

ATRN-119, a Novel Macrocyclic ATR Inhibitor, in Patients with Advanced Solid Malignancies: A Phase 1/2a Trial (ABOYA-119)

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INTRODUCTION

DNA damage and response (DDR) and Ataxia Telangiectasia and Rad3-related (ATR) kinase

- DDR is key to a cell's ability to maintain genomic stability, and its deficiency has been characterized in many cancer types^{1,2}
- ATR kinase is a principal regulator of DDR; when activated, it stabilizes stressed DNA replication forks and prevents their collapse into DNA double-strand breaks
- ATR inhibition may be a promising strategy in cancer therapeutics^{1,3}
- Advanced solid tumors with dysregulated DDR or oncogenic stress may be more susceptible to ATR kinase inhibition

ATRN-119

- ATRN-119 is a differentiated macrocyclic and highly selective, potent, oral ATR inhibitor with antitumor activity correlated with specific DDR, tumor suppressor, or oncogene alterations
- Preliminary studies showed improved selectivity correlated with increased tolerability²
- Both *in vitro* and *in vivo* studies of ATRN-119 demonstrate significant anticancer effects in DDR deficient models, both alone and in combination with other targeted treatments, such as PARP inhibitors and WEE1 inhibitors^{2,4}
- Here, we summarize preliminary results of this ongoing first-in-human study of ATRN-119

STUDY OBJECTIVES

Primary objectives and endpoints

- Evaluate the safety profile of escalating doses of ATRN-119
- Determine the MTD and RP2D
- Characterize the PK profile of ATRN-119 and its active metabolite ATRN-157

Secondary objectives and endpoints

- Evaluate antitumor activity of ATRN-119 in various solid tumors
- Exploratory objective
 - Explore the association between mutations identified in tumor tissue and clinical outcomes

KEY ELIGIBILITY CRITERIA

Inclusion criteria

- ≥ 12 years old with advanced solid tumor harboring ≥ 1 NGS-confirmed specific DDR mutation:
 - Any mutation in *ARID1A*
 - Any missense mutation in *KRAS* at *Gly12* or *Gly13*
 - Probable loss of function mutation in select genes
 - Amplification (>4 genomic copies in total) of *MYC*, *CCNE1* or *CCNE2*
- Merkel cell carcinoma regardless of known mutation status
- Measurable disease per RECIST v1.1 (PCWG3 criteria for mCRPC)
- Failed ≥ 1 approved SOC therapy
- ECOG PS ≤ 1

