



Copies of this poster obtained through Quick Response (QR) Code are for personal use only and may not be reproduced without permission from WILEY or the author of this poster.

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

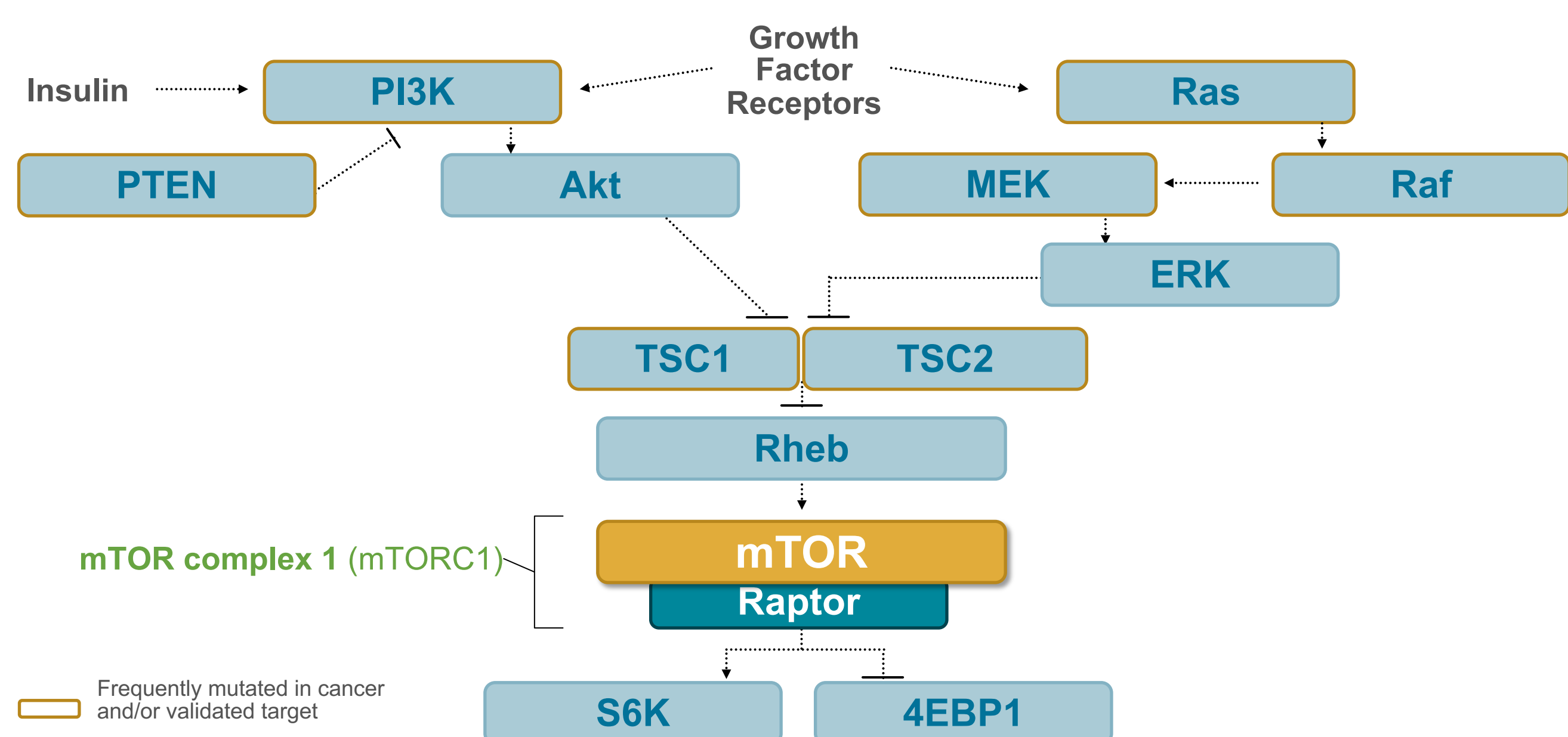
Gopa Iyer, MD¹; Michael J. Demeure, MD²; Li Ding, MS, MA³; Anita N. Schmid, PhD³; Willis H. Navarro, MD³; David J. Kwiatkowski, MD, PhD⁴; Jordi Rodon Ahnert, MD, PhD⁵

