.1%)

2 (4.4%)

4 (7.4%)

# Grade 3–5 infections by chemo and by phase

<i>n (%) of pts reporting ≥1 event</i>	<i>R-benda, n=338</i>	<i>G-benda, n=338</i>	<i>R-CHOP, n=203</i>	<i>G-CHOP, n=193</i>	<i>R-CVP, n=56</i>	<i>G-CVP, n=61</i>
All study periods	66 (19.5)	89 (26.3)	25 (12.3)	23 (11.9)	7 (12.5)	8 (13.1)
Induction	26 (7.7)	27 (8.0)	13 (6.4)	14 (7.3)	4 (7.1)	3 (4.9)
Maintenance	<b>39 (13.0)</b>	<b>51 (16.7)</b>	11 (5.9)	7 (3.9)	1 (2.5)	5 (8.8)
Observation	12 (3.8)	28 (8.8)	6 (3.1)	3 (1.6)	3 (5.7)	1 (1.7)
<i>N (%) of pts receiving G-CSF prophylaxis</i>	48 (14.2)	54 (16.0)	108 (53.2)	112 (58.0)	13 (23.2)	10 (16.4)

Comparisons confounded by imbalances in baseline patient and disease characteristics between chemo groups

\*Safety population

# Lenalidomide produces responses in patients with R/R FL

## Lenalidomide monotherapy

- 22 with FL Grade 1/2<sup>1</sup>
- Median (range) prior systemic therapies: 3 (1–17)<sup>1</sup>

ORR: 27% in Grade 1/2 FL<sup>1</sup>



## Lenalidomide combinations

- Subsequent studies focused on lenalidomide and rituximab combinations in relapsed and resistant disease<sup>2,3</sup>

ORR: 65–77%<sup>2,3</sup>

1. Witzig TE, et al. *J Clin Oncol* 2009
2. Chong EA, et al. *Clin Cancer Res* 2015;
3. Tuscano JM, et al. *Br J Haematol* 2014

# R2 in first-line FL

	MDACC <sup>1</sup>	CALGB <sup>2</sup>
N (enrolled)	50	66
Lenalidomide	20 mg d <sub>1-21</sub> x 6	20 mg d <sub>1-21</sub> x 12
Rituximab	375 mg/m <sup>2</sup> QD 4 wks x 6	375 mg/m <sup>2</sup> weekly x 4 + Cycle 4, 6, 8, 10
Median age <sup>a</sup>	58	53
FLIPI ≥2 <sup>a</sup> (%)	78	69
RR (%)	98	93
CR (%)	87	72
2-year PFS (%)	89	—

Is R<sup>2</sup> as active as R-chemo?



# Alliance study

- Untreated Follicular lymphoma
  - Grade 1-3a
  - Stage 2 bulky, 3, 4
  - FLIPI 0-2 risk factors
- No CNS involvement
- ANC  $\geq$  1,000/ $\mu$ L
- Plts  $\geq$  75,000/ $\mu$ L
- CrCl  $\geq$  30 mL/min
- T bili  $\leq$  2 x ULN
- No HBV, HCV

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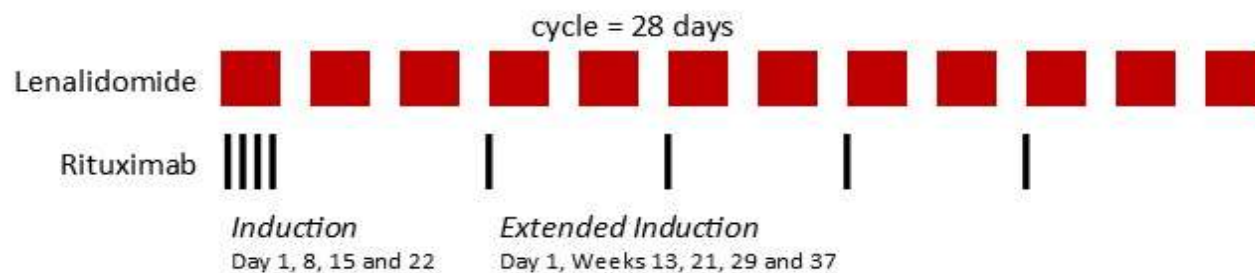
Characteristics	N = 66
Age	53 years (32-79)
Sex M vs. F	32 vs. 34
FLIPI 0-1 vs. 2 vs. 3	21 vs. 43 vs. 2
Grade 1 vs. 2. vs 3a	39 vs. 21 vs. 4
Non-bulky vs. bulky	50 vs. 15
FCGR3A 158F vs. 158F/V vs 158V	22 vs. 27 vs. 10
FCGR2A 131H vs. 131R/H vs 131R	18 vs. 28 vs. 13

**Registration**

Registered to CALGB 50803  
N = 66

Excluded (N=1)  
• Never started treatment (n=1)

