

Accelerating Small Molecule Drug Discovery via Al-Powered Kinetic Ensemble® Platform

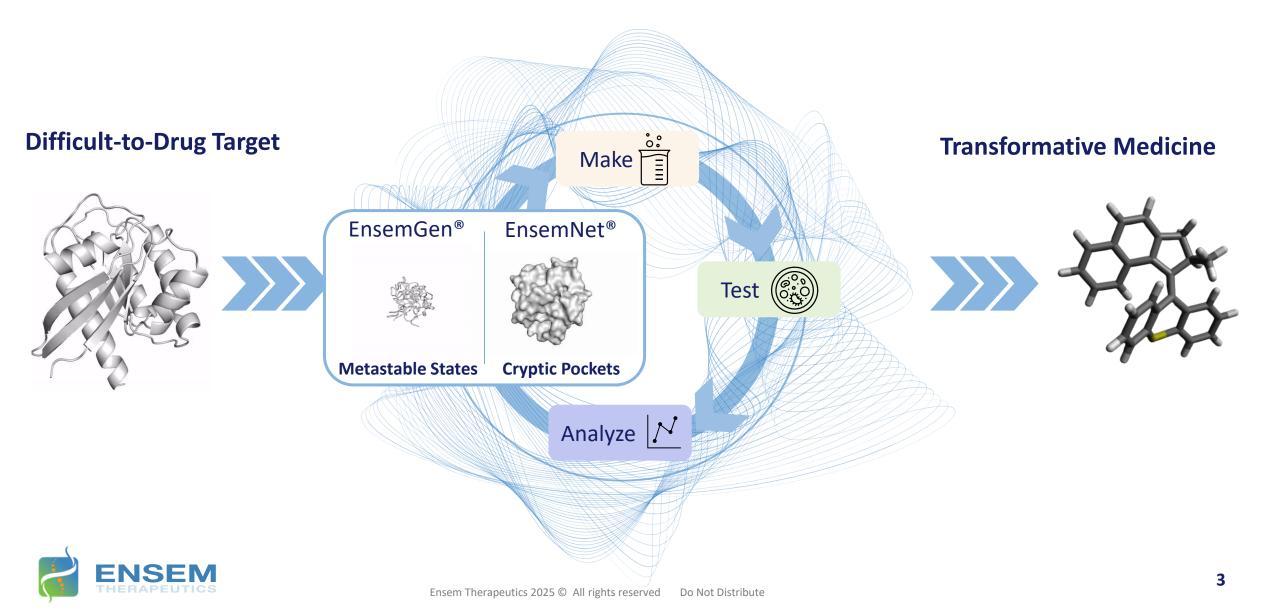
Shengfang Jin, PhD CEO and Co-Founder

ENSEM is Expanding the Universe of Druggable Targets to Bring Transformative Therapies to Patients

- Substantial challenges exist in small molecule drug discovery in high-value but difficult-to-drug targets
- Kinetic Ensemble[®], ENSEM's deep learning AI platform, addresses these challenges and has enabled a clinical-stage proprietary precision medicine pipeline in three years
 - CDK2 inhibitor in Phase 1 (partnered with BeiGene, with a total deal value ~\$1.33B + royalty)
 - Pan-mutant PI3Ka **dual** inhibitor/degrader to file IND and initiate Phase 1 in 1H 2025
- Seasoned industry leaders and "drug hunters"
- Strong financial backing: \$120M* raised since inception in 2021
- Multiple value inflection points in the next 24 months



Kinetic Ensemble® Delivers Solutions for Difficult-to-Drug Targets



Rapid Development of Precision Medicine Pipeline in 3 Years Derisked with Biologically Validated and Clinically Translatable Programs

TARGET	PROGRAM	INDICATION	PRECLINICAL	IND	PHASE I	GLOBAL RIGHTS
CDK2	ETX-197/ BG-68501	Oncology				BeiGene
Pan Mutant PI3Kα	ETX-636	Oncology & Rare Disease				
Synthetic Lethal Target	undisclosed	Oncology				
Oncogenic Driver	undisclosed	Oncology				