Exclusion criteria

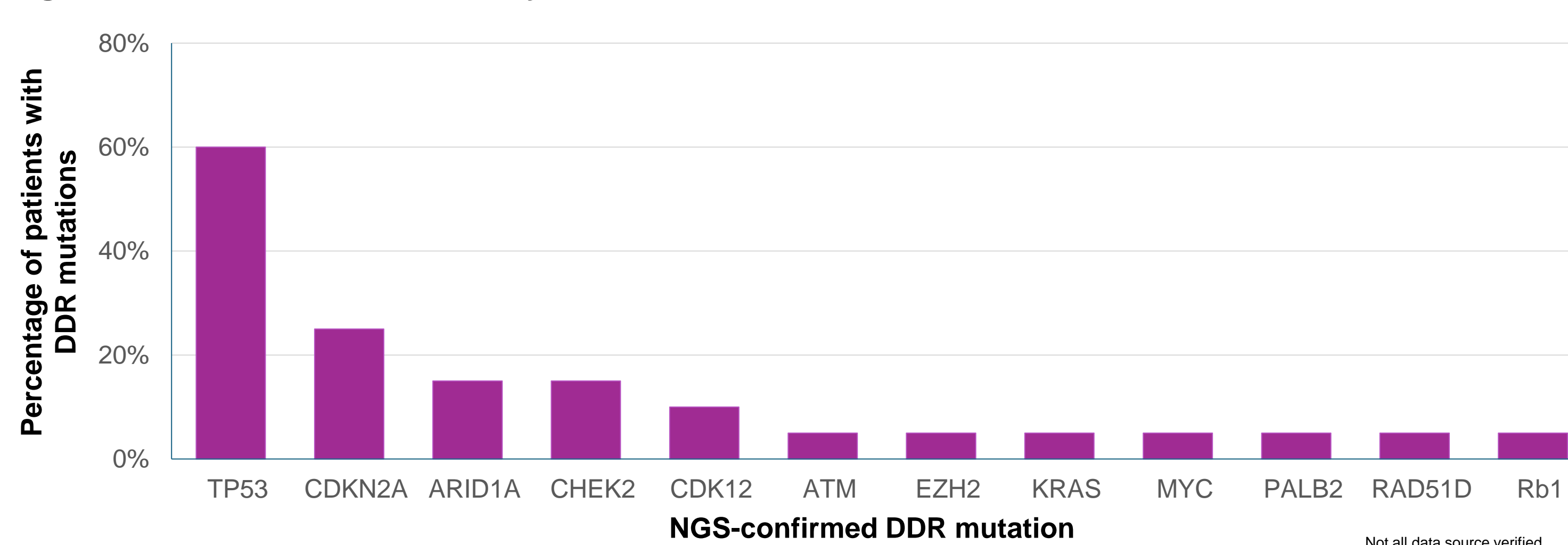
- Known additional malignancy that is progressing or requires active treatment (with some exceptions)
- Known CNS metastases/involvement that is not treated and stable for the previous 1 month
- Cytotoxic chemotherapy, immunotherapy, radiotherapy, or targeted therapies within 4 weeks or ≥ 5 half-lives, and all prior therapy-related AEs are not at baseline/stable
- Concomitant treatment with strong inhibitors or inducers of CYP3A4 or CYP2D6

PATIENT DEMOGRAPHICS

Table 1. Baseline demographics

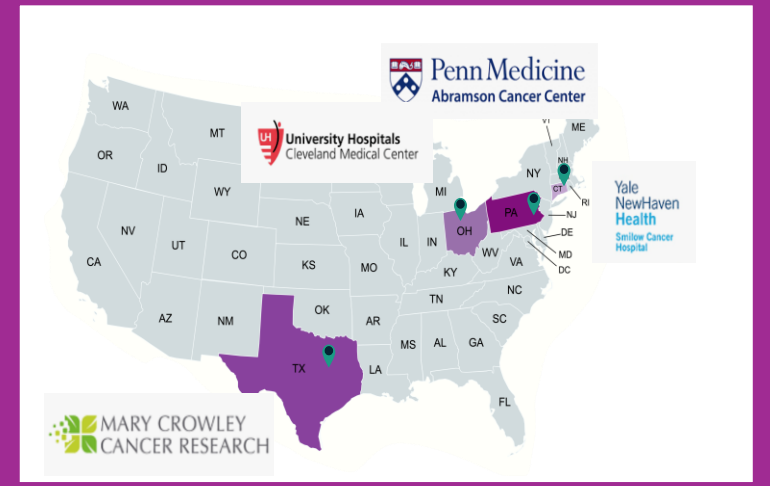
Characteristic	Study patients (n=20)	Characteristic	Study patients (n=20)
Sex, n (%)		Tumor type, n (%)	
Male	7 (35%)	Colorectal carcinoma	5 (25%)
Female	13 (65%)	Breast cancer	3 (15%)
Median age (range), years	62 (42 - 79)	Lung cancer	2 (10%)
Race, n (%)		Pancreatic cancer	2 (10%)
White	15 (75%)	Adrenal cortical carcinoma	1 (5%)
Black or African American	5 (25%)	Appendiceal adenocarcinoma	1 (5%)
ECOG PS, n (%)		Duodenal cancer	1 (5%)
0	5 (25%)	Endometrial cancer	1 (5%)
1	15 (75%)	Fallopian tube adenocarcinoma	1 (5%)
Prior lines of systemic chemotherapies, n (%)		Ovarian cancer	1 (5%)
< 2	2 (10%)	Prostate cancer	1 (5%)
2 - 3	9 (45%)	Adenocarcinoma of unknown primary	1 (5%)
≥ 4	9 (45%)		
Prior systemic therapy, n (%)			
Platinum-based chemotherapy	16 (80%)		
Immuno-oncology	4 (20%)		
PARP inhibitor	2 (10%)		

