



Driver alterations in *FBXW7*, *PPP2R1A*, *CCNE1* confer a poor prognosis in patients with metastatic gynecologic cancers

Alison M. Schram¹, Elizabeth K. Lee², Ying L. Liu¹, Yi Xu³, Julia Yang³, Sunantha Sethuraman³, Paul Basciano³, Maria Koehler³, Ian M. Silverman³

¹Medical Oncology, Memorial Sloan Kettering Cancer Center, New York, NY, USA; ²Medical Oncology, Dana-Farber Cancer Institute, Boston, MA, USA; ³Repare Therapeutics Inc., Cambridge, MA, USA

Introduction

- The combination of lunresertib (Lunre), a PKMYT1 inhibitor, and camonsertib, an ATR inhibitor, is currently being studied in patients harboring lunre-sensitizing biomarkers (Lunre BM) including *CCNE1* amplifications or mutations in *FBXW7* or *PPP2R1A* in the phase I MYTHIC trial (NCT04855656)
- CCNE1* amplifications, which occur in ~30% of platinum-resistant ovarian cancers,¹⁻³ are well established as a poor prognostic indicator in ovarian cancer,⁴⁻⁷ but little is known about other Lunre BM in ovarian and endometrial cancers

Key translational questions

- What is the distribution of Lunre BM in ovarian and endometrial cancers?
- What is the prognosis for gynecological malignancies with Lunre BM?
- Is presence of a Lunre BM independently prognostic or associated with other poor outcomes indicators?

Lunre-sensitizing alterations are enriched in gynecological malignancies

- Lunre-sensitizing alterations are most highly enriched in gynecological malignancies (Figure 1A)
- In MYTHIC, the distribution of individual biomarkers enrolled within gynecological malignancies was consistent with prevalence estimates from TCGA¹ (Figure 1B)

Figure 1: (A) Estimated prevalence of Lunre BM from TCGA. (B) Distribution of enrollment biomarkers in ovarian and endometrial patients enrolled in the MYTHIC trial. (C) Distribution of histologies in endometrial patients enrolled in the MYTHIC trial. *Includes only high-grade serous ovarian patients. †Uterine Endometrial Carcinoma and Uterine Mixed Endometrial Carcinoma ‡Soft-tissue sarcoma only. §Squamous histology of non-small cell lung cancer only.

