



20TH

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SYMPOSIUM**

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**FL- Current Gaps of
Treatment**

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Disclosures

I have received honorarium from:

Abbvie, ADC Therapeutics, BMS, Caribou Biosciences, Daiichi Sankyo, DeNovo, Genentech, Genmab, Gilead/Kite, Janssen, Interius Bio, MEI, Merck, Novartis, and Takeda

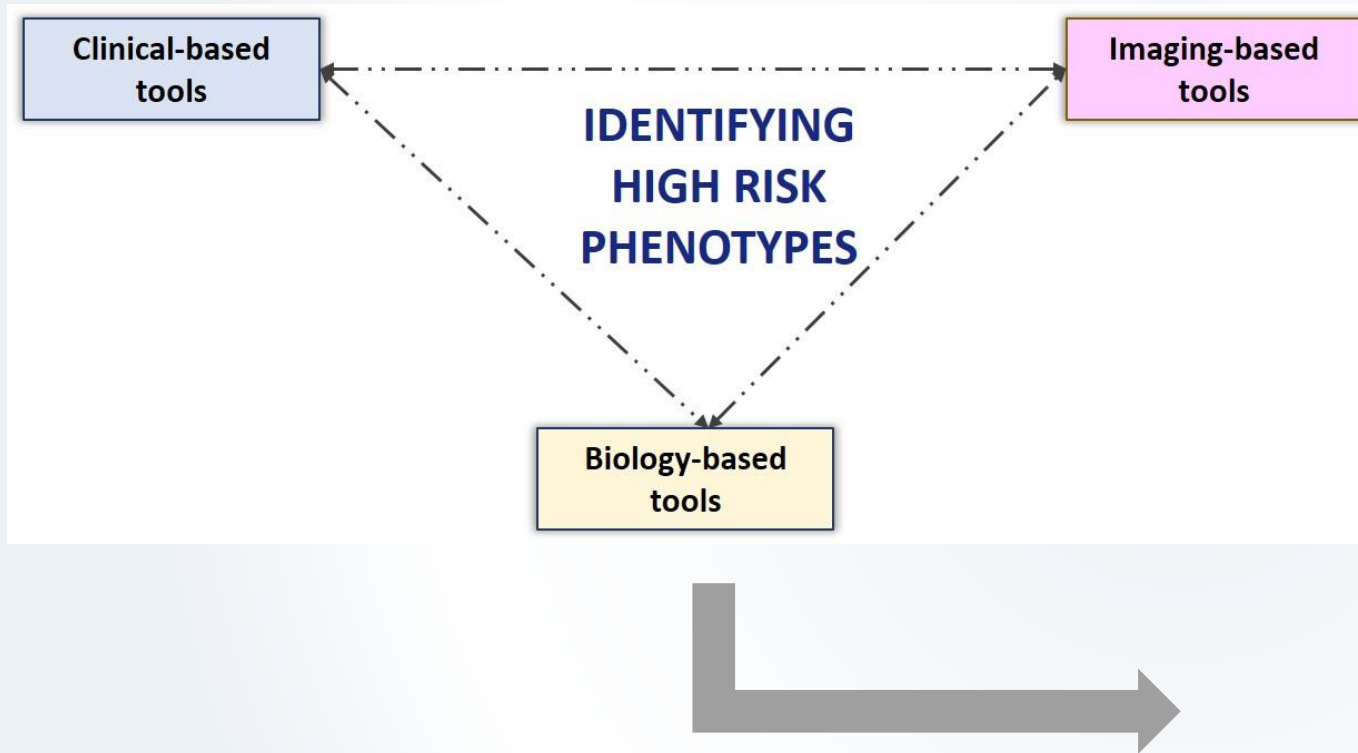
I have received research support from:

BMS, Caribou Biosciences, Daiichi Sankyo, Genentech, Genmab, Gilead/Kite, Janssen, IGM Biosciences, Novartis, and Takeda

What are the Gaps in FL?

1. Can we cure FL?
2. Risk Stratification
3. Predictive biomarkers
4. Optimal Sequencing
5. Toxicity Mitigation

GAP- Risk Stratification



At Diagnosis/Pre-treatment:

- Clinical: FLIPI, FLIPI2, PRIMA-PI, FLEX
- Biology: m7-FLIPI, PRIMA 23-gene, PD-L2
- Imaging: Baseline PET metrics

At End of Induction/After therapy:

- Imaging: EOI PET
- Biology: MRD (*not standard*)
- Response-based: POD24, Transformation

At Intervals/Dynamic:

- Biology: circulating tumor DNA (*Research tools*)

Clinical Prognostic Indices: FLIPI, FLIPI-2, PRIMA-PI and FLEX

RISK FACTORS

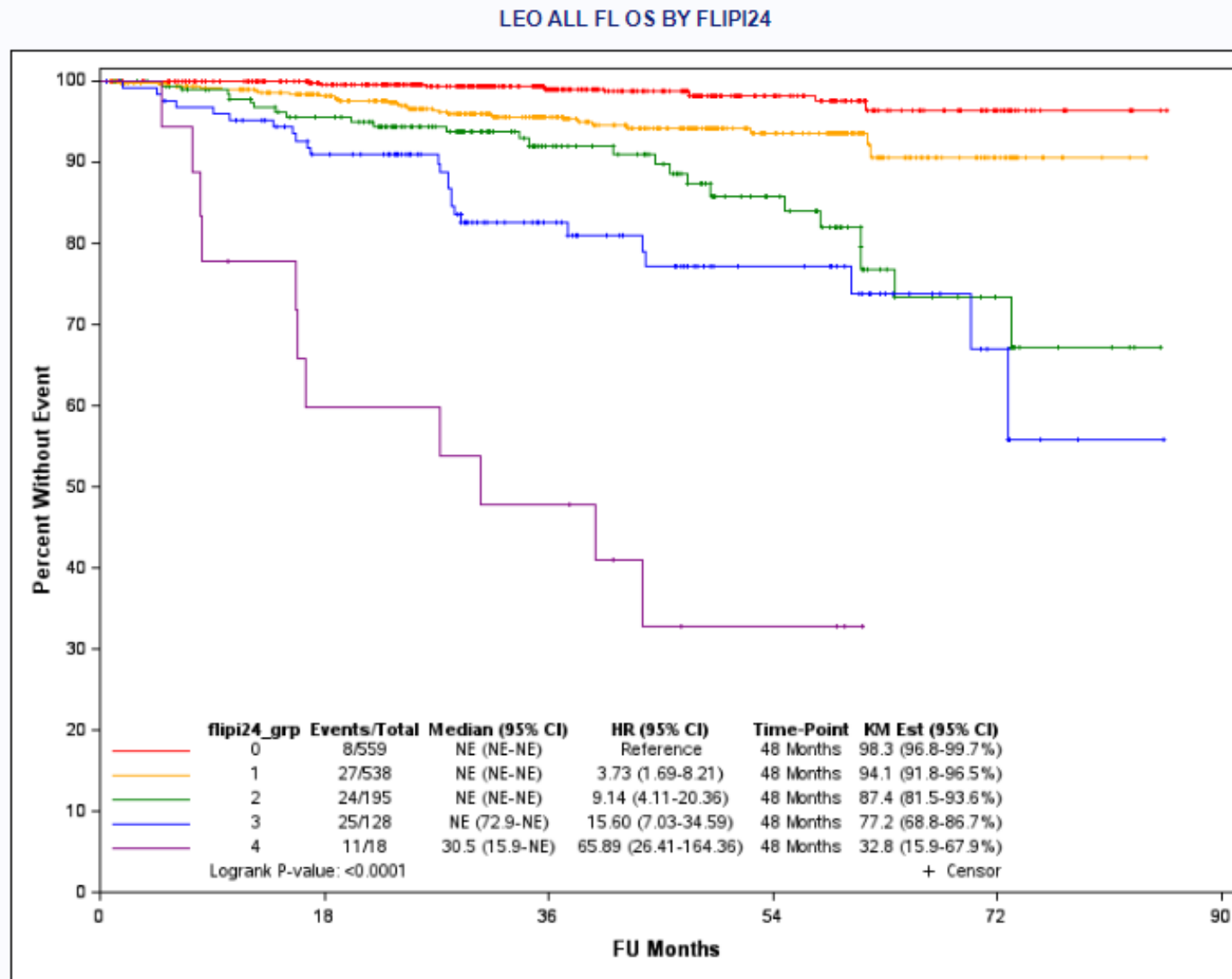
| FLIPI ¹ | FLIPI-2 ² | PRIMA-PI ³ | FLEX ⁴ |
|--|---|---|--|
| Age > 60 yrs | Age > 60yrs | BM involvement | Male sex |
| Stage III/IV | BM involvement | β_2 -M > 3mg/L | SPD in the highest quartile |
| LDH > normal | β_2 -M > normal | | LDH > normal |
| Hb <120g/L | Hb <120g/L | | Hb <120g/L |
| >4 nodal sites | Tumour mass > 6cm | | >2 extranodal sites |
| | | | Histologic grade 3A PS > 1; NK cell count < 100/ μ L; β_2 -M > normal |
| <i>Pre-rituximab era</i> | <i>R-treated</i> | <i>R-treated</i> | <i>R-treated</i> |
| Predicts OS | Predicts PFS and OS | Predicts PFS | Predicts PFS |
| Low: 5-yr OS 91% Inter: 5-yr OS 78% High: 5-yr OS 53% | Low: 5-yr PFS 80% Inter: 5-yr PFS 51% High: 5-yr PFS 19% | Low: 5-yr PFS 69% Inter: 5-yr PFS 55% High: 5-yr PFS 37% | Low: 3-yr PFS 86% High: 3-yr PFS 68% |

