

#CT057

Phase 2, Multicenter Open-label Basket Trial of *nab*-Sirolimus for Patients With Inactivating Alterations in TSC1 or TSC2 (PRECISION I)

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rapamycin (mTOR) complex 1 activation¹ (**Figure 1**)

Figure 1. PI3K-Akt-mTOR Pathway^{2,3}



4EBPI, eukaryotic translation initiation factor 4E-binding protein; Akt, protein kinase B; Erk, extracellular signal-regulated kinase; Mek, mitogen-activated protein kinase kinase; mTOR, mechanistic target of rapamycin; mTORC1, mTOR complex 1; nab, nanoparticle albumin-bound; PI3K, phosphatidylinositol 3-kinase; PTEN, phosphatase and tensin homolog; Raf, rapidly accelerated fibrosarcoma; Raptor, regulatory-associated protein of mTOR; Ras, rat sarcoma virus homolog; Rheb, Ras homolog enriched in brain; S6K, ribosomal S6 kinase; TSC1/TSC2, tuberous sclerosis complex subunit 1 or 2.

• Inactivating alterations in TSC1 and/or TSC2 have been observed in several types of cancer, but no treatment options exist specifically for patients with *TSC1* or *TSC2* inactivating alterations (**Table 1**)

Table 1. Estimated Frequency of Definite Impact TSC1 or TSC2 Alterations			
Tumor Type	TSC1 Alterations, % ^a	TSC2 Alterations, %	TSC1 or TSC2 Combined, %
Bladder	6.33	1.70	8.03
Hepatobiliary	1.27	3.31	4.58
Endometrial	2.10	1.22	3.32
Soft tissue sarcoma	1.28	1.71	2.99
Ovarian	1.85	0.92	2.77
Esophagogastric	0.65	1.46	2.11
Non-small cell lung cancer	0.77	1.16	1.93
Colorectal carcinoma	0.99	0.39	1.38
Breast	0.41	0.10	0.51

^aThe proportion of patients with definite impact alterations (ie, alterations known to have a biological impact, including frameshift, nonsense, and splice-site mutations and deep deletions) derived from analysis of TCGA and cBioPortal by Gulati et al. (Data on file). TSC1, TSC2, tuberous sclerosis complex subunit 1, 2

- The utility of oral mTOR inhibitors (mTORis), such as sirolimus, as pan-cancer agents may be restricted by low bioavailability and dose-limiting toxicity 2,4
- To improve the pharmacologic properties of sirolimus, *nab*-sirolimus, a nanoparticle form of human albumin-bound sirolimus, was developed for intravenous (IV) use
- In preclinical animal models, *nab*-sirolimus demonstrated significantly higher intratumor drug concentrations, greater tumor growth inhibition, improved survival, and greater inhibition of the downstream marker of mTOR activity, phosphorylated-S6 (pS6) ribosomal protein, relative to equal weekly doses of sirolimus and everolimus (Figure 2)

Figure 2. (A) Intratumor Drug Concentration, (B) Tumor Growth Inhibition, (C) Survival, and (D) mTOR Activity



AUC, area under the curve; IV, intravenous, nab, nanoparticle albumin-bound; PO, per os (orally); pS6, ribosomal protein S6.





- results from the AMPECT trial (NCT02494570)⁶ alterations (**Figure 3**)^{6,7}
- Most treatment-emergent adverse events (TEAEs) were grade 1/2 and were manageable for long-term treatment; no grade ≥4 treatment-related TEAEs were observed
- The overall safety profile was consistent with other mTORis, with no new or unexpected safety signals emerging in the AMPECT trial

Figure 3. Percent Change in Target Tumor Measurements Over Time



- Patients with various malignancies bearing TSC1 or TSC2 alterations treated with *nab*-sirolimus as part of the expanded access program (NCT03817515) showed evidence of response (partial response in 5/7 patients) and manageable toxicities⁸
- The phase 2 PRECISION I trial was initiated to evaluate the potential of mTOR inhibition with nabsirolimus for the treatment of patients with solid tumors harboring TSC1 or TSC2 inactivating alterations

STUDY DESIGN

with pathogenic inactivating alterations in *TSC1* (Arm A) or *TSC2* (Arm B) (**Figure 4**)

