# Efficacy of RLY-2608, a mutant-selective PI3Kα inhibitor in patients with *PIK3CA*-mutant HR+/HER2- advanced breast cancer: ReDiscover trial

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#### INTRODUCTION

- Oncogenic *PIK3CA* mutations constitutively activate PI3Kα and drive ~40% of cases of HR+/HER2- BC<sup>1,2</sup>
- Approved therapies (alpelisib, inavolisib, everolimus, capivasertib) are non-selective PI3K pathway inhibitors with off-target toxicity that limits their tolerability (hyperglycemia, rash, diarrhea, stomatitis) and efficacy (mPFS of ~5.5–8 months with endocrine therapy)<sup>3-9</sup>
- RLY-2608 is the first pan-mutant-selective inhibitor designed to overcome these limitations
- RLY-2608 selectively targets mutant PI3Kα, via binding to a novel pocket, distinct from approved orthosteric inhibitors and emerging inhibitors that target only H1047R (Figure 1)10
- ReDiscover is a multi-arm, open-label, FIH study designed to evaluate RLY-2608 in patients with advanced solid tumors that have PIK3CA mutations present in blood and/or tumor per local assessment (Figure 2)11
- Previously we demonstrated robust and continuous target coverage and promising initial efficacy in phase 1
- Here we present the efficacy and safety of RLY-2608 + fulvestrant in patients with PIK3CA-mutant, HR+/HER2- BC treated
- Monotherapy and triplet (RLY-2608 + fulvestrant + ribociclib; RLY-2608 + fulvestrant + atirmociclib) arms are ongoing and will be presented separately

# Figure 1. RLY-2608 is the first pan-mutant-selective PI3Kα inhibitor LY-2608, a pan-mutant selective PI3Ka inhibitor, binds a novel allosteric site ve PI3Kα inhibitors lpelisib, inavolisib) bind the orthosteric (active Non-selectively inhibit wild type and mutant PI3Kα nerging H1047R-specific PI3K $\alpha$ inhibitors Non-selective downstream pathway inhibit apivasertib, everolimus) do not bind PI3Ko

#### Figure 2. ReDiscover study design



# **METHODS**

- 118 patients with advanced *PIK3CA*-mutant, HR+/HER2- BC received doublet therapy with RLY-2608 (100–1000 mg BID under fasting conditions) + fulvestrant in 28-day cycles: 64 patients received the RP2D across dose-escalation and -expansion phases (**Table 1**)
- Efficacy is presented for 52 patients without baseline PTEN or AKT1 E17K co-mutation who received the RP2D
- Key endpoints included determination of MTD and RP2D, safety and tolerability, PK/PD, and preliminary efficacy per RECIST 1.1
- Key inclusion criteria for the RLY-2608 doublet therapy arm included<sup>11</sup>:
- Histologically or cytologically confirmed HR+/HER2- unresectable or metastatic BC not amenable to curative therapy
- One or more documented primary oncogenic PIK3CA kinase or non-kinase mutation(s) in blood and/or tumor per local assessment
- Evaluable disease per RECIST 1.1 No prior PI3K/mTOR inhibitor
- ≥1 CDK4/6i in either the adjuvant and/or metastatic setting
- ≥1 anti-estrogen therapy

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≤1 line of chemotherapy in the metastatic setting

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# Figure 3. Potent and mutant-selective target inhibition at the recommended dose Mutant-selective inhibition with minima Rapid decline in mutant ctDNA at RP2D across mutation impact on glucose homeostasis at RP2D 600 mg BID (N=64) Non-kinase and C2D1. Notably, 12 patients exhibited Grade 3<sup>†</sup> (250–500 mg/dL) 5 patients without *PTEN/AKT1* E17K co-alterations have detectable ESR1 mutations at baseline. Paired ctDNA samples were collected at C1D1 Grade 1 (115–160 mg/dL) and C2D1. Notably, 10 patients exhibited multiple Time since first dose <sup>†</sup>Per CTCAE version 4.0.