Each index segregates patients into 2 or 3 risk groups:

Low, **Intermediate** and **High** risk

1. Solal-Céligny P et al. *Blood*. 2004;104:1258-1265. 2. Federico M et al. *J Clin Oncol*. 2009;27:4555-4562. 3. Bachy E et al. *Blood*. 2018;132:49-58. 4. Mir et al. *Am J Hematol*. 2020;95:1503-1510.

FLIPI-24, Another Prognostic Tool



Variable

Age

LDH/ULN

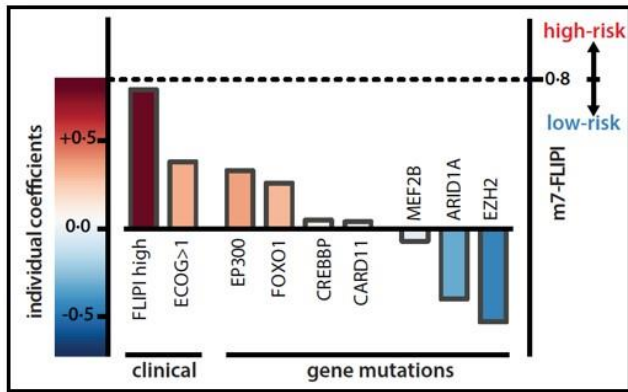
B2M

HGB

WBC

Biological Prognostic Tools: Incorporating Tumor Genotype and TME

m7-FLIPI

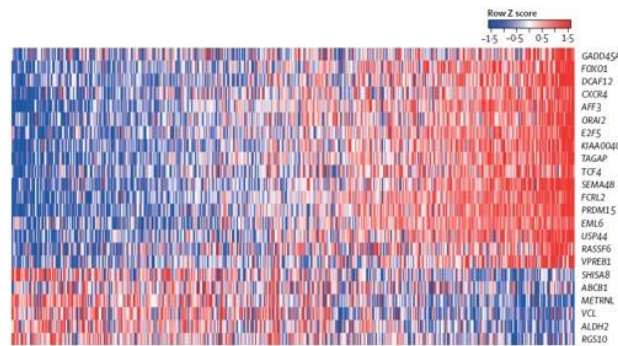


Model: 7 genes + PS + FLIPI score

| | 5 yr PFS | 5 yr OS |
|--------------------|----------|---------|
| High risk m7-FLIPI | 38.29% | 65.25% |
| Low risk m7-FLIPI | 77.21% | 89.98% |

Pastore A et al. *Lancet Oncol.* 2015;16:1111-1122.

PRIMA 23-Gene expression



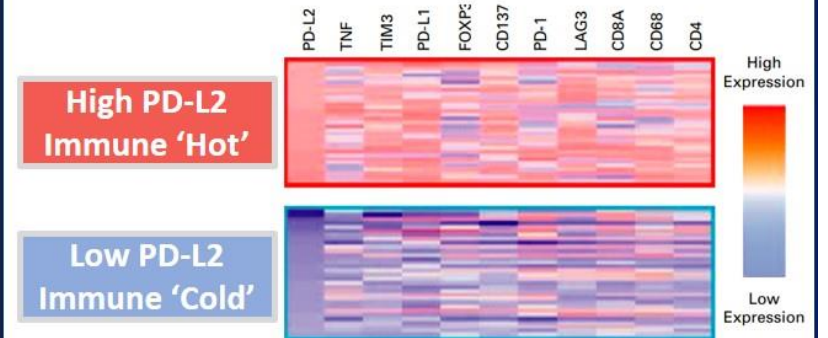
N = 134 patients (Training cohort)
Confirmed in 3 validation cohorts

| | 5 yr PFS |
|-----------|----------|
| High risk | 26% |
| Low risk | 73% |

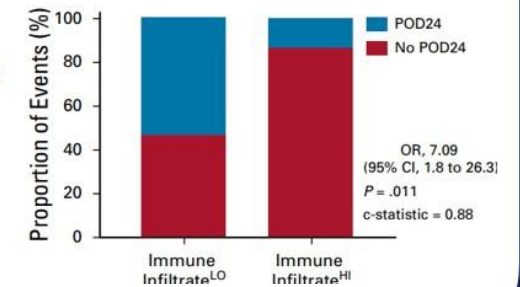
No effect
on OS

Huet S et al. *Lancet Oncol.* 2018;19:549-561.

PD-L2 expression



Immune 'cold'
= higher
POD24



Tobin JWD et al. *J Clin Oncol.* 2019;37:3300-3309.



Can We Routinely Use These Tools to Guide Clinical Decision-Making?

ABILITY TO IDENTIFY POD24 PATIENTS?

