

# A dose escalation and cohort expansion study of the CDK9 inhibitor KB-0742 in relapsed, refractory and transcriptionally addicted solid tumors

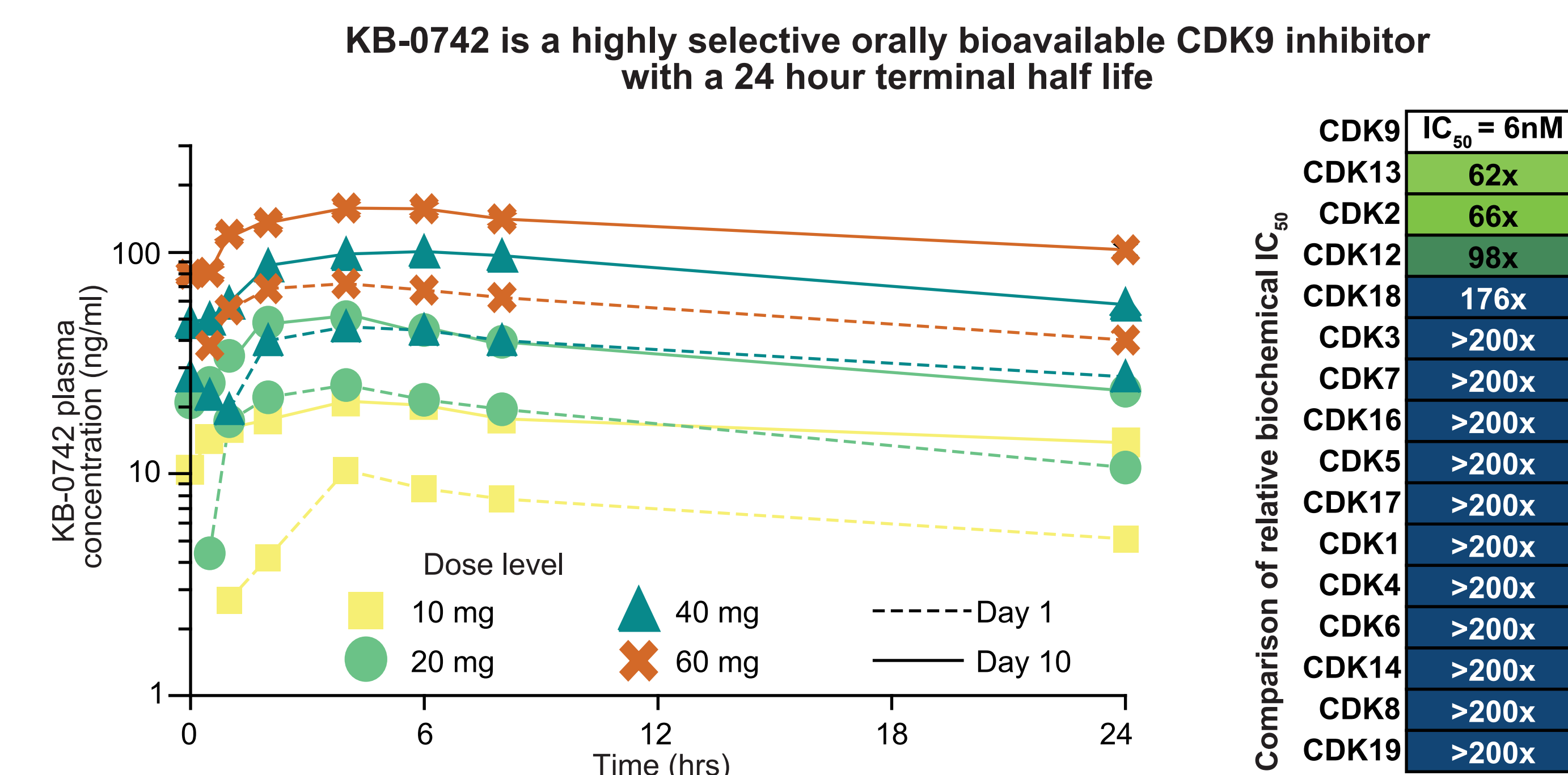
Miguel Villalona-Calero<sup>1</sup>, Mark Agulnik<sup>1</sup>, Monica Mita<sup>2</sup>, Alain Mita<sup>2</sup>, Noah Federman<sup>3</sup>, Drew W. Rasco<sup>4</sup>, David Spigel<sup>5</sup>, Jia Luo<sup>6</sup>, Glenn Hanna<sup>6</sup>, Gregory M. Cote<sup>7</sup>, Mohamed Adham Salkeni<sup>8</sup>, Rashmi Chugh<sup>9</sup>, Natraj J. Ammakannavar<sup>10</sup>, Satish A. Shah<sup>11</sup>, Amol Rao<sup>12</sup>, Kamallesh Kumar Sankhala<sup>13</sup>, Kathleen Guindon<sup>14</sup>, Richard E. Cutler<sup>14</sup>, Tressa R. Hood<sup>14</sup>, Luis A. Carvajal<sup>14</sup>, Charles Y. Lin<sup>14</sup>, Jorge F. DiMartino<sup>14</sup>, Elizabeth A. Olek<sup>14</sup>, Brian A. Van Tine<sup>15</sup>