Figure 4. Study Design



^aCentral confirmation of TSC1 and TSC2 pathogenic inactivating alterations is via evaluation of NGS reports. Patients will be enrolled only after central confirmation of eligibility. TSC1 and TSC2 alterations should be identified using NGS in tumor tissue or liquid biopsy and must be determined by analytically validated NGS tests performed in CLIA-certified laboratories. ^bFollow-up is for survival and initiation of anticancer therapy. Follow-up is initiated after the EOS visit. CLIA, Clinical Laboratory Improvement Amendments; DCR, disease control rate; DOR, duration of response; ECOG, Eastern Cooperative Oncology Group; EOS, end of study; IRR, interrater reliability; KPS, Karnofsky Performance Scale; LPPS, Lansky Play-Performance Scale; mTORi, mechanistic target of rapamycin inhibitor; *nab*, nanoparticle albumin-bound; NGS, next-generation sequencing; ORR, overall response rate; OS, overall survival; PFS, progression-free survival; PS, performance status; QOL, quality of life; RECIST, Response Evaluation Criteria for Solid Tumors; TSC1/TSC2, tuberous sclerosis complex 1 or 2. Reproduced from Schulte et al.⁷

- *nab*-Sirolimus is a novel albumin-bound mTOR inhibitor (mTORi) and is approved in the United States for the treatment of adult patients with locally advanced, unresectable or metastatic malignant perivascular epithelioid cell tumor (PEComa)⁵ based on clinical efficacy and safety
- An exploratory biomarker analysis from AMPECT demonstrated that known inactivating alterations in TSC1 or TSC2 were associated with response to *nab*-sirolimus in PEComa, suggesting *nab*sirolimus might be beneficial for patients with other solid tumors harboring TSC1 or TSC2

• PRECISION I (NCT05103358) is a prospective, phase 2, open-label, multi-institution basket trial to determine the efficacy and safety profile of *nab*-sirolimus in patients with malignant solid tumors

Table 2. Key Inclusion and Exclusion Criteria Key Inclusion Criteria

≥12 years of age

Metastatic or locally advance pathogenic inactivating TSC: surgical resection is not an o severe morbidity

Received appropriate standar tolerate or derive clinically me standard therapy as determin

ECOG performance status of play-performance scale for pe

≥1 measurable target lesion k

The full list of inclusion and exclusion criteria is available at https://www.clinicaltrials.gov/ct2/show/NCT05103358

Partnerships



SUMMARY

tissue biopsy

- alterations in *TSC1* and/or *TSC2*

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	Key Exclusion Criteria
	Prior treatment with mTOR inhibitor
d malignant solid tumor with a or <i>TSC2</i> alteration, where otion, or likely to result in	Recent infection requiring systemic anti- infective treatment
rd treatments or be unlikely to eaningful benefit from led by the investigator	Primary brain tumor or PEComa
⁷ 0 or 1, KPS ≥80, or Lansky ediatric patients ≥80	Severe and/or uncontrolled medical or psychiatric conditions
by CT scan or MRI	For patients on strong inhibitors, inducers, and known CYP3A4 substrates, discontinuation is required \geq 5 half-lives prior to receiving the first dose of <i>nab</i> -sirolimus
	T05400050

CT, computed tomography; CYP3A4, cytochrome P450 3A4; ECOG, Eastern Cooperative Oncology Group; KPS, Karnofsky performance status; mTOR, mechanistic target of rapamycin; MRI, magnetic resonance imaging; nab, nanoparticle albumin-bound; PEComa, perivascular epithelioid tumor; TSC1/TSC2, tuberous sclerosis complex subunit 1 or 2.

 Partnerships with leading next-generation sequencing companies (Foundation Medicine, Tempus, and Caris) and US Oncology will facilitate identification of patients with qualifying inactivating TSC1 or TSC2 alterations and expand access to the study through just-in-time trial locations and accelerated site activation (Figure 5)

Figure 5. Expanding PRECISION I Access and Enrollment Through Strategic



PRECISION I site, and site notified of eligible patient





enrolled in trial

• Available data from the AMPECT exploratory analysis and an expanded access program suggest acceptable efficacy and safety of *nab*-sirolimus, an mTORi with enhanced antitumor activity, in patients with solid tumors harboring inactivating

• PRECISION I is a registrational basket trial for patients with solid tumors driven by *TSC1* or *TSC2* alterations; enrollment began in March 2022

• This trial is designed to evaluate the efficacy, safety, and tolerability of *nab*-sirolimus in a patient population with advanced malignancies and limited therapeutic options

 Collaboration with leading next-generation sequencing vendors will expedite the identification of patients with qualifying TSC1 or TSC2 alterations; study access will be facilitated through a "just-in-time" approach to trial location activation