Table 1. *CCNE1* amplification frequency in ovarian cancer

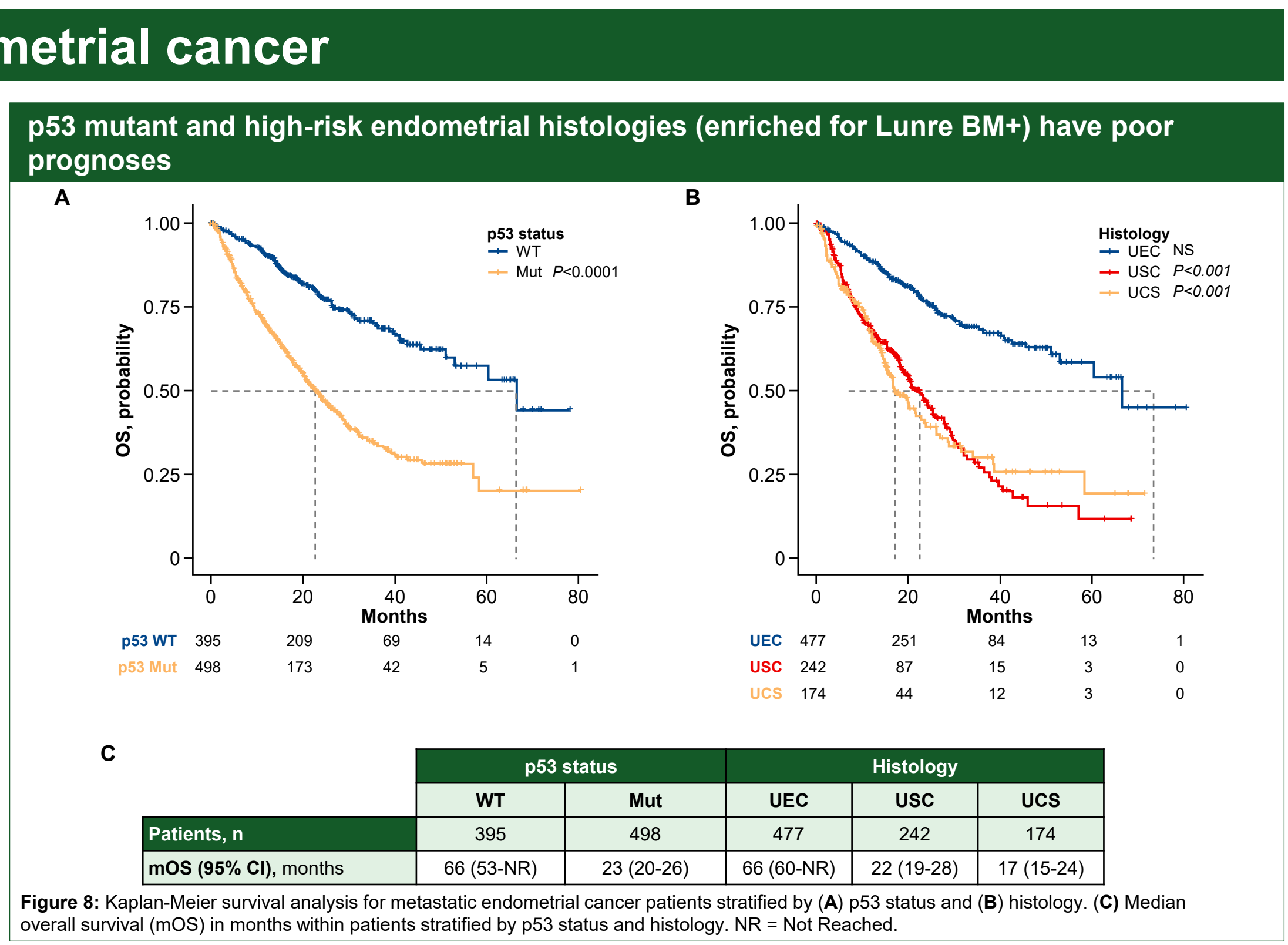