¹Memorial Sloan Kettering Cancer Center, New York, NY, USA; ²Hoag Memorial Hospital Presbyterian, Newport Beach, CA, USA; ³Aadi Bioscience, Pacific Palisades, CA, USA; ⁴Brigham and Women's Hospital, Boston, MA, USA; ⁵MD Anderson Cancer Center, Houston, TX, USA

INTRODUCTION

- Tuberous sclerosis complex subunit 1 and 2 (*TSC1*, *TSC2*) form a protein complex and together are critical negative regulators of mechanistic target of rapamycin (mTOR) complex 1 activation¹ (**Figure 1**)

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



4EBP1, 4E binding protein 1; Akt, Akt serine/threonine protein kinase; ERK, extracellular signal-regulated kinases; MEK, mitogen-activated protein kinase; mTOR, mechanistic target of rapamycin; PI3K, phosphatidylinositol 3-kinase; PTEN, phosphatase and tensin homolog; Raf, rapidly accelerated fibrosarcoma protein; Raptor, regulatory-associated protein of mTOR; Ras, rat sarcoma virus homolog; Rheb, Ras homolog enriched in brain; S6K, ribosomal protein S6 kinase; TSC, tuberous sclerosis complex; TSC1/TSC2, tuberous sclerosis complex 1/2.
1. Saxton RA, et al. *Cell*. 2017;169(6):960-976. 2. Hou S, et al. Poster presented at: AACR-NCI-EORTC Molecular Targets and Cancer Therapeutics, October 7-10, 2021 [virtual]. Poster P138. 3. Hay N, et al. *Genes Dev*. 2004;18(16):1926-1945. 4. Mossmann D, et al. *Nat Rev Cancer*. 2018;18(12):744-757.

- 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**)

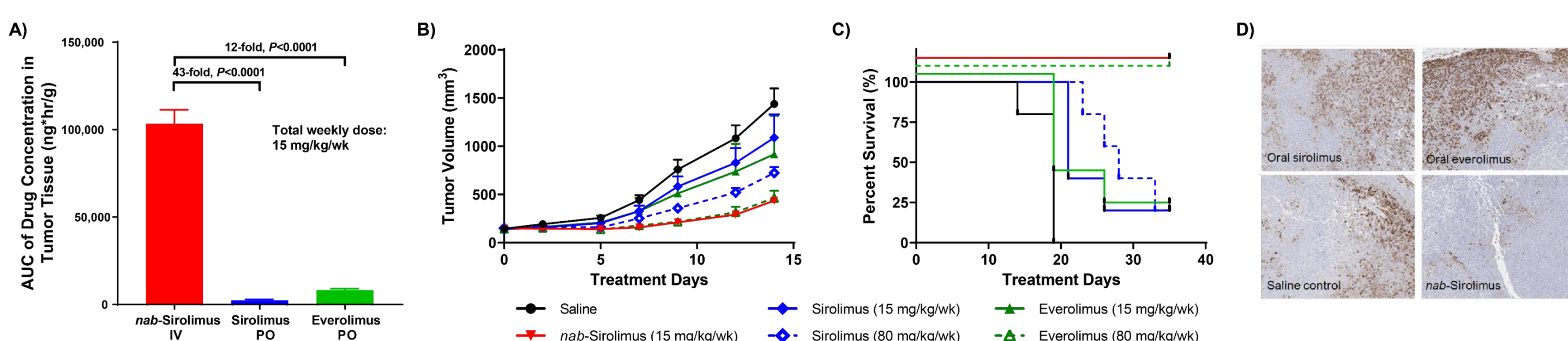
Table. Estimated Frequency of Definite Impact *TSC1* or *TSC2* Alterations

Tumor Type	<i>TSC1</i> Alterations, % ^a	<i>TSC2</i> Alterations, %	<i>TSC1</i> or <i>TSC2</i> 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.