### RESULTS

#### Table 1. Patient demographics and baseline characteristics

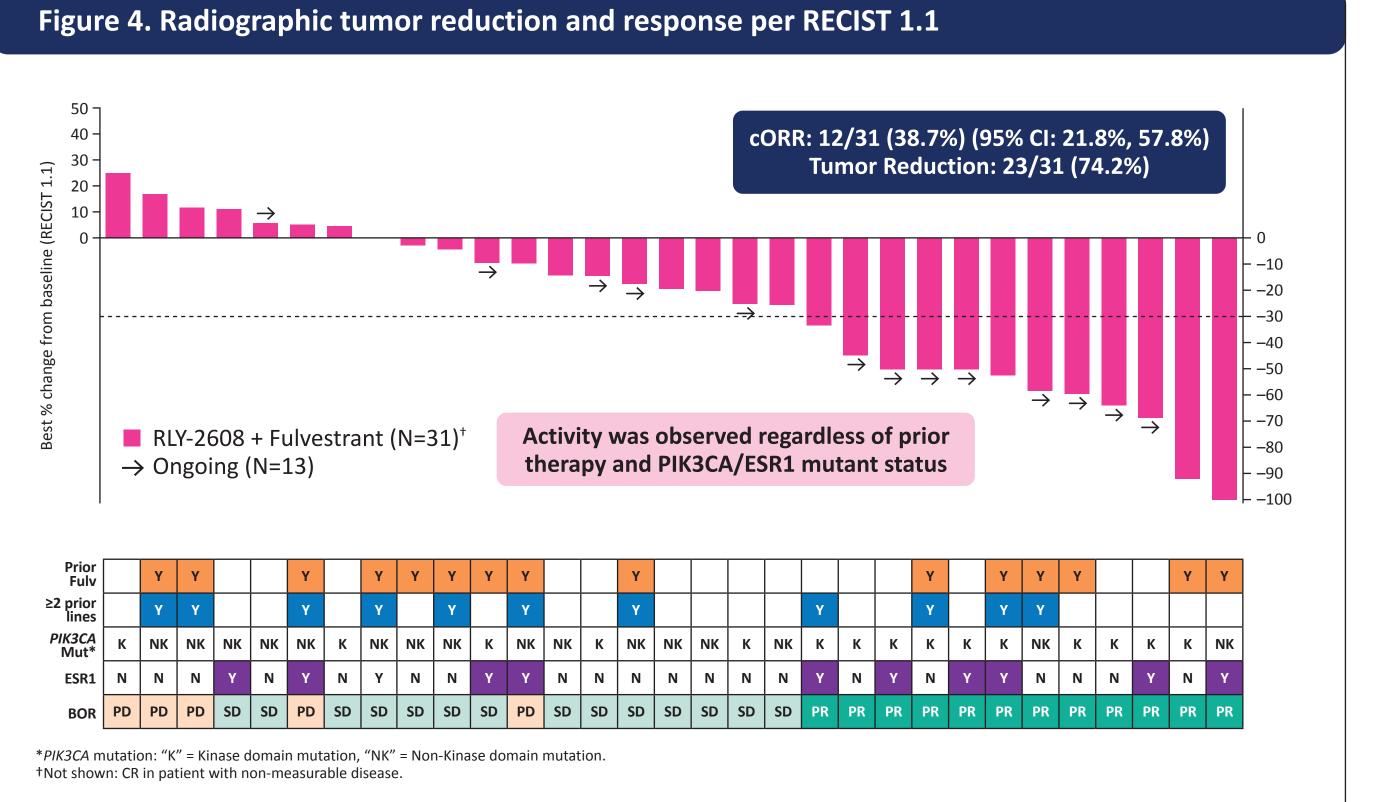
	RLY-2608 +			
	All Patients (N=118)	600 mg BID (RP2D, N=64)		
Age, Median (Range), Years ECOG, 0 / 1, n (%) Local <i>PIK3CA</i> Baseline Results	59.0 (34, 85) 69 (58.5) / 49 (41.5)	59.0 (34, 80) 38 (59.4) / 26 (40.6)	600 mg BID cohort (N=64) Still on treatment: 28 (43.8%) Discontinued: 36 (56.3%)	
Kinase Mutation, n (%) Non-Kinase Mutations, n (%) BMI >30 or HbA1c >5.7%, n (%) Measurable Disease, n (%) Patients with Visceral Metastases, n (%)* Prior Lines of Therapy in Advanced Setting	57 (48.3) 61 (51.7) 44 (37.3) 83 (70.3) 75 (63.6)	31 (48.4) 33 (51.6) 22 (34.4) 42 (65.6) 38 (59.4)	<ul> <li>Progressive disease: 30 (46.9%)</li> <li>Patient withdrew consent: 3 (4.7%)</li> <li>Due to adverse event: 2 (3.1%)</li> </ul>	
1, n (%) 2+, n (%)	62 (52.5) 56 (47.5)	38 (59.4) 26 (40.6)	<ul><li>Physician decision: 1 (1.6%)</li></ul>	
Prior Therapies in Advanced Setting				
CDK4/6 inhibitor, n (%) <sup>†</sup> Fulvestrant or Novel SERD, n (%) Chemo / ADC, n (%)  ESR1 Mutation (Central Read), n (%) <sup>‡</sup>	118 (100.0) 66 (55.9) 30 (25.4) 40 (35.4)	64 (100.0) 33 (51.6) 16 (25.0) 18 (28.6)		
PTEN or AKT1 E17K Mutation, n (%)	25 (21.2)	12 (18.8)		

