

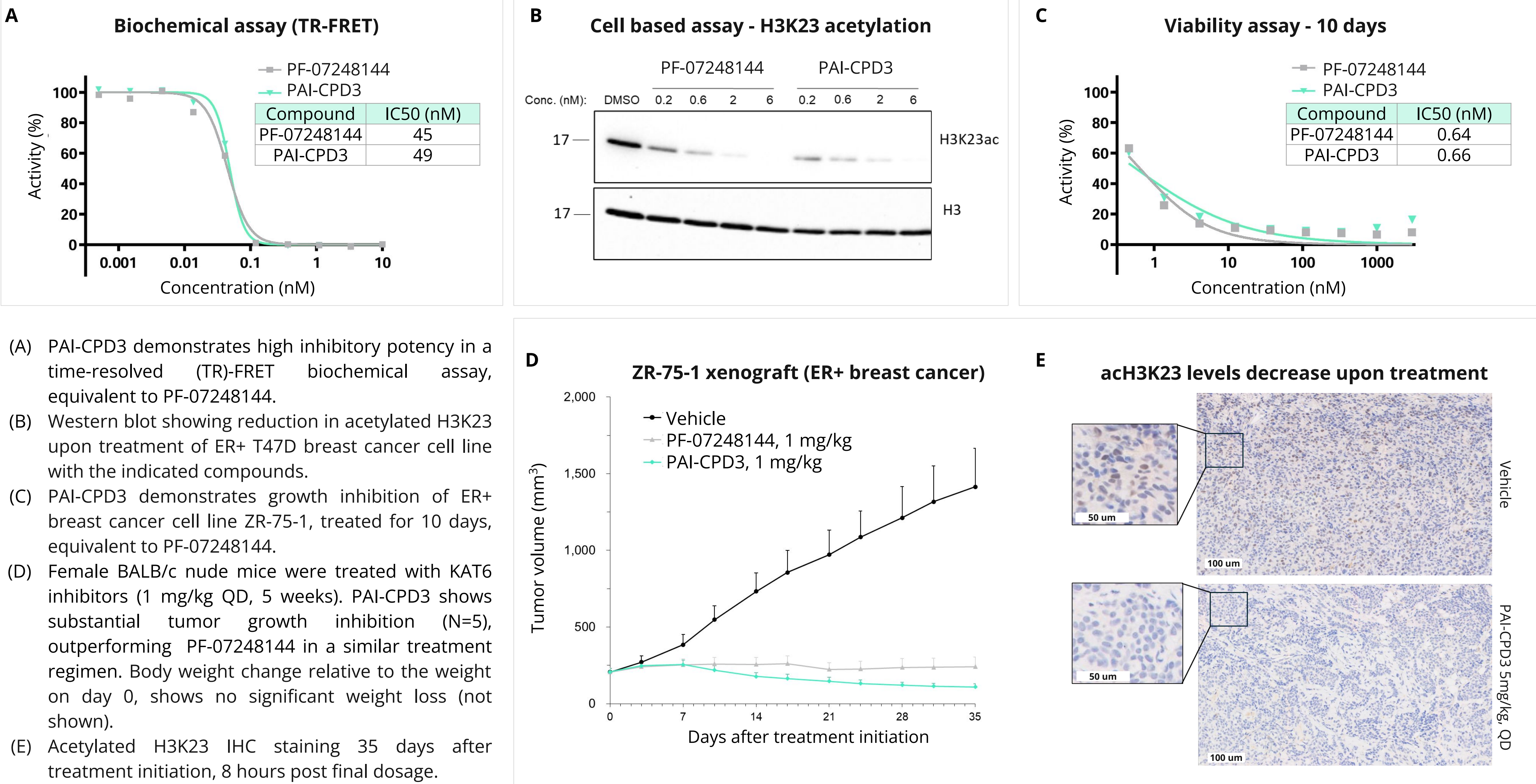
Discovery of a novel KAT6A/B inhibitor with anti-tumor activity coupled with novel response biomarkers

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Introduction

- Lysine histone acetyltransferase 6 (KAT6A/B) is a MYST family member of histone acetyltransferases (HATs), known to regulate gene expression by acetylation of both histone and non-histone substrates.
- KAT6A/B is a promising cancer drug target, specifically in estrogen receptor positive (ER+) breast cancer, but anti-cancer activity in other cancer indications has also been shown, suggesting a potential benefit for wider patient populations¹.
- We discovered a novel, highly potent KAT6A/B inhibitor, showing anti tumor activity both *in-vitro* and in xenograft models, as well as improved selectivity compared to the benchmark compound PF-07248144.
- Combining drug sensitivity with basal and dynamic proteomics data, Protai's AIMSTM platform² discovered a novel biomarker for sensitive cells outside ER+ breast cell lines.