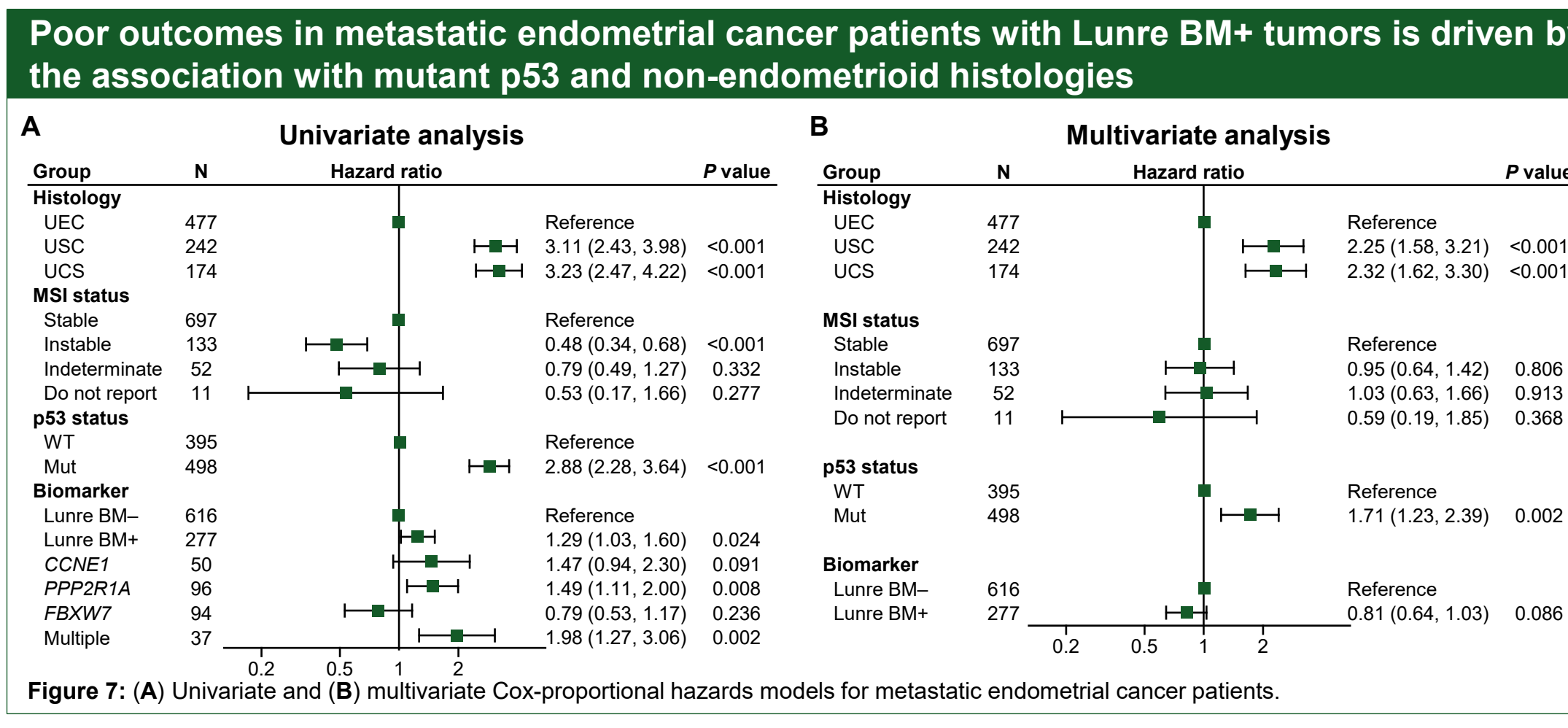
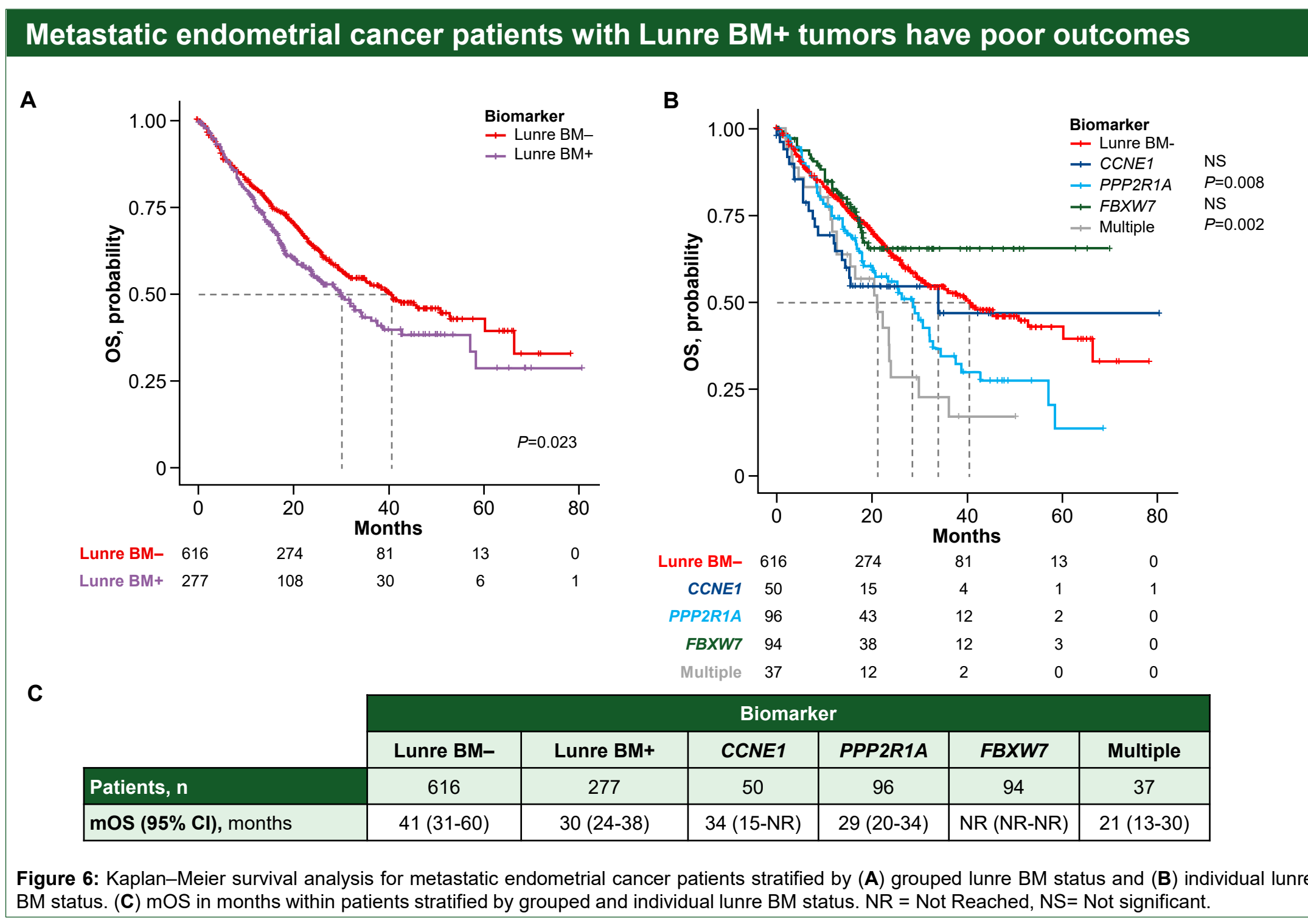
