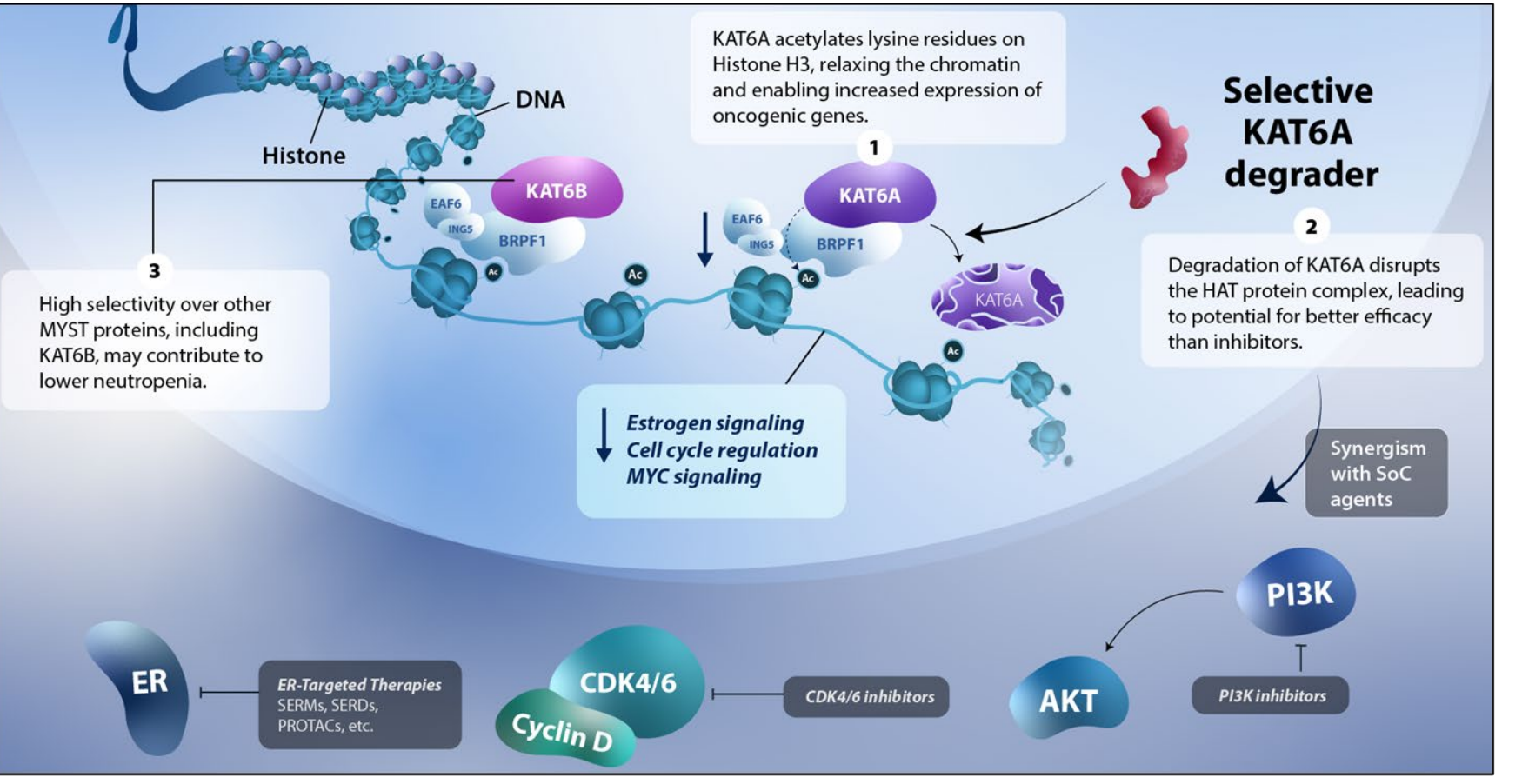


First-in-class potent and selective oral KAT6A degrader development candidate, PRT13722, drives complete tumor regressions as a monotherapy with an improved preclinical hematological safety profile

Monisha Sivakumar¹, Sarah Pawley¹, Corey Basch¹, Jimin Park¹, Justin Kurian¹, Anthony Reichelderfer¹, Yue Zou¹, Kirsten Gallagher¹, Miles Cowart¹, Joy Cote¹, Alexander Grego¹, Jessica Burtell¹, Amy Crossan¹, Michael Hulse¹, Anjana Agarwal¹, Arpita Mondal¹, Chun Chen¹, Vijay Devannah¹, Sina Rezazadeh¹, Quincy Lewis¹, Patrick Wen¹, Ken Ray¹, Raul Leal¹, Daniel Porreca¹, Ganfeng Cao¹, Neha Bhagwat¹, Shanthi Ganesan¹, Stefan Ruepp¹, Min Wang¹, Joseph Rager¹, Koichi Ito¹, Sandy Geeganage¹, Andrew Combs¹, Peggy Scherle¹, Andrew Buesking¹, Jack Carter¹

¹Prelude Therapeutics Incorporated, Wilmington, DE; contact: jdcarter@preludetx.com

KAT6A Degradation As a Novel Therapeutic Modality for Cancer



- MYST proteins like KAT6A, its paralog KAT6B, and KAT7 are histone acetyltransferases (HAT) that epigenetically regulate chromatin accessibility.^{1,2}
- KAT6A is highly expressed in breast cancer and is associated with cancer growth¹⁻² via catalytic and non-catalytic functions.⁴⁻⁵
- Dual KAT6A/B inhibition is a clinically-validated approach, achieving promising efficacy in ER+/HER2- breast cancer patients and currently under Phase 3 clinical evaluation, albeit with on-target safety considerations like neutropenia.^{3,8}
- KAT6A is the dominant cancer-driving paralog, while MYST proteins co-regulate normal hematopoiesis, which may have safety implications for inhibitors co-targeting KAT6B/KAT7.⁶
- KAT6A, mutually exclusive from KAT6B, interacts with a HAT complex to enhance and define its gene regulation, and expression of complex proteins correlate with KAT6A cancer dependency.^{1,7}
- Unlike the expanding field of dual KAT6A/B and triple KAT6A/B/KAT7 inhibitors, targeted protein degradation is a highly differentiated approach to improve KAT6A selectivity while disrupting the KAT6A protein complex to drive deeper pathway suppression and efficacy.⁷
- PRT13722, a KAT6A-selective degrader development candidate, is shown herein and demonstrates differentiated preclinical efficacy and safety when compared to a dual KAT6A/B inhibitor, prifetrastat.