ETX-197: Best-in-Class CDK2 Inhibitor

Concept to IND in 2 Years

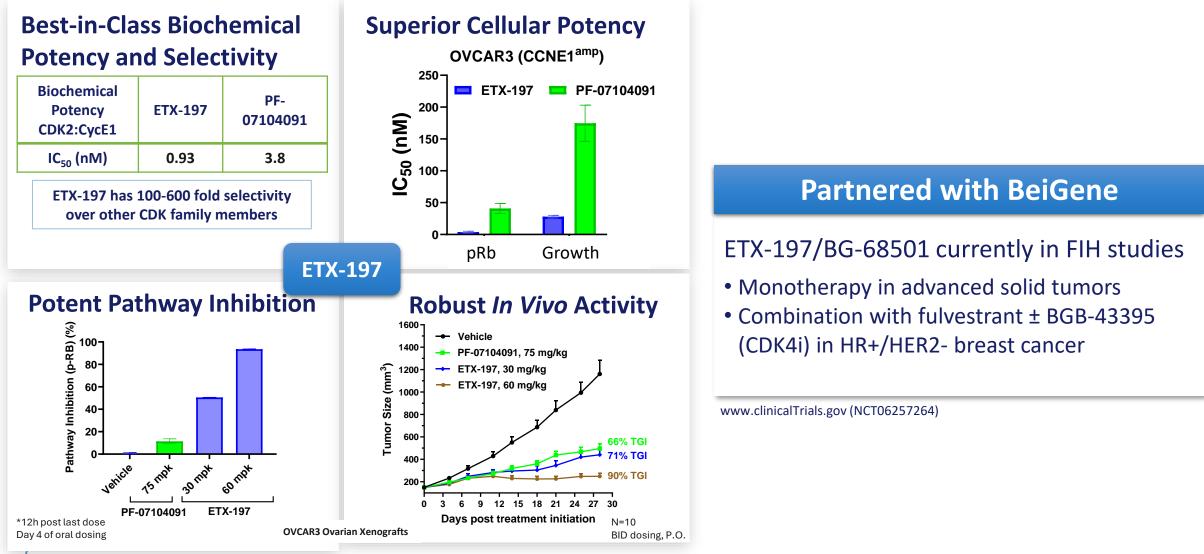
- 1st Program discovered using Kinetic Ensemble®
- Overcomes suboptimal potency, selectivity, and/or druglike properties of current and emerging CDK2 therapies
- Partnered with BeiGene





THFRAPFUTICS

ETX-197: Potentially Best-in-Class CDK2 Inhibitor in the Clinic



ETX-636: Next Generation Pan-Mutant Selective Allosteric Inhibitor and Degrader of PI3Kα

