

Preliminary Phase 1 Data From Ongoing First-in-Human Phase 1/2 TRESR Study of RP-3500

Virtual Investor Update
October 8, 2021



Today's agenda

Brief introduction

Lloyd M. Segal &
Maria Koehler, MD, PhD

President & CEO, Repare Therapeutics
EVP & CMO, Repare Therapeutics

Summary of AACR-NCI-EOTRC data and select case studies

Timothy Yap, MBBS, PhD, FRCP

Medical Director, Institute for Applied Cancer
Science, Associate Professor, Department
of Investigational Cancer Therapeutics, Division
of Cancer Medicine, MD Anderson Cancer Center

RP-3500 next steps

Maria Koehler, MD, PhD
Timothy Yap, MBBS, PhD, FRCP

EVP & CMO, Repare Therapeutics
MD Anderson Cancer Center

Concluding remarks

Lloyd M. Segal &
Maria Koehler, MD, PhD

President & CEO, Repare Therapeutics
EVP & CMO, Repare Therapeutics

Q&A

REPARE participants



Lloyd M. Segal
President & CEO



Maria Koehler, MD, PhD
Chief Medical Officer

Joining for Q&A



Mike Zinda, PhD
Chief Scientific Officer



Steve Forte
Chief Financial Officer

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Statements contained in this presentation regarding matters that are not historical facts are “forward-looking statements” within the meaning of the Private Securities Litigation Reform Act of 1995, as amended. Words such as "anticipates," "believes," "expects," "intends," “plans,” “potential,” "projects,” “would” and "future" or similar expressions are intended to identify forward-looking statements. Each of these forward-looking statements involves substantial risks and uncertainties that could cause actual results to differ significantly from those expressed or implied by such forward-looking statements. Forward-looking statements contained in this presentation include, but are not limited to, statements regarding the initiation, timing, progress and results of our current and future preclinical studies and clinical trials; the expected timing of program updates and data disclosures; the timing of filing INDs and other regulatory documents; the timing and likelihood of seeking regulatory approval for our product candidates; the competitive landscape for our product candidates; our ability to identify and develop additional product candidates using our SNIPRx platform; and our estimates regarding expenses, future revenue, capital requirements and needs for additional financing.

These forward-looking statements reflect our current beliefs and expectations. Many factors may cause differences between current expectations and actual results, including unexpected safety or efficacy data observed during preclinical or clinical studies, clinical site activation rates or clinical trial enrollment rates that are lower than expected, changes in expected or existing competition, changes in the regulatory environment, and unexpected litigation or other disputes. These and other risks are described more fully in our filings with the Securities and Exchange Commission (“SEC”), including the “Risk Factors” section of our Quarterly Report on Form 10-Q filed with the SEC on August 12, 2021, and other documents we subsequently filed with or furnished to the SEC. All forward-looking statements contained in this presentation speak only as of the date on which they were made. Except as required by law, we assume no obligation to update any forward-looking statements contained herein to reflect any change in expectations, even as new information becomes available.

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Leading clinical-stage precision oncology company focused on synthetic lethality



Lead clinical-stage candidate **RP-3500**, a **potential best-in-class ATR inhibitor**; currently in Phase 1/2 monotherapy and combination therapy



Robust pipeline of SL-based therapeutics; including **RP-6306**, our **PKMYT1 inhibitor** currently in Phase 1, and our **Polθ inhibitor**



Proprietary **genome-wide CRISPR-enabled SNIPRx platform**, focused on genomic instability and DNA damage repair



Powerful SL-based approach and proprietary platform provides **differentiated patient selection insights**



Cash, restricted cash and marketable securities of \$301 million as of June 30, 2021











Expert participant: Timothy Yap, MBBS, PhD, FRCP



Medical Oncologist and Physician-Scientist at the University of Texas, MD Anderson Cancer Center

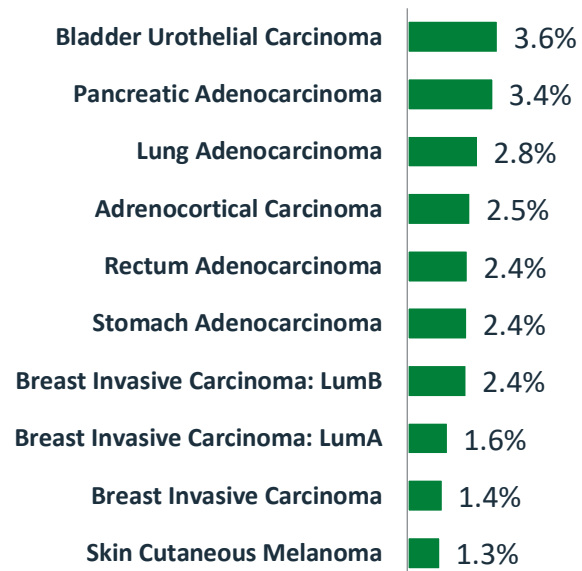
- Associate Professor, Department for Investigational Cancer Therapeutics
- Medical Director of the Institute for Applied Cancer Science
- Associate Director of Translational Research in the Institute for Personalized Cancer Therapy
- Primary research focuses on the first-in-human and combinatorial development of molecularly targeted agents and immunotherapies, and their acceleration through clinical studies using novel predictive and pharmacodynamic biomarkers
- Main interests include the targeting of the DNA damage response with novel therapeutics, such as ATR and PARP inhibitors, as well as the development of novel immuno-therapeutics
- BSc degree in Immunology and Infectious Diseases and MD from Imperial College London, UK

Robust pipeline of SL-based precision oncology therapeutics

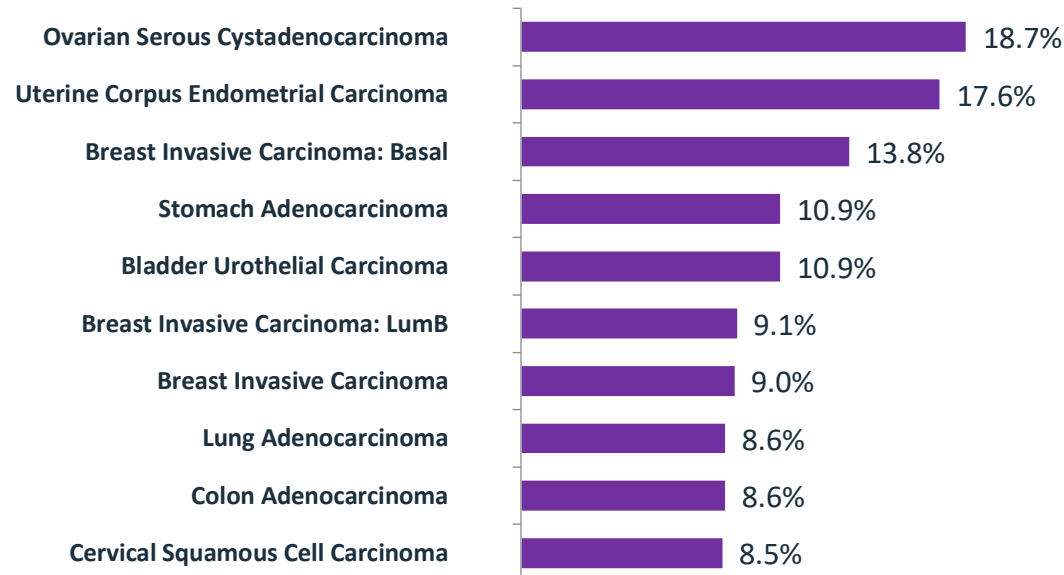
		SL Pair		Discovery	IND-Enabling	Phase 1/2	Registration-directed	Anticipated milestones	Rights
Clinical	ATR inhibitor RP-3500	ATM + 16 STEP ² lesions	ATR					<ul style="list-style-type: none"> Q2 22 TRESR final MonoRx data Q3 22 PARP combo early data 	
	PKMYT1 inhibitor RP-6306	CCNE1, FBXW7 + others	PKMYT1					H2 22 early Phase 1 readout	
Preclinical	Polθ inhibitor	BRCA1/2 + others	Polθ					IND-enabling studies in H1 22	 
Discovery	SNIPRx [®] platform	8 additional SL targets							
		Discovery and validation of new SL precision oncology targets							 