<sup>1</sup>City of Hope National Medical Center, Duarte, CA; <sup>2</sup>Cedars-Sinai Cancer Institute, Los Angeles, CA; <sup>3</sup>UCLA Jonsson Comprehensive Cancer Center, Los Angeles, CA; <sup>4</sup>START Center for Cancer Care, San Antonio, TX; <sup>5</sup>Sarah Cannon Research Institute at Tennessee Oncology, Nashville, TN; <sup>6</sup>Dana Farber Cancer Institute, Boston, MA; <sup>7</sup>Mass General Cancer Center, Boston, MA; <sup>8</sup>Virginia Cancer Specialists, Fairfax, VA; <sup>9</sup>University of Michigan, Ann Arbor, MI; <sup>10</sup>Community Cancer Center North, Indianapolis, IN; <sup>11</sup>Pennsylvania Cancer Specialists and Research Institute, Gettysburg, PA; <sup>12</sup>Memorial Care, Fountain Valley, CA; <sup>13</sup>NextGen Oncology, Beverly Hills, CA; <sup>14</sup>Kronos Bio, Inc., San Mateo, CA; <sup>15</sup>Washington University in St. Louis, St. Louis, MO

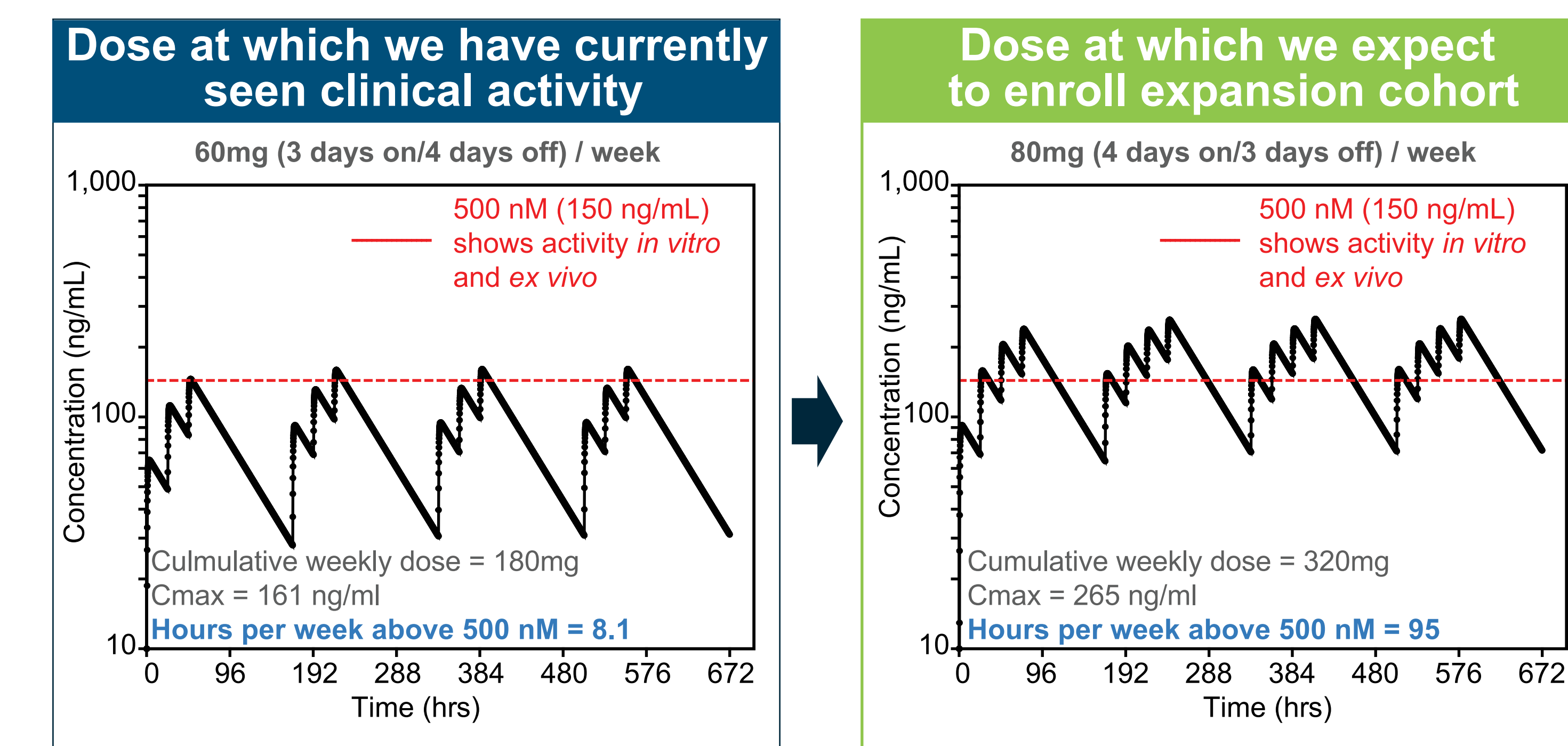
## Background and rationale for targeting CDK9 in transcriptionally addicted tumors

- Two categories of transcriptionally addicted tumors are sarcomas with transcription factor fusions and tumors with MYC deregulation.
- We have previously shown clinical benefit in myxoid liposarcomas with TF fusions (Villalona-Calero et al., abstract B159, AACR-NCI-EORTC, 2023).
- Analysis of a real-world solid tumor cohort including NSCLC (n= 20,470), SCLC (n=1,517), TNBC (n=2,576) and ovarian cancer (n=1,667) demonstrated MYC family overexpression and/or genomic amplification in 47% to 87% of patients.
- Both TF-fusion and MYC deregulated tumors exhibit high dependency on cyclin-dependent kinase 9 (CDK9) activity.
- KB-0742 is a potent, selective, and orally bioavailable inhibitor of CDK9 with a long plasma half-life. KB-0742 is being evaluated in an ongoing Phase 1/2 study in advanced solid tumors (NCT04718675).
- KB-0742 is dosed orally once daily for three consecutive days followed by four days off in 28-day cycles until unacceptable toxicity or disease progression. Based on pharmacokinetic data, alternate dosing schedules are being explored with KB-0742 dosed four consecutive days on, followed by three days off treatment schedule.

