

***Therapeutic News
from the World of
Waldenstrom's
Macroglobulinemia***

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DISCLOSURES

Company

Abbvie/Pharmacyclics

Research Support

Beigene

Research Support, Consulting

Schrodinger

Consulting

Eli Lilly

Research Support

Johnson and Johnson

Research Support



**Sixth International Workshop on Waldenstrom's Macroglobulinemia
Giampaolo Merlini, Enrica Morra, Steve Treon (Co-Organizers)
Venice 2010**



SOMEDAY,
ALL THIS
WILL BE YOURS.

Pre-MYD88

WM
Treatment
Approach

*“Hand me
down
medicines”*

guy &
rodd

Mutated MYD88 Pro-survival Signaling in WM

The NEW ENGLAND JOURNAL of MEDICINE

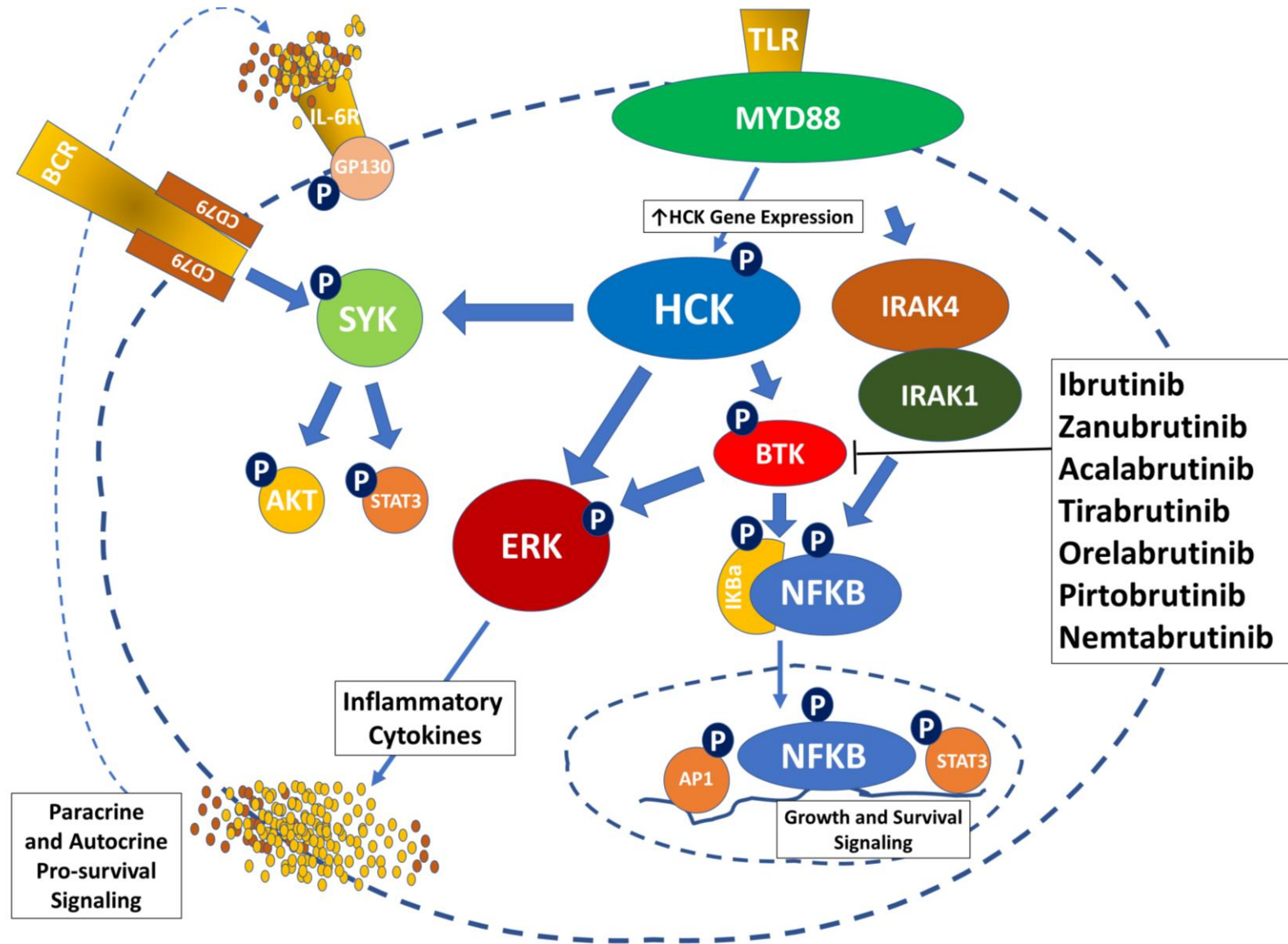
ORIGINAL ARTICLE

MYD88 L265P Somatic Mutation in Waldenström's Macroglobulinemia

Steven P. Treon, M.D., Ph.D., Lian Xu, M.S., Guang Yang, Ph.D., Yangsheng Zhou, M.D., Ph.D., Xia Liu, M.D., Yang Cao, M.D., Patricia Sheehy, N.P., Robert J. Manning, B.S., Christopher J. Patterson, M.A., Christina Tripsas, M.A., Luca Arcaini, M.D., Geraldine S. Pinkus, M.D., Scott J. Rodig, M.D., Ph.D., Aliyah R. Sohani, M.D., Nancy Lee Harris, M.D., Jason M. Laramie, Ph.D., Donald A. Skifter, Ph.D., Stephen E. Lincoln, Ph.D., and Zachary R. Hunter, M.A.

MYD88 mutations occur in 95-97% WM Patients

Treon, et al. N Engl J Med. 2012;367(9):826-833.
 Yang, et al. Blood. 2013;122(7):1222-1232.
 Hodge, et al. Blood. 2014;123(7):1055-1058.
 Yang, et al. Blood. 2016;127(25):3237-3252.
 Chen, et al. Blood. 2018;131(18):2047-2059.
 Liu, et al. Blood Adv. 2020;4(1):141-153.
 Munshi, et al. Blood Cancer J. 2020;10:12.
 Munshi, et al. Blood Adv. 2022.



BTK-Inhibitor Trials in WM

| Study | Cohort | Agent (s) | N= | Time to Major Resp. | ORR/Major RR | ≥VGPR | PFS |
|---------------------------|---------|------------------------|-----|---------------------|--------------|-------|--------------|
| Pivotal Study | R/R | Ibrutinib | 63 | 2 mo. | 91% / 79% | 30% | 54% @ 60 mo. |
| INNOVATE Arm C | R/R | Ibrutinib | 31 | 2 mo. | 87% / 77% | 29% | 40% @ 60 mo. |
| Phase 2 | TN | Ibrutinib | 30 | 1.9 mo. | 100% / 87% | 30% | 76% @ 48 mo. |
| INNOVATE Arms A, B | TN, R/R | Ibrutinib Rituximab | 150 | 3 mo. | 92% / 76% | 31% | 68% @ 54 mo. |

*Median ORR: 93%; Major RR: 81%; ≥VGPR: 30%;
PFS 76% @ 4 yrs*

| | | | | | | | |
|-----------------------------|---------|---------------|-----|----------------|--|--|---|
| (MYD88^{WT}) | | | | | | | |
| Phase 2 | TN, R/R | Acalabrutinib | 106 | N/A | 94% / 81% | 39% | 84% TN / 52% R/R (@ 66 mo.) |
| Phase 2 | TN, R/R | Tirabrutinib | 27 | 1.9 TN 2.1 R/R | 96% / 93% | 33% | 93% @ 24 mo. |
| Phase 2 | R/R | Pirtobrutinib | 80 | N/A | 81% / 67% (prior cBTKi) 88% / 88% (cBTKi naïve) | 24% (prior cBTKi) 29% (cBTKi naïve) | 57% @ 18 mo. (for prior cBTKi) N/A for cBTKi naïve. |

CXCR4 mutations impact WM disease presentation and treatment response.

Plenary Paper

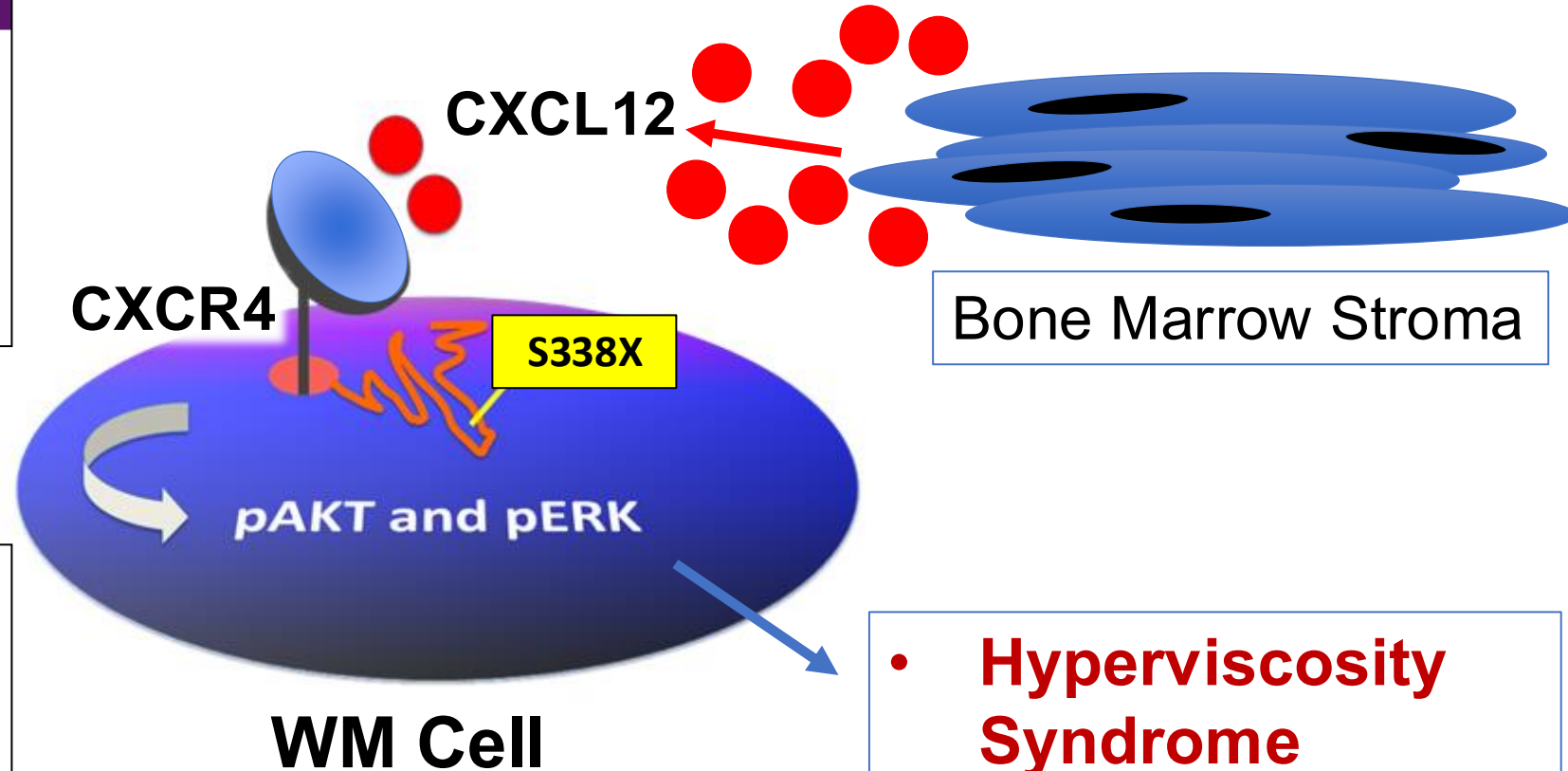
LYMPHOID NEOPLASIA

The genomic landscape of Waldenström macroglobulinemia is characterized by highly recurring MYD88 and WHIM-like CXCR4 mutations, and small somatic deletions associated with B-cell lymphomagenesis

Zachary R. Hunter,^{1,2} Lian Xu,¹ Guang Yang,¹ Yangsheng Zhou,¹ Xia Liu,¹ Yang Cao,¹ Robert J. Manning,¹ Christina Tripsas,¹ Christopher J. Patterson,¹ Patricia Sheehy,¹ and Steven P. Treon^{1,3}

¹Bing Center for Waldenström's Macroglobulinemia, Dana-Farber Cancer Institute, Boston, MA; ²Department of Pathology and Laboratory Medicine, Boston University School of Graduate Medical Sciences, Boston, MA; and ³Harvard Medical School, Boston, MA

- 30-40% of WM patients are CXCR4 mutated.
- >40 different CXCR4 NS and FS mutations
- Most common is S338X.



CXCR4 Impact on BTK-Inhibitor Outcomes in WM

| Study | Patient Population | Agent (s) | Time to Major Response (CXCR ^{Mut} vs. WT) | Major Response Rate (CXCR ^{Mut} vs. WT) | ≥VGPR (CXCR ^{Mut} vs. WT) | PFS (CXCR ^{Mut} vs. WT) |
|----------------|--------------------|-----------|---|--|------------------------------------|----------------------------------|
| Pivotal | R/R | Ibrutinib | 4.7 vs. 1.8 mo. | 68% vs. 97% | 9% vs. 47% | 38% vs. 70% (@ 60 mo.) |

CXCR4^{Mut} vs CXCR4^{WT}

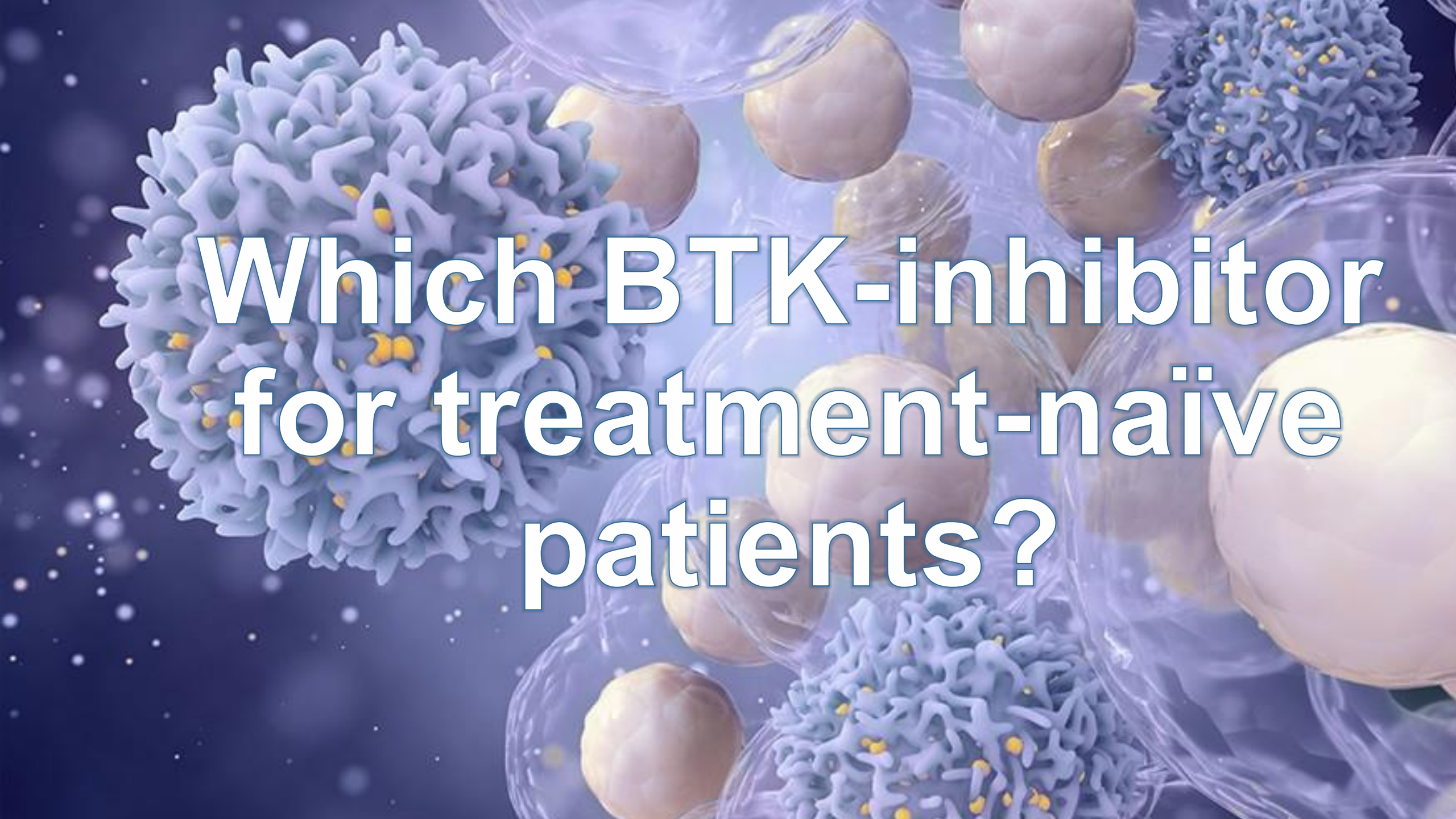
Median Time to Major Response: (4.2 vs. 1.9 mos)

Median Major RR: 71% vs. 87%

Median ≥VGPR: 14% vs. 41%

PFS: 59% vs. 75% @4 years

| | | | | | | |
|-----------------------|---------|--------------|------------------|-------------|-------------|------------------------|
| ASPEN Cohort 1 | TN, R/R | Ibrutinib | 6.6 vs. 2.8 mos. | 65% vs. 82% | 10% vs. 24% | 49% vs. 75% (@ 42 mo.) |
| | TN, R/R | Zanubrutinib | 3.4 vs. 2.8 mos. | 70% vs. 82% | 18% vs. 34% | 73% vs. 81% (@ 42 mo.) |

A 3D scientific illustration featuring several blue, complex, branched molecular structures, likely representing BTK-inhibitors, and several brown, spherical particles. The background is a dark blue gradient with small white specks, suggesting a microscopic or molecular environment. The text is overlaid in the center in a white, bold, sans-serif font with a thin blue outline.