**Treatment**



**Follow-up**

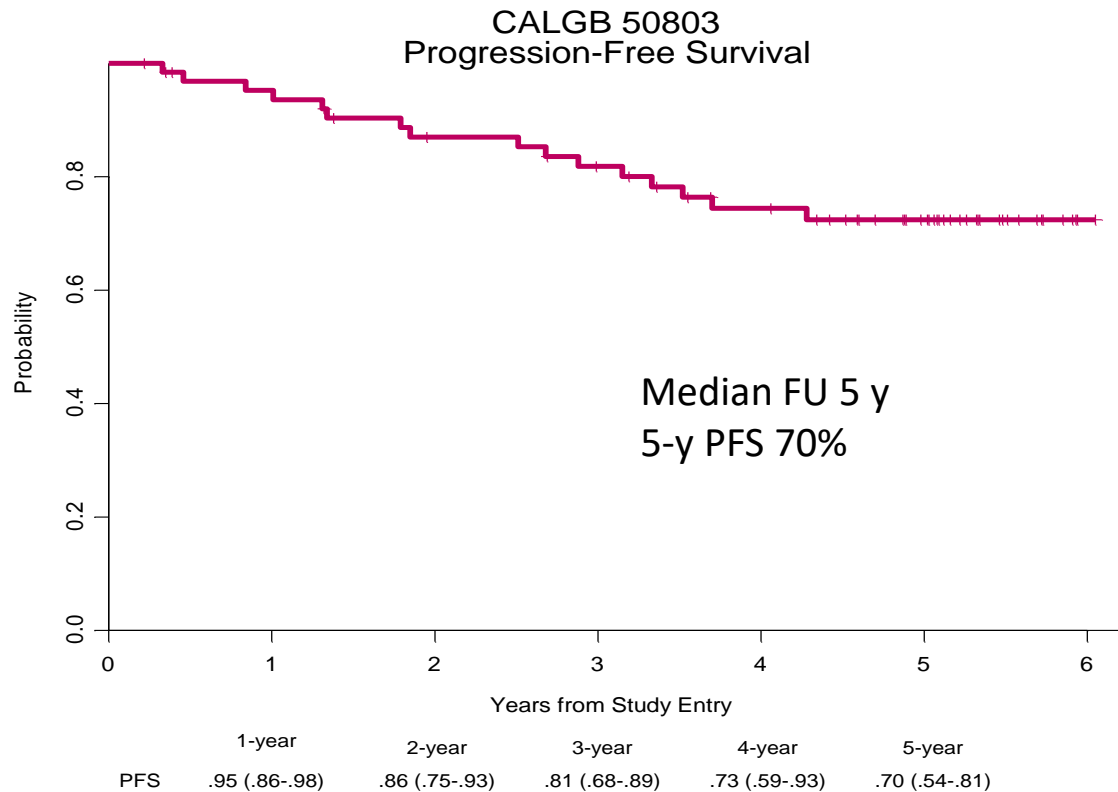
Discontinued treatment early (n=14)  
• Progression (n=2)  
• Adverse Event (n=6)  
• Refused (n=6)

**Analysis**

Included in response analysis  
N = 65

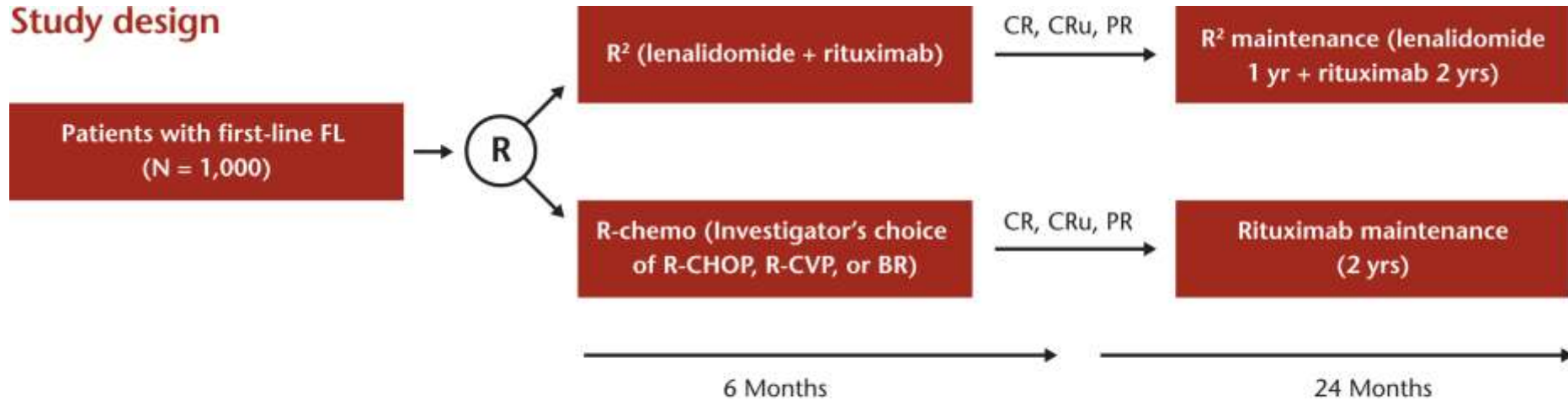
# Efficacy

Best Response	FLIPI 0-1, N=21	FLIPI 2-3, N=44*	Overall (N=65)
ORR	94%	96%	95%
CR	15 (71%)	32 (73%)	47 (72%)
PR	5 (23%)	10 (23%)	15 (23%)
Stable	0	1 (2%)	1 (2%)
Not evaluated – AE	1 (5%)	1 (2%)	1 (2%)



