REPARE THERAPEUTICS

Insight that enriches. Precision that empowers.

Corporate Presentation November 2024

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These forward-looking statements reflect our current beliefs and expectations. Many factors may cause differences between current expectations and actual results, including the duration and impact of the COVID-19 pandemic on our business and market volatility, 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

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Developing Next-Generation Precision Oncology Medicines



- Lunresertib: First-in-class oral PKMYT1i
- Camonsertib: Potential best-in-class ATRi
- RP-1664: First-in-class selective PLK4i
- RP-3467: Potential best-in-class Pol0 ATPase inhibitor

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)

Multiple clinical catalysts in 2024 and 2025

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

Strong balance sheet

- Cash and investments of ~\$180M¹ fund operations into second half of 2026
- Multiple clinical catalysts in that timeframe



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¹ As of September 30, 2024.

Targeting the untargetable through synthetic lethality





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

PROGRAM	TUMOR LESION	DRUG TARGET	PRECLINICAL	PHASE 1	PHASE 2	PHASE 3	RIGHTS
Lunresertib (RP-6306) ¹	CCNE1, FBXW7 + PPP2R1A	PKMYT1	Camonsertib Combination Chemotherapy Combinations (FC Debio 0123 WEE1i Combination	OLFIRI/Gemcitabine)			REPARE
Camonsertib (RP-3500)	ATM + 16 STEP ² lesions	ATR	Monotherapy NSCLC Expansion Other Combinations (PARP Inhib	pitors/Gemcitabine)			REPARE
RP-1664	TRIM37-high	PLK4	Monotherapy (LIONS)				REPARE
RP-3467	BRCA1/2	Polθ ATPase	Monotherapy & PARPi Combo (P	POLAR)			REPARE THERAPEUTICS
SNIPRx®	Additional SL targe	ts in advanced stage	es of development				REPARE
Platform	Discovery and validation of new SL precision oncology targets					therapeutics	



Proven experience in drug discovery and development









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 PKMYT1 inhibitor into the clinic

Supported preclinical synergy hypothesis and patient selection approach from proprietary SNIPRx platform



POC, proof of concept; RP2D, recommended Phase 2 dose.

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

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Evolving broad trial program: sponsored and collaborative

Lunresertib Combination Therapy





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.

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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 lun+cam updates since ENA; 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



Recent MYTHIC publications



Successful approach to mitigating mechanismbased anemia while maintaining clinical benefit of L+C in broader MYTHIC clinical trial



LINK TO DATA at EORTC-NCI-AACR Symposium on Molecular Targets and Cancer Therapeutics, October 2024

Significant survival disparities, inherent chemotherapy resistance, and lack of treatment options for metastatic gynecologic cancer patients with our target biomarkers



LINK TO DATA at AACR's Ovarian Cancer Research Symposium, September 2024



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.



Lun+cam combination effective in gynecologic tumors

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



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 lun+cam anemia

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



	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 lunre+camo combination



Lun+FOLFIRI promising with excellent safety despite full dose FOLFIRI

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
- Prolonged PFS observed well beyond FOLFIRI benchmark in both FOLFIRI-naïve and experienced patients







Lun+FOLFIRI: efficacy profile warrants Phase 2 investigation

Prolonged clinical benefit and robust anti-tumor activity observed, including in patients with prior irinotecan exposure



Data presented at ESMO GI, June 2024. ^aCBR defined as best overall response of CR or PR according to RECIST 1.1 criteria or duration of treatment ≥16 weeks (dashed line). ^bORR defined as best response of confirmed CR or PR, unconfirmed CR or PR, or tumor marker response according to RECIST v1.1 criteria. ^cctDNA MR was defined as a ≥50% decline in ctDNA (dashed line). For DOT and tumor reduction data as of 6June24 and represent the efficacy evaluable population (≥1 post-baseline tumor assessment; n=33). ctDNA MR data as of 07May2024 using the Tempus xF+ liquid biopsy panel. Patients with no variants detected at baseline were deemed as non-monitorable for this analysis (n=7). BL, baseline; CBR, clinical benefit rate; CRC, colorectal cancer; DOT, duration of treatment; iri, irinotecan; MRR, molecular response rate;
 mVAFR, mean variant allele frequency rate.