Results - In Vitro

Assay	PRT13722	prifetrastat
KAT6A HiBIT nM DC ₅₀	0.3	-
KAT6B HiBIT nM DC ₅₀ (D _{max})	>999 (<18%)	-
KAT5, KAT7, KAT8 degradation nM DC ₅₀ (D _{max})	>999 (<22%)	-
T47-D CTG nM EC ₅₀ (E _{max}) ^a	0.3 (95%)	0.7 (61%)
T47-D <i>in vivo</i> efficacy (TGI at 1 mg/kg, PO, QD; Day 42)	102%	76%
Mouse plasma C _{avg} μM at similar TGI ^b	0.09	5.5
Mouse, rat, dog, cyno F% ^c	69, 23, 38, 32	-
CYP inhibition panel, hERG inhibition (μM)	All >30, >30	-

Figure 1. PRT13722 is a Potent and Highly Selective KAT6A Degradation

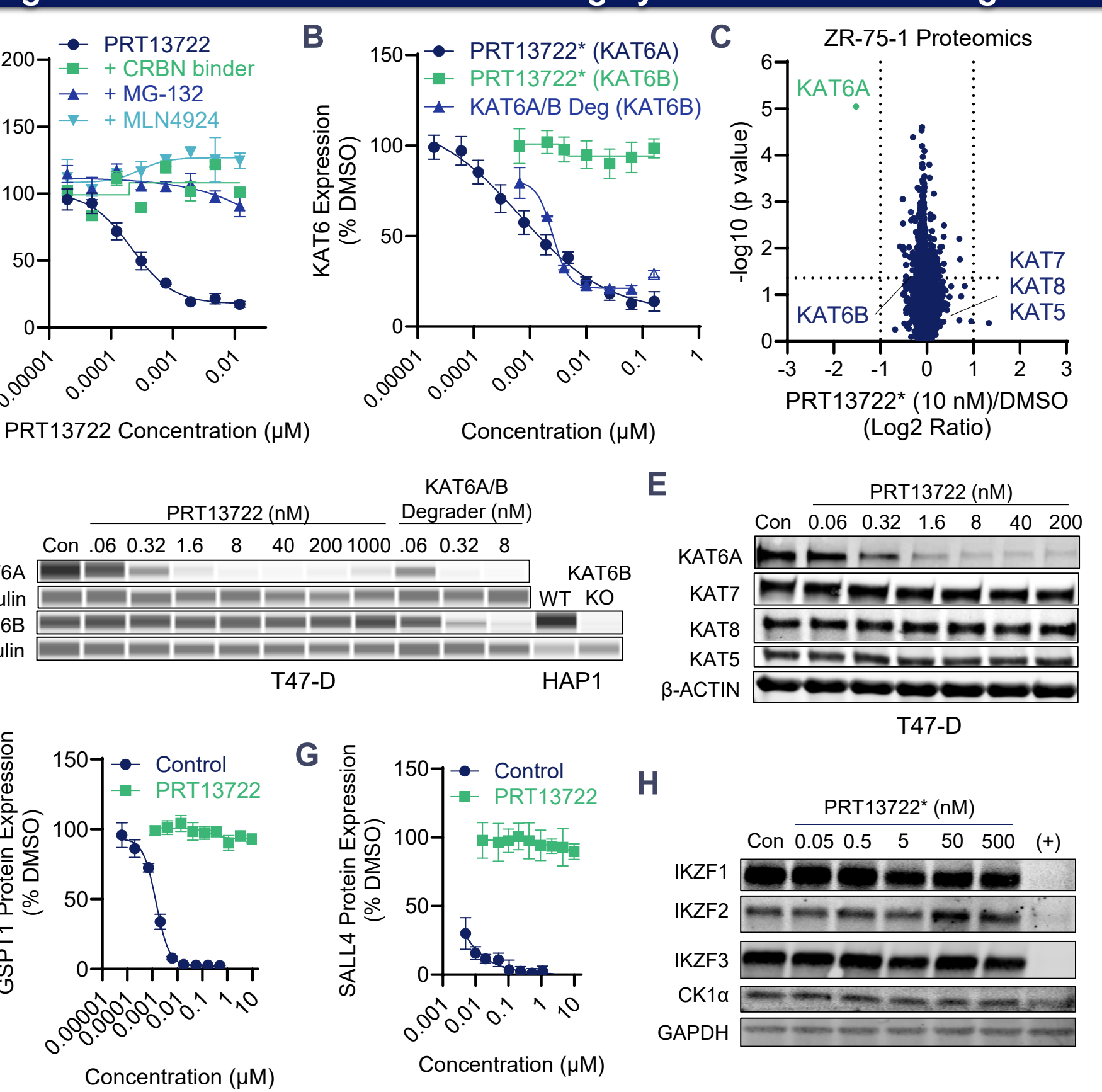


Figure 1. (A) KAT6A expression in KAT6A HiBIT cells treated with PRT13722 alone or in competition with a CRBN binder, proteasome inhibitor (MG-132), or neddylation inhibitor (MLN4924). (B) Expression of KAT6A or KAT6B in KAT6A or KAT6B HiBIT cells treated with PRT13722 or a KAT6A/B degrader for 24 h. (C) Global proteomics in ZR-75-1 cells treated with PRT13722 or a KAT6A/B degrader for 24 h. (D) T47-D cells treated with PRT13722 or a KAT6A/B degrader for 24 h. (E) Western blot of related MYST proteins in T47-D cells following 24 h treatment with PRT13722. (F) SPT1 expression assessed via Western blot after treatment with PRT13722 or CC-885 for 24 h. (G) SALL4 expression assessed via Western blot after treatment with PRT13722 or CC-885 for 24 h. (H) Expression of additional CRBN neoblasts after treatment with PRT13722 for 24 h, compared to a positive control (+) (CC-885, Pomalidomide).