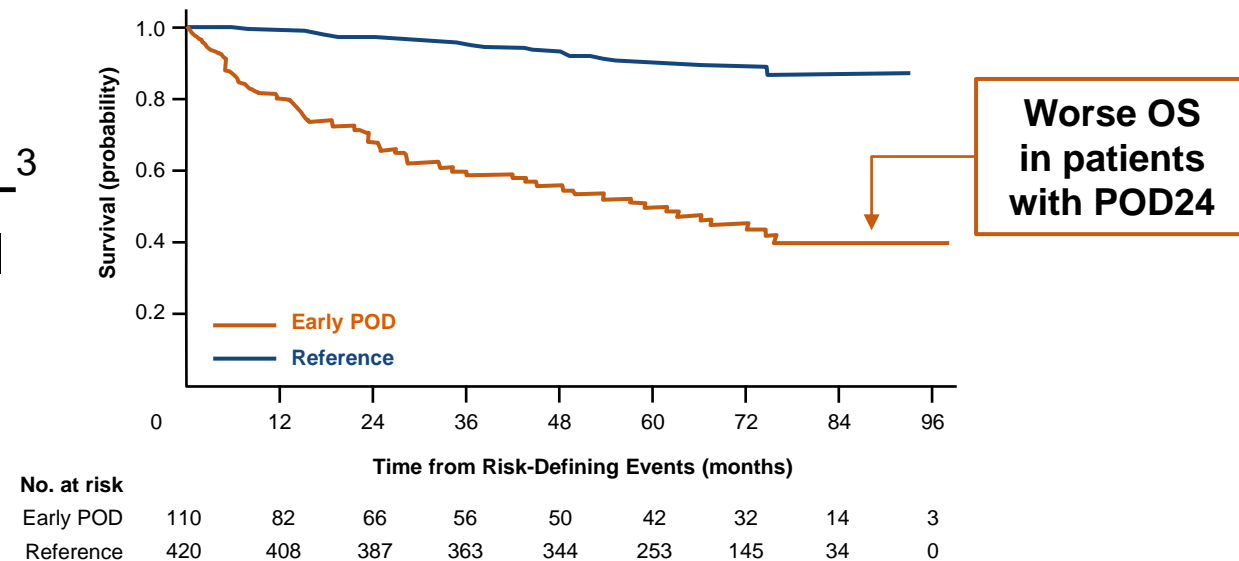
- What is the prognostic **ACCURACY**? →

| POD24 | High-risk FLIPI | High-risk FLIPI2 | PRIMA-PI | FLEX | M7-FLIPI | 23-gene ^{high} risk | PD-L2 |
|-------------|-----------------|------------------|----------|------|----------|------------------------------|--------|
| Sensitivity | 53-78% | 53% | 69% | 60% | 43-61% | 43% | 66-74% |
| Specificity | 56-62% | 59-76% | 48% | 68% | 77-86% | 79% | 60-62% |

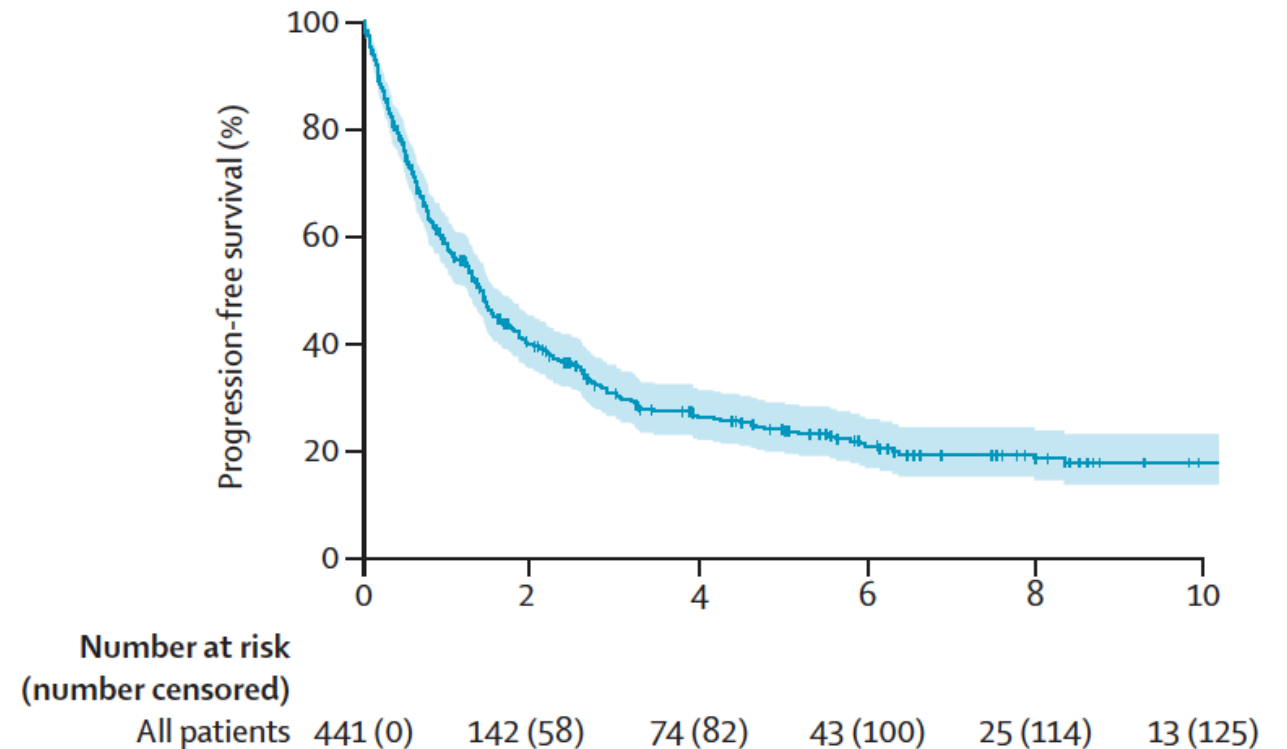
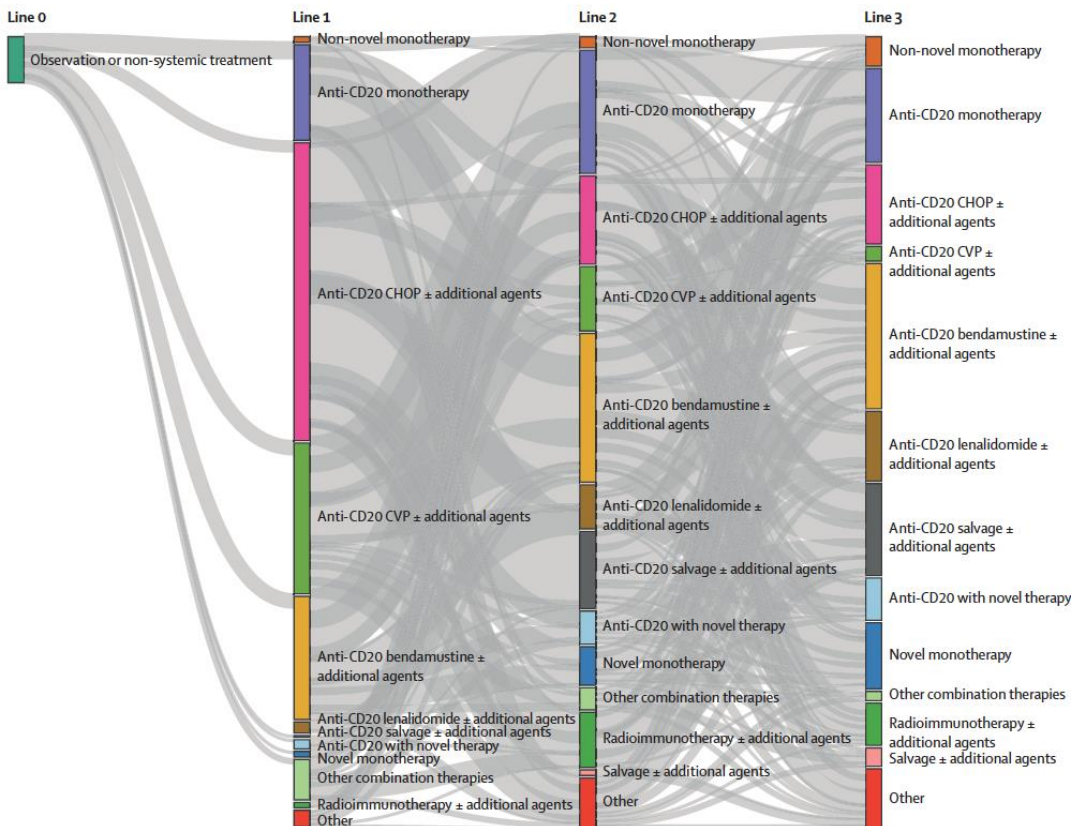
- Is it widely **ACCESSIBLE/AFFORDABLE**? → Clinical prognostic tools – straightforward
Genotyping/gene expression - research tools
- Is it **ACTIONABLE**? Guide therapy selection? → **None guide treatment**