PAI-CPD3 shows high potency *in-vitro* and *in-vivo*



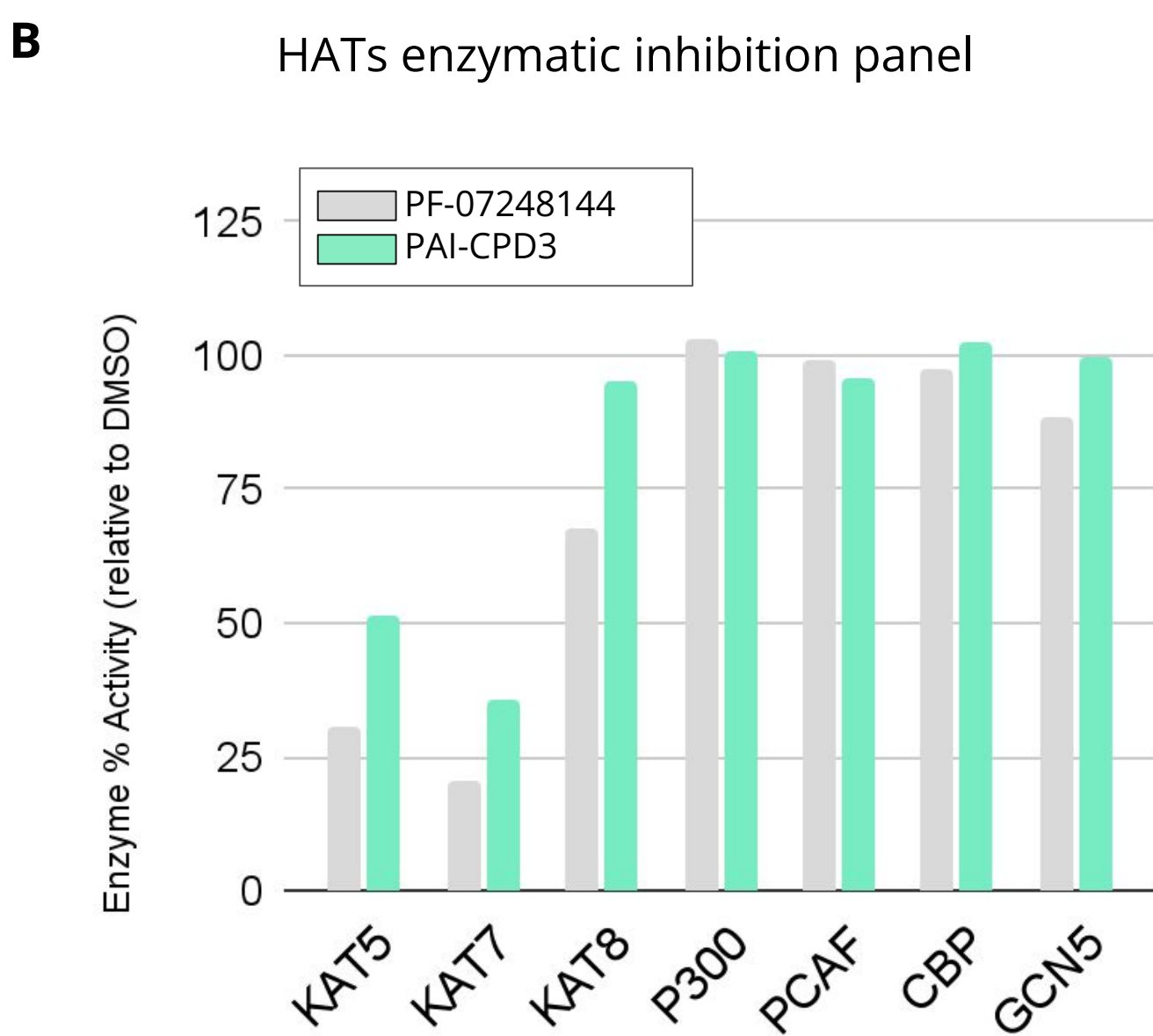
AIMSTM informs PAI-CPD3 favorable selectivity profile

