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Efficacy, molecular activity and safety from the pivotal PYNNACLE Phase 2 trial assessing rezatapopt, a p53 reactivator, in *TP53* Y220C-positive advanced solid tumors

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Rezatapopt is an investigational drug that has not been approved by any health authority

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Disclosure information

Alison M. Schram

I have the following relevant financial relationships to disclose:

- Advisory board: Relay Therapeutics, Mersana, Merus, Partner Therapeutics, PMV Pharmaceuticals, Inc., Schrödinger, Repare Therapeutics, Revolution Medicines, Endeavor Biomedicines, Day One Biopharmaceuticals, Transcode Therapeutics, Guardant, Boehringer Ingelheim
- Advisory role: Merus, Pfizer, PMV Pharmaceuticals, Inc., Schrödinger, Repare Therapeutics, Relay Therapeutics
- Research funding to institution: AstraZeneca, ArQule, BeiGene/SpringWorks, Black Diamond Therapeutics, Boehringer Ingelheim, Debiopharm, Elevar Therapeutics, Elevation Oncology, Kura, Lilly, Merus, Northern Biologics, Partner Therapeutics, Pfizer, Pheon Therapeutics, PMV Pharmaceuticals, Inc., Relay Therapeutics, Repare Therapeutics, Revolution Medicines, Surface Oncology

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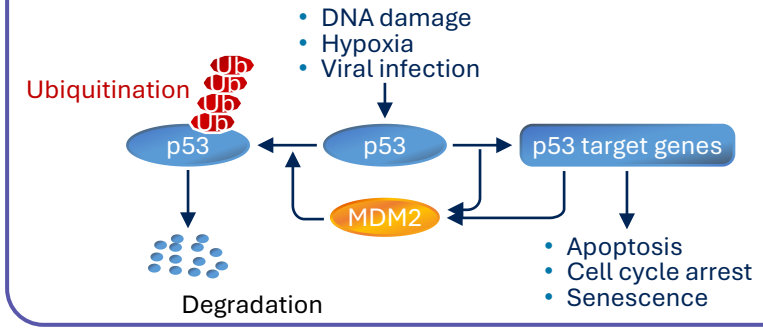
Full author disclosures can be found at the QR code.

Targeting p53 with rezatapopt, a selective p53 Y220C reactivator

p53 is a key player in the body's defense against cancer



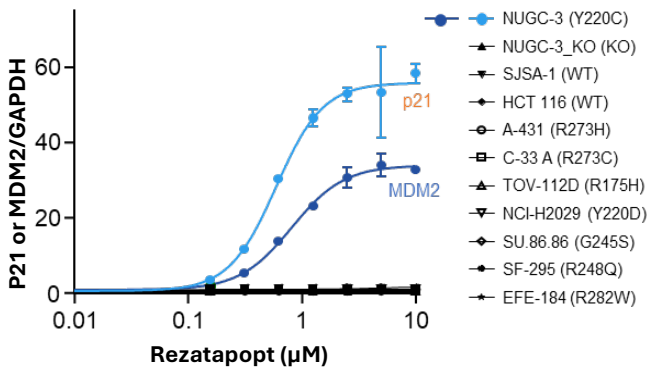
- *TP53*: most frequently mutated gene across all cancers[1–3]
- p53: binds DNA; key role in cell cycle arrest, DNA repair and apoptosis[2–5]
- *TP53* mutations: inactivation of p53, contributing to oncogenesis[2–5]
- *TP53* Y220C: in ~1.2% of solid tumors, ~3.2% of ovarian cancer and ~3.7% of HGSOC[6]
- *TP53* Y220C: destabilizes p53, resulting in loss of tumor suppressor function[1,7,8]

Rezatapopt is a selective p53 Y220C reactivator

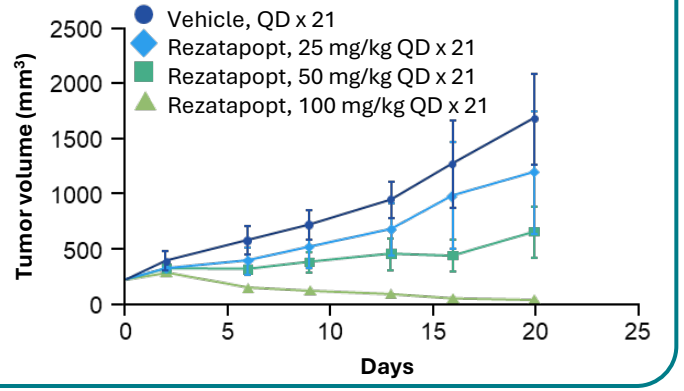
- Rezatapopt: investigational, first-in-class, selective p53 reactivator specific to the *TP53* Y220C mutation[7–9]
- Rezatapopt: stabilizes mutated Y220C-p53 in wild-type p53 conformation by binding to the pocket created in the protein, restoring p53 tumor suppressor activity[7–9]



p53 transcriptional activity restored[10]



Tumor regression in vivo[10]



HGSOC, high-grade serous ovarian cancer; KO, knockout; MDM2, mouse double minute 2 homolog; QD, once daily; UB, ubiquitin; WT, wild-type.

1. Baugh EH, et al. *Cell Death Differ*. 2018;25:154–160; 2. Kastenhuber ER, et al. *Cell*. 2017;170:1062–1078; 3. Levine AJ. *Nat Rev Cancer*. 2020;20:471–480; 4. Hassin O, et al. *Nat Rev Drug Discov*. 2023;22:127–144; 5. Chillemi G, et al. *Cold Spring Harb Perspect Med*. 2017;7:a028308; 6. FoundationInsights™. A proprietary database (FoundationCore® Q1-2026) used under license with review and approval from Foundation Medicine®. Available at: Biopharma Services Overview | Foundation Medicine. Deduplicated data accessed May 2026; 7. Vu BT, et al. *ACS Med Chem Lett*. 2025;16:34–39; 8. Puzio-Kuter AM, et al. *Cancer Discov*. 2025;15:1159–1179; 9. PYNNACLE Study. Available at: https://pynnaclestudy.com/. Accessed November 2026; 10. Dumble M, et al. *Cancer Res*. 2021;81(13_Suppl):Abstract LB006.