Expanding RP-3500 patient opportunity with STEP² selection tools*

Top 10 tumor types with highest prevalence of ATM deficiency



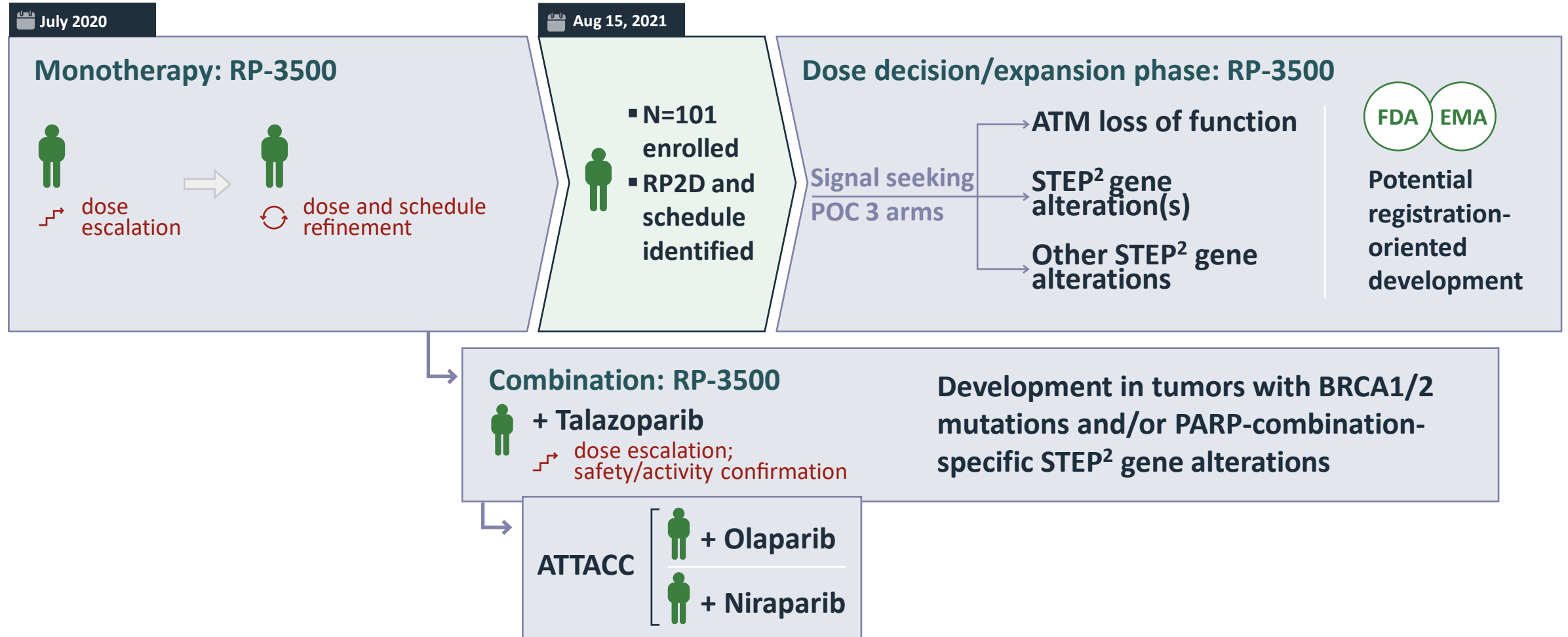
Top 10 tumor types with highest prevalence of ATM deficiency or STEP² genomic alterations



- Beyond ATM, 16 of 19 additional, mutually exclusive genomic alterations identified as SL with RP-3500 are eligible for recruitment into the ongoing trial
- Represents expanded, clinically relevant populations with unmet medical needs
- Average prevalence of ~2% (ATM) to ~10% (STEP² genes) across multiple tumors

RP-3500 clinical progress to-date

Global multicenter study designed for patients with recurrent tumors with ATM loss or loss of any of the additional 16 STEP² genes



Summary of AACR-NCI-EORTC data and select case studies



AACR-NCI-EORTC Virtual International Conference on

MOLECULAR TARGETS AND CANCER THERAPEUTICS

October 7-10, 2021



First-in-Human biomarker-driven Phase I TRESR trial of ATR inhibitor RP-3500 in patients with advanced solid tumors harboring synthetic lethal genomic alterations

Timothy A. Yap¹, Elizabeth Lee², David Spigel³, Elisa Fontana⁴, Martin Hojgaard⁵, Stephanie Lheureux⁶, Niharika Mettu⁷, Louise Carter⁸, Ruth Plummer⁹, Victoria Rimkunas¹⁰, Ian M. Silverman¹⁰, Adrian J. Fretland¹⁰, Danielle Ulanet¹⁰, Peter Manley¹⁰, Ezra Rosen¹¹

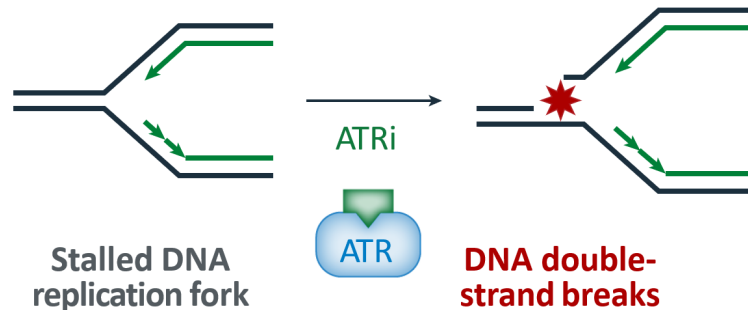
¹University of Texas MD Anderson Cancer Center, Houston, TX; ²Dana-Farber Cancer Institute, Boston, MA; ³Sarah Cannon Research Institute, Nashville, TN; ⁴Sarah Cannon Research Institute, London, UK; ⁵Copenhagen University Hospital, Herlev, Denmark; ⁶Princess Margaret Cancer Centre, Toronto, Canada; ⁷Duke University Medical Center, Durham, NC; ⁸The Christie NHS Foundation Trust, Manchester, UK; ⁹Newcastle upon Tyne Hospitals NHS Foundation Trust, Newcastle upon Tyne, UK; ¹⁰Repare Therapeutics, Cambridge, MA; ¹¹Memorial Sloan Kettering Cancer Center, New York, NY

Speaker Disclosures: Timothy Yap, MD

I have the following financial relationships to disclose:

- **Employee of:** University of Texas MD Anderson Cancer Center, where I am Medical Director of the Institute for Applied Cancer Science, which has a commercial interest in DDR and other inhibitors.
- **Grant/Research support (to Institution) from:** Repare, AstraZeneca, Artios, Bayer, Beigene, BioNTech, BMS, Clovis, Constellation, Cyteir, Eli Lilly, EMD Serono, Forbuis, F-Star, Artios, GlaxoSmithKline, Genentech, Haihe, ImmuneSensor, Ionis, Ipsen, Jounce, Karyopharm, KSQ, Kyowa, Merck, Novartis, Pfizer, Ribon Therapeutics, Regeneron, Rubius, Sanofi, Scholar Rock, Seattle Genetics, Tesaro and Vivace.
- **Consultant for:** Repare, AstraZeneca, Almac, Aduro, Artios, Athena, Atrin, Axiom, Bayer, Bristol Myers Squibb, Calithera, Clovis, Cybrexa, EMD Serono, F-Star, GLG, Guidepoint, Ignyta, I-Mab, ImmuneSensor, Jansen, Merck, Pfizer, Roche, Schrodinger, Seattle Genetics, Varian, Zai Labs and ZielBio
- **Stockholder in:** Seagen
- **I will discuss the following off label use and/or investigational use in my presentation:** RP-3500

RP-3500: a potential best-in-class, highly selective inhibitor of ATR kinase

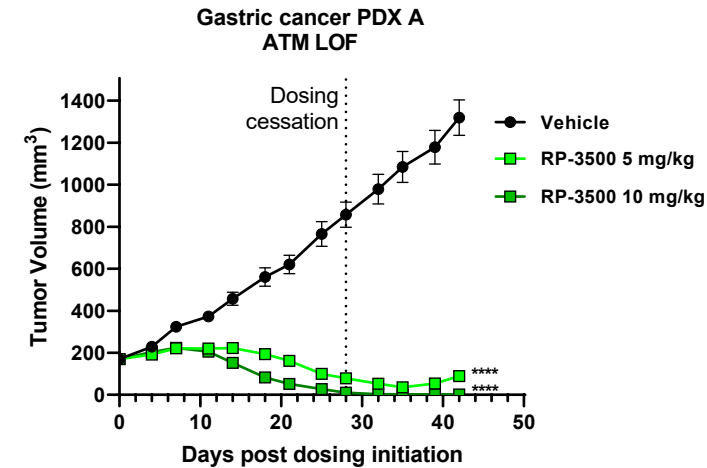


ATR inhibition is synthetically lethal with genomic alterations affecting DNA damage response

- ATR is a key mediator of cellular DNA damage response (DDR) and is activated in response to DNA replication stress
- A genome-wide CRISPR-based screening platform identified multiple synthetic lethal genomic alterations that predict for sensitivity to RP-3500 (STEP²* genes)
 - ATM, ATRIP, BRCA1/2, CHEK2, CDK12, CHTF8, FZR1, MRE11, NBN, PALB2, RAD17, RAD50, RAD51B/C/D, REV3L, RNASEH2A/B, SETD2

*STEP² = SNIPRX** targeted expansion of patient populations

**SNIPRX = SyNthetic lethal Interactions for Precision Rx



RP-3500 is a potent and highly selective inhibitor of ATR

- Low nanomolar potency in biochemical (1.0 nM) and cell-based assays (0.33 nM)
- >2,000-fold selectivity over ATM, DNA-PK and PI3Ka
- Single agent activity in tumor models of different histologies and DDR defects

Roulston et al., ENA 2021, #P054

First-in-human Phase 1/2 TRESR study design

Phase 1/2 TRESR (Treatment Enabled by SNIPRx) study

NCT04497116
(accruing)

Inclusion Criteria

- Patients ≥18yo with solid tumors resistant, refractory, and/or intolerant to standard therapy
- Tumors with *centrally reviewed** deleterious STEP² alterations
- ECOG PS 0 or 1
- Hgb ≥9.5g/dL, Platelets ≥140K/uL, ANC ≥1.7K/uL

Module 1: single agent RP-3500

Primary endpoints:

- Safety and tolerability
- Recommended Phase 2 dose (RP2D), schedule

Other endpoints

- Pharmacokinetics
- Pharmacodynamics in paired tumor biopsies
- Preliminary antitumor activity
- Kinetics of circulating tumor DNA (ctDNA)