Biochemical HAT enzymatic activity shows selectivity advantage to PAI-CPD3

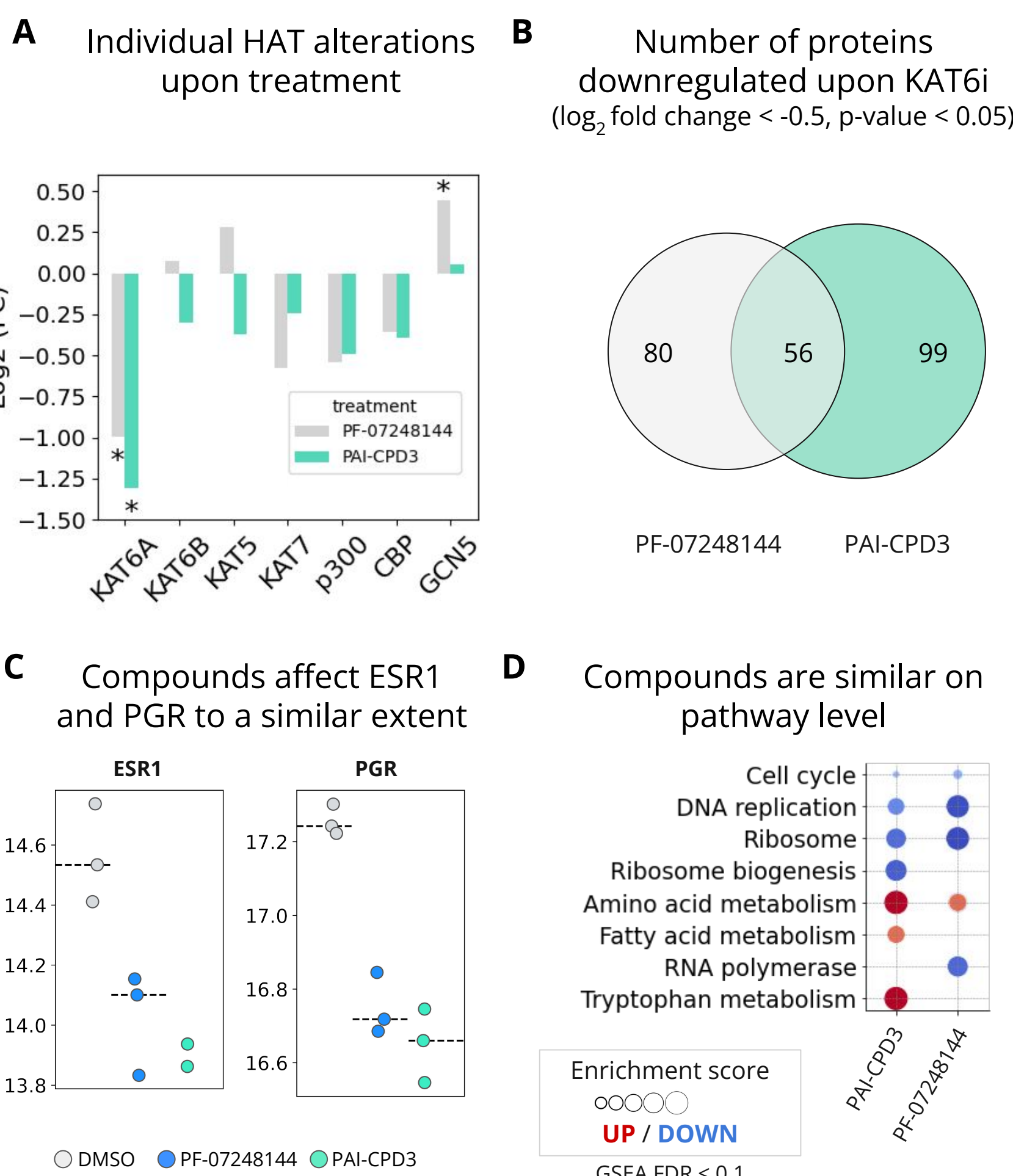
Biochemical assays comparing PAI-CPD3 to PF-07248144 show equivalent potency in KAT6A/B inhibition (10 dose IC50, 3-fold serial dilution starting at 30μM) (A). Both compounds also show equivalent selectivity over other HATs, with a slight advantage of PAI-CPD3 in KAT8 and KAT5 (single dose, duplicate, 10μM) (B).