## KB-0742 plasma concentrations at day 10 of dosing across the 10mg to 60mg cohorts



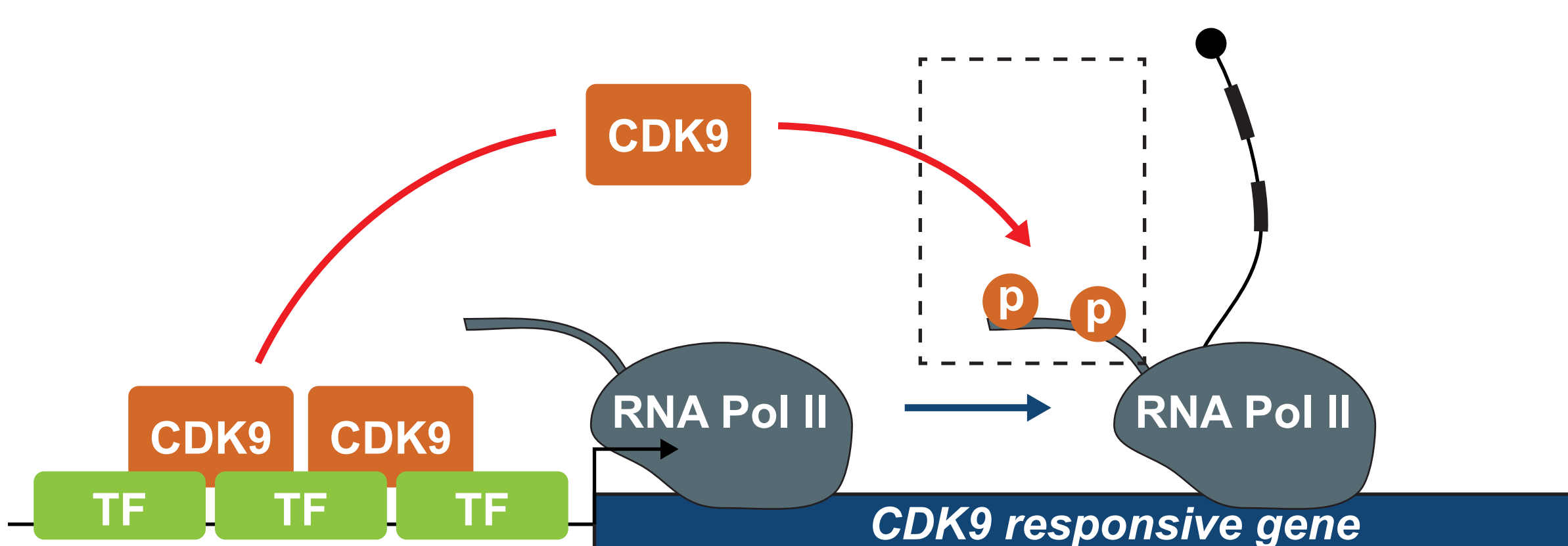
## KB-0742 long half-life and kinase selectivity allows selective targeting of transcriptional addiction