Presentation of **early data** from this ongoing study:

- TRESR initiated in July 2020
- Data cut-off date: August 15, 2021
- 101 patients included in this early dataset

Phase 1/2 TRESR: patient characteristics

All patients		N = 101
	Male	42
	Female	59
Median age, years (range)		63 (33-77)
	≥65 yrs	46
ECOG status		
	0	48
	1	53
Lines of prior therapy		
	1-3	51
	4 or more	45
	Pending	5
Prior Platinum		62
Prior PARP inhibitor		28
Prior PD-1/L1 inhibitor		20

Tumor types

Ovarian	19
Prostate	18
Breast	13
Pancreas	8
Sarcoma	8
Other*	35

*other tumor types:

- CRC
- Bile Duct
- GI
- Endometrial
- Lung
- Ampullary
- Appendix
- HNSCC
- Melanoma
- Mesothelioma
- Skin

Most common genotypes

ATM	37
BRCA1	31
BRCA2	13
CDK12	7
Other STEP2**	23

**STEP² genotypes:

- CHEK2
- NBN
- PALB2
- RAD51C/B
- RNASEH2
- SETD2
- FZR1

RP-3500 MTD/RP2D established at 160mg QD, 3d on/4d off schedule

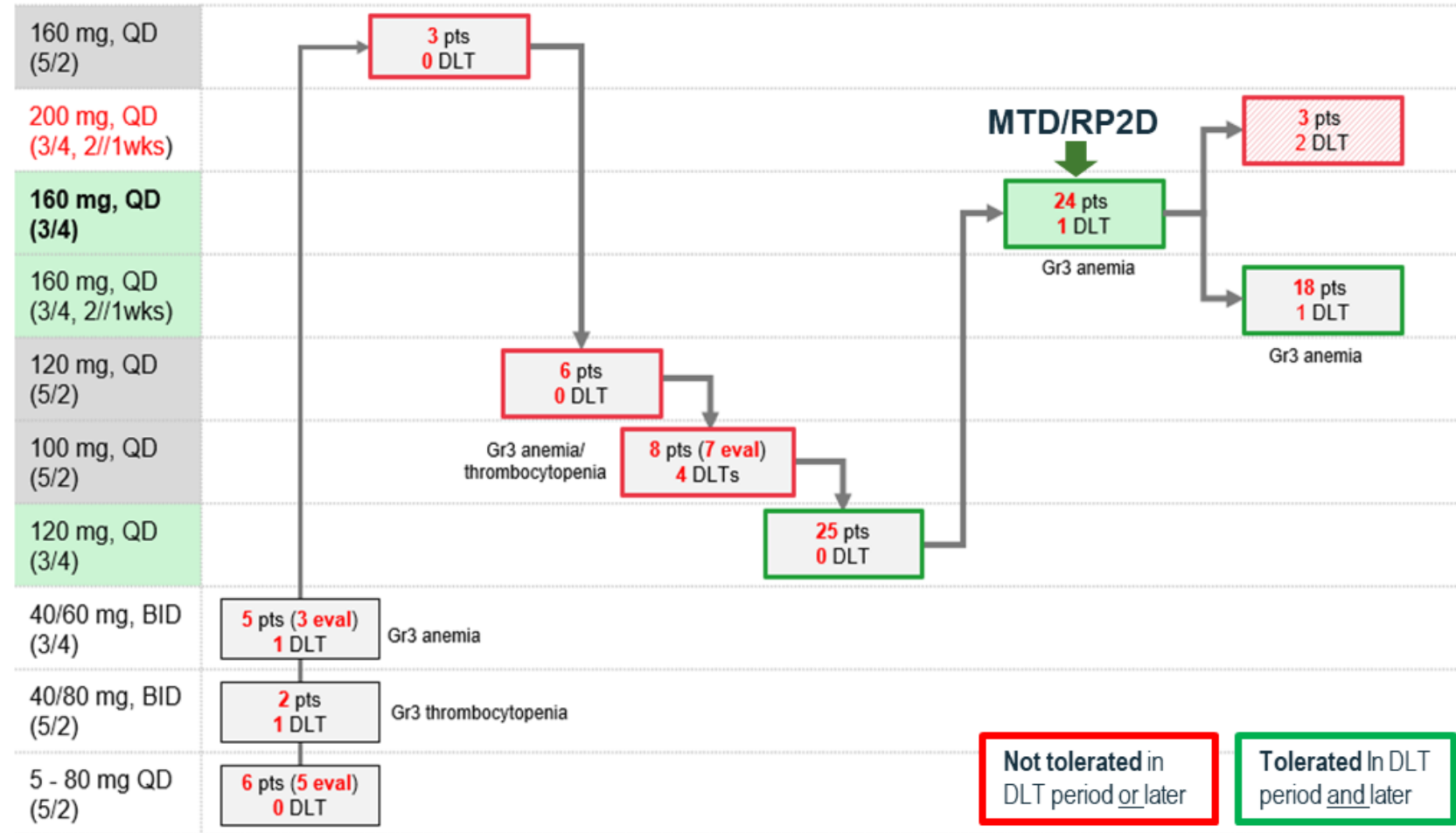
Comprehensive assessment for RP-3500 MTD/RP2D

Single agent RP-3500 tested at multiple doses and schedules

- Adaptive BOIN design and sufficient cohort sizes to ensure confidence in MTD/RP2D decision
- Once daily (QD) and twice daily (BID)
- 5d on/2d off and 3d on/4d off; continuously and 2w on/1w off

DLTs: anemia, thrombocytopenia

MTD/RP2D of RP-3500:
160mg QD, 3d on/4d off



RP-3500 treatment emergent adverse events

All grades, occurring in ≥10% of patients

Preferred term	5/2 Schedule (N=25)			3/4 Schedule (N=76)			All Patients (N=101)		
	All Grade n (%)	Grade 3 n (%)	Grade 4 n (%)	All Grade n (%)	Grade 3 n (%)	Grade 4 n (%)	All Grade n (%)	Grade 3 n (%)	Grade 4 n (%)
Any TEAE	25 (100)	15 (60)	2 (8)	58 (76.3)	19 (25.0)	1 (1.3)	83 (82.2)	34 (33.7)	3 (3.0)*
Anemia	19 (76)	11 (44)	0	40 (52.2)	11 (14.5)	0	59 (58.4)	22 (21.8)	0
Fatigue	9 (36)	1 (4)	0	19 (25.0)	2 (2.6)	0	28 (27.7)	3 (3.0)	0
Decreased appetite	6 (24)	0	0	17 (22.4)	0	0	23 (22.8)	0	0
Nausea	6 (24)	0	0	16 (21.1)	1 (1.3)	0	22 (21.8)	1 (1.0)	0
Neutrophil count decreased	5 (20)	2 (8)	0	14 (18.4)	4 (5.3)	0	19 (18.8)	6 (5.9)	0
Platelet count decreased	7 (28)	2 (8)	1 (4)	12 (15.8)	3 (3.9)	1 (1.3)	19 (18.8)	5 (5.0)	2 (2.0)*
Diarrhea	3 (12)	0	0	14 (18.4)	0	0	17 (16.8)	0	0
Abdominal pain	3 (12)	0	0	8 (10.5)	1 (1.3)	0	11 (10.9)	1 (1.0)	0