Figure 2. PRT13722 Drives Superior Depth and Breadth of Anti-Cancer Activity in HR+/HER2- Breast Cancer Compared to prifetrastat

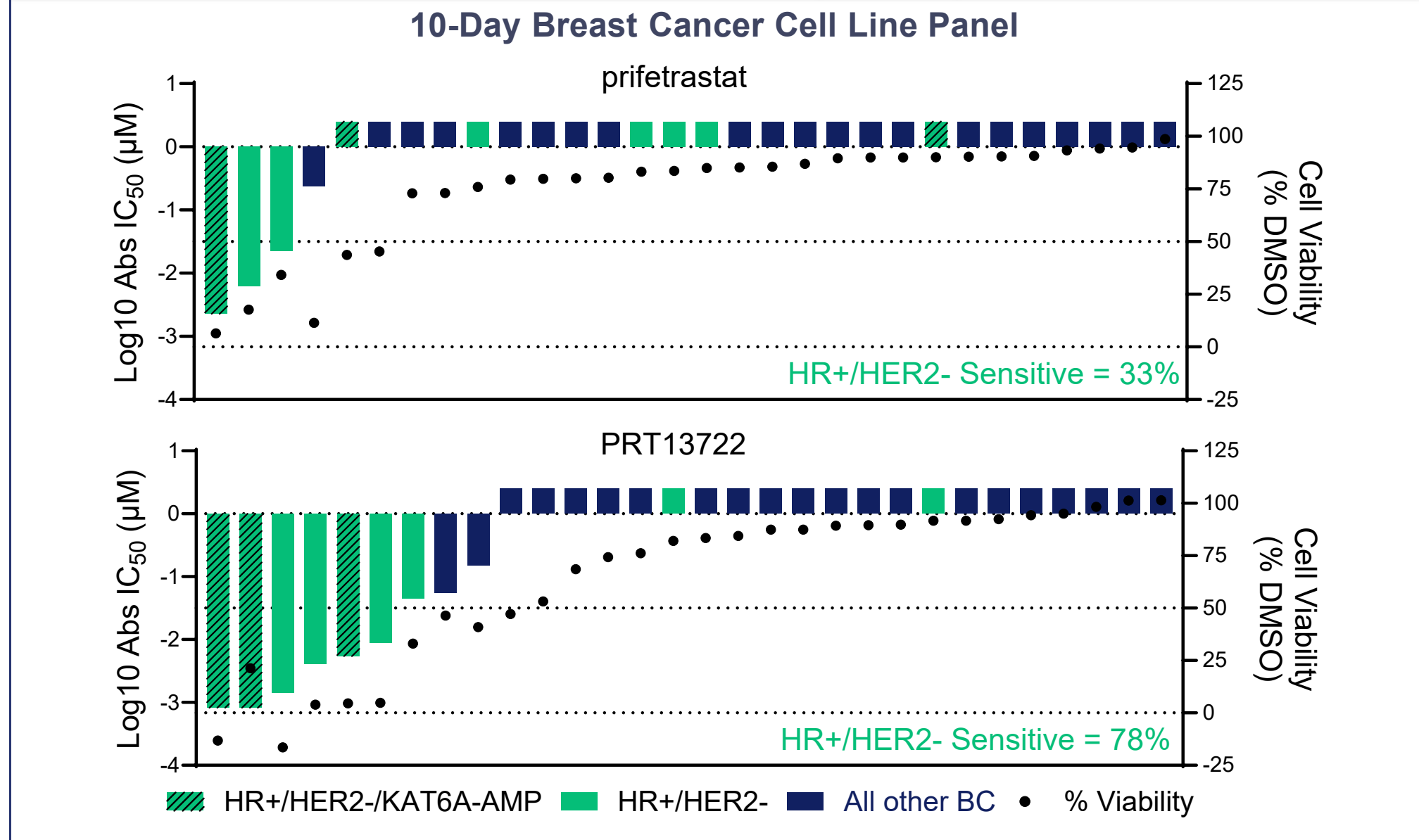


Figure 2. A panel of 29 breast cancer cell lines were assessed for anti-proliferative activity of prifetrastat and PRT13722 head-to-head in 10-day CTG assays up to 2.5 μM. Bars represent potency and black dots represent cell viability reduction.

Figure 3. PRT13722 Has Potent Anti-Cancer Activity and Synergy with Endocrine and CDK4/6-Targeted Therapies in Breast Cancer Cells