# THE RELEVANCE TRIAL

## Study design

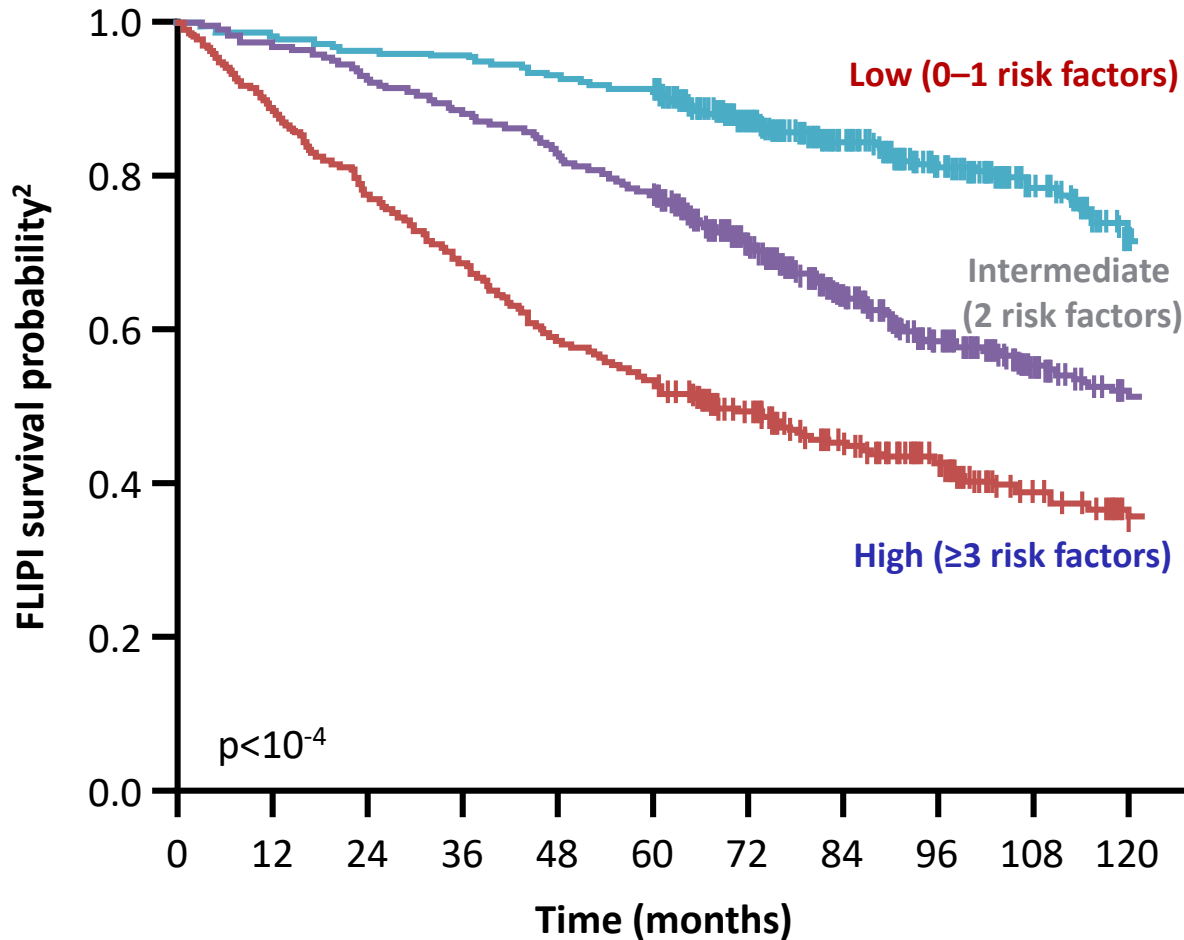


BR = bendamustine, rituximab; CR = complete response; CRu = complete response, unconfirmed; FL = follicular lymphoma; PR = partial response; R = randomize; R<sup>2</sup> = lenalidomide, rituximab; R-CHOP = rituximab, cyclophosphamide, doxorubicin, vincristine, prednisone; R-CVP = rituximab, cyclophosphamide, vincristine, prednisone



