

Survival Outcomes of Multiple Myeloma Patients Previously Exposed to BCMA-Targeted Therapies in the HONEUR European Network

Roman Hájek¹, Jiri Minarik², Martin Stork³, Alexandra Jungova⁴, Giselle Lostaunau Costa⁵, Yuwei Wang⁶, Solenn Salaun⁷, Nolen J Peruaila⁸, Guillaume Azarias⁹, Markus Rückert⁹

¹University Hospital Ostrava and Faculty of Medicine, University of Ostrava, Ostrava, Czech Republic; ²University Hospital Olomouc and Palacky University, Olomouc, Czech Republic; ³Hematology and Oncology, University Hospital Brno and Masaryk University, Brno, Czech Republic; ⁴Charles University Hospital Pilsen, Pilsen, Czech Republic; ⁵Johnson & Johnson-Cilag, Madrid, Spain; ⁶Johnson & Johnson, Breda, Netherlands; ⁷Johnson & Johnson, Allschwil, Switzerland; ⁸Johnson & Johnson, Beerse, Belgium; ⁹TriNetX Oncology GmbH, Freiburg, Germany

Key Takeaway



Despite new treatment advances, such as anti-BCMA therapy, this real-world analysis indicates that there is still an unmet need in heavily pretreated patients with MM

Conclusions



Real-world data from 3 European registries indicate that BCMA-exposed patients are heavily pretreated and have limited survival outcomes, with a median OS of 15.8 months and a median TTNT of 5.2 months



With a median of 5 prior lines, 93% were triple-class refractory and 54.5% were penta-drug refractory, highlighting significant treatment resistance within this relapsed/refractory MM population



The persistent prescription of chemotherapy and IMiD/PI triplets as subsequent treatment underscores a critical gap in treatment options and the need for new therapeutic targets to be adopted in clinical practice



Please scan QR code

<https://www.congresshub.com/ASH2025/Oncology/Talquetamab/Hajek>



Poster

The QR code is intended to provide scientific information for individual reference, and the information should not be altered or reproduced in any way.

Acknowledgments

We thank the patients who participated in the study and their families and caregivers, and the physicians, nurses, and staff members involved in data collection and analyses. This study was funded by Johnson & Johnson Innovative Medicine EMEA. Medical writing support was provided by Teja Rus, PhD, of Eloquent, part of Envision Spark, an Envision Medical Communications agency, a part of Envision Pharma Group, and funded by Johnson & Johnson Innovative Medicine EMEA.

Disclosures

RH reports a consulting/advisory role for AbbVie, Amgen, BMS, Celgene, Johnson & Johnson, Novartis, PharmaMar, and Takeda; honoraria from Amgen/BMS, Celgene, Johnson & Johnson, PharmaMar, and Takeda; and research funding from Amgen, BMS, Celgene, Johnson & Johnson, Novartis, and Takeda.

Introduction

- Survival rates for patients with multiple myeloma (MM) have recently improved,^{1,2} thanks to advances in treatment options that target B-cell maturation antigen (BCMA) receptors, including antibody-drug conjugates (ADCs), bispecific antibodies (BsAbs), and chimeric antigen receptor (CAR)-T cell therapies¹⁻⁴
- However, this progress introduces a new population of patients with disease refractory to both standard-of-care and novel BCMA therapies, thereby limiting treatment options at relapse
- Therefore, understanding real-world outcomes of BCMA-exposed patients is essential

Methods

- Data from 3 MM European registries were included in the analysis: IUCT (France), TriNetX (Germany), and RMG (Czech Republic) (**Figure 1**)
- Locally collected patient-level data were analyzed uniformly and site-specific results were aggregated centrally using the HONEUR federated data network

Results

Study population

- A total of 242 patients with MM who had received an anti-BCMA–based regimen were analyzed from 3 European registries
 - Most cases came from IUCT (82.2%), with TriNetX contributing to 10.3% of cases and RMG contributing to 7.4%
- Baseline characteristics are shown in **Table 1**
 - Most patients were ≥65 years of age
 - Half of patients (51.2%) were ISS stage II/III, and 23.6% presented with high-risk cytogenetics
 - Most patients (70.2%) had previously undergone stem cell transplant (SCT)