PYNNACLE: Pivotal Phase 2 study design assessing rezatapopt

PYNNACLE (NCT04585750): Pivotal Phase 2, global, multi-cohort clinical trial assessing rezatapopt in patients with solid tumors with a *TP53* Y220C mutation and *KRAS* wild-type[1,2]

Key eligibility criteria

- Aged ≥12 years[a,b]
- Locally advanced or metastatic solid tumors, excluding primary CNS tumors
- Documented somatic *TP53* Y220C mutation and *KRAS* wild-type[c]
- Prior standard therapy or ineligible for SoC

Basket N≈200

Patient cohorts defined by tumor type

Rezatapopt 2000 mg QD with food

- Cohort 1: Ovarian cancer (platinum-resistant/refractory)
- Cohort 2: Lung cancer
- Cohort 3: Breast cancer
- Cohort 4: Endometrial cancer
- Cohort 5: All other solid tumors

Endpoints

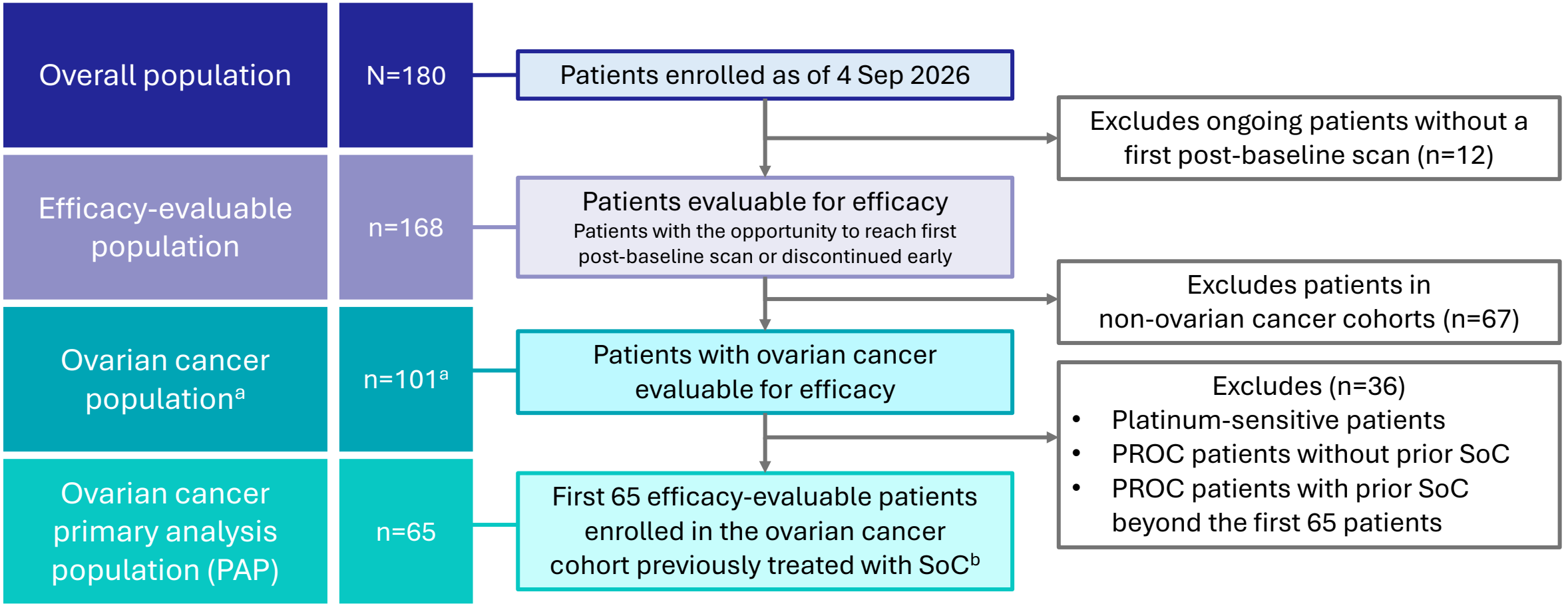
Primary: ORR (RECIST v1.1) per BICR across cohorts and in the ovarian cancer cohort

Key secondary: ORR per investigator, TTR, DoR, DCR, PFS per BICR and investigator, OS, safety

Exploratory: Changes in ctDNA dynamics and mutational/genomic profile (e.g., *TP53* Y220C, other *TP53* mutations, non-*TP53* mutations, known oncogenic drivers) before and after treatment

[a] Adults aged ≥18 years for all global sites except Singapore (must be ≥21 years of age) and the Republic of Korea (must be ≥19 years of age). [b] Adolescents aged 12–17 years if weighing ≥40 kg (in Australia, the Republic of Korea [12–18 years of age] and the USA only). [c] Test results confirmed from tissue analysis; exception: ovarian cancer patients with a known *KRAS* single nucleotide variant mutation are permitted. BICR, blinded independent central review; CNS, central nervous system; ctDNA, circulating tumor DNA; DCR, disease control rate; DoR, duration of response; ORR, overall response rate; OS, overall survival; PFS, progression-free survival; QD, once daily; RECIST, Response Evaluation Criteria in Solid Tumors; SoC, standard of care; TTR, time to response. 1. ClinicalTrials.gov. Available at: https://clinicaltrials.gov/study/NCT04585750. Accessed November 2026; 2. PYNNACLE Study. Available at: https://pynnaclestudy.com/. Accessed November 2026.