Early Relapse (POD24)

- Biopsy recommended to detect histologic transformation of FL, which is reported to occur at a rate of 2% per year^{1,2}
 - Particularly for BR treated patients, transformation rates are higher
- Early progression of disease (≤ 2 years) after frontline chemoimmunotherapy (POD24) occurs in approximately 10-20% of patients
 - Associated with a poor prognosis and represents an unmet medical need in FL³
 - Represents a population requiring novel intervention with non-chemoimmuntherapeutic agents

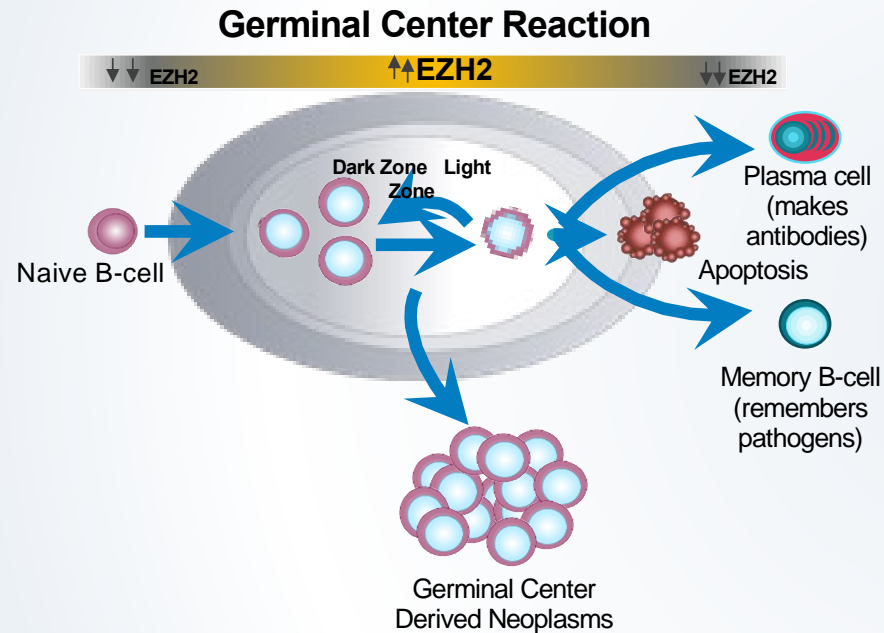


Outcomes in FL: Third Line and Beyond

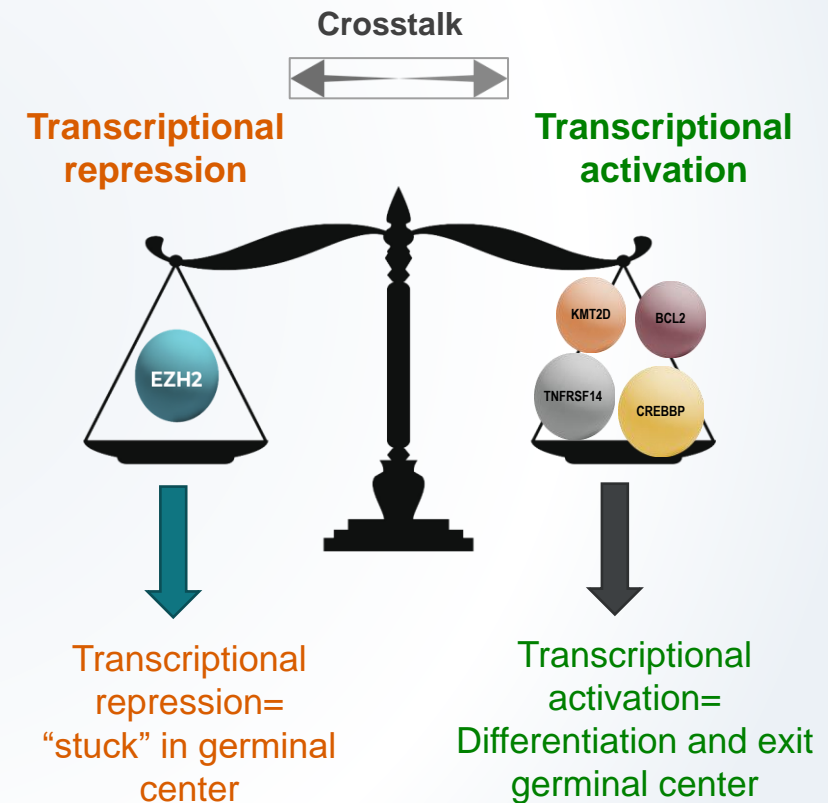


GAP- Predictive Biomarkers

Tazemetostat: Follicular Lymphoma and *EZH2*

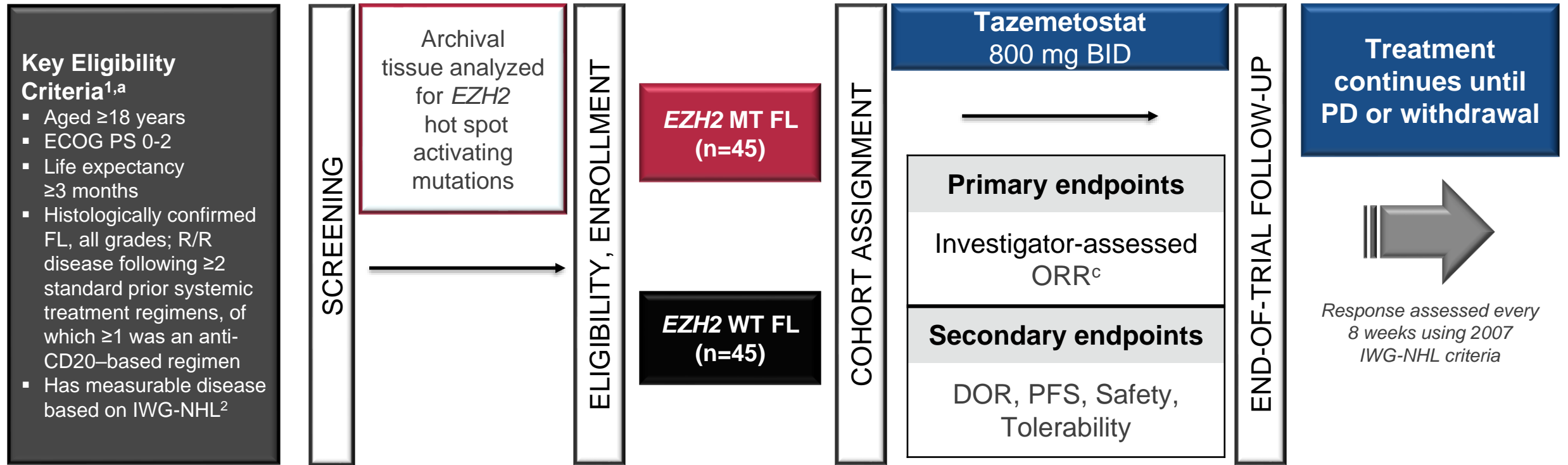


- EZH2 an epigenetic regulator of gene expression and cell fate decisions¹
- EZH2 is required for normal B-cell biology and germinal center
 - Oncogenic mutations in EZH2 suppress exit from germinal state and “lock” B cells in this state thereby transforming into a cancer²
- formation²



