



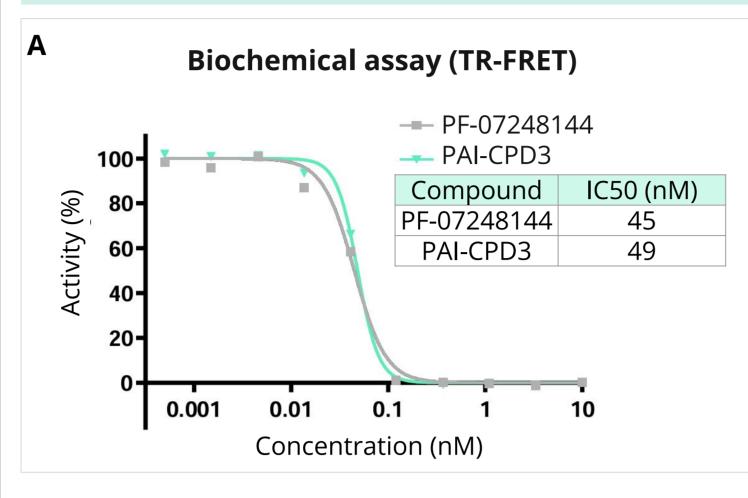
# Discovery of a novel KAT6A/B inhibitor with anti-tumor activity coupled with novel response biomarkers

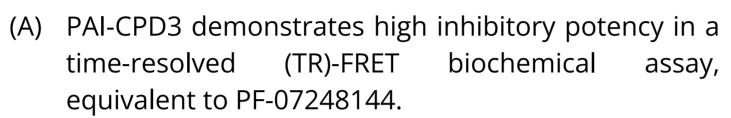
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### Introduction

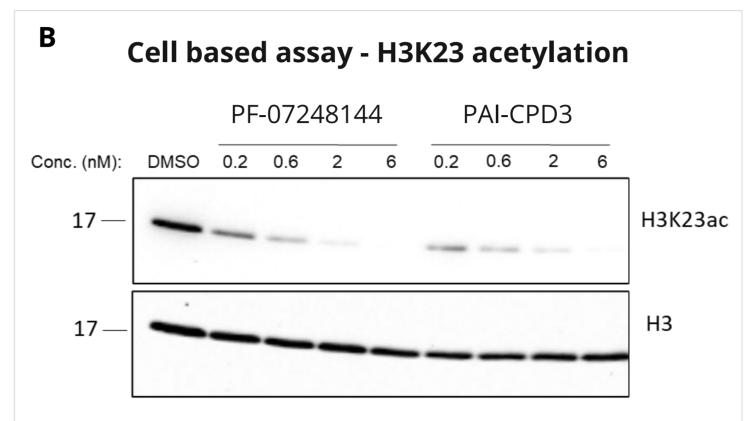
- histone acetyltransferase 6 (KAT6A/B) is a MYST family member of histone acetyltransferases (HATs), known to regulate gene expression by acetylation histone and non-histone substrates.
- KAT6A/B is a promising cancer drug target, specifically in estrogen receptor positive (ER+) breast cancer, anti-cancer activity in other cancer has also been shown, indications suggesting a potential benefit for wider patient populations<sup>1</sup>.
- 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 AIMS<sup>TM</sup> platform<sup>2</sup> discovered a novel biomarker for sensitive cells outside ER+ breast cell lines.