Analysis populations

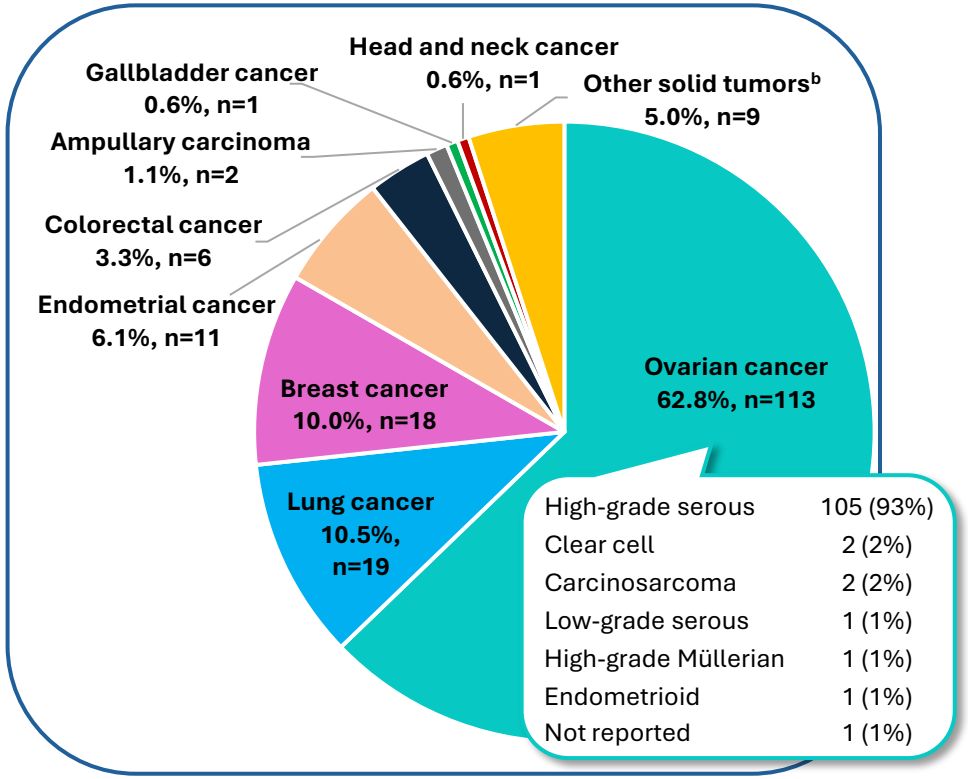


[a] In total, 113 patients with ovarian cancer were enrolled; 101 patients with ovarian cancer were efficacy evaluable (includes patients with first post-baseline tumor assessment and patients who discontinued early). [b] As defined by the protocol. Seamless adaptive clinical trial design: patient enrollment continues beyond the initial 65 patients with ovarian cancer for the primary analysis to collect supportive data. PROC, platinum-resistant/refractory ovarian cancer; SoC, standard of care.

Patient demographics and baseline characteristics

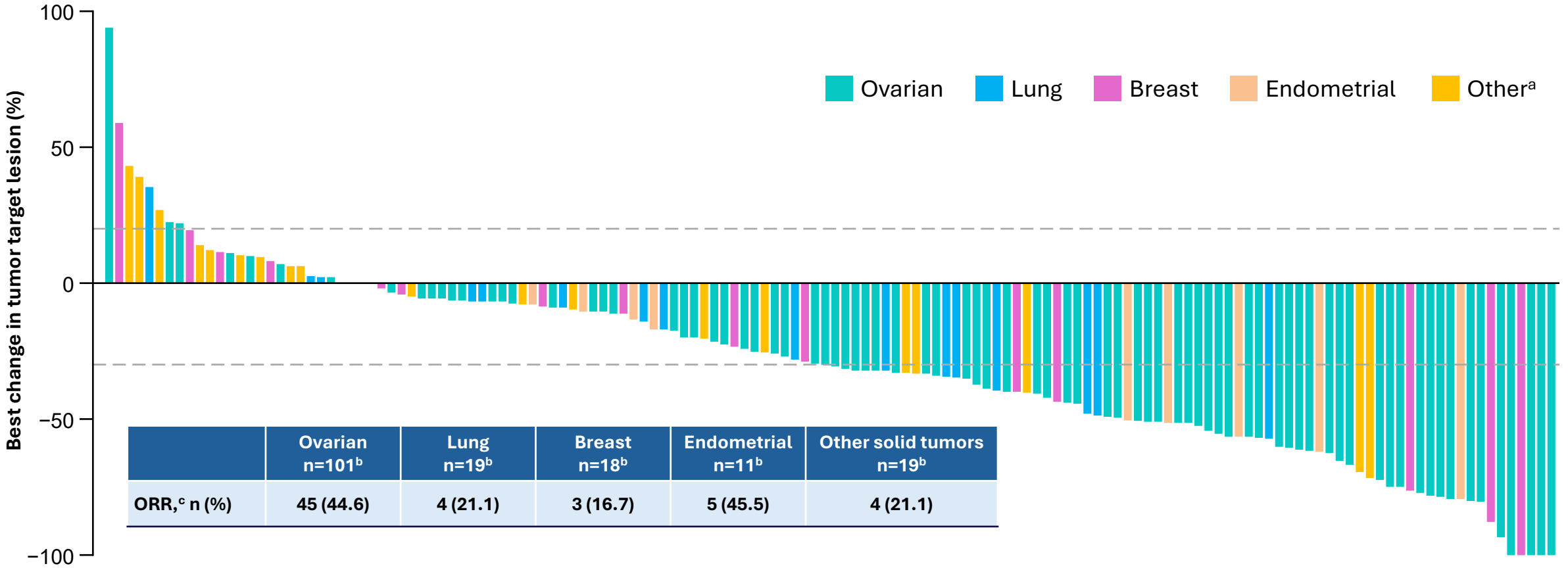
	Overall population N=180	Ovarian cancer n=113	Ovarian cancer PAP n=65
Median age, years (min–max)	65 (33–91)	66 (33–91)	65 (46–84)
ECOG status,[a] n (%)	n=179		
0/1	76 (42)/103 (58)	50 (44)/63 (56)	33 (51)/32 (49)
Platinum status, n (%)			
Sensitive	N/A	3 (3)	N/A
Resistant	N/A	71 (63)	44 (68)
Refractory	N/A	39 (35)	21 (32)
Primary platinum refractory	N/A	9 (8)	5 (8)
Prior lines of systemic therapy, n (%)	n=168		
1/2	24 (14)/37 (22)	9 (8)/21 (19)	1 (2)/6 (9)
≥3	102 (61)	83 (73)	58 (89)
Median (min–max)	3 (0–9)	4 (1–9)	4 (1–9)
Prior therapies, n (%)			
Bevacizumab	N/A	93 (82)	59 (91)
PARPi	N/A	59 (52)	43 (66)
Mirvetuximab soravtansine	N/A	26 (23)	17 (26)
HRD+, n (%)	N/A	n=105 41 (39)	n=63 27 (43)
Somatic *BRCA* 1/2 mutation,[a] n (%)	N/A	13 (12)	5 (8)
FOLR1+,[a] n (%)	N/A	n=95 46 (48)	n=56 25 (45)
PD-L1 CPS+,[a] n (%)	N/A	n=72 28 (39)	n=49 18 (37)
***CCNE1* amplification,[a] n (%)**	N/A	n=90 17 (19)	n=57 12 (21)

Patients were heavily pretreated across a broad spectrum of tumor types



Data cutoff: 4 Sep 2026. [a] Status was reported post data cutoff. [b] Includes gastric cancer (n=2), sarcoma (n=2), small intestine cancer (n=1), hepatocellular carcinoma (n=1), pancreatic cancer (n=1), thymic carcinoma (n=1) and esophagueal carcinoma (n=1).
BRCA, breast cancer gene; *CCNE1*, cyclin E1; ECOG, Eastern Cooperative Oncology Group; FOLR1, folate receptor 1; HRD, homologous recombination deficiency; PAP, primary analysis population; PARPi, poly (ADP-ribose) polymerase inhibitor; PD-L1 CPS, programmed death-ligand 1 combined positive score.