MTD/RP2D established at 160mg QD, 3d on/4d off

3 days on/4 days off schedule preferred

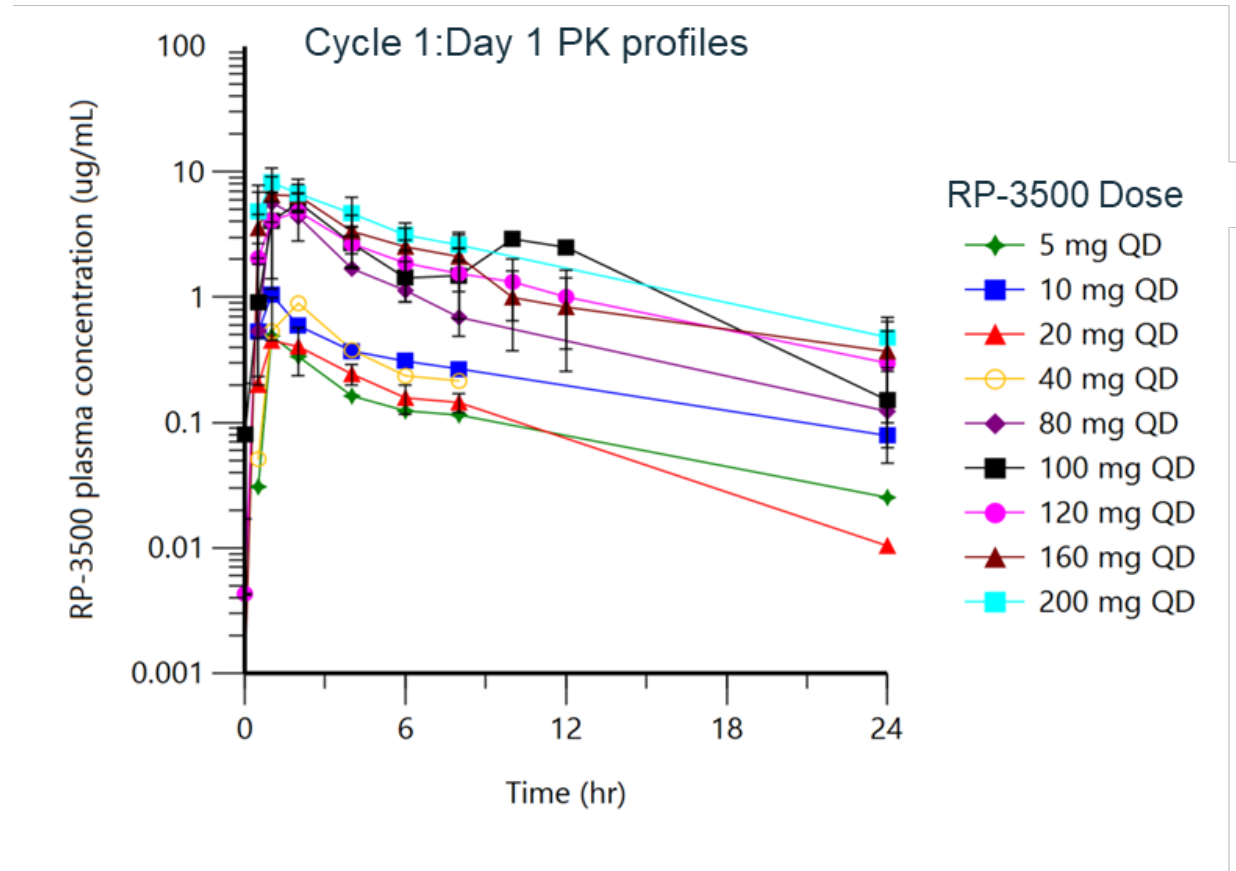
Manageable impact of on-target anemia

- Anemia is the most common cause of dose interruptions, modifications
- At the preferred 3/4 schedule, dose interruptions, reductions and transfusions were infrequent
- No discontinuations related to RP-3500 emergent adverse events

	5/2 Schedule ≥100 mg/day* (N=18)	3/4 Schedule ≥100 mg/day* (N=75)
# Cycles, mean (SD) [Range]	3.9 (2.62) [0, 10]	1.9 (1.72) [0, 8]
Subjects Exposed to RP-3500 n (%)		
≥1 cycle	17 (94.4%)	57 (76.0%)
≥2 cycles	14 (77.8%)	37 (48.7%)
Interruptions n (%)		
1	8 (44.0%)	16 (21.3%)
2	2 (11.1%)	3 (4.0%)
≥3	5 (27.8%)	2 (2.7%)
Dose Reductions, n (%)		
1	6 (33.3%)	10 (13.3%)
2	3 (16.7%)	1 (1.3%)
Transfusions, n (%)		
Cycle 1	4 (22.2%)	6 (8.0%)
Cycles 1-2	9 (50.0%)	9 (12.0%)
Cycles 1-3	12 (66.7%)	10 (13.2%)

Pharmacokinetic profile RP-3500 exposures meet efficacy targets at $\geq 100\text{mg}$

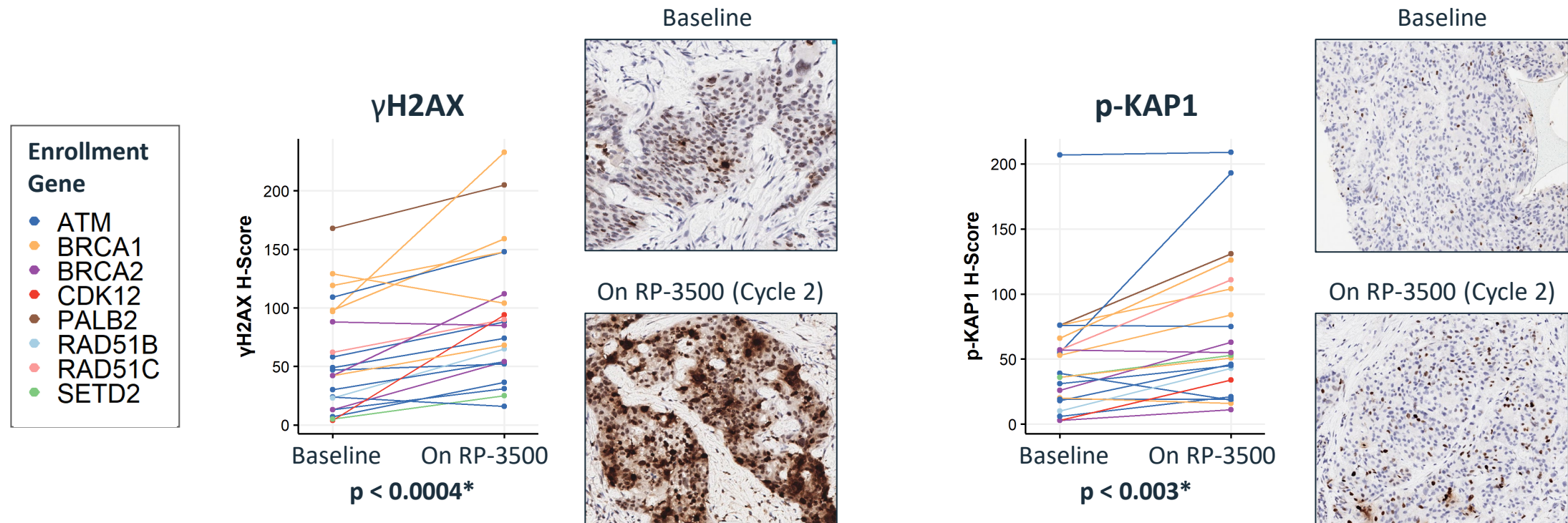
- Half-life is ~ 6 hrs
- Linear Cmax and AUC, consistent across all doses
- Doses ≥ 100 mg QD achieve predicted efficacious RP-3500 exposures
 - Once daily (QD) regimen selected over twice daily (BID) based on minimal differences in target coverage between these 2 regimens
 - Based on the results of the RP-3500 food effect study, RP-3500 can be given with and without food



RP-3500 Pharmacodynamics

Robust pathway modulation in paired tumor biopsies

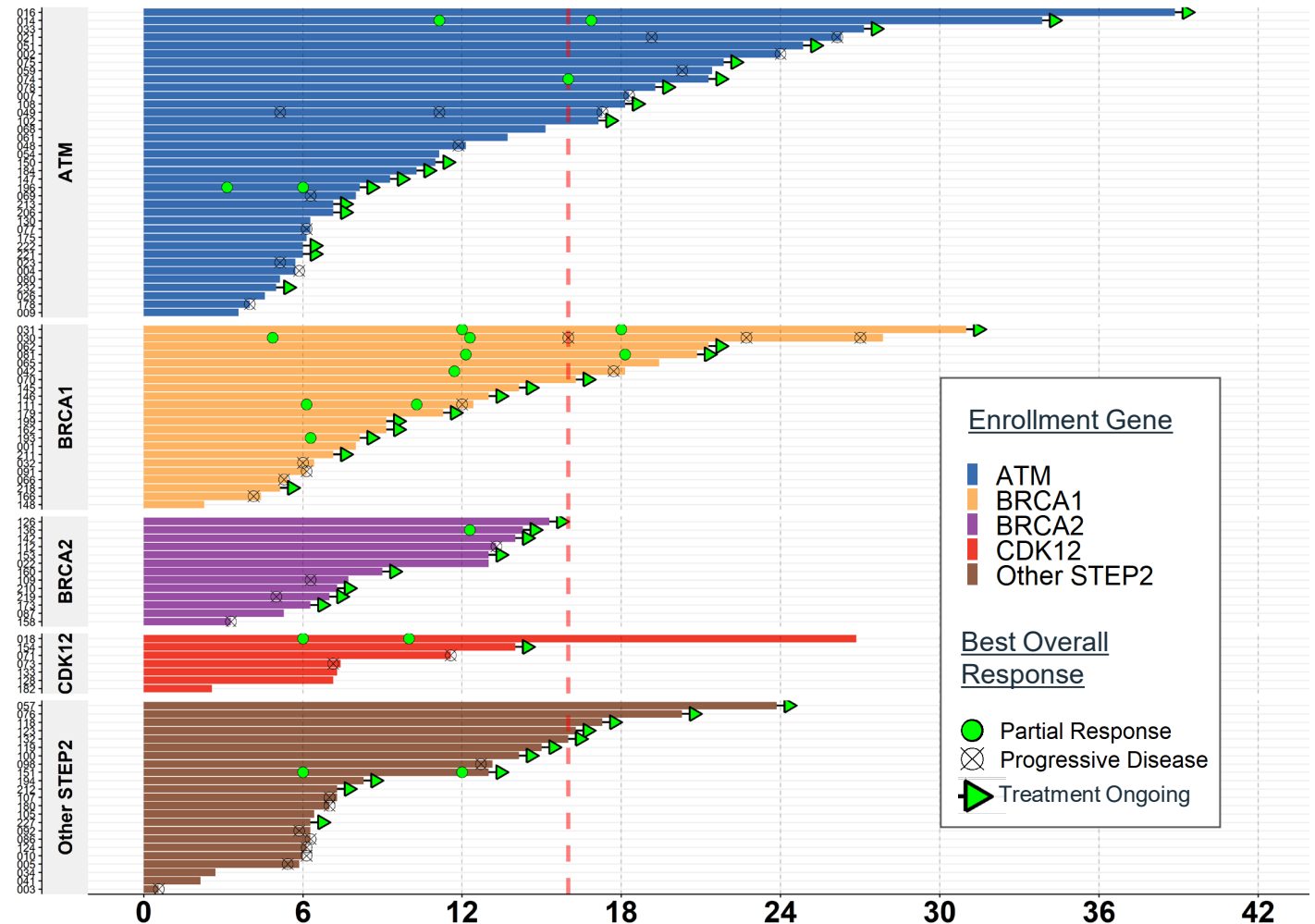
Consistent and statistically significant on-treatment increases
in DNA damage-induced γ H2AX and p-KAP1 across tumor genotypes (N=21)