A KAT6A over KAT6B selectivity

Compound	KAT6A IC50 (nM)	KAT6B IC50 (nM)
PF-07248144	2.37	19.38
PAI-CPD3	1.46	24

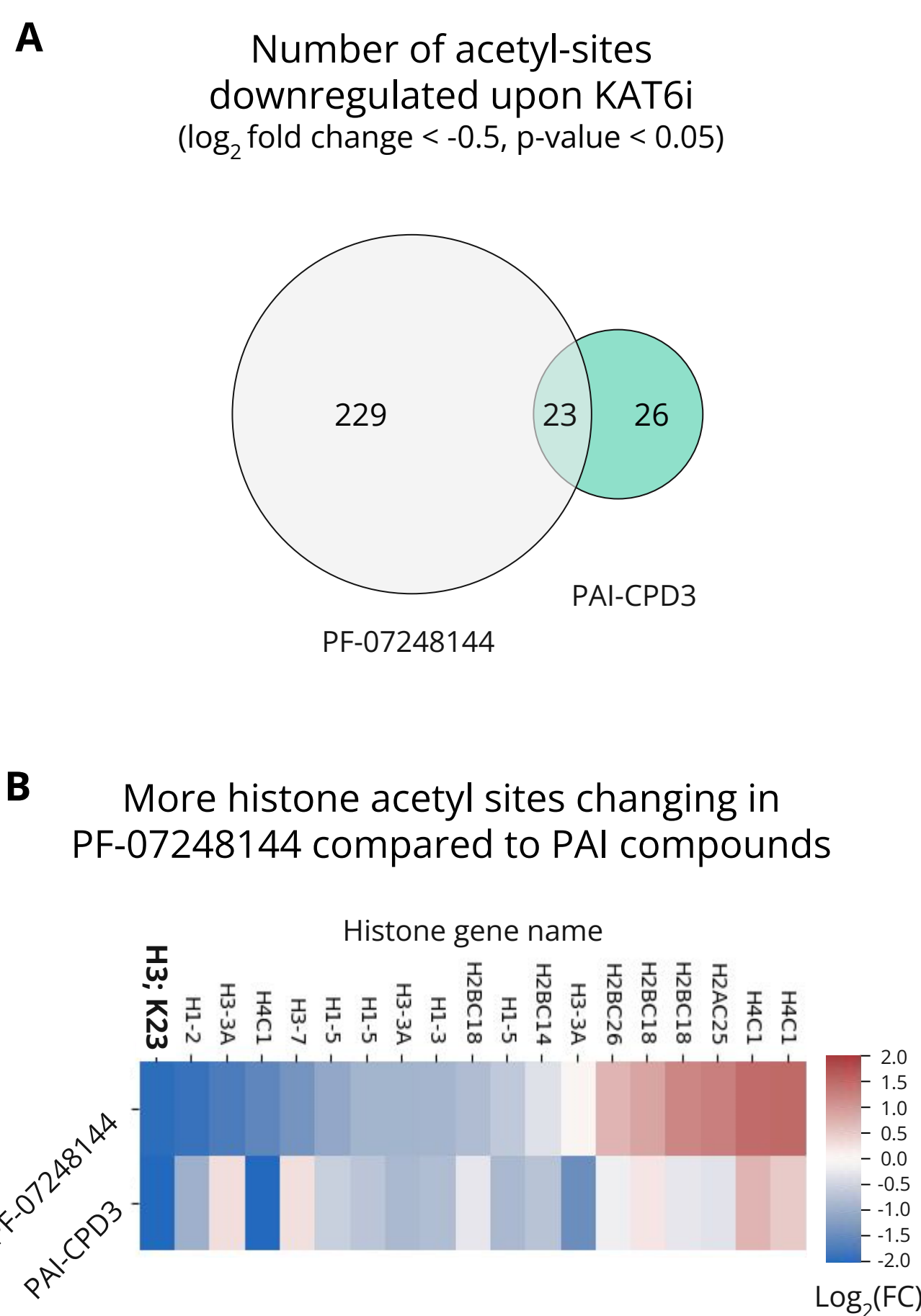


Protein dynamics show similar post-treatment biology



Protein dynamics compound comparison was carried out on ZR-75-1 cell lines at 0.1uM for 24h. Proteomic analysis of downregulated features showed significant inhibition of KAT6A and minimal inhibition of other HATs (A); and overall similar dynamics of both compounds in the number of proteins downregulated upon treatment (B), downregulation of ESR1 and PGR (C), and significantly altered pathways (D).