Deep tumor shrinkage observed across tumor types following treatment with rezatapopt – overall population



	Ovarian n=101[b]	Lung n=19[b]	Breast n=18[b]	Endometrial n=11[b]	Other solid tumors n=19[b]
ORR,[c] n (%)	45 (44.6)	4 (21.1)	3 (16.7)	5 (45.5)	4 (21.1)

Data cutoff: 4 Sep 2026. n=144 patients with a post-baseline assessment. [a] Responses observed in head and neck cancer, colorectal cancer, gallbladder cancer and ampullary carcinoma. [b] Efficacy-evaluable population (n=168) includes patients with first post-baseline tumor assessment and who discontinued early. [c] ORR is calculated using CR, PR and uPR. CR, complete response; ORR, overall response rate; PR, partial response; uPR, unconfirmed partial response.

Clinically meaningful response rates observed across tumor types and ovarian cancer populations

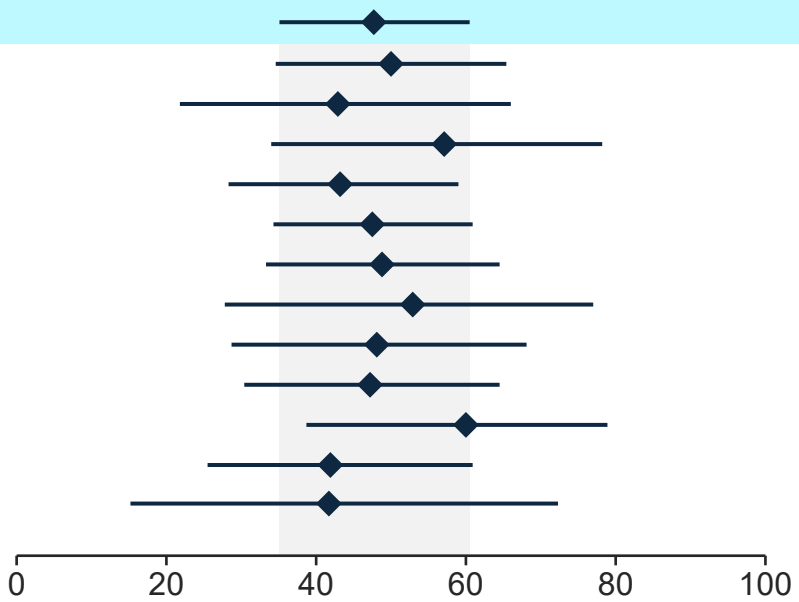
Investigator-assessed ORR per RECIST v1.1			
Patients, n (%)	**Overall population n=168[a]**	**Ovarian cancer n=101[a]**	**Ovarian cancer PAP n=65[a]**
ORR,[b]	**61 (36.3)**	**45 (44.6)**	**31 (47.7)**
(95% CI)	29.0-44.1	34.7-54.8	35.1-60.5
CR	4 (2.4)	4 (4.0)	1 (1.5)
PR	53 (31.5)	37[c] (36.6)	30 (46.2)
uPR	4 (2.4)	4 (4.0)	–
SD	69 (41.1)	36 (35.6)	23 (35.4)
Disease control rate[d]	**130 (77.4)**	**81 (80.2)**	**54 (83.1)**

Data cutoff: 4 Sep 2026. [a] Efficacy-evaluable population includes patients with first post-baseline tumor assessment and patients who discontinued early. [b] ORR is calculated using CR, PR and uPR. [c] Post data cutoff 1 patient with a PR subsequently converted to a uCR. [d] Disease control rate is calculated using CR, PR, uPR and SD. CI, confidence interval; CR, complete response; ORR, overall response rate; PAP, primary analysis population; PR, partial response; SD, stable disease; uCR, unconfirmed complete response; uPR, unconfirmed partial response.

Consistent response rates observed across subgroups in the ovarian cancer PAP

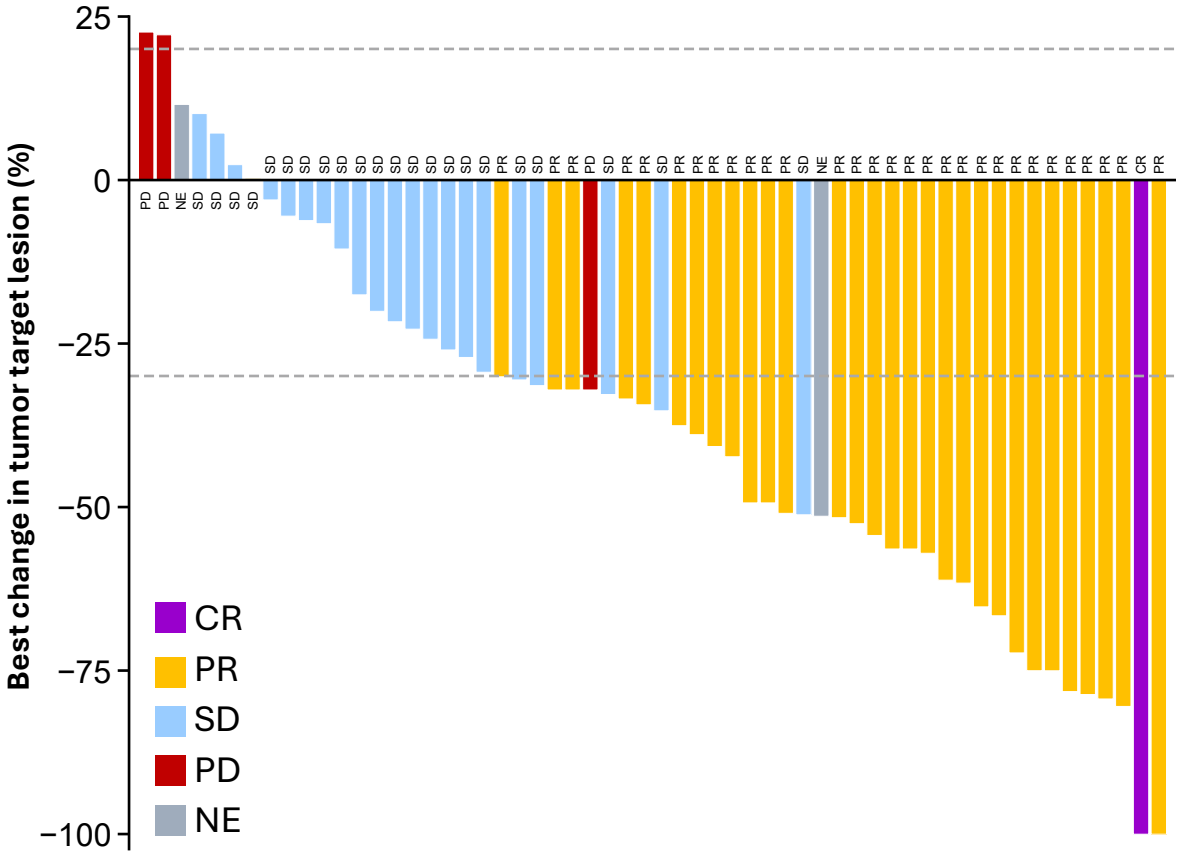
Investigator-assessed ORR[a] per RECIST v1.1 by subgroup