Table 1: Baseline patient characteristics

Characteristics, n (%)	N=242
Sex	
Female	115 (47.5)
Male	127 (52.5)
Age at line of treatment initiation, years	
Median (range)	68 (43–88)
≤64 years	78 (32.2)
65–74 years	109 (45)
≥75 years	55 (22.7)
ISS stage ^a	
I	35 (14.5)
II	68 (28.1)
III	56 (23.1)
Unavailable	83 (34.3)
M protein ^a	
Non-IgG positive	50 (20.7)
IgG positive	88 (36.4)
Unavailable	104 (43)
Cytogenetic risk ^{a,b}	
High risk	57 (23.6)
Standard risk	122 (50.4)
Unavailable	63 (26)
Prior line received SCT	
No	72 (29.8)
Yes	170 (70.2)
Prior lines	
Median (range)	5 (3–12)
3 or 4 lines	73 (30.2)
>4 lines	169 (69.8)

^aLOCF values used. ^bHigh risk defined as any presence of del(17p), and/or t(4;14), and/or t(14;16). Standard risk=no high risk. Ig, immunoglobulin.

References

- National Comprehensive Cancer Network (NCCN). Multiple Myeloma. (Version 2.2026).
- Jagannath S, et al. *J Clin Oncol* 2025;43:2766-71.
- Tan CR, et al. *Blood Cancer J* 2025;15:53.
- Zheng H, et al. *J Hematol Oncol* 2025;18:23.

- Inclusion criteria:
 - Patients diagnosed with MM
 - At least 18 years of age at start of frontline treatment
 - Received ≥3 prior lines
 - Quad-class exposed (received at least 1 proteasome inhibitor [PI], 1 immunomodulatory drug [IMiD], 1 anti-CD38 antibody, and 1 BCMA-targeted therapy)
 - First eligible line started ≥2020
- Exclusion criteria: Patients receiving retreatment with anti-BCMA immunotherapy were excluded from the analysis
- Statistical analyses:
 - Patient characteristics and treatment patterns were descriptively analyzed
 - Index date was considered as the date of initiating the subsequent therapy after anti-BCMA treatment
 - Time to next treatment (TTNT), used as a proxy for progression-free survival, and overall survival (OS) were analyzed using the Kaplan-Meier method
 - The last observation carried forward (LOCF) method was employed for the International Staging System (ISS), M protein, and cytogenetics data

Baseline treatment patterns

- With a median of 5 prior lines of treatment, only 30.2% of patients received 3 or 4 prior lines of treatment (**Table 1**)
- Regarding refractoriness, 93% of patients were triple-class refractory and 54.5% were penta-drug refractory, highlighting the significant treatment resistance within this relapsed/refractory MM population observed in real-world clinical practice (**Table 2**)
- At BCMA exposure, data reflect the real-world clinical setting, with low CAR-T cell therapy utilization (**Table 2**)

Table 2: Baseline treatment characteristics

Characteristics, n (%)	N=242
Exposed to 2 PIs, 2 IMiDs, and 1 anti-CD38 antibody	
No	66 (27.3)
Yes	176 (72.7)
Exposed to CAR-T	
No	223 (92.1)
Yes	19 (7.9)
Exposed to BsAb	
No	140 (57.9)
Yes	102 (42.1)
Exposed to ADC	
No	119 (49.2)
Yes	123 (50.8)
Refractory status	
Triple-refractory ^a	16 (6.6)
Quad-refractory ^b	65 (26.9)
Penta-refractory ^c	132 (54.5)
Other	29 (12.0)
Triple-class refractory	
No	17 (7.0)
Yes	225 (93.0)

^aRefractory to 1 IMiD, 1 PI, and 1 anti-CD38 mAb. ^bRefractory to ≥2 IMiDs, 1 PI, and 1 anti-CD38 mAb or ≥2 PIs, 1 IMiD, and 1 anti-CD38 mAb. ^cRefractory to ≥2 IMiDs, ≥2 PIs, and 1 anti-CD38 mAb. mAb, monoclonal antibody.

