Ixazomib, pomalidomide and dexamethasone in relapsed or refractory multiple myeloma characterized with highrisk cytogenetics: the IFM 2014-01 study

The definition of high-risk (HR) multiple myeloma (MM) is constantly evolving,¹ the presence of a deletion of 17p13 (del[17p]) and/or translocation (4;14) (t[4;14]), which represents approximately 20% to 25% of patients in the series remains the main definition of HR MM.² These patients are characterized with shorter survivals, related to an early relapse rate (median time to progression [TTP] <4 months) and rapid development of mechanisms of resistance to treatments.² The multicenter phase II, IFM 2010-02 study, evaluated pomalidomide plus dexamethasone (Pd) in RRMM characterized with del(17p) and/or t(4:14) demonstrated limited activity, with an median TTP at 3.0 months, at 7.3 and 2.8 months in case of del(17p) and of t(4;14), respectively.³ The development of triplet or quadruplet-based regimens has improved survival, with limitations to these treatment regimens because of the safety profile issues that can be experienced as difficult.⁴⁻⁶ We hypothesized that addition of ixazomib (a boronate acid oral proteasome inhibitor) at increased dose density to Pd (IxPd) in HR RRMM would provide a triplet-based treatment approach with improved convenience and safety profile, and thus adherence to treatment to ease the continuous treatment until progression.

This study is a multicenter, open-label, single arm, phase II study of IxPd in RRMM with adverse genomic abnormalities. The main eligibility criteria were RRMM in line 2 or 3, refractory to lenalidomide, but not to pomalidomide and ixazomib, a measurable disease,⁷ a platelet count \geq 75×10⁹/L, a neutrophil count $\geq 1.0 \times 10^{9}$ /L, and creatinine clearance (MDRD) >30 mL/ min. The cytogenetic analysis was performed centrally by Pr Avet Loiseau/Pr Corre Jill on sorted bone marrow plasma cells. HR was define as presence of either del(17p) and/or t(4:14) since diagnosis or at study entry to compare IxPd to Pd from the IFM 2010-02 study. The study was sponsored by the Intergroupe Francophone du Myélome (IFM) in accordance with national regulations in France (Eudract number 2016-002650-20). This trial is registered at clinicaltrials gov Identifier: NCT03683277. The patients received 17 induction 21-days'-based cycles, consisting of ixazomib 3 mg/day (days 1, 4, 8 and 11), pomalidomide 4 mg/day (days 1 to 14) and weekly dexamethasone (40 mg day 1, 8 and 15, 20 mg \geq 75 years old), followed by a maintenance phase of 28-day-based cycles with ixazomib 4 mg/day (days 1, 8 and 15) and pomalidomide 4 mg/day (days 1 to 21), given until progression. The primary endpoint was time to progression (TTP).7 Secondary endpoints included the safety profile of the regimen using the National Cancer Institute Common Terminology Criteria for Adverse Events (version 4.0), the response rates, response durations, overall survival time (OS) and progression-free survival (PFS). Sparse pharmacokinetic (PK) samples were collected from patients on days 1 and 11 of cycle 1, then pre-dose for cycles 2 to 5. We studied whether PK analysis would differ in HR RRMM from previous studies using the same dose schedule of Ixazomib (MLN2238). We considered a 100% improvement in median TTP from the median TTP at 3 months in IFM 2010-02 as clinically relevant in IFM 2014-01. The number of patients to be included is 26 patients if using a two-sided test with a type i error of 5% and ensuring a power of 80%.8 Considering the median TTP would differ across HR RRMM,⁹ and under the assumption that we would study separately the two populations of HR RRMM, the statistical design considered the inclusion of 70 patients with a power slightly above 90, assuming a minimum of 40% of patients would be included in the least represented population.⁸ We recruited 26 patients then the study was terminated prematurely on December 31, 2022 due to lack of recruitment. A safety analysis was performed when 15 patients had completed at least one cycle at ANSM (agence nationale de securite du medicament) request. The data monitoring committee (DMC) confirmed the treatment was safe and allowed the study to fully recruit. The primary analysis was conducted on an intention-to-treat basis (ITT). All statistical analyses were performed with SAS version 9.4 (SAS Institute, Cary, NC) and R software version 4.0.4 (The R Foundation for Statistical Computing, Vienna, Austria).