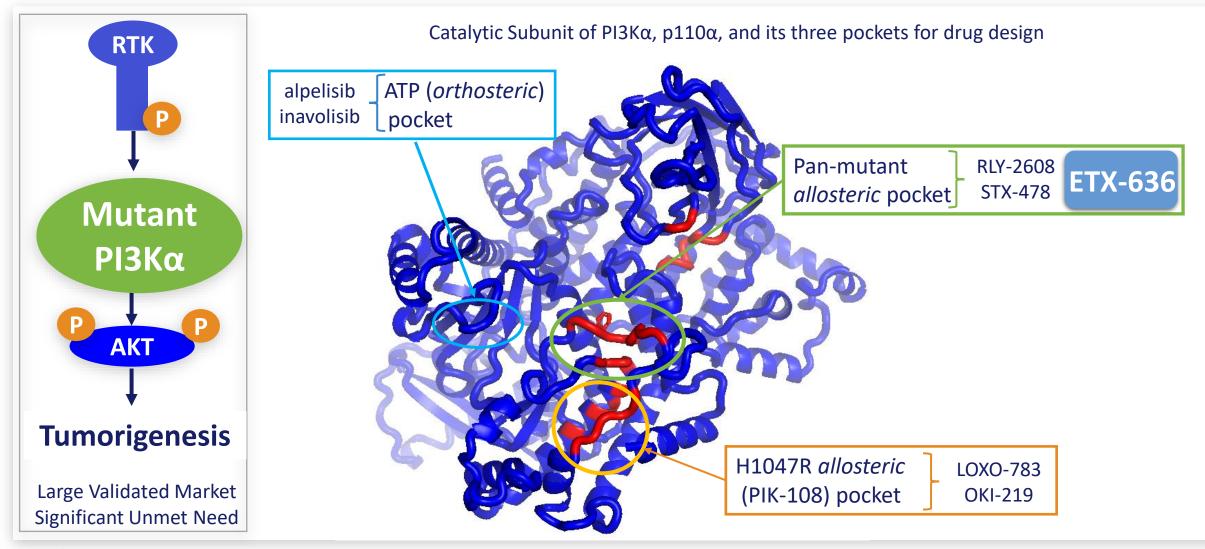
Concept to IND submission in 3 years

Allosteric Inhibition + Protein Degradation = Best-in-Class





Kinetic Ensemble[®] Prioritizes a Specific PI3Kα Allosteric Pocket





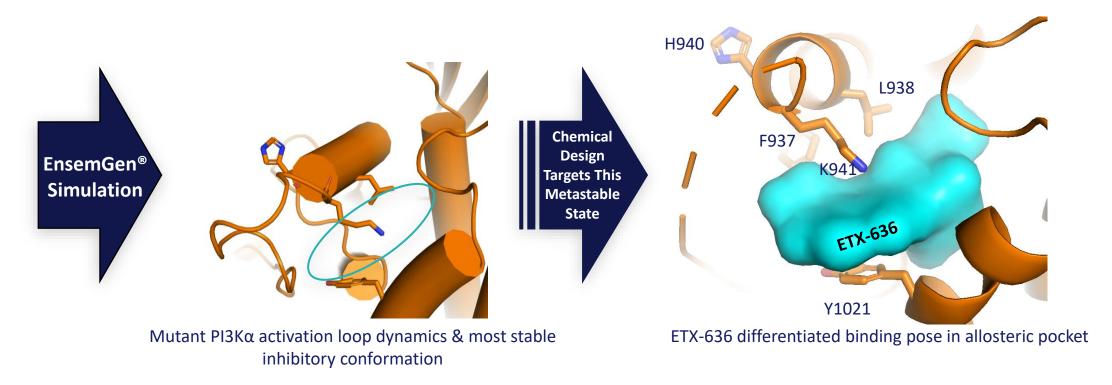
ETX-636 Delivers Solutions to Deficiencies of PI3K α Clinical Agents

Issues with currently approved & emerging PI3Kα inhibitors	Desired Target Drug Profile
 Orthosteric ATP-pocket inhibitors (e.g., alpelisib, inavolisib) Lack mutant/wildtype selectivity which undercuts efficacy Low tolerability (e.g., hyperglycemia), limited clinical utility 	 >10x selectivity for mutant vs wild type PI3Kα to improve tolerability, efficacy and broaden patient population
 Allosteric inhibitors (e.g., RLY-2608, STX-478, OKI-219) Suboptimal <i>in vitro</i>, <i>in vivo</i> potency and/or drug properties 	 Improved efficacy resulting from superior potency and better drug properties
 PI3Kα inhibition triggers compensatory feedback loop ATP pocket mutations create resistance to orthosteric inhibitors 	 DUAL allosteric inhibition and degradation of mPI3Kα to abrogate compensatory feedback and overcome drug resistance

ETX-636 Is a Differentiated Next Generation Dual PI3Kα Inhibitor & Degrader - Allosteric & Pan-Mutant Selective