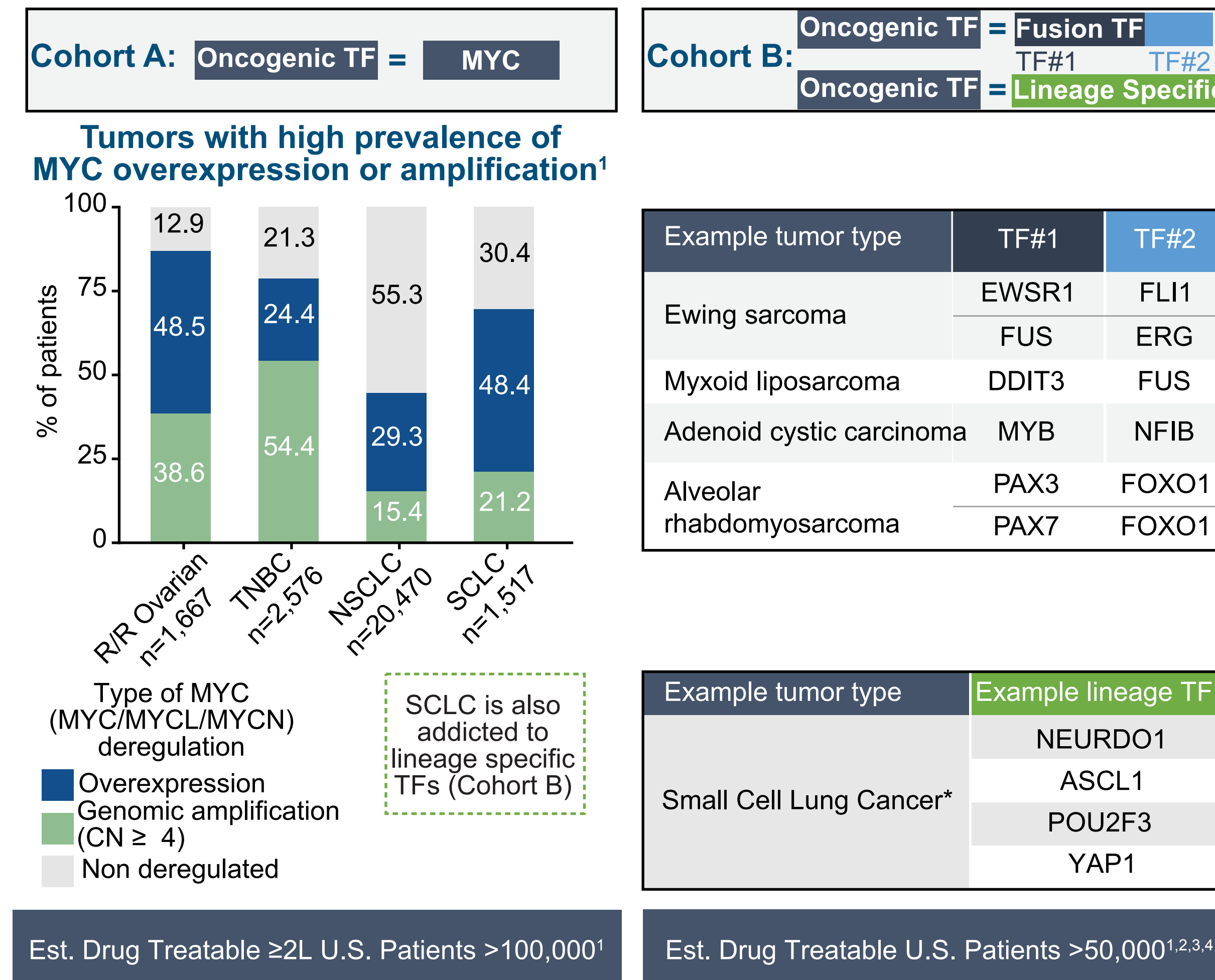
## CDK9 is a critical cofactor of transcriptionally addicted tumors

**CDK9** is a kinase that is recruited to the genome by transcription factors and drives transcription elongation at many oncogenes

Phosphorylated serine 2 (pSER2) of the RNA polymerase II C-terminal domain is a key CDK9 kinase substrate