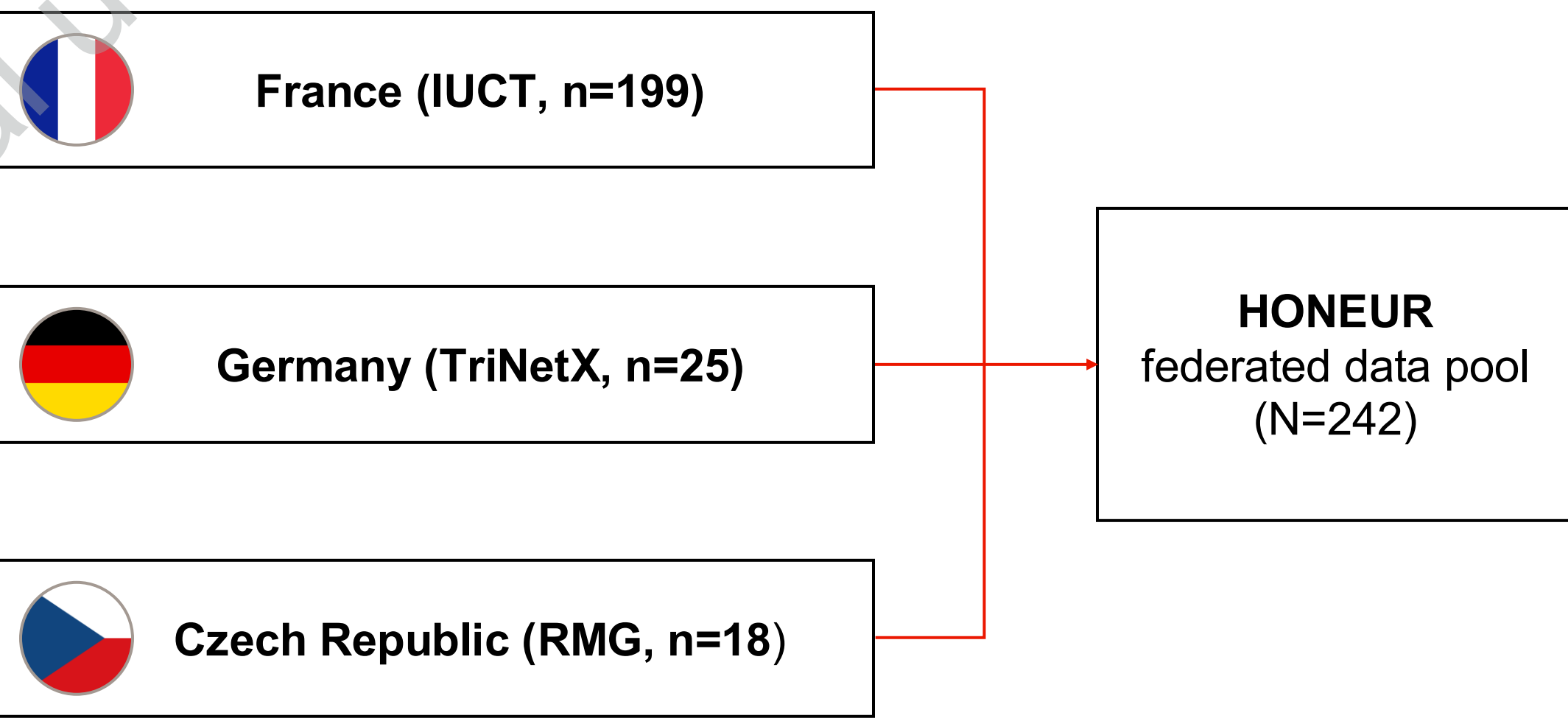
Post-BCMA treatment

- Following BCMA exposure, the available treatment regimens predominantly included chemotherapy doublet (34.7%) IMiD/PI-based triplet (17.8%), IMiD/PI-based doublet (12.4%), and 4.1% anti-CD38–based triplet (**Table 3**)

Efficacy outcomes

- With a median follow-up of 9.1 months, median OS was 15.8 months (95% CI, 10.91–31.08 months) (**Figure 2**), and median TTNT was 5.2 months (95% CI, 3.98–6.24 months) (**Figure 3**)

Figure 1: European registries included in analysis



HONEUR, Haematology Outcomes Network in Europe; IUCT, Institut Universitaire du Cancer de Toulouse; RMG, The Registry of Monoclonal Gammopathies.

Table 3: Post-BCMA treatment

Drug combination, n (%)	N=242
Chemotherapy doublet ^a	84 (34.7)
PI/IMiDs doublet ^b	30 (12.4)
PI/IMiDs triplet ^c	43 (17.8)
CD38 triplet ^d	10 (4.1)
Unknown or Others	75 (31.0)