EnsemGen® Elucidated a Unique PI3Ka Conformation Enabling ETX-636 Design

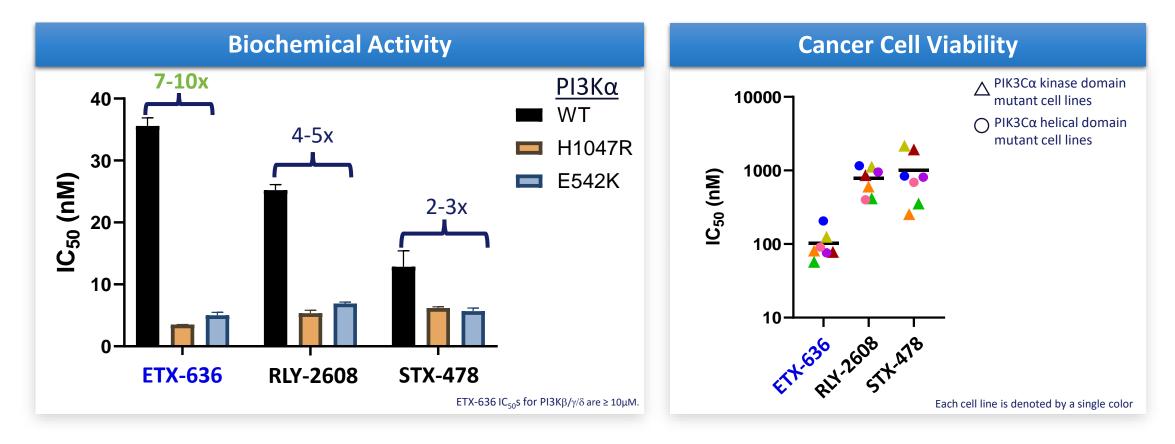


Delivers **tight binding to a unique metastable state** -> slowest off-rate and longest residence time

2 Selectively captures mutant PI3K α in the **metastable** state -> **degradation** of mutant PI3K α in cells



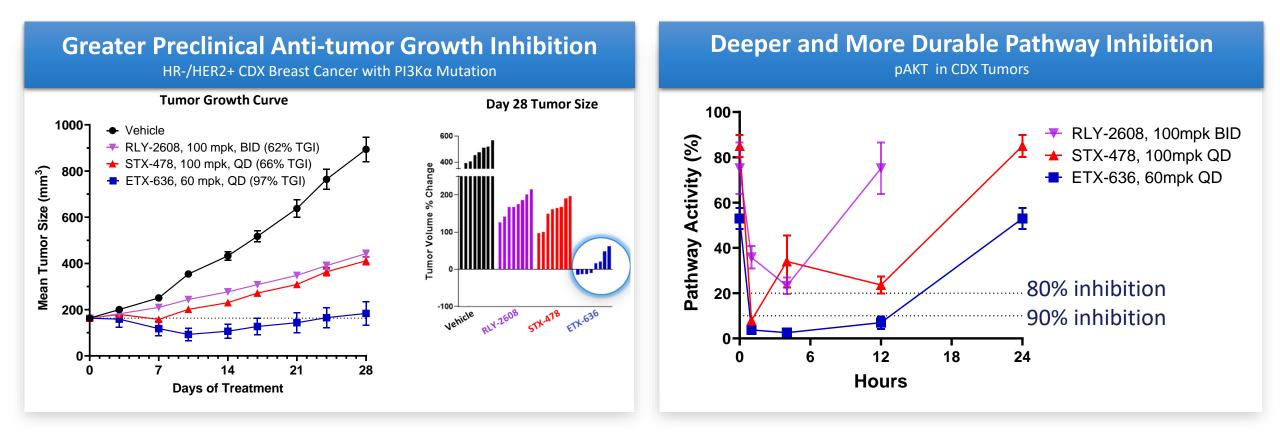
ETX-636 Demonstrates Superior Potency and Selectivity for Mutant PI3Kα



- ETX-636 potently inhibits both kinase and helical domain hotspot mutations and other PI3Kα activating mutations
- ETX-636 demonstrates good concordance with its biomarker (pAKT) and cell growth (CTG) potencies



ETX-636 Shows Superior In Vivo Efficacy and PI3Kα Pathway Inhibition

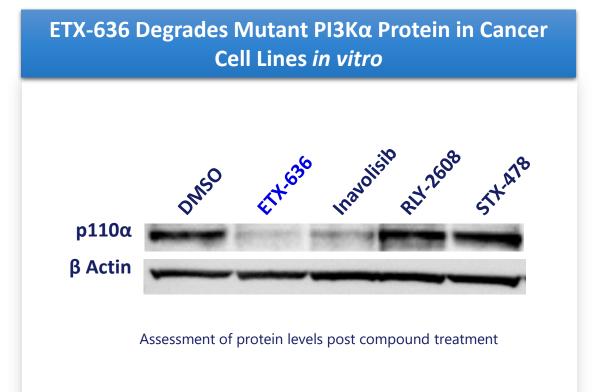


- ETX-636 does not impact glucose homeostasis at efficacious doses in experimental animals
- ETX-636 demonstrates good concordance of drug exposure, biomarker (pAKT) and tumor growth inhibition (TGI)