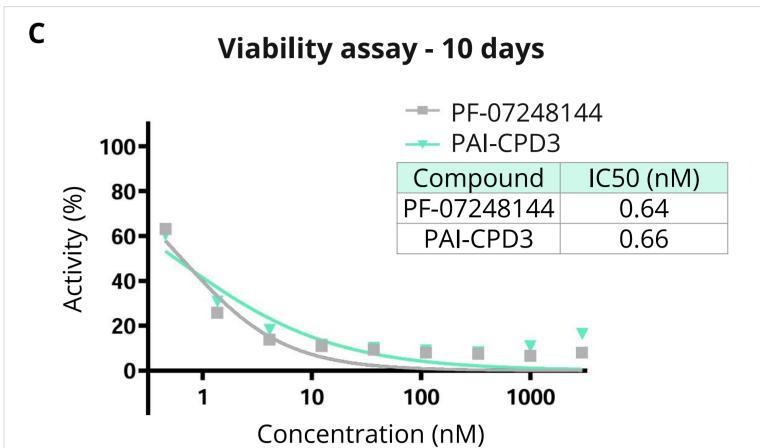
### PAI-CPD3 shows high potency in-vitro and in-vivo

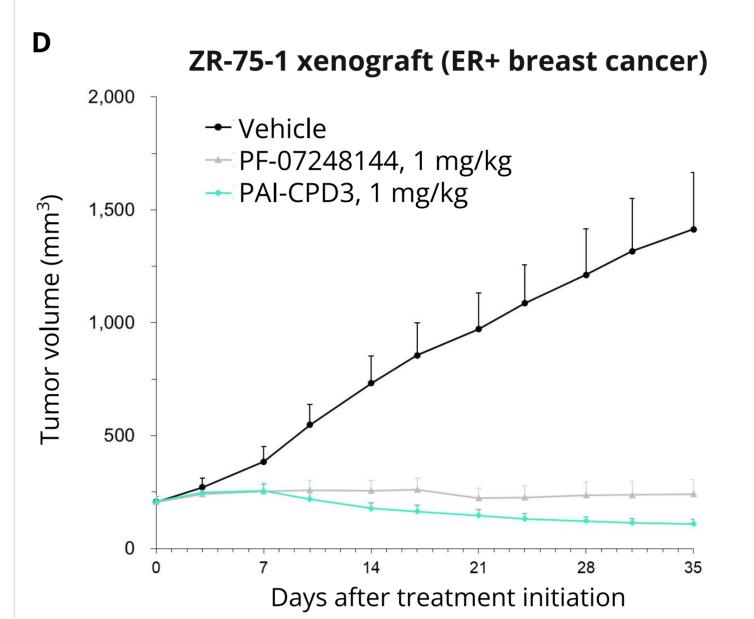


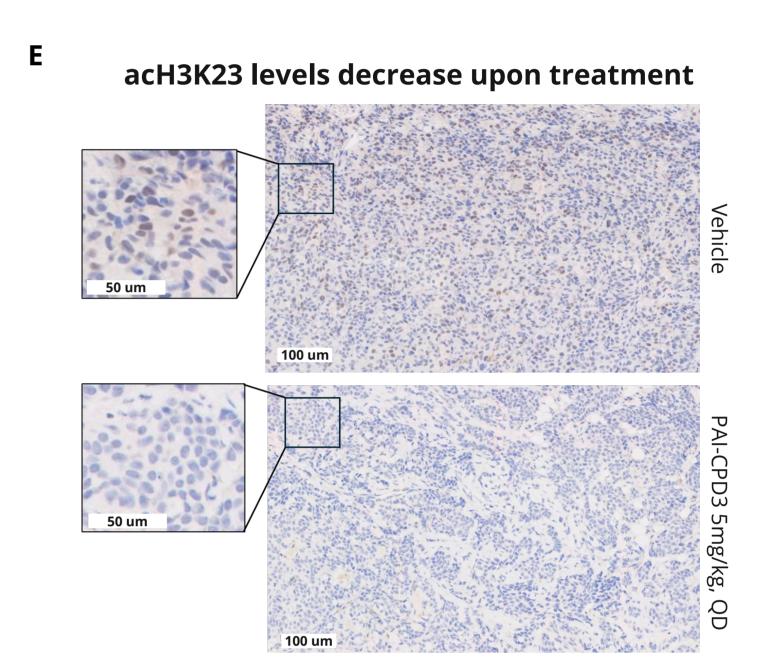


- (B) Western blot showing reduction in acetylated H3K23 upon treatment of ER+ T47D breast cancer cell line with the indicated compounds.
- (C) PAI-CPD3 demonstrates growth inhibition of ER+ breast cancer cell line ZR-75-1, treated for 10 days, equivalent to PF-07248144.
- (D) Female BALB/c nude mice were treated with KAT6 inhibitors (1 mg/kg QD, 5 weeks). PAI-CPD3 shows substantial tumor growth inhibition (N=5), outperforming PF-07248144 in a similar treatment regimen. Body weight change relative to the weight on day 0, shows no significant weight loss (not shown).