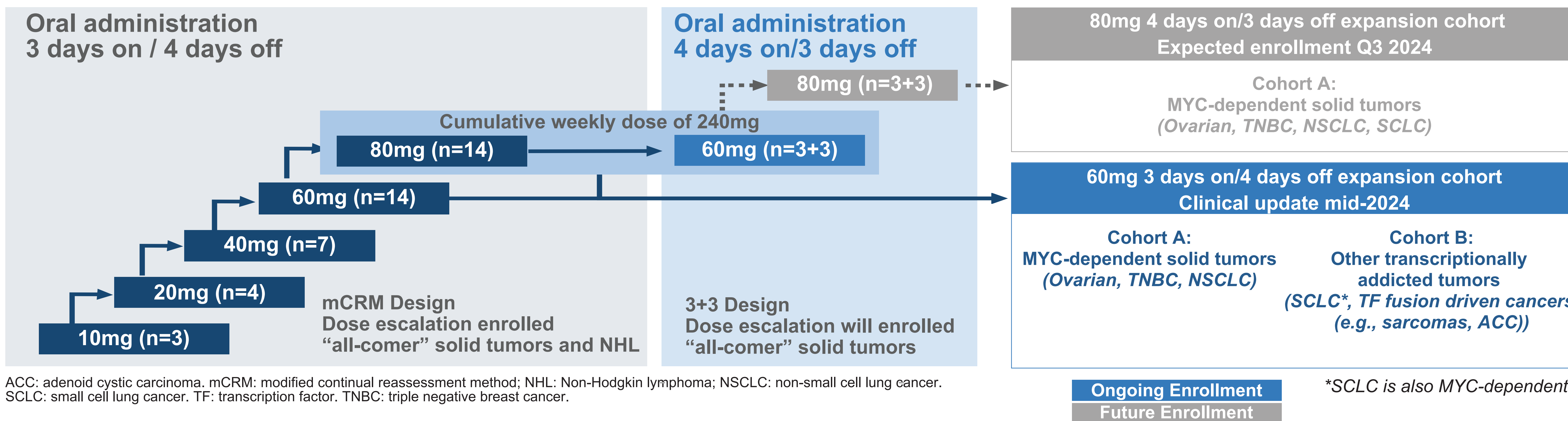


## Cohorts with CDK9-dependent transcriptional addiction



## KB-0742 – 1001 overall trial design (NCT04718675)

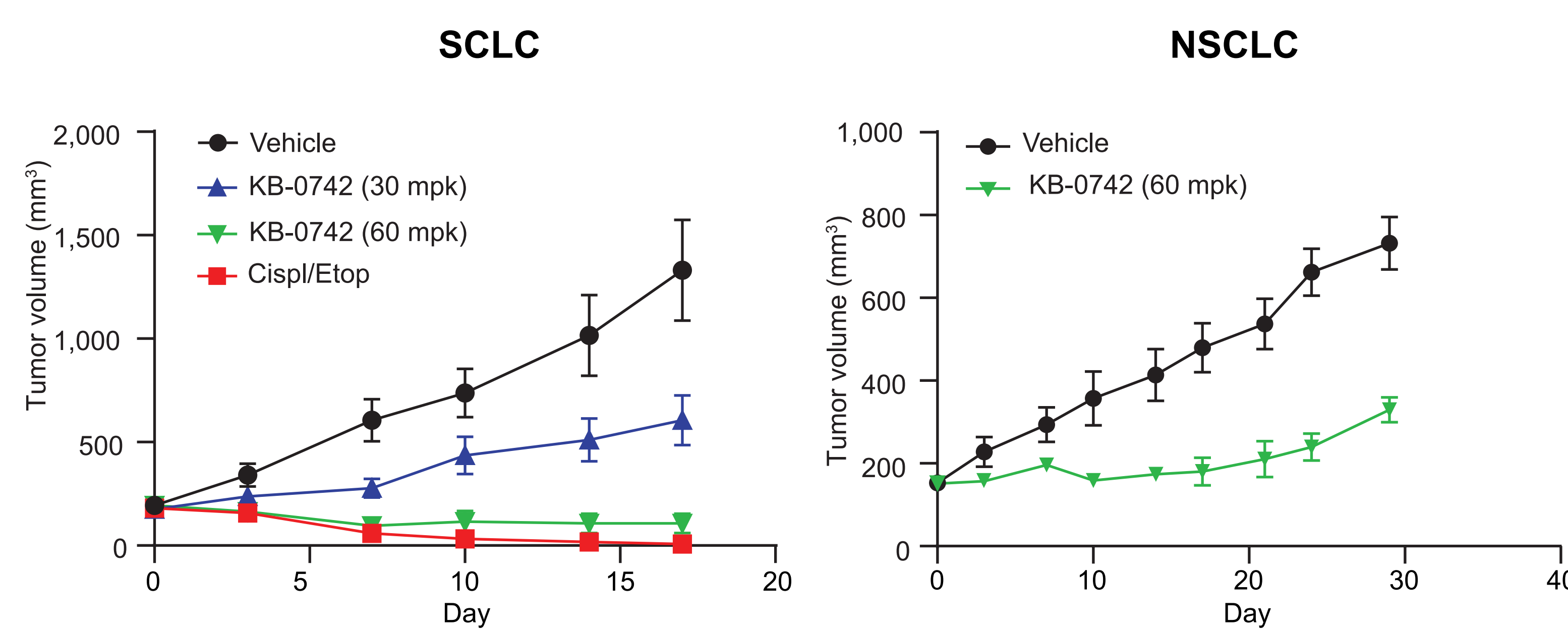
Phase 1/2, first-in-human, open-label dose escalation and expansion study of KB-0742 in patients with relapsed or refractory solid tumors or NHL