Figure 1. DDR mutations at study enrollment



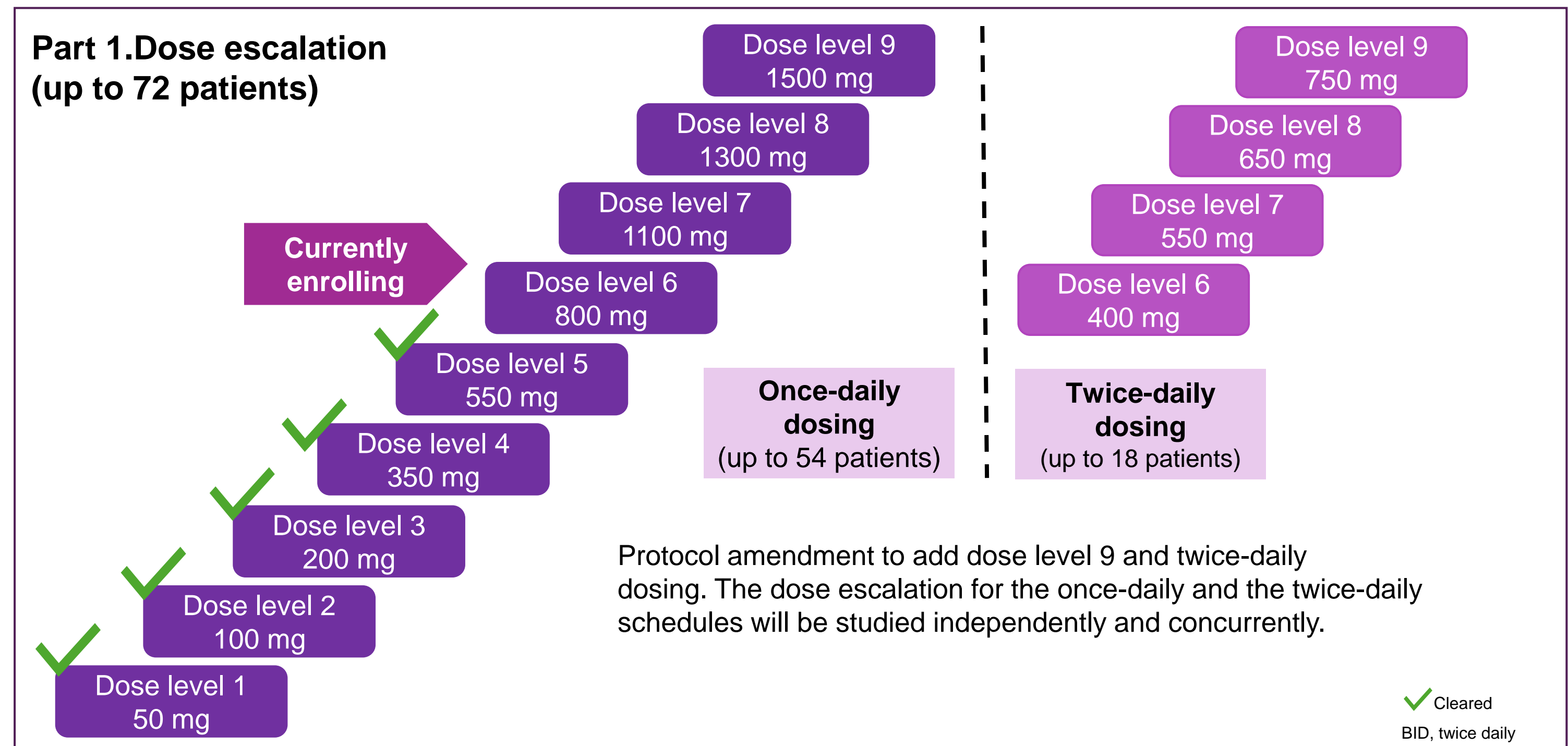
SUMMARY

- In this ongoing first-in-human DNA study of oral ATRN-119 in patients with advanced solid tumors harboring ≥ 1 specific DDR mutation, continuous daily administration up to and including 800 mg is shown to be safe and tolerable
- As of Oct 2, 2024, most AEs are Grade 1 or 2 and there are no DLTs, no SAEs, and no ATRN-119-related AEs ≥ Grade 4
- PK studies demonstrate increasing exposure with escalating doses