**Which BTK-inhibitor
for treatment-naïve
patients?**

Long-term f/u for Phase III ASPEN study in WM

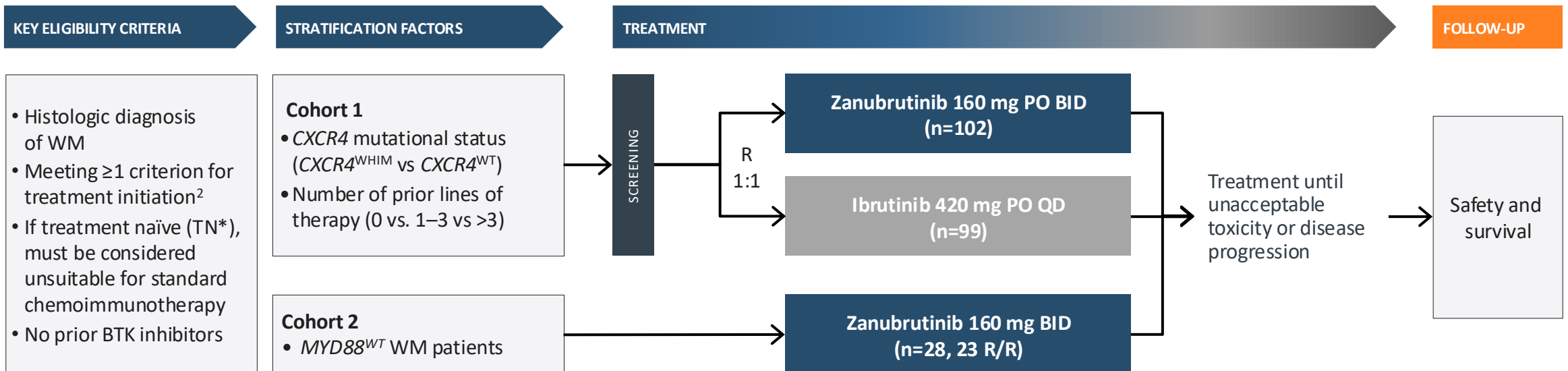
PHASE 3

Study Identifier: BGB-3111-302,
NCT03053440

Primary Endpoint: CR/VGPR rate

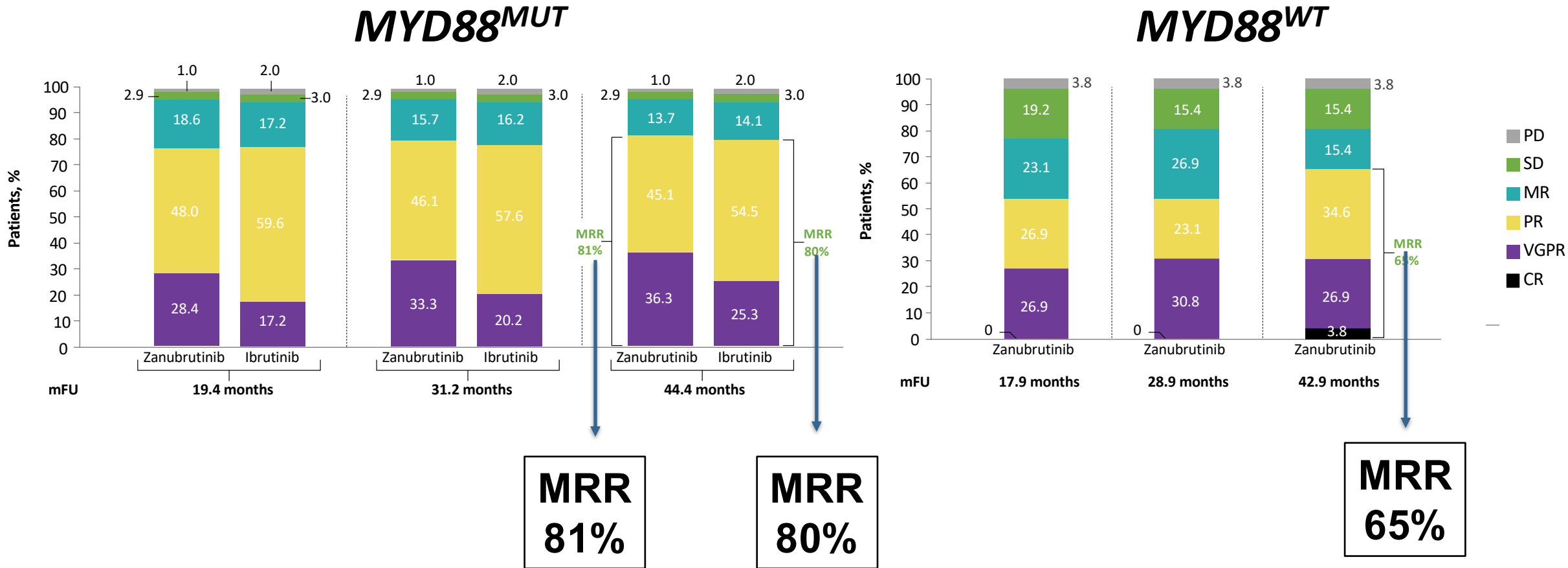
Key Secondary Endpoints: MRR (\geq PR), PFS, OS, DOR, symptom resolution, safety

Exploratory Endpoints: PK, QoL



Best Overall Responses by Investigators

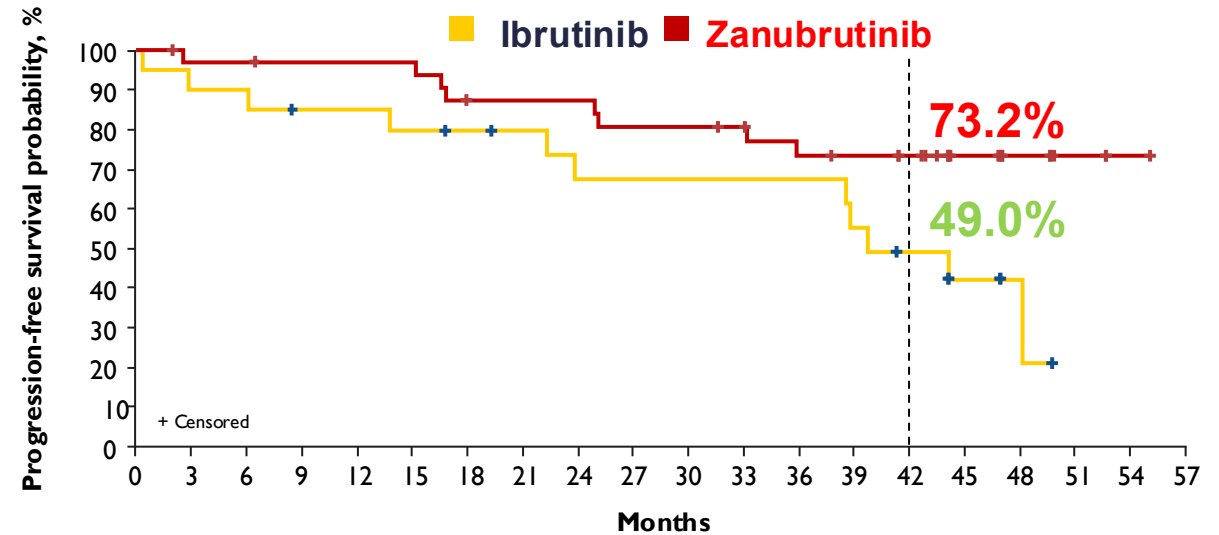
ASPEN – Long-term follow-up



Response Assessment by CXCR4 Mutation Status

ASPEN – Long-term follow-up (Cohort 1)

| | <i>CXCR4</i> ^{MUT} | | <i>CXCR4</i> ^{WT} | |
|--|-----------------------------|---------------------|----------------------------|---------------------|
| | Ibrutinib (n=20) | Zanubrutinib (n=33) | Ibrutinib (n=72) | Zanubrutinib (n=65) |
| VGPR or better | 2 (10.0) | 7 (21.2) | 22 (30.6) | 29 (44.6) |
| Major response | 13 (65.0) | 26 (78.8) | 61 (84.7) | 54 (83.1) |
| Overall response | 19 (95.0) | 30 (90.9) | 68 (94.4) | 63 (96.9) |
| Time to major response, median (months) | 6.6 | 3.4 | 2.8 | 2.8 |
| Time to VGPR, median (months) | 31.3 | 11.1 | 11.3 | 6.5 |



| | Zanubrutinib | Ibrutinib |
|----------------------|-------------------|-----------|
| Events, n (%) | 8 (24.2) | 11 (55.0) |
| HR (95% CI) | 0.50 (0.20, 1.29) | |

PFS impacted by MYD88^{WT} or TP53^{ALT} mutation status following treatment with Zanubrutinib in ASPEN Study

| | Cohort 1/MYD88 MUT (n=102) | Cohort 2/MYD88 WT (n=26) |
|--|-------------------------------|-----------------------------|
| 60-mo event-free rate for PFS, % (95% CI) | 74.8 (64.5, 82.5) | 39.3 (20.0-58.1) |
| <i>CXCR4^{WHIM}^a</i> | 70 (50.1-83.2) | NE |
| <i>CXCR4^{WT}^b</i> | 77.4 (64.2-86.3) | 31.6 (11.4-54.3) |
| <i>TP53^{MUT}^c</i> | 57.3 (35-74.4) | NE |
| <i>TP53^{WT}^d</i> | 81.2 (69.2-88.9) | 33.8 (11.8-57.5) |
| Unknown ^e | 75.0 (12.8-96.1) | 66.7 (19.5-90.4) |
| 60-mo event-free rate for OS, % (95% CI) | 82.8 (73.5-89.1) | 79.0 (56.4-90.8) |

Adverse Events of Interest (Cohort 1)

ASPEN – Long-term follow-up

| AEs, ^a n (%) | All grades | | Grade ≥3 | |
|--|-----------------------|-------------------------|---------------------|-------------------------|
| | Ibrutinib (n=98) | Zanubrutinib (n=101) | Ibrutinib (n=98) | Zanubrutinib (n=101) |
| Infection | 78 (79.6) | 80 (79.2) | 27 (27.6) | 22 (21.8) |
| Bleeding | 61 (62.2) | 56 (55.4) | 10 (10.2) | 9 (8.9) |
| Diarrhea | 34 (34.7) | 23 (22.8) | 2 (2.0) | 3 (3.0) |
| Hypertension* | 25 (25.5) | 15 (14.9) | 20 (20.4)* | 10 (9.9) |
| Atrial fibrillation/flutter* | 23 (23.5)* | 8 (7.9) | 8 (8.2)* | 2 (2.0) |
| Anemia | 22 (22.4) | 18 (17.8) | 6 (6.1) | 12 (11.9) |
| Neutropenia*^b | 20 (20.4) | 35 (34.7)* | 10 (10.2) | 24 (23.8)* |
| Thrombocytopenia | 17 (17.3) | 17 (16.8) | 6 (6.1) | 11 (10.9) |
| Second primary malignancy/ Non-Skin Cancers | 17 (17.3)/ 6 (6.1) | 17 (16.8)/ 6 (5.9) | 3 (3.1)/ 3 (3.1) | 6 (5.9)/ 4 (4.0) |

A 3D scientific illustration of B cells and plasma cells. The B cells are shown as large, spherical cells with a highly textured, blue, spiky surface and small yellow dots representing nuclei. The plasma cells are depicted as larger, more rounded cells with a smoother, light-colored surface and a prominent, dark nucleus. The background is a dark blue gradient with small white specks, suggesting a microscopic or cellular environment.

**BTK-inhibitors or
chemo-immunotherapy
for treatment-naïve
patients?**



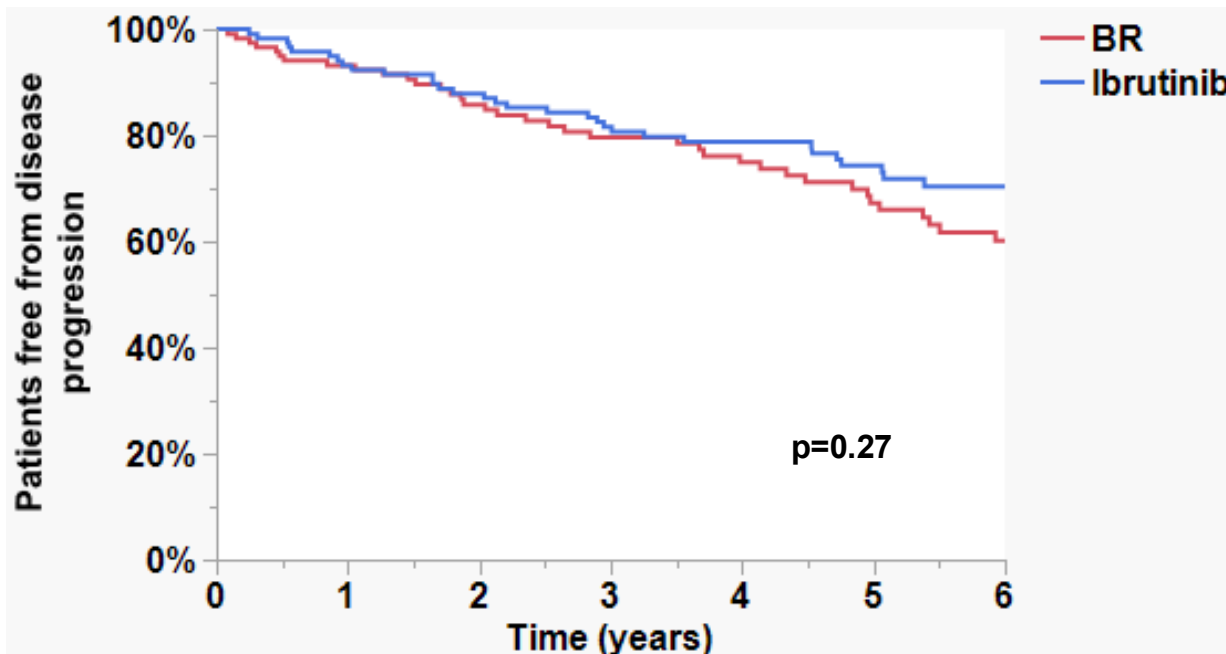
Benda-R vs. Ibrutinib in Treatment-Naïve WM (Multinational Retrospective Study)