- Acetylated H3K23 IHC staining 35 days after treatment initiation, 8 hours post final dosage.









### AIMS™ informs PAI-CPD3 favorable selectivity profile

#### **Biochemical HAT enzymatic activity shows selectivity advantage to PAI-CPD3**

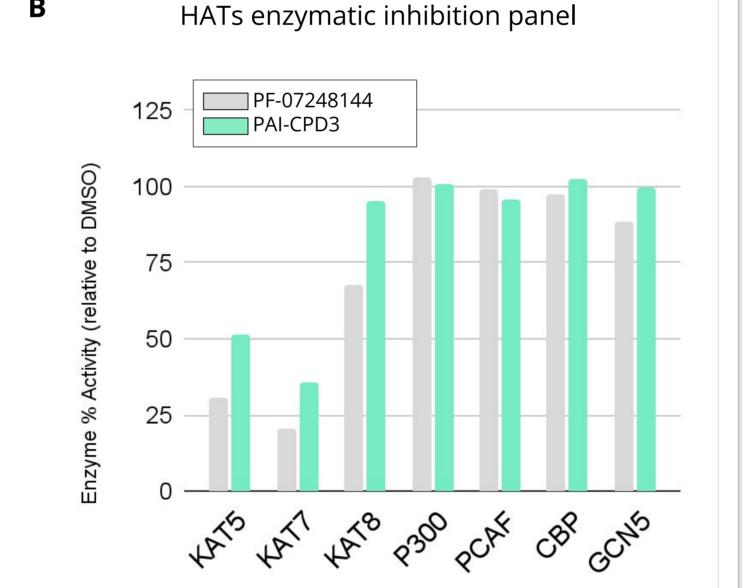
Biochemical assays comparing PAI-CPD3 to PF-07248144 show equivalent potency in KAT6A/B inhibition (10 dose IC50, 3-fold serial dilution starting at 30µM) (A).

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**).

KAT6A over KAT6B selectivity

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

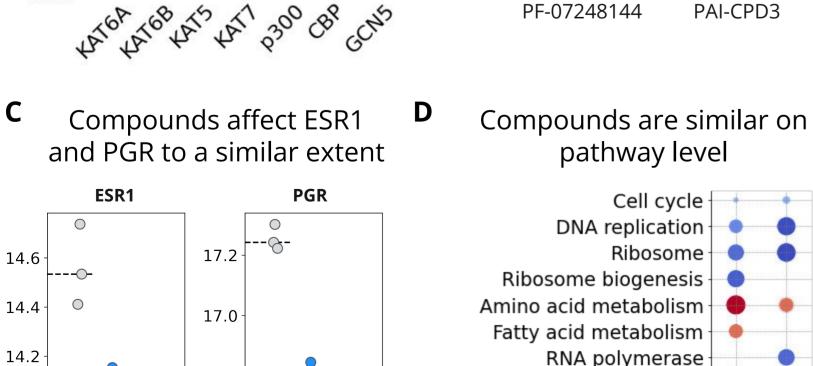
Α



downregulated upon KAT6i upon treatment  $(\log_2 \text{ fold change} < -0.5, p-value} < 0.05)$ 0.50 0.25 26 −0.50 −0.75 56 PF-07248144 -1.00