Of the 32 screened patients, 26 were included and started the protocol induction phase, eight received the maintenance according to protocol (*Online Supplementary Figure S1*). At end of study, 25 patients had discontinued the study treatment, 22 for progressive disease, 15 patients died, all of them related to MM, and one patient remains on treatment with commercial products. At study entry, 12 patients presented with del(17p), 9 with t(4;14), and five with del(17p) and t(4;14). Fifteen patients had two prior lines. All patients were refractory to lenalidomide. The patients' characteristics are summarized in Table 1.

With a median follow-up time of 27 months (interquartile range [IQR], 23-29), the median TTP time improved by more than three times at 10.2 months (95% confidence interval [CI]: 4.4-13.0). The median OS time was 23.7 (95% CI: 12.2-not evaluable [NE]) months. Survival times are shown in Figure 1. The patients with t(4;14) had longer median survival times compared to IFM 2010-02, the benefit appeared limited in other subgroups of interest. The response rates were similar at the end of induction and at study completion (*Online*)

Supplementary Table S1). The median duration of response was 10.0 (IQR, 3.4-13.6) months, similar across HR subgroups. Dose-normalized PK concentrations were comparable across patient response subgroups; a slight trend toward higher values was observed in patients with partial response/very good partial response (PR/VGPR) (Figure 1E).

At the data cutoff, 25 patients (96%) had discontinued treatment and the most common reason was disease progression (N=18, 70%). All patients experienced at least one AE, with a median frequency of ten occurrences per patient

Table 1. Patients' characteristics, according to their abnormalities, t(4;14) and del(17p), at inclusion, median (Q1-Q3) or N (%) and description of prior lines of therapy, N (%). Data presented as a whole and according to their abnormalities, t(4;14) and del(17p), at inclusion.

	All N=26	t(4;14) N=9	Del(17p) N=12	Del(17p) and t(4;14) N=5
Age in years ≥75 years	72 (67-78) 12 (46)	67 (67-76) 4 (44)	74 (67-80) 6 (50)	73 (68-75) 2(40)
Two prior lines**	15 (58)	7 (78)	6 (50)	2 (40)
Time from diagnosis to study entry	4 (2-5)	4 (3-5)	3 (2-6)	3 (1-4)
Time in days from relapse to treatment start	34 (26-48)	29 (21-35)	42 (26-67)	47 (34-47)
ß2-microglobulin mg/L*	5 (4-7)	5 (4-8)	4 (3-5)	6 (4-8)
ISS stage III*	9 (36)	4 (44)	3 (25)	2 (50)
LDH abnormal*	6 (27)	1 (12)	3 (30)	2 (50)
Circulating plasma cells**+	5 (19)	2 (22)	3 (25)	0 (0)
Cytogenetic* t(4;14) del(17p) del(17p) and t(4;14) +1q gain +del1p32	7 (27) 7 (27) 3 (11) 4 (15) 4 (15)	7 (78) 0 (0) 0 (0) 2 (22) 2 (22)	0 (0) 7 (70) 0 (0) 1 (8) 2 (17)	5 (100) 3 (60) 3 (60) 1 (20) 0 (0)
Plasmacytomas** Paraskeletal (PMD) Soft tissue (EMD)	4 (15) 2 (8)	1 (11) 0 (0)	2 (17) 2 (17)	1 (20) 0 (0)
Line 1 Bortezomib-based* VMP VCyd VTd Lenalidomide-based** Rd RCyd VRd DRd AntiCD38VRd Autologous transplantation-based	26 6 (23) 5 (19) 3 (12) 4 (15) 1 (4) 4 (1) 1 (4) 2 (8) 7 (27)	9 3 (33) 2 (22) 1 (11) 1 (11) 0 (0) 2 (22) - - 3 (33)	12 3 (25) 2 (17) 2 (17) 2 (17) 0 (0) 1 (8) 1 (8) 1 (8) 1 (8) 4 (33)	5 0 (0) 1 (20) 0 (0) 1 (20) 1 (20) 1 (20) 0 (0) 1 (20) -
Line 2 VMP Rd VRd DRd KRd	15 1 4 3 6 1	7 1 1 3 1	6 0 2 1 3 0	2 0 1 1 0 -