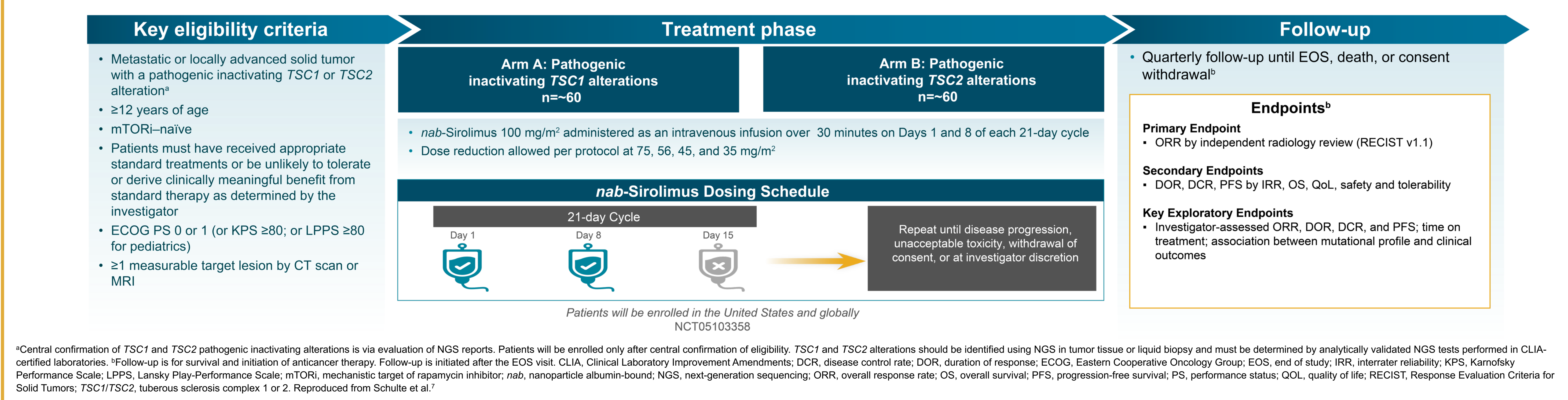
- The utility of oral mTOR inhibitors (mTORis), such as sirolimus, as pan-cancer agents may be restricted by low bioavailability and dose-limiting toxicity^{2,4}
- To improve the pharmacologic properties of sirolimus, *nab*-sirolimus, a nanoparticle form of human albumin-bound sirolimus, was developed 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 everolimus³ (**Figure 2**)

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



Results from *PTEN*-null UMC3 bladder cancer model. AUC, area under the curve; IV, intravenous, *nab*, nanoparticle albumin-bound; PO, per os (orally); pS6, ribosomal protein S6.