Lunre+Debio 0123 1st clinical trial inhibiting PKMYT1 + 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...

...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.



Benada et al., NAR Cancer, 2023.

Platinum-resistant ovarian cancer lun+cam market opportunity



Unmet need remains significant for platinum-resistant patients



Note: Majority of ovarian cancer patients treated in MYTHIC as of ENA 2023 data cut-off were post-bevacizumab.

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 lun+cam cancer market opportunity

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



Single Agent Chemo^{1,2} (non biomarkerdefined, excludes endometroid)

15 - 16% ORR **3.8 - 4.0 mo. PFS** 12.0 - 12.3 mo. OS

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

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1 Lenvatinib/Pembrolizumab vs Single Agent Chemo (KEYNOTE-775); Source: Lenvatinib FDA Label. 2 Ixabepilone vs Paclitaxel or Doxorubicin; McMeekin S. Gynecologic Oncology 2015 https://doi.org/10.1016/j.ygyno.2015.04.026. 3 Eligible Patients in US and EU4/UK Based On Company Estimates from TCGA and GENIE, 2L+ (2L - 5L).

Metastatic CRC is a large market opportunity for lun+FOLFIRI

Unmet need in 2L+ oxaliplatin-treated mCRC patients



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)

Camonsertib:

Potential best-in-class ATR inhibitor X

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

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**





NSCLC, non-small cell lung cancer.

Potential across significant additional patient populations



Top 10 Tumor Types* with Highest

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





18.7%

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

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Promising Camonsertib mPFS in NSCLC



* mPFS of 4.6 months reported for both primary resistance and acquired resistance cohorts in the biomarker non-matched group, as reported in Besse, B. et al. Biomarker-directed targeted therapy plus durvalumab in advanced non-small-cell lung cancer: a phase 2 umbrella trial. Nature Medicine. 13 February 2024 (HUDSON trial).

Camonsertib NSCLC market opportunity

Significant unmet need for non-biomarker driven NSCLC patients



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





RP-1664

First-in-class, oral PLK4 inhibitor Highly potent, selective and bioavailable PLK4

and bioavailable PLK4 inhibitor synthetically lethal with TRIM37 gain of function Strong, dose-dependent anti-tumor activity observed as monotherapy across preclinical models

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



*Based on estimated number of pts available for 1st line treatment in the advanced setting (CancerMPact®, Patient Metrics, 2022; accessed 8/18/23) and lesion prevalence (TCGA; GENIE-Neuroblastoma Only).¹ Represents only gene amplification for high risk Neuroblastoma; ² Non-Squamous subtype of Non-Small Cell Lung Cancer only; ³ Squamous subtype of Non-Small Cell Lung Cancer only; ³ Squamous subtype of Non-Small Cell Lung Cancer only

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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





	Key Parameter	RP-1664
	PLK4 Enzyme IC ₅₀	1 nM
O	PLK4 cell binding IC ₅₀	3 nM
vitr	Cell proliferation in MCF7 / T47D (TRIM37 amp) EC ₅₀	51 / 17 nM
Ē	Cell-base selectivity vs AurA, AurB	>2000-fold
	Kinome screen at 90x PLK4 IC ₅₀	8/280 kinases >50% inh
	Human Honatocyto Clearance (ul /min/106 colls)	2 2
ME		2.2
ADN	Rat PK (%F, $t_{1/2}$)	28%, 4h
	Monkey PK (%F, t _{1/2})	96%, 9h

- 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



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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

Potential best-inclass Polθ ATPase inhibitor

FPI in 2H 2024



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

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



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





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





- 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



Multiple potential market opportunities for near term milestones





1 Eligible Patients Based On Company Estimates from TCGA and GENIE; Source: Drug Treated patients from CernerEnviza/CancerMpact 46

2H 2024	2025
RP-3467 Ph1 clinical trial initiation	Initiate first pivotal trial for lun+cam in 2025
Lun+cam expansion cohort data in ovarian and endometrial in Q4	Lunresertib + Debio 0123 combination data
	Camonsertib monotherapy data in NSCLC



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- Lunresertib: First-in-class oral PKMYT1i
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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)

Multiple clinical catalysts in 2024 and 2025

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

Strong balance sheet

- Cash and investments of ~\$180M¹ fund operations into second half of 2026
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