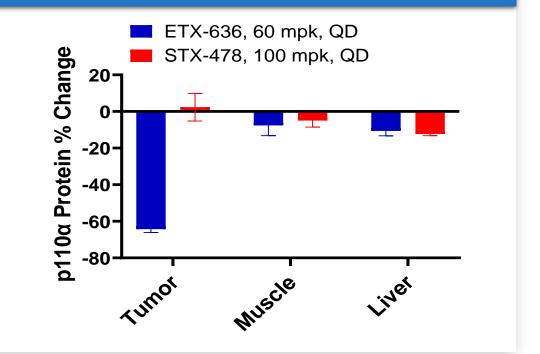
Studies conducted using HCC1954, an HR-/HER2+ Breast Cancer cell line harboring PIK3CA kinase domain (H1047R) mutation, similar observation in helical domain mutants. CDX = cell line-derived tumor xenograft. mpk=milligram per kilogram; QD = daily dosing; BID=twice a day dosing



ETX-636 Selectively Degrades Mutant PI3Kα, Sparing Wildtype



ETX-636 Degrades Mutant PI3Kα in CDX Tumors but not Wildtype PI3Kα in Healthy Tissues *in vivo*

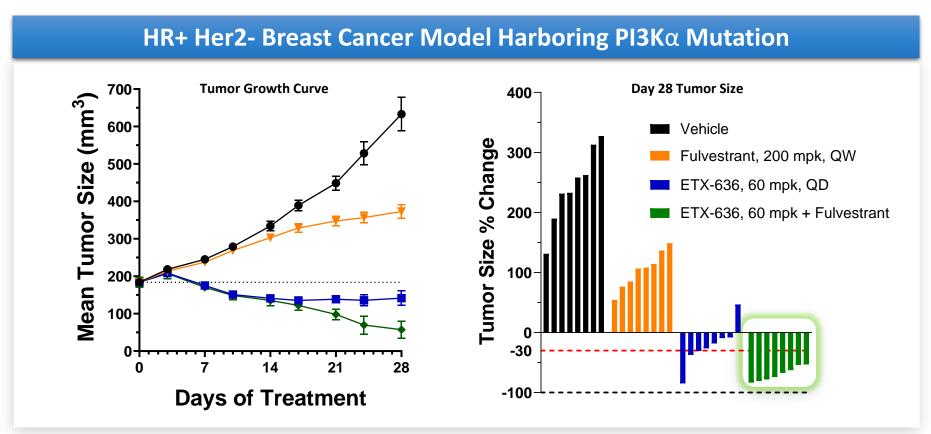


Loss of Mutant PI3Kα contributes to superior depth and durability of pathway abrogation (pAKT) and concordant tumor growth inhibition

HCC1954, HR-/HER2+ PIK3CA H1047R breast cancer cell line, with a mutant variant allele frequency of 0.79 Similar observations are seen in PIK3CA helical domain mutant cancer cells. mpk=milligram per kilogram; QD = daily dosing



ETX-636 Combination with Fulvestrant Demonstrates Synergy and Supports Targeting 2nd Line Advanced Breast Cancer



Study conducted with MCF7, an HR+/HER2- breast cancer cell line harboring PIK3CA helical domain (E545K) mutation mpk=milligram per kilogram; QD = daily dosing; QW = once a week dosing



ETX-636 Is a Differentiated Next Generation Dual PI3Kα Inhibitor & Degrader Allosteric & Pan-Mutant Selective