Early analysis of treatment duration

Therapy ongoing in 54* (54%) of 101 patients

- Early analysis of therapy duration shows clinical activity across tumor types and STEP2 alterations
- Responses** included:
 - CRPC (ATM, CDK12)
 - Ovarian, post-PARPi (4 BRCA1, 1 RAD51C)
 - ER+ breast (BRCA1)
 - HNSCC (BRCA1)
 - Melanoma (BRCA2)
- Copy number analysis of enrolled gene alterations is ongoing



Early efficacy observed with RP-3500 ≥ 100 mg/day

Broad spectrum of efficacy observed

- **Meaningful clinical benefit in 34 (49%) of 69 evaluable patients**
- **Across STEP2 gene alterations**
- **Across schedules & after PARPi failure**
 - **12 responses:** 8 RECISTv1.1 cPR/uPR, 2 pts by PCWG3 and 2 pts by GCIG
(ATM, CDK12, BRCA1, BRCA2, RAD51B, RAD51C and FZR1)
 - **14 patients** ongoing SD ≥ 16 weeks
 - **8 patients** <16 w on study: early significant decreases in tumor markers and tumor shrinkage ($<30\%$)
- **Late responses also observed: initial RECISTv1.1 partial response (PR) seen at week 16**

	5/2 Schedule ≥ 100 mg/day (N=18)	3/4 Schedule ≥ 100 mg/day (N=76)	All patients ≥ 100 mg/day (N=94)
Evaluable pts (≥ 1 post baseline scan)	17	52	69
Best response	4	8	12
RECISTv1.1	2 cPR*; 1uPR **	2 cPR; 3 uPR [#]	4 cPR; 4uPR
PCWG3 PSA	1	1	2
GCIG CA125	-	2	2
SD (≥ 16 w)	6	8	14
SD (≥ 6 w) ^{&}	6	23	29
PD	6	21	27
Data pending	1	0	1
Discontinued w/o scan	1	3	4
On treatment w/o scan	0	21	21

*1 patient with cPR required radiotherapy to brain lesions early in trial. No brain scan at study entry.

**Patient started on 5/2 schedule and changed to 3/4, later achieved uPR response on 3/4

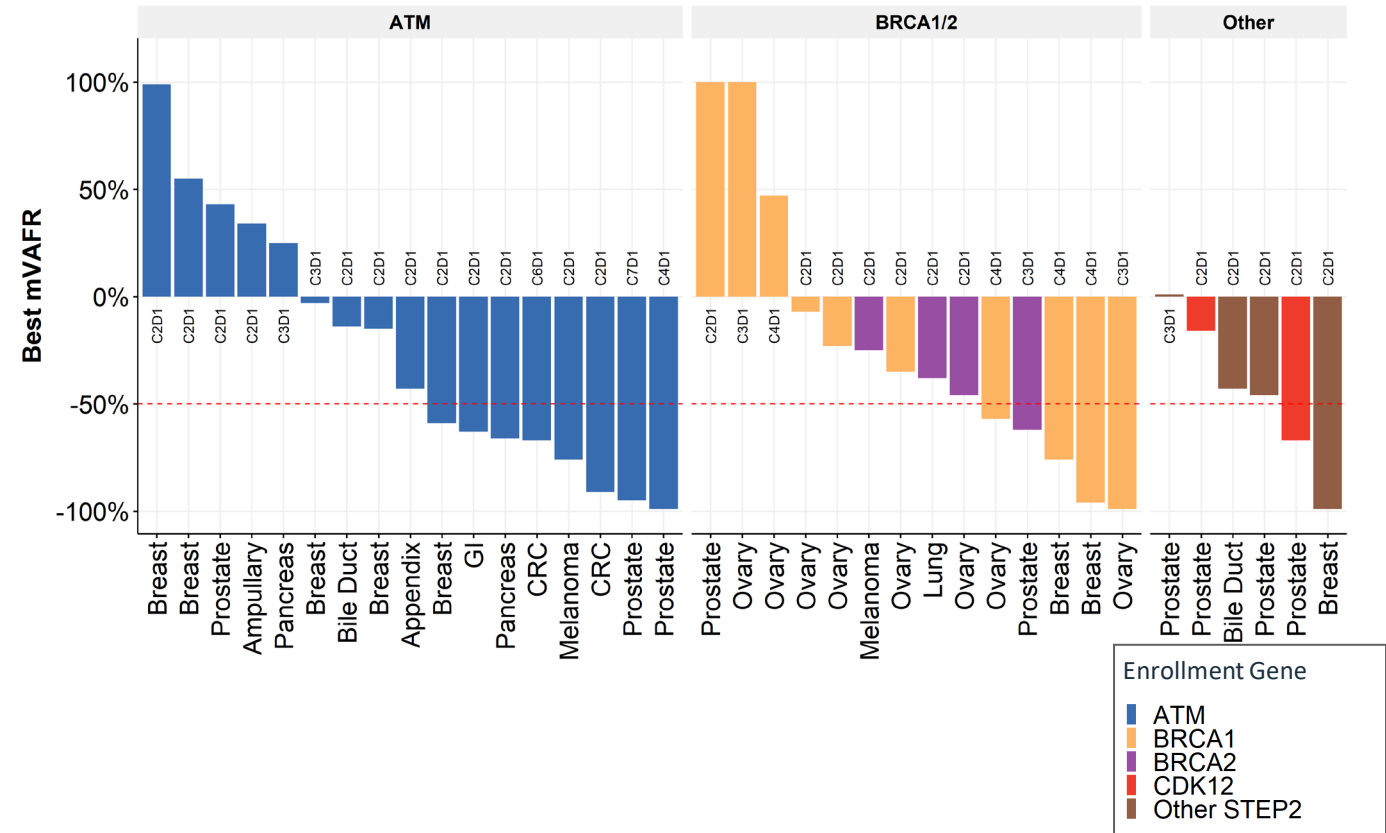
[#]1 patient began on PARPi+RP3500 for 2 weeks, before transitioning to RP-3500 monotherapy.

[&]includes the SD >16 w patients

Deep molecular responses in TRESR

Circulating tumor DNA (ctDNA) measured serially in 37 pts

- ctDNA, fragmented tumor DNA in circulating blood, may reflect the entire tumor genome as “liquid biopsies” at various time points to monitor tumor during treatment
- Published data suggest that early molecular responses in ctDNA may be correlated with patient benefit during treatment with anticancer drugs
- ctDNA best mean variant allele frequency (mVAF*) measures change in tumor burden
- RP-3500 data show early and significant decrease (>50%) in mVAFR** in tumors with multiple genotypes