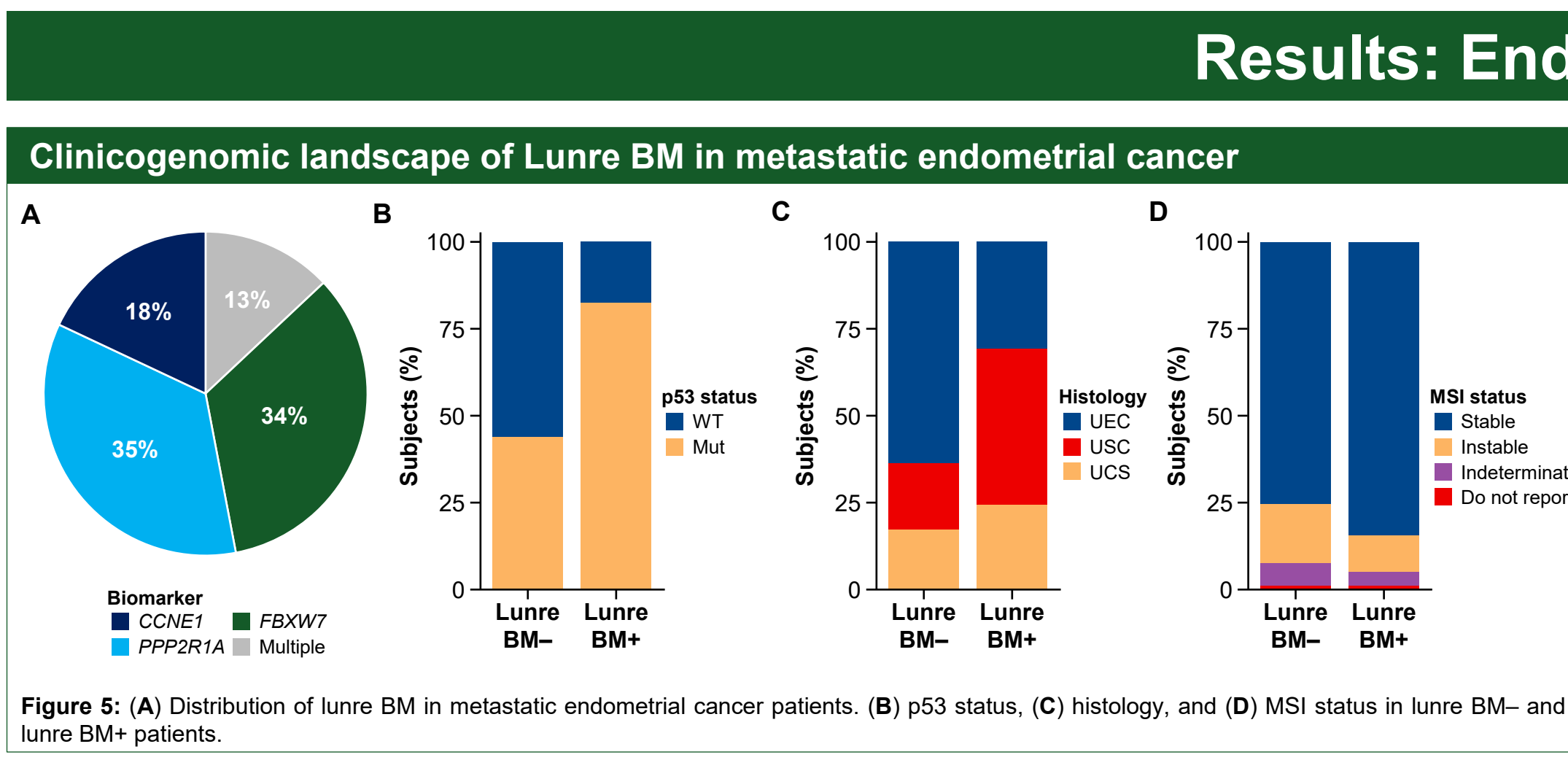
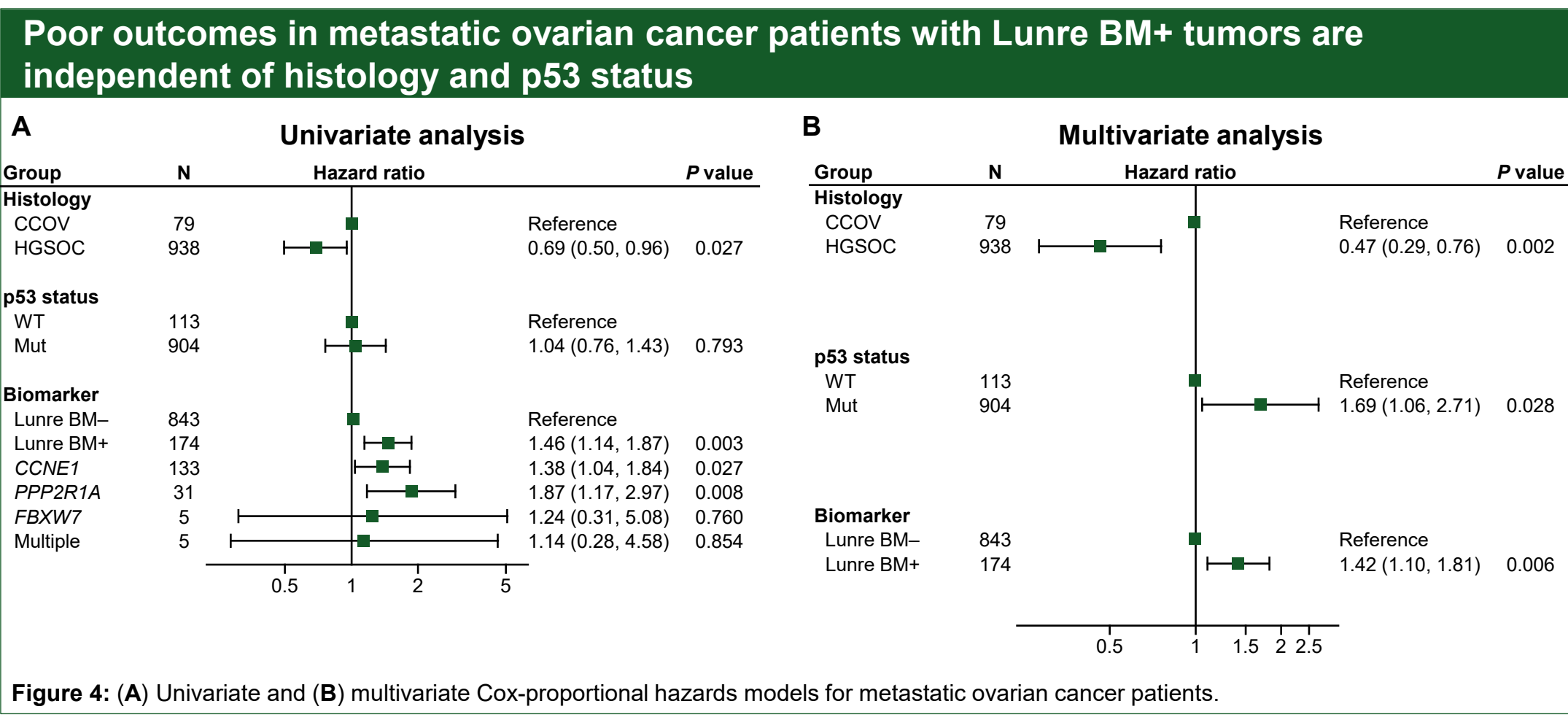
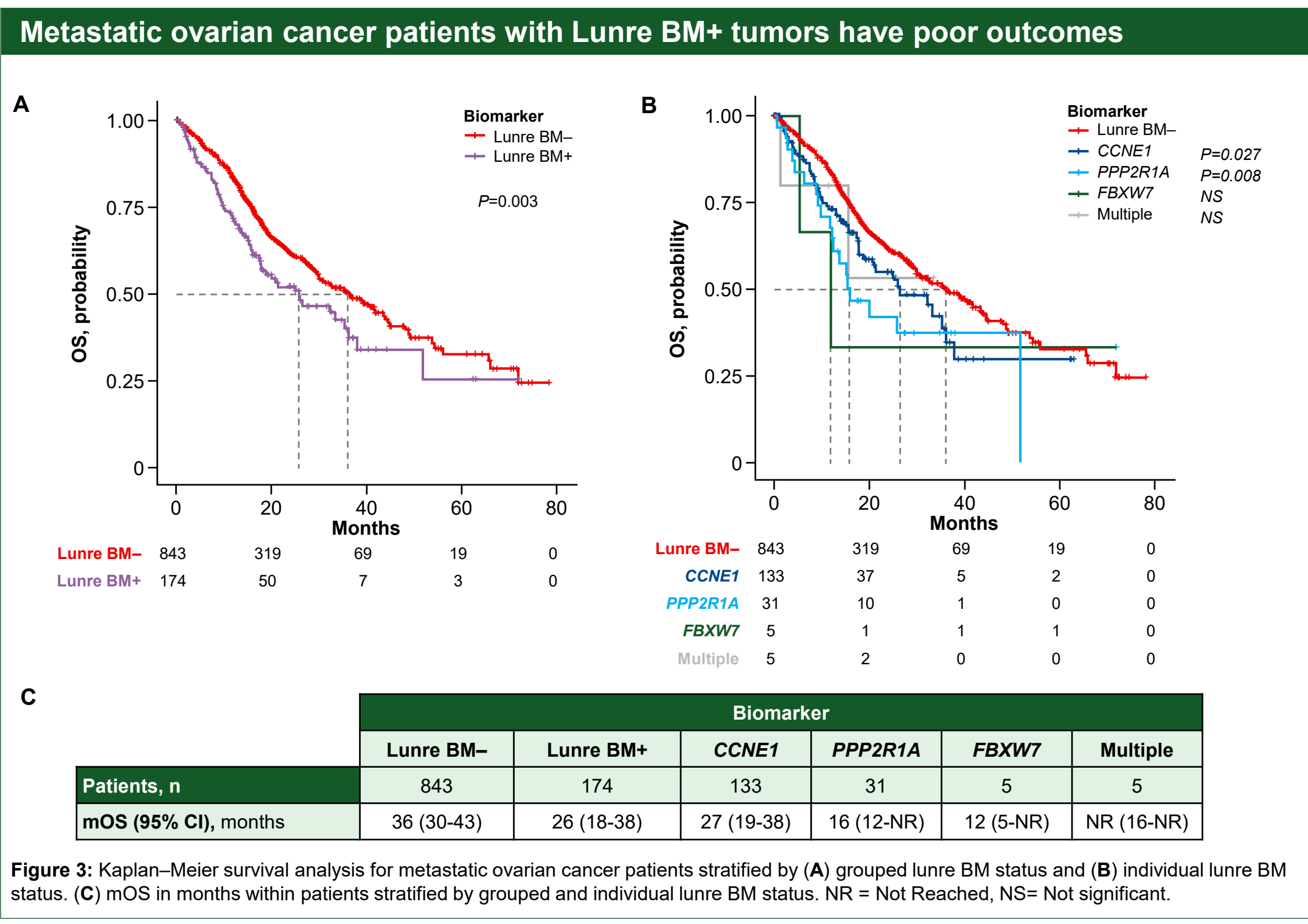
