# Switching from Covalent BTKi to BCL2i is Associated with Improved Clinical **Outcomes Compared to Switching to a** Different Covalent BTKi in Patients with CLL/SLL Treated in the Real-world Setting

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# OBJECTIVE

To assess real-world treatment outcomes of patients with CLL/SLL who switched to a cBTKi- or a BCL2i-based regimen following discontinuation of a cBTKi-based regimen

# CONCLUSIONS

Patients who switched from a cBTKi to a BCL2i-based regimen had significantly greater odds of responding to therapy, a significantly lower hazard of progression/death, and a trending lower hazard of starting next treatment /death compared to those who switched to cBTKi-based regimens

This study provides real-world evidence supporting the effectiveness of BCL2i-based regimens in improving treatment outcomes post cBTKi-based regimens compared to sequencing through cBTKis

In light of the dynamic CLL treatment landscape, it is vital to consider the impact of switching to agent(s) with a different MOA versus retreating with agent(s) in the same class as a means of optimizing clinical care

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### INTRODUCTION

• Covalent Bruton tyrosine kinase (cBTKi) and B-cell lymphoma 2 (BCL2i) inhibitors have demonstrated improved treatment efficacy and safety compared to chemotherapy/chemoimmunotherapy (CT/CIT) for patients with chronic lymphocytic leukemia and small lymphocytic lymphoma (CLL/SLL)<sup>1-3</sup>

• Although some patients progress and discontinue their cBTKi-based regimens prematurely (12-17%), a substantial proportion (24-62%) discontinue due to reasons other than disease progression (PD) and often switch to another targeted agent with the same or different mechanism of action (MOA)<sup>4-9</sup>

## RESULTS

#### **Patient characteristics**

• The study included 121 BCL2i-naïve patients who discontinued their first cBTKi-based regimen; of them, 44 patients (36.4%) switched to a cBTKi- (cBTKi cohort) and 77 patients (63.6%) to a BCL2i-based regimen (BCL2i cohort)

 Patients in both cohorts had similar characteristics, such as median age at initiation (cBTKi: 71.5 years, BCL2i: 69.0 years), male sex (cBTKi: 63.6%, BCL2i: 68.8%), unmutated IGHV (cBTKi: 47.7%, BCL2i: 37.7%), and with del(17p)/TP53 mutation (cBTKi: 20.5%) BCL2i: 27.3%) (Table 1)

 However, the cBTKi cohort had a greater proportion of patients with ECOG Performance Status 0 (cBTKi: 50.0%, BCL2i: 27.3%) and a lower proportion with elevated LDH (cBTKi: 29.5%, BCL2i: 46.8%)

#### Table 1. Demographic characteristics for the cBTKi and **BCL2i cohorts**

| Patient characteristics  | cBTKi<br>N = 44      | BCL2i<br>N = 77      | p-value |
|--|----------------------|----------------------|---------|
| Age at cBTKi or BCL2i initiation<br>(years), Mean± SD [Median] | 70.2 ± 9.2<br>[71.5] | 68.7 ± 9.6<br>[69.0] | 0.41    |
| Male sex, N (%)  | 28 (63.6)            | 53 (68.8)            | 0.56    |
| Race, N (%)  |                      |                      | 0.98    |
| White  | 36 (81.8)            | 63 (81.8)            |         |
| Other  | 6 (13.6)             | 10 (13.0)            |         |
| Comorbidities  |                      |                      |         |
| Cardiovascular condition                                       | 23 (52.3)            | 38 (49.4)            | 0.76    |
| Endocrine/metabolic condition                                  | 14 (31.8)            | 19 (24.7)            | 0.40    |
| Musculoskeletal condition                                      | 9 (20.5)             | 7 (9.1)              | 0.08    |
| Renal condition  | 6 (13.6)             | 16 (20.8)            | 0.33    |
| Neurologic condition   | 6 (13.6)             | 9 (11.7)             | 0.75    |
| Respiratory condition  | 5 (11.4)             | 10 (13.0)            | 0.79    |
| Rai staging, N (%)   |                      |                      | 0.98    |
| Stage 0  | 5 (11.4)             | 9 (11.7)             |         |
| Stage I - IV   | 28 (63.6)            | 50 (64.9)            |         |
| ECOG, N (%)  |                      |                      | 0.03*   |
| Grade 0  | 22 (50.0)            | 21 (27.3)            |         |
| Grade 1 - 3  | 14 (31.8)            | 42 (54.5)            |         |
| Jnmutated IGHV, N (%)  | 21 (47.7)            | 29 (37.7)            | 0.38    |
| Elevated LDH, N (%)  | 13 (29.5)            | 36 (46.8)            | 0.04*   |
| Nith del(17p)/TP53 mutation                                    | 9 (20.5)             | 21 (27.3)            | 0.59    |