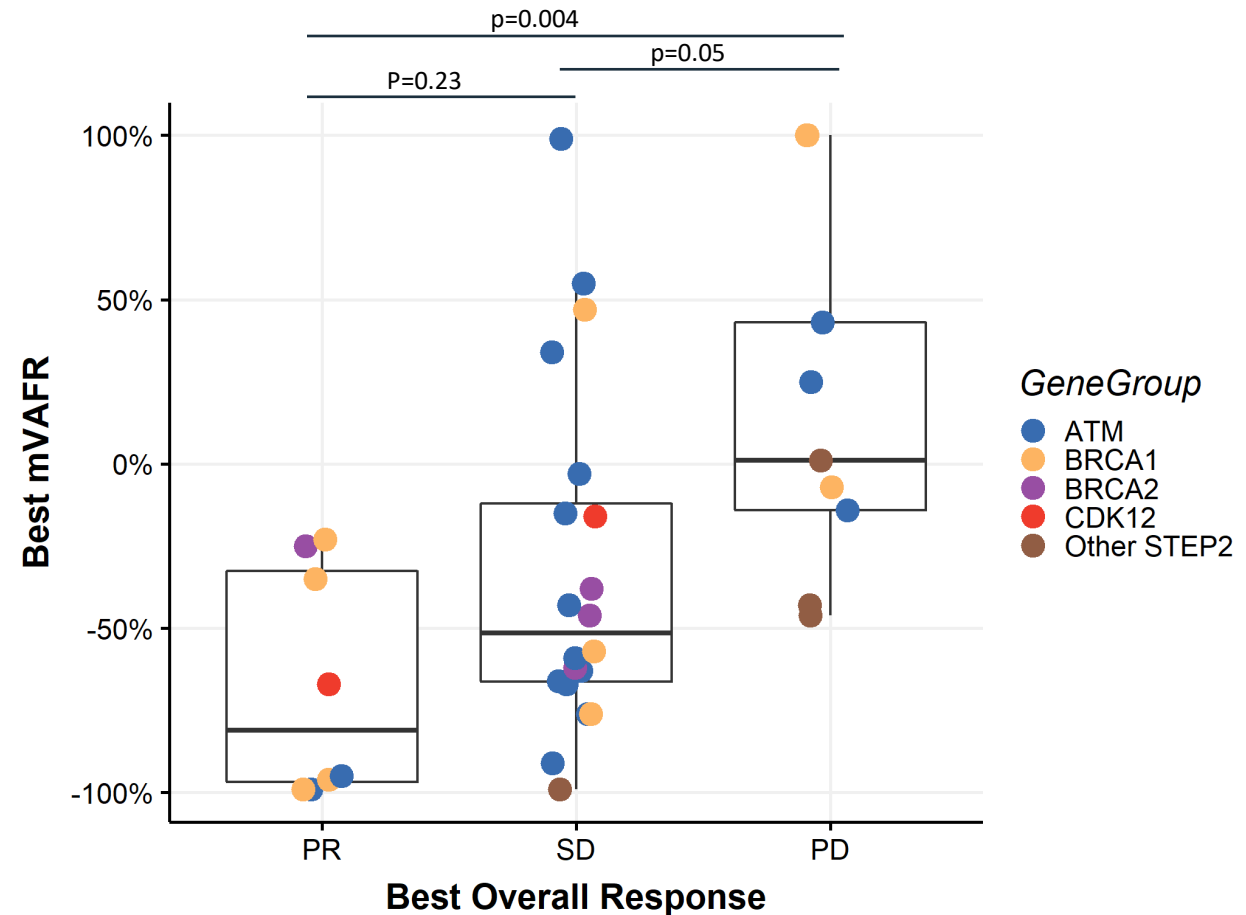


*mVAF calculated as % decrease from baseline

**mVAFR is the mean variant allele frequency ratio (relative to baseline). mVAFR capped at +100%

Preliminary analysis suggests ctDNA response may predict clinical benefit

- RP-3500 patients with PRs showed early and frequently significant (>50%) reductions in mVAFR in ctDNA
- Analyses are ongoing to correlate ctDNA responses with clinical efficacy in order to evaluate the predictive power of serial ctDNA measurements

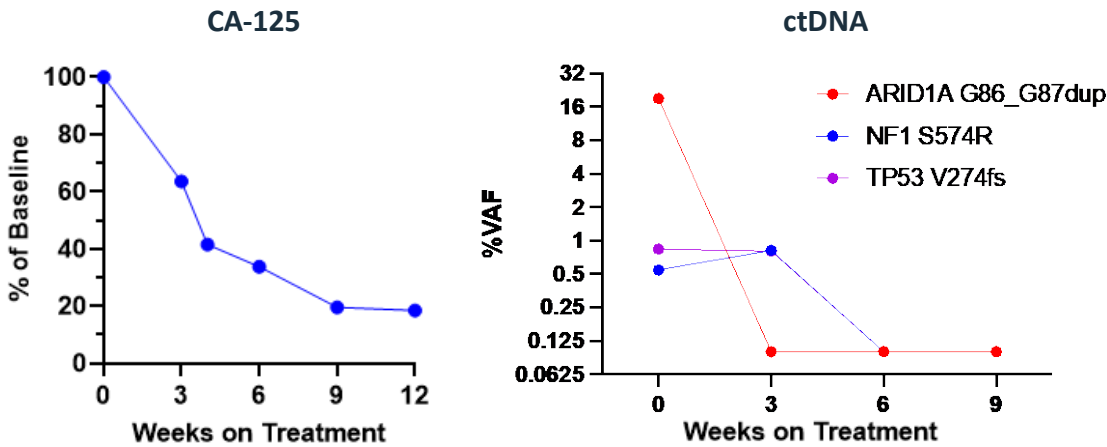


- ctDNA cohort = 37/101 enrolled patients
- mVAFR = mean variant allele frequency ratio as an indirect measure of tumor burden

Ovarian cancer with gBRCA1 mutation

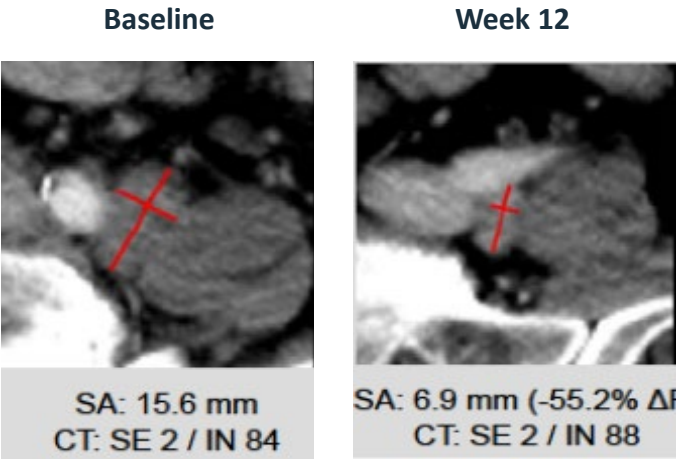
Ongoing RECIST cPR*

Ongoing GCIIG CA125 response

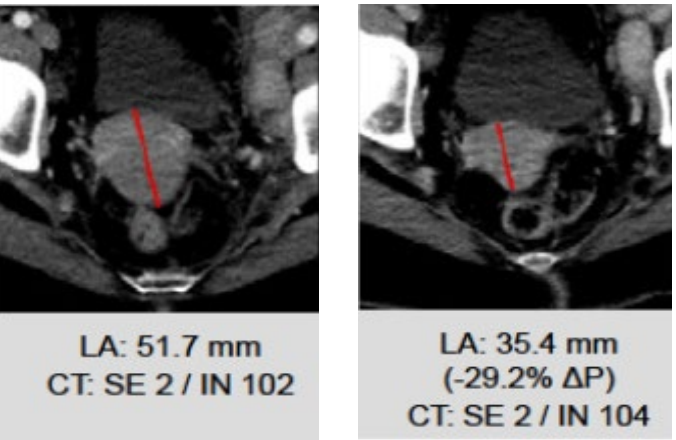


59 y/o female with 4 prior therapeutic regimens for metastatic ovarian cancer: Prior platinum, previous failure of PARP inhibitor (best response PD) and docetaxel + avastin (best response PD)

T01 Lymph Node
common iliac left
Baseline: 15.6 mm
Restage: 6.9 mm



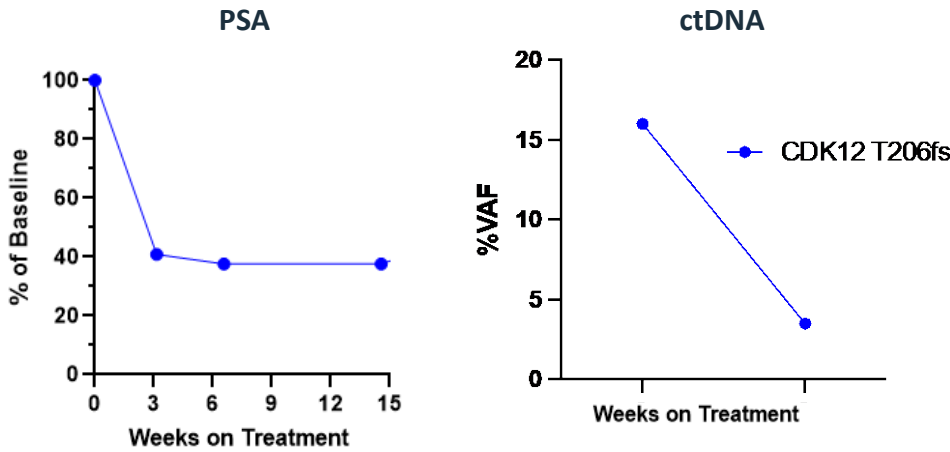
T03 Vaginal Cuff
Baseline: 51.7 mm
Restage: 35.4 mm



CRPC with CDK12 mutation

Ongoing RECIST cPR

Ongoing PCWG3 PSA response

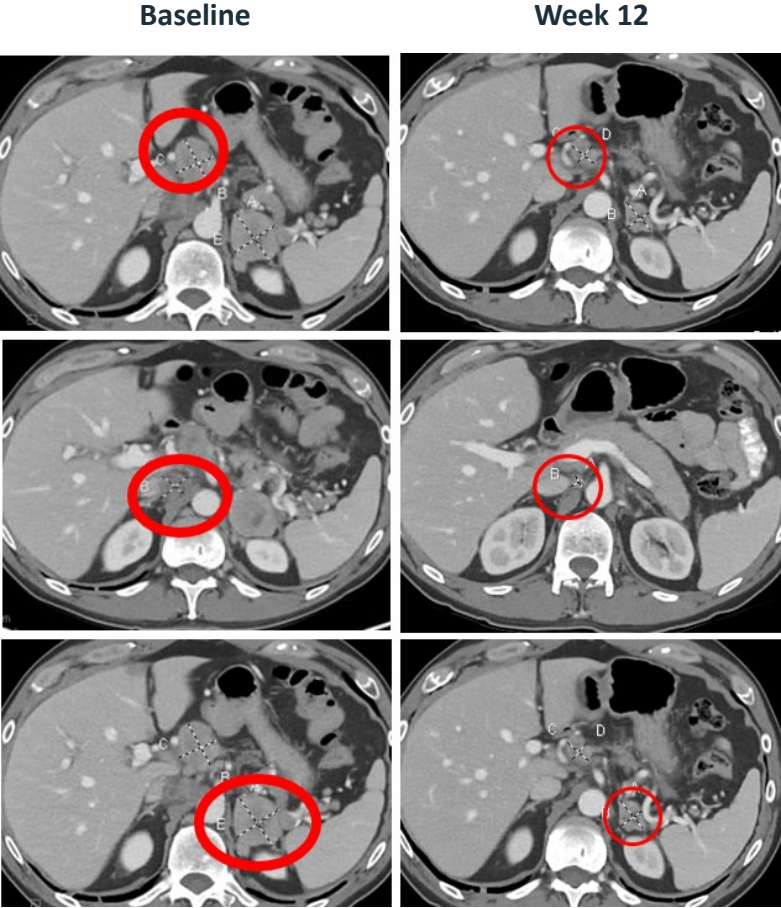


57 y/o male with CRPC received 6 prior regimens, incl. best response of PD on docetaxel, platinum/etoposide and lpi/Nivo