Abeykoon et al, IWWM-12, 2024

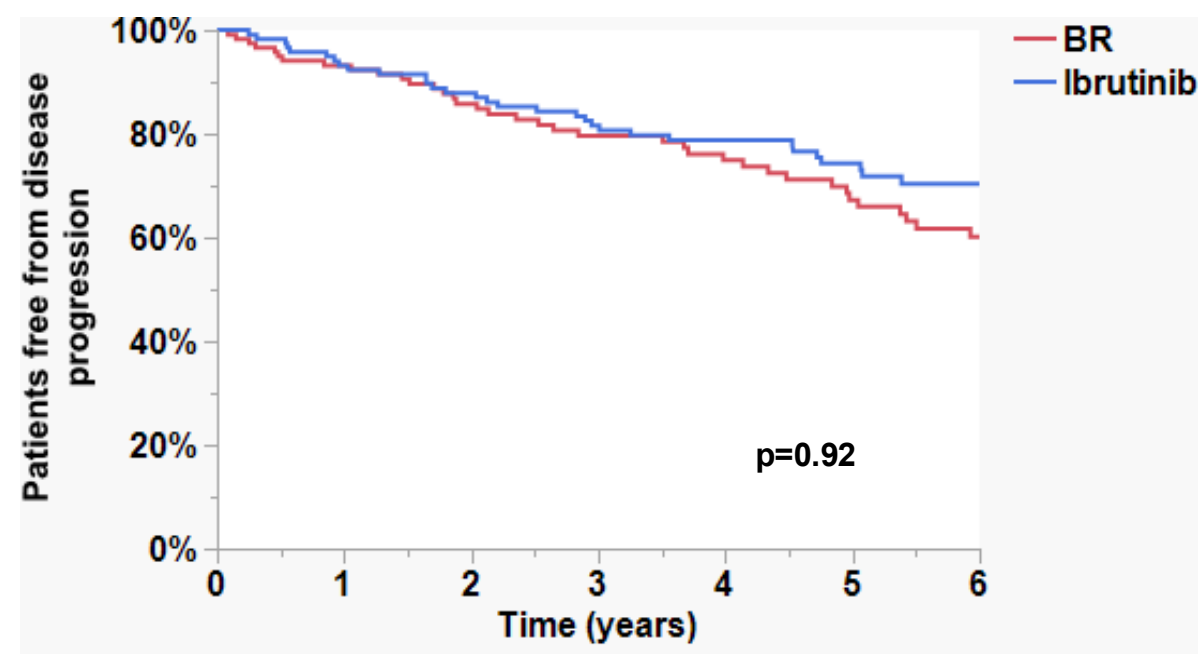
| Variable | BR (n=123) | Ibrutinib (n=123) | p- value |
|---------------------------|---------------------------|----------------------|------------------|
| Median Follow-up (yrs) | 6.0 (5.1-6.6) | 6.0 (5.4-6.6) | 0.89 |
| Median Age (yrs) | 68 (40-86) | 68 (39-86) | 0.9 |
| IPSS, % | | | 0.63 |
| Low | 11 | 17 | |
| Intermediate | 33 | 33 | |
| High | 56 | 48 | |
| Median Cycles | 6 (1-6) >4 cycles, 79% | 54 (1-114) | |
| ORR (%) | 95 | 93 | 0.47 |
| MRR (%) | 93 | 82 | 0.014 |
| CR (%) | 17 | 2 | <0.001 |
| ≥VGPR, % | 50 | 33 | 0.008 |

Benda-R vs. Ibrutinib: Time-to-event analyses

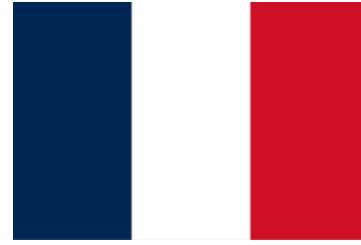
Progression Free Survival



Overall Survival



| Variable | BR (n=123) | Ibrutinib (n=123) | p-value |
|---------------|------------|-------------------|---------|
| 6-year PFS, % | 58 | 70 | 0.27 |
| 6-year OS, % | 80.5 | 84 | 0.92 |



BENDAMUSTINE AND RITUXIMAB (BR) IN WALDENSTROM MACROGLOBULINEMIA (WM): LONG-TERM RESULTS

Eveillard JR, Chaoui D, Cavalieri D, Dartigeas C, Willems L, Le Calloch R, Roos-Weil D, Merabet F, Roussel X, Bareaux B, Tricot S, Dupuis J, Poulain S, Laribi K, Leblond V

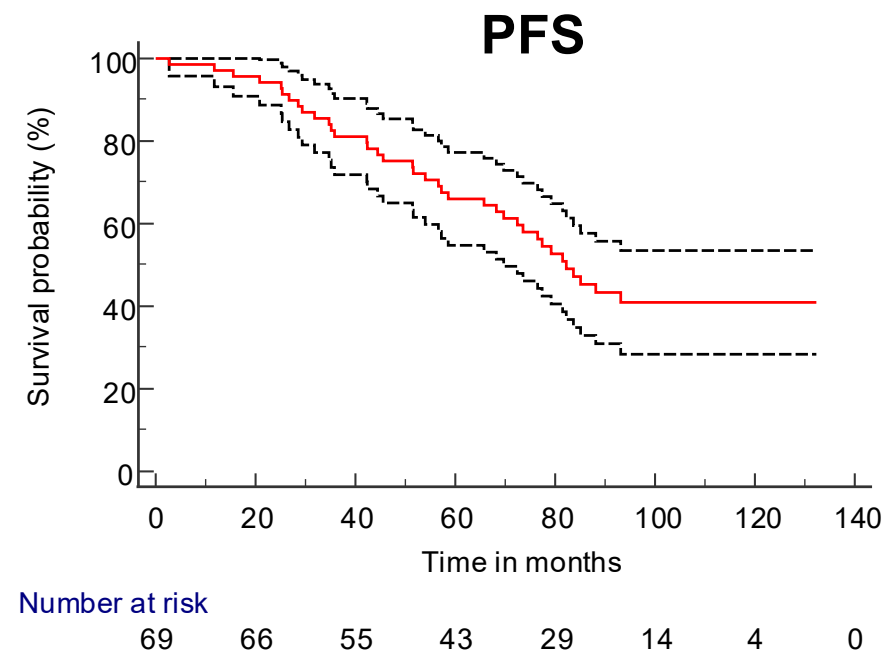


Leblond et al, IWWM-12, 2024

LONG TERM RESPONSE DATA FOR BENDA-R

| Response | N | % |
|------------------|-------|-----|
| ORR | 68/69 | 97% |
| MRR | 67/69 | 96% |
| VGPR (Neg IFX) | 13 | 19% |
| VGPR (Pos IFX) | 26 | 37% |
| Partial response | 28 | 40% |
| Minor response | 1 | |

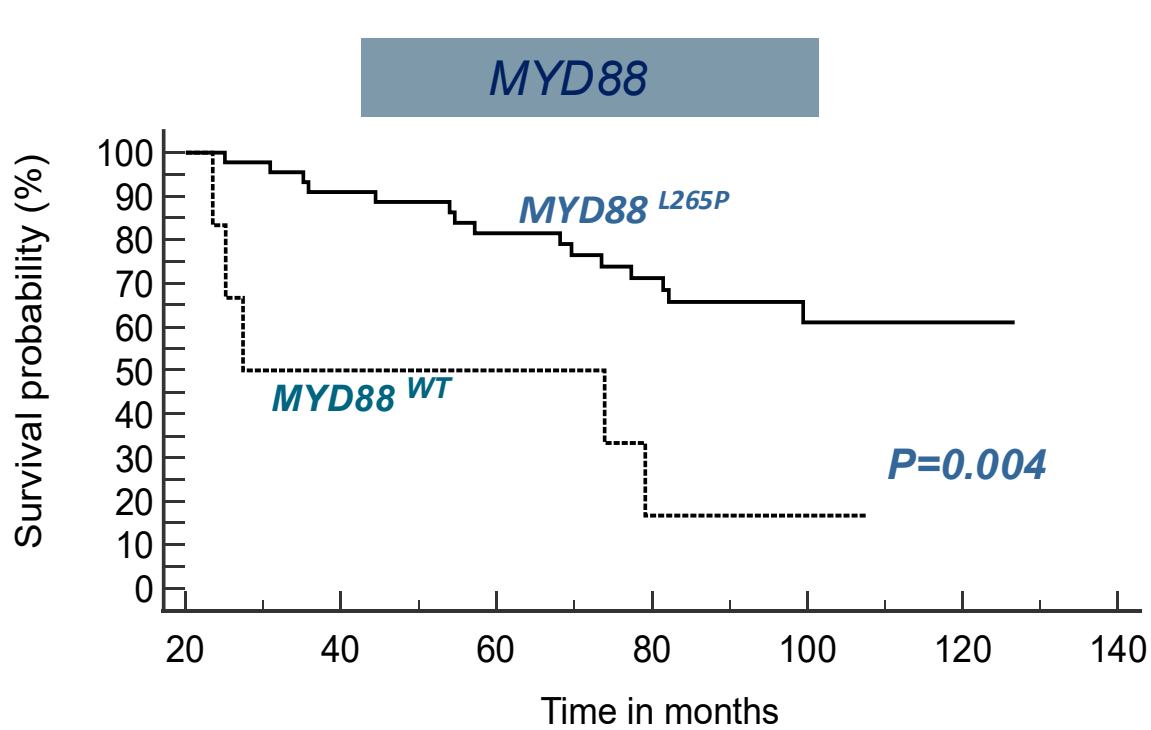
} **56%**



**Median PFS: 82.2 months
(69.7-93.1)**

IMPACT OF MUTATIONS ON OVERALL SURVIVAL

MYD88^{WT} patients



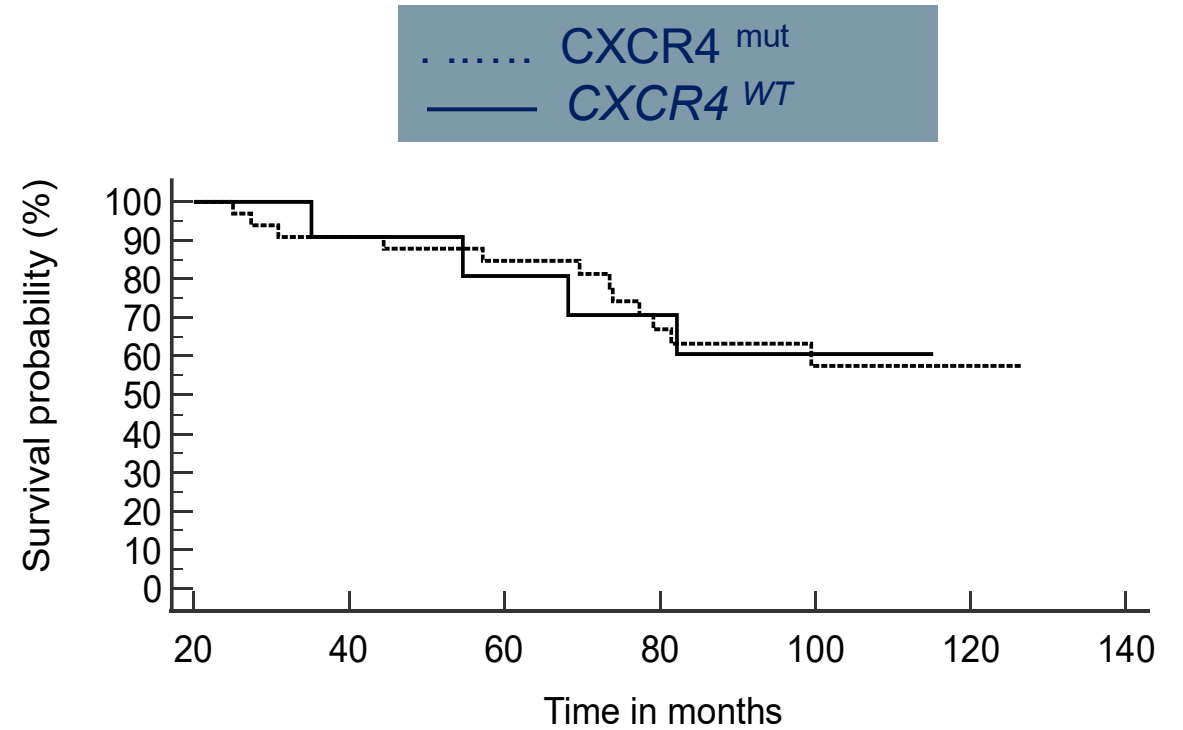
Number at risk

Group: 0

| | | | | | | |
|---|---|---|---|---|---|---|
| 6 | 3 | 3 | 1 | 1 | 0 | 0 |
|---|---|---|---|---|---|---|

Group: 1

| | | | | | | |
|----|----|----|----|----|---|---|
| 45 | 40 | 34 | 26 | 13 | 3 | 0 |
|----|----|----|----|----|---|---|



Number at risk

Group: 0

| | | | | | | |
|----|----|----|----|----|---|---|
| 33 | 30 | 27 | 18 | 10 | 3 | 0 |
|----|----|----|----|----|---|---|

Group: 1

| | | | | | | |
|----|----|---|---|---|---|---|
| 11 | 10 | 8 | 7 | 4 | 0 | 0 |
|----|----|---|---|---|---|---|

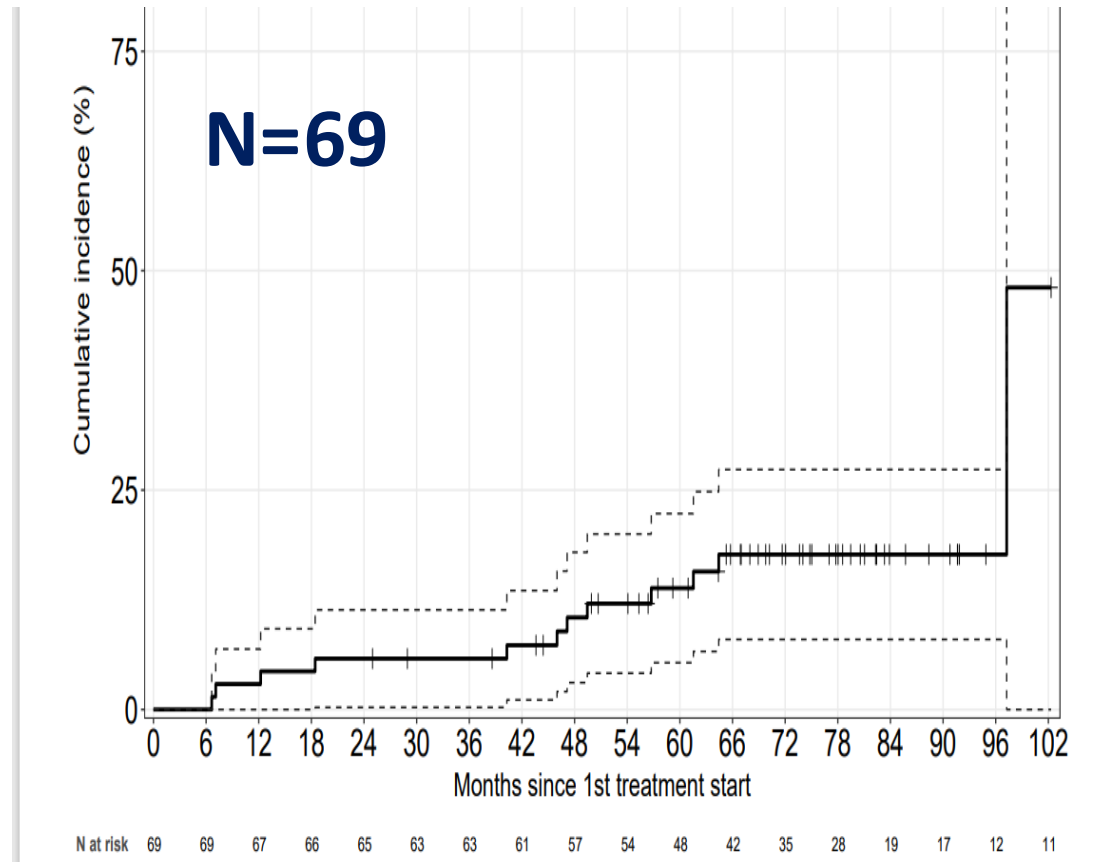
LATE-ONSET TOXICITIES FOR BENDA-R

| Type of Cytopenia | N | % | Duration (median) |
|-------------------|----|-----|-------------------|
| Neutropenia | 26 | 38% | 9m (3-24) |
| Anemia | 17 | 25% | 6m (3-36) |
| Thrombocytopenia | 11 | 16% | 9m (3-36) |

➤ Long-lasting cytopenia 35 pts (51%)

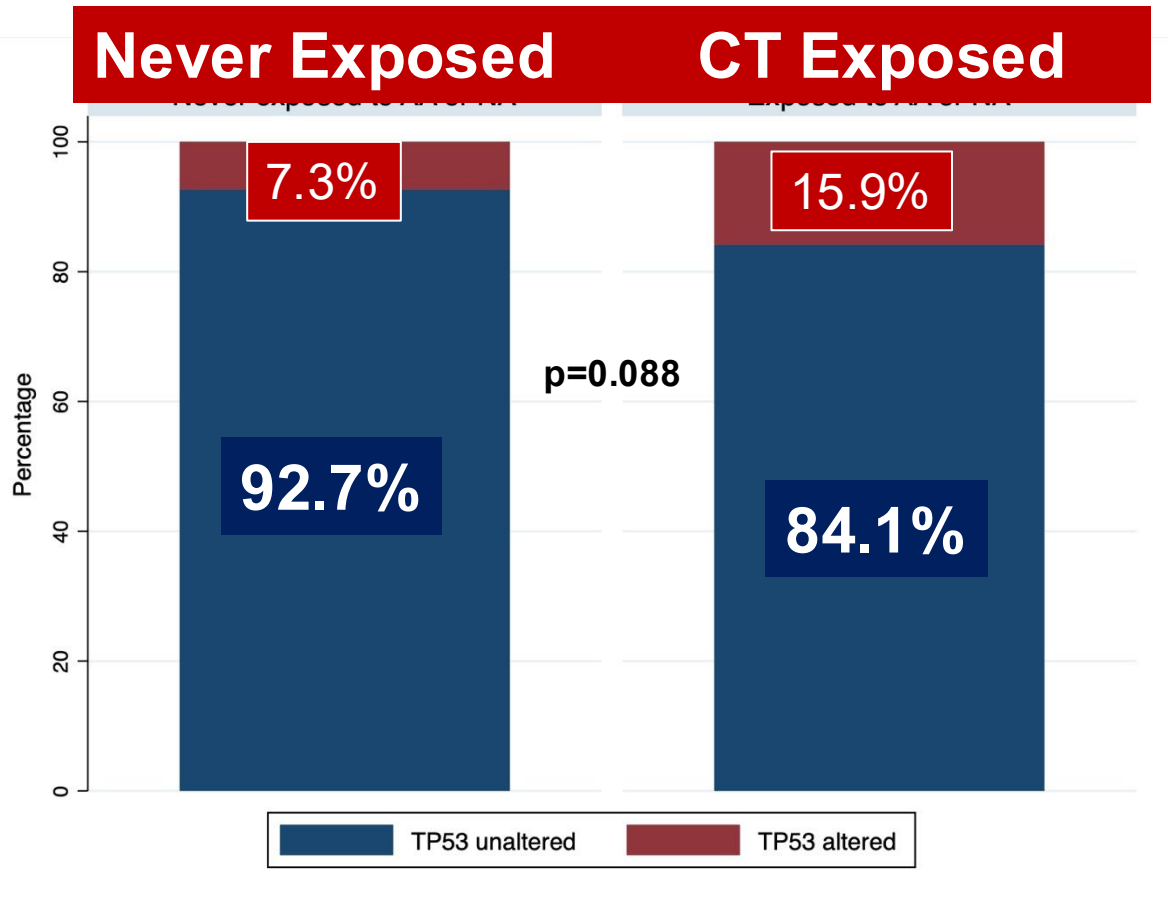
➤ Second malignancies: 12 patients

- 9 solid tumors (2 pancreas , 2 gastric, 1 colic, 1 oesophagus 1 lung, 1 skin, 1 breast)
- 3 MDS with 2 AML



Cumulative incidence of second malignancies of 17.66% [7.99-27.64] at 66 months.