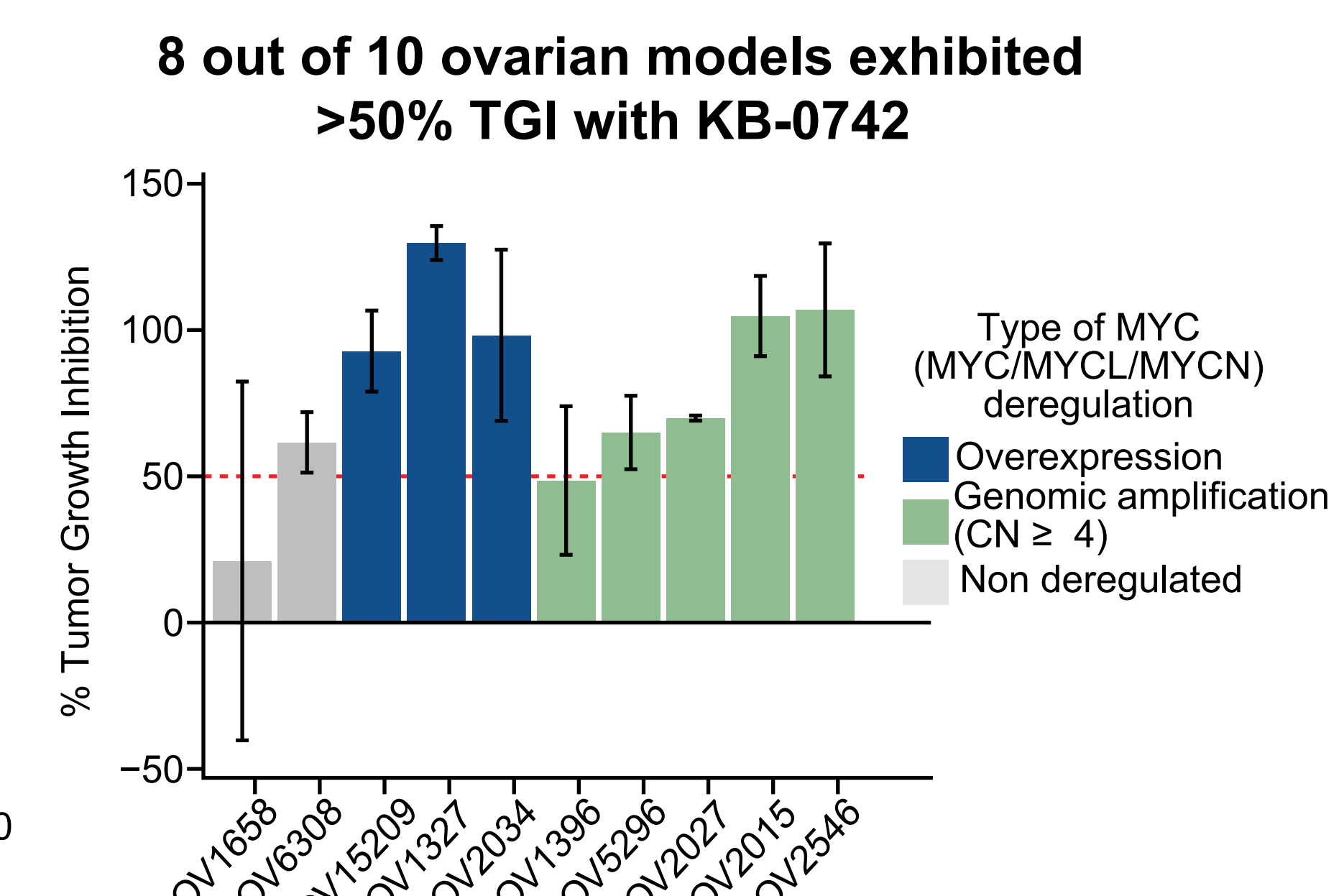
ACC: adenoid cystic carcinoma, mCRM: modified continual reassessment method; NHL: Non-Hodgkin lymphoma; NSCLC: non-small cell lung cancer. SCLC: small cell lung cancer. TF: transcription factor. TNBC: triple negative breast cancer.