*At diagnosis; **at inclusion. *Circulating plasma cells were identified and counted by regular flow analysis. Quantitative data were expressed as means ± standard deviation (SD) or median (25th-75th percentiles). Qualitative variables are given as number (percentage) of patients. ISS: International Staging System; LDH: lactate dehydrogenase; PMD: paramedullary disease; EMD: extramedullary disease; VMP: bortezomib plus melphalan and prednisone; VCyd: bortezomib plus cyclophosphamide and dexamethasone; VTd: bortezomib, thalidomide, and dexamethasone; Rd: lenalidomide and dexamethasone; VRd: bortezomib, lenalidomide, and dexamethasone; DRd: daratumumab, lenalidomide, and dexamethasone; KRd: carfilzomib, lenalidomide and dexamethasone; Q: quartile.



Figure 1. Survival from study entry (N=26) - time to progression. (A) Overall population. (B) According to high-risk relapsed/refractory multiple myeloma (HR RRMM) population, either del(17p), or t(4:14) or del(17p) and t(4:14). (C) Overall, population as a whole. (D) According to HR RRMM population, either del(17p), or t(4;14) or del(17p) and t(4;14). The median time to progression (TTP) and overall survival (OS) were at 10.5 (95% confidence interval [CI]: 7.9-not estimable [NE]) and NE (95% CI: 27.1-NE), as a whole, respectively. Kaplan-Meier method was used to analyze time-to-event data. Results were expressed as median time-to-event in months and 95% CI. Pharmacokinetic (PK) analysis (N=26). (E) The patients are stratified by best response over the course of the study. Sparse pharmacokinetic (PK) samples were collected from patients on days 1 and 11 of cycle 1, then pre-dose for cycles 2 to 5. The objective was to study whether PK analysis would differ in HR RRMM from previous studies using the same dose schedule of ixazomib (MLN2238). The systemic ixazomib concentrations were quantified from patient plasma samples using a validated liquid chromatography/tandem mass spectrometry assay with a range of 0.5 to 500 ng/mL. Given that ixazomib exhibits dose-proportional PK, all PK concentrations in this analysis were normalized by the corresponding dose administered; these normalized values enable the comparison across studies and dose levels. The dose-normalized PK concentrations in our study were comparable with a previous single-agent dose escalation trial which utilized the same dose regimen (clinicaltrials gov Identifier: NCT00932698) and the dose-normalized PK concentrations were comparable across patient response subgroups. These results are consistent with previous model-driven exposure-response analyses that concluded ixazomib exposures were not a statistically significant predictor of complete response (CR), \geq very good partial response (≥VGPR), or ≥ partial response (≥PR) rate. Despite a slight trend toward higher values observed in patients who experienced a PR/VG-PR response, this data supports that HR RRMM patients don't seem to have different PK concentrations' of ixazomib, independently of the dose density or intensity of the drug. This data does not support, therefore, the use of the serum concentration of ixazomib as a potential biomarker to tailor treatment schema of ixazomib and better determine dose density over intensity in order to optimize the activity of Ixazomib in RRMM HR. MR: minor response; NA: not applicable; PD: progressive disease; SD: stable disease; Conc: concentration.

(IQR, 4-15), and 85% (N=22) experienced AE of grade \geq 3, with a median frequency of two occurrences per patient (IQR, 1-4) (*Online Supplementary Table S2*). There were no observed differences across study phases, induction and maintenance, and across HR subgroups. The summary of the dose relative intensity of ixazomib and pomalidomide is provided in Table 2.