Acetylation dynamics reveals potential cleaner profile of PAI-CPD3

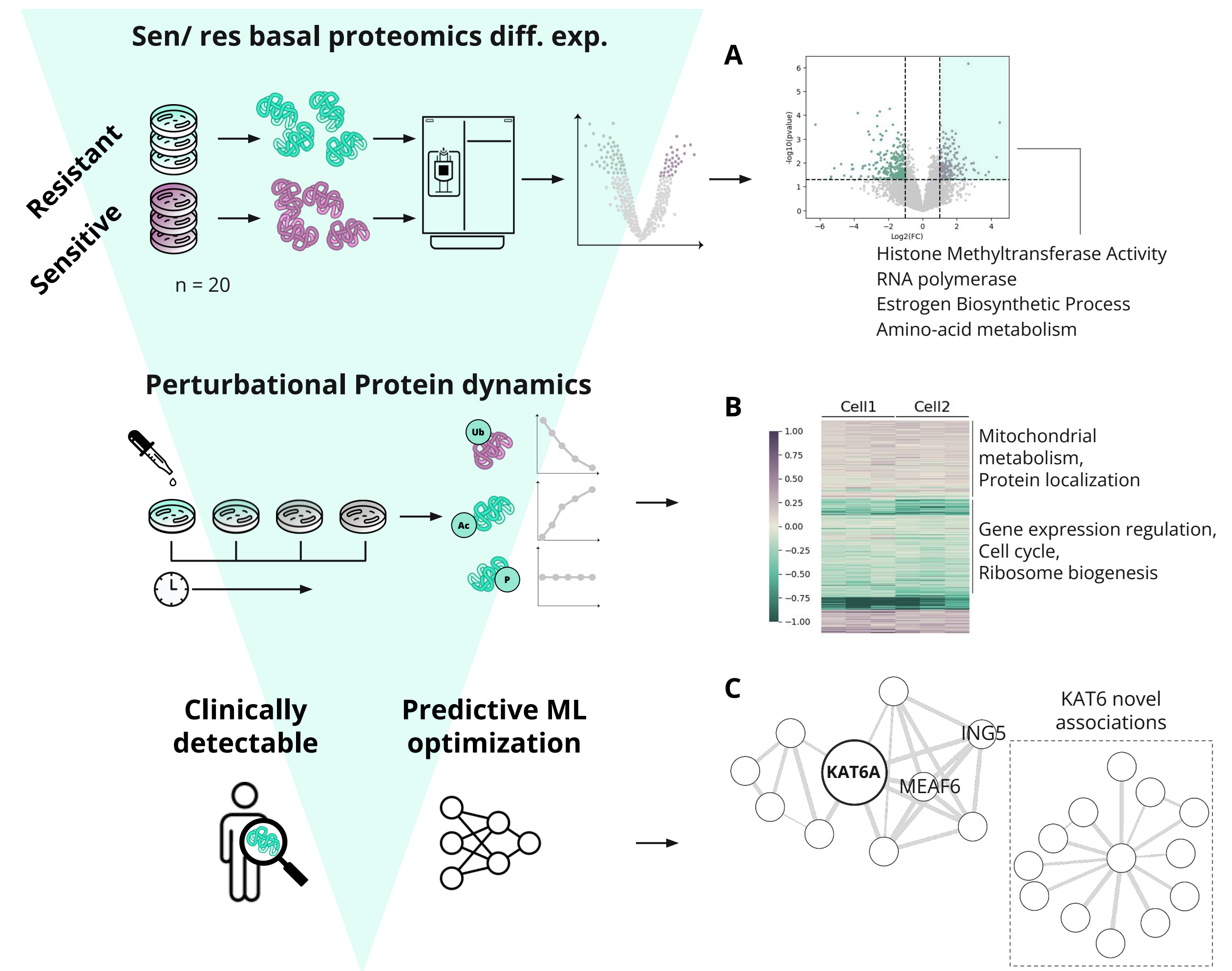


Acetylation dynamics compound comparison was carried out on ZR-75-1 cell line at 0.1uM for 2h. Acetyl-dynamics shows potential selectivity advantage of PAI-CPD3, evident in the total number of regulated acetyl sites (A) and specifically in histone sites, higher in PF-07248144 (B). The KAT6A substrate H3K23 is marked on the left.

