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# Obinutuzumab and idelalisib in symptomatic patients with relapsed/refractory Waldenström macroglobulinemia

Cécile Tomowiak,<sup>1,\*</sup> Stéphanie Poulain,<sup>2,\*</sup> Charles Herbaux,<sup>3</sup> Aurore Perrot,<sup>4</sup> Béatrice Mahé,<sup>5</sup> Pierre Morel,<sup>6</sup> Thérèse Aurrant,<sup>7</sup> Olivier Tournilhac,<sup>8</sup> Stéphane Leprêtre,<sup>9</sup> Souad Assaad,<sup>10</sup> Bruno Villemagne,<sup>11</sup> Olivier Casasnovas,<sup>12</sup> Delphine Nollet,<sup>13</sup> Damien Roos-Weil,<sup>14</sup> Sylvie Chevret,<sup>15</sup> and Véronique Leblond,<sup>14</sup> on behalf of the FILO Group

<sup>1</sup>Hematology Department and Centre d'Investigations Cliniques (CIC) 1402 INSERM, University Hospital, Poitiers, France; <sup>2</sup>INSERM 1277–Centre National de la Recherche Scientifique 9020 Unite Mixte de Recherche 12, University of Lille, Laboratory of Hematology, Biology, and Pathology Center, Centre Hospitalier Universitaire de Lille, Lille, France; <sup>3</sup>Department of Blood Diseases, University Hospital, Lille, France; <sup>4</sup>Hematology Department, University Hospital, Institut Universitaire du Cancer–Oncopole, Université de Toulouse, Université Paris-Saclay, Toulouse, France; <sup>5</sup>Hematology Department, University Hospital, Nantes, France; <sup>6</sup>Hematology Department, University Hospital Amiens-Picardie, Amiens, France; <sup>7</sup>Department of Hematology, Institut Paoli Calmettes, Marseille, France; <sup>8</sup>Hematology Department, University Hospital Estaing, University Clermont Auvergne EA 7453, CIC1405, Clermont-Ferrand, France; <sup>9</sup>INSERM U1245 and Department of Hematology, Centre Henri Becquerel and Normandie University UNIROUEN, Rouen, France; <sup>10</sup>Centre Léon Bérard, Lyon, France; <sup>11</sup>Hematology Department, Centre Hospitalier Départemental, La Roche sur Yon, France; <sup>12</sup>Department of Hematology, University Hospital Dijon Bourgogne and INSERM 1231, Dijon, France; <sup>13</sup>Hematology Department, University Hospital Bretonneau, Tours, France; <sup>14</sup>Hematology Department, Sorbonne Université, Hôpital Pitié Salpêtrière APHP, Paris, France; and <sup>15</sup>U1153 Centre de Recherche en Épidémiologie et Statistiques, Equipe de Recherche en Biostatistiques et Épidémiologie Clinique

## Key Points

- Fixed-duration chemotherapy-free treatment (obinutuzumab + idelalisib) is effective in R/R WM, although toxicity is a limitation.

We present the results of a phase 2 study evaluating the combination of obinutuzumab + idelalisib in relapsed/refractory (R/R) Waldenström macroglobulinemia (WM). The goal was to determine the safety and efficacy of a fixed-duration chemotherapy-free treatment. During the induction phase, patients received idelalisib + obinutuzumab for 6 cycles, followed by a maintenance phase with idelalisib alone for  $\leq 2$  years. Forty-eight patients with R/R WM were treated with the induction combination, and 27 patients participated in the maintenance phase. The best responses, reached after a median of 6.5 months (interquartile range, 3.4–7.1; range, 2.6–22.1 months), were very good partial response in 5 patients, partial response in 27 patients, and minor response in 3 patients, leading to overall response rate and major response rate estimates of 71.4% (95% confidence interval [CI], 56.7–83.4) and 65.3% (95% CI, 50.4–78.3), respectively. With a median follow-up of 25.9 months, median progression-free survival was 25.4 months (95% CI, 15.7–29.0). Univariate analysis focusing on molecular screening found no significant impact of *CXCR4* genotypes on responses and survivals but a deleterious impact of *TP53* mutations on survival. Although there was no grade 5 toxicity, 26 patients were removed from the study because of side effects; the most frequent were neutropenia (9.4%), diarrhea (8.6%), and liver toxicity (9.3%). The combination of idelalisib + obinutuzumab is effective in R/R WM. Nonetheless, the apparent lack of impact of genotype on outcome could give new meaning to targeting of the phosphatidylinositol 3-kinase pathway in WM. This trial was registered at [www.clinicaltrials.gov](http://www.clinicaltrials.gov) as #NCT02962401.

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\*C.T. and S.P. are joint first authors and contributed equally to this work.

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Data sharing requests should be sent to Cécile Tomowiak ([ceciletomowiak@gmail.com](mailto:ceciletomowiak@gmail.com)).

The full-text version of this article contains a data supplement.

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## Introduction

Waldenström macroglobulinemia (WM) is characterized by the malignant accumulation of immunoglobulin M (IgM) that is caused by IgM-secreting lymphoplasmacytic lymphoma cells in bone marrow (BM).<sup>1</sup> Since the initial development of treatment recommendations at the Second International Workshop on Waldenström Macroglobulinemia,<sup>2</sup> treatment options for WM have evolved from immunotherapy to new compounds, such as Bruton tyrosine kinase (BTK) inhibitors, either alone or in combination with rituximab.<sup>3</sup> The B-cell receptor (BCR) has been identified as a potential therapeutic target in many mature B-cell malignancies with distinct and/or heterogeneous physiopathology or clinical course. BCR-associated kinases, such as BTK or phosphatidylinositol 3-kinase  $\delta$  (PI3K $\delta$ ), are critical signaling transducers of BCRs supporting the survival and growth of malignant B cells.<sup>4</sup>

Mutation of *MYD88*<sup>L265P</sup> was described by Treon and colleagues in 90% of WM patients; it is responsible for the proliferation and survival of WM cells by activating the NF- $\kappa$ B pathway via BTK.<sup>5,6</sup> As a result, targeting of BCR kinases is emerging as a new treatment paradigm for numerous B-cell malignancies. Ibrutinib, a selective BTK inhibitor, has shown high activity and very promising results in patients with WM. Most chemotherapy-free regimen data have been reported in relapsing or refractory (R/R) patients treated with BTK inhibitors, alone or in association with rituximab.<sup>7-10</sup> However, lower response rates and longer time to reach a response have been reported in *MYD88*<sup>WT</sup> and *CXCR4*<sup>MUT</sup> genotype patients treated with ibrutinib.<sup>7,11</sup>

Along with ibrutinib, idelalisib, an orally active selective PI3K $\delta$  inhibitor, represents a new class of agents targeting signal transduction downstream from BCR in malignant B cells. *MYD88*<sup>L265P</sup> also promotes activation of the PI3K pathway constitutively activated in WM.<sup>12</sup> In preclinical studies, idelalisib was shown to abrogate AKT activation, and it may target the BCR-controlled microenvironmental interaction of WM cells.<sup>13</sup> Idelalisib has manifested antitumor activity in patients with previously treated indolent non-Hodgkin lymphomas.<sup>14</sup> Compared with placebo + rituximab, the combination of idelalisib + rituximab appeared to shorten the period of lymphocytosis and improved progression-free survival (PFS), overall response rate (ORR), and overall survival (OS) in patients with relapsed chronic lymphocytic leukemia and clinically significant coexisting medical conditions, with an acceptable safety profile.<sup>15</sup> However, idelalisib has been associated with significant and limiting adverse effects (AEs), particularly hepatic and infectious effects.<sup>16</sup>

Obinutuzumab, a glycoengineered type II anti-CD20 monoclonal antibody (mAb), has been described as being more effective than rituximab in low-grade lymphoid malignancies, including in unfit treatment-naïve patients with chronic lymphocytic leukemia and in follicular lymphoma, resulting in longer PFS than with rituximab-based therapy.<sup>17,18</sup> Use of a chemotherapy-free treatment, anti-CD20 mAb in combination with oral idelalisib, should allow some patients with WM, especially those who are medically unfit, to achieve a response and experience improved outcome and quality of life.

We present the results of a single-arm multicentric phase 2 study evaluating the combination of obinutuzumab + idelalisib in R/R WM.

## Patients and methods

The trial was approved by an institutional review board of national ethical committees, Agence Nationale de Sécurité du Médicament et des Produits de Santé and Comité de Protection des Personnes. All patients provided written consent. This trial was registered at [www.clinicaltrials.gov](http://www.clinicaltrials.gov) as #NCT02962401. Enrollment lasted from March 2017 until July 2018. Data updates were performed in April 2020.

Patients were eligible if they met the Second International Workshop on Waldenström's Macroglobulinemia diagnostic criteria<sup>1</sup> and criteria to treat.<sup>19</sup> Patients were relapsing (progressing >6 months after the last administration of first-line or subsequent treatment) or refractory (progressing  $\leq$ 6 months after first-line or subsequent treatment) after 1 to 3 prior regimens/lines.

The primary end point was PFS, calculated from the date of inclusion to progression or death. Secondary end points included ORR, response rates (according to the V1th International Workshop on Waldenström's Macroglobulinemia criteria<sup>20</sup>) at months 8 and 24, OS, time to at least a minor response (MR), time to treatment failure (TTF), and time to new treatment. Secondary objectives were to assess safety and tolerability and the effect of *MYD88* mutations (*MYD88*<sup>MUT</sup>) and *CXCR4* mutations (*CXCR4*<sup>MUT</sup>) on response and PFS.

Measurable disease was required (ie, serum IgM  $\geq$ 2 times the upper limit of normal [ULN]). Patients were required to meet the following pretreatment laboratory criteria: platelet count  $\geq 75 \times 10^9/L$  and absolute neutrophil count  $\geq 1.5 \times 10^9/L$  if not the result of BM infiltration; creatinine clearance  $\geq 40$  mL/min; total bilirubin  $\leq 2$  times the ULN, unless clearly related to the disease or Gilbert syndrome; serum alanine aminotransferase levels  $\leq 2.5$  times the ULN; aspartate aminotransferase  $\leq 2.5$  times the ULN; and Eastern Cooperative Oncology Group Performance Status  $\leq 2$ . Patients with central nervous system involvement or with previous liver disease were excluded.

Serum IgM levels and complete blood cell counts were obtained at the beginning of each cycle for 6 cycles and then every 2 cycles thereafter. BM biopsies and computed tomography scans (if extramedullary disease was present at baseline) were repeated at cycles 6 and 24.

### Drugs

During induction, idelalisib was given continuously (150 mg, twice a day orally) in association with IV obinutuzumab (100 mg on day 1, 900 mg on day 2, and 1000 mg on days 8 and 15 of cycle 1 and, thereafter, on day 1 of cycles 2 to 6 (28-day cycle). During the maintenance phase, idelalisib was given alone for  $\leq 2$  years.

### Safety

Patients were closely monitored for infusion-related reactions. AEs were graded per Common Terminology Criteria for Adverse Events v.4.0. Idelalisib was held for absolute neutrophil count  $< 0.5 \times 10^9/L$ , grade  $\geq 2$  diarrhea, pneumonitis (of any grade), or grade  $\geq 3$  liver toxicity. Full-dose retreatment was permitted after toxicity recovery from the first drug hold; for subsequent events, reduction to 100 mg, twice a day, followed by discontinuance were required. Filgrastim or transfusion support was permitted. Herpesvirus and *Pneumocystis* prophylaxis were mandatory. All patients were tested

before the initiation of treatment for antibody-response status (IgG and IgM) against cytomegalovirus (CMV) antigens (serology). CMV pp65 antigen or polymerase chain reaction (PCR) testing was obtained at screening in CMV-seropositive patients, followed by every month during the induction phase and every 2 months during the maintenance phase.

### Gene mutation analysis

Next-generation sequencing (NGS) was performed centrally on DNA extracted from cells following Ficoll-Paque gradient centrifugation of BM samples collected at inclusion using the Ion PGM Torrent platform (Thermo Fisher Scientific). The custom gene panel included *MYD88*, *CXCR4*, *TP53*, *BTK*, and *PLCG2*. A mean coverage of 2000 to 2500 $\times$  was obtained for all targeted regions, allowing analysis of subclonal variation with a limit of detection of 0.8%.<sup>21</sup> In addition, droplet digital PCR was used to screen *MYD88*<sup>L265P</sup> and *CXCR4*<sup>S338X</sup> variants, with a limit of detection of 0.05%, according to the manufacturer's recommendations (Bio-Rad).

### Statistical analysis

The study aimed to assess whether PFS in advanced WM is improved by the combination of obinutuzumab + idelalisib compared with the combination applied in the usual therapy. The main chemotherapy-free approaches available when we designed the study were rituximab monotherapy or the combination of bortezomib, rituximab, and dexamethasone. We wished to demonstrate a median PFS  $\geq$ 25 months compared with 15 months under the null hypothesis. With  $\alpha = 0.05$ , power = 0.90, 24-month enrollment, and an assumed dropout rate at 0.010, the number of patients required was 50, based on a 1-sample log-rank test.<sup>22</sup> All summary statistics are medians (interquartile range [IQR]) or percentages. Swimmer plots were plotted to display individual patient data over the trial. Analysis of efficacy used an intent-to-treat principle, based on all enrolled patients. PFS, OS, and TTF were estimated by the Kaplan-Meier method. Cumulative incidence of new treatment used deaths free of treatment change as competing risk. The safety population considered patients who received  $\geq$ 1 dose of idelalisib. Data Safety Monitoring Board (DSMB) meetings were planned after the first 5 patients had received 3 cycles, after 25 enrolled patients, and after induction of treatment in 50 patients. All AEs and SAEs were submitted to the DSMB by the study statistician, independent of the sponsor and the investigators. Bayes estimation of the probability of grade  $\geq$ 3 AEs was computed, using a  $\beta$  binomial model with noninformative uniform priors. All analyses were performed using R 3.5.1 (<http://www.R-project.org>). Two-sided *P* values  $\leq .05$  denoted statistical significance.

## Results

### Patients and disease characteristics

Fifty patients were enrolled across 21 sites. After 1 screen failure, 49 patients were actually included. Baseline characteristics of patients and mutations are listed in Table 1. No mutation in *BTK* or in *PLCG2* was found in 3 patients previously treated with ibrutinib.

### Treatment

The study flowchart is displayed in Figure 1. Among the 49 enrolled patients, 1 died (unknown cause) before any treatment and 48 patients received the combination induction. Among the 48 treated

**Table 1. Baseline characteristics of enrolled patients (n = 49)**

Characteristics	Values
Age, y	68.5 (50-83)
Males/females, n	36/13
ECOG PS, 0/1/2/unknown, n	14/26/8/1
Adenopathy/splenomegaly/hepatomegaly, n	18/6/3
White blood cells, $\times 10^9/L$	5.25 (4.1-7.3)
Hemoglobin, g/dL	10 (8.9-11.1)
Platelets, $\times 10^9/L$	189 (94-2751)
Serum IgM, g/L	24.6 (13.5-48.2)
$\beta 2$ -microglobulin, mg/L	3.37 (2.7-4.7)
BM infiltration, %	55 (25-80)
<b>Genomic (NGS), n (%)</b>	
<i>MYD88</i> <sup>MUT</sup>	48 (97.9)
<i>MYD88</i> <sup>L265P</sup> / <i>MYD88</i> <sup>WT</sup>	46 (93.9)/1 (0.02)
<i>CXCR4</i> <sup>MUT*</sup>	27 (56.2)
<i>CXCR4</i> <sup>NS</sup> / <i>CXCR4</i> <sup>FS</sup>	20 (41.6)/7 (14.5)
<i>TP53</i> <sup>MUT†</sup>	11 (24.4)
<b>No. of previous treatments, median</b>	
1 line/2 lines/3 lines	30/12/6
DRC/other R-chemotherapy/ibrutinib	19/25/3

Data are median (IQR), unless otherwise noted. Median (IQR) and percentages were calculated for the entire cohort (49 patients), unless otherwise noted. Among the 3 cases of non-*MYD88*<sup>L265P</sup>, NGS identified 1 case with the *MYD88*<sup>M232I</sup> variant and 1 had a TG $\rightarrow$ CT substitution at position 38182641, which predicted the *MYD88*<sup>L265P</sup> variant.

DRC, dexamethasone-rituximab-cyclophosphamide; R-chemotherapy, rituximab chemotherapy.

\*Forty-eight patients were evaluated for *CXCR4* mutation.

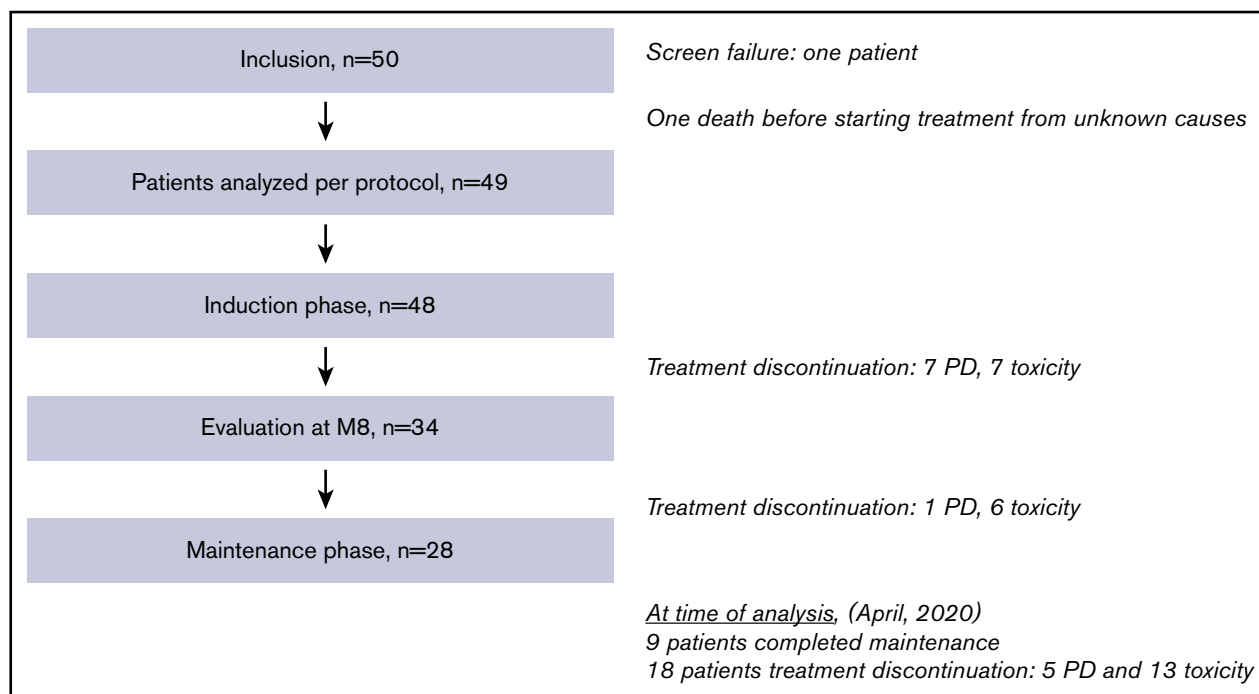
†Forty-five patients were evaluated for *TP53* mutation.

patients, induction was interrupted in 14 (7 for progressive disease and 7 for AEs). Maintenance was started in 27 patients. At the time of analysis (20 May 2020), 9 patients had completed maintenance and 18 patients had stopped treatment (13 for intolerance and 5 for progression).

### Efficacy

**Responses.** Forty-three patients were evaluable after induction for response assessment at month 8, with the 6 nonassessed patients considered failures. The best responses, reached after a median of 6.5 months (IQR, 3.4-7.1; range, 2.6-22.1 months), were very good partial response (VGPR) in 5 patients, partial response (PR) in 27 patients, and MR in 3 patients, leading to ORR and major response rate (MRR) estimates of 71.4% (95% confidence interval [CI], 56.7-83.4) and 65.3% (95% CI, 50.4-78.3), respectively. Stable disease and progressive disease were observed in 5 patients and 3 patients, respectively.

**Survival.** With a median follow-up time of 25.9 months (IQR, 22.3-30.5), median PFS was 25.4 months (95% CI, 15.7-29.0) and the PFS rates at 12 and 24 months were 75.5% (95% CI, 64.4-88.6) and 55% (95% CI, 42.7-70.9), respectively (Figure 2A). Median TTF was 25.6 months (95% CI, 19.9-30.4), and the cumulative incidence of new treatment at 12 and 24 months was 22.5% (95% CI, 11.9-35.0) and 36.8% (95% CI, 23.4-50.2), respectively. OS rates at 12 and 24 months were 97.8% (95% CI,



**Figure 1.** Study flowchart, patient disposition, and treatment exposure. M8, month 8; PD, progressive disease.

94.1-100) and 89.8% (95% CI, 81.7-98.7), respectively (Figure 2B). Altogether, 28 patients progressed. The outcome of the 3 patients previously exposed to ibrutinib is summarized in supplemental Figure 2. Among the 24 patients who received a new treatment, 14 received a chemotherapy-free regimen (ibrutinib [n = 10], 9 still in response; bortezomib + dexamethasone + rituximab [n = 1]; rituximab [n = 1]; idelalisib [n = 1]; bispecific mAb [n = 1]), and 10 received an immunochemotherapy combination. Of note, IgM rebound was observed after idelalisib discontinuation for toxicity in 4 patients who became symptomatic again and received a new treatment at 2, 8, 12, or 14 months after progression. Two other patients completed maintenance and presented an IgM rebound treated by the reintroduction of idelalisib and ibrutinib. Finally, 6 patients died, 2 free from relapse (1 from unknown cause before protocol therapy was started and 1 from infection after protocol ending), 1 from progression and 3 from infection (n = 2) or from thrombosis (n = 1) after progression.

#### **Impact of molecular profile on response and survival.**

Univariate analysis focusing on molecular screening found no significant impact on response rate, depth of response, time to response and PFS of either *CXCR4* (whatever the type of mutation) or *TP53* genotypes (Figures 2C-D and 3; supplemental Figure 1; supplemental Table 1). The only predictive factor of response was a high level of BM infiltration that was associated with a decreased occurrence of MRR. No prognostic factors for survival were found, except a deleterious effect of *TP53* mutation that was nonetheless not statistically significant in the Cox model (supplemental Table 1).

**Safety.** After enrollment of the first 25 patients, the DSMB validated the study continuation based on safety data and the safety profile of the combination treatment, without a need for any unplanned additional meetings. Altogether, 124 grade  $\geq 3$  AEs and

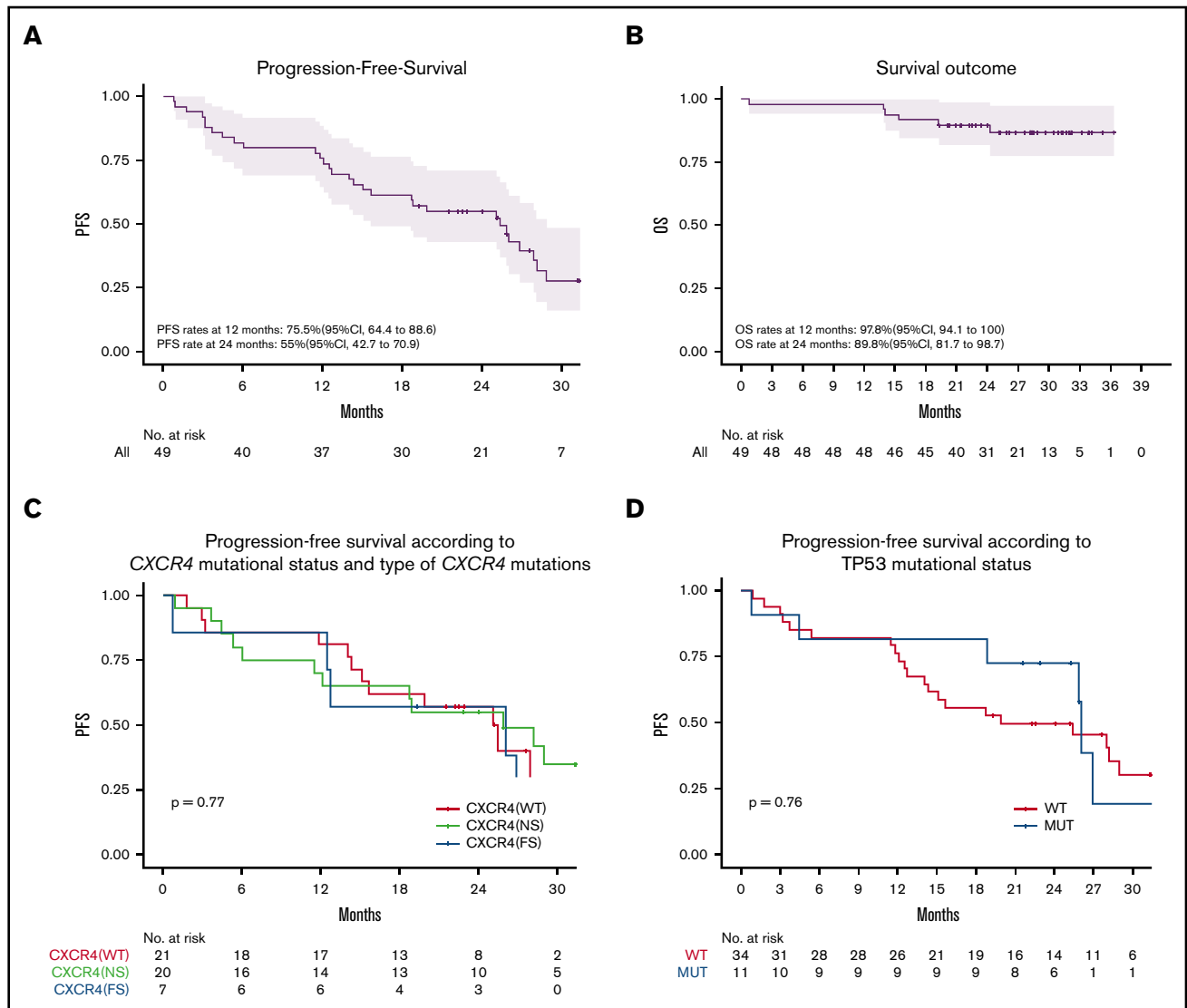
56 SAEs were reported in 43 and 36 patients, respectively. The most prevalent AEs are reported in Table 2. Based on the Bayesian inference, the posterior probability of experiencing  $\geq 1$  grade 3-4 AE was 87.0% (95% credibility interval, 76.4-94.7), with a 23% probability that 9 of 10 patients would experience such events. The probability that  $>3$  of 4 patients would experience grade 3-4 AEs was 98%.

No grade 5 toxicity, tumor lysis syndrome, or flare was reported. Four grade 2 CMV reactivations were reported, leading to introduction of an empiric antiviral treatment with a temporary discontinuation of idelalisib. Twenty-six patients (53%) discontinued idelalisib for toxicity after a median of 3 cycles (IQR, 2-4; range, 1-5). Median duration of idelalisib exposure was 8.1 months (IQR, 3.2-15.3). Dose reduction was required in 21 (43%) patients.

## **Discussion**

We report a prospective study of the combination of idelalisib + obinutuzumab in symptomatic R/R WM patients with a 2-year fixed duration of idelalisib. We aimed to demonstrate that median PFS could be improved up to 25 months. This hypothesis was confirmed by our study, with an estimated median PFS of 25.4 months, although likely values given by the 95% CI ranged from 15.7 to 29.0 months. In previously treated WM patients, reuse of a rituximab-based regimen with cyclophosphamide, bendamustine, or proteasome inhibitor was a therapy option if the response lasted  $> 3$  years, whereas ibrutinib alone, especially in rituximab-refractory patients, or in combination with rituximab are also acceptable options.<sup>23</sup> Ibrutinib is used in R/R WM patients with a high response rate and a median PFS not reached at 5 years.<sup>24</sup> There is actually no consensus in this setting with regard to the duration of treatment (ie, fixed vs until progression).





**Figure 2. OS and PFS according to molecular profile.** PFS (A), OS (B), PFS according to *CXCR4* mutational status and the type of *CXCR4* mutation (frameshift vs non-sense) (C), and PFS according to *TP53* mutational status (D). (C) Red line, absence of *CXCR4* mutation [CXCR4(WT)] (left panel), green line, *CXCR4* mutation non-sense [CXCR4(NS)]; blue line, *CXCR4* mutation frameshift [CXCR4(FS)]. (D) Red line, absence of *TP53* mutation (WT); blue line, *TP53* mutation (MUT).

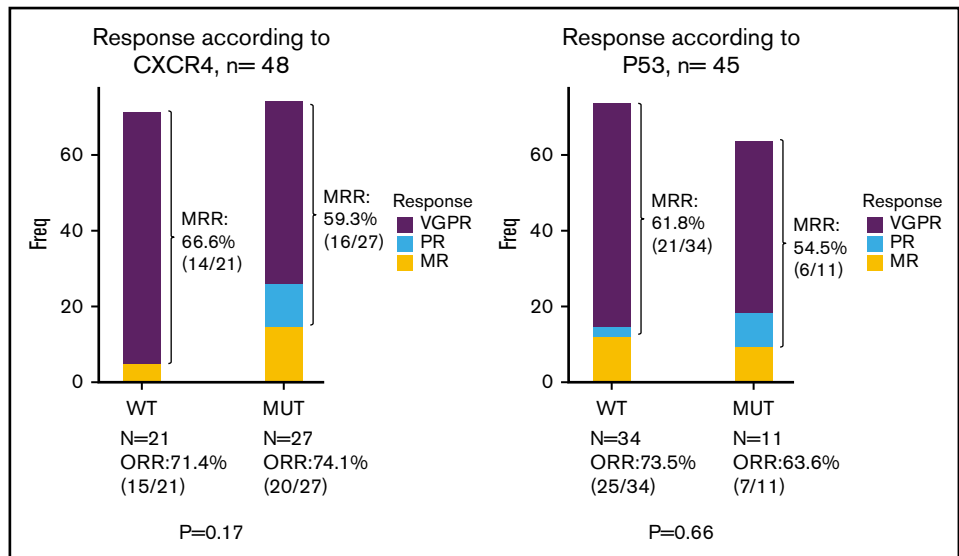
Idelalisib reversibly inhibits the p110 $\delta$ -PI3K isoform, leading to decreased phosphorylation of several downstream targets, including Akt, directly inducing apoptosis and ultimately disrupting interactions between tumor cells and the BM microenvironment.<sup>13,25</sup> The PI3K/AKT signaling pathways are hyperactivated in WM, highlighting the interest in PI3K inhibitors for WM.<sup>26</sup> Considering its combination with obinutuzumab, PI3K $\delta$  inhibition by idelalisib has minimal impact in vitro on the immune effector function of obinutuzumab and rituximab, as measured in natural killer cell-mediated antibody-dependent cellular cytotoxicity, macrophage-mediated antibody-dependent cellular cytotoxicity (ADCP), and whole-blood B-cell depletion.<sup>27</sup>

Data on efficacy and toxicity in WM patients are scarce, with no data available on the combination with obinutuzumab in this setting. The largest study reported 80% ORR in 10 patients with lymphoplasmacytic lymphoma/WM among 125 low-grade lymphomas that was refractory to anti-CD20 and alkylating agents and

treated with idelalisib (150 mg twice a day by mouth).<sup>14</sup> These results were updated in 2019 with a median duration of therapy in WM of 29 months (range, 6.4-51). Of the 9 responders, 7 had PR, 1 had MR, and 1 had SD. Median PFS was 22 months, and the median duration of response was 20 months. All patients experienced  $\geq 1$  AE related to the treatment, and  $>65\%$  experienced a grade  $\geq 3$  AE. The most common grade  $\geq 3$  AEs included neutropenia (28%), diarrhea (28%) and increased liver enzymes (16%). Among patients on therapy for 6 months, AE led to dose reductions for 19.2%, treatment discontinuation for 23.1%, and death for 10.3%.<sup>28</sup> A phase 2 study aimed at evaluating idelalisib, 150 mg twice a day by mouth, in R/R WM patients was terminated early because of grade  $\geq 3$  increased liver enzyme levels in 3 of the first 5 enrolled patients.<sup>16</sup>

The pivotal ibrutinib study, which was recently updated,<sup>28</sup> and other trials with new BTK inhibitors (monotherapy-like zanubrutinib<sup>29</sup> and

**Figure 3. Best responses according to *CXCR4* (left panel) and *TP53* (right panel) mutational status (n = 49 and n = 45, respectively).**



acalabrutinib<sup>30</sup>) have provided evidence that BTK inhibitors produce long-term disease control as single-agent therapy with manageable safety profiles in treatment-naïve or R/R WM patients; most clinicians consider them the treatment of choice in this patient group. Overall, MRR was 80%, with a median follow-up of 27 to 59 months. With regard to BTK inhibitors, the impact of *MYD88* and *CXCR4* mutations on the quality and duration of the response remains controversial. The impact of *MYD88*<sup>MUT</sup> and *CXCR4*<sup>MUT</sup> status was reported in patients treated with ibrutinib. Response rates, depth of response, and time to response were highly dependent on the *MYD88* and *CXCR4* genotype. *MYD88*<sup>WT</sup> patients appear to have a lower response rate to ibrutinib.<sup>7,11,28</sup> However, the results of the iNNOVATE study combining ibrutinib + rituximab questioned the impact of *MYD88*<sup>MUT</sup> status on the response, as well as the second generation of BTK inhibitors, such as acalabrutinib or zanubrutinib, with a high response rate in *MYD88*<sup>WT</sup> patients in the ASPEN study.<sup>8,10,31</sup> More data are necessary to confirm whether or not *MYD88*<sup>MUT</sup> genotype has

a negative impact. Of 49 included patients, only 1 had a confirmed *MYD88*<sup>WT</sup> genotype, which is insufficient to draw any conclusion for this subgroup.

Ibrutinib-treated *CXCR4*<sup>MUT</sup> patients experienced a longer median time to reach MR and lower MRR, including attainment of VGPR, in previously treated WM.<sup>11,32,33</sup> Faster IgM response kinetics were also observed for patients with rituximab-refractory *CXCR4*<sup>WT</sup> disease who received ibrutinib monotherapy.<sup>9</sup> The frequency of *CXCR4*<sup>MUT</sup> patients in our trial was higher than previously described (56%), likely as a result of the sensitivity of the deep NGS strategy and digital PCR, as well as the relapse setting.<sup>34,35</sup> We did not find any difference in time to response, quality of response, or PFS according to *CXCR4* mutation profile or type of mutation (frameshift [FS] vs nonsense [NS]), in contrast to previously described patients treated with ibrutinib.<sup>35</sup> *TP53* mutations were observed in 24% of the patients in our R/R cohort, which is also higher than previously described at diagnosis.<sup>21,36</sup> Sequential *TP53*

**Table 2. Primary AEs observed in patients**

AE	Total AEs (N = 279)	Grade 1-2 (n = 155) (55.5%)	Grade 3-4 (n = 124) (44.5%)	SAEs (n = 56)
<b>Hematologic</b>	<b>82 (29.4)</b>	<b>24 (8.6)</b>	<b>58 (20.8)</b>	<b>2 (3.6)</b>
Neutropenia	54 (19.4)	9 (3.2)	45 (16.1)	1 (1.8)
Anemia	16 (5.7)	9 (3.2)	7 (2.5)	1 (1.8)
Thrombocytopenia	12 (4.3)	6 (2.2)	6 (2.2)	0
<b>Nonhematologic</b>	<b>197 (70.6)</b>	<b>131 (46.9)</b>	<b>66 (23.6)</b>	<b>54 (96.4)</b>
Diarrhea	24 (8.5)	9 (3.2)	15 (5.3)	12 (21.4)
Hepatic cytolysis	26 (9.3)	7 (2.5)	19 (6.8)	4 (7.1)
Infections	52 (18.6)	44 (15.7)	8 (2.8)	13 (23.2)
Skin	16 (5.7)	11 (3.9)	5 (1.8)	3 (5.3)
IRR	15 (5.4)	15 (5.3)	0	1 (1.8)
Others	64 (22.9)	45 (16.4)	19 (6.8)	21 (37.5)

Only AEs with rates  $\geq 5\%$  are reported. Some patients exhibited grade 1-2 and grade 3-4 AEs, so that patient numbers do not add up to the total by row. Eight other malignancies were observed: 5 skin cancer (3 basocellular, 2 epidermoid), 1 prostatic adenocarcinoma, 1 myelodysplastic syndrome, and 1 glioblastoma.

IRR, infusion-related reaction.

analysis was recently reported in a cohort, including 14 patients with WM, and the acquisition of *TP53* variation over time was observed in 29% of cases.<sup>37</sup> Selection of the *TP53* clone is well described under chemotherapy pressure in other B-cell lymphoproliferative disorders pretreated with genotoxic agents.<sup>22,36</sup> The poor prognostic impact of the *TP53* mutation has primarily been described in a retrospective analysis of untreated patients or after immunochemotherapy but not in patients treated with BCR signaling inhibitors.<sup>21,36</sup> In our study, there was no evidence of any predictive value of *TP53* mutation for response or PFS in this small subgroup of patients.

This study has limitations but suggests new treatment opportunities. First, with a median PFS of 25.5 months, we reached our primary goal of demonstrating a median PFS  $\geq 25$  months compared with 15 months with usual therapy. Second, the sample size does not allow firm conclusions about the impact of genotype on efficacy, because the study was not designed for this end point. Our results suggest that this combination might be of interest in *CXCR4*<sup>MUT</sup> WM patients with advanced disease, because they demonstrate reduced response to ibrutinib. Compensatory PI3K/Akt signaling may contribute to ibrutinib resistance in the *CXCR4*<sup>MUT</sup> WM model, as well as in the *CXCR4*<sup>WT</sup> WM model, resulting in synthetic lethality toward the PI3K/Akt inhibitor.<sup>38-40</sup> Therefore, targeting of the PI3K/Akt pathway could be an option in BTK inhibitor-resistant WM patients and appears to be a proof of concept regarding the efficacy of this class of compound in WM patients. In addition, for the first time, we provide an estimate of the response rate or PFS achieved with a combination regimen including inhibitors of BCR-associated kinases in *TP53*<sup>MUT</sup> WM patients; however, the number of patients was small, and a nonstatistically significant deleterious effect of *TP53* mutation was observed. Third, the safety of the combination with high liver and gastrointestinal toxicity led to early treatment discontinuation and/or a dose reduction in half of the patients. Using Bayesian analysis, which quantified, through probabilistic statements, the proportion of treated patients who experienced grade 3 or 4 AEs, it appeared more than likely (98%) that more than three fourths of patients could be impacted by such events. These results did not differ substantially from those observed by Gopal et al<sup>14</sup> and Wagner-Johnston et al.<sup>28</sup> It is a limitation of further clinical use. However, considering the potential efficacy of this class of agents, next-generation PI3K inhibitors associated with obinutuzumab could overcome the toxicity profile observed in this phase 2 trial and may be a reasonable treatment option when standard therapies have been exhausted.

In summary, our study demonstrates interesting efficacy of the combination of idelalisib + obinutuzumab in symptomatic R/R WM patients, although toxicity is a major limitation. Nevertheless, there is a need to develop new approaches in patients who are refractory to BTK inhibitors. This population may benefit from this therapeutic option, which could be investigated in further clinical trials.

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## Authorship

Contribution: C.T., S.P., S.C., and V.L. conceived and designed the study, analyzed and interpreted data, and wrote the manuscript, all authors collected and assembled data and approved the final version of the manuscript; and C.T., S.P., D.N., S.C., and V.L. are accountable for all aspects of the work.

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ORCID profiles: C.T., 0000-0003-4093-6463; C.H., 0000-0003-4910-476X; A.P., 0000-0003-0131-8689; O.T., 0000-0002-9438-621X; S.L., 0000-0003-4282-2160; O.C., 0000-0002-1156-8983; D.R.-W., 0000-0002-7767-755X.

Correspondence: Cécile Tomowiak, Service d'Hématologie et Thérapie Cellulaire, PRC, Hôpital de La Milétrie, Faculté de médecine and INSERM CIC 1402, CHU, 2 Rue de la Milétrie, 86021 Poitiers Cedex, France; e-mail: cecile.tomowiak@chu-poitiers.fr.



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