- Preliminary signs of clinical benefit have been observed in two patients treated at the 50 mg and 200 mg dose level
- This study is currently in the dose escalation phase with planned addition of a twice-daily ATRN-119 dosing regimen
- Active enrollment is ongoing at four sites in the U.S. (NCT04905914)



STUDY SCHEMA

Figure 2. 3+3 dose escalation and dose expansion (up to 132 patients)



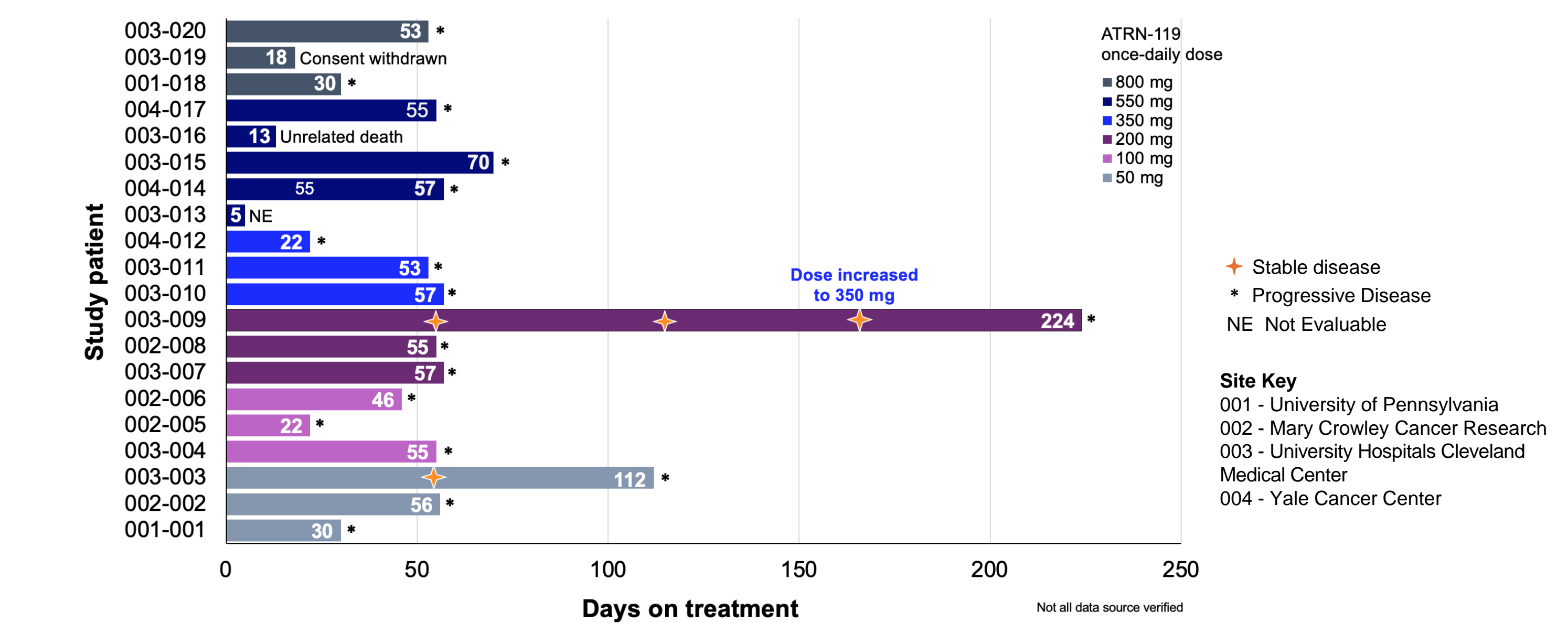
Part 2. Dose expansion (up to 60 patients)

