

UPDATED RESULTS OF THE ASPEN TRIAL FROM A COHORT OF PATIENTS WITH MYD88 WILD-TYPE WALDENSTRÖM MACROGLOBULINEMIA

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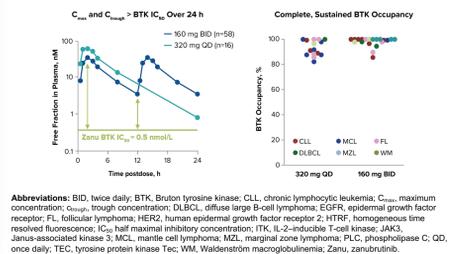
INTRODUCTION

- Bruton tyrosine kinase (BTK) plays a critical role in B-cell receptor signaling, which mediates B-cell proliferation, migration, and adhesion¹⁻³
- First-generation BTK inhibitor ibrutinib has shown activity in Waldenström macroglobulinemia (WM) and has become a standard of care⁴
 - However, lower response rates,⁵ no major responses,^{5,6} and shorter survival⁷ have been reported in patients who lack *MYD88*^{265P} or other activating mutations (*MYD88*^{WT})
- Zanubrutinib is a next-generation BTK inhibitor designed to maximize BTK occupancy and minimize off-target inhibition of TEC- and EGFR-family kinases (Figure 1)
 - Potent, selective, and irreversible⁸
 - Equipotent against BTK compared with ibrutinib; higher selectivity versus EGFR, ITK, JAK3, HER2, and TEC⁹
 - Advantageous pharmacokinetic/pharmacodynamic properties: complete and sustained BTK occupancy in peripheral blood mononuclear cells and lymph nodes⁸
 - Favorable drug-drug interaction properties: can be coadministered with strong/moderate CYP3A inhibitors at a reduced dose, proton pump inhibitors, acid-reducing agents, and anti-thrombotic agents^{10,11}

Figure 1a. Zanubrutinib: A Potent and Selective BTK Inhibitor^{8,9}

| ON TARGET | Targets | Assays | Zanubrutinib IC ₅₀ (nM) | Ibrutinib IC ₅₀ (nM) | Ratio (Zanubrutinib/Ibrutinib) |
|-----------|--------------------------------|--------|------------------------------------|---------------------------------|--------------------------------|
| BTK | BTK-pT232 Cellular Assay | 1.8 | 3.5 | 0.5 | |
| | Rec-1 Proliferation | 0.36 | 0.34 | 1.1 | |
| | BTK Occupation Cellular Assay | 2.2 | 2.3 | 1 | |
| | BTK Biochemical Assay | 0.22 | 0.2 | 1.1 | |
| EGFR | p-EGFR HTRF Cellular Assay | 606 | 101 | 6 | |
| | A43 Proliferation | 3210 | 323 | 9.9 | |
| | ITK Occupation Cellular Assay | 3265 | 189 | 17 | |
| | p-PLCγ1 Cellular Assay | 3433 | 77 | 45 | |
| ITK | IL-2 Production Cellular Assay | 2536 | 260 | 9.8 | |
| | ITK Biochemical Assay | 30 | 0.9 | 33 | |
| | JAK3 Biochemical Assay | 200 | 3.9 | 51 | |
| | HER2 Biochemical Assay | 661 | 9.4 | 70 | |
| TEC | TEC Biochemical Assay | 1.9 | 0.8 | 2.4 | |

Figure 1b. Complete, Sustained BTK Occupancy With BID or QD Dosing^{8,9}



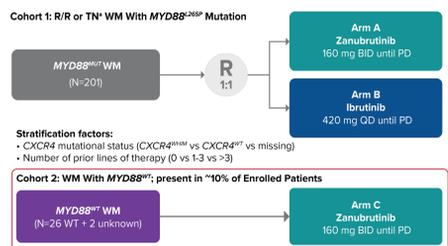
OBJECTIVES

- To assess the safety and efficacy of zanubrutinib in WM patients with *MYD88*^{WT} from an exploratory cohort of the ongoing phase 3 study of zanubrutinib versus ibrutinib in patients with WM (ASPEN; NCT03053440)

METHODS

- ASPEN is an open-label, multicenter, randomized, phase 3 study of zanubrutinib versus ibrutinib in patients with WM (Figure 2)

Figure 2. Phase 3 ASPEN Trial Design



EUDRACT 2016-002980-33; NCT03053440. *TN must be unsuitable for standard chemioimmunotherapy. Abbreviations: BID, twice daily; CXCR4, C-X-C motif chemokine receptor 4; MYD88, myeloid differentiation primary response gene 88; PD, progressive disease; QD, once daily; R/R, relapsed/refractory; TN, treatment-naïve; WM, Waldenström macroglobulinemia; WT, wild-type.