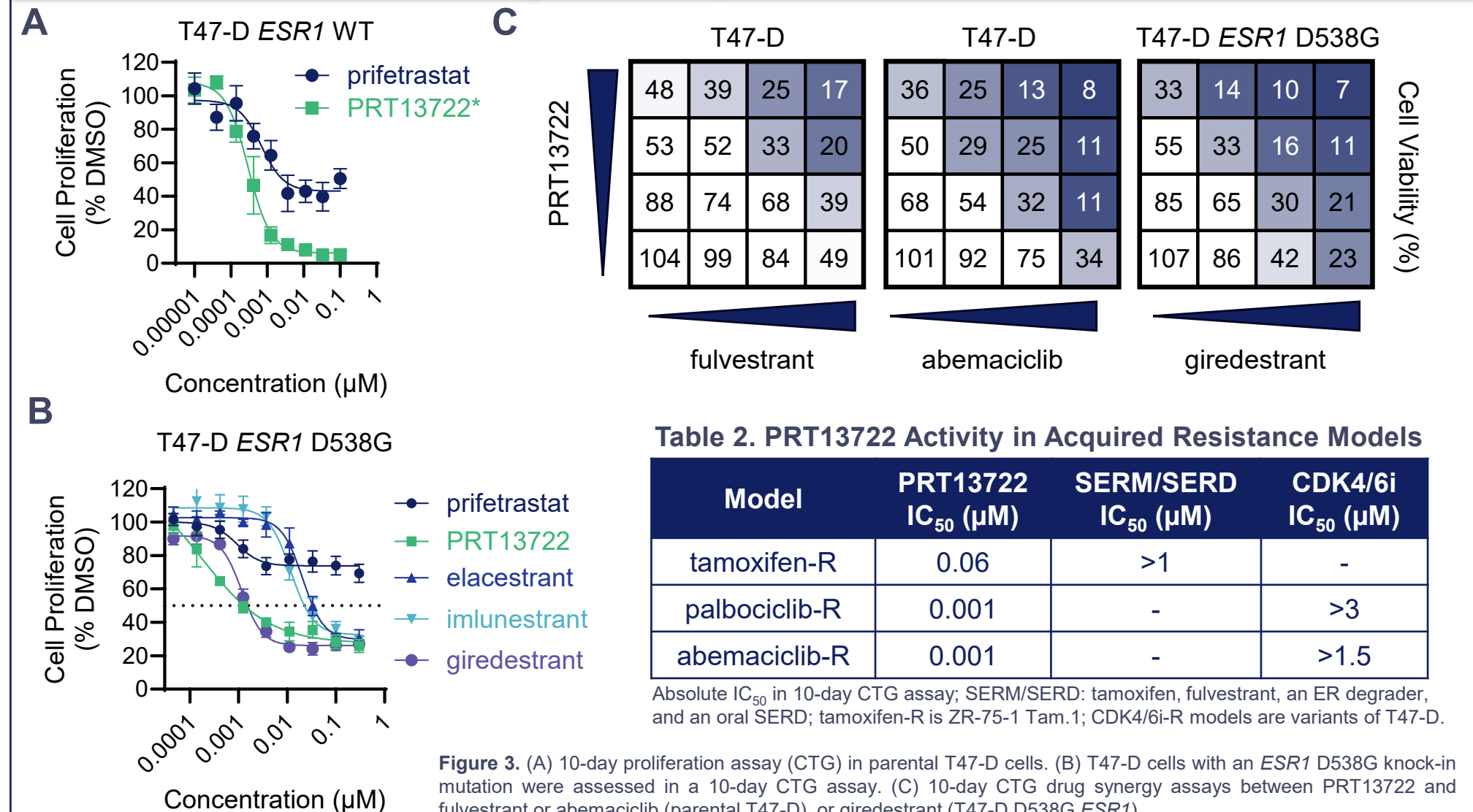


Figure 4. PRT13722, But Not KAT6A/B Inhibitors, Disrupt KAT6A Complexes, Inducing Deeper Molecular Perturbations in Breast Cancer Cells