- Single-agent ATRN-119 after MTD/RP2D is established
- Potential indications: colorectal, prostate, gastric, endometrial

DURATION OF TREATMENT

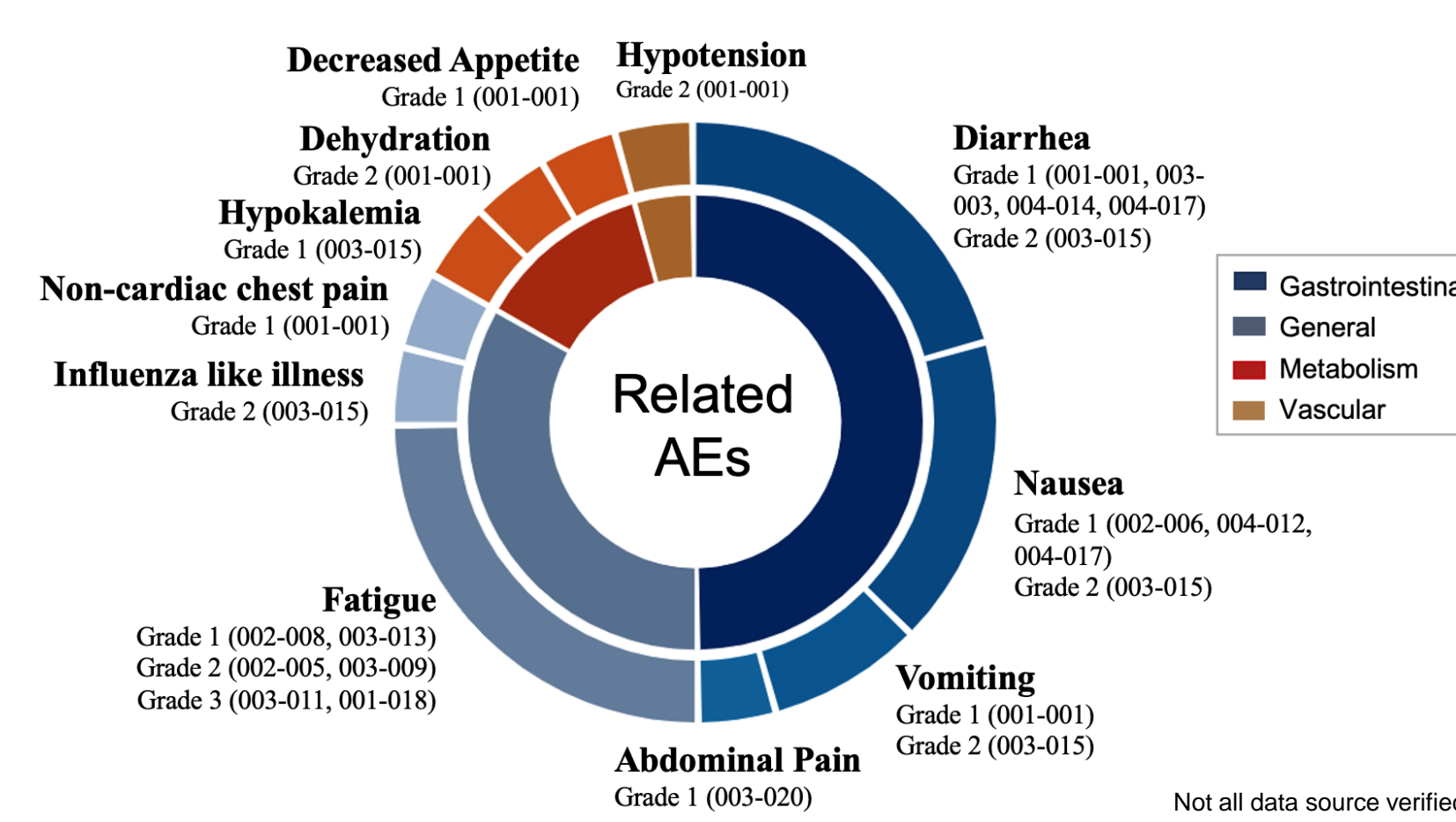
Figure 3. Summary of duration of treatment with once-daily ATRN-119