Our study met its primary endpoint, however, it is fair to conclude that the benefit brought to these patients is limited with our IxPd regimen. The other approved regimens available to these patients lenalidomide refractory, across high-risk and even non-high risk RRMM patients, also showed limited activity, with median PFS from 7.8 months to 28.1 months.¹⁰ These data confirm that we have recruited patients that were truly high-risk being lenalidomide refractory RRMM, even in early relapse L2 and L3, and on top cytogenetic high-risk, thus characterized with a very poor outcome. Our results demonstrate that new drug developments are needed for this population.

Our study results appear to confirm the added value of a proteosome inhibitor (PI) in HR MM with t(4;14) compared to Pd doublet-based regimen. This raises questions on whether the effect seen with IxPd was only explained by the triplet-based regimen, triplet expectedly performing better, or whether there is a true biological effect, yet to be demonstrated. In our study, not all t(4;14) HR MM benefited from the study treatment, also questioning on certain t(4;14) being of lesser HR profile compared to other.¹¹

There are three different PI available in MM in various indications, ixazomib and bortezomib that belong to the boronate acid-based family and carfilzomib an epoxyketone-based PI. The bortezomib's safety profile remains difficult for a long period of time in HR MM, who tend to relapse rapidly when treatments are stopped, despite the subcutaneous and weekly developments.^{12,13} When IFM 2014-01 started, carfilzomib was available only as a bi-weekly schema, with an increasing risk of cardio vascular safety concerns. Ixazomib appeared an interesting PI to study given its oral bioavailability and its favorable safety profile. Ixazomib was therefore the most appealing PI to improve the adherence of patients to a prolonged study treatment for responders and to combine with immunomodulatory drugs.¹⁴

We believe our study results support our hypothesis that the twice-weekly ixazomib concentrations with 12 mg total given per 21 days-based cycle of ixazomib, with reduced doses (3 mg at days 1, 4, 8 and 11), in the induction phase might be of importance in HR RRMM, compared to Tourmaline-MM1 study.¹⁵ Our study population was harder to treat, and we have observed overall response rate (ORR) and VGPR rates at 60% at end of induction phase, while ORR and \geq VGPR rates were 78% and 48% in Tourmaline-MM1 study. It would

Table 2. Number of cycles, relative dose intensity of ixazomib and pomalidomide, and patients with more than 12 cycles of treatment, as a whole and according to their abnormalities, t(4;14) and del(17p), at inclusion.

	All N=26	t(4;14) N=9	Del(17p) N=12	Del(17p) and t(4;14) N=5			
Ixazomib							
Number of cycles, median (Q1-Q3) All study long Induction Maintenance*	9 (4-19) 9 (4-17] 4.5 (2-7)	13 (6-20) 13 (6-17) 6 (3-8)	9.5 (3.5-19.5) 9.5 (3.5-17) 4.5 (2.5-7)	8 (5.5-12) 8 (5.5-12) 1 (1-1)			
Relative dose intensity, median (Q1-Q3) Induction Maintenance*	80.0 (69.2-90.6) 69.4 (60.0-74.2)	79.4 (69.0-87.7) 72.1 (64.9-73.0)	77.7 (68.3-94.2) 66.2 (56.3-87.5)	87.5 (71.9-95.9) 66.7, N=1			
Patients with >12 cycles, N (%)	12 (46)	5 (56)	6 (50)	1 (20)			
Pomalidomide							
Number of cycles, median (Q1-Q3) All study long Induction Maintenance*	9 (4-19) 9 (4-17) 4.5 (2-7)	13 (6-20) 13 (6-17) 6 (3-8)	9.5 (3.5-19.5) 9.5 (3.5-17) 4.5 (2.5-7)	8 (5.5-12) 8 (5.5-12) 1 (1-1)			
Relative dose intensity, median (Q1-Q3) Induction Maintenance*	89.6 (73.7-97.9) 89.3 (58.1-100.0)	84.7 (71.1-97.8) 97.6 (85.1-112.5)	87.3 (71.9-98.9) 84.3 (58.1-100.0)	91.7 (82.5-94.2) 0, N=1			
Patients with >12 cycles, N (%)	12 (46)	5 (56)	6 (50)	1 (20)			

*The number of patients analyzed at maintenance phase were overall 8 patients, breakdown as t(4;14) N=3, del(17p) N=4, and del(17p) and t(4;14) N=1. The relative dose intensity of pomalidomide and ixazomib drugs' administrations was calculated as the dose concentration received by patients over the protocol-planned doses concentrations across the study induction and maintenance phases. Q: quartile.

be of interest to further develop this design in future and larger ixazomib studies.