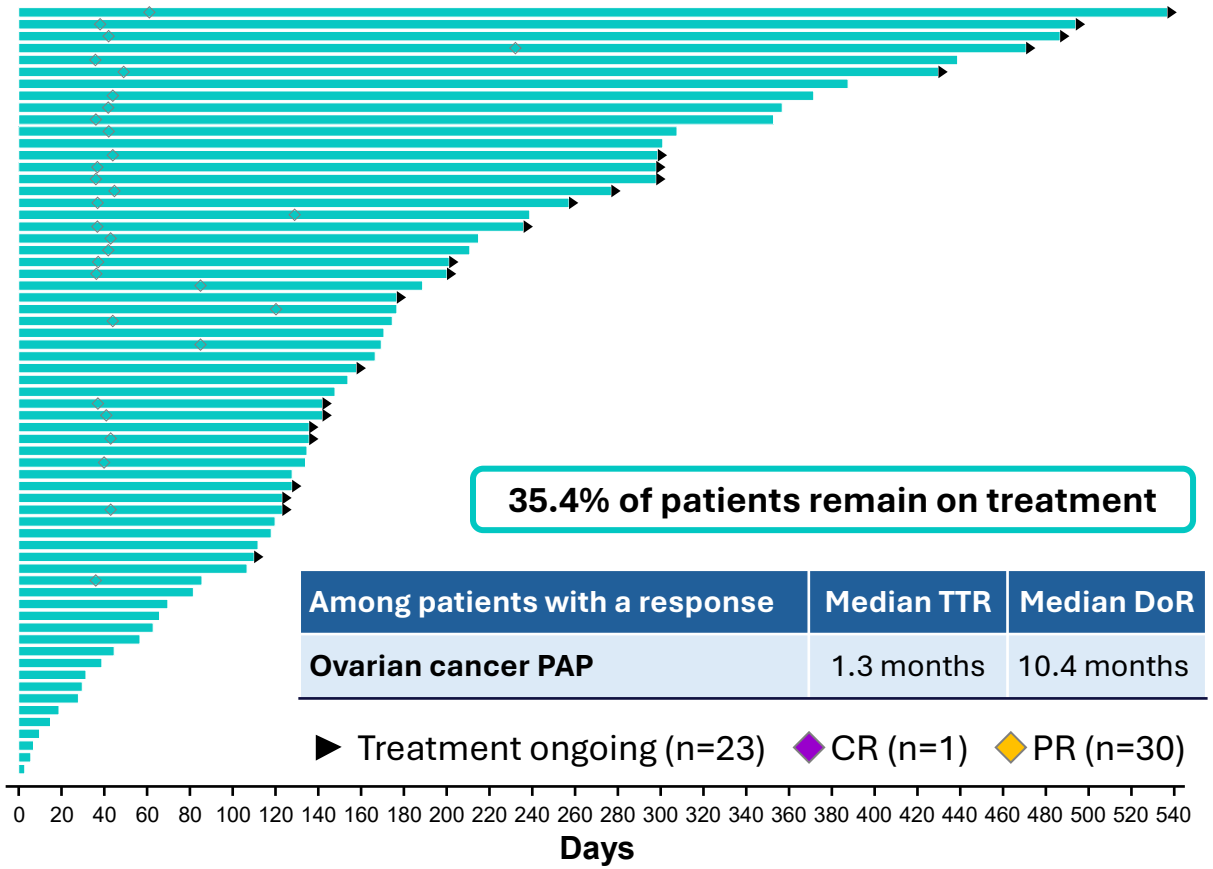
Subgroup	Responders/n	ORR, % (95% CI)
Ovarian cancer PAP[b]	**31/65**	**47.7 (35.1–60.5)**
Platinum resistant	22/44	50.0 (34.6–65.4)
Platinum refractory	9/21	42.9 (21.8–66.0)
1–3 prior lines of therapy	12/21	57.1 (34.0–78.2)
>3 prior lines of therapy	19/44	43.2 (28.3–59.0)
Prior bevacizumab	28/59	47.5 (34.3–60.9)
Prior PARPi	21/43	48.8 (33.3–64.5)
Prior mirvetuximab soravtansine	9/17	52.9 (27.8–77.0)
HRD+	13/27	48.1 (28.7–68.1)
HRD−	17/36	47.2 (30.4–64.5)
FOLR1+	15/25	60.0 (38.7–78.9)
FOLR1−	13/31	41.9 (25.5–60.9)
CCNE1 amplification	5/12	41.7 (15.2–72.3)



Exploratory subgroup analyses – data cutoff: 4 Sep 2026. [a] ORR is calculated using CR, PR and uPR. [b] Same efficacy-evaluable population as the previous slide – efficacy-evaluable population includes patients with first post-baseline tumor assessment and patients who discontinued early. *CCNE1*, cyclin E1; CI, confidence interval; CR, complete response; FOLR1, folate receptor 1; HRD, homologous recombination deficiency; ORR, overall response rate; PAP, primary analysis population; PARPi, poly (ADP-ribose) polymerase inhibitor; PR, partial response; uPR, unconfirmed partial response.