^aCyclophosphamide/dexamethasone; bendamustine/dexamethasone; melphalan/prednisone; vincristine/dexamethasone; cyclophosphamide/doxorubicin.
^bBortezomib/dexamethasone; carfilzomib/dexamethasone; pomalidomide/dexamethasone; thalidomide/dexamethasone.
^cCarfilzomib/cyclophosphamide/dexamethasone; bortezomib/cyclophosphamide/dexamethasone; bortezomib/melphalan/prednisone; ixazomib/cyclophosphamide/dexamethasone; ixazomib/lenalidomide/dexamethasone; cyclophosphamide/pomalidomide/dexamethasone; cyclophosphamide/thalidomide/dexamethasone; elotuzumab/pomalidomide/dexamethasone; ixazomib/pomalidomide/dexamethasone; melphalan/pomalidomide/dexamethasone; carfilzomib/lenalidomide/dexamethasone; bortezomib/lenalidomide/dexamethasone; selinexor/bortezomib/dexamethasone; bortezomib/cyclophosphamide/thalidomide; bortezomib/pomalidomide/dexamethasone.
^dDaratumumab/pomalidomide/dexamethasone; daratumumab/lenalidomide/dexamethasone; daratumumab/carfilzomib/dexamethasone; isatuximab/pomalidomide/dexamethasone; isatuximab/thalidomide/dexamethasone; isatuximab/carfilzomib/dexamethasone.