# Prognostic assessment

# FLIPI



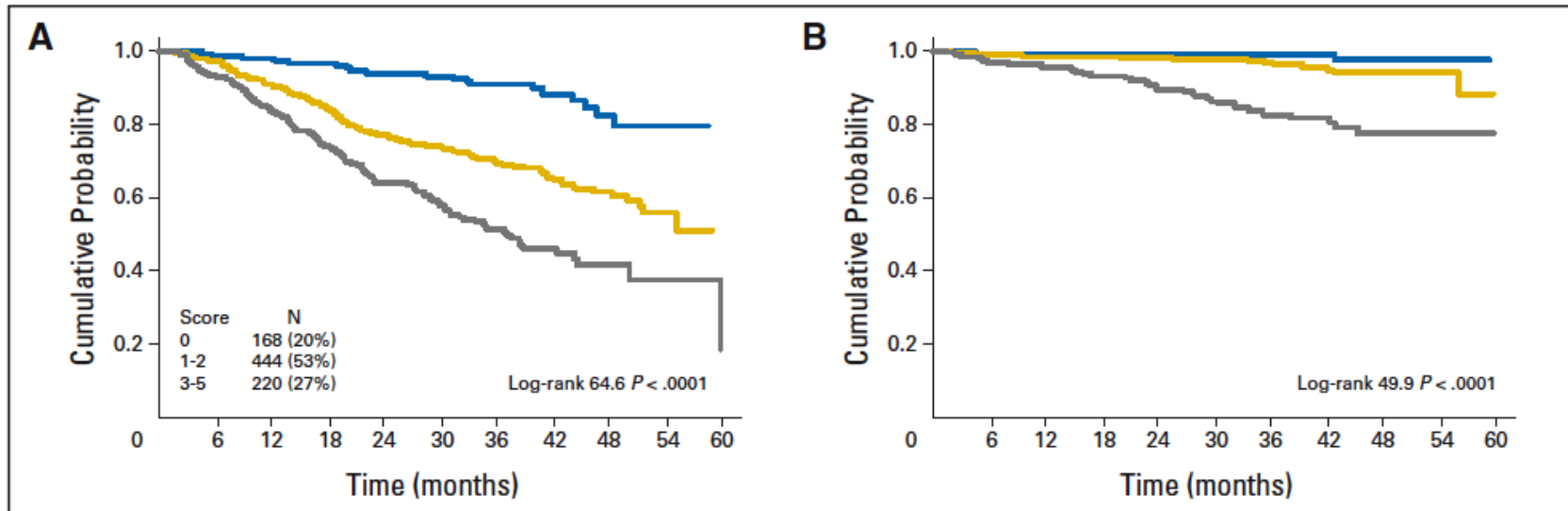
Distribution of patients	5-year OS	10-year OS
36%	90.6%	70.7%
37%	77.6%	50.9%
27%	52.5%	35.5%

Risk factors: age ( $\geq 60$  vs  $< 60$  years), Ann Arbor stage (III–IV vs I–II), number of nodal areas involved ( $> 4$  vs  $\leq 4$ ), lactate dehydrogenase serum level (above normal vs normal or below)  
 FLIPI: Follicular Lymphoma International Prognostic Index; OS: overall survival

1. Smith SM, et al. *Hem Am Soc Hematol Educ Program* 2013; 2013;561–567.  
 2. Solal-Céligny P, et al. *Blood* 2004;104:1258–1265.

# FLIPI2 – multivariate analysis of PFS

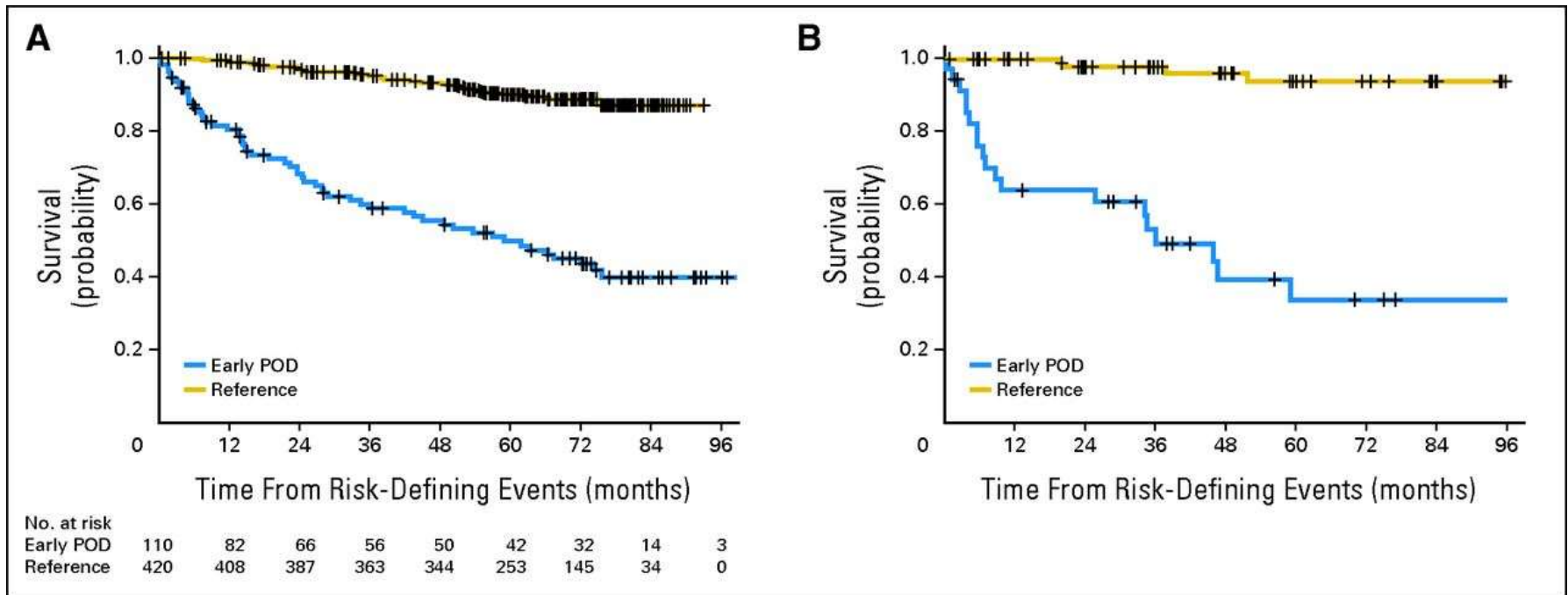
	HR	P
B2M	1.47	0.004
Hb	1.55	0.003
Age	1.43	0.005
BM	1.56	0.001
LoDLIN <sup>a</sup>	1.43	0.007