As of October 7, 2024



ADVERSE EVENTS

Figure 4. AEs at least possibly related to ATRN-119



All-cause TEAEs reported at ATRN-119 dose levels 50-800 mg once-daily

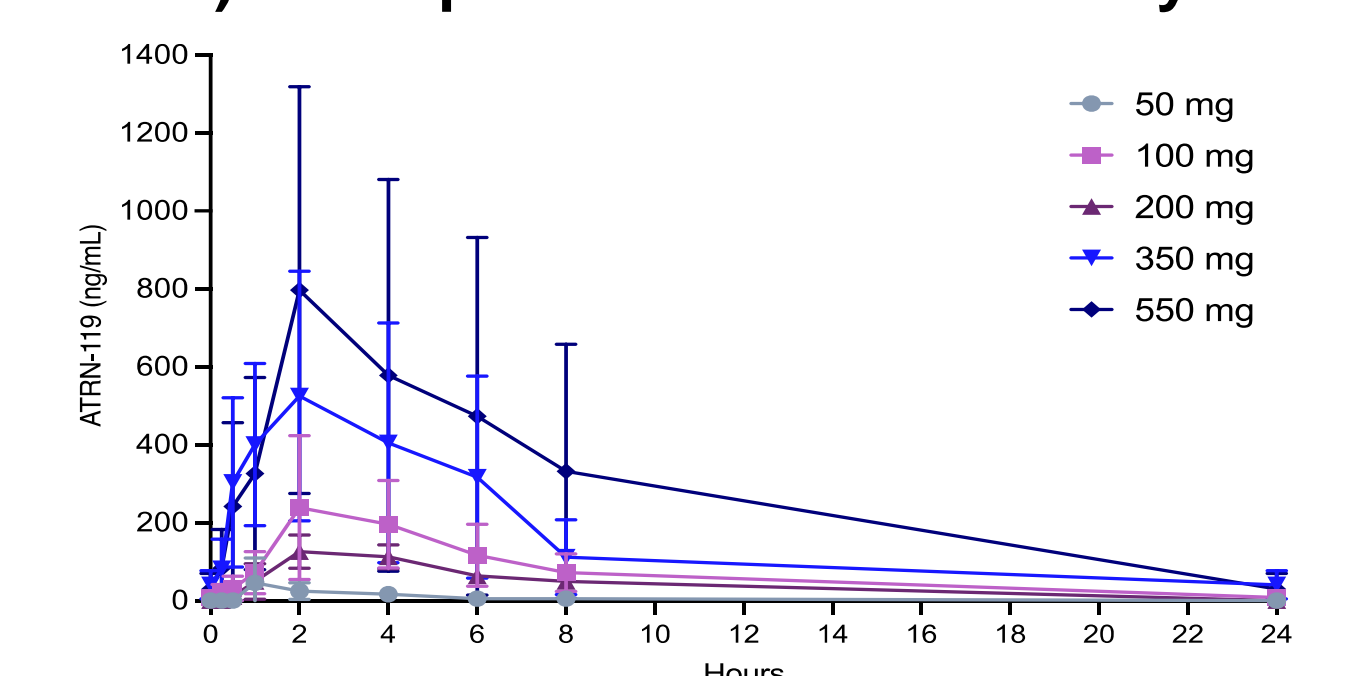
- As of Oct 2, 2024, 14 of 20 patients experienced AEs considered to possibly/probably related to ATRN-119
- No related SAE or grade 4-5 AEs have been observed
- There is no clear dose-relationship in AE
- There is no clear target organ toxicity
- No signs of hematological toxicity have been registered and no DLTs have been observed to date