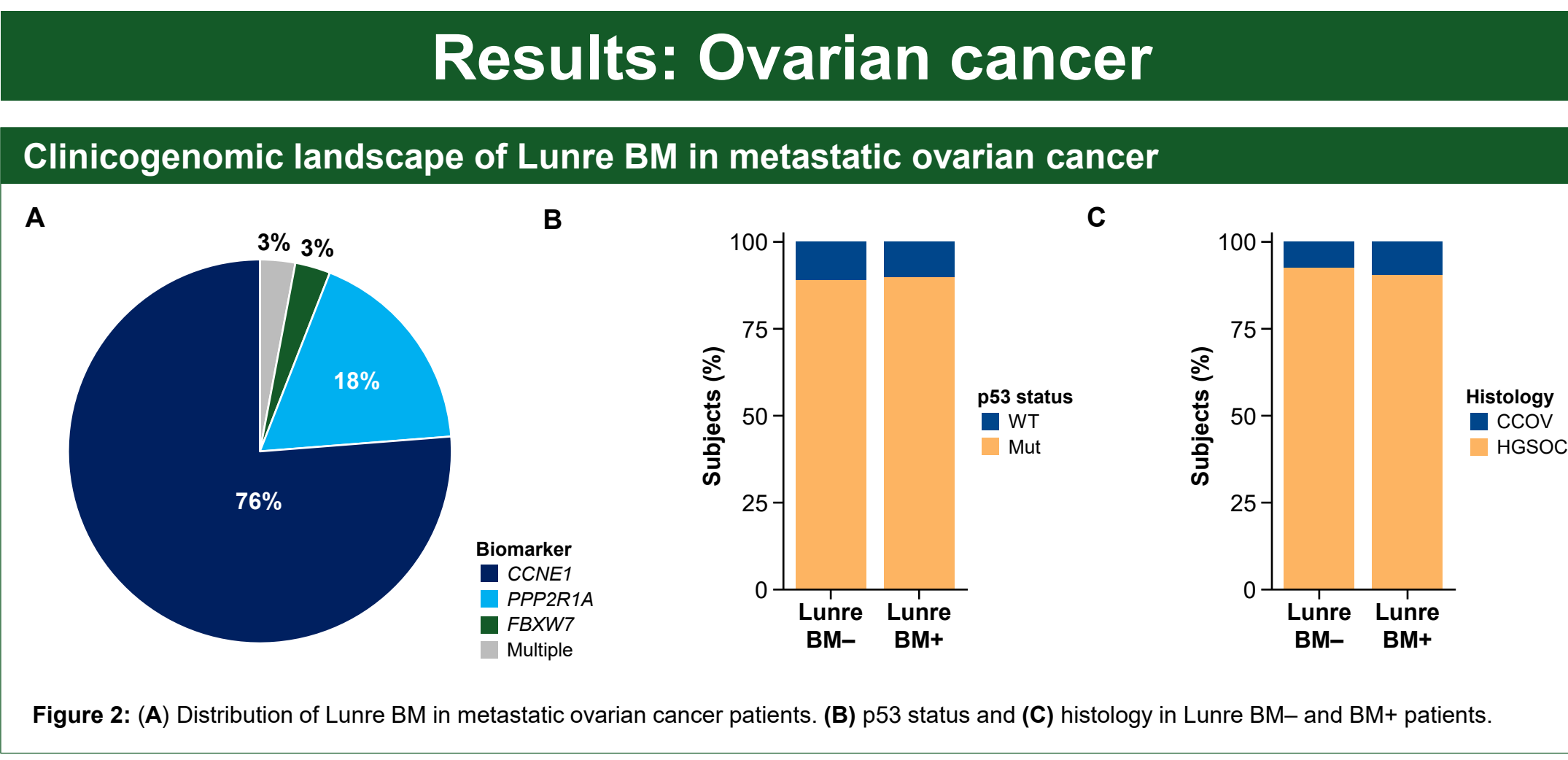
Platinum status	Refractory ^a	Resistant ^b	Sensitive
TCGA. <i>Nature</i> . 2011 ¹		27.6% (n=58)	13.1% (n=130)
Patch A-M, et al. <i>Nature</i> . 2015 ²	41% (n=12)	29.7% (n=37)	6.5% (n=31)
Smith P, et al. <i>Nat Commun</i> . 2023 ³	58% (n=12)		8.8% (n=114)