<sup>a</sup>LoDLIN: longest diameter of largest lymph node ( $\geq 6$  cm).

Federico M, et al. *J Clin Oncol.* 2009;27(27):4555-4562.

**(A) Overall survival (OS) from a risk-defining event after diagnosis in patients who received rituximab with cyclophosphamide, doxorubicin, vincristine, and prednisone (R-CHOP) chemotherapy in the National LymphoCare Study group.**





# M-7 FLIPI

Integration of gene mutations in risk prognostication for patients receiving first-line immunochemotherapy for follicular lymphoma: a retrospective analysis of a prospective clinical trial and validation in a population-based registry

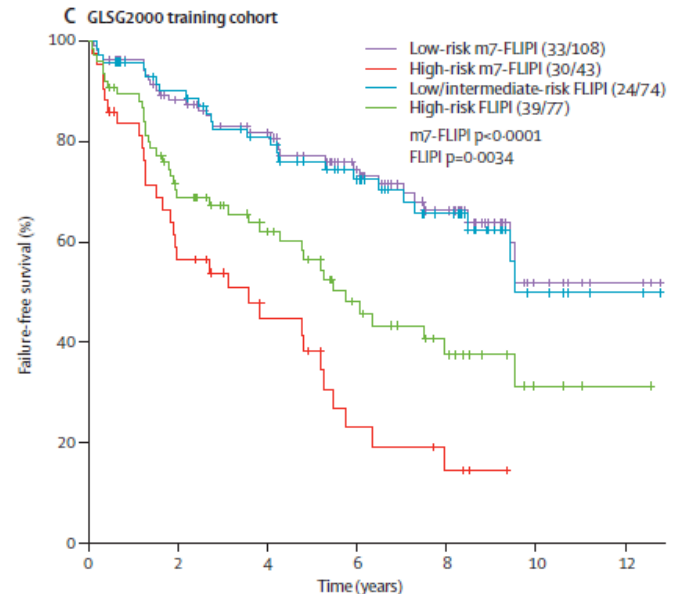
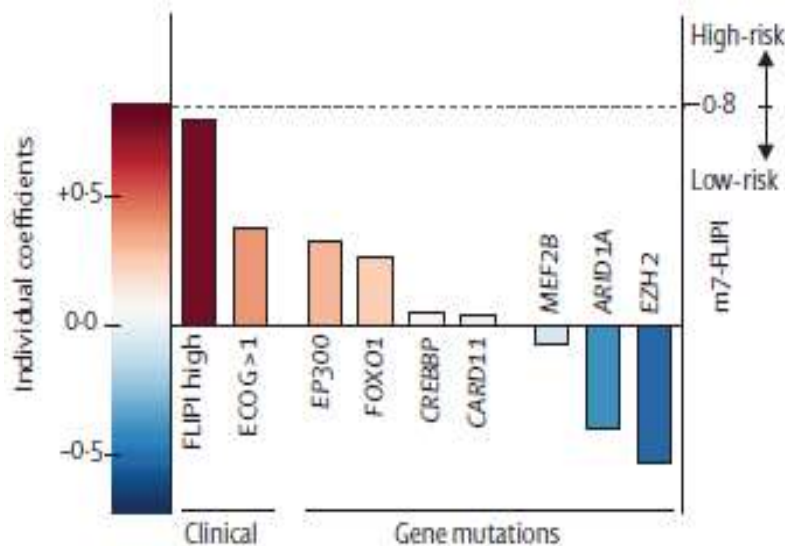


