A phase 1/2, first-in-human study of STX-478 monotherapy or in combination with fulvestrant in patients with breast cancer or other advanced solid tumors (trial in progress)

Alberto J. Montero,¹ Douglas Orr,² Pamela Munster,³ Patricia LoRusso,⁴ Komal Jhaveri,⁵ Timothy Pluard,⁶ Ira Winer,⁷ Michael Streit,⁸ Monica Patterson,⁸ Simon Roberts,⁸ Stefani Corsi-Travali,⁸ Courtney Ewert,⁸ Dejan Juric⁹

¹Department of Medicine, Division of Hematology and Oncology, University Hospitals/Seidman Cancer Center, Case Western Research, Dallas, TX, USA; ³Division of Hematology and Oncology, University of California San Francisco, CA, USA; ⁴Medical Oncology, Yale University, Cleveland, OH, USA; ³Division of Hematology and Oncology, University of California San Francisco, CA, USA; ⁴Medical Oncology, Yale University, Cleveland, OH, USA; ⁴Medical Oncology, Yale University, Cleveland, OH, USA; ⁴Medical Oncology, Cancer Research, Dallas, TX, USA; ⁴Med New Haven, CT, USA; ⁵Memorial Sloan Kettering Cancer Center, New York, NY, USA; ⁶Saint Luke's Cancer Institute, Kansas City, MO, USA; ⁷Karmanos Cancer Institute and Wayne State University, Detroit, MI, USA; ⁸Scorpion Therapeutics, Inc., Boston, MA, USA; ⁹Massachusetts General Hospital Cancer Center, Boston, MA, USA;

Background

- Phosphoinositide 3-kinase α (PI3K α) is one of the most commonly mutated genes across multiple cancers^{1,2}
- Mutations occur throughout the PI3Kα gene, but hotspot mutations predominate in the helical and kinase domains^{1,2}
- Activating PI3K α mutations are most prevalent in breast, gynecological, and squamous cell cancers of the head and neck³⁻⁶
- The therapeutic benefit of isoform-selective PI3Kα inhibition was established with alpelisib (PIQRAY[®]), which has equipotent activity against the wild-type (WT) and mutant enzyme⁷
- Inhibition of WT PI3K α is associated with severe toxicities, including hyperglycemia, rash, and diarrhea that limit widespread alpelisib use⁷
- By selectively targeting mutant PI3K α , a potent small-molecule inhibitor could demonstrate an improved safety- and anti-tumor efficacy profile

STX-478: Profile

- STX-478 binds to a novel, cryptic allosteric site present in both mutant and WT PI3Kα isoforms⁸
- STX-478 has activity against the most common PI3Kα kinase-domain mutation H1047R (inhibitory concentration $[IC_{50}] = 9.4$ nmol/L), with 14-fold selectivity over WT PI3K α (IC₅₀ = 131 nmol/L)⁸
- In a 900-tumor cell line screen, STX-478 selectively inhibited the *in vitro* proliferation of cell lines with kinase-domain and helical-domain mutations compared with cells expressing WT PI3K α^8
- In vivo, STX-478 monotherapy or STX-478 in combination with fulvestrant and/or cyclin-dependent kinase 4/6 inhibitors showed robust and durable tumor regres- sion in human tumor xenograft models harboring PI3K α kinase- or helical-domain mutations⁸
- No toxicities indicative of WT PI3Kα inhibition were observed in preclinical animal models within the expected therapeutic STX-478 dose range
- STX-478 entered clinical development with the potential to provide the bestin-class treatment for patients with PI3K α mutant cancers

Methods

Study Overview

- Multi-part Phase 1/2, first-in-human study ongoing in the USA with expansion into Europe
- Study Design: Traditional 3+3 design for dose escalation with (i) back-fill cohorts and (ii) dose expansion cohorts
- Intra-patient dose escalation allowed
- Study conduct guided by Safety Evaluation Team of participating investigators and sponsor personnel

Study Objectives

- Primary Objective: Determine the Optimal Biologic Dose and Maximum Tolerated Dose and select the Recommended Phase 2 Dose of STX-478 administered as (i) monotherapy and (ii) in combination with fulvestrant
- Secondary Objectives: Characterize the overall safety, tolerability, pharmacokinetics, and preliminary antitumor activity (including exploratory circulating tumor DNA measurements) of STX-478 administered as (i) monotherapy and (ii) in combination with fulvestrant