TP53 ALT Were More Common in CT- vs. Non-CT-Exposed Patients

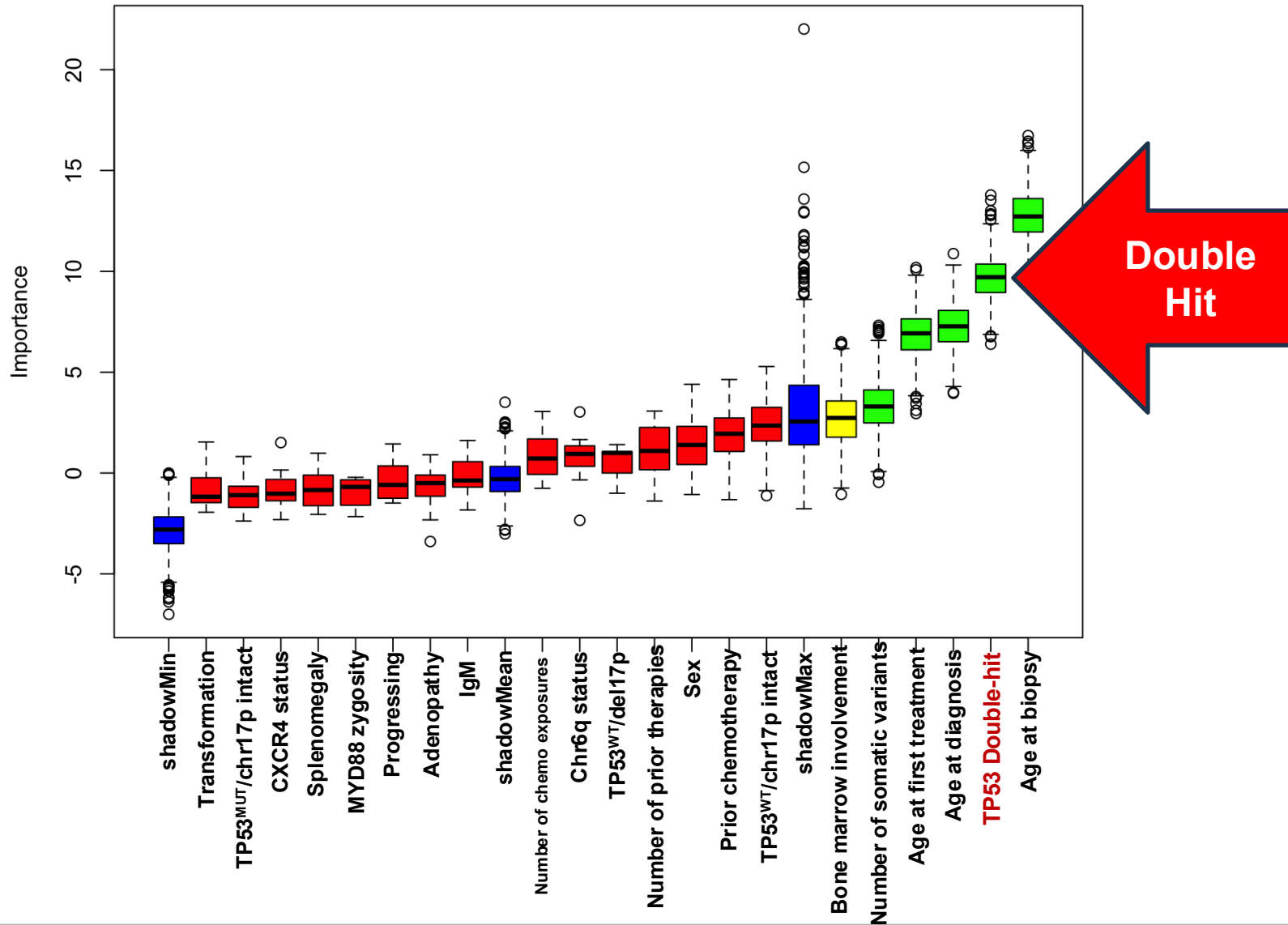


Multivariate Logistic Regression

| Term | p-value | OR |
|----------------------|---------|------|
| Prior AA or NA | 0.102 | 2.89 |
| CXCR4 Mutated | 0.118 | 2.28 |
| Age at Biopsy | 0.589 | 1.01 |
| >1 Prior Therapy | 0.945 | 0.96 |
| Actively Progressing | 0.909 | 1.06 |

Double-hit TP53^{ALT} were more common in patients who received both AA and NA (18.8%) versus either an AA or NA (6.1%) or no CT (3.6%); p=0.069 for three-way comparison.

Boruta Identifies TP53 Double-hit Status as One of the Most Relevant Predictors of OS



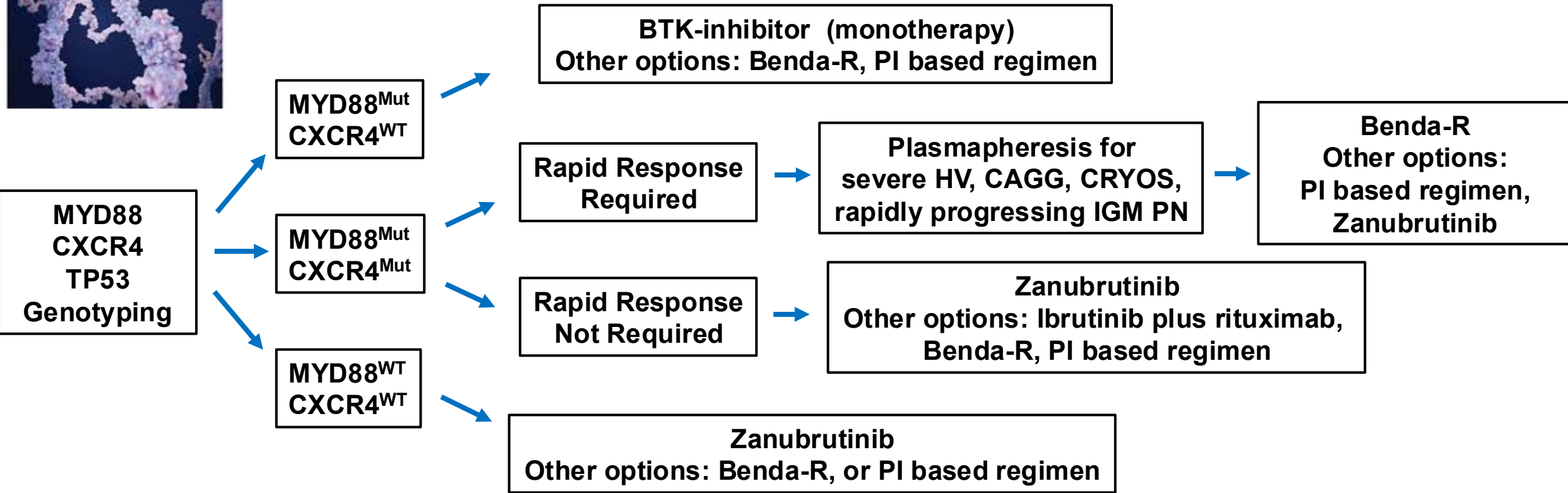
Cox proportional hazards model

| Term | p-value | HR |
|------------------------|--------------|-------------|
| Sex | 0.022 | 2.14 |
| Age at biopsy | <0.001 | 1.08 |
| CXCR4 Mutated | 0.389 | 1.30 |
| TP53 Single-hit | 0.737 | 0.78 |
| TP53 Double-hit | 0.001 | 3.85 |
| >1 Prior Therapy | 0.388 | 1.31 |

A Cox proportional hazards model adjusting for sex, age, CXCR4 status, amount of TP53 alterations, and number of prior therapies identified sex, age, and TP53 Double-hit as significantly impacting OS



Treatment Algorithm for Symptomatic Treatment-Naïve WM

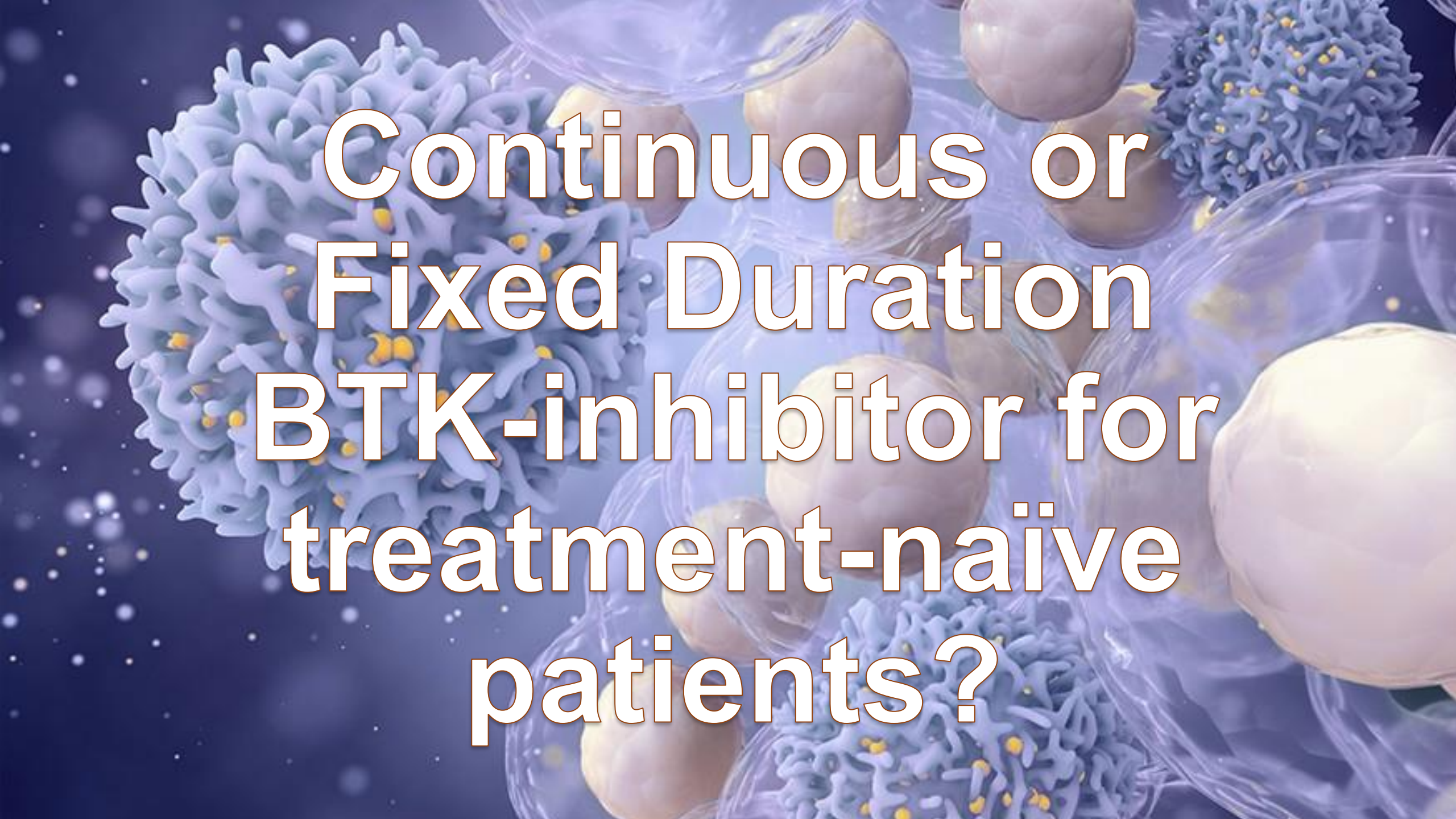


¹Zanubrutinib is recommended for those with MYD88^{WT}, CXCR4^{Mut} or TP53^{Alt}

²Bendamustine-R is recommended for those with bulky disease

³Bendamustine-R or Proteasome Inhibitors for symptomatic amyloid (with ASCT for eligible patients)

⁴BTK-inhibitor is recommended for BNS



**Continuous or
Fixed Duration
BTK-inhibitor for
treatment-naïve
patients?**

Ibrutinib and Venetoclax (IVEN) for Treatment Naive WM



**STUDY
TERMINATED**

Ibrutinib 420 mg (24 Months)

Venetoclax Ramp-up (1 Month), 400 mg (23 Months)

Median Time on Treatment: 10.2 months

Major RR 95%

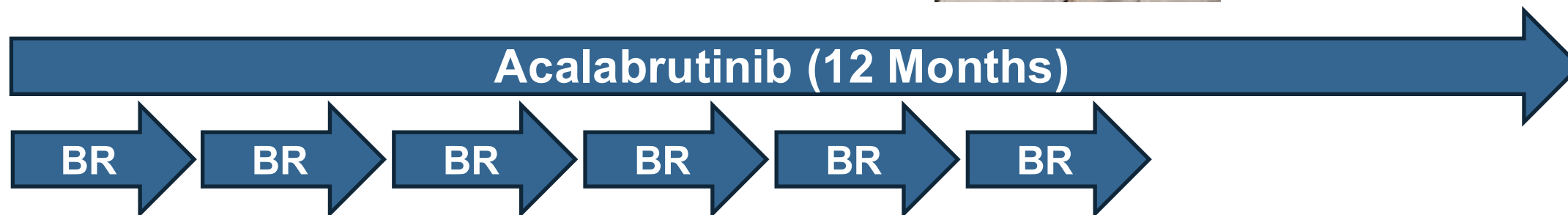
VGPR 42% (29% CXCR4^{Mut}; 50% CXCR4^{WT})

No CRs

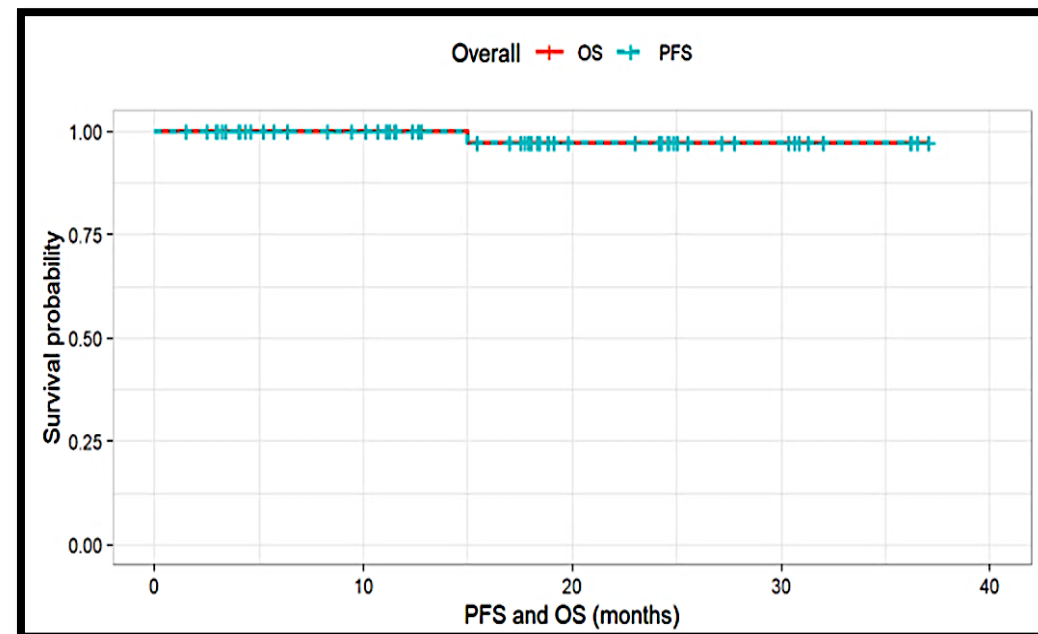
3 yr PFS: 51%. Not impacted by CXCR4 status