**Protein dynamics show similar post-treatment biology** 

Individual HAT alterations



14.0 DMSO PF-07248144 PAI-CPD3

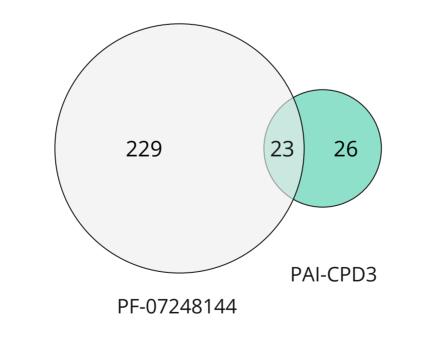
pathway level Cell cycle DNA replication + Ribosome + Ribosome biogenesis Amino acid metabolism 🗕 🌑 Fatty acid metabolism | • RNA polymerase Tryptophan metabolism 🗐 **Enrichment score** 0000 **UP / DOWN** GSEA FDR < 0.1

Number of proteins

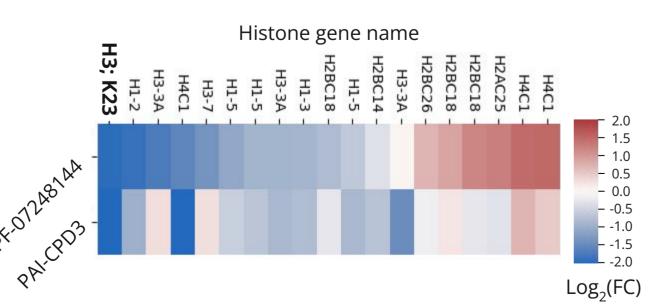
Protein dynamics compound comparison was carried out on ZR-75-1 cell lines at 0.1uM for 24h. Proteomic analysis of downregulated features showed significant inhibition of KAT6A and minimal inhibition of other HATs (A); and overall similar dynamics of both compounds in the number of proteins downregulated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

#### **Acetylation dynamics reveals potential** cleaner profile of PAI-CPD3

Number of acetyl-sites downregulated upon KAT6i  $(\log_2 \text{ fold change} < -0.5, p\text{-value} < 0.05)$ 



More histone acetyl sites changing in PF-07248144 compared to PAI compounds

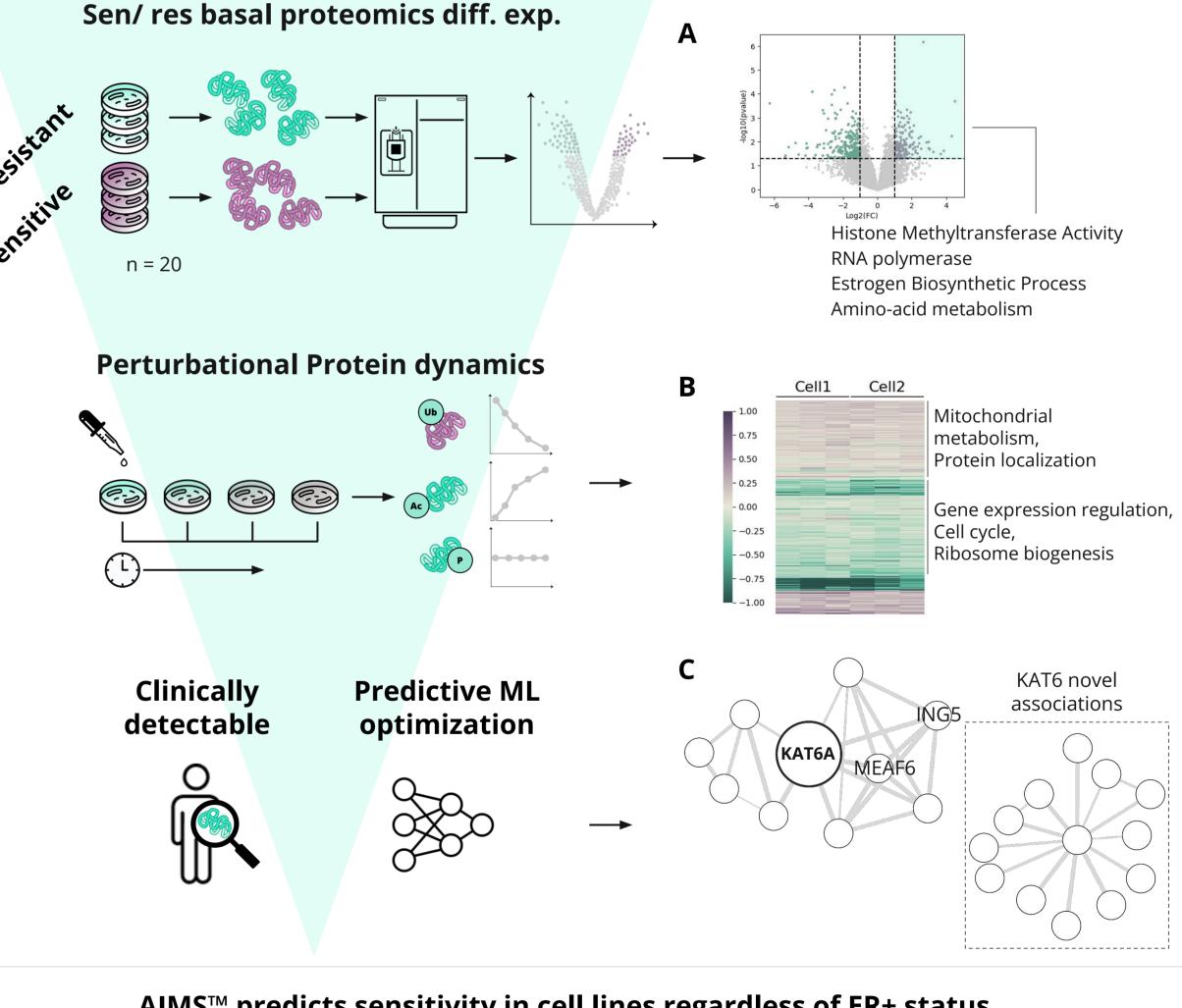


Acetylation dynamics compound comparison was carried out on ZR-75-1 cell line at 0.1uM for 2h. Acetyl-dynamics shows potential selectivity advantage of PAI-CPD3, evident in the total number of regulated acetyl sites (A) and specifically in histone sites, higher in PF-07248144 (**B**). The KAT6A substrate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

## Proteomic biomarker identifies sensitivity beyond ER+ breast cancer

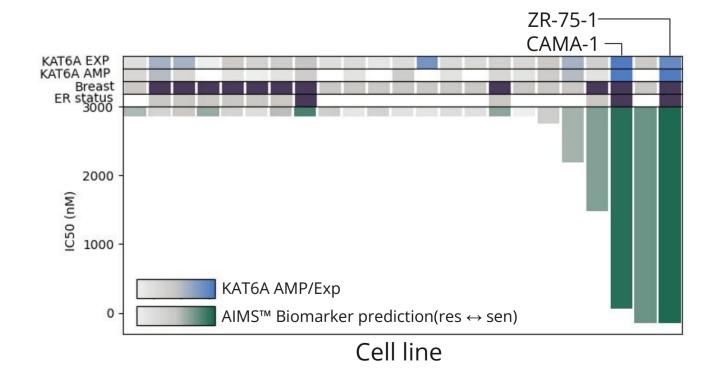
### **AIMS™** biomarker discovery workflow

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).



#### **AIMS™** predicts sensitivity in cell lines regardless of ER+ status

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

- 1. Mukohara, et al. Nature Medicine
- (2024)2. Alchanati et al. AACR (2024)