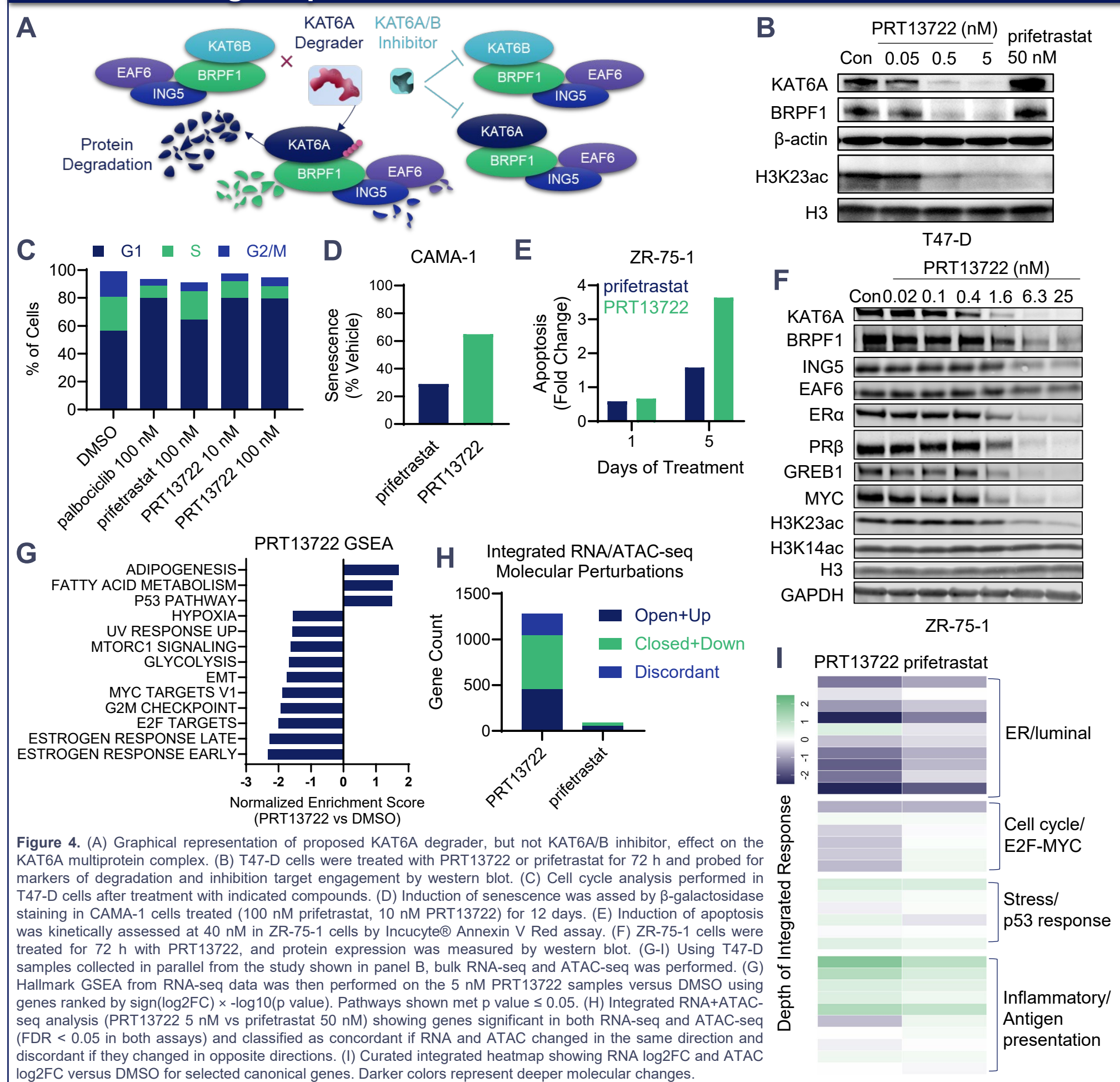


Figure 4. (A) Graphical representation of proposed KAT6A degrader, but not KAT6A/B inhibitor, effect on the KAT6A multiprotein complex. (B) T47-D cells were treated with PRT13722 or prifetrastat for 72 h and probed for markers of degradation and inhibition target engagement by western blot. (C) Cell cycle analysis performed in T47-D cells after treatment with indicated compounds. (D) Induction of apoptosis was assessed by β-galactosidase staining in CAMA-1 cells treated (100 nM prifetrastat, 10 nM PRT13722) for 12 days. (E) Induction of apoptosis was kinetically assessed at 40 nM in ZR-75-1 cells by Incubycell Annexin V Red assay. (F) ZR-75-1 cells were treated for 72 h with PRT13722, and protein expression was measured by western blot. (G-H) Using T47-D samples collected in parallel from the study shown in panel B, bulk RNA-seq and ATAC-seq was performed. (G) Hallmark GSEA from RNA-seq data was then performed on the 5 nM PRT13722 samples versus DMSO using genes ranked by signal(FDR) > -log10(p-value). Pathways shown met p-value ≤ 0.05. (H) Integrated RNA-seq/ATAC-seq analysis (PRT13722 5 nM vs prifetrastat 50 nM) showing genes significant in both RNA-seq and ATAC-seq (FDR < 0.05 in both assays) and classified as concordant if RNA and ATAC changed in the same direction and discordant if they changed in opposite directions. (I) Darker integrated heatmap showing RNA logFC and ATAC log2FC versus DMSO for selected canonical genes. Curved colors represent deeper molecular perturbations.

Results - In Vivo

Figure 5. PRT13722 Achieves Selective KAT6A Degradation and Target Engagement In Vivo

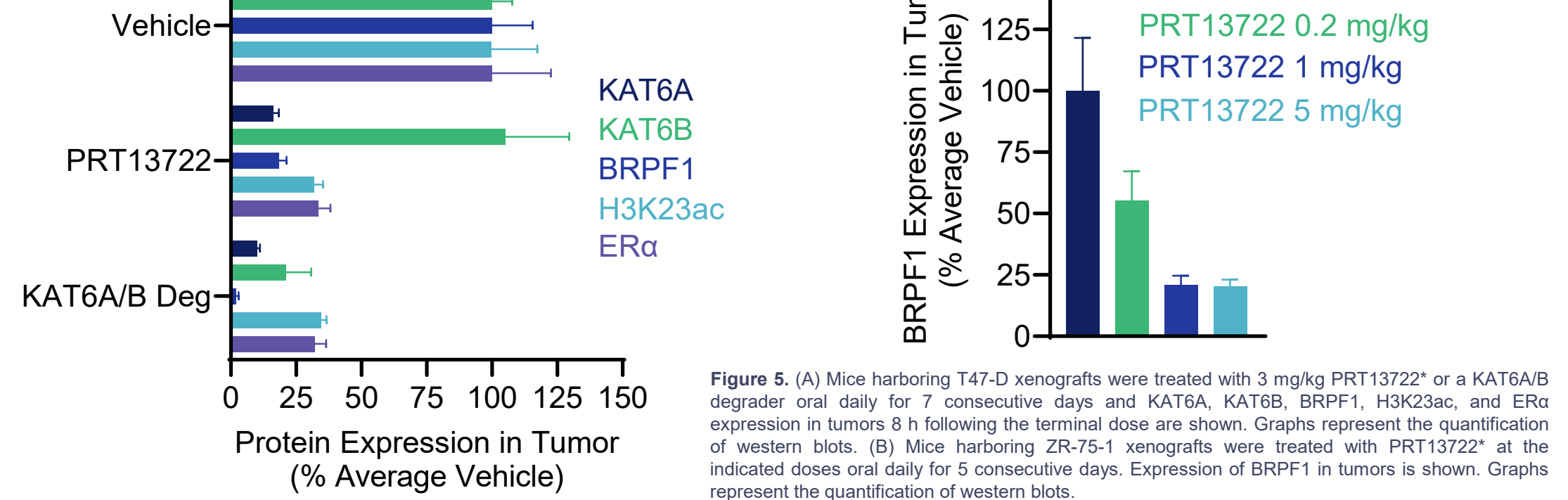


Figure 6. PRT13722 Monotherapy Drives Deep and Complete In Vivo Tumor Regressions in Multiple KAT6A-Amplified Breast Cancer Xenografts