V-fib (9%; 2 G5 events)

Bendamustine, Rituximab and Acalabrutinib (BRAWM) for Treatment Naïve WM

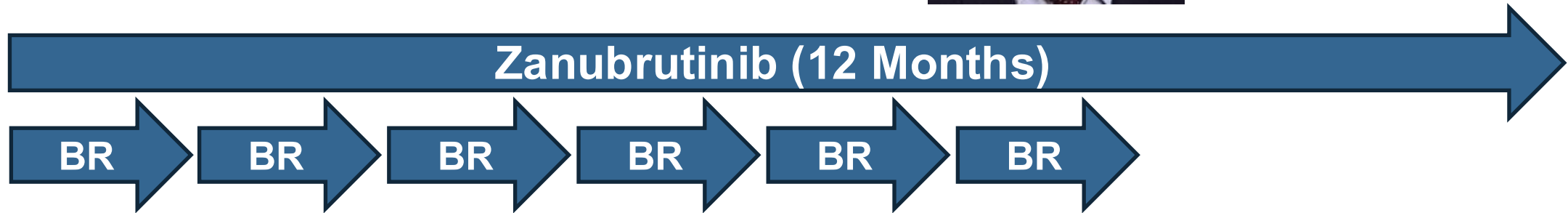


Major RR 95%
VGPR 38%
CR 18%
MRD 23% (18 mos)

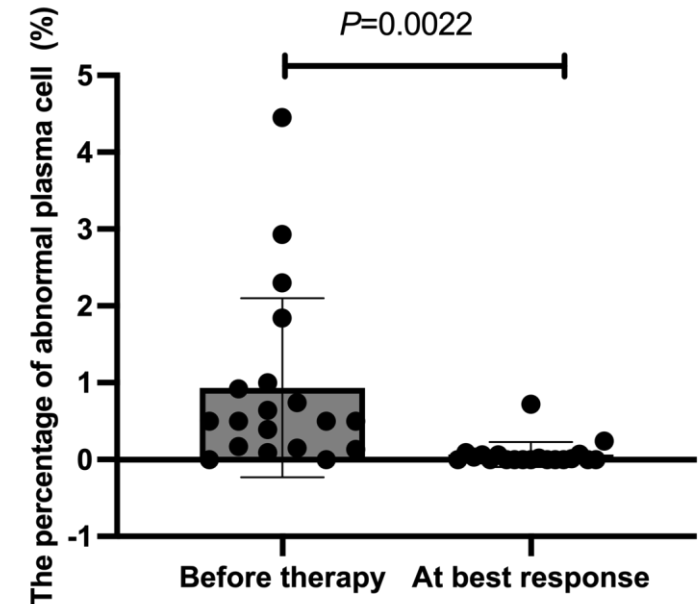
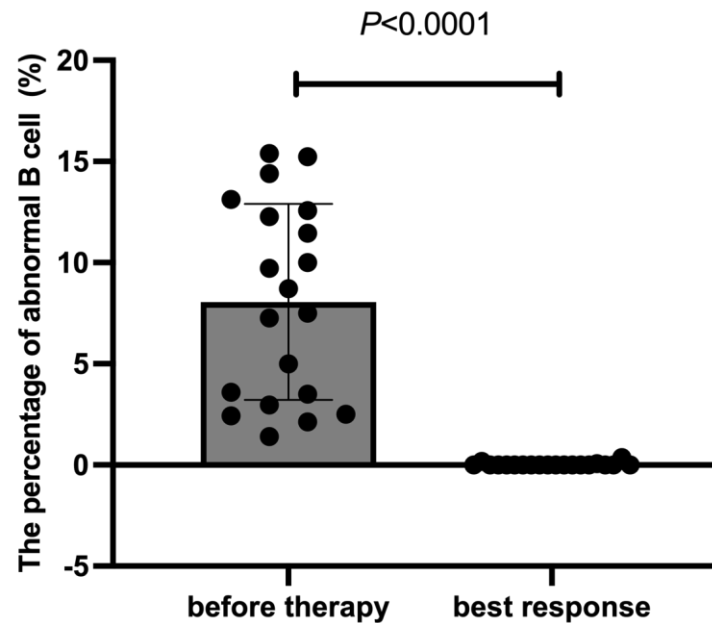


Berenstein et al, IWWM-12, 2024

Zanubrutinib, Bendamustine and Rituximab (ZBR) for Treatment Naïve WM

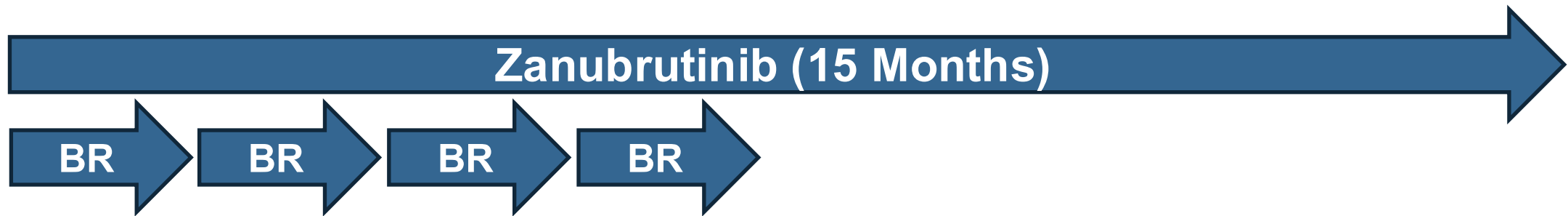


Major RR 95%
VGPR 65%
MRD (50% of evaluable patients)



Zanubrutinib, Bendamustine and Rituximab (ZeBRa) for Treatment Naïve WM

www.clinicaltrials.gov NCT06561347



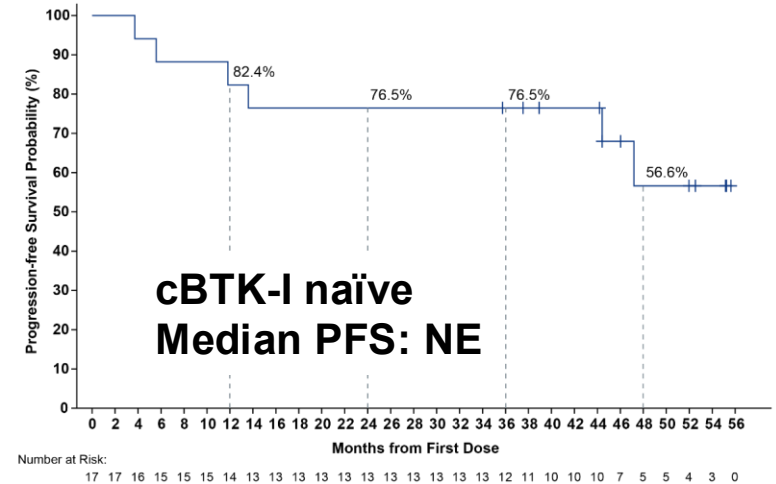
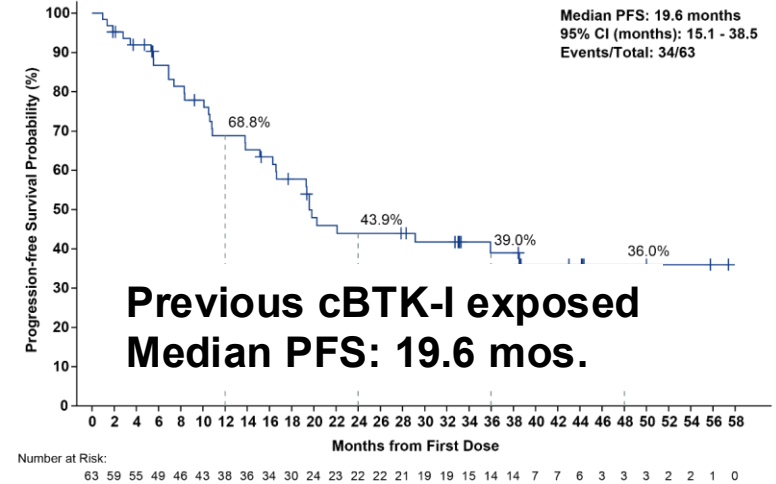
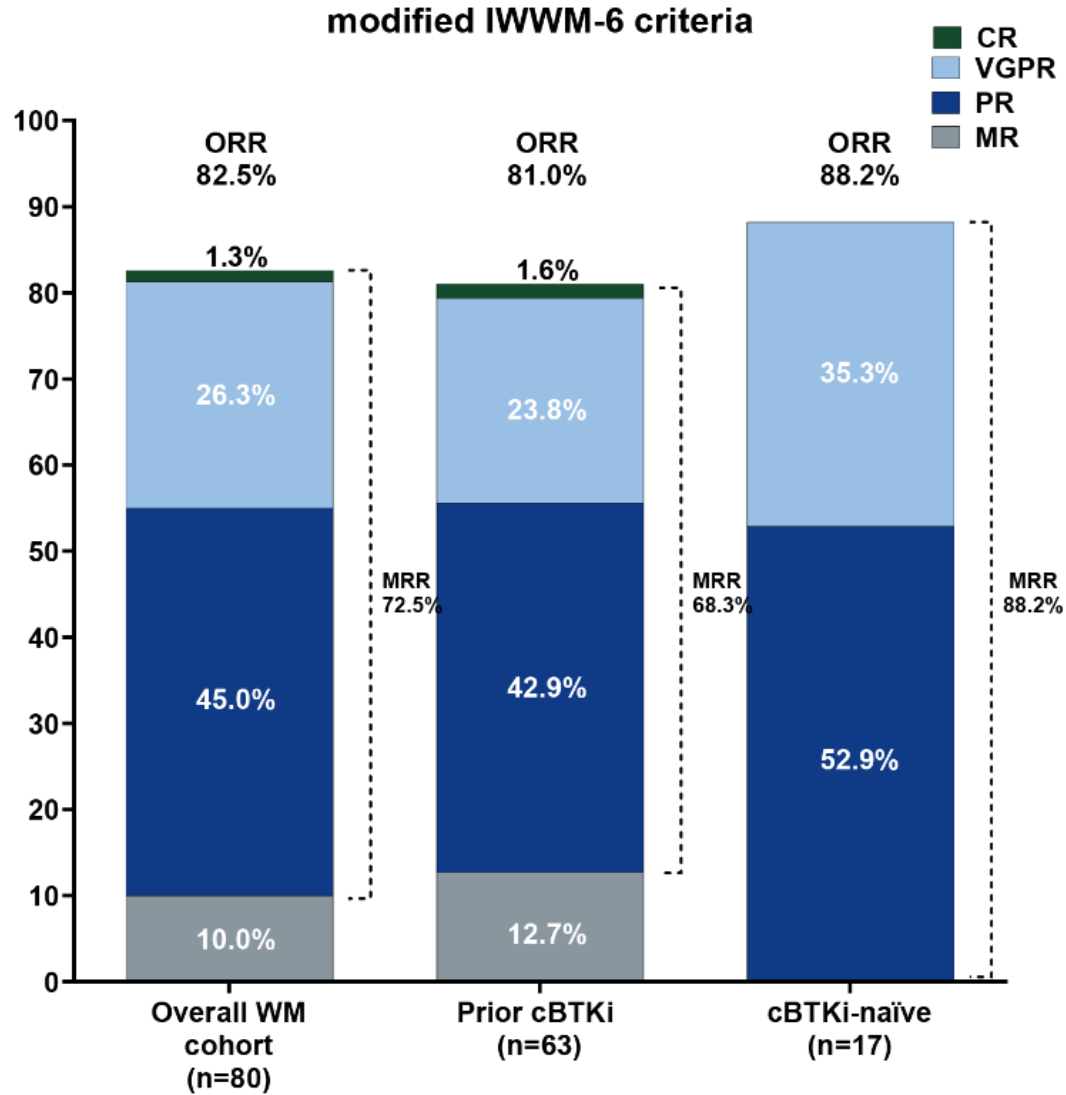
Inclusion Criteria:

- Presence of any MYD88 and CXCR4 mutation status
- Meeting criteria for treatment per IWWM2 criteria.
- Participants must meet the following organ and marrow function as defined below:
 - $ANC \geq 500/mcL$ believed to be caused by WM BM involvement.
 - Platelets $\geq 30,000/mcL$ believed to be caused by WM BM involvement. Hemoglobin ≥ 7 g/dL.
 - Total bilirubin ≤ 1.5 X institutional ULN
 - $AST/ALT \leq 2.5 \times$ institutional ULN
 - Creatinine clearance ≥ 30 mL/min using the Cockcroft-Gault formula



**How do we manage
covalent BTK-inhibitor
resistant disease?**

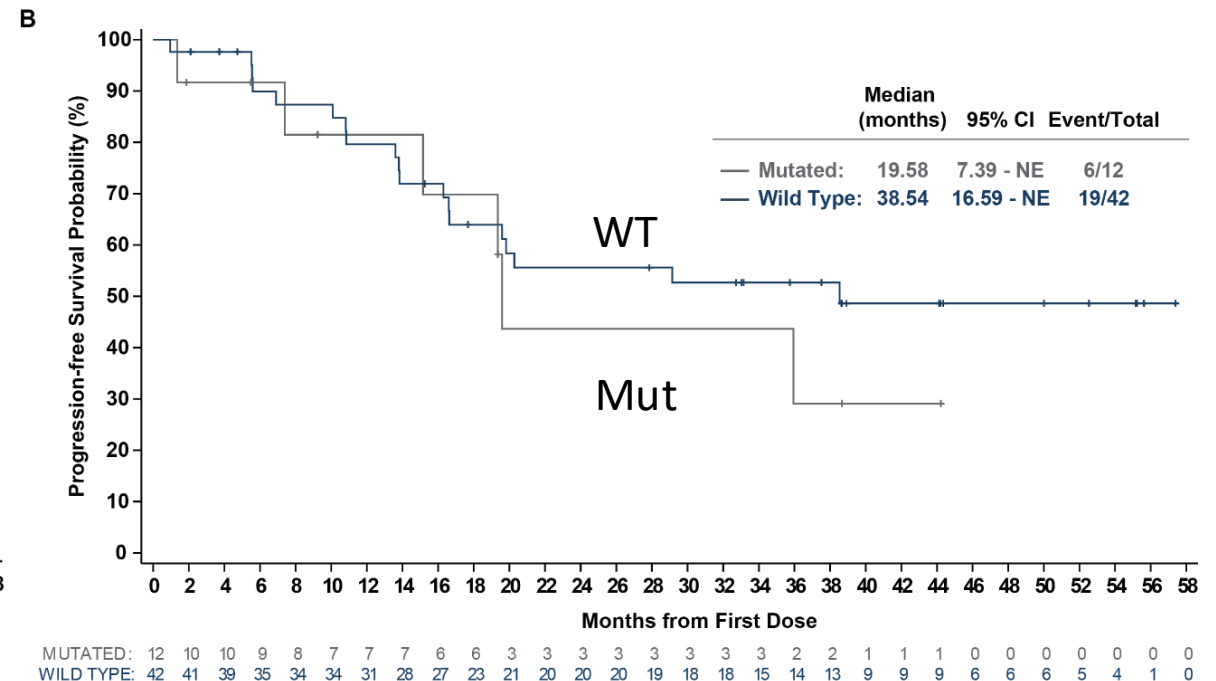
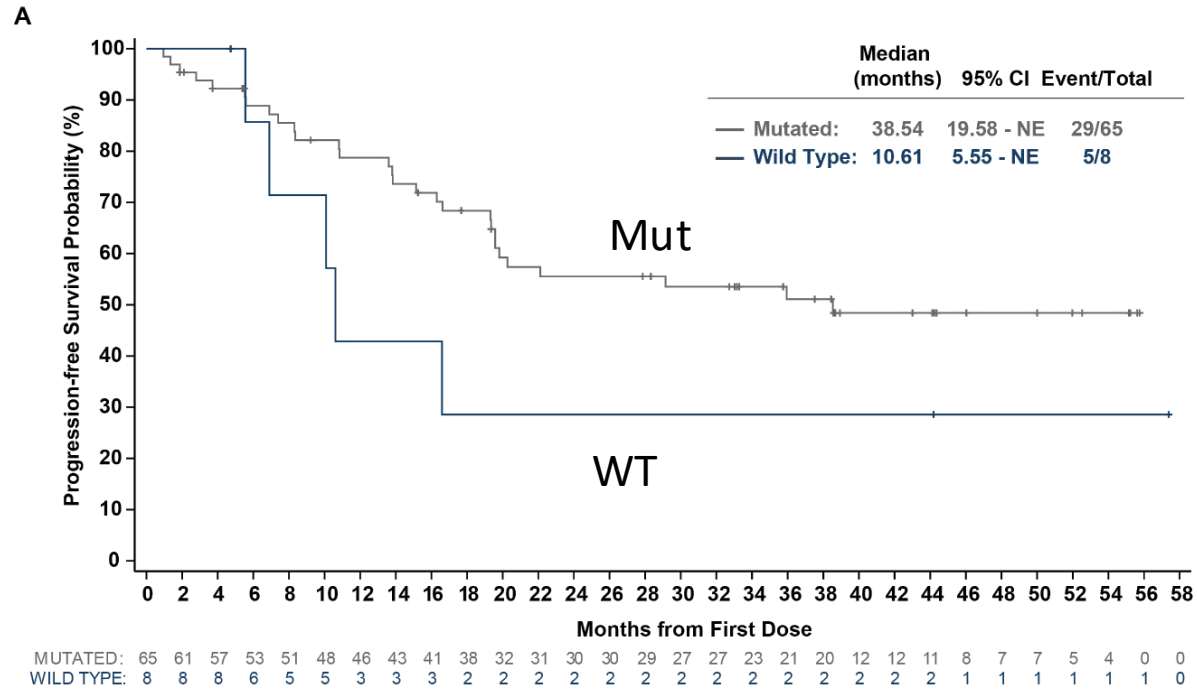
Pirtobrutinib in Previously Treated WM