As a conclusion, the study IFM 2014-01, a phase II study of the triplet-based combination of ixazomib plus pomalidomide and dexamethasone, in high-risk early RRMM refractory to lenalidomide, met its primary end point objective. We have observed an increase of the median TTP with the addition of ixazomib to pomalidomide and dexamethasone in this very hard to treat population characterized with a very poor outcome. This data confirms the importance of introducing a proteasome inhibitors in HR RRMM treatment's regimens, particularly in t(4;14). This phase II study needs confirmation in a larger cohort.

Authors

Arthur Bobin,¹ Salomon Manier.² Joe de Keizer,³ Jaydeep K. Srimani,⁴ Cyrille Hulin,⁵ Lionel Karlin,⁶ Denis Caillot,⁷ Ingrid Lafon,⁷ Clara Mariette,⁸ Carla Araujo,⁹ Bertrand Arnulf,¹⁰ Benoît Bareau,¹¹ Karim Belhadj,¹² Lofti Benboubker,¹³ Thorsten Braun,¹⁴ Claire Calmettes,¹⁵ Olivier Decaux,¹⁶ Mamoun Dib,¹⁷ Hélène Demarquette,¹⁸ Caroline Jacquet,¹⁹ Cécile Sonntag,²⁰ Sophie Godet,²¹ Arnaud Jaccard,²² Pascal Lenain,²³ Margaret Macro,²⁴ Valentine Richez-Olivier,²⁵ Mourad Tiab,²⁶ Laure Vincent,²⁷ Hacene Zerazhi,²⁸ Marie-Odile Pétillon,²⁹ Sandrine Rollet,²⁹ Helene Gardeney,¹ Geraldine Durand,¹ Anthony Levy,³⁰ Cyrille Touzeau,³¹ Aurore Perrot,³² Philippe Moreau,³¹ Thierry Facon,² Jill Corre,³³ Stephanie Ragot,³ Herve Avet-Loiseau³³ and Xavier Leleu¹

¹Poitiers University, U1313, U1402, Poitiers, France; ²Department of Hematology, University Hospital and INSERM UMR-S1277, Lille, France; ³Clinical Investigation Center CIC INSERM 1402, University Hospital, Poitiers, France; ⁴Takeda Development Center Americas, Cambridge MA, USA; ⁵Department of Hematology, Haut-Lévêque Hospital, Pessac, France; ⁶Department of Clinical Hematology, Civils Hospices, Lyon, France; ⁷Department of Oncology Hematology, Bourgogne Cancerology Institute, Dijon, France; ⁸Department of Hematology, University Hospital, Grenoble, France; ⁹Department of Hematology, Côte Basque Hospital, Bayonne, France; ¹⁰Department of Hematology, Saint-Louis Hospital, Paris, France; ¹¹Department of Hematology, Private Hospital, Cesson-Sévigné, France; ¹²Lymphoid Hemopathies Unit, Henri Mondor University Hospital, Créteil, France; ¹³Department of Hematology and Cell Therapy, Tours University Hospital, France; ¹⁴Department of Clinical Hematology, Avicenne Hospital, Bobigny, France; ¹⁵Department of Onco-Hematology, Périgueux Hospital, Périgueux, France; ¹⁶Department of Clinical Hematology, UMR U1236 INSERM, Rennes University, Rennes, France; ¹⁷Blood Disease Department, University Hospital, Angers, France; ¹⁸Department of Hematology, Dunkerque Hospital, Dunkerque, France; ¹⁹Department of Hematology, University Hospital - Brabois Hospitals, Vandoeuvre les Nancy, France; ²⁰Department of Hematology, ICANS University Hospital, Strasbourg, France; ²¹Department of Clinical Hematology, University Hospital, Robert Debré Hospital, Reims, France; ²²Department of Clinical Hematology and Cell Therapy, University