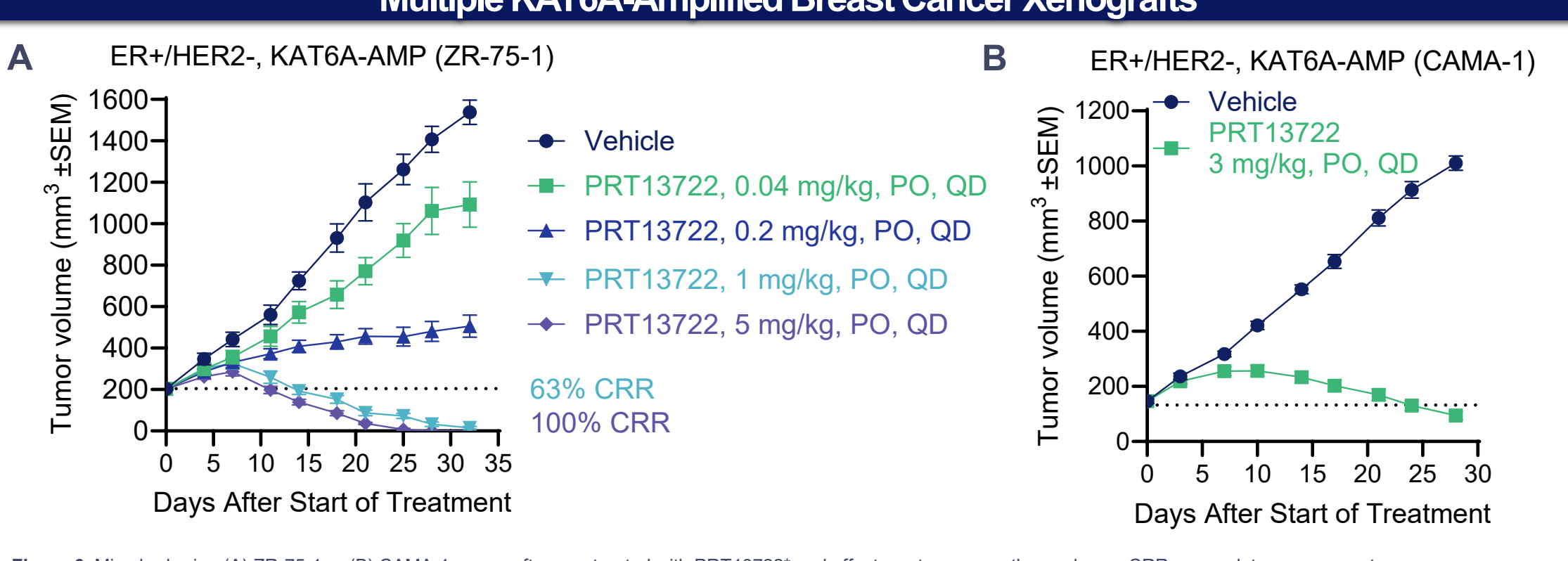


Figure 7. PRT13722 Monotherapy Safely Drives Deep and Durable In Vivo Tumor Regressions in Breast Cancer Xenografts with Superior Efficacy and Pharmacology Than prifetrastat

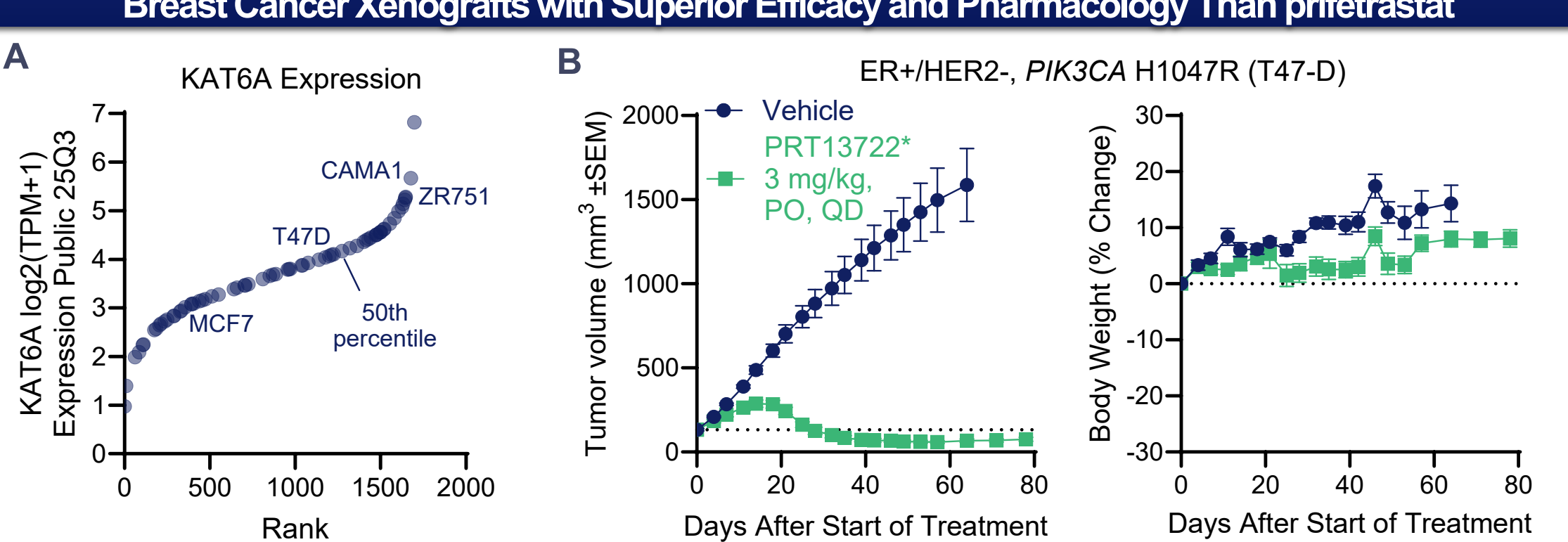


Figure 8. PRT13722 Monotherapy Durably Eradicates Post Endocrine Therapy Patient-Derived Tumors