Alessandro Pastore\*, Vindijurinic\*, Robert Kridel\*, Eva Hoster\*, Annette M Staiger, Monika Szczepanowski, Christiane Pott, Nadja Kopp, Mark Murakami, Heike Horn, Ellen Leich, Aiden A Moccia, Anja Mottok, Ashwini Sunkarwalli, Paul Van Hummelen, Matthew Ducar, Daisuke Ennishi, Hennady P Shulha, Christoffer Hother, Joseph M Connors, Laurie H Sehn, Martin Dreyling, Donna Neuberg, Peter Möller, Alfred C Feller, Martin L Hansmann, Harald Stein, Andreas Rosenwald, German Ott, Wolfram Klapper, Michael Unterhalt, Wolfgang Hiddemann, Randy D Gascoyne\*, David M Weinstock\*, Oliver Weigert\*

Lancet Oncol. 2015; 16(9):1111-1122.

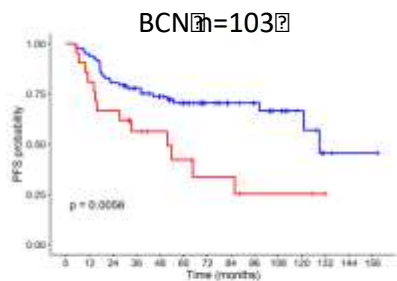
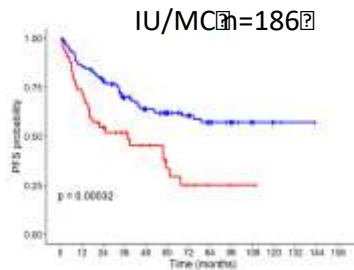
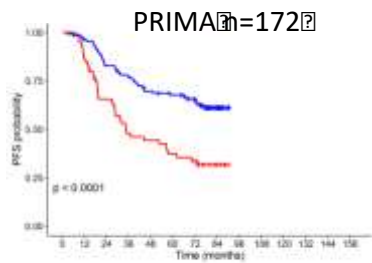
- m7-FLIPI as a clinicogenetic risk model, that included:
- the mutation status of seven genes (*EZH2*, *ARID1A*, *MEF2B*, *EP300*, *FOXO1*, *CREBBP*, *CARD11*)
  - FLIPI score
  - ECOG PS

DNA deep sequencing to retrospectively analyse the mutation status of 74 genes in 151 follicular lymphoma biopsy specimens