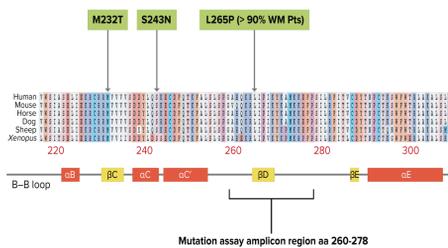
Eligibility

- Clinical and definitive histologic diagnosis of WM, with measurable disease (serum IgM >0.5 g/dL), and meeting ≥1 criterion for treatment according to consensus panel criteria from the Seventh International Workshop on WM¹²
- If treatment naïve, must be considered by treating physician unsuitable for standard chemioimmunotherapy regimens
- Eastern Cooperative Oncology Group performance status 0-2
- Absolute neutrophil count ≥750/μL, platelets ≥50000/μL (independent of growth factor/transfusions)
- Adequate renal, hepatic, and coagulation function
- No significant cardiac disease, active central nervous system involvement, or prior BTK inhibitors

Cohort Assignment

- Bone marrow *MYD88* and *CXCR4* mutations were assessed centrally at study entry (NeoGenomics Laboratory, Aliso Viejo, CA, USA)¹³
 - The *MYD88* mutation assay used detects all mutations in the region encompassing amino acid Ala²⁶⁰-Pro²⁷⁸, which includes the predominant mutation in WM, *MYD88*^{265P} (Figure 3)
- Patients were assigned to Cohort 1 (*MYD88* mutated; randomized) or exploratory Cohort 2 (*MYD88*^{WT} or *MYD88* unknown, nonrandomized) based on the central laboratory *MYD88* mutation assay results

Figure 3. *MYD88*-Activating Mutations in Patients With WM



Adapted from Leon et al.¹⁴ Abbreviations: LOD, limit of detection; MYD88, myeloid differentiation primary response gene 88; WM, Waldenström macroglobulinemia.

- Detection in the *MYD88* amplicon (Ala²⁶⁰-Pro²⁷⁸) by the NeoGenomics LDT assay includes a wild-type-allele-blocking approach (limit of detection [LOD], 0.5%)¹³ versus standard polymerase chain reaction/bidirectional Sanger sequencing assay used to detect *CXCR4* mutations (LOD, 10%-15%)
- For *MYD88*^{WT} patients with available samples (12 of 26), *MYD88* mutations were also evaluated by next-generation sequencing (200x; LOD, 5%); no other activating mutations were detected

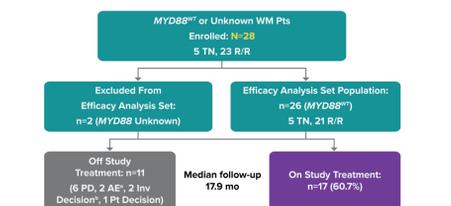
Exploratory Endpoints for Cohort 2

- Responses were assessed monthly by immunoglobulin M (IgM) with extramedullary disease assessment every 3 months, according to response criteria in the National Comprehensive Cancer Network WM guidelines¹⁵ and modified Owen criteria¹⁶ as assessed by the independent review committee
- Efficacy: response rates (overall and major response rate), duration of response, progression-free survival, and overall survival; safety assessed according to National Cancer Institute Common Terminology Criteria for Adverse Events version 4.03

RESULTS

- In total, 28 patients (n=26 *MYD88*^{WT}; n=2 *MYD88* mutation status unknown) were enrolled into Cohort 2
- The safety analysis set includes all 28 patients, and the efficacy analysis set includes 26 *MYD88*^{WT} patients, with a median follow-up of 17.9 months (range, 2.3-27.8; Figure 4 and Table 1)

Figure 4. Disposition of Patients in Cohort 2



Data cutoff date: 31 August 2019. *Grade 4 subdural hemorrhage; grade 3 diarrhea; †Investigator decided no further treatment needed (n=1); ‡Discharged to hospice for palliative care (n=1). Abbreviations: AE, adverse event; inv, investigator; MYD88, myeloid differentiation primary response gene 88; PD, progressive disease; Pt, patient; R/R, relapsed/refractory; TN, treatment-naïve; WT, wild-type.