Deep tumor shrinkage with rapid and durable responses in the ovarian cancer PAP





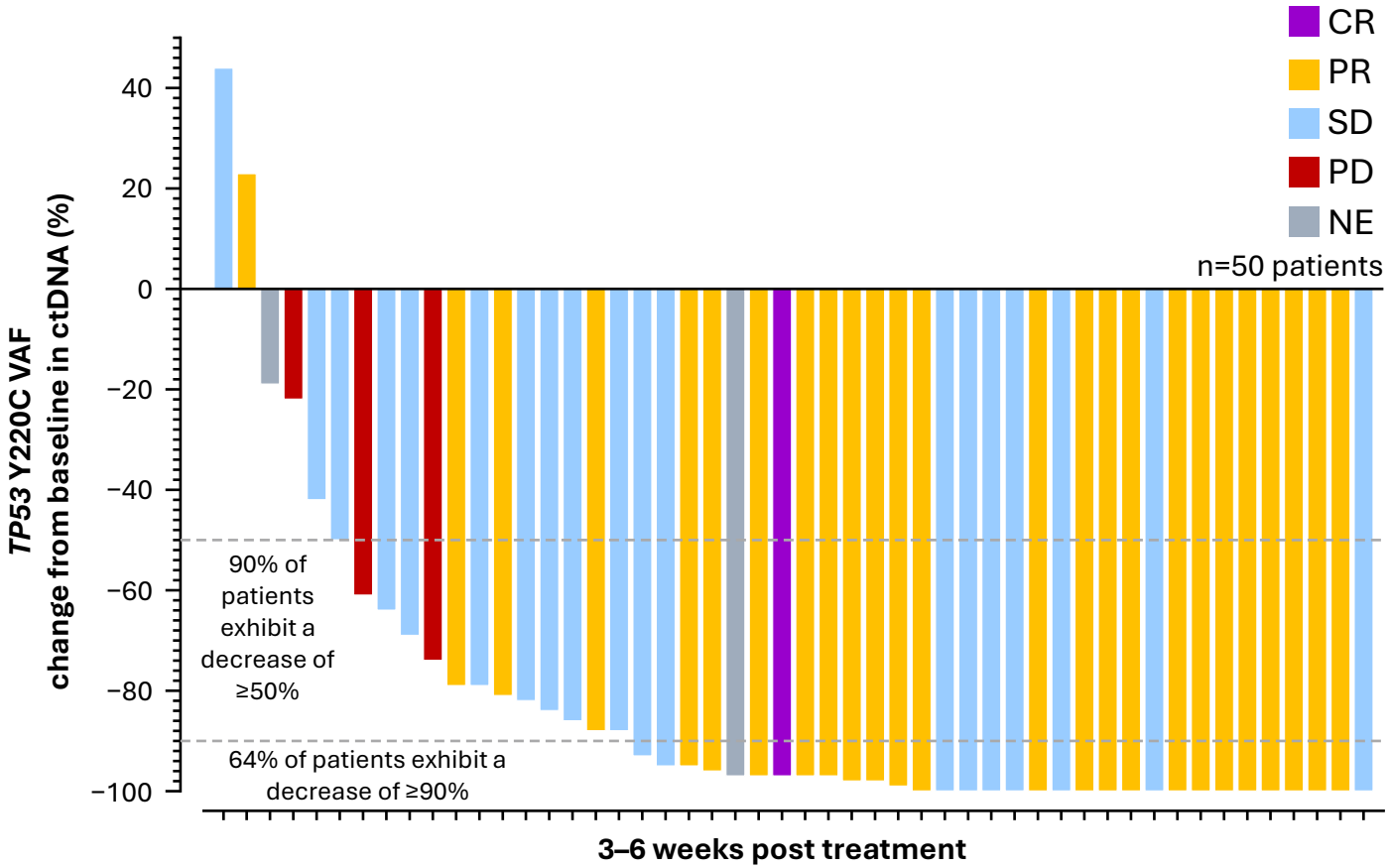
35.4% of patients remain on treatment

Among patients with a response	Median TTR	Median DoR
Ovarian cancer PAP	1.3 months	10.4 months

▶ Treatment ongoing (n=23) ◆ CR (n=1) ◆ PR (n=30)

Data cutoff: 4 Sep 2026. [a] n=58 patients with a post-baseline assessment.
CR, complete response; DoR, duration of response; NE, non-evaluable; PAP, primary analysis population; PD, progressive disease; PR, partial response; SD, stable disease; TTR, time to response.

Rezatapopt on-target activity in the ovarian cancer PAP was supported by decreases in ctDNA *TP53* Y220C VAF

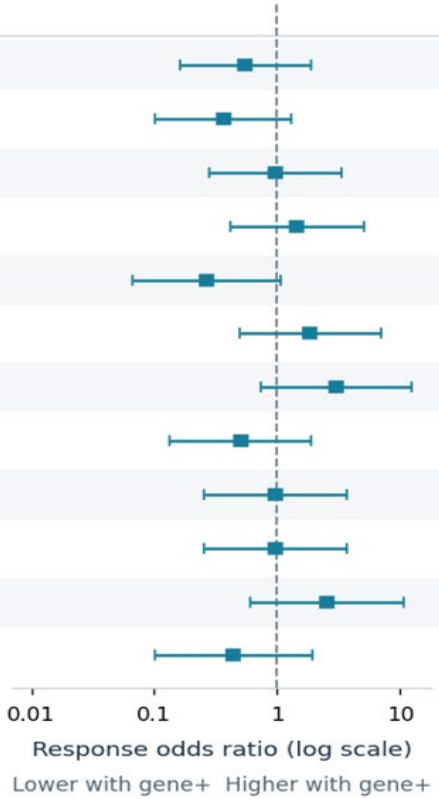


- 50 patients in the ovarian cancer PAP had available ctDNA *TP53* Y220C VAF data at baseline and following treatment (3–6 weeks)
- All patients except one experiencing a response had a reduction in *TP53* Y220C VAF
- 90% of patients achieved ≥50% reduction in ctDNA *TP53* Y220C VAF
- 64% of patients achieved ≥90% reduction in ctDNA *TP53* Y220C VAF

Data cutoff: 4 Sep 2026. CR, complete response; ctDNA, circulating tumor DNA; NE, non-evaluable; PAP, primary analysis population; PD, progressive disease; PR, partial response; SD, stable disease; VAF, variant allele frequency.