Progression-free survival for previously treated WM Patients treated with Pirtobrutinib by MYD88 and CXCR4 Mutation Status

MYD88 Status

CXCR4 Status

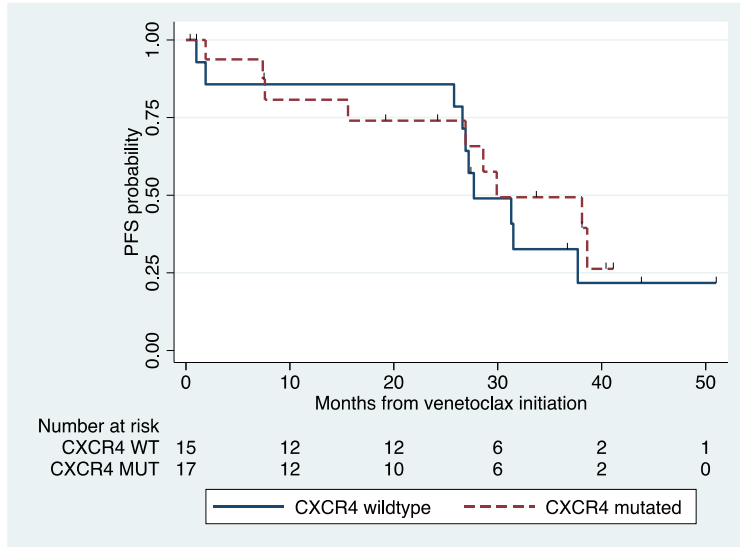
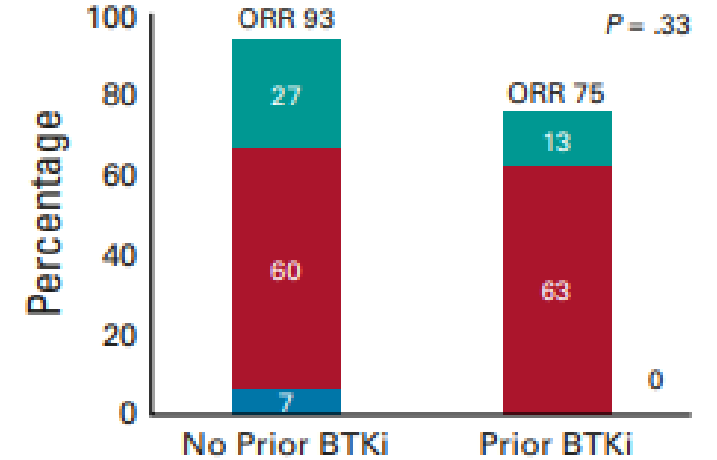


Venetoclax for Previously Treated WM

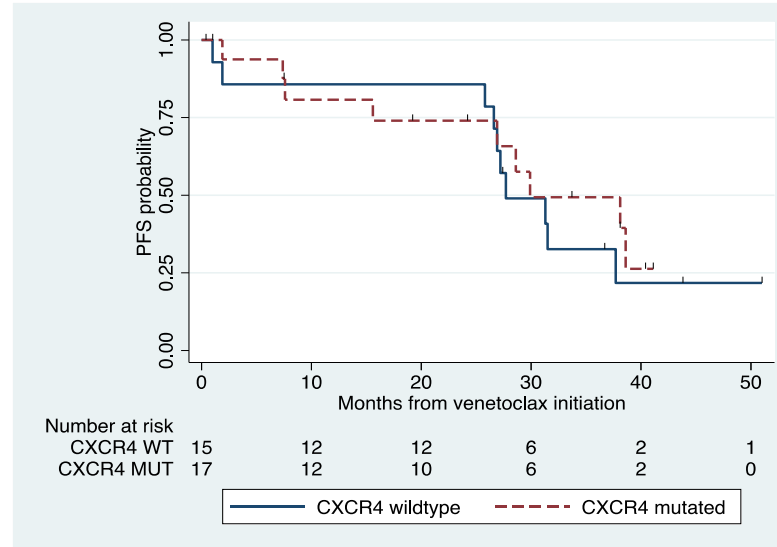
Dose escalation to 800 mg/day, 2 years treatment



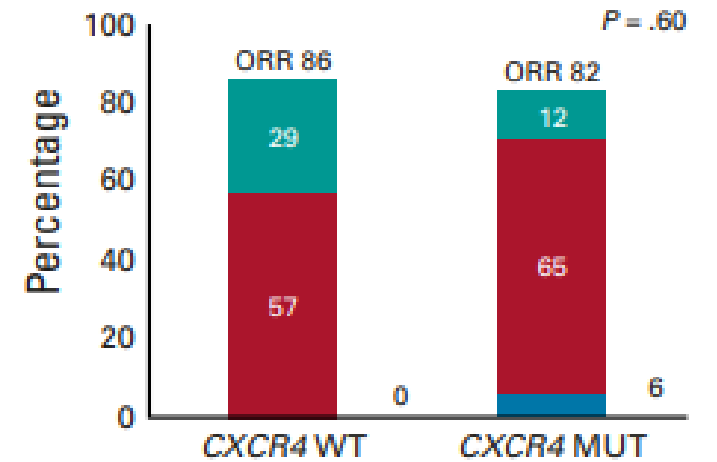
ORR 84%; Major RR 81%
Median PFS: 30 mos.
Not impacted by CXCR4 mutation status.
Grade ≥ 3 neutropenia: 45%



PFS for All Pts



PFS by CXCR4 Mut Status



MR PR VGPR

Pirtobrutinib and Venetoclax (PVEN) Treatment schema



Key inclusion criteria

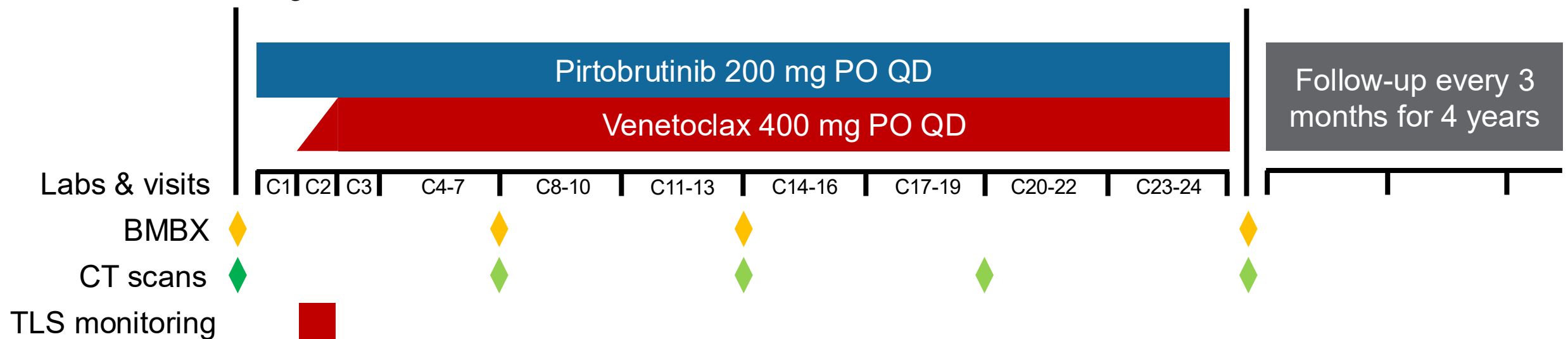
- 18+ years
- Diagnosis and need for treatment per IWWM2
- MYD88 L265P
- 1+ previous therapy

Key exclusion criteria

- CNS involvement
- Pregnancy
- Active HIV, HBV, HCV infection
- Previous non-covalent BTK inhibitor

Screening

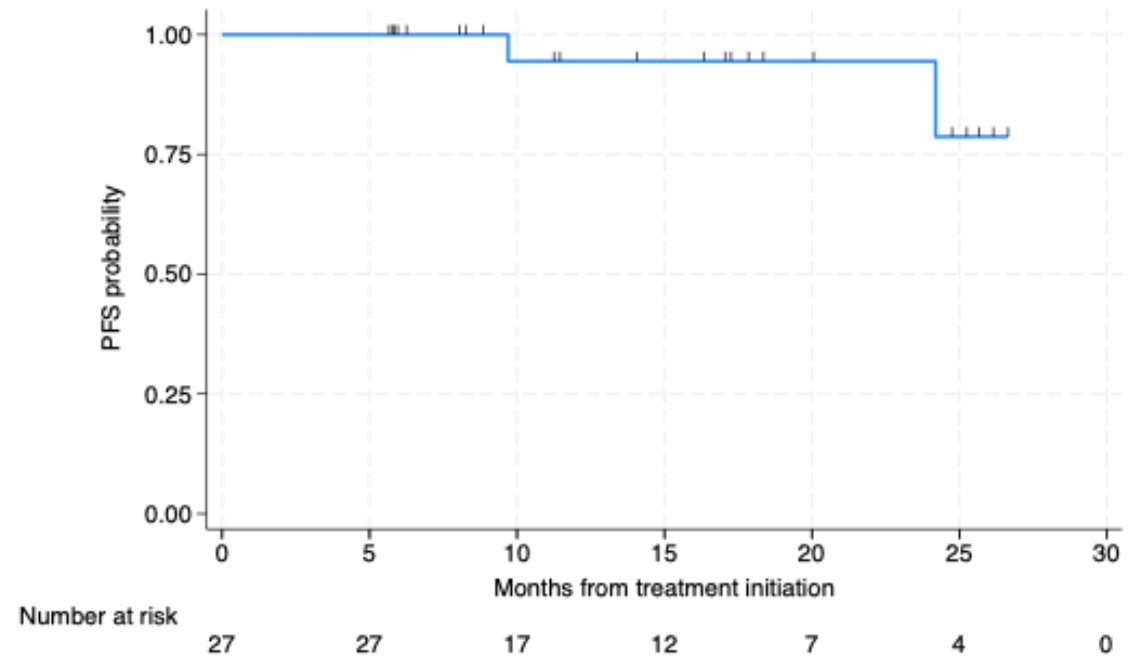
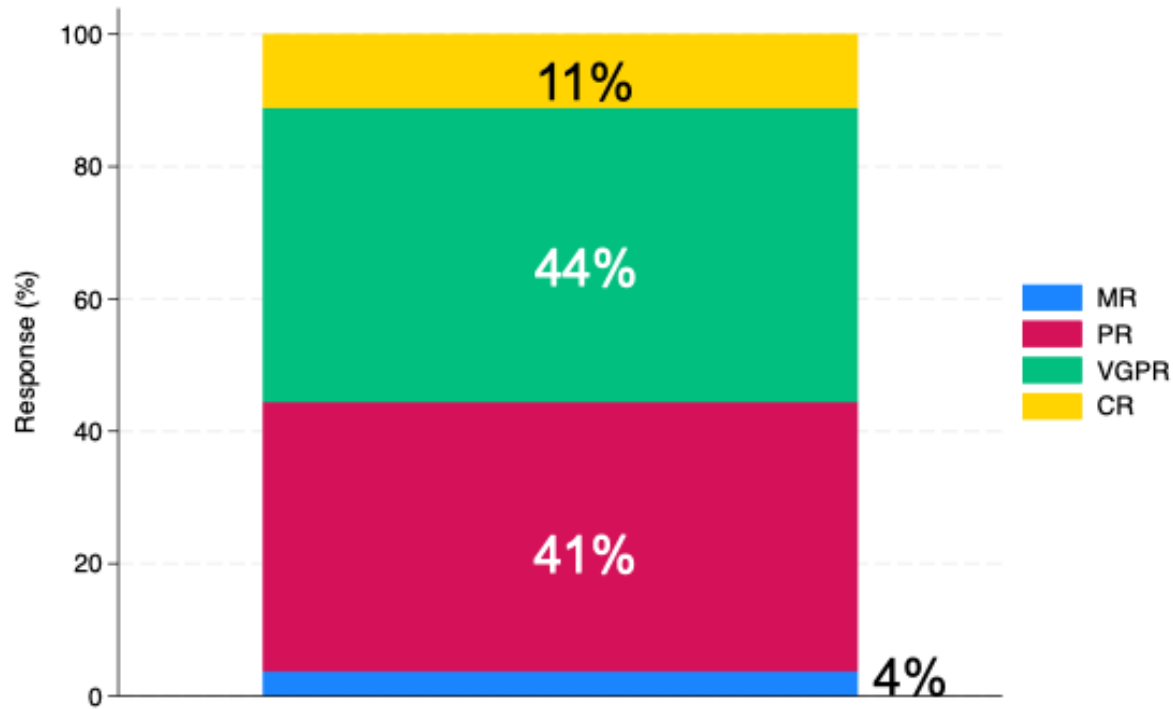
EOT



PVEN Responses

ORR 100%
Major RR 96%
VGPR/CR 11%

12 mo. PFS rate 94%
(95% CI 67-99%)



Median follow-up: 11 months (95% CI 8-18)

Castillo et al, ASH 2025

THE BTK DEGRADER BGB-16673 IS HIGHLY ACTIVE IN R/R WM

- Responses were observed at the lowest dose (100 mg; 7/9) and in patients with prior cBTK inhibitor (22/27) or ncBTK inhibitor (4/4)

| | Total ^a (N=27) |
|---|---------------------------|
| Best overall response, n (%) | |
| VGPR | 7 (25.9) |
| PR | 13 (48.1) |
| MR | 2 (7.4) |
| SD | 3 (11.1) |
| Not evaluable | 1 (3.7) |
| Discontinued prior to first assessment | 1 (3.7) |
| ORR, n (%)^b | 22 (81.5) |
| Major response rate, n (%)^c | 20 (74.1) |
| DCR, n (%)^d | 25 (93.0) |
| Follow-up, median (range), months | 5.0 (0.8-24.6) |
| Time to first response, median (range), months^e | 1.0 (0.9-3.7) |

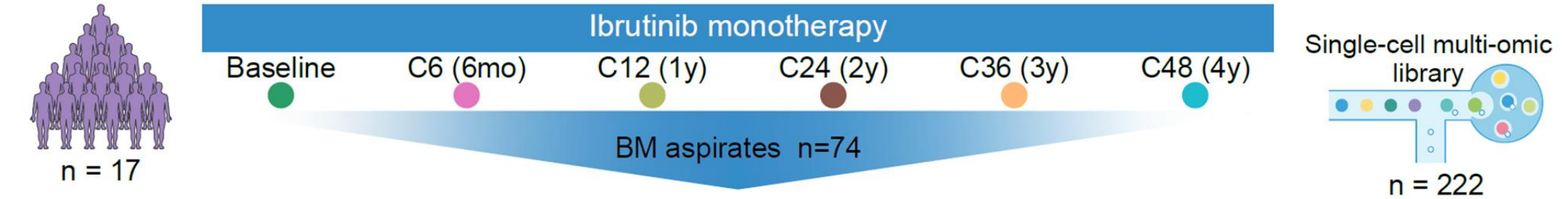
^a cBTK, covalent BTK; DCR, disease control rate; MR, minor response; ncBTK, noncovalent BTK; VGPR, very good partial response.