PHARMACOKINETICS

Table 2. ATRN-119 Cycle 1 Day 7 (steady state) PK parameters

Dose Level mg, once daily	N	AUC _{0-24hr} (ng*h/mL)	C _{max} (ng/mL)	Half-life (hours)
		Mean (SD)	Mean (SD)	Mean (SD)
50	3	180 (143)	57 (56)	2.1 (1.4)
100	3	1771 (920)	277 (151)	3.8 (1.6)
200	3	1024 (162)	149 (9.2)	3.2 (0.5)
350	3	5252 (4362)	525 (320)	5.9 (0.5)
550	3	6899 (6058)	797 (522)	5.5 (1.4)

Figure 5. ATRN-119 Cycle 1 Day 7 (steady state) mean plasma concentration by dose



References

1. Blackford AN, Jackson SP, ATM, ATR, and DNA-PK: The Trinity at the Heart of the DNA Damage Response. *Mol Cell* 2017;66(6):801-817; 2. Krijnenburg TA, Wang L, Zimmermann MT, et al. Genomic and Molecular Landscape of DNA Damage Repair Deficiency across The Cancer Genome Atlas. *Cell Rep* 2018;23(1):239-254 e6; 3. Leona E, Fernandez-Capitillo O. Targeting ATR in cancer. *Nat Rev Cancer* 2018;18(9):586-595; 4. Vacca et al. *Cancer Res* 2020;80(7):Supply/Abstract nr 6177.

Abbreviations

AE, adverse event; ARID1A, AT-rich interactive domain 1A; ATR, Ataxia Telangiectasia and Rad3-related; AUC, area under the curve; BID, twice daily; CHEK, checkpoint kinase; DDR, DNA damage response; CCRK, Cyclin C; CCRK, Cyclin C; CCRK, cyclin-dependent kinase; CDKN2A, cyclin-dependent kinase inhibitor 2A; C_{max}, maximum plasma concentration; CNS, central nervous system; CYP, cytochrome P450; DNA, deoxyribonucleic acid; DNA damage and response; ECOG PS, Eastern Cooperative Oncology Group performance status; EMT, enhancer of zeste 2 polycomb repressive complex 2 subunit; Gly, glycine; KRAS, Kirsten rat sarcoma viral oncogene homolog; mCRPC, metastatic castration-resistant prostate cancer; MTD, maximum tolerated dose; MYC, myelocytomatosis oncogene; NGS, next-generation sequencing; PARP, paricicic acid; PARP, poly ADP-ribose polymerase; PCWG3, Prostate Cancer Clinical Trial Working Group 3; PK, pharmacokinetics; RAD51, RAD51 paring 1; Rb1, retinoblastoma 1; RECIST, Response Evaluation Criteria in Solid Tumors; RP2D, recommended phase 2 dose; SD, standard deviation; SOC, standard of care; TP53, tumor protein 53; WEE1, Wee1-like protein kinase.

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