In ovarian cancer, responses occurred regardless of co-occurring genomic alterations at baseline

Gene[a]	Response[a]	OR (95% CI)[b]	P value[c]	q[d]
PRKCI	5/13	0.55 (0.16–1.87)	0.375	0.643
TERC	4/13	0.37 (0.10–1.32)	0.137	0.643
CCNE1	6/12	0.97 (0.28–3.33)	1.000	1.000
MYC	7/12	1.45 (0.41–5.04)	0.754	1.000
NF1	3/12	0.27 (0.07–1.08)	0.065	0.643
MAP3K1	7/11	1.86 (0.50–6.99)	0.516	0.774
NOTCH3	8/11	3.02 (0.73–12.44)	0.191	0.643
ATR	4/11	0.50 (0.13–1.89)	0.346	0.643
ARID1A	5/10	0.97 (0.26–3.67)	1.000	1.000
BCL2L1	5/10	0.97 (0.26–3.67)	1.000	1.000
ROS1	7/10	2.56 (0.61–10.77)	0.309	0.643
PIK3CA	3/9	0.44 (0.10–1.92)	0.309	0.643

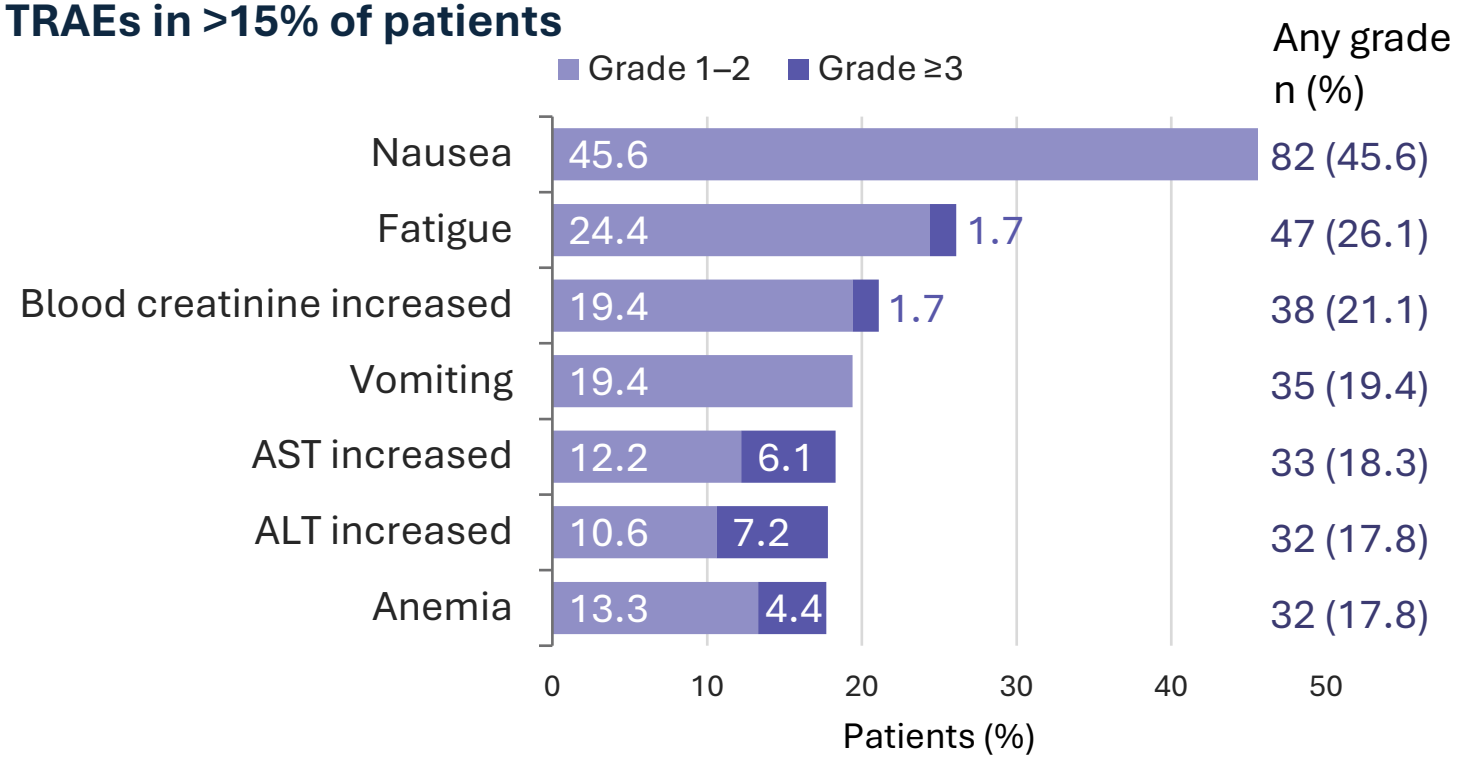


- 75 patients with ovarian cancer had tumor genomic profiling and response data available
- No association between co-occurring alterations at baseline and response
- Four patients had additional co-occurring *TP53* mutations
  - CR (n=1; two additional *TP53* mutations)
  - PR (n=1); SD (n=1); PD (n=1)

Data cutoff: 4 Sep 2026. [a] Responders/gene-positive patients; gene presence counted once per patient. [b] Bars: unadjusted approximate log-Wald 95% CIs. [c] Two-sided Fisher exact P. [d] Benjamini–Hochberg false discovery rate-adjusted P value (q) across 12 tests. *ARID1A,* AT-rich interaction domain 1A; *ATR*, ATR checkpoint kinase; *BCL2L1*, BCL2 like 1; *CCNE1*, cyclin E1; CI, confidence interval; CR, complete response; *MAP3K1*, mitogen-activated protein kinase kinase kinase 1; *MYC*, MYC proto-oncogene, bHLH transcription factor; *NF1*, neurofibromin 1; *NOTCH3,* Notch receptor 3; OR, odds ratio; PD, progressive disease; *PIK3CA*, phosphatidylinositol-4,5-bisphosphate 3-kinase catalytic subunit alpha; PR, partial response; *PRKCI,* protein kinase C iota; *ROS1*, ROS proto-oncogene 1, receptor tyrosine kinase; SD, stable disease; *TERC*, telomerase RNA component.

Rezatapopt displays a favorable safety and tolerability profile in the overall population

Total safety population (N=180)
TRAEs in >15% of patients



- TRAEs were mostly Grade 1/2
- Laboratory abnormalities were manageable/monitorable, with most cases being transient and reversible
- No Hy's law cases were observed
- Similar safety profile in patients with ovarian cancer
- 11 patients (6.1%) discontinued treatment due to TRAEs, including 7 (6.2%) with ovarian cancer

Data cutoff: 4 Sep 2026. Bars show the maximum CTCAE grade per patient; one patient experienced a Grade 4 TRAE of AST increased; no Grade 5 TRAEs were observed.
ALT, alanine aminotransferase; AST, aspartate aminotransferase; CTCAE, Common Terminology Criteria for Adverse Events; TRAE, treatment-related adverse event.