STUDY DESIGN

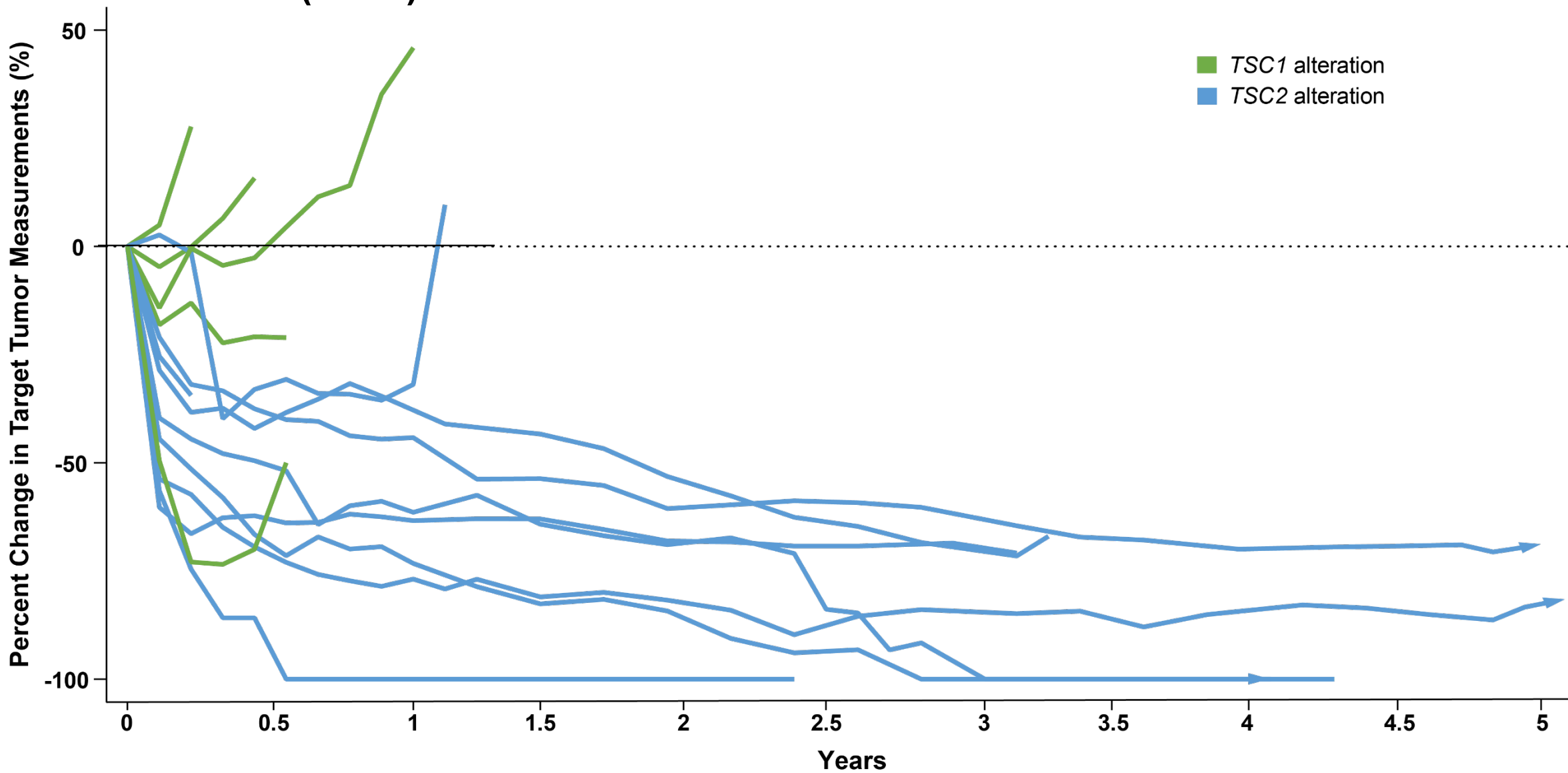


^aCentral confirmation of *TSC1* and *TSC2* pathogenic inactivating alterations is via evaluation of NGS reports. Patients will be enrolled only after central confirmation of eligibility. *TSC1* and *TSC2* alterations should be identified using NGS in tumor tissue or liquid biopsy and must be determined 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

INTRODUCTION (cont.)

- nab*-Sirolimus is a novel albumin-bound 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 (overall response rate 38.7%) and safety results from the AMPECT trial (NCT02494570)⁶
- Results from the AMPECT exploratory biomarker analysis demonstrated rapid and durable responses in patients with *TSC1* or *TSC2* inactivating alterations and suggested significant clinical benefit (**Figure 3**)^{6,7}
 - The most common nonhematologic TRAEs were stomatitis (28/34 [82%]) and fatigue and rash (21/34 [62%] each), and the most common hematologic TRAEs were anemia (18/34 [53%]) and thrombocytopenia (12/34 [35%])
 - 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
- 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 *nab*-sirolimus for the treatment of patients with solid tumors harboring *TSC1* or *TSC2* inactivating alterations

Figure 3. Exploratory Biomarker Analysis From AMPECT Demonstrated Rapid and Durable Responses in Patients With *TSC1* or *TSC2* Alteration Treated With *nab*-Sirolimus (n=14)