Hospital, Limoges, France; ²³Department of Hematology, Henri Becquerel Center, Rouen, France; ²⁴Department of Cinical Hematology, University Hospital, Caen, France; ²⁵Department of Clinical Hematology, Archet Hospital, Nice, France; ²⁶Department of Onco-Hematology, Vendée Departmental Hospital, La Roche/Yon, France; ²⁷Department of Clinical Hematology, Montpellier University, Montpellier, France; ²⁸Department of Onco-Hematology, Henri Duffaut Hospital, Avignon, France; ²⁹Intergroupe Francophone du Myélome, Paris, France; ³⁰Department of Onco-Hematology, Vichy Hospital, Vichy, France; ³¹Department of Clinical Hematology, University Hospital, Hôtel Dieu, Nantes, France; ³²Department of Hematology, CRCT INSERM 1037, University Hospital, IUCT-O, Toulouse, France and ³³Unit for Genomics in Myeloma, CRCT INSERM 1037, University Hospital, IUCT-O, Toulouse, France

Correspondence:

X. LELEU - xavier.leleu@chu-poitiers.fr

https://doi.org/10.3324/haematol.2024.285916

Received: June 14, 2024. Accepted: November 5, 2024. Early view: November 14, 2024.

©2025 Ferrata Storti Foundation Published under a CC BY-NC license 🖭 😨

Disclosures

AB discloses consultancy for Sanofi, Janssen and Amgen. SM discloses consultancy for Abbvie, Adaptive Biotechnology, Amgen, Celgene/BMS, GlaxoSmithKline, Janssen, Novartis, Pfizer, Regeneron, Roche, Sanofi and Takeda. LK discloses honoraria from and advisory board participation at Amgen, Celgene-BMS, Janssen, Takeda, Abbvie, Sanofi and Pfizer. CM discloses advisory board participation at Janssen, Menarini, BMS and Sanofi. KB discloses advisory board participitation at Amgen; discloses honoriara from Amgen, BMS, Janssen, and Sanofi. OD discloses honoraria from Janssen, Celgene/BMS, Amgen, Takeda, GSK, Sanofi, Abbvie, Roche, The Binding Site, Sebia, Menarini-Stemline and Pfizer. CS discloses consulting fees and honoraria from Takeda, BMS, Janssen and Amgen. MM discloses research funding from Janssen and Takeda; discloses honoraria from Amgen, BMS, Janssen, Sanofi and Takeda. LV discloses advisory board participitation at Takeda and BMS. CT discloses advisory board participation at and honoraria from Takeda and Celgene/BMS. AP discloses honoraria from Abbvie. Amgen, BMS, Janssen, Pfizer, Sanofi, and Takeda. PM discloses honoraria from and advisory board participitation at Janssen, Celgene/BMS, Takeda, Amgen, Sanofi, Pfizer and Abbvie. JC discloses advisory board participation at Sanofi and Bristol Myers Squibb; discloses consultancy for Janssen, Sanofi, Bristol Myers Squibb, Pfizer and Adaptive; discloses research support from Sanofi and Bristol Myers Squibb. XL discloses honoraria and consultancy from Abbvie, BMS, GSK, Janssen, Novartis, Roche, Pfizer, Sanofi, Amgen, Takeda, Regeneron and Kite. All other author have no conflicts of interest to disclose.

Contributions

Conception and design by XL, AB, HAL, PM and TF. Collection and assembly of data as well as data analysis and interpretation by all authors. Manuscript writing by XL, AB, JdK and SR.

Acknowledgments

The authors wish to thank the patients and their families. IFM (Intergroupe Francophone of Multiple Myeloma) project's team that conducted the study. A special thanks to the project manager Miss S Rollet from IFM. A special thanks to Pr Jean-Yves Mary who has always overviewed the study from its conception to its end. A special thanks to the members of the DMC, Pr Chantal Doyen (Belgique), Pr Jean-Pierre Marolleau (CHU Amiens, France) et Dr Marie-Christine Perrault-Pochat (CHU Poitiers, France). BMS provided pomalidomide and Takeda ixazomobid.