| Mutation status, n/N tested (%) | Total ^a (N=27) |
|---------------------------------|---------------------------|
| BTK | |
| Mutated | 10/11 (90.9) |
| Unmutated | 11/14 (78.6) |
| Unknown | 1/2 (50.0) |
| MYD88 | |
| Mutated | 20/24 (83.3) |
| Unmutated | 1/2 (50.0) |
| Unknown | 1/1 (100) |
| CXCR4 | |
| Mutated | 11/12 (91.7) |
| Unmutated | 10/13 (76.9) |
| Unknown | 1/2 (50.0) |
| TP53 | |
| Mutated | 12/13 (92.3) |
| Unmutated | 9/12 (75.0) |
| Unknown | 1/2 (50.0) |

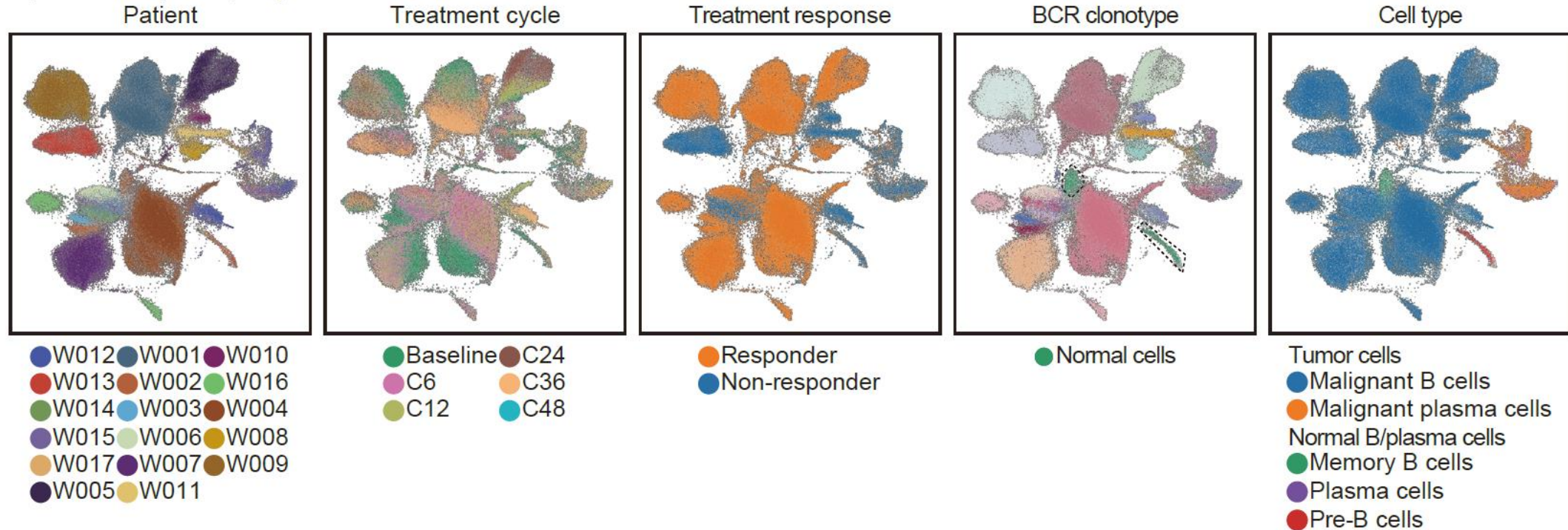
best overall response of SD or better.

Seymour et al, IWWM-12, 2024

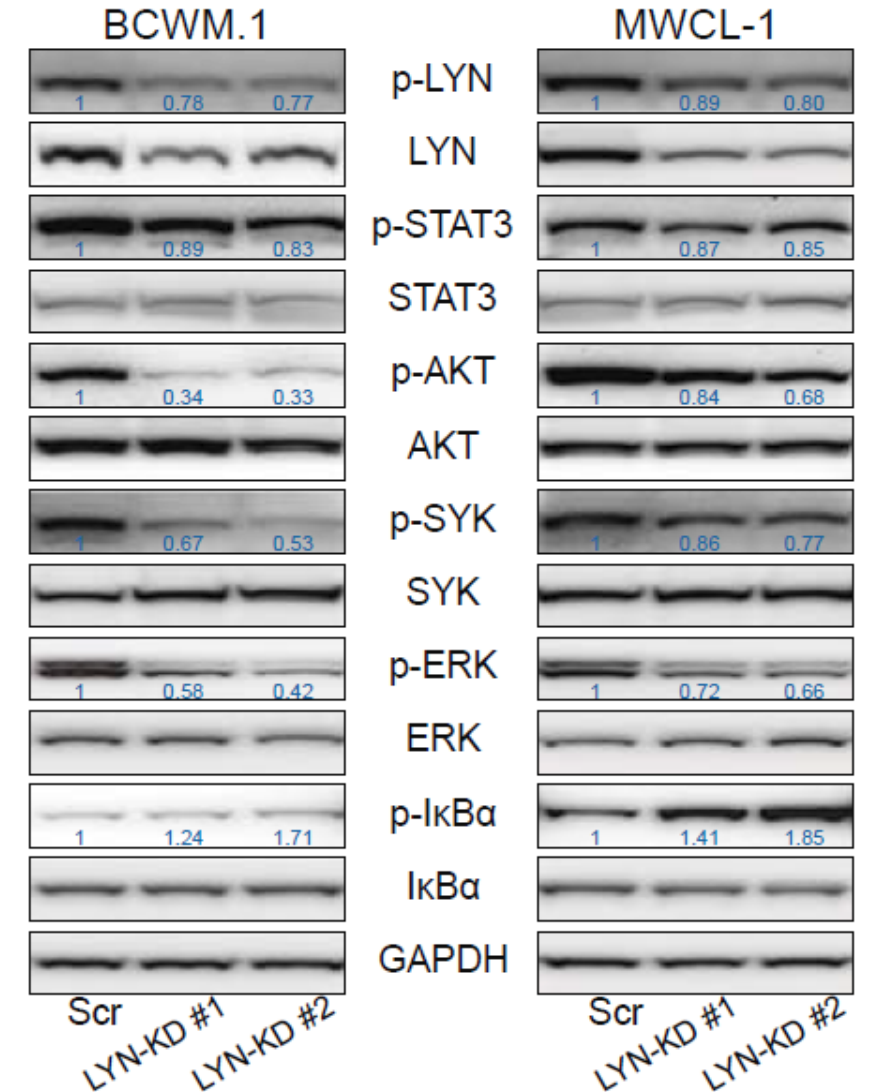
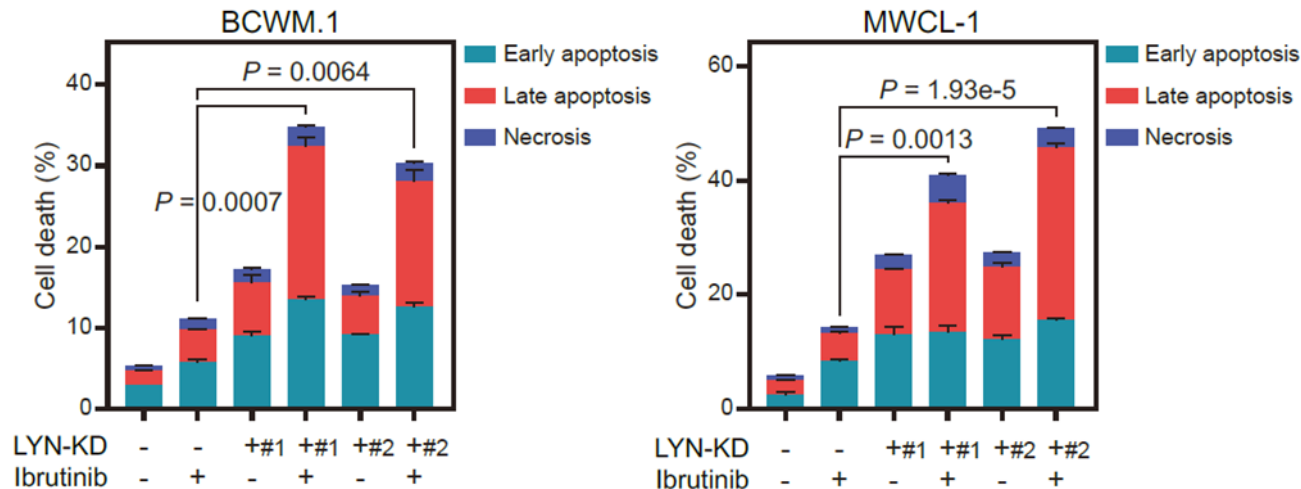
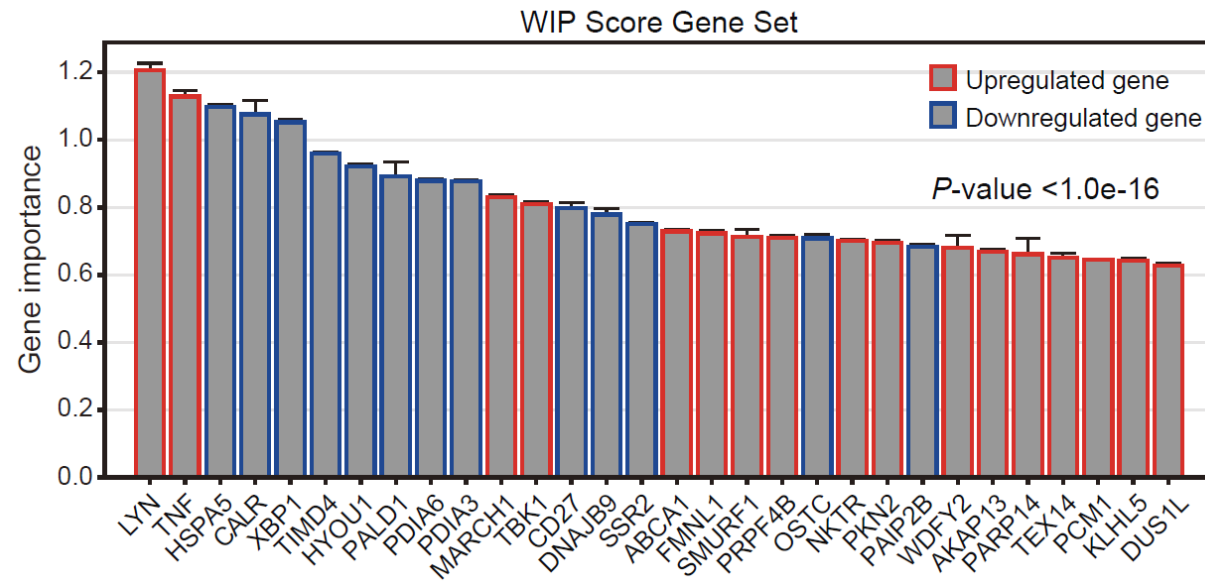
Single-cell sequencing of longitudinal BM aspirates from WM patients treated with ibrutinib monotherapy with serial 5-year tumor collection and follow-up.



B/plasma cells (n = 148,458)



The SRC family member LYN emerged as a top gene in ibrutinib-resistant WM disease, and LYN knockdown decreased pro-survival signaling and increased ibrutinib sensitivity.

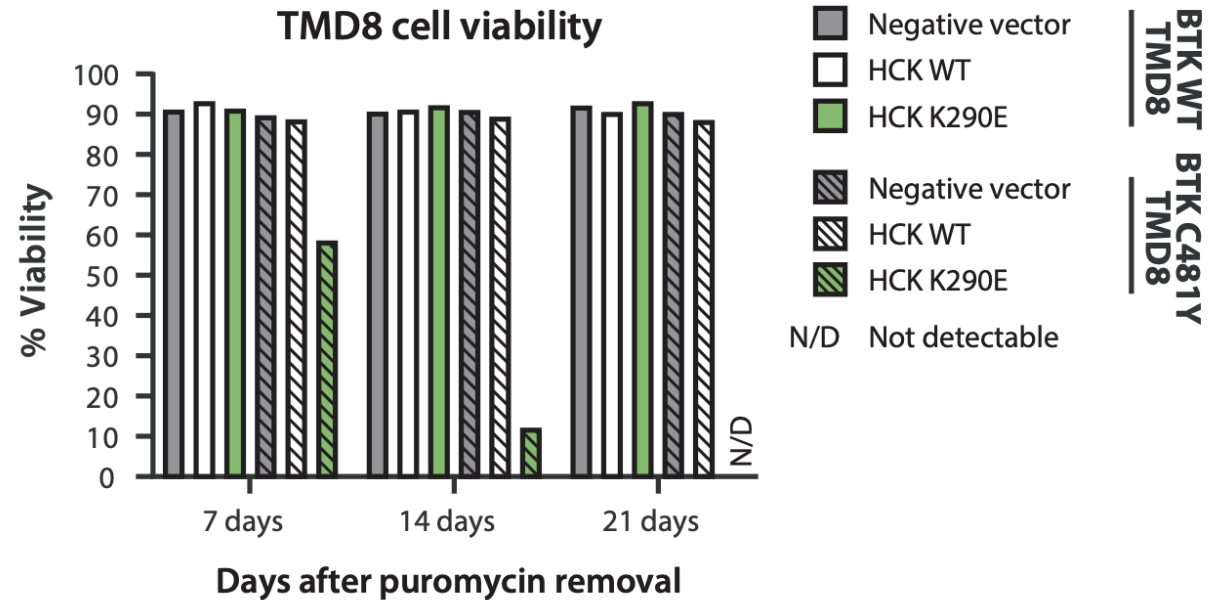
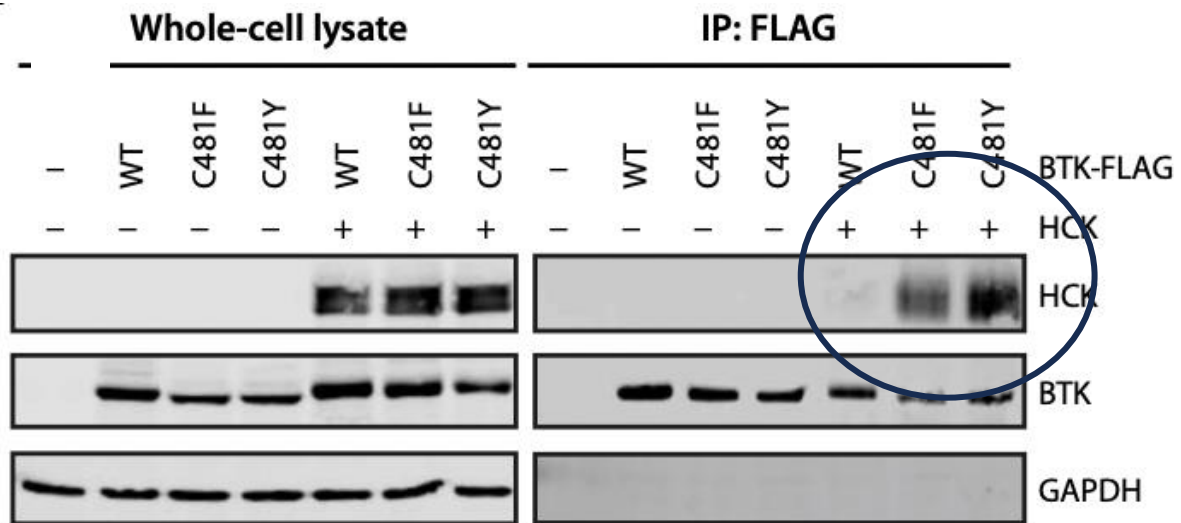


CANCER

Kinase-deficient BTK mutants confer ibrutinib resistance through activation of the kinase HCK