Common hepatic lymph node
Baseline: 32mm
Restage: 17mm

Retrocaval lymph node
Baseline: 22mm
Restage: 9mm

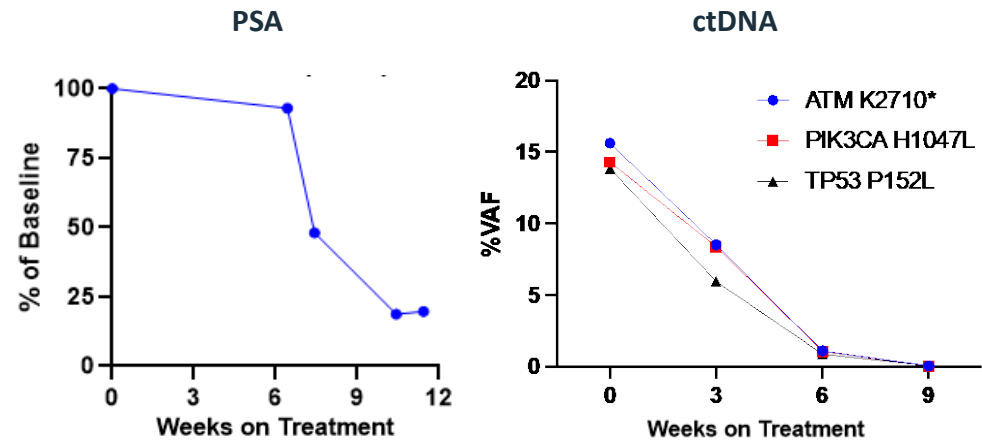
Left adrenal gland mass
Baseline: 44mm
Restage: 27mm



CRPC with ATM mutation

Ongoing PCWG3 PSA response

Ongoing RECISTv1.1 uPR, confirmation of response awaited



68 y/o male with CRPC, multiple bone metastases and lymph node disease, with disease progression on 2 prior regimens

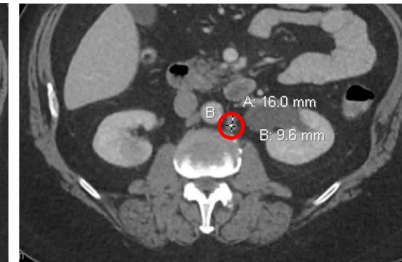
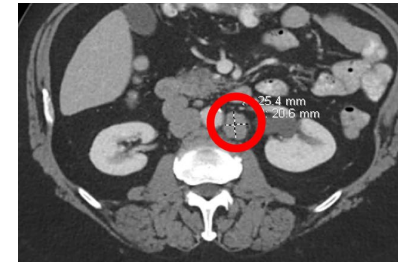
Left para-aortic LN

Baseline: 21mm

Restage: 10mm

Baseline

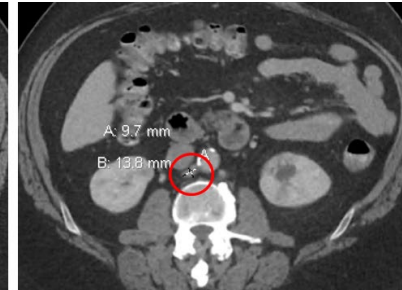
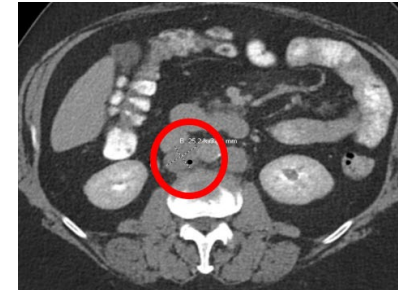
Week 16



Retrocaval LN

Baseline: 25mm

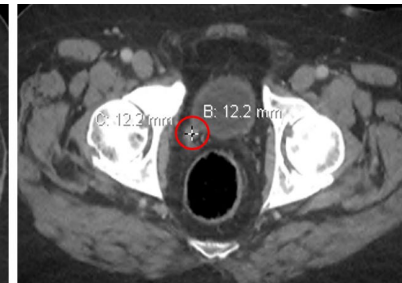
Restage: 9mm



Right pelvic nodule along bladder wall

Baseline: 17mm

Restage: 13mm



Summary – Early Phase 1/2 TRESR trial data

RP-3500 is safe and well tolerated, with mainly G1-2 anemia (only 21.8% G3 overall and 14.5% in chosen schedule)

- Likely off-target toxicities, e.g., fatigue or GI events, are infrequent (only up to 3% G3)

RP-3500 RP2D for further monotherapy evaluation is 160mg QD 3d on/4d off

Biomarker data confirm proof-of-mechanism across different tumors with multiple molecular backgrounds

Preliminary antitumor activity observed in patients with cancers harboring selected genomic alterations at doses $\geq 100\text{mg}$

- **Meaningful clinical benefit in 34 (49%) of 69 patients**
 - **12 patients** with objective tumor responses:
 - 8 RECIST V1.1: cPR/uPR (7 confirmed, 1 unconfirmed)
 - 2 PCWG3 PSA responses
 - 2 GCIG CA125 responses
 - **14 patients** with ongoing RECIST v1.1 SD for at least 16 weeks
 - **8 patients** <16 weeks on study: SD with early significant decreases in tumor markers and <30% tumor shrinkage

Conclusions

RP-3500 is in development as a potent, potentially best-in-class, highly selective ATR inhibitor

The TRESR Phase 1/2 study is the largest biomarker-selected trial testing a single agent ATR inhibitor (N=101) and continues to enroll

Early TRESR data provide clinical POC and validate Repare Therapeutics' SNIPRx platform for molecular selection of tumors for ATR inhibitor therapy

Favorable and differentiated safety profile, along with promising and distinct early efficacy, offer a clear direction for further development of RP-3500

Next steps include:

- TRESR Phase 2 expansion cohorts to open imminently
- Combination studies are ongoing or will open shortly

Acknowledgements

The patients and their families who make this trial possible

Participating TRESR Sites for their work and contribution

▪ Billy Hoadley; Christian Brown; Sandra Montez	University of Texas MD Anderson Cancer Center
▪ Emily Gibson; Mary Liebers	Dana Farber Cancer Institute
▪ Danielle McCreary	Sarah Cannon Research Institute/Tennessee Oncology
▪ Alissa Casas, Arousiak Kazarian	Memorial Sloan Kettering Cancer Center
▪ Paula Lee	Duke Cancer Institute
▪ Jocelyn Hubbard	Massachusetts General Hospital
▪ Rachael DiGiulio; Brenton Travers	Rhode Island Hospital/Lifespan
▪ Rachel Wildman	Princess Margaret Cancer Center
▪ Terry Wood; Sinead O'Reilly	The Christie NHS Foundation Trust
▪ Helen Blair	Newcastle Hospital NHS Foundation Trust
▪ Victoria Hall; Victoria Borja Bernal	Sarah Cannon Research Institute UK
▪ Cecilia Sonander Westphal; Emma-Sofie Sønderskov Darre	Righospitalet, Denmark

Precision Oncology Decision Support (PODS) Group at the University of Texas MD Anderson Cancer Center

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Treximo (part of the ProPharma Group)

This Study is funded by Repare Therapeutics

RP-3500 next steps



Update on TRESR (as of Oct 6th)

Update on TRESR Module 1

Enrollment to the main component of TRESR Module 1 (RP2D determination) is closed



Recruitment to Module 1 is now closed (goal 120 pts)



Unchanged safety/tolerability profile as of Oct 6th



Update of 4 patients with uPR reported as of AACR-NCI-EORTC 2021 meeting: 3 cPR, 1 uPR



Final results of Module 1 expected to be reported in 2Q 2022 with expected 3-4 months follow up of all M1 patients

Additional Modules in TRESR anticipated to open imminently

Additional details on clinical responders

Tumor type	Genotype	Enrolled Dose Level	Schedule Modifications	Best Response (Aug 15)	Update on uPRs (Oct 6)
Prostate	CDK12	120 QD <u>(5/2)</u>	PR at 6w. Switched to 3/4 after 11w	cPR	
Ovarian	BRCA1	160 QD (3/4)		cPR	
Ovarian	BRCA1	160 QD <u>(5/2)</u>	Switched to 3/4 after 6w	cPR	
Breast	BRCA1	160 QD (3/4)		uPR	Unconfirmed
Prostate	ATM	120 QD <u>(5/2)</u>	Switched to 3/4 after 7w	uPR	Confirmed
Ovarian	BRCA1	120 QD (3/4)*	Switched to monotherapy after 2w	cPR	
Melanoma	BRCA2	120 QD (3/4)		uPR	Confirmed
SCC	BRCA1	160 QD (3/4)		uPR	Confirmed
Prostate	ATM	120 QD <u>(5/2)</u>	Switched to 3/4 after 17w	cPSA**	
Prostate Nonmeasurable disease	ATM	160 QD (3/4)		cPSA	
Ovarian	BRCA1	120 QD (3/4)		cCA-125	
Ovarian	RAD51C	160 QD (3/4)		cCA-125***	