Tazemetostat for R/R FL

Phase 2, Open-Label, Multicenter Study



^aFor a full list of study eligibility criteria, please see [Clinicaltrials.gov/ct2/show/NCT01897571](https://clinicaltrials.gov/ct2/show/NCT01897571). ^bActual enrollment: n=54. ^cORR defined as the number of participants with a best objective response of CR or PR.

Tazemetostat for R/R FL

Phase 2, Open-Label, Multicenter Study

Response in the MT *EZH2* Cohort

| Response in MT <i>EZH2</i> (n=45) | IRC | INV |
|---|---------------------|---------------------|
| ORR, n (%) [95% CI ^a] | 31 (69) [53, 82] | 35 (78) [63, 89] |
| CR, n (%) | 6 (13) | 4 (9) |
| PR, n (%) | 25 (56) | 31 (69) |
| SD, n (%) | 13 (29) | 10 (22) |
| PD, n (%) | 1 (2) | 0 |

- 44 of 45^b (98%) patients with evidence of tumor reduction, by IRC
- mPFS, 13.8 mos (95% CI, 10.7-22.0)

^aBy Brookmeyer and Crowley method. ^b4 subjects with missing post-baseline values and 1 subject with poor image. ^cBest overall response based on Cheson (2007) criteria for lymphomas.

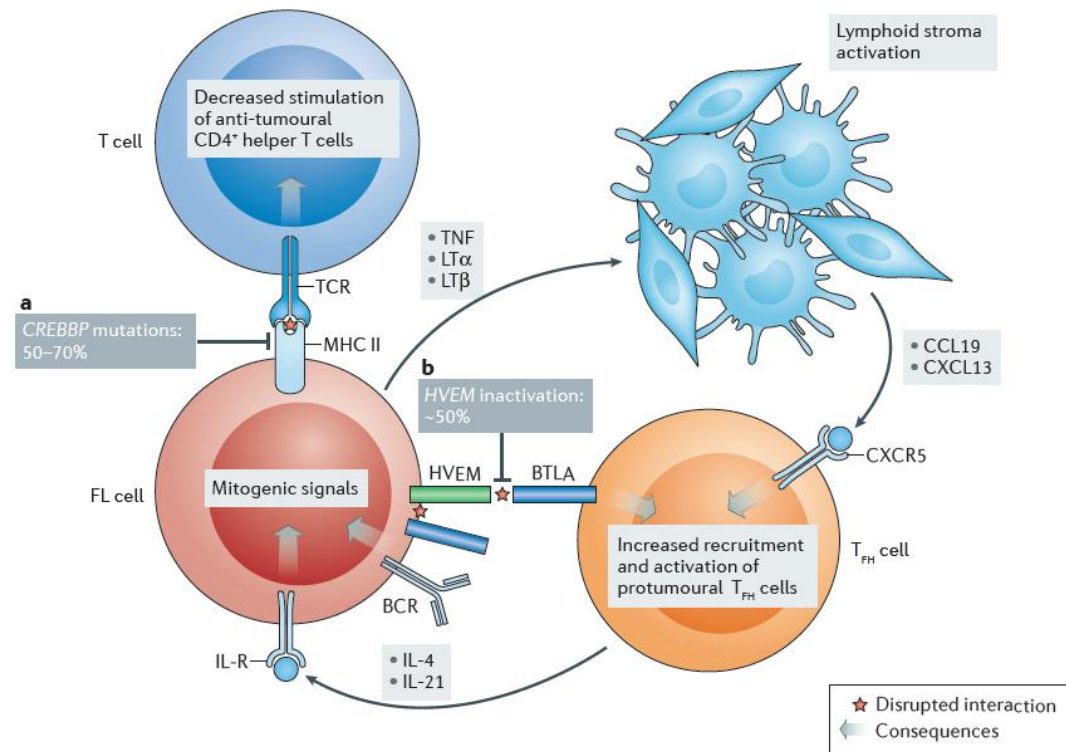
Response in the WT *EZH2* Cohort

| Response in WT <i>EZH2</i> (n=54) | IRC | INV |
|--|---------------------|---------------------|
| ORR, n (%) [95% CI ^a] | 19 (35) [23, 49] | 18 (33) [21, 48] |
| CR, n (%) | 2 (4) | 3 (6) |
| PR, n (%) | 17 (31) | 15 (28) |
| SD, n (%) | 18 (33) | 16 (30) |
| PD, n (%) | 12 (22) | 16 (30) |
| NE/missing/unknown,^b n (%) | 5 (9) | 4 (7) |

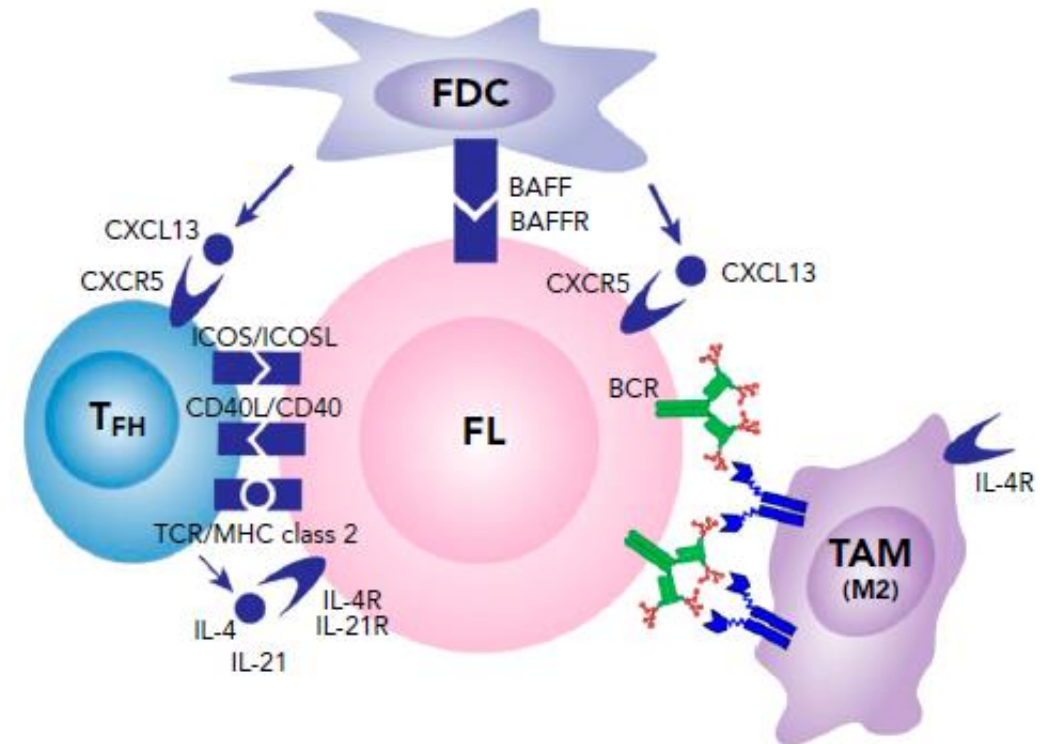
- 37 of 49^c (69%) patients with evidence of tumor reduction, by IRC
- mPFS, 11.1 mos (95%CI, 3.7-`14.6)

Influences of the Microenvironment on FL cells

Recurrent genetic alterations allow immune escape, shifting immune and stromal cells towards a supportive phenotype.