References

- D'agostino M, Cairns DA, Lahuerta JJ, et al. Second Revision of the International Staging System (R2-ISS) for overall survival in multiple myeloma: a European Myeloma Network (EMN) report within the HARMONY Project. J Clin Oncol. 2022;40(29):3406-3418.
- 2. Perrot A, Lauwers-Cances V, Tournay E, et al. Development and validation of a cytogenetic prognostic index predicting survival in multiple myeloma. J Clin Oncol. 2019;37(19):1657-1665.
- Leleu X, Karlin L, Macro M, et al. Pomalidomide plus low-dose dexamethasone in multiple myeloma with deletion 17p and/or translocation (4;14): IFM 2010-02 trial results. Blood. 2015;125(9):1411-1417.
- 4. Schmidt J, Braggio E, Kortuem KM, et al. Genome-wide studies in multiple myeloma identify XPO1/CRM1 as a critical target validated using the selective nuclear export inhibitor KPT-276. Leukemia. 2013;27(12):2357-2365.
- 5. Vincent Rajkumar S. Multiple myeloma: 2013 update on diagnosis, risk-stratification, and management. Am J Hematol. 2013;88(3):226-235.
- 6. Avet-Loiseau H, Magrangeas F, Moreau P, et al. Molecular heterogeneity of multiple myeloma: Pathogenesis, prognosis, and therapeutic implications. J Clin Oncol. 2011;29(14):1893-1897.
- Sonneveld P, Avet-Loiseau H, Lonial S, et al. Treatment of multiple myeloma with high-risk cytogenetics: a consensus of the International Myeloma Working Group. Blood. 2016;127(24):2955-2962.
- 8. Lachin JM. Biostatistical methods: the assessment of relative

Funding

The study was funded by IFM, Takeda and BMS.

Data-sharing statement

Data supporting this article are part of an ongoing clinical trial and are not publicly available. Data will be considered for sharing, with restriction due to data privacy regulations, and the informed consent. Requests for de-identified patient data by researchers with proposed use of the data can be made to the corresponding author with specific data needs, analysis plans and dissemination plans. Those requests will be reviewed by a study steering committee (IFM group) and the study sponsor for release upon publication. Response will typically be given in 3 months.

risks. 2nd ed. Wiley; 2014.

- 9. Reece D, Song KW, Fu T, et al. Influence of cytogenetics in patients with relapsed or refractory multiple myeloma treated with lenalidomide plus dexamethasone: adverse effect of deletion 17p13. Blood. 2009;114(3):522-525.
- 10. Dimopoulos MA, Moreau P, Terpos E, et al. Multiple myeloma: EHA-ESMO clinical practice guidelines for diagnosis, treatment and follow-up. Ann Oncol. 2021;32(3):309-322.
- Stong N, Ortiz-Estévez M, Towfic F, et al. The location of the t(4;14) translocation breakpoint within the NSD2 gene identifies a subset of patients with high-risk NDMM. Blood. 2023;141(13):1574-1583.
- 12. Delforge M, Bladé J, Dimopoulos MA, et al. Treatment-related peripheral neuropathy in multiple myeloma: the challenge continues. Lancet Oncol. 2010;11(11):1086-1095.
- Mateos MV, San Miguel JF. Safety and efficacy of subcutaneous formulation of bortezomib versus the conventional intravenous formulation in multiple myeloma. Ther Adv Hematol. 2012;3(2):117-124.
- 14. Chauhan D, Tian Z, Zhou B, et al. In vitro and in vivo selective antitumor activity of a novel orally bioavailable proteasome inhibitor MLN9708 against multiple myeloma cells. Clin Cancer Res. 2011;17(16):5311-5321.
- Moreau P, Masszi T, Grzasko N, et al. Oral ixazomib, lenalidomide, and dexamethasone for multiple myeloma. N Engl J Med. 2016;374(17):1621-1634.