^aRefractory: Defined as disease progression while on primary platinum treatment or within one month of end of primary platinum treatment. ^bResistant: Defined as a platinum-free interval of ≥6 months.

Methods

Analysis of MSK MetTropism^{8,9} dataset

- Clinical and genomic data were obtained from the MSK MetTropism^{8,9}
- Patients with metastatic endometrial cancer (UEC, n=478; USC, n=243; UCS, n=174), HGSOE (n=949), and CCOV (n=80) were included (Figures 2A and 5A)
- Overall survival (OS) was measured from the time of NGS testing to death and was censored at the last time reported alive
- OS was modeled on Lunre BM status alone or concurrently with p53 status, tumor histology, and MSI status (endometrial only) using multivariate Cox proportional-hazards models



Potential limitations of this analysis include:

- Analysis of data from a single academic center is subject to center-specific biases
- CCNE1* amplification copy number thresholds from MSK-IMPACT[®] may differ from other commonly used NGS assays
- Important ovarian cancer co-variables, such as platinum sensitivity and HRD status, were not available and likely impact prevalence and clinical outcomes
- Survival analysis was anchored on NGS date; date of diagnosis or relapse may more accurately reflect true prognosis

Conclusions

- In both endometrial and ovarian cancers, Lunre BM+ tumors had worse prognoses compared to Lunre BM- tumors
- Lunre BM+ status was independently associated with poor prognosis in ovarian cancer
- In endometrial cancers, Lunre BM+ status was indirectly associated with poor prognosis due to the enrichment for patients with high-risk histologies (UCS and USC) and genotype (p53 mutant)
- MSI was largely distinct from the Lunre BM+ population and was associated with better prognosis due to its enrichment in UEC
- Additional treatment solutions are a critical unmet need for this very high-risk population with adverse outcomes

References

- Cancer Genome Atlas Research Network. *Nature*. 2011;474(7353):609-615.
- Patch A-M, et al. *Nature*. 2015;521(7553):489-494.
- Smith P, et al. *Nat Commun*. 2023;14(1):4387.
- Chan AM, et al. *J Pathol Clin Res*. 2020;6(4):252-262.
- Kang E-Y, et al. *Cancer*. 2023;129(5):697-713.
- Nakayama N, et al. *Cancer*. 2010;116(11):2621-2634.
- Stronach EA, et al. *Mol Cancer Res*. 2018;16(7):1103-1111.
- Nguyen B, et al. *Cell*. 2022;185(3):563-575.e11.

Acknowledgments and Disclosures

Alison M. Schram reports advisory board compensation from Repare Therapeutics and Merck. Elizabeth K. Lee has received research funding from Merck and consulting funding from Aadi Biosciences. Ying L. Liu reports research funding from AstraZeneca, GSK, and Repare Therapeutics and advisory board for Myriad Laboratory, Yi Xu, Sunantha Sethuraman, Paul Basciano, Maria Koehler, and Ian M. Silverman are employees of Repare Therapeutics and may hold stock and/or stock options. Funding for this analysis and poster including medical writing support from Justin L. Eddy, Ph.D., of Repare Therapeutics, and Onyx, a Prime Global Company, including editorial support from Rosie Henderson and figure redraws by was funded by Repare Therapeutics.

Abbreviations

ATR, ataxia telangiectasia and Rad-3 related inhibitor; CI, confidence interval; CN, copy number; HGSOE, high-grade serous ovarian carcinoma; HRD, homologous recombination deficient; Lunre, lunresertib; Lunre BM, lunre-sensitizing biomarkers; MSI, microsatellite instability; MSK-IMPACT, Memorial Sloan Kettering - Integrated Mutation Profiling of Actionable Cancer Targets; MSK-MetTropism, Memorial Sloan Kettering - Metastatic Events and Tropisms; mOS, median overall survival; mut, mutated; MYTHIC, PKMYT1 inhibition for the treatment of Cancers; NGS, next-generation sequencing; NR, not reached; CCOV, ovarian clear cell carcinoma; OS, overall survival; PKMYT1, membrane-associated tyrosine- and threonine-specific, Cdc2-inhibitory kinase; strata, stratified; UCS, uterine carcinosarcoma; UEC, uterine endometrial cancer; USC, uterine serous carcinoma; WT, wild-type.

Contact: ssethuraman@reparex.com