#### Table 2. RLY-2608 has a favorable safety profile consistent with mutant-selective PI3Kα inhibition

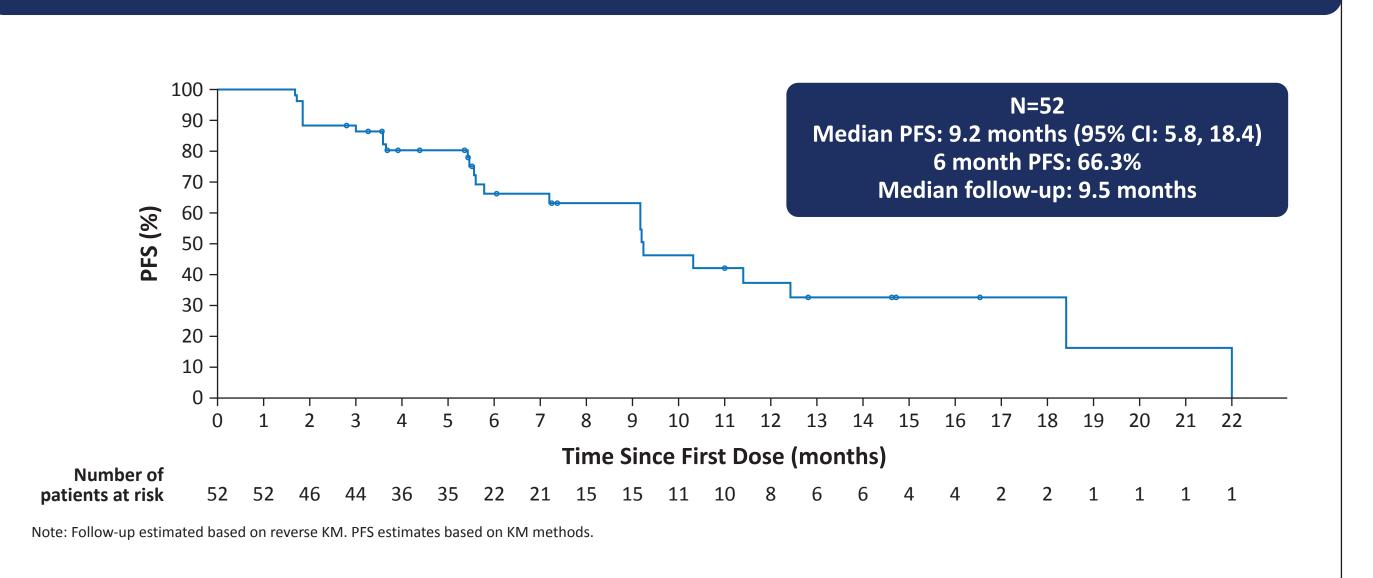
		All Patients (N=118)		600mg BID (RP2D, N=64)		
		All Grades	Grade 3	All Grades	Grade 3	27% Grade 1
Any TRAE, %		92.4	25.4	93.8	31.3	hyperglycemia
TRAEs ≥15% of 600 mg BID	Hyperglycemia*	42.4	2.5	46.9	3.1	(no intervention required)  No Grade 4/5 TRAEs observed
	Nausea	41.5	0.8	50.0	1.6	
	Fatigue*	40.7	8.5	35.9	9.4	
	Creatinine Increased	34.7	0.8	34.4	1.6	
	Diarrhea	30.5	1.7	35.9	3.1	
	Decreased Appetite	16.9	0	20.3	0	
	Headache	15.3	0.8	20.3	0	
	Hypokalemia*	15.3	1.7	17.2	1.6	
	Vomiting	12.7	0	15.6	0	
Other select TRAEs	Rash*	11.9	0.8	10.9	1.6	
	Stomatitis	3.4	0.8	4.7	0	
	No	Grade 4–5 TRAE	<b>S</b>			