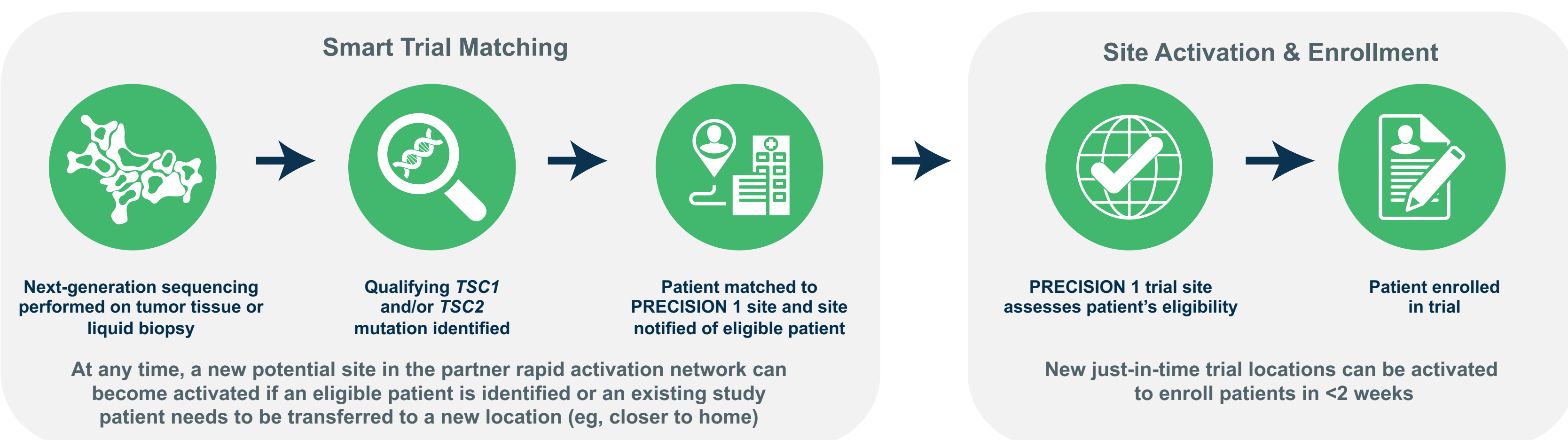
REFERENCES

- He Y, et al. *Signal Transduct Target Ther*. 2021;6(1):425. 2. Saxton RA, et al. *Cell*. 2017;169(2):361-371. 3. Hou S, et al. AACR-NCI-EORTC, October 7-10, 2021 [virtual]. Poster P138. 4. Pavavra F, et al. *Oxid Med Cell*. 2017;9820181. 5. FYARRO (sirolimus albumin-bound particles for injectable suspension). Package insert. Aadi Bioscience, Inc.; Pacific Palisades, CA; 2021. 6. Wagner AJ, et al. *J Clin Oncol*. 2021;39(33):3660-3670. 7. Schulte B, et al. CTOS, November 16-19, 2022, Vancouver, BC. Poster 313. 8. Dickson MA, et al. ASCO, June 4-8, 2021, Virtual.

STUDY DESIGN AND FUTURE PARTNERSHIPS

- 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 with pathogenic inactivating alterations in *TSC1* (Arm A) or *TSC2* (Arm B) (**Study Design inset**)
- 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 4**)

Figure 4. Expanding PRECISION I Access and Enrollment Through Strategic Partnerships



SUMMARY

- nab*-Sirolimus is an mTORi utilizing *nab* technology to enhance antitumor activity in non-clinical animal models
- Available data from the AMPECT exploratory analysis and an expanded access program suggest *nab*-sirolimus will provide clinically relevant benefit with a manageable safety profile in patients with solid tumors harboring inactivating alterations in *TSC1* and/or *TSC2*
- 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

DISCLOSURES: GI: no conflicts of interest. MJD: Consulting fees: Bayer, Eli Lilly, On Cusp Therapeutics, and TD2. LD, ANS, WHN: Employment and stock ownership: Aadi Bioscience, Inc. DJK: Research funding: Aadi Bioscience, Inc., Genentech, Revolution Medicines; consulting fees: Aadi Bioscience, Inc., Bridgebio, Genentech, Guidepoint. JRA: Research funding: Bayer, Novartis; personal fees: Eli Lilly, Peptomyc, and Servier.

ACKNOWLEDGMENTS: Medical writing and editorial assistance were provided by Cynthia D. Gioiello, PharmD, and Stephen Bublitz, ELS, of MedVal Scientific Information Services, LLC, and were funded by Aadi Bioscience, Inc.