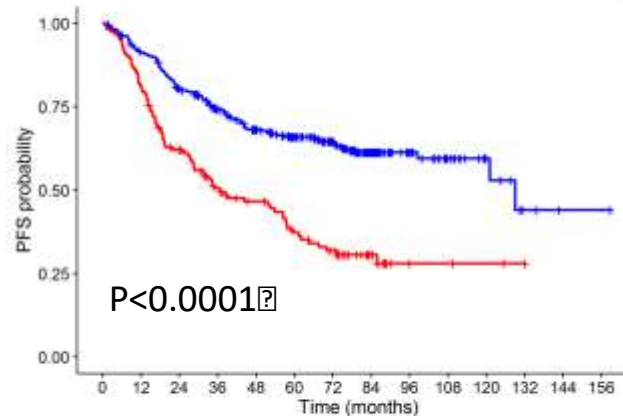


# GENE-EXPRESSION PROFILING PREDICTS DISEASE PROGRESSION IN FOLLICULAR LYMPHOMA

## VALIDATION COHORTS - RESULTS

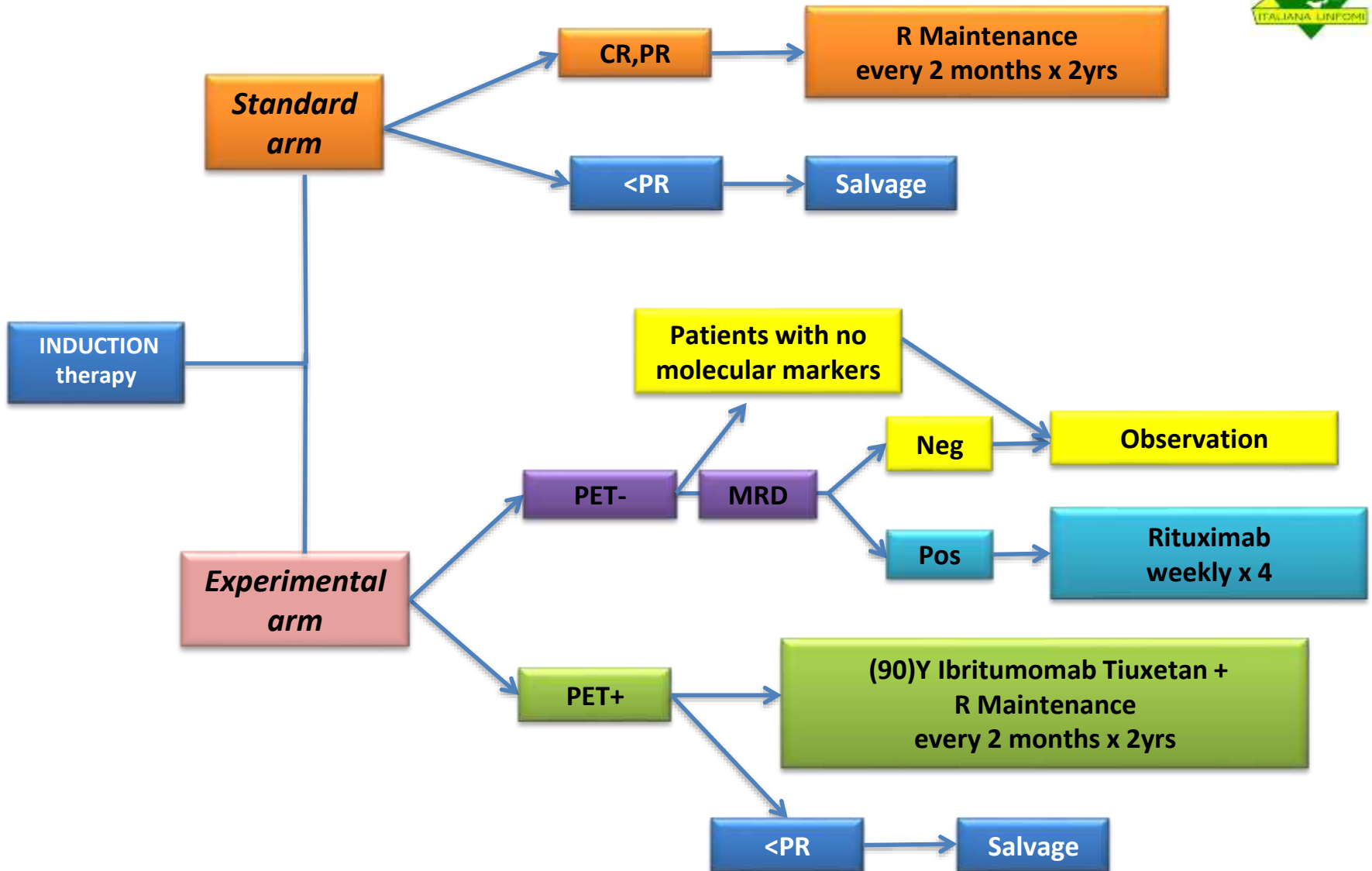


All validation cohorts  $n=461$



- HR=2.30 (1.72-3.08)
- C-index: 0.628
- Median time to progression: 3.1 vs 10.8 years
- POD24 : 38% (high score) vs 19% (low score)
- Multivariate analysis: the predictor score was independent of FLIPI and maintenance