Table 1. Patient and Disease Characteristics

| Characteristics, n (%) | Total (N=28) |
|--|--------------|
| Age median (range), y | 70.1 (39-87) |
| >65 y | 19 (67.9) |
| >75 y | 12 (42.9) |
| Sex, n (%) | |
| Male | 14 (50) |
| Female | 14 (50) |
| IPSSWM, n (%) | |
| Low | 5 (17.9) |
| Intermediate | 11 (39.3) |
| High | 12 (42.9) |
| Prior treatment status, n (%) | |
| R/R, n (%) | 23 (82.1) |
| No. of prior therapies for R/R patients, median (range) | 1 (1-5) |
| Extramedullary disease present at baseline by IRC, n (%) | 21 (75.0) |
| Genotype, n (%) | |
| <i>MYD88</i> ^{WT} / <i>CXCR4</i> ^{WT} | 23 (82.1) |
| <i>MYD88</i> ^{WT} / <i>CXCR4</i> ^{Mut} | 1 (3.6) |
| <i>MYD88</i> ^{Mut} / <i>CXCR4</i> ^{WT} | 2 (7.1) |
| <i>MYD88</i> unknown/ <i>CXCR4</i> unknown | 2 (7.1) |
| Hemoglobin ≤110 g/L, n (%) | 15 (53.6) |

Abbreviations: CXCR4, C-X-C motif chemokine receptor 4; IPSSWM, International Prognostic Scoring System Waldenström macroglobulinemia; IRC, independent review committee; MYD88, myeloid differentiation primary response gene 88; R/R, relapsed/refractory; WT, wild-type.

Safety (N=28)

Table 2. AE Overview

| Treatment-Emergent AE | n (%) |
|---|----------------------|
| Patients with ≥1 AE grade ≥3 | 18 (64.3) |
| Patients with ≥1 serious AE | 11 (39.3) |
| AE leading to death | 0 |
| AE leading to treatment discontinuation | 2 ^a (7.1) |
| AE leading to dose reduction | 2 ^b (7.1) |

^aGrade 4 subdural hemorrhage (related) and grade 3 diarrhea (related). ^bGrade 3 pneumonitis resolved and followed by grade 2 pneumonia (n=1); grade 1 diarrhea (n=1). Abbreviation: AE, adverse event.

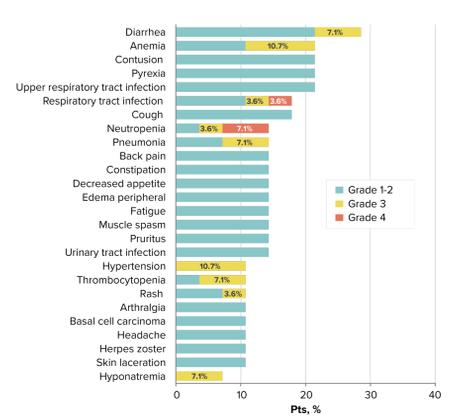
- No treatment-emergent adverse events (AEs) leading to death (Table 2)
- Two patients discontinued because of AEs
 - Grade 4 subdural hemorrhage
 - Grade 3 diarrhea
- Major hemorrhage occurred in two patients (Table 3)
 - Gastric ulcer hemorrhage
 - Periorbital hematoma, subdural hematoma, and subdural hemorrhage; treatment was permanently discontinued per protocol
- Atrial fibrillation/flutter occurred in one patient (grade 1)
- Most common AEs (in >15% patients) were diarrhea, anemia, contusion, pyrexia, upper respiratory tract infection, respiratory tract infection, and cough (Figure 5)

Table 3. AE Categories of Interest (BTKI Class AEs)

| AE Categories (Pooled Terms), n (%) | All Grade | Grade ≥3 |
|-------------------------------------|-----------|----------|
| Atrial fibrillation/flutter | 1 (3.6) | 0 |
| Diarrhea (PT) | 8 (28.6) | 2 (7.1) |
| Hemorrhage | 11 (39.3) | 2 (7.1) |
| Major hemorrhage ^a | 2 (7.1) | 2 (7.1) |
| Hypertension | 3 (10.7) | 3 (10.7) |
| Neutropenia ^b | 5 (17.9) | 3 (10.7) |
| Infection | 21 (75.0) | 8 (28.6) |
| Second malignancy ^c | 4 (14.3) | 0 |

No tumor lysis syndrome or opportunistic infection was reported. ^aDefined as any grade ≥3 hemorrhage or any grade central nervous system hemorrhage, gastric ulcer hemorrhage, and 1 patient had periorbital hematoma, subdural hematoma, and subdural hemorrhage. ^bIncluding PT terms of neutropenia, neutrophil count decreased, febrile neutropenia, agranulocytosis, neutropenic infection, and neutropenic sepsis. ^cBasal cell carcinoma (n=3) and Queyret erythroplasia (n=1). Abbreviations: AE, adverse event; BTKI, Bruton tyrosine kinase inhibitor; PT, preferred term.