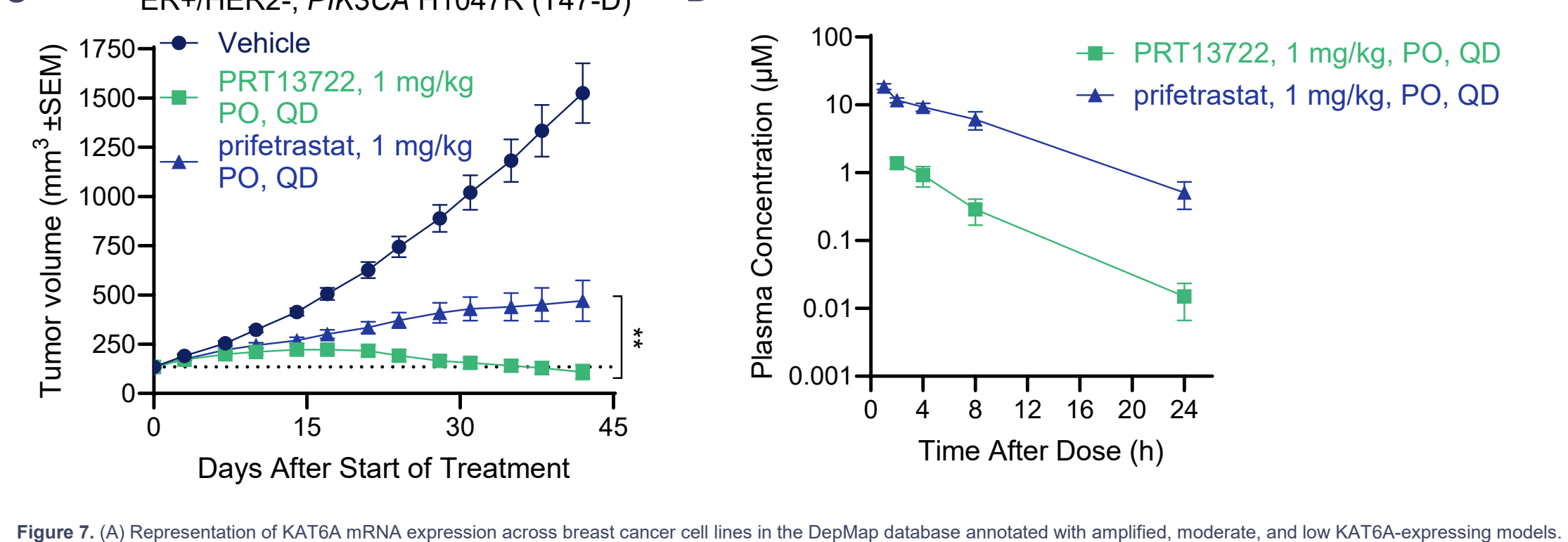


Figure 9. PRT13722 Demonstrates Robust In Vivo Combination with Current and Advanced Next-Generation Clinical Endocrine-, CDK4/6-, and PI3Kα-Targeted Therapies

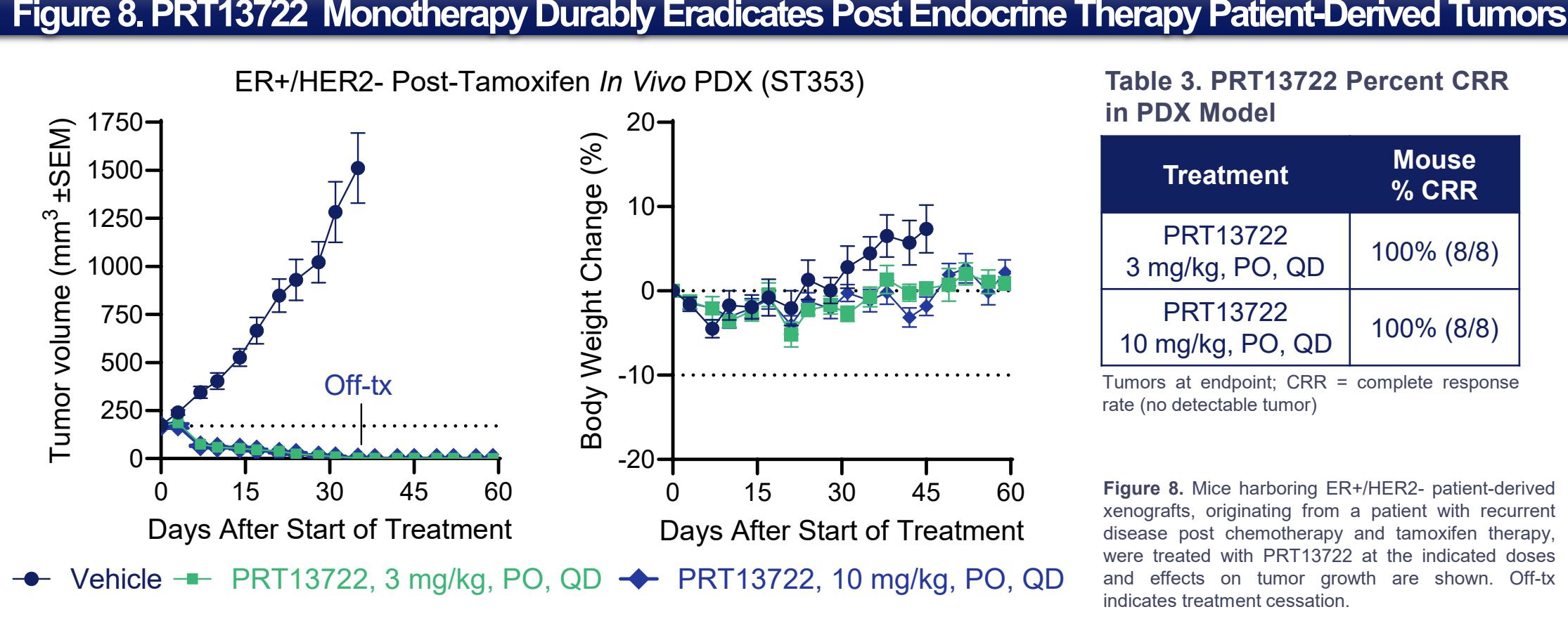


Figure 9. (A-B) T47-D xenografts were treated with PRT13722 and various combinations of endocrine, CDK4/6, or PI3Kα inhibitors and changes in tumor volume from baseline is shown. (C-D) Mouse body weight change from the study in (A) panel A and (D) panel B.