Case study: A heavily pretreated patient with primary platinum-refractory HGSOC achieved a partial response by Week 6

73-year-old white woman with primary platinum-refractory HGSOC

- HRD+, FOLR1−
- *CCNE1* and *FGFR1* amplification

Three lines of prior therapy

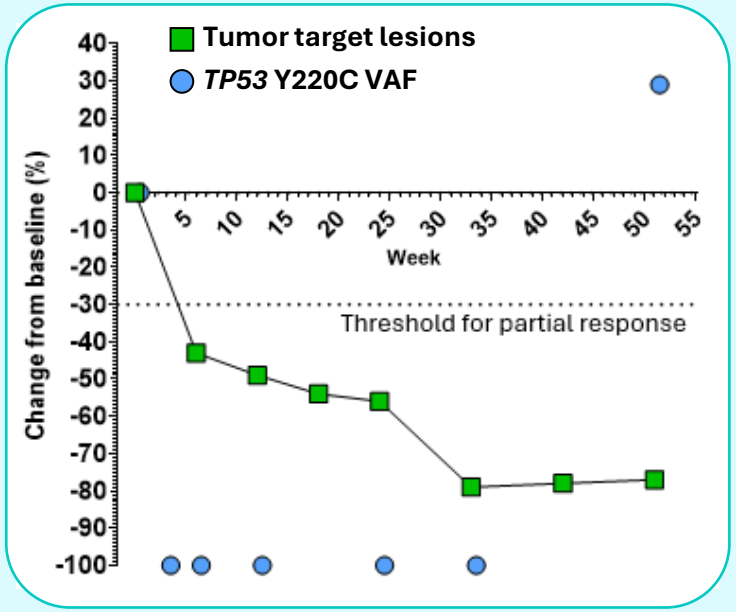
- Carboplatin/paclitaxel
- Pegylated liposomal doxorubicin
- Letrozole

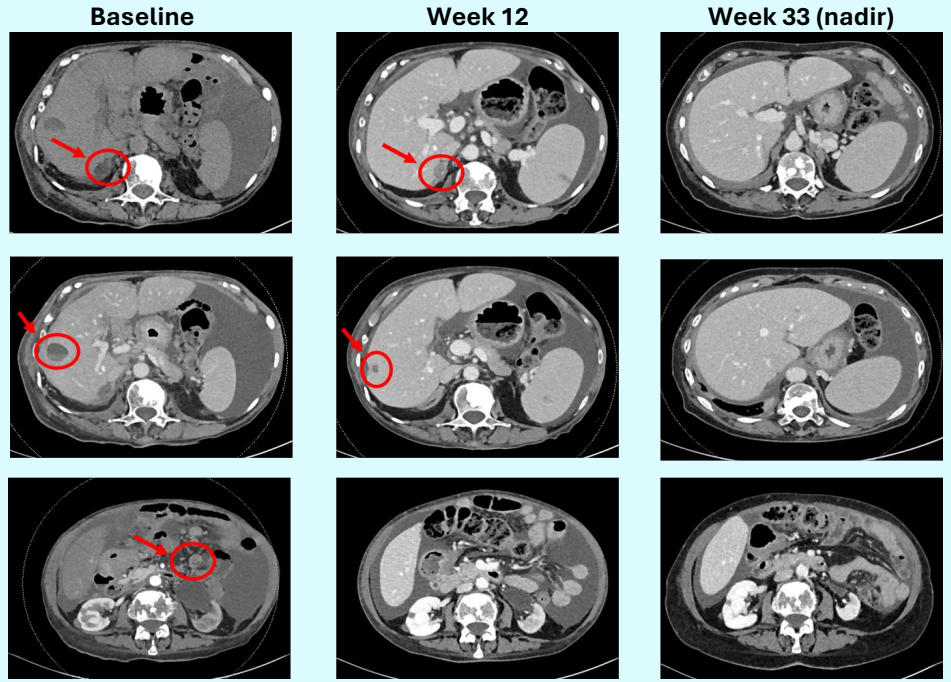
Rezatapopt 2000 mg QD

- No SAEs or AESIs
- One dose reduction to 1500 mg QD at Cycle 3 due to Grade 2 nausea
- Treatment duration: 14 months
- EOT: due to clinical progression

Response to rezatapopt

- Target tumor reduction (baseline: 107 mm) – Week 6: 61 mm (43%); Week 33 (nadir): 23 mm (78.5%)
- CA125 levels (baseline: 944 U/mL) – Week 6: 270 U/mL; Week 33: 108 U/mL; EOT: 379 U/mL
- DoR: 10.4 months; TTR: 1.6 months





Data cutoff: 4 Sep 2026. AESI, adverse event of special interest; *CCNE1*, cyclin E1; DoR, duration of response; EOT, end of treatment; *FGFR1*, fibroblast growth factor receptor 1; FOLR1, folate receptor 1; HGSOC, high-grade serous ovarian cancer; HRD, homologous recombination deficiency; QD, once daily; SAE, serious adverse event; TTR, time to response; VAF, variant allele frequency.

Case study provided courtesy of Anna Passarelli

PYNNACLE
NCT04585750

Conclusions

- Rezatapopt, a first-in-class, selective p53 Y220C reactivator, demonstrated efficacy and a manageable safety profile as an oral, chemotherapy-free monotherapy in heavily pretreated patients with *TP53* Y220C-positive solid tumors, including PROC
  - Overall population: ORR 36%
  - Ovarian cancer primary analysis population: ORR 48%
  - Mostly Grade 1/2 TRAEs; 6.1% discontinuation rate due to TRAEs
- Consistent response rates were seen across all ovarian cancer subgroups, regardless of co-occurring genomic alterations at baseline

To view the presentation please scan the QR code

ORR, overall response rate; PROC, platinum-resistant/refractory ovarian cancer; TRAE, treatment-related adverse event. https://clinicaltrials.gov/study/NCT04585750.

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- Investigators and research staff
- PPD, part of Thermo Fisher Scientific
- Foundation Medicine

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PYNNACLE Phase 2 clinical trial: Sites and Principal Investigators

USA
- Ikechukwu Akunyili: WellSpan York Cancer Center
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- Andrew L. Coveler: University of Washington, Fred Hutch Cancer Center
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