## KB-0742 drives tumor growth inhibition in MYC deregulated *in vivo* tumor models

### KB-0742 shows monotherapy anti-tumor activity in lung patient-derived xenografts



### KB-0742 activity in ovarian cancer patient-derived xenograft (PDX) models



Left: KB-0742 showed antitumor activity in PDX models of SCLC and NSCLC. PDX models were treated with KB0742 at either 30 and 60mg/kg on a dosing schedule of 3 days on/4 days off. KB-0742 showed antitumor activity in a dose-dependent manner with 92% and 57% TGI in SCLC and NSCLC at 60mg/kg, respectively. Right: Ovarian cancer mouse clinical trial results demonstrating broad single agent activity in MYC family amplified or overexpressed subtypes. Three animals per model were enrolled in the study.<sup>6</sup>

- Eligibility criteria**
- Any R/R solid tumor with readily accessible biopsy sites and consenting to 1 baseline and 1 on-treatment biopsy
  - Age ≥ 18, acceptable organ function and ECOG PS < 2
  - Tumor types of interest include:
    - Small cell lung cancer
    - Ovarian, TNBC, or NSCLC
    - Diffuse large B-cell lymphoma or Burkitt's lymphoma with MYC translocation
    - Sarcoma of histologic subtype known to be associated with TF Fusion
    - Chordoma, nut midline carcinoma, adenoid cystic carcinoma

- Eligibility criteria**
- Solid tumors that have failed, are intolerant to, or are ineligible for standard of care anticancer treatments
  - Cohort A:** Tumor types with high prevalence of MYC deregulation: ovarian, triple negative breast cancer, non small cell lung cancer
  - Cohort B:** Small cell lung cancer and transcriptionally addicted tumor types including sarcomas of histologic subtypes associated with TF fusion, chordoma, ACC and nut midline carcinoma

## Study objectives

- Primary**
- Evaluate safety and tolerability of KB-0742 in defined patient cohorts
- Secondary**
- Characterize the PK of KB-0742
  - Characterize anti-tumor activity of KB-0742

## Study endpoints

- Primary**
- Descriptive safety analysis of treatment-emergent adverse events (TEAEs), serious adverse events (SAEs), dose limiting toxicities (DLTs), and laboratory assessments
- Secondary**
- Evaluation of the pharmacokinetic profile, progression free survival (PFS), overall response rate (ORR) and duration of response (DOR)

## Summary

- KB-0742 is an oral highly selective CDK9 inhibitor with an attractive profile including a long half-life.
- Preliminary data have been previously reported; Villalona-Calero et al., Abstract B159, AACR-NCI-EORTC, 2023.
- The ongoing trial KB-0742-1001 will continue to enroll and examine additional dosage/schedules (NCT04718675).