Interactive loop between FL cells and macrophages in FL tissue provides a persistent low-level signal essential for survival.

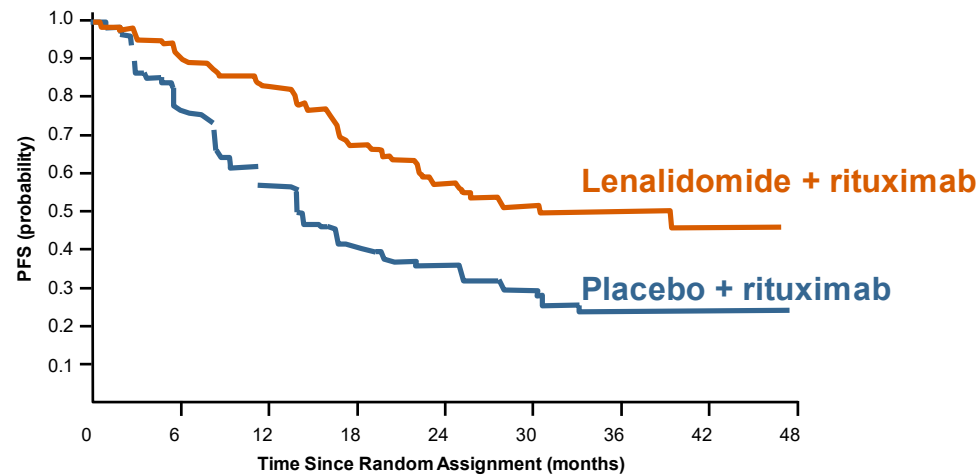


R² vs R in R/R FL and MZL

Phase III AUGMENT Study: PFS, OS

Progression-free Survival

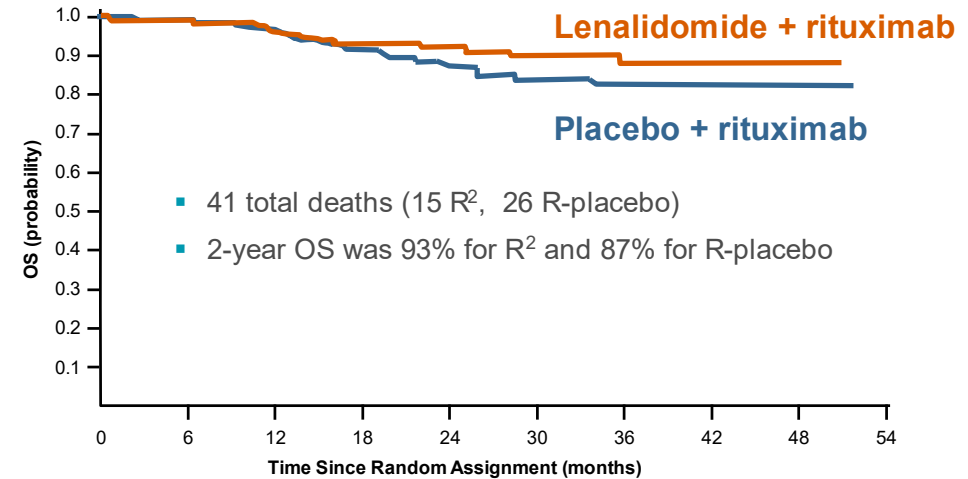
HR, 0.46 (95% CI, 0.34 to 0.62)
P < 0.0001



| No. at risk | 0 | 6 | 12 | 18 | 24 | 30 | 36 | 42 | 48 |
|--------------------------|-----|-----|-----|----|----|----|----|----|----|
| Lenalidomide + rituximab | 178 | 148 | 124 | 91 | 59 | 39 | 20 | 7 | 0 |
| Placebo + rituximab | 180 | 132 | 92 | 58 | 40 | 26 | 10 | 4 | 0 |

Overall Survival

HR, 0.61 (95% CI, 0.33 to 1.13)



| No. at risk | 0 | 6 | 12 | 18 | 24 | 30 | 36 | 42 | 48 | 54 |
|--------------------------|-----|-----|-----|-----|-----|----|----|----|----|----|
| Lenalidomide + rituximab | 178 | 167 | 155 | 143 | 122 | 80 | 44 | 15 | 1 | 0 |
| Placebo + rituximab | 180 | 176 | 167 | 145 | 116 | 79 | 40 | 14 | 3 | 0 |

| Median PFS | R ² (n=178) | R-Placebo (n=180) | HR | P Value |
|---------------------|---------------------------|----------------------|------------------|---------|
| By IRC, mo (95% CI) | 39.4 (22.9-NE) | 14.1 (11.4-16.7) | 0.46 (0.34-0.62) | <0.0001 |
| By INV, mo (95% CI) | 25.3 (21.2-NE) | 14.3 (12.4-17.7) | 0.51 (0.38-0.69) | <0.0001 |

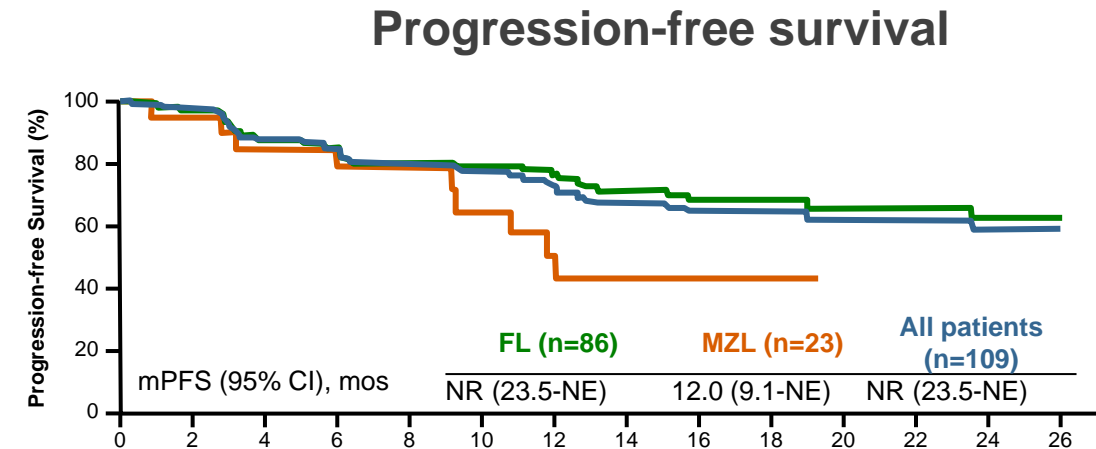
Median follow-up: 28.3 months

Leonard J et al. *J Clin Oncol*. 2019;37:1188-1199.