Study Therapy

- STX-478 once daily oral dosing
- STX-478 in combination with intramuscular administration of fulvestrant (500 mg) on Days 1, 15, 29 and once monthly thereafter

STX-478: Mechanism of Action

Figure 1. STX-478 – Molecular structure⁸

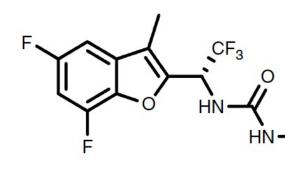


Figure 2. X-ray structure of STX-478 bound in a novel, allosteric site within PI3K α (p110 purple/silver; p85 orange). Inavolisib bound in the adenosine triphosphate site⁸

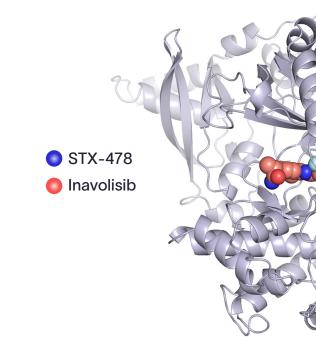
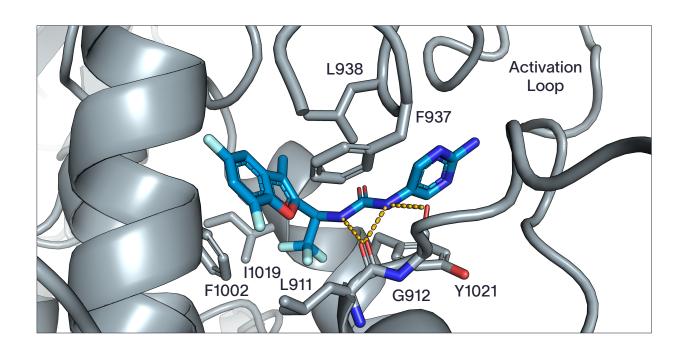


Figure 3. X-ray structure showing facilitated STX-478 binding to the allosteric site by mutation in the kinase domain of PI3K α^8





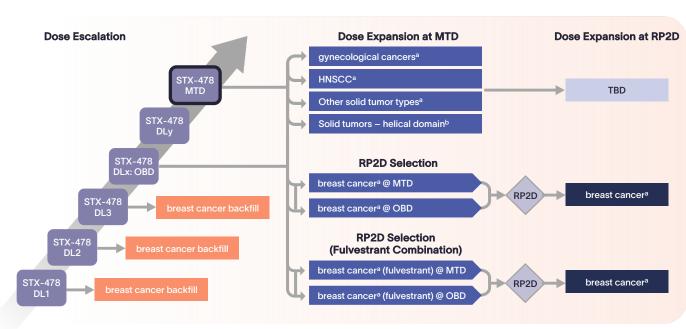
• Expansion to Europe is underway

Study Information

Enrollment

- Clinical Trials.gov: https://clinicaltrials.gov/study/NCT05768139
- Contact clinicaltrials@scorpiontx.com for additional information

Figure 4. Phase 1/2 Study Design



^a PI3Kα kinase domain mutations ^b PI3Kα helical domain mutations

Abbreviations: DL, dose level; HNSCC, head and neck squamous cell carcinoma; MTD, maximum tolerated dose; OBD, optimal biologically active dose; RP2D, recommended Phase 2 dose; TBD, to be determined.

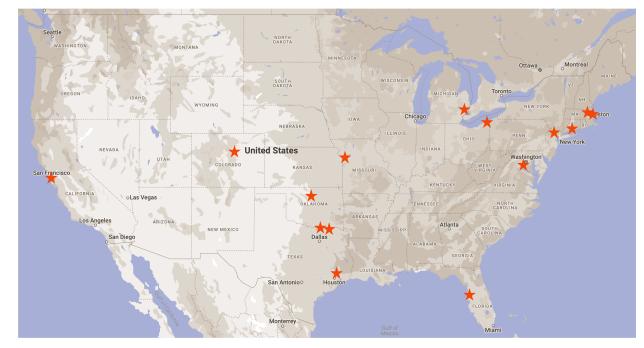
Table1. Key eligibility criteria

Kev inclusion criteria

- Metastatic or locally advanced and unresectable solid tumor
- Confirmed PI3kα-mutation in the kinase and/or helical domain
- Received standard therapy option(s)
- Measurable or evaluable disease
- The Eastern Cooperative Oncology Group (ECOG) performance status score of 0 to 1
- Adequate hematologic and organ function, including fasting glucose <140 mg/dL (7.7 mmol/L) and hemoglobin A1c (glycated hemoglobin; HbA1c) <7.0%