Figure 5. Common AEs (Any Grade >10% or Grade ≥3 in >1 Pt), Regardless of Causality

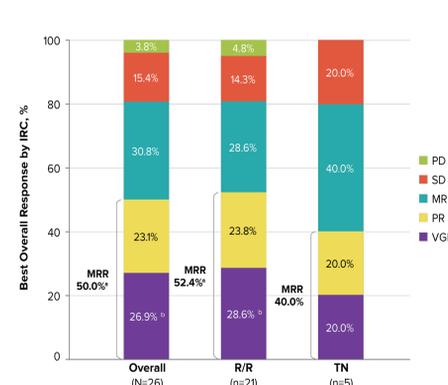


Abbreviations: AE, adverse event; Pt, patient.

Efficacy (n=26)

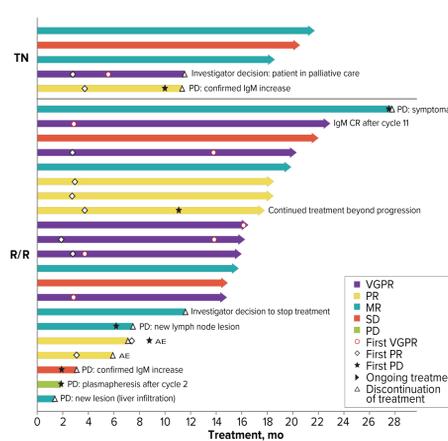
- Major response rate of 50.0% including 26.9% with very good partial response (Figure 6)
- Median time to first major response (partial response or better, requiring reduction in extramedullary disease if present at baseline) was 2.9 mo (range, 1.9-16.1; Figure 7)
- IgM complete response (requiring normal IgM and immunofixation negative) was achieved in one patient
- Median progression-free and overall survival were not yet reached (Figure 8)

Figure 6. Best Responses by IRC in Patients With *MYD88*^{WT} WM



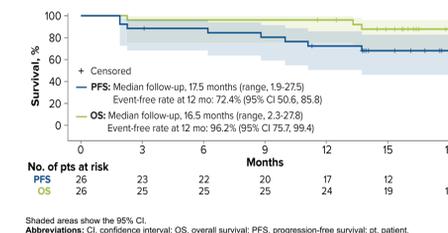
including patients confirmed by next-generation sequencing of no other activating *MYD88* mutations. ^aOne patient achieved IgM complete response (normalized IgM and negative immunofixation since cycle 11, with bulky extramedullary disease improving). Abbreviations: CR, complete response; IgM, immunoglobulin M; IRC, independent review committee; MR, minor response; MRR, major response rate (MRR); MYD88, myeloid differentiation primary response gene 88; PD, progressive disease; PR, partial response; R/R, relapsed/refractory; SD, stable disease; TN, treatment-naïve; VGPR, very good PR; WM, Waldenström macroglobulinemia; WT, wild-type.

Figure 7. Responses Over Time on Treatment



Note: color of bars represents the best response for each patient. Abbreviations: AE, adverse event; IgM, immunoglobulin M; MR, minor response; PD, progressive disease; PR, partial response; R/R, relapsed/refractory; SD, stable disease; TN, treatment-naïve; VGPR, very good PR.

Figure 8. Progression-Free and Overall Survival



CONCLUSIONS

- Largest cohort of patients with WM with confirmed *MYD88*^{WT} (n=26) studied in terms of safety and efficacy of BTK inhibitor treatment
- Single-agent zanubrutinib resulted in major responses (including very good partial response)
 - Major response rate of 50.0% including 26.9% with very good partial response
 - Median time to first major response was 2.9 months (range, 1.9-16.1)
- Zanubrutinib was well tolerated
 - Primary reason for discontinuation was progressive disease (3 of 6 within first 3 cycles)
 - Discontinuation because of AEs occurred in 7.1% of patients (2/28)
 - No fatal AEs reported
 - Low incidences of atrial fibrillation
 - AE profile is consistent with Cohort 1 finding in the ASPEN study

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