Figure 2: OS first eligible lines

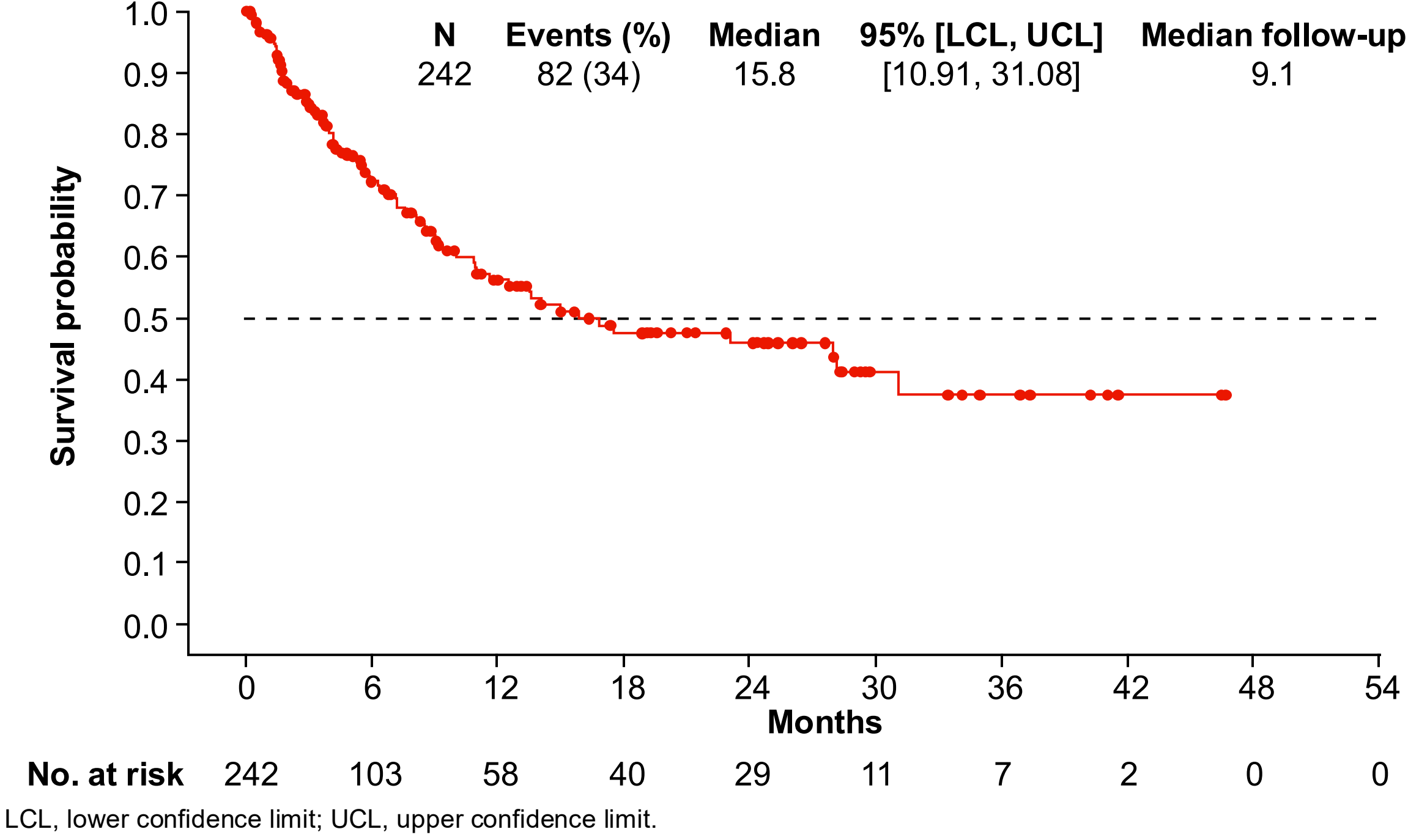
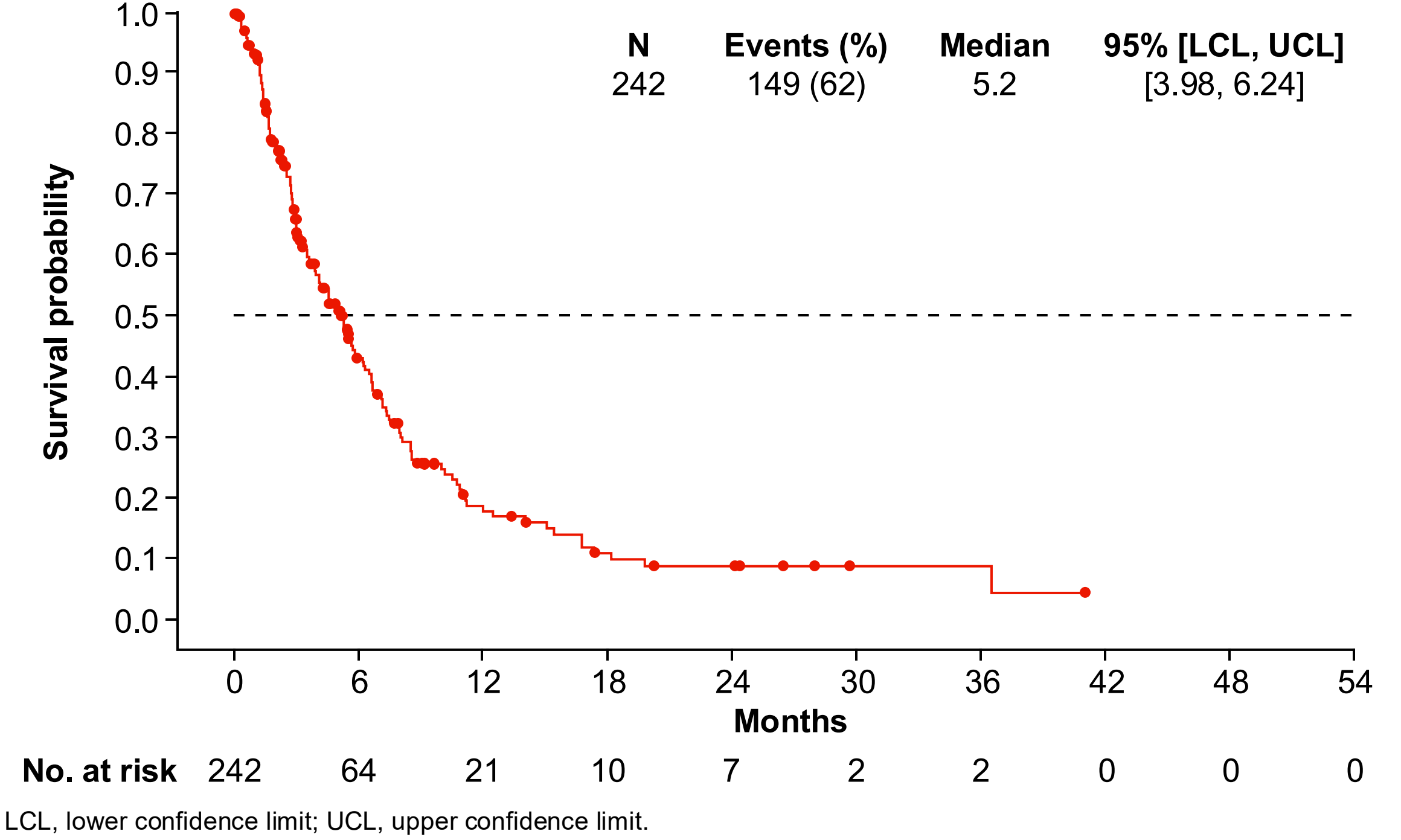


Figure 3: TTNT first eligible lines



Multiple Myeloma