Table 4. Summary of Percent of Tumors Regressed with PRT13722 Monotherapy and Combinations at 1 Month

Class	Treatment	% Tumors Regressed
Mono-Tx	prifetrastat, 1 mg/kg	0%
	PRT13722, 0.3 mg/kg	25%
	PRT13722, 1 mg/kg	50%
+ ET	PRT13722, 3 mg/kg	75%
	prifetrastat, 1 mg/kg + fulvestrant	13%
	PRT13722, 1 mg/kg + fulvestrant	88%
+ CDK4/6i	PRT13722, 0.3 mg/kg + imlunestrant	100%
	PRT13722, 0.3 mg/kg + giredestrant	100%
	prifetrastat, 1 mg/kg + abemaciclib	38%
+ PI3Kαi	PRT13722, 1 mg/kg + abemaciclib	100%
	PRT13722, 0.3 mg/kg + ribociclib	100%
	PRT13722, 1 mg/kg + alpelisib	88%
	PRT13722, 0.3 mg/kg + torisolisib	63%
	PRT13722, 0.3 mg/kg + inavolisib	75%

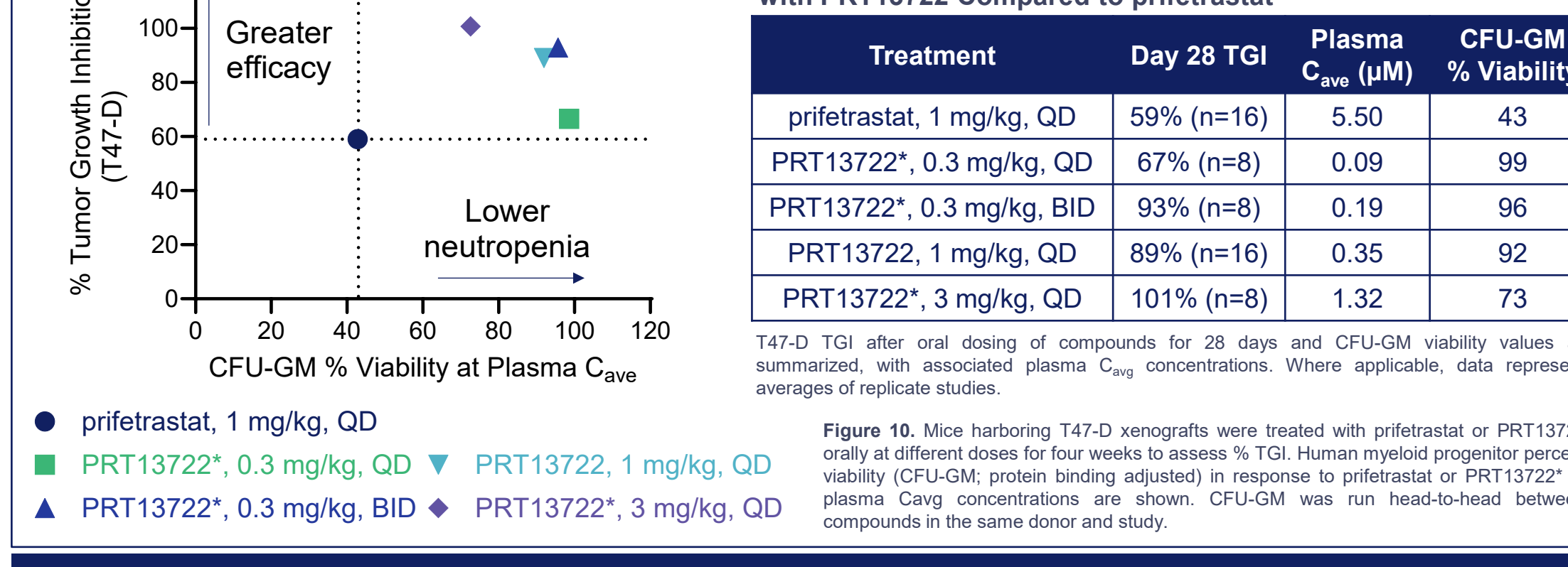
% T47-D tumors regressed below baseline by Day 31-32. Combination agents given at doses with 0% regression. All agents besides fulvestrant are PO QD; giredestrant 1 mg/kg; imlunestrant 3 mg/kg; abemaciclib 25 mg/kg; ribociclib 25 mg/kg; alpelisib 10 mg/kg; inavolisib 5 mg/kg; torisolisib 25-100 mg/kg, PO, QD, SC, QW + LD

Table 5. Summary of Efficacy, Exposure, and CFU-GM Viability with PRT13722 Compared to prifetrastat

Treatment	Day 28 TGI	Plasma C _{avg} (μM)	CFU-GM % Viability
prifetrastat, 1 mg/kg, QD	59% (n=16)	5.50	43
PRT13722*, 0.3 mg/kg, QD	67% (n=8)	0.09	99
PRT13722*, 0.3 mg/kg, BID	93% (n=8)	0.19	96
PRT13722, 1 mg/kg, QD	89% (n=16)	0.35	92
PRT13722*, 3 mg/kg, QD	101% (n=8)	1.32	73

T47-D TGI after oral dosing of compounds for 28 days and CFU-GM viability values are summarized, with associated plasma C_{avg} concentrations. Where applicable, data represents averages of replicate studies.

Figure 10. PRT13722 Has Potential for an Improved Hematological Safety Profile vs prifetrastat



Conclusions

- PRT13722 is a highly differentiated, first-in-class, orally bioavailable, potent and highly selective KAT6A degrader development candidate.
- PRT13722, by degrading KAT6A, drives more complete disruption of KAT6A regulatory pathways than dual KAT6A/B inhibitors, resulting in more robust depth and breadth of efficacy in HR+/HER2- breast cancer.
- PRT13722 drives durable complete tumor regressions in HR+/HER2- CDX and PDX xenograft models (both ET sensitive and experienced) at well-tolerated doses, as a monotherapy.
- PRT13722 is synergistic with ET, CDK4/6i, and PI3Kαi while maintaining monotherapy and combination activity across HR+ BC models, including ESR1 mutated and acquired therapy-resistant cancer cells.
- PRT13722 has an improved preclinical hematological safety profile compared to prifetrastat, which may enable combinations with SOC agents in HR+ BC.
- PRT13722 is on track for IND filing in mid-2026.

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