#### **Abbreviations**

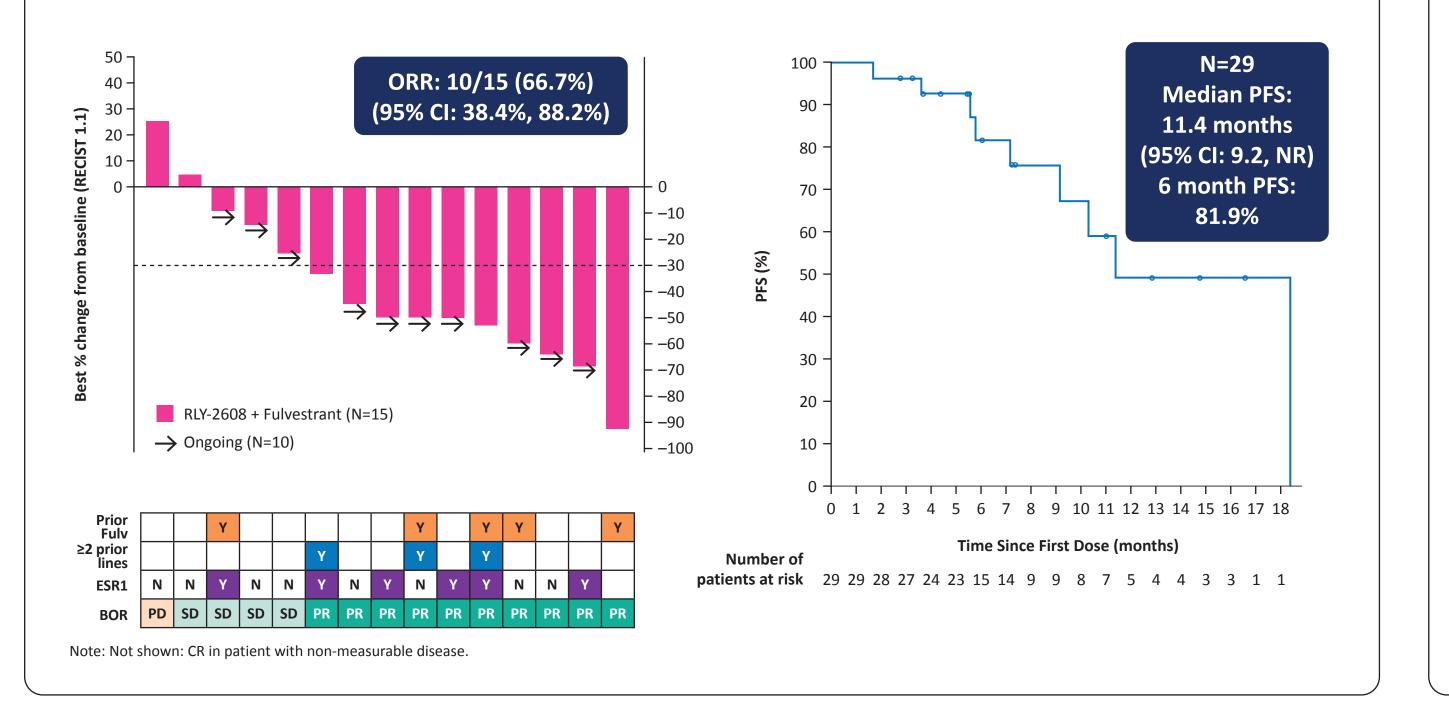
2L, second-line; ADC, antibody-drug conjugate; AE, adverse event; BC, breast cancer; BID, twice per day; BMI, body mass index; C2D1, cycle 2 day 1; CBR, clinical benefit rate; CDK4/6i, cyclin-dependent kinase 4/6 inhibitor; CR, complete response; ctDNA, circulating tumor DNA; DCR, disease control rate; DoR, duration of response; ESR1, estrogen receptor alpha gene; FIH, first-in-human; HbA1c, hemoglobin A1c; HER2-, human epidermal growth factor receptor 2-negative; HR+, hormone receptor-positive; KM, Kaplan Meier; MedDRA, Medical Dictionary for Regulatory Activities; MTD, maximum tolerated dose; mut, mutations; ORR, objective response rate; PD, progressive disease; (m)PFS, (median) progression-free survival; PI3Kα, phosphatidylinositol 3-kinase alpha; PIK3CA, phosphatidylinositol-4,5-bisphosphate 3-kinase, catalytic subunit alpha; PK, pharmacokinetics; PR, partial response; RECIST 1.1, Response Evaluation Criteria in Solid Tumors version 1.1; RP2D, recommended Phase 2 dose; SD, stable disease; SERD, selective estrogen receptor degrader; SOC, standard-of