ZUMA-5 Study of Axi-cel in Relapsed/Refractory FL and MZL

| Characteristic | FL n=124 | MZL N=24 | All Patients N=148 |
|-----------------------------|-------------|-------------|-----------------------|
| Median age (range) | 60 (53-67) | 65 (61-72) | 61 (53-68) |
| FLIPI 3-5 | 54 (44%) | N/A | N/A |
| High tumor burden (GELF) | 64 (52%) | 10 (42%) | 74 (50%) |
| Median prior tx (IQR) | 3 (2-4) | 3 (2-5) | 3 (2-5) |
| Refractory to last tx | 84 (68%) | 18 (75%) | 102 (69%) |
| POD24 | 68 (55%) | 13 (57%) | 81 (55%) |

| | All patients (n=109) | FL (n=86) | MZL (n=23) |
|-----|-------------------------|--------------|---------------|
| ORR | 92% | 94% | 83% |
| CRR | 76% | 79% | 65% |



| AEs of Special Interest (n=148) | |
|---------------------------------|-----|
| Cytokine Release Syndrome | |
| Any grade | 82% |
| Grade \geq 3 | 7% |
| Neurologic Events | |
| Any grade | 59% |
| Grade \geq 3 | 19% |

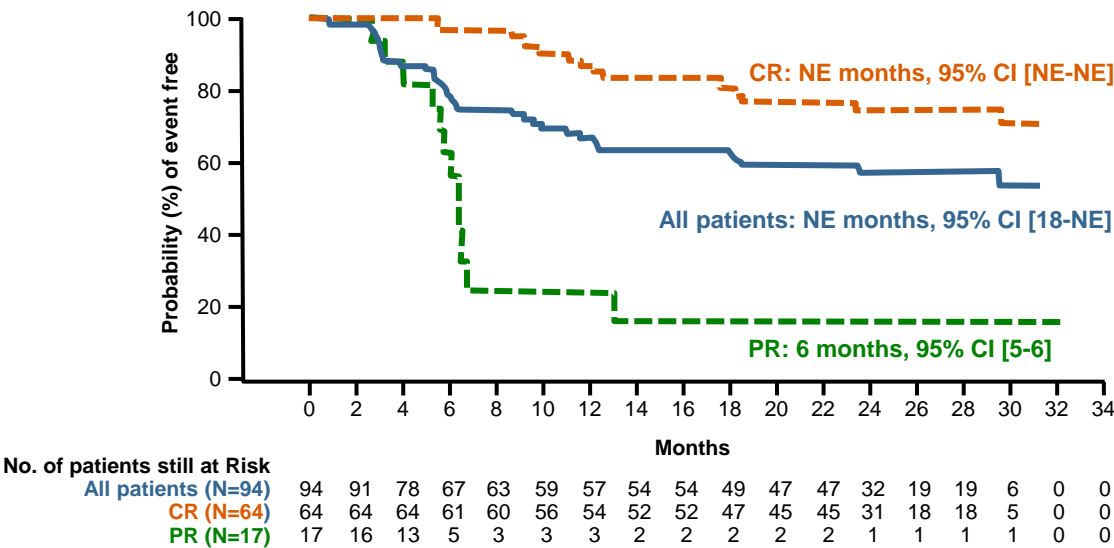
ELARA Study of Tisa-cel in Relapsed/Refractory FL

| Characteristic | n=97 |
|-------------------------|------------|
| Median age (range) | 57 (49-64) |
| Median prior tx (range) | 4 (2-13) |
| Refractory | 78% |
| POD24 | 63% |

| | All patients n=94 |
|-----|----------------------|
| ORR | 86% |
| CRR | 69% |

| AEs of Special Interest (n=97) | |
|--------------------------------|-----|
| Cytokine Release Syndrome | |
| Any grade | 49% |
| Grade ≥ 3 | 0% |
| ICANS | |
| Any grade | 4% |
| Grade ≥ 3 | 1% |

Progression-free Survival



Median PFS not reached

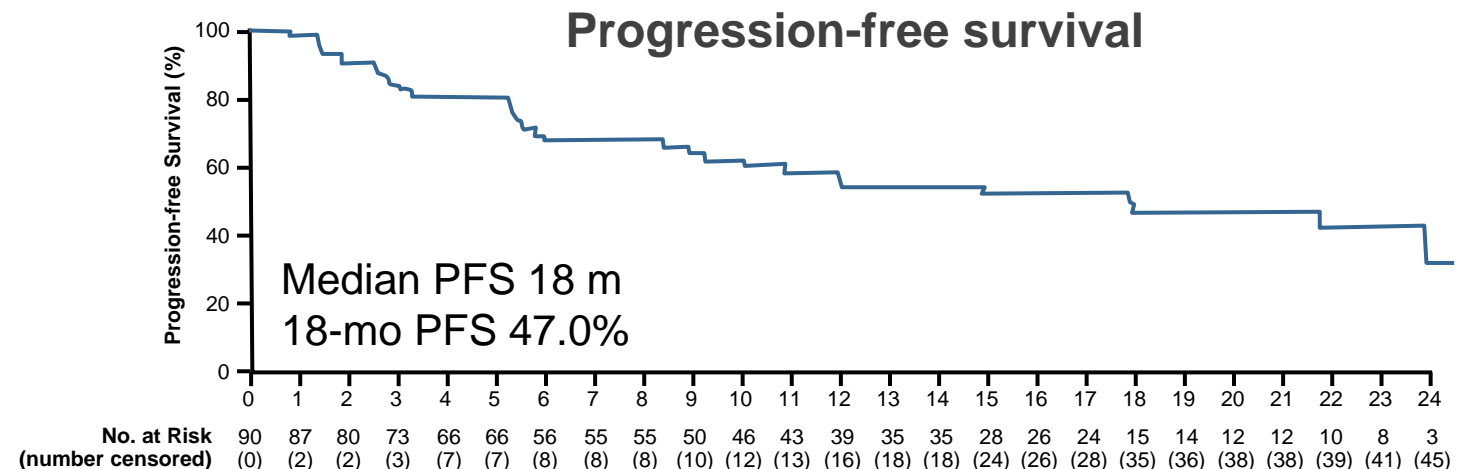
| Event-free Probability | % (95% CI) |
|------------------------------|------------|
| 24-month PFS, all patients | 57 (46-67) |
| 24-month PFS, patients in CR | 75 (62-84) |

The New Kid on the Block: Mosunetuzumab, an Anti-CD20/CD3 Bispecific Antibody, in Relapsed/Refractory FL

- Mosunetuzumab IV 1 mg on cycle 1 day 1, 2 mg on day 8, 60 mg on day 15 and cycle 2 day 1, and 30 mg on day 1 of cycle 3 onwards
- Total of 8 cycles for patients in CR, 17cycles for patients in PR

| Characteristic | n=90 |
|-----------------------|------------|
| Median age (range) | 60 (53-67) |
| FLIPI 3-5 | 46% |
| Median prior tx (IQR) | 3 (2-4) |
| Refractory to last tx | 69% |
| POD24 | 52% |

| Response | n=90 |
|------------|------|
| ORR | 80% |
| CRR | 60% |
| Median DOR | 23 m |
| 18-mo DOR | 57% |