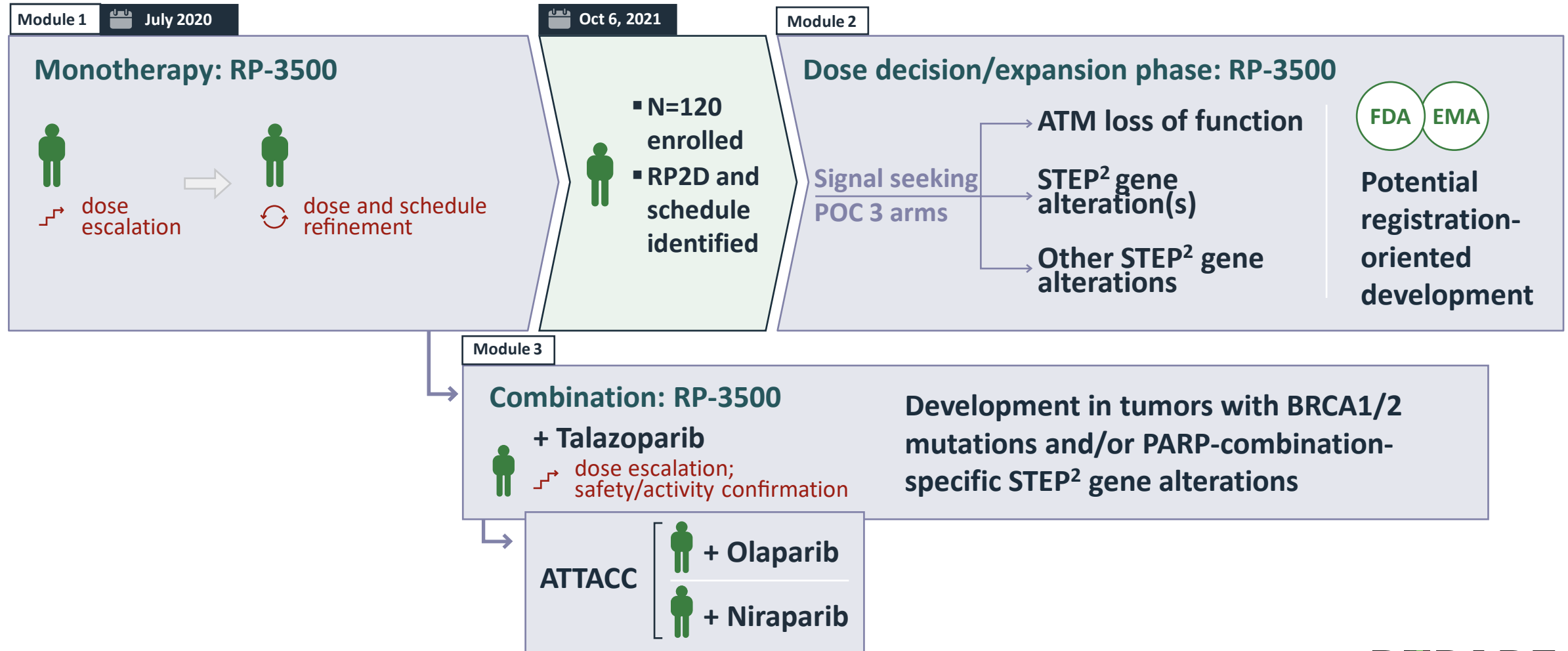
* Initial 32 weeks on talazoparib combination, 23 w on therapy

** Approx. 8 Mo on therapy (SD last scan **-26% on Sept 23rd**)

*** On-Tx 15weeks: SD **(-28.7% at last scan on 07Sep)**

RP-3500 clinical program as of Oct 6th

Global multicenter study designed for patients with any recurrent tumor with ATM loss or loss of any of the additional 16 STEP² genes



RP-3500 TRESR study – Updates to upcoming Module 2

Global multi-center study:

- Phase 2 study with 3 distinct arms designed to deliver PoC
- Expected 4Q 2021 start
- Results will inform discussion with the FDA/EMA on next steps

ATM loss of function

- Selected ATM LoF patients
- Less heavily pre-treated patients than in Module 1

STEP² gene alteration(s)

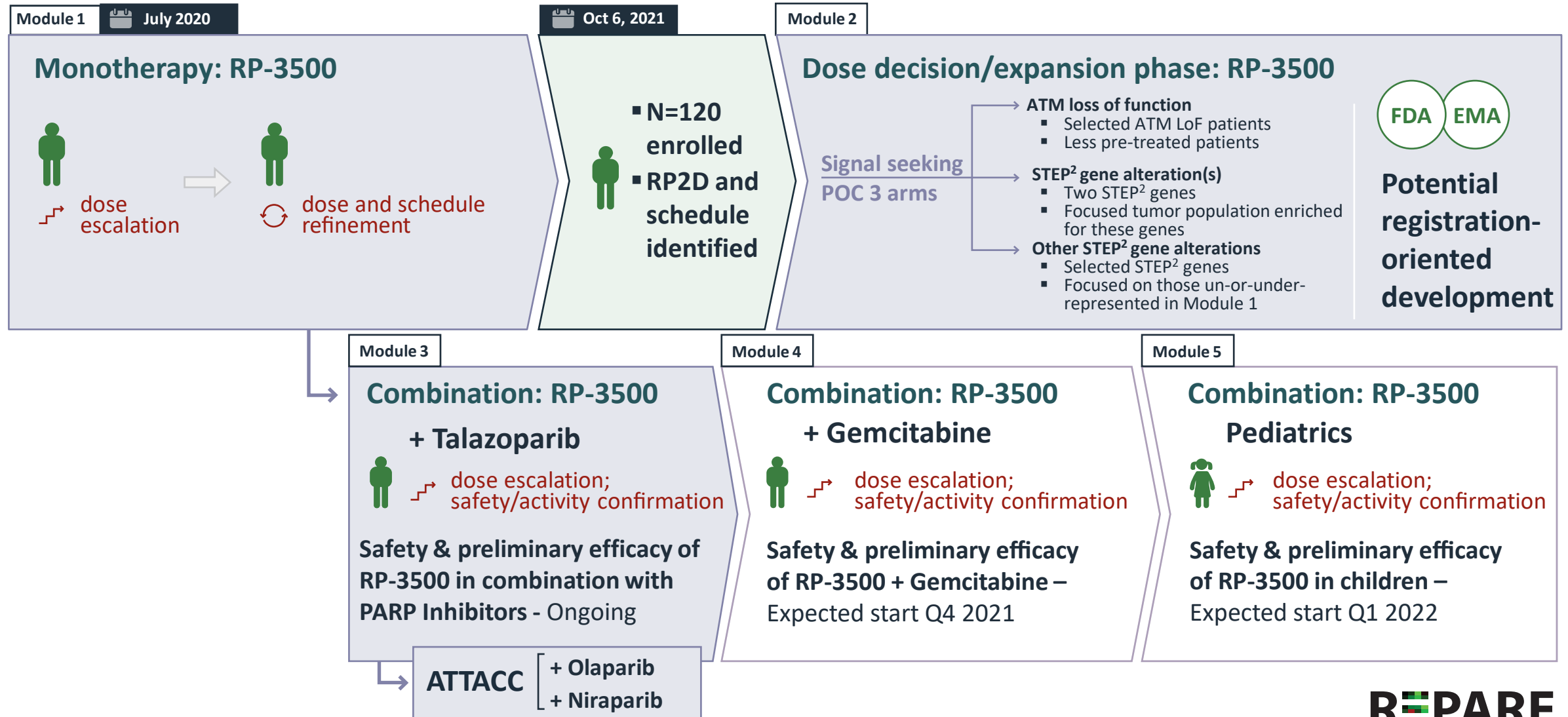
- Two STEP² genes
- Focused tumor population enriched for these genes

Other STEP² gene alteration(s)

- Selected STEP² genes
- Focused on those un/under-represented in Module 1

RP-3500 updated clinical trial program: additional modules

Trial results to date support expanded clinical development



RP-3500 key clinical summary

Early TRESR data provide clinical POC and validate Repare Therapeutics' SNIPRx platform

Favorable and differentiated safety profile and distinct early activity, offer a clear direction for further development

RP-3500 RP2D for further monotherapy evaluation is 160mg QD 3d on/4d off

Biomarker data confirm proof-of-mechanism across different tumors with multiple molecular backgrounds

Preliminary results observed in tumors with selected genomic alterations continue to mature

Final efficacy and safety analysis from Module 1 expected in 2Q 22

Meaningful clinical benefit in 34 (49%) of 69 patients as of 15th August:

- **12 patients** with objective tumor responses:
 - 8 RECIST V1.1 cPR/uPR- 1 unconfirmed PR
 - 2 PCWG3 PSA responses
 - 2 GCIG CA125 responses
- **14 patients** with ongoing RECIST v1.1 SD for at least 16 weeks
- **8 patients** <16w on study: SD with early significant decreases in tumor markers and <30% tumor shrinkage

Recent progress and upcoming anticipated milestones

2021

2022

[illegible]

Q&A Session