# Maintenance

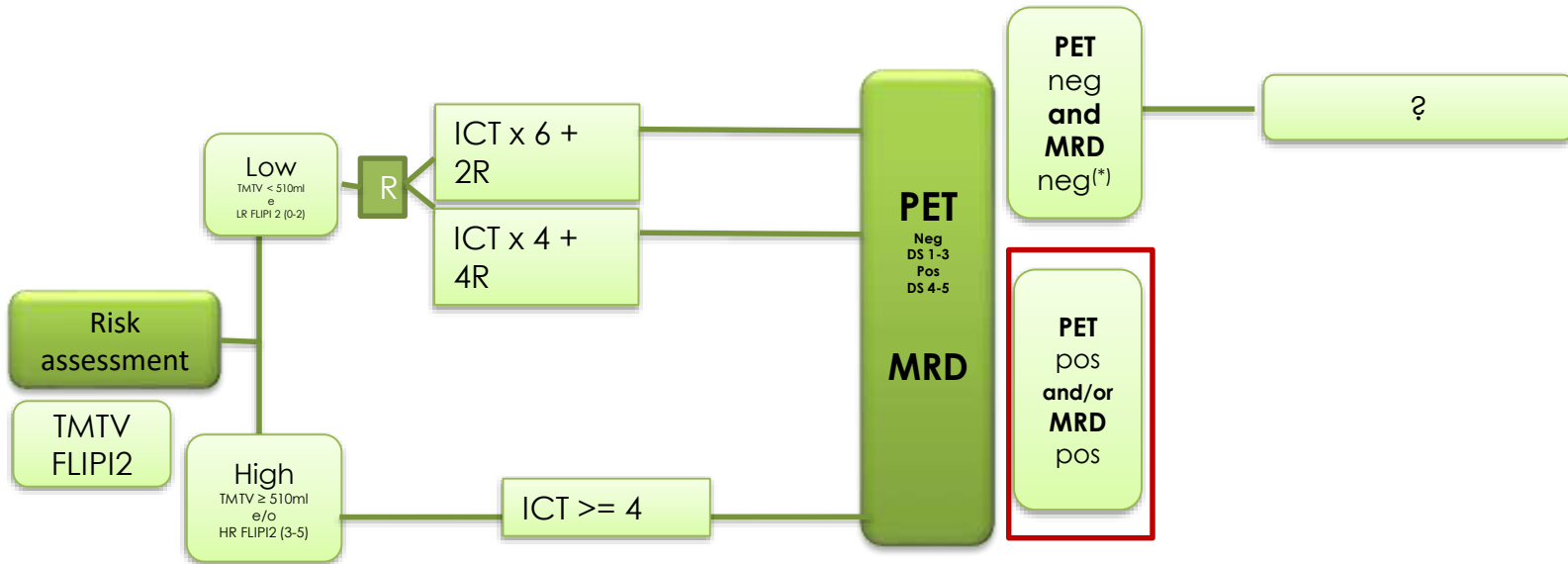


# Prognostic factors in FL

- Baseline
  - FLIPI and/or FLIPI2
  - m7FLIPI
- Post-induction
  - MRD
  - PET response
  - TMTV

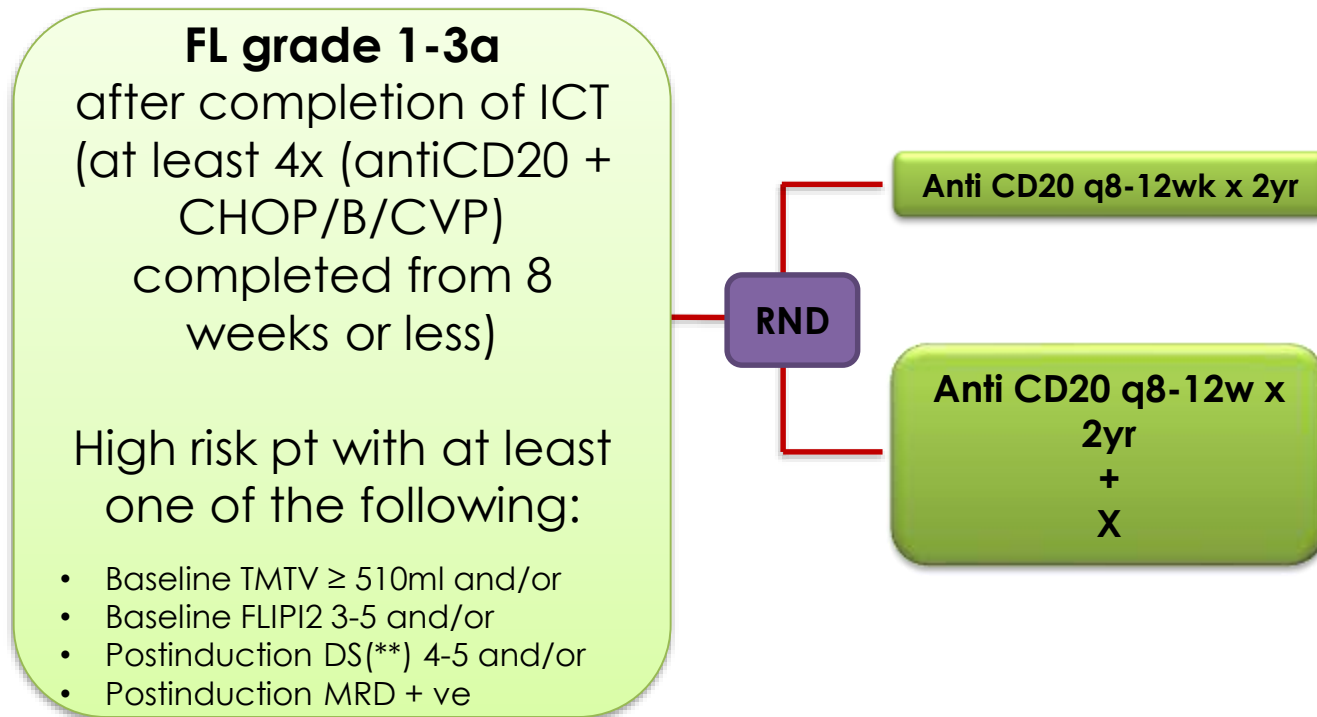
# Prognostic factors in FL

- Baseline
  - FLIPI and FLIPI2
  - ~~– m7FLIPI~~
- Post-induction
  - MRD
  - PET response
  - TMTV



ICT: AntiCD20 + CHOP or Benda or CVP  
 \* Include pz senza marcatore molecolare

# Trial Outline: X-R vs R as postinduction therapy in high risk FL responding to induction CT



Legend to figure: FL: Follicular lymphoma; TMTV Total Metabolic Tumor volume (Meignan et al JCO 2016); FLIPI2 Follicular Lymphoma Internationale Prognostic Index 2 (Federico et al JCO 2009) MRD Minimal Residual Disease; DS Deauville score (Meignan et al Leuk Lymph 2009)