Proteomic biomarker identifies sensitivity beyond ER+ breast cancer

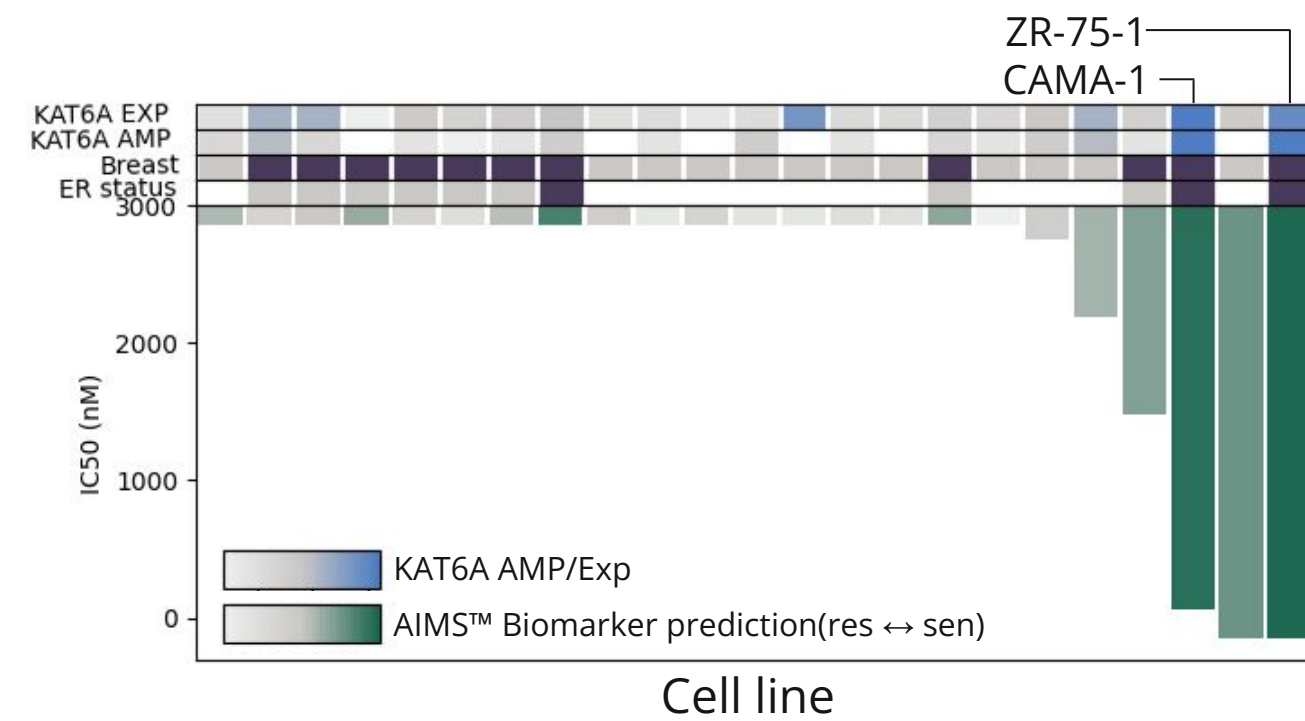
AIMSTM biomarker discovery workflow

Predictive biomarker discovery for KAT6i treatment is composed of several data layers. Differential analysis of both basal-state proteomics (A) and on-treatment dynamics (B) of KAT6i sensitive and resistant cell lines (both in and outside breast cancer) highlights several candidates for KAT6 predictive biomarker, both KAT6A-related and novel KAT6A associations (C, KAT6A PPI network).



AIMSTM predicts sensitivity in cell lines regardless of ER+ status

Applying the selected biomarkers to internal proteomic datasets coupled with KAT6i IC50 values (barplot) demonstrates that biomarker candidates predict KAT6 sensitivity beyond ER status and KAT6A copy number amplification/protein expression.



Summary

- PAI-CPD3, a novel KAT6A/B inhibitor, exhibits potency and *in-vivo* efficacy in ER+ breast cancer.
- PAI-CPD3 demonstrates selective acetylation impact, both on histones and non histone sites.
- KAT6A/B directed treatment represents opportunity for ER+ breast cancer patients as well as other tumor types, supported by a proteomic biomarker.

References

- Mukohara, et al. *Nature Medicine* (2024)
- Alchanati et al. *AACR* (2024)