Group; IGHV, immunoglobulin heavy-chain variable region gene; LDH, lactate dehydrogenase; SD, standard deviation \*Significant at p<0.05

#### **cBTKi** discontinuation and subsequent treatments

 Patients in the cBTKi cohort were treated longer with their first cBTKibased regimen before discontinuation and had longer time from CLL/SLL diagnosis to cBTKi-based regimen initiation than those in the BCL2i cohort (**Table 2**)

 The most common reason for cBTKi-based regimen discontinuation was intolerance (80.2%; n=97)

### **METHODS**

- observational study involving 23 centers

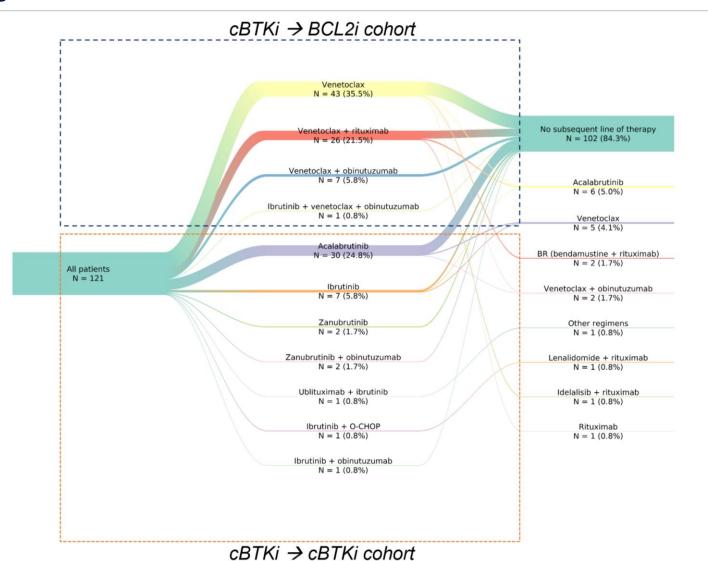
### Table 2. Treatment characteristics for the cBTKi and BCL2i cohorts

| Treatment characteristics   | cBTKi<br>N = 44  | BCL2i<br>N = 77           | p-value |
|---|--|---------------------------|---------|
| Time from CLL/SLL diagnosis to cBTKi<br>or BCL2i initiation (months), Mean± SD<br>[Median]  | $\begin{array}{c} 98.4 \pm 54.6 \\ [91.0] \end{array}$ | $76.4 \pm 66.7 \\ [62.6]$ | 0.07    |
| Time from first cBTKi discontinuation to cBTKi/BCL2i initiation (months), Mean± SD [Median] | 7.3 ± 11.0<br>[0.51]                                   | 7.5 ± 11.7<br>[1.64]      | 0.94    |
| Time to first cBTKi discontinuation<br>(months), Mean± SD [Median]                          | 22.8 ± 19.8<br>[19.6]                                  | 14.3 ± 17.1<br>[10.8]     | 0.01*   |
| Duration of follow-up (months), Mean±<br>SD [Median]  | 16.1 ± 11.7<br>[11.6]                                  | 20.1 ± 14.2<br>[16.4]     | 0.13    |
| Reasons for cBTKi discontinuation   |  |                           |         |
| Intolerance   | 35 (79.5%)   | 62 (80.5%)                | -       |
| Patient request   | 2 (4.5%)   | 3 (3.9%)                  | -       |
| Watchful waiting due to low or no disease activity  | 1 (2.3%)   | 5 (6.5%)                  | -       |
| Disease transformation (diffuse large B-<br>Cell or Hodgkin lymphoma)                       | 1 (2.3%)   | 0 (0.0%)                  | -       |
| Economic reasons  | 0 (0.0%)   | 2 (2.6%)                  | _       |
| Other   | 7 (15.9%)  | 7 (9.1%)                  | -       |
| Number of prior lines, Mean± SD<br>[Median]   | 1.7 ± 1.0<br>[1.0]                                     | 1.7 ± 0.8<br>[2.0]        | 0.98    |
| Number of prior lines, N (%)  | _ <u>-</u>   |                           | 0.44    |
| 1   | 25 (56.8)  | 37 (48.1)                 |         |
| 2   | 11 (25.0)  | 28 (36.4)                 |         |
| 3+  | 8 (18.2)   | 12 (15.6)                 |         |

ard deviation, SLL, small lymphocytic lymphoma Significant at p<0.05

• Most common treatments following the cBTKi-based regimen discontinuation included acalabrutinib (25.2%) and ibrutinib (5.9%) in the cBTKi cohort; and venetoclax (36.1%) and venetoclax+rituximab (21.8%) in the BCL2i cohort (Figure 1)

#### Figure 1. Treatments following cBTKi-based regimen discontinuation<sup>1-3</sup>



BCL2i, B-cell lymphoma 2 inhibitor; cBTKi, covalent Bruton tyrosine kinase inhibitor; <sup>1</sup>The most common first cBTKi-based egimens received by patients were ibrutinib-based (118/121 [97.5%]) and acalabrutinib-based (3/121 [2.5%]) regimens; <sup>2</sup>Among atients who did not initiate a subsequent line of therapy, a greater proportion were from the BCL2i cohort (67/102 [65.7%]) elative to the cBTKi cohort (35/102 [34.3%]); <sup>3</sup>One patient who was treated with both a cBTKi and a BCL2i as part of the same egimen was included in the BCL2i cohort

 Data were obtained as part of the CLL Collaborative Study of Real-World Evidence (CORE), an international, retrospective

 Adult patients were included if they were diagnosed with CLL/SLL, were naïve to BCL2i, discontinued their first treatment with a cBTKi-based regimen in any line of therapy for reasons other than PD or therapy completion, and initiated a cBTKi or BCL2i-based regimen after their first

cBTKi-based regimen discontinuation. Then, 2 mutually exclusive cohorts were derived based on the treatments these patients received: cBTKi cohort (i.e., cBTKi→cBTKi) or BCL2i cohort (i.e., cBTKi→BCL2i)

 Treatment outcomes included overall response (OR; i.e., complete [CR] or partial response [PR]), progression-free survival (PFS; i.e., time from initiation to PD/death (event) with censoring at last follow-up), and time to next treatment or death (TTNT-D; i.e., time from initiation to start of next treatment/death (event), with censoring at last follow-up)

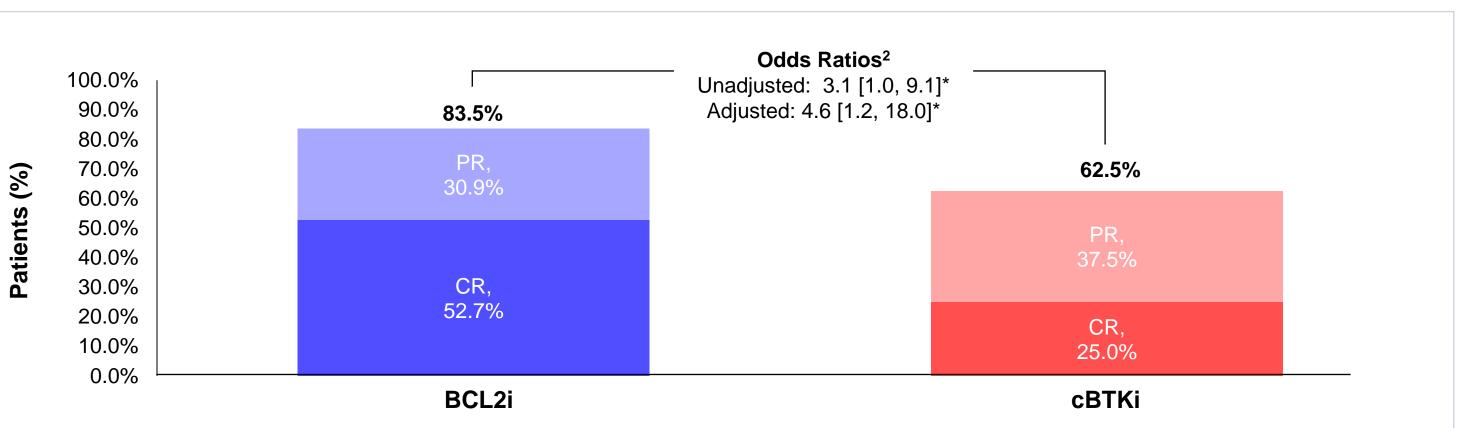
#### **Overall response (OR)**

• Of the 79 patients with recorded response, the BCL2i cohort had significantly greater odds of achieving a response compared to the cBTKi cohort (**Figure 2**)

#### Progression-free survival (PFS)

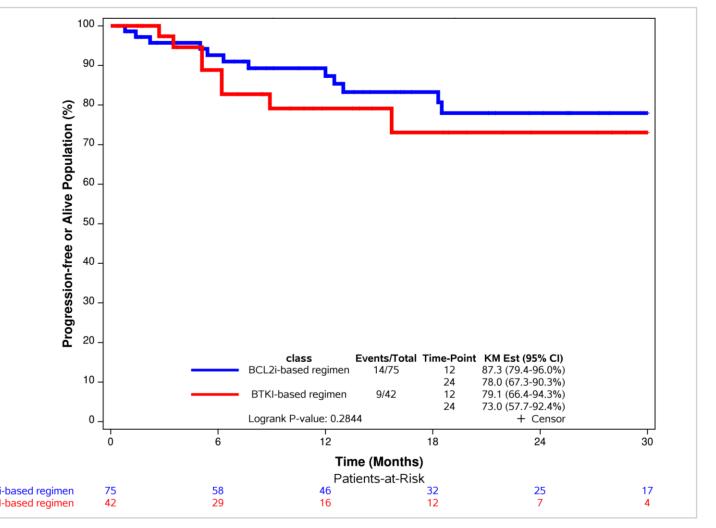
- The BCL2i cohort had higher PFS KM estimates than cBTKi cohort at 12 months (cBTKi: 79.1%, BCL2i: 87.3%) (Figure 3)
- The BCL2i cohort had a 70% lower hazard of progression/death

#### Figure 2. Overall response to therapy after the first cBTKi<sup>1</sup>



CR. complete remission: PD. progressive disease: PR. partial remission: SD. stable disease Responses were dichotomized into responder (CR [BCL2i: 52.7%; cBTKi: 25.0%] + PR [BCL2i: 30.9%; cBTKi: 37.5%]) and non-responder (SD [BCL2i: 12.7%; cBTKi: 25.0%] + PD [BCL2i: 3.6%; cBTKi: 12.5%] <sup>2</sup>Overall response was compared between cohorts using a logistic regression model, adjusting for age at initiation, sex, time from CLL/SLL diagnosis to initiation, duration of discontinued cBTKi therapy, ECOG, and LDH status Significant at p<0.05

#### Figure 3. PFS for cBTKi vs BTCL2i cohorts<sup>1-2</sup>



BCL2i, B-cell lymphoma 2 inhibitor; cBTKi, covalent Bruton tyrosine kinase inhibitor; KM, Kaplan Meier; PFS, progression free survival; RMST, restricted mean survival time <sup>1</sup>The estimated RMST at Tau = 39.7 months was 33.1 months (standard error = 2.7) for the BCL2i cohort and 31.1 months (standard error = 1.7) for the cBTKi cohort. <sup>2</sup>Curves are limited to the time point with at least 10% of the starting population at-risk in both groups.

#### LIMITATIONS

- As this was a retrospective chart review across multiple centers, there may be differences in clinical data interpretation and possibility of data entry errors
- Findings should be interpreted with care given the limited durations of follow-up, especially in the cBTKi cohort

#### Time to next treatment or death (TTNT-D)

• Relative to cBTKi, the BCL2i cohort had higher, but not statistically significant, TTNT-D KM estimates (12 months: cBTKi: 81.1%, BCL2i: 84.7%), and lower hazards of time to next treatment/death (unadjusted HR: 0.8 [CI: 0.4; 1.9]; adjusted HR: 0.4 [CI: 0.1; 1.0], p=0.055) (Figure 4)

3CL2i-based regime

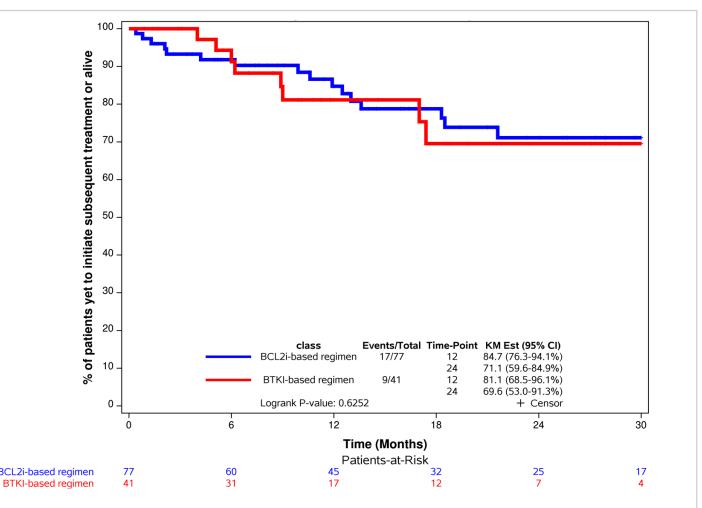
• Logistic regression models assessed the odds of achieving an OR; Cox proportional hazards (PH) regressions assessed PFS and TTNT-D

- All models controlled for age at initiation, sex, time from CLL/SLL diagnosis to initiation, duration of discontinued cBTKi therapy, ECOG, and lactate dehydrogenase (LDH)

 Kaplan-Meier (KM) estimates were also used to estimate PFS and TTNT-D rates at 12 months

(unadjusted hazard ratio [HR]: 0.6 [confidence interval [CI]: 0.3; 1.5]; adjusted HR: 0.3 [CI: 0.1; 0.8], p=0.023) compared to cBTKi cohort in the Cox PH regression

#### Figure 4. TTNT-D for cBTKi vs BCL2i cohorts<sup>1-2</sup>



BCL2i, B-cell lymphoma 2 inhibitor; cBTKi, covalent Bruton tyrosine kinase inhibitor; KM, Kaplan Meier; RMST, restricted mean survival time; TTNT-D, time to next treatment or death The estimated RMST at Tau = 43.3 months was 33.6 months (standard error = 2.0) for the BCL2i cohort and 33.4 months (standard error = 3.0) for the cBTKi cohort. <sup>2</sup>Curves are limited to the time point with at least 10% of the starting population at-risk in both groups.

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