BTK^{Cys481Phe}
BTK^{Cys481Tyr}

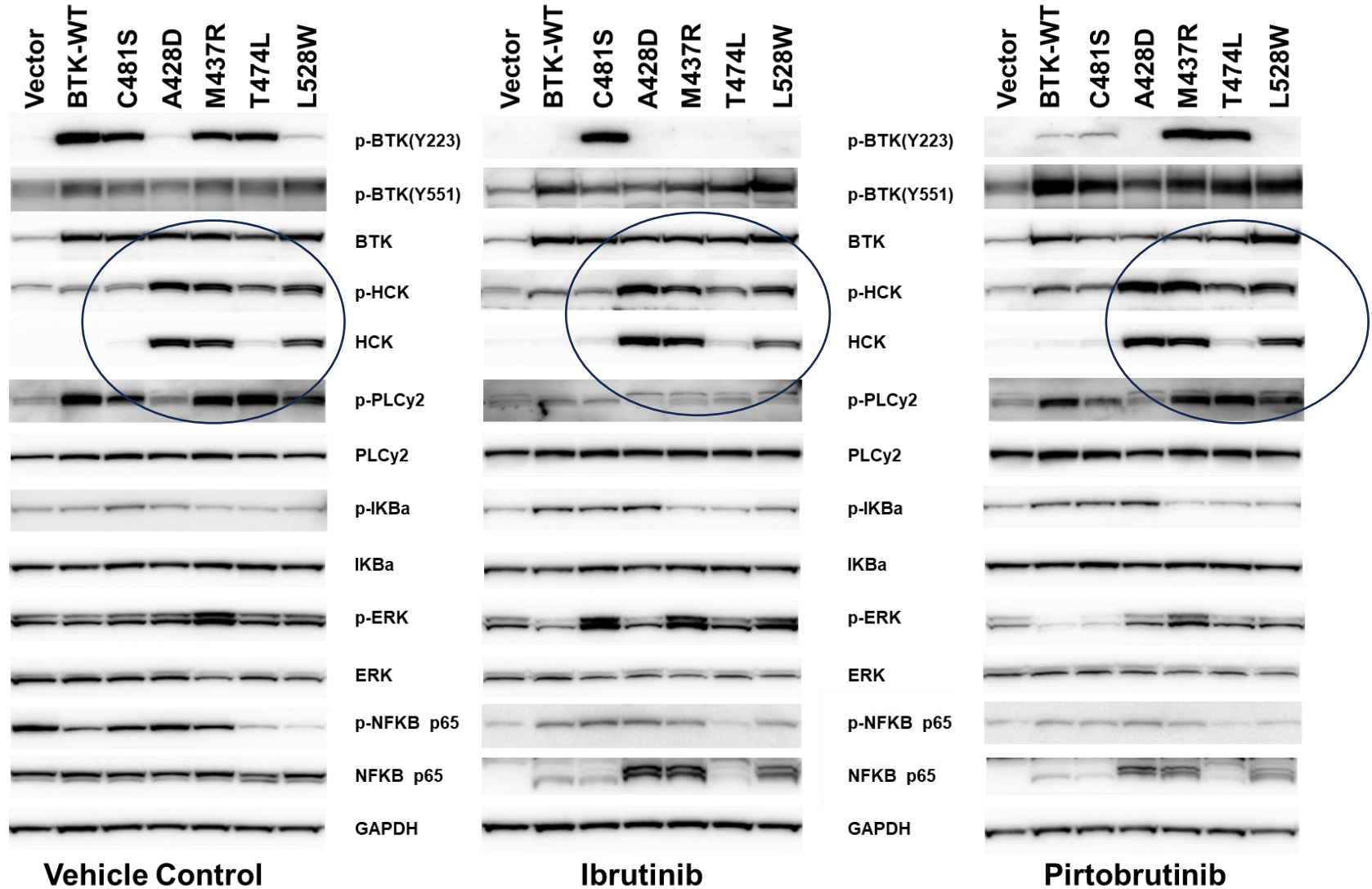
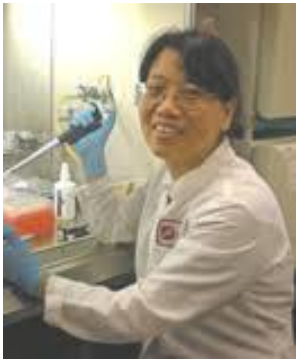
Kamaldeep Dhama¹, Anirban Chakraborty¹, Tarikere L. Gururaja¹, Leo W.-K. Cheung^{1,2},
 Chaohong Sun², Felix DeAnda¹, XiaoDong Huang^{1*}



Acquired BTK mutations related to BTK-inhibitor resistance for agents used to treat WM

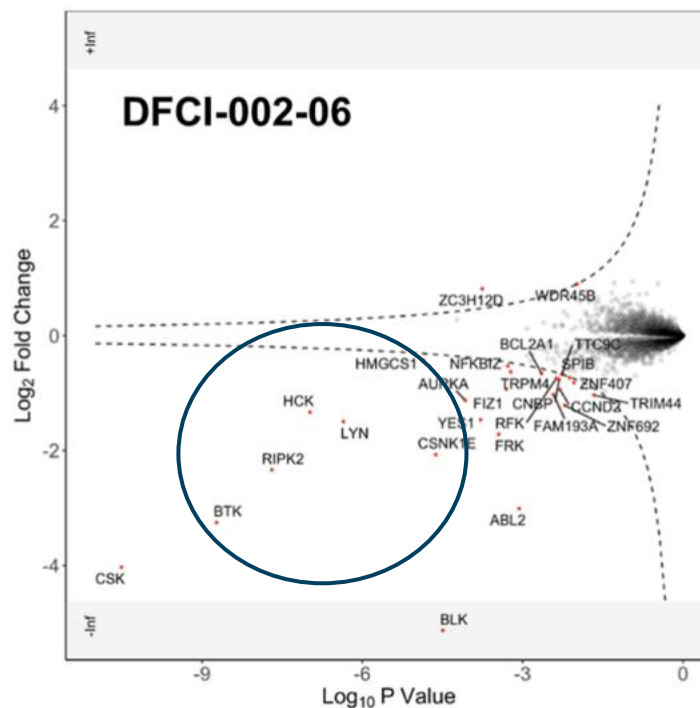
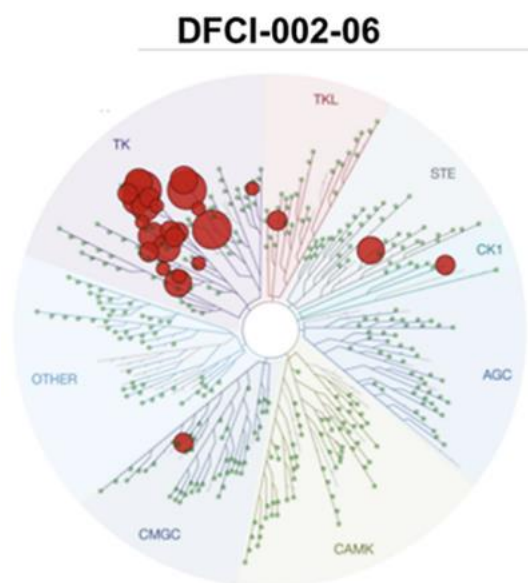
| BTK Mutation | BTK-Inhibitor Class | | |
|--------------------------|---------------------|--------------|--------|
| | Covalent | Non-Covalent | PROTAC |
| BTK ^{Cys481Ser} | ● | | |
| BTK ^{Cys481Arg} | ● | | |
| BTK ^{Cys481Tyr} | ● | | |
| BTK ^{Val416Leu} | | ● | |
| BTK ^{Ala428Asp} | ● | ● | ● |
| BTK ^{Thr474Leu} | | ● | |
| BTK ^{Leu528Trp} | ● | ● | |
| BTK ^{Met437Arg} | | ● | |

HCK is activated in BCWM.1 WM Cells expressing BTK mutations associated with BTK non-covalent kinase inhibitors and PROTACS

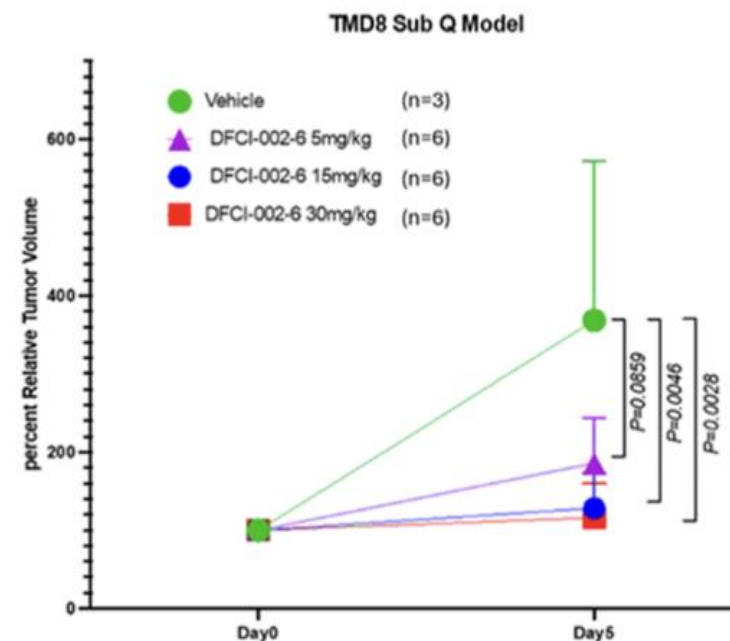
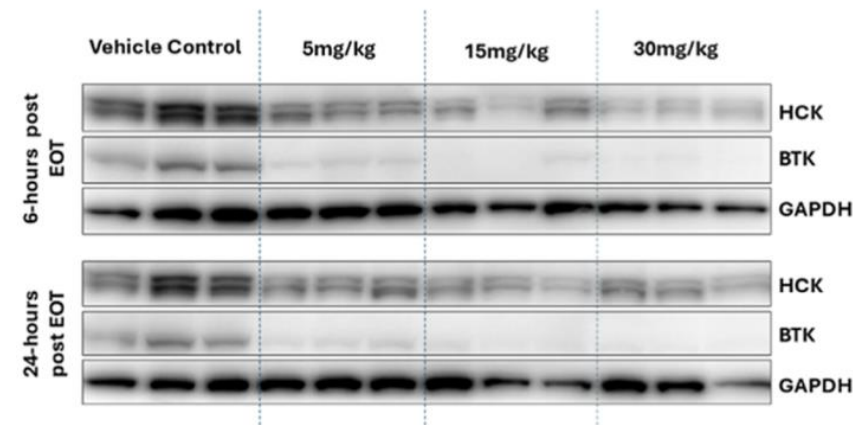


Discovery and Characterization of a First-In-Class HCK/BTK PROTAC DFCI-002-06 for the Treatment of MYD88 Mutated B Cell Malignancies

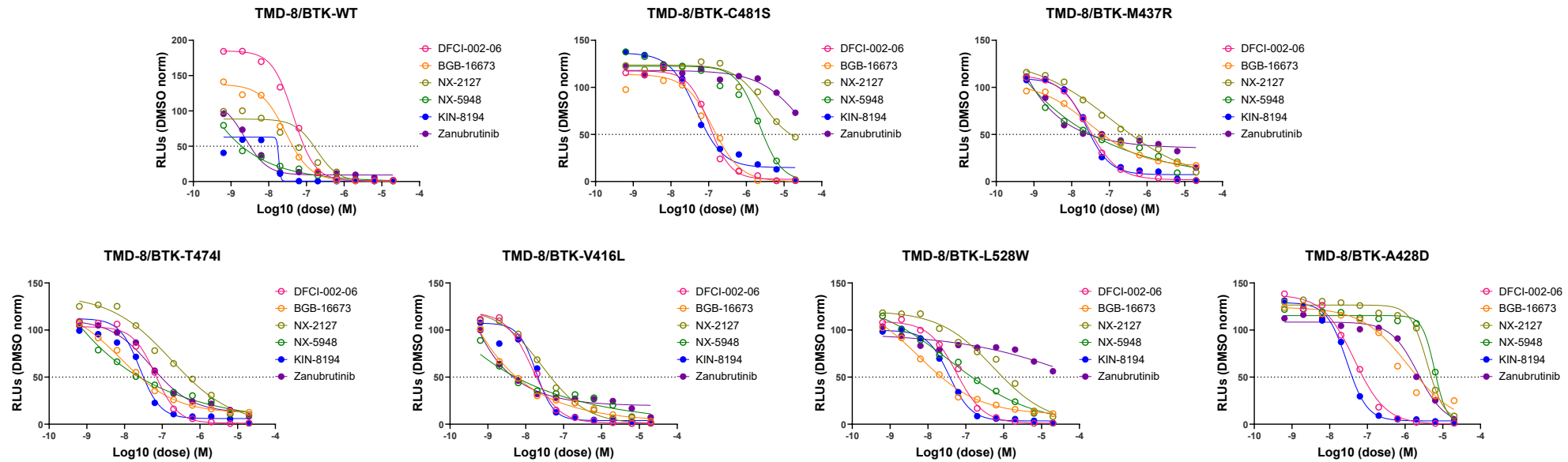
John M. Hatcher,¹ Shirong Liu,¹ Amanda Kofides, Alexa Canning, Dominic Pizzarella, Xia Liu, Nickolas Tsakmaklis, Maria Guerrero, Christopher Patterson, Alberto Guijosa, Prafulla Gokhale, Zachary Hunter, Shayna Sarosiek, Jorge Castillo, Jinhua Wang, Sara J. Buhrlage, and Steven P. Treon*



TMD8 Xenograft Model



Dose-Response Curves for BTK and HCK Targeting Compounds in MYD88 mutated TMD-8 cells expressing mutated BTK



| | BTK-WT | BTK-C481S | BTK-V416L | BTK-A428D | BTK-M437R | BTK-T474I | BTK-L528W |
|---------------------|----------|-----------|-----------|-----------|-----------|-----------|-----------|
| DFCI-002-06 | 4.5E-08 | 1.05E-07 | 1.45E-08 | 4.49E-08 | 2.76E-08 | 6.59E-08 | 5.57E-08 |
| BGB-16673 | 2.82E-08 | 1.3E-07 | 6.29E-25 | 3.08E-07 | 3.05E-08 | 8.19E-09 | 5.6E-09 |
| NX-2127 | 1.87E-07 | 2.82E-06 | 2.7E-08 | 3.84E-06 | 1.14E-07 | 1.72E-07 | 5.39E-07 |
| NX-5948 | 2E-15 | 2.17E-06 | 3.24E-33 | 6.07E-06 | 2.76E-30 | 1.9E-26 | 3.82E-08 |
| KIN-8194 | 1.85E-08 | 4.66E-08 | 1.87E-08 | 2.86E-08 | 2.44E-08 | 2.56E-08 | 3.64E-08 |
| Zanubrutinib | 2.34E-09 | 3.93E-05 | 2.24E-20 | 2.18E-06 | - | 5.09E-08 | 0.000149 |



A 3D scientific illustration of biological structures. On the left, a large, complex, light blue structure resembling a DNA molecule or a protein complex is shown. To its right and in the foreground, several translucent, spherical cells are depicted, some containing smaller, brownish-yellow spherical structures. The background is a dark blue gradient with small white dots, suggesting a microscopic or molecular environment.

**How do we manage
TP53ALT WMM disease?**

Phase II Study of the CD19-ADC Loncastuximab in Previously Treated WM

Shayna Sarosiek, PI. NCT05190705



| | N (range or %) |
|---------------------------|-----------------|
| Median age, y | 71 (53-78) |
| Male sex | 9 (64%) |
| Median previous therapies | 4 (2-10) |
| Median serum IgM, mg/dL | 2146 (723-5955) |
| Median hemoglobin, g/dL | 9.5 (8.5-12.6) |
| Median BM involvement, % | 53 (4-90) |
| MYD88 MUT | 12 (86%) |
| CXCR4 MUT (n=13) | 8 (62%) |
| TP53 MUT | 8 (54%) |
| LN ≥1.5 cm or EMD | 3 (21%) |
| Splenomegaly ≥15 cm | 1 (7%) |

| | MYD88 ^{Mut} | CXCR4 ^{Mut} | TP53 ^{ALT} | RESPONSE |
|-------|----------------------|----------------------|---------------------|----------|
| WM-1 | YES | YES | NO | VGPR |
| WM-2 | YES | YES | NO | PD |
| WM-3 | YES | NO | NO | PR |
| WM-4 | YES | NO | NO | VGPR |
| WM-5 | NO | NO | NO | VGPR |
| WM-6 | NO | NO | NO | MR |
| WM-7 | YES | YES | YES | CR |
| WM-8 | YES | YES | YES | VGPR |
| WM-9 | YES | YES | YES | VGPR |
| WM-10 | YES | NO | YES | VGPR |
| WM-11 | YES | YES | YES | VGPR |
| WM-12 | YES | YES | YES | VGPR |
| WM-13 | YES | UNK | YES | VGPR |
| WM-14 | YES | YES | YES | PR |

Prospective Cellular Therapies Trials for WM

| Agent | Target | Investigator | Response Activity | Trial Status |
|-------------|--------|--------------------------------------|----------------------------|---|
| Epcoritimab | CD20 | Von Keudell (Harvard) | NA | In progress |
| Epcoritamab | CD20 | Vos (Amsterdam) | NA | In progress |
| Surovatamig | CD19 | Sarosiek Mouhaddiene (Harvard) | NA | In progress |
| MB106 CAR T | CD20 | Shadman (Fred Hutch) | MRR 67% CR 33% (N=6) | Enrollment stopped for lack of sponsor funding |

13th International Workshop on WM

October 13-17, 2026

Ranjana Advani, Steve Ansell, Prashant Kapoor, Steve Treon
Co-Organizers

PALM SPRINGS

Palm Springs, California

<http://waldenstromsworkshop.org>