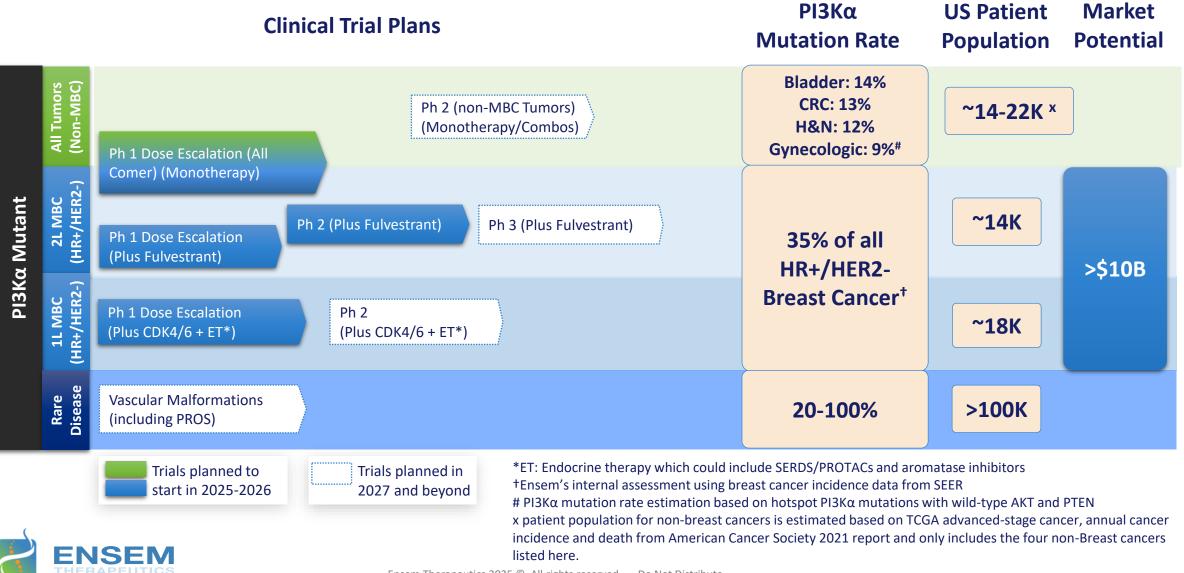
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Projected to achieve PI3Kα target/pathway coverage above IC₉₀ for up to 24hrs <u>without</u> impacting glucose homeostasis in the clinic

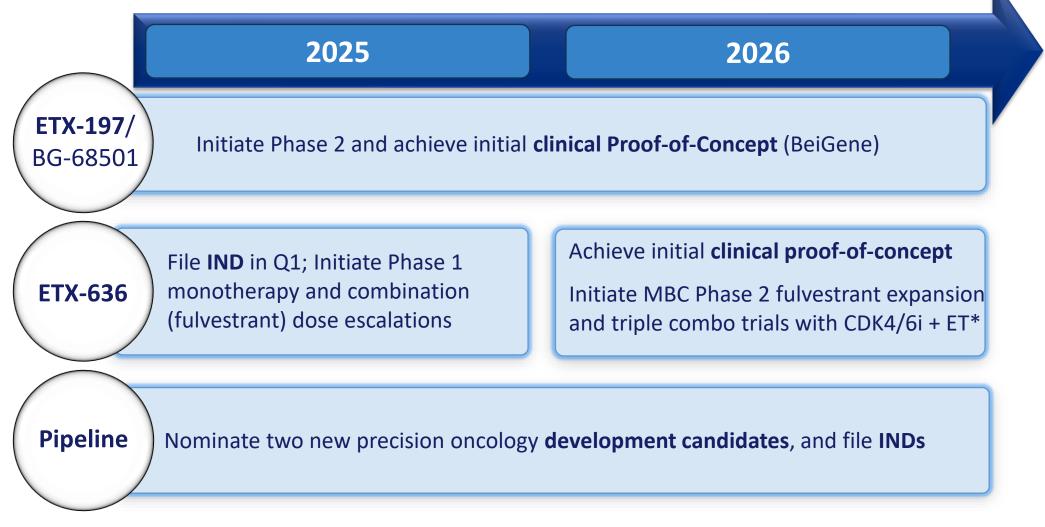
ETX-636: Potential to Achieve Superior Clinical Activity Over All Competitors PI3Kαi



ETX-636: Excellent Potential for Diseases Driven by PI3Kα Mutations



Multiple Value Inflection Points in 2025 and Beyond





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Dedicated to Making an Impact on Patient's Lives via Transformative Medicines



Thank You

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ENSEM THERAPEUTICS