

Insight that enriches. Precision that empowers.

Corporate Presentation May 2024



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Developing Next-Generation Precision Oncology Medicines

Differentiated, proprietary clinical pipeline

- Lunresertib: First-in-class oral PKMYT1 inhibitor (Phase 1/2)
- Camonsertib: ATR inhibitor (Phase 1/2)
- RP-1664: First-in-class selective PLK4 inhibitor (Phase 1)

Multiple clinical catalysts expected in 2024

- Key readouts from ongoing trials
- New clinical entries (PLK4 and Pol0 ATPase inhibitors)



Proprietary CRISPRenabled SNIPRx platform

- Focused on genomic instability and DNA damage repair
- Clinical trials enriched for patients with tumors carrying a network of synthetic lethal alterations (STEP²)

Strong balance sheet

- Cash and investments of ~\$237M¹ fund operations to mid-2026
- Multiple clinical catalysts in that timeframe

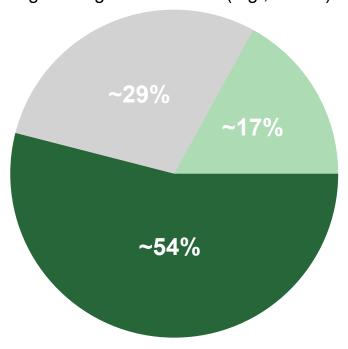


Targeting the untargetable through synthetic lethality



Precision oncology last 20 years:

Targetable gain of function (e.g., EGFR)



REPARE

T H E R A P E U T I C S

Focused on 71% untapped target space, conventionally untargetable

- Gain of function (e.g., CCNE1, 17%)
- Loss of function (no known driver; e.g., BRCA1, 54%)



Specifically targeting and disrupting genes essential for cancer cell survival



SNIPRx identifies and targets necessary genes to induce synthetic lethality

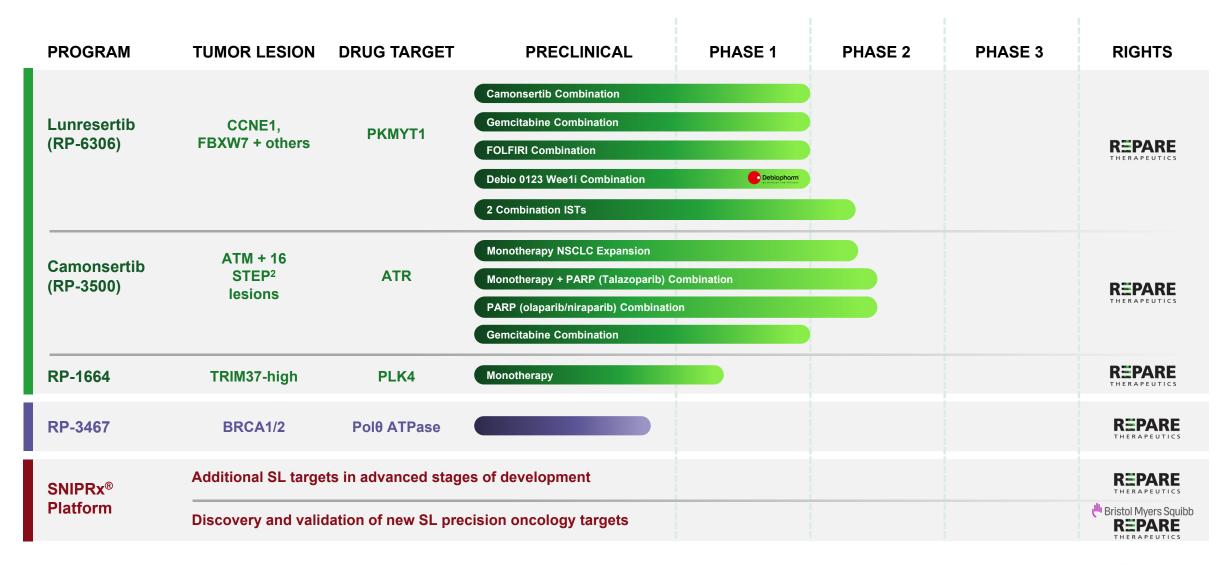
- Highly targeted & tumor-type agnostic approach
- Exploiting cancer cell genomic instability, including DNA damage repair



Platform validated with established and expanding clinical-stage pipeline



Expanding pipeline of precision oncology therapeutics





Proven experience in drug discovery and development



Leadership Team



Lloyd M. Segal President & CEO

McKinsey & Company







Steve Forte, CPA Chief Financial Officer

clementia APTALIS





Michael Zinda, PhD Chief Scientific Officer

AstraZeneca Lilly





Maria Koehler MD, PhD Chief Medical Officer









Cameron Black, PhD Head of Discovery







Philip Herman Chief Commercial, Portfolio Development Officer











Kim A. Seth, PhD Chief Business Officer







Daniel Bélanger Head of Human Resources





Scientific Founders



Daniel Durocher, PhD

- Developed CRISPR SL platform
- Deep DNA repair knowledge
- Lunenfeld-Tanenbaum Research Institute (LTRI) & professor at University of Toronto



Agnel Sfeir, PhD

- DDR and cancer pathway investigator
- Pioneer in Polθ, genome instability
- Professor, MSKCC



Frank Sicheri, PhD

- Globally recognized structural biologist, expert in eukaryotic cell signaling, drug mechanism of action
- LTRI & professor at University of Toronto



Lunresertib (RP-6306)





Lunresertib:

First-in-class, oral, small molecule, PKMYT1 inhibitor

Large, genomically defined potential addressable patient population of ~90k

 50% RECIST response in camonsertib combination in gynecological tumors



Anti-tumor activity observed

- Across multiple tumor types and genotypes
- POC in patients established
- FDA agreed with RP2D; safe and well tolerated

Repare discovered synthetic lethality of PKMYT1 inhibition

- Initially identified CCNE1 amplification
- STEP² screen identified additional genes
 FBXW7 and PPP2R1A
 - First and currently the only PKMYT1 inhibitor in clinical trials

Supported preclinical synergy hypothesis and patient selection approach

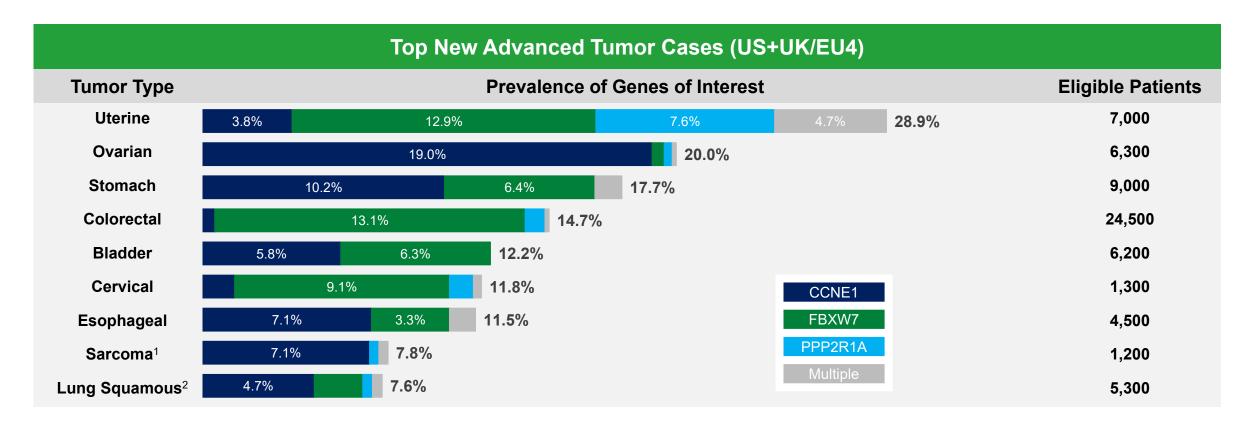
from proprietary SNIPRx platform





Large, genomically defined potential patient population

~90K addressable patients including ~65K among top tumors with genetic alterations largely mutually exclusive



^{*} Based on estimated number of pts US+UK/EU4 treated in 1st line, advanced setting for diagnosed and new recurrent patients (CancerMPact®, Treatment Architecture, United States, 2021; accessed 5/19/23) and lesion prevalence (TCGA). ¹ Soft Tissue Sarcoma only; ² Squamous subtype of Non-Small Cell Lung Cancer only



Evolving broad trial program: sponsored and collaborative





Key inclusion criteria:

Recurrent solid tumors

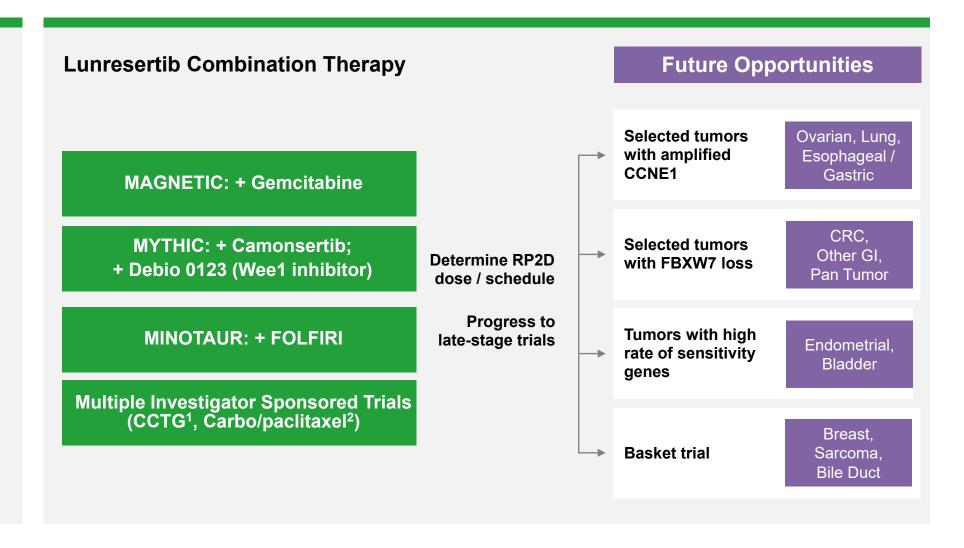
CCNE1

amplification or

PPP2R1A

FBXW7

inactivating mutations



¹ Canadian Clinical Trial Group (CCTG) collaborations include NCT05605509 and NCT05601440.



² Standard of care ("SOC") for 1st line ovarian cancer is carbo/paclitaxel (6 cycles) + PARPi maintenance therapy or carbo/paclitaxel with bevacizumab + bev maintenance therapy; this IST supports future potential 1st line combination studies as triplet therapy in patients with CCNE1 amplified tumors.



Lunresertib:

MYTHIC Preliminary Phase 1 Trial Results (M1: Monotherapy) (M2: Camonsertib Combination Therapy)

CAMONSERTIB COMBINATION THERAPY

Safe, well tolerated and promising anti-tumor activity observed across tumors and all lunresertib-sensitizing genomic alterations (N=59)

23.6% OR; 41.8% CBR in efficacy-evaluable patients (N=55)

33.3% OR; 50.0% CBR at preliminary RP2D range, across all tumors (N=18)

38.5% OR; 57.7% CBR in patients with heavily pre-treated gynecologic cancers (N=26); **50% RECIST response** at preliminary RP2D (N=10)

Dose/schedule optimization complete; RP2D of lunresertib 80mg twice daily and camonsertib 80mg once daily

MONOTHERAPY

Safe, well tolerated and anti-tumor activity observed (N=67)

Recommended Phase 2 dose: 80 mg twice daily in intermittent schedule



Key updates since ENA 2023; registrational decision on track



Registrational decision on track in gyn expansions in Q4 2024

Continuing trends of patient response and benefit

Grade 3 anemia reduced from 45% to 25% at RP2D with updated dosing

2 weeks on / 1 week off for patients with low Hg, otherwise weekly

FDA agreed with RP2D

Efficacy assessment is ongoing, continues to be **promising and on track** to be shared by end of Q4 2024

Data is expected to include ~20-30 patients per histology (ovarian and endometrial) at RP2D



Lun + cam responses across tumor types and genotypes

RECIST and tumor marker responses occurred early despite heavily pre-treated, relapsed/refractory patient population

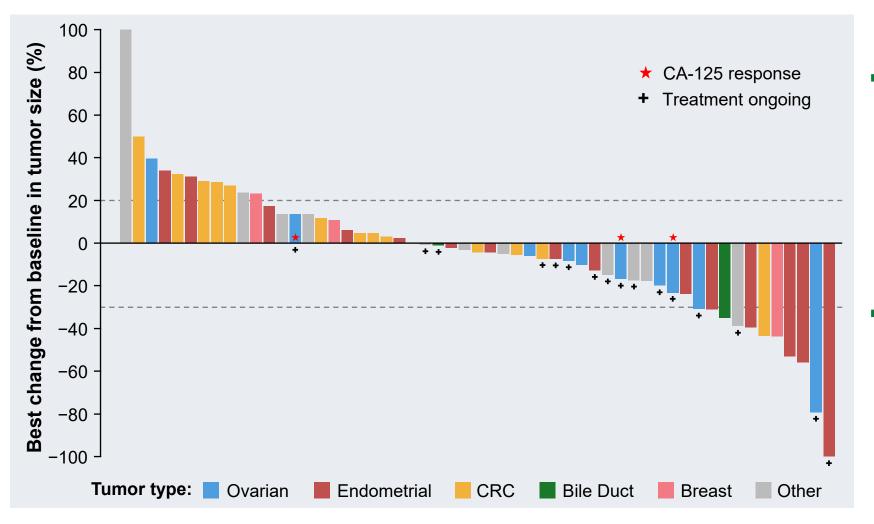
Tumor type	Genotype	Response	Best % change in TL from BL	Therapy (weeks)	Lines of prior Tx/ prior platinum
	PPP2R1A/FBXW7	cPR	-55.9	30.4	3/Y
	PPP2R1A/CCNE1	cPR	-53.0	18.1	2/Y
Endometrial	FBXW7	cPR*	-100.0	11.1+	3/Y
	FBXW7	uPR	-39.6	16.0	3/Y
	FBXW7	uPR*	-44.7	11.4+	3/Y
	CCNE1	cPR*	-70.2	21.4+	2/Y
	CCNE1†	cPR*	-30.8	12.6+	3/Y
Ovarian	CCNE1	CA-125	-16.9	29.0+	9/Y
	CCNE1	CA-125	-23.1	37.0+	2/Y
	CCNE1	CA-125	13.6	12.9+	5/Y
Cervical	PPP2R1A	cPR*	-44.4	11.0+	1/Y
Colorectal	FBXW7	cPR	-43.3	27.6	3/Y
Bile duct	CCNE1	cPR	-35.0	28.1	2/Y
Breast	FBXW7‡	uPR	-43.8	18.1	2/N

^{*} One response evaluable patient became uPR and four patients had responses confirmed after the Sept. 5, 2023 cutoff, data as of Oct. 6, 2023. Relevant patient tumor comutations †BRCA1 rearrangement and †BRCA2 biallelic loss. +Treatment ongoing. BL, baseline; cPR, confirmed partial response; RECIST, Response Evaluation Criteria in Solid Tumors; TL, target lesion; Tx, treatment; uPR, unconfirmed partial response.



Frequent and deep tumor reductions observed with lun + cam





In evaluable patients*, across all tumors/doses:

- OR: 23.6% (n=55)

CBR: 41.8% (n=55)

- MRR: 50.0% (n=24)

At preliminary RP2D, across all tumors:

- OR: 33.3% (n=18)

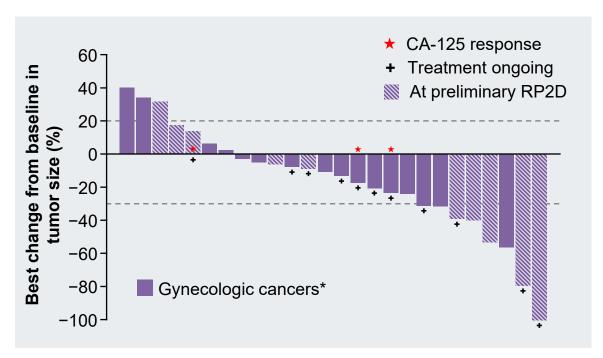
CBR: 50.0% (n=18)

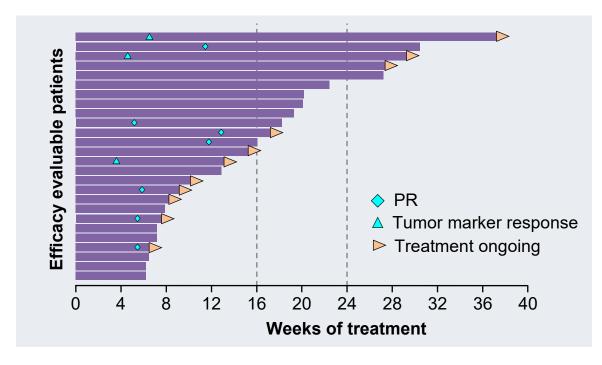


^{*}Efficacy evaluable patients only (≥1 post-baseline tumor assessment). Other tumor types include cervical (n=1), esophageal (n=1), GI (n=1), liver (n=1), lung (n=3), melanoma (n=1), osteosarcoma (n=1), pancreatic (n=2), and upper GI (n=1). CBR: overall response or time on treatment ≥ 16 wk w/o progression; CRC, colorectal cancer; RECIST, Response Evaluation Criteria in Solid Tumors (RECIST) Gynecologic Cancer InterGroup (GCIG); MRR, molecular response rate; OR, overall response based on RECIST or GCIG CA-125 response; RP2D, recommended phase 2 dose; lun, lunresertib.

Combination treatment effective in gynecologic tumors

Meaningful tumor reductions, durable clinical benefit observed in heavily pre-treated patients to date





Across all doses (n=26):

Overall response: 38.5%; RECIST Response: 26.9%

CBR: 57.7%; MRR: 8/10 (80%)

At preliminary RP2D (n=10):

Overall response: 60%; RECIST Response: 50%

CBR: 70%

Patients had 1-9 prior therapies (median=3); treatment ongoing in 11 patients; enrollment in expansion cohorts at RP2D continues

^{*} Gynecologic cancers: ovarian, endometrial, and cervical cancers. Data represent the efficacy evaluable population (≥1 post-baseline tumor assessment). CBR, clinical benefit rate; OR, overall response based on RECIST or GCIG CA-125 response; MRR, molecular response rate; PR, partial response; RECIST, Response Evaluation Criteria in Solid Tumors; Gynecological Cancer InterGroup (GCIG); RP2D, recommended phase 2 dose.



Significant improvement in anemia observed with updated dosing

RP2D: lunresertib 80mg BID + camonsertib 80mg QD 3d/4d

	RP2D (ENA Cutoff) ^a N=20		RP2D (Cohort Post ENA) N=44			
Selected hematologic TRAEs, n (%)	All Grades	Gr3	Gr4	All Grades	Gr3	Gr4
Anemia	13 (65.0)	9 (45.0)	0	29 (65.9)	11 (25.0)	0
Leukopenia	3 (15.0)	0	0	9 (20.5)	3 (6.8)	0
Neutropenia	3 (15.0)	2 (10.0)	0	7 (15.9)	5 (11.4)	0
Thrombocytopenia	0	0	0	0	0	0

Updated dosing strategy reduced Grade 3 anemia by ~half Hematologic safety profile similar to commercial SL agents No thrombocytopenia observed



Continued favorable safety profile observed to date

Lun+Cam RP2D

	N=65 ^a			
TRAEs in ≥10% of patients, n (%)	All Grades	Gr3	Gr4	
Nausea/Vomiting	34 (52.3)	0	0	
Rash ^a	26 (40.0)	1 (1.5)	0	
Fatigue	18 (27.7)	1 (1.5)	0	
Stomatitis	18 (27.7)	4 (6.2)	0	
Decreased appetite	13 (20.0)	0	0	
Diarrhea	10 (15.4)	0	0	
Headache	7 (10.8)	0	0	
Constipation	5 (7.7)	0	0	

- Patient demographics remain comparable:
 - Entry Hg
 - Gender and age
 - Prior lines and therapies
 - ECOG
 - Histologies and DOT
- Differences in anemia rates likely a result of the updated dosing strategy

FDA agreement on RP2D

No FDA comments raised about safety profile observed in lun + cam combination

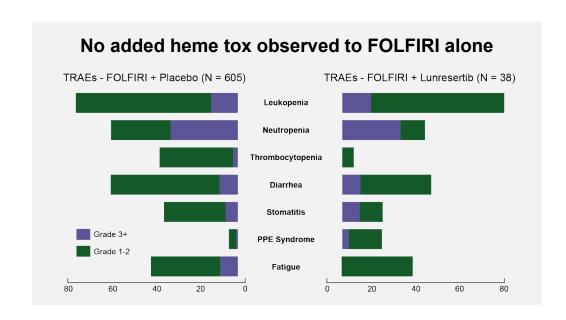


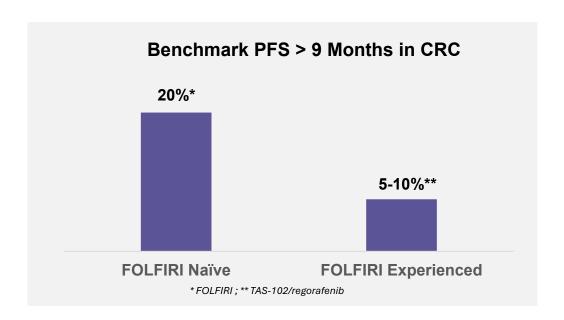
Lun + FOLFIRI combination promising



MINOTAUR demonstrating overall favorable tolerability and early signal in CRC and other GI tumors

- Favorable tolerability: lunresertib given continuously daily (QD), demonstrating promising tolerability profile
 vs. other FOLFIRI combinations
- No new toxicities observed; no treatment withdrawals at RP2D
- Focus on potential for duration of treatment advantage in both FOLFIRI-naïve and experienced patients





Full data to be shared at ESMO GI in June 2024



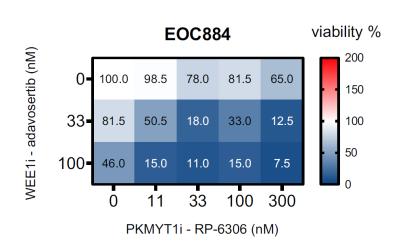
First clinical trial inhibiting both PKMYT1 and WEE1





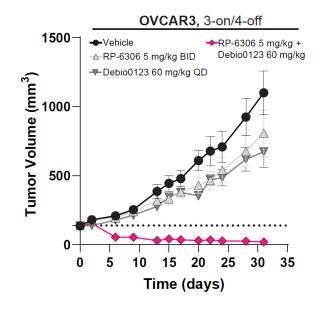
Strong preclinical evidence of PKMYT1 + WEE1 inhibitor combination potential; Ph1/1b now enrolling





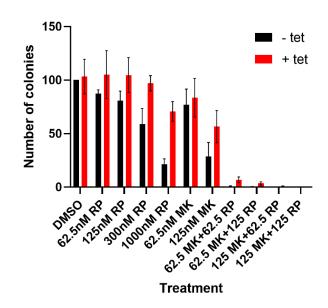
Combination synergistically eradicates ovarian cancer cells and organoid models at a low doses...

Benada et al., NAR Cancer, 2023.



...drives tumor regressions on intermittent schedule at doses below monotherapy EC₅₀ ...

Gallo et al., ANE 2023, Poster #A023.



... and overcomes resistance to MK-1775 (adavosertib) mediated by tet-induced Myt1 upregulation

Sokhi et al., AACR 2023, Poster #5511.



Platinum-resistant ovarian cancer (PROC) market opportunity



Unmet need remains significant for platinum resistant patients



Chemo + Bevacizumab¹

28% ORR 6.8 mo. PFS 16.6 mo. OS Single Agent Chemo^{1,2}

13 - 16% ORR 3.4 - 4.0 mo. PFS 12.8 - 13.3 mo. OS Mirvetuximab²

42% ORR **5.6 mo. PFS** 16.5 mo. OS ~2,900 2L+ Patients
with CCNE13



¹ Chemo+Bevacizumab vs Chemo (AURELIA); Source: Bevacizumab FDA Label

² Mirvetuximab vs Chemo (MIRASOL); Source: Mirvetuximab FDA Label, ASCO 2023. Mirvetuximab is approved for ~1/3 of PROC patients who are folate receptor positive.

³ Eligible Patients in US and EU4/UK Based On Company Estimates from TCGA and GENIE, 2L+ (2L - 5L) PFI, progression-free interval.

Endometrial cancer market opportunity



Evolving 1L SOC towards Chemo + ICI creating large unmet need in future 2L+ setting

1st Line Therapy

Chemo +/- Immune Checkpoint Inhibitor ("ICI")

No Prior ICI

Prior ICI

2nd Line+ Therapy

ICI or Single Agent Chemo

No SOC (various chemotherapies)

Unmet Need

Single Agent Chemo^{1,2}

15 - 16% ORR

3.8 - 4.0 mo. PFS

12.0 - 12.3 mo. OS

~3,600 2L+ Patients with CCNE1, FBXW7, or PPP2R1A³



Metastatic CRC is a large market opportunity for MINOTAUR



Unmet need in 2L+ oxaliplatin-treated mCRC patients

1st Line Therapy

Oxaliplatin-based Chemotherapy +/VEGF

FOLFIRI+ VEGF^{1,2,4}

13 - 20% ORR

5.7 - 9.2 mo. PFS

13.3 - 21.4 mo. OS

FOLFIRI^{1,2,3}

11 - 15% ORR

4.5 - 5.6 mo. PFS

11.7 - 13.8 mo. OS

2nd Line+ Therapy

Irinotecan-based Chemotherapy +/- VEGF KRAS Inhibitors (KRAS G12C) Encorafenib + Cetuximab (BRAF V600)

Unmet Need

~11,300 2L+ Patients

with FBXW7⁵ (~13% of CRC)

G7 Colorectal Cancer market:

>\$8B today (>\$10B by 2032)



¹ FOLFIRI+Aflibercept vs FOLFIRI (VELOUR); Source: Aflibercept FDA Label

² FOLFIRI+Ramucirumab vs FOLFIRI (RAISE); Source: Ramucirumab FDA Label, Lancet 2015

³ Napabucasin+FOLFIRI vs FOLFIRI+Bevacizumab (CanStem303C); Source: Shah M. Clinical Colorectal Cancer 2022

⁴ Panitumumab+FOLFIRI vs FOLFIRI+Bevacizumab (SPIRITT); Source: Hecht JR. Clinical Colorectal Cancer 2015

⁵ Eligible Patients in US and EU4/UK Based On Company Estimates from TCGA and GENIE. 2L+ (2L - 5L)

Camonsertib (RP-3500)







Demonstrated synthetic lethal interaction of ATR and a network of genes identified by SNIPRx and

STEP² process

Proof of concept established in Phase 1/2 monotherapy trial



Potential best-in-class ATR inhibitor

Durable antitumor activity in combination with PARPi; meaningful clinical benefit observed in ovarian cancer Global development and commercialization rights **wholly-owned** by Repare

- Rapid monotherapy signal confirmation in **NSCLC**



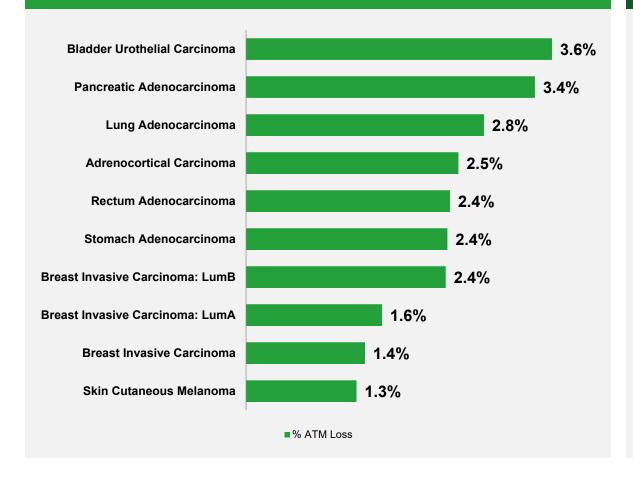




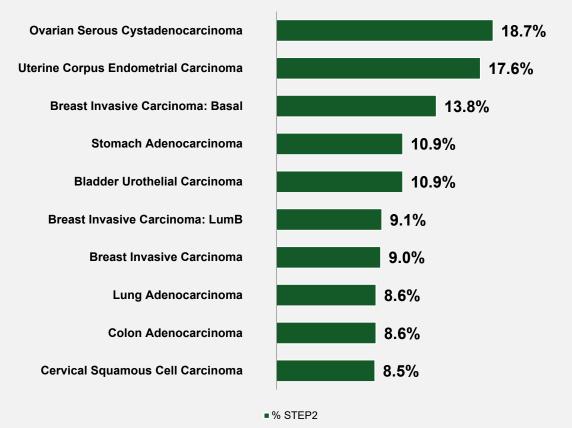
Potential across significant additional patient populations







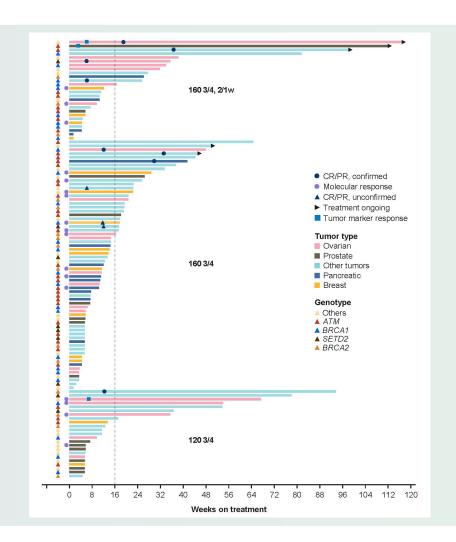
Top 10 Tumor Types* with Highest Prevalence of ATM Deficiency or STEP² Genomic Alterations





Updated camonsertib monotherapy data in ATMm tumors

- Updated data continues to support ATR-ATM synthetic lethality thesis across various tumor types and genotypes
- 114 total efficacy evaluable patients treated at 3 efficacious dose levels
- 36 patients enrolled with ATM alterations
 - 4 with responses and treatment durations 41-112+
 weeks
 - 9/36 (25%) total with Tx duration >6 months

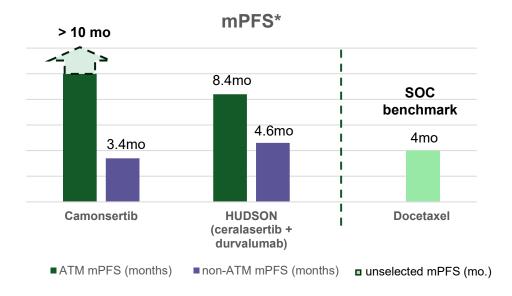




Camonsertib: rapid monotherapy signal confirmation in NSCLC

- >12 months durability observed in >20% of patients with ATMm tumors treated with camonsertib monotherapy
- ATMm NSCLC (~4% of NSCLC) an attractive opportunity
 - Camonsertib monotherapy signal potentially offers rapid and costefficient path to PoC with ~15-20 more patients within TRESR
 - 11 NSCLC patients (4 with ATMm) highlight improved mPFS in ATMm NSCLC vs non-ATMm
 - AstraZeneca HUDSON Ph2 data subset (ATR + PD-L1 post IO) further supports ATMm hypothesis in NSCLC
 - ATMm tumors do not have better outcomes in NSCLC
- TRESR open to enrollment; data expected in 2025, with potential for expansion
- IO collaborations beyond monotherapy an obvious, substantial opportunity

Promising Camonsertib mPFS in NSCLC





Camonsertib NSCLC market opportunity



Significant unmet need for non-biomarker driven NSCLC patients

Non-Biomarker Driven Patients

1st Line Therapy

Chemoimmunotherapy Immunotherapy (PD-L1)

2nd Line+ Therapy

Single Agent Chemotherapy +/- VEGF Immunotherapy (PD-L1)

Unmet Need

Biomarker Driven Patients (EGFR, ALK, KRAS, ROS, etc.)
1+ Lines of Biomarker Driven Therapies

Ramucirumab + Chemo²
23% ORR

4.5 mo. PFS

10.5 mo. OS

Single Agent Chemo^{1,2}

13 - 14% ORR

3.0 - 4.0 mo. PFS

9.1 - 9.6 mo. OS

~5,300 2L+ Patients



¹ Atezolizumab vs Chemo (OAK); Source: Atezolizumab FDA Label

² Ramucirumab+Chemo vs Chemo (REVEL); Source: Ramucirumab FDA Label

³ Eligible Patients in US and EU4/UK Based On Company Estimates from TCGA and GENIE. 2L+ (2L - 5L)

RP-1664







Highly potent, selective and bioavailable PLK4 inhibitor synthetically lethal with TRIM37 gain of function

Strong, dose-dependent anti-tumor activity observed as monotherapy across preclinical models

RP-1664

First-in-class, oral PLK4 inhibitor

Actively enrolling in solid tumors and neuroblastoma

~63K addressable patient population with limited treatment options; potential across multiple tumor types

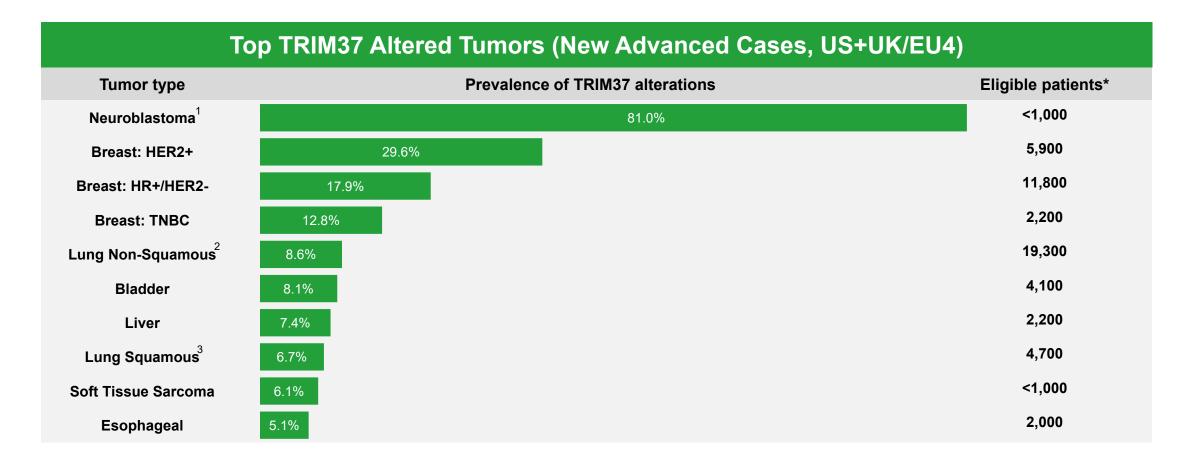






High prevalence in patient populations with limited treatment options

~63K patients with TRIM37 amplification or overexpression, with ~53K among top tumors





Compelling synthetic lethal rationale for targeting PLK4



Biomarker-driven patient selection hypothesis for development of oral PLK4i for TRIM37-high tumors

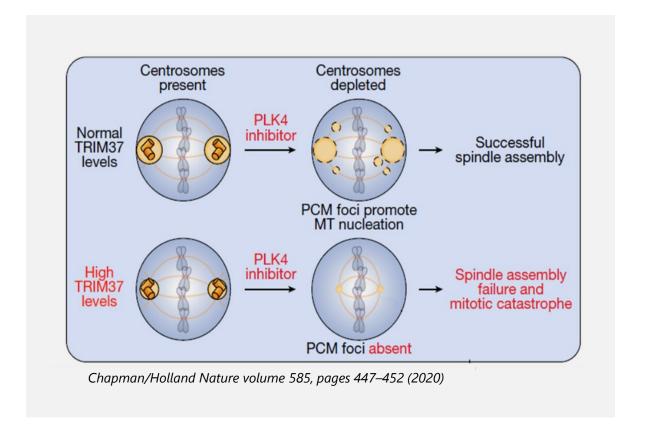
Centrosomes use centrioles and pericentriolar material (PCM) for mitotic spindle formation

Polo-Like Kinase 4 (PLK4) required for centriole creation in S-phase

TRIM37 (an E3 Ligase) reduces PCM stability; excess TRIM37 depletes PCM, increasing cell reliance on centrioles for spindle assembly

Thus, PLK4 inhibition is harmful in cells with high TRIM37 and low PCM

Validated in two 2020 Nature publications





Potential first-in-class oral PLK4 inhibitor

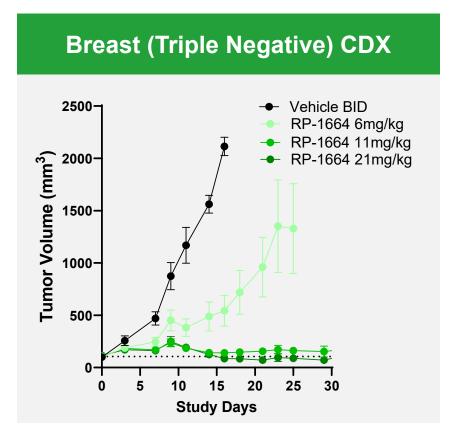
	Key Parameter	RP-1664
	PLK4 Enzyme IC ₅₀	1 nM
0	PLK4 cell binding IC ₅₀	3 nM
In vitro	Cell proliferation in MCF7 / T47D (TRIM37 amp) EC ₅₀	51 / 17 nM
드	Cell-base selectivity vs AurA, AurB	>2000-fold
	Kinome screen at 90x PLK4 IC ₅₀	8/280 kinases >50% inh
	Human Hepatocyte Clearance (µL/min/10 ⁶ cells)	2.2
ADME	Rat PK (%F, t _{1/2})	28%, 4h
АБ	Monkey PK (%F, t _{1/2})	96%, 9h
	WIOTINGY I IX (701, L _{1/2})	JU /0, JII

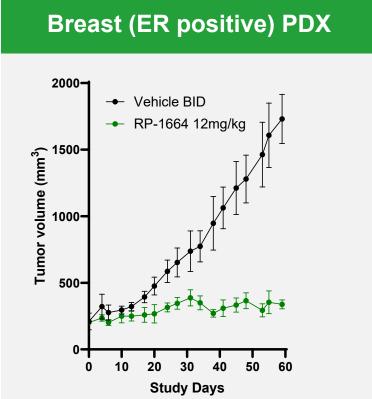
- Highly potent, selective and orally bioavailable PLK4 inhibitor
 - − ~10x more potent than competitor molecules¹ with vastly improved selectivity vs AurB
- Clean in PanLabs safety pharmacology screen

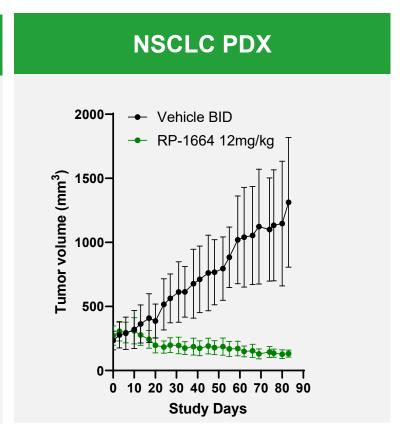


Robust monotherapy efficacy across solid tumor PDX/CDX models

Monotherapy drives tumor stasis to regression in TRIM37-high models





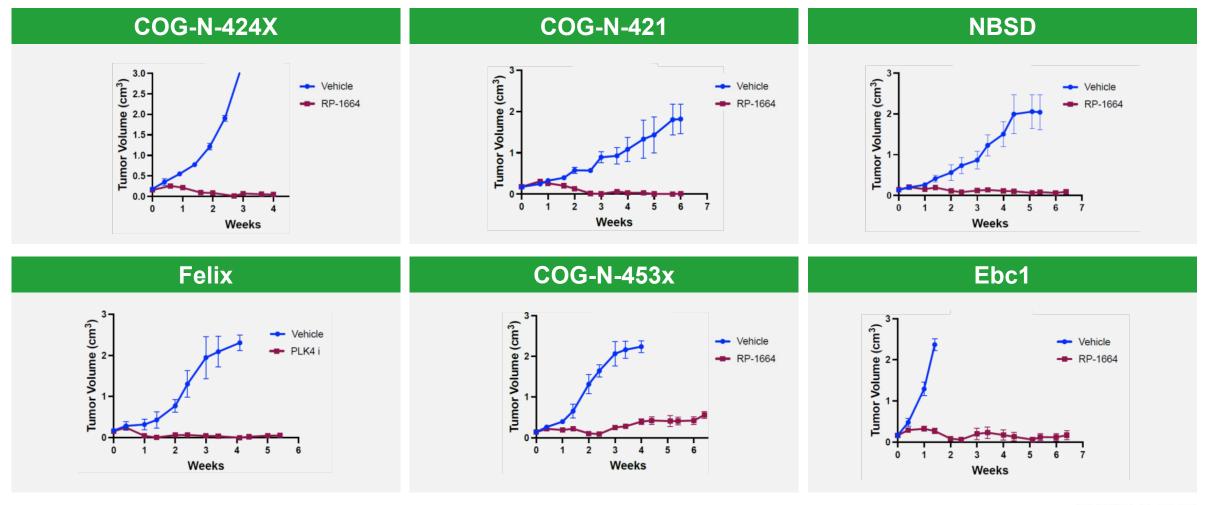




Highly efficacious as monotherapy in neuroblastoma models



Neuroblastoma PDX and CDX models (all TRIM37-high) conducted at Children's Hospital of Philadelphia demonstrate deep and prolonged monotherapy regressions in 5 of 6 evaluable models

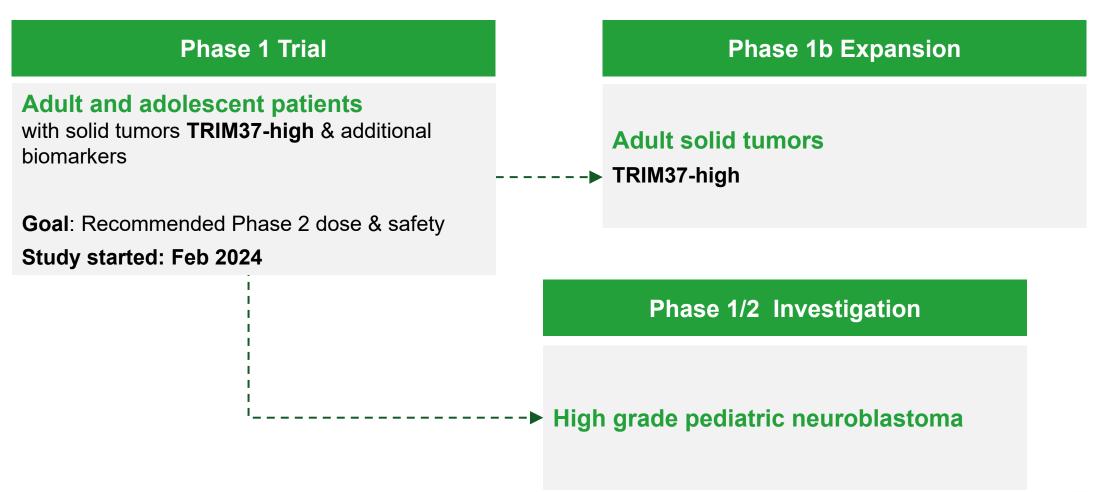




RP-1664 Phase 1/2 monotherapy clinical development plan



Efficient RP-1664 Phase 1 plan enables early start for pediatric dose finding study in neuroblastoma and clear view on adult solid tumor opportunity





RP-3467







Highly potent, selective
Polθ ATPase inhibitor;
inhibits DNA repair and is
synthetic lethal with
BRCA loss

Demonstrates compelling potential for combination efficacy without added toxicity



Potential best-inclass Polθ ATPase inhibitor

FPI in 2H 2024

RP-3467 capable of complete regressions and synergies in PARPi resistance preclinical models

Global market segments comprise \$16 billion in PARP inhibitors, RLT, and chemotherapy









RP-3467 clinical plan: multiple potential Phase 1/2 trials



Phase 1 Trial

Initiation expected

Goal: PK, safety and RP2D

in H2 2024

PARPi combination – PARP1/2 or PARP1

Phase 1/2 Trials

Deep/durable complete responses preclinically, with no additional toxicity

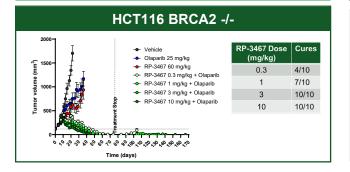
RLT combination

Survival benefit preclinically in unselected tumor backgrounds, with no additional toxicity

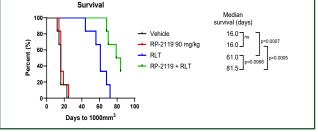
Chemotherapy / ADC Payloads

Well tolerated preclinically in combination with carboplatin and irinotecan, including topoisomerase ADC payloads

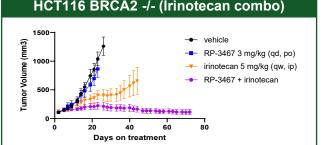
Preclinical Results



Combination Survival Benefit Survival



HCT116 BRCA2 -/- (Irinotecan combo)



Global Market Segment

~\$3 Billion

~\$8 Billion

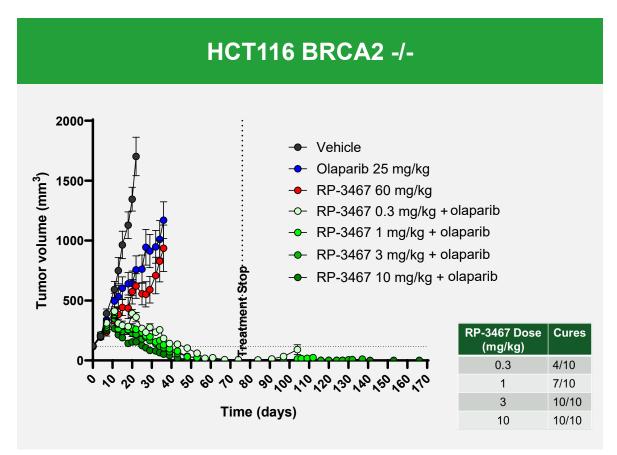
~\$5 Billion

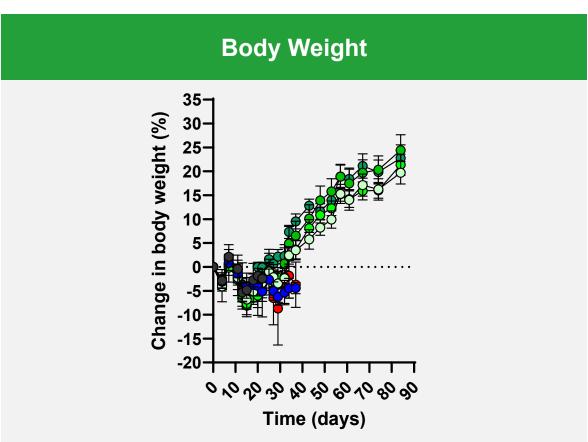


Profound, durable synergy observed with PARP inhibition



Deep/durable complete regressions observed across a wide dose range and well tolerated







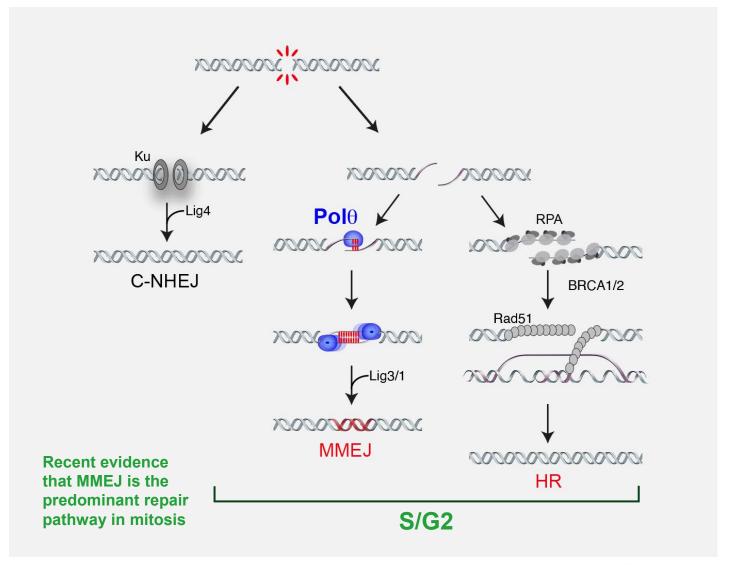
Polθ: uniquely promising therapeutic target

Polθ is a unique multifunctional DNA polymerase with ATP-dependent DNA helicase activity

Required for microhomology-mediated end joining (MMEJ), a **key mechanism** of double-strand DNA break repair

Uniquely active to repair double-strand DNA breaks during mitosis

Minimally expressed in normal tissue and knockout animals have no significant phenotype





Target profile: potent, tolerable, capable of complete regressions

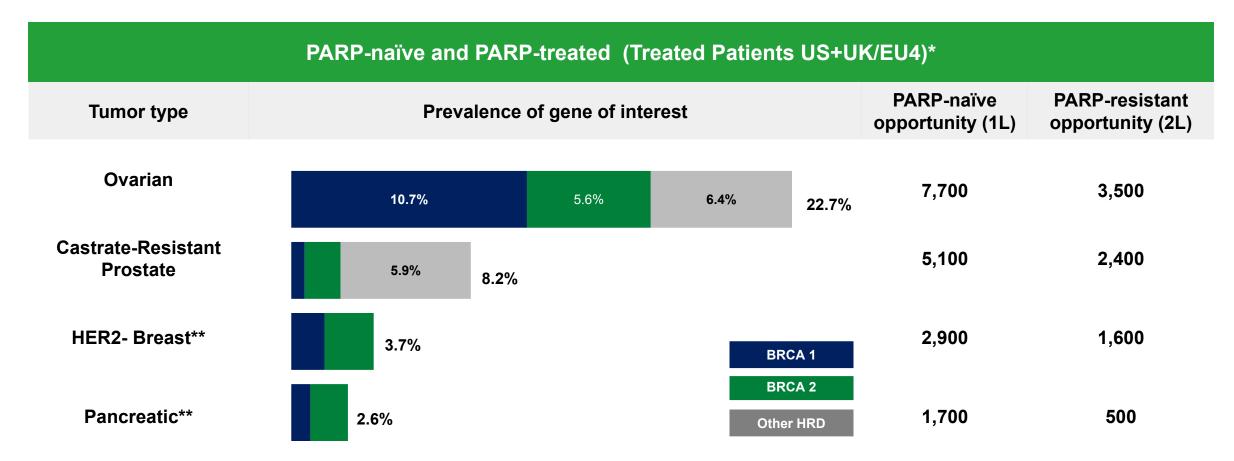
	Parameter	RP-3467	Complete regressions in PDX models at low doses			
	Polθ ATPase Enzyme IC ₅₀	<0.25 nM				
Potency	CETSA cellular target engagement IC ₅₀	5 nM	HBCx-22 (BRCA2null)	HBCx-10 (BRCA2null)		
	Cell proliferation DLD1 / HCT116 (BRCA2mt) EC ₅₀	4 / 7 nM	1750-	2000 – 1750 –		
lectiv	Off-target ATPase (HELQ, WRN, BLM) IC ₅₀	> 10 µM	© 1250- © 1000-	1500- 1250- 12		
	Off-target Polθ polymerase domain IC ₅₀	> 100 µM	1000 - Vehicle - 50 mg/kg Olaparib - 3 mg/kg RP-3467 - 0.3 mg/kg RP-3467 + olaparib - 3 mg/kg RP-3467 + olaparib - 3 mg/kg RP-3467 + olaparib	25 mg/kg RP-3467 3 mg/kg RP-3467 + olaparib 3 mg/kg RP-3467 + olaparib 3 mg/kg RP-3467 + olaparib		
	Human Hepatocyte Clearance (μL/min/10 ⁶ cells)	2.1	250	250-		
	Rat PK (%F, t _{1/2})	123%, 6h	0 10 20 30 40 50 60 70 Days on treatment	o 10 20 30 40 50 60 Days on treatment		
	Monkey PK (%F, t _{1/2})	60%, 3h				

- Highly potent, selective and orally bioavailable Polθ ATPase inhibitor; clean PanLabs safety pharmacology screen
- RP-3467 demonstrated complete regressions in BRCA1/2 null PDX models, also synergy in a PARPi resistance model



Addressing unmet need in critical patient populations

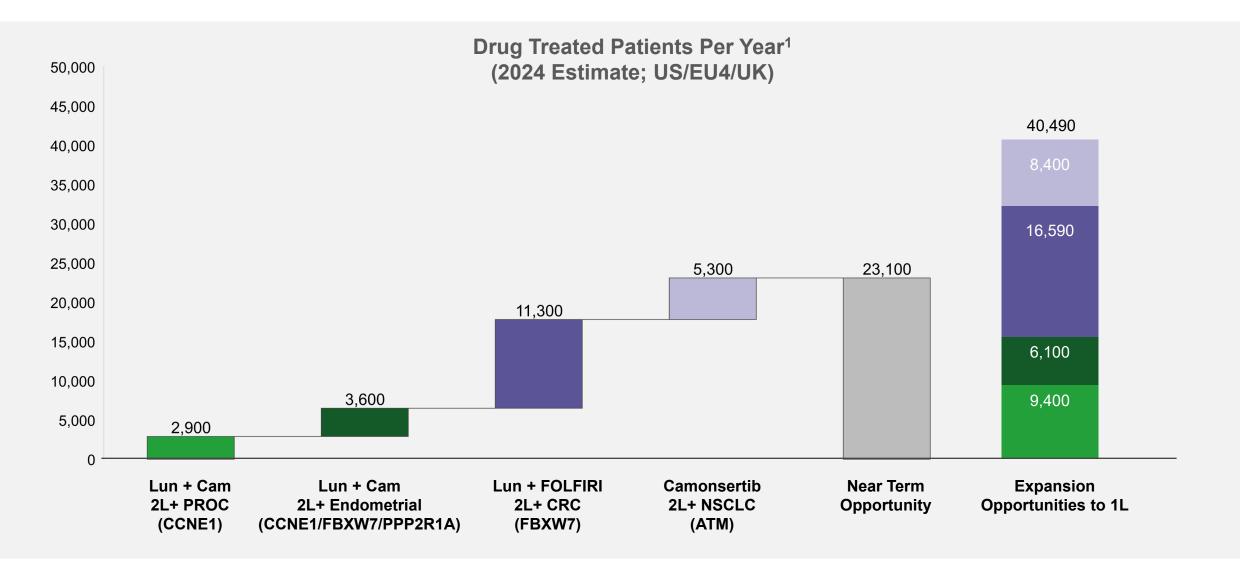
~26K among patients with PARP-naïve and PARP-treated tumors



^{*} Based on estimated number of drug treated pts in the advanced setting likely to be naïve to PARP inhibitor treatment or previously treated with a PARP inhibitor (CancerMPact®, Patient Metrics, 2022; accessed 9/25/23) and lesion prevalence (TCGA; Riaz, N. et al. Pan-cancer analysis of bi-allelic alterations in homologous recombination DNA repair genes. Nat Commun 8, 857 (2017)). Other HRD gene mutations include: BRIP1, ATM, RAD51B, RAD51C, RAD51D, PALB2, BARD1, CDK12, CHEK1, CHEK2, FANCL, RAD54L. ** Includes germline BRCA1/2 only









Recent and expected milestones



1H 2024

2H 2024

2025

- ✓ RP-1664 (PLK4i) clinical trial initiation
- ✓ Lunresertib + Debio 0123 combination Ph1/1b clinical trial initiation
- ✓ Regained camonsertib rights

Initial **lunresertib + FOLFIRI**combination Ph1 data at
ESMO GI in June

Camonsertib monotherapy expansion to NSCLC in TRESR

RP-3467
Ph1 clinical trial initiation

Lunresertib + camonsertib
expansion cohort data in ovarian
and endometrial in Q4

Lunresertib + Debio 0123 combination data

Camonsertib monotherapy data in NSCLC

Initiate first **pivotal trial** for **lun+cam** in 2025



Developing Next-Generation Precision Oncology Medicines

Differentiated, proprietary clinical pipeline

- Lunresertib: First-in-class oral PKMYT1 inhibitor (Phase 1/2)
- Camonsertib: ATR inhibitor (Phase 1/2)
- RP-1664: First-in-class selective PLK4 inhibitor (Phase 1)

Multiple clinical catalysts expected in 2024

- Key readouts from ongoing trials
- New clinical entries (PLK4 and Pol0 ATPase inhibitors)



Proprietary CRISPRenabled SNIPRx platform

- Focused on genomic instability and DNA damage repair
- Clinical trials enriched for patients with tumors carrying a network of synthetic lethal alterations (STEP²)

Strong balance sheet

- Cash and investments of ~\$237M¹ fund operations to mid-2026
- Multiple clinical catalysts in that timeframe





Insight that enriches. Precision that empowers.

Corporate Presentation May 2024