Key exclusion criteria

- Known phosphatase and tensin homolog (PTEN) mutation/deletion or activating AKT mutation
- No prior use of PI3K/AKT/mTOR inhibitors (except prior intolerance)
- Type 1 diabetes or uncontrolled type 2 diabetes
- Symptomatic brain or spinal metastases



Clinical Sites

- 8. Buckbinder L, St Jean DJ, Tieu T, et al. STX-478, a mutant-selective, allosteric PI3Ka inhibitor spares meta-bolic dysfunction and improves therapeutic response in PI3Ka-mutant xenografts. Cancer Discov. 2023.

involved in this study. and the author of this poster.

PO1-20-04

San Antonio **Breast Cancer Symposium[®]** December 5-9, 2023

Figure 5. Participating sites in the USA

- California: University of California, San Francisco
- Colorado: University of Colorado Anschutz Medical Center
- **Connecticut:** Yale University
- Massachusetts: Massachusetts General Hospital; Dana Farber Cancer Center
- Michigan: Karmanos Cancer Institute
- Missouri: Saint Luke's Cancer Institute
- New York: Memorial Sloan Kettering Cancer Center
- Ohio: University Hospitals/Seidman Cancer Center Case Western Reserve University **Oklahoma:** University of Oklahoma
- Texas: Mary Crowley Cancer Research; Texas Oncology-Baylor Charles A. Sammons Cancer Center; The University of Texas MD Anderson Cancer Center
- Virginia: NEXT Virginia
- Florida: Moffitt Cancer Center

References

- 1. Mayer IA, Arteaga CL. The PI3K/AKT pathway as a target for cancer treatment. Annu Rev Med. 2016;67:11-28. 2. Fruman DA, Chiu H, Hopkins BD, Bagrodia S, Cantley LC, Abraham RT. The PI3K pathway in human disease. Cell. 2017;170(4):605-35.
- 3. Zhang Y, Kwok-Shing Ng P, Kucherlapati M, et al. A pan-cancer proteogenomic atlas of PI3K/AKT/mTOR bathway alterations. Cancer Cell. 2017;31(6):820-32 e3.
- 4. Martinez-Saez O, Chic N, Pascual T, et al. Frequency and spectrum of PIK3CA somatic mutations in breast cancer Breast Cancer Res 2020-22(1)-45
- 5. Cerami E, Gao J, Dogrusoz U, et al. The cBio cancer genomics portal: an open platform for exploring
- multidimensional cancer genomics data. Cancer Discov. 2012;2(5):401-4. 6. 6. Arafeh R, Samuels Y. PIK3CA in cancer: The past 30 years. Semin Cancer Biol. 2019;59:36-49.
- 7. Andre F, Ciruelos E, Rubovszky G, et al. Alpelisib for PIK3CA-mutated, hormone receptor-positive advanced breast cancer. N Engl J Med. 2019;380(20):1929-40.

Disclosures

Alberto J. Montero has served in a consulting/advisory role for New Century Health, Welwaze, and Paragon Healthcare; has received honoraria from Celgene, AstraZeneca, and OncoSec; and has received research funding from F. Hoffmann-La Roche Ltd., Basel, Switzerland.

Acknowledgements

- The authors thank the patients and their caregivers, investigators, and study personnel
- This work was supported by Scorpion Therapeutics, Inc.
- Medical writing support was provided by Shereen Cynthia D'Cruz, PhD, of Certara Synchrogenix, under the direction of the authors in accordance with Good Publication Practice guidelines
- (Ann Intern Med 2022;175:1298-1304) and was funded by Scorpion Therapeutics, Inc. Copies of this poster obtained through Quick Response (QR) Code are for personal use only and may not be reproduced without permission from SABCS®
- This presentation is the intellectual property of the author/presenter. Contact them at Alberto.Montero@UHhospitals.org for permission to reprint and/or distribute.