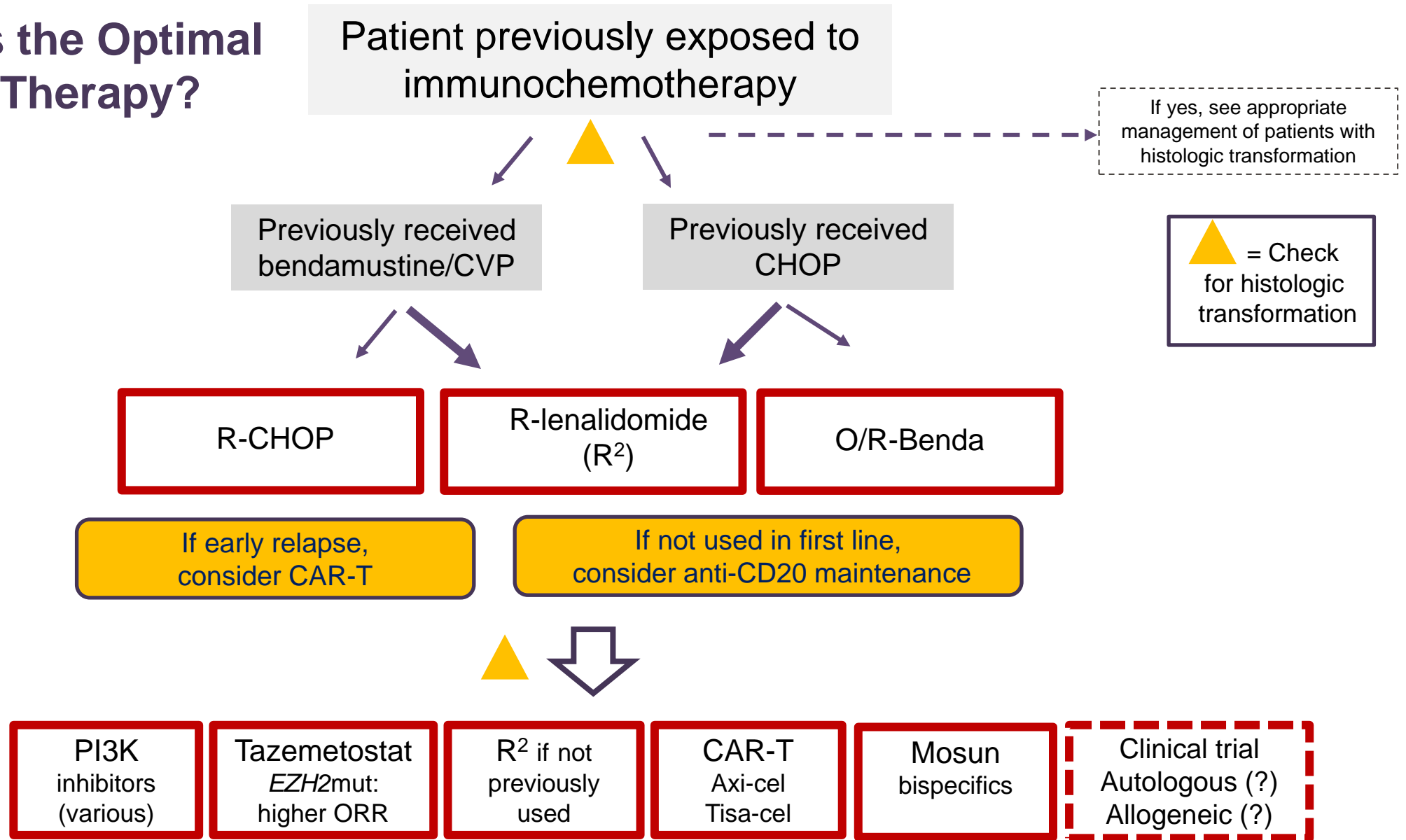


| AEs of Special Interest (n=90) | |
|--------------------------------|-----|
| Cytokine Release Syndrome | |
| Any grade | 44% |
| Grade ≥ 3 | 2% |
| Neurologic Events | |
| Any grade | 5% |
| Grade ≥ 3 | 0% |

GAP- What is the Optimal Sequence of Therapy?

2nd line

3rd line and +



ZUMA-5 CRS and Neurologic Events

| Parameter | CRS ^a | | | Neurologic Events ^a | | |
|---|-------------------------|---------------|---------------------------|--------------------------------|---------------|-------------------------|
| | FL (n=124) | MZL (n=22) | All Patients (N=146) | FL (n=124) | MZL (n=22) | All Patients (N=146) |
| Any grade | 97 (78) | 22 (100) | 119 (82) | 70 (56) | 17 (77) | 87 (60) |
| Grade ≥3 | 8 (6) | 2 (9) | 10 (7) | 19 (15) | 9 (41) | 28 (19) |
| Most common CRS symptoms of any grade, n/n (%) | | | | | | |
| Pyrexia | 94/97 (97) | 20/22 (91) | 114/119 (96) | – | – | – |
| Hypotension | 39/97 (40) | 10/22 (45) | 49/119 (41) | – | – | – |
| Most common neurologic events of any grade, n/n (%) | | | | | | |
| Tremor | – | – | – | 36/70 (51) | 9/17 (53) | 45/87 (52) |
| Confusional state | – | – | – | 28/70 (40) | 7/17 (41) | 35/87 (40) |
| Tocilizumab use, n (%) | 56 (45) | 15 (68) | 71 (49) | 7 (6) | 2 (9) | 9 (6) |
| Corticosteroid use, n (%) | 19 (15) | 6 (27) | 25 (17) | 38 (31) | 14 (64) | 52 (36) |
| Median time to onset (range), days | 4 (1–15) | 4 (1–9) | 4 (1–15) | 7 (1–177) | 7 (3–19) | 7 (1–177) |
| Median duration of events (range), days | 6 (1–27) | 6 (2–14) | 6 (1–27) | 14 (1–452) | 10 (2–81) | 14 (1–452) |
| Patients with resolved events, n/n (%) | 96/97 (99) ^b | 22/22 (100) | 118/119 (99) ^b | 67/70 (96) | 14/17 (82) | 81/87 (93) |

Predictors of Response and Toxicity with CD19 Auto CARs

Improved Response

PATIENT

- Low tumor burden, low lactate dehydrogenase
- Low pretreatment inflammatory markers
- Absence of medical comorbidities
- Lack of need for bridging therapy

T CELLS

- Proportion of CCR7+ and other early memory T cells in the CAR product
- Faster doubling time in vitro
- Higher CAR T-cell peak to tumor burden ratio

TUMOR

- Low tumor myeloid-derived suppressor cells
- High tumor-infiltrating lymphocytes
- Absence of *MYC* overexpression
- Absence of *CD58* mutations

Increased Toxicity

PRE-TREATMENT

- High tumor burden, elevated pretreatment lactate dehydrogenase
- High pretreatment inflammatory markers
- ? High pretreatment monocyte levels

POST-TREATMENT

- High peak CAR T-cell levels
- High peak cytokine levels
- Markers of disseminated intravascular coagulation (including fibrinogen levels)
- Early cytokine release syndrome

What do we know about response/risk of toxicity with bispecifics?

Conclusions

- Outcomes for the majority of patients with FL are favorable.
 - Can we do a better job with risk stratification?
- Balancing the goals of therapy with patient specific characteristics generally informs treatment selection,
 - Can we be more scientific about this?
- An unmet need is identifying optimal sequencing of therapy.
 - We need more randomized trials and biomarker exploration.
- The goal of treatment is to achieve a normal life expectancy without negatively impacting quality of life.
 - Is functional cure as important as curative intent with a given line of therapy?

Acknowledgements

Patients and their loved ones

Indolent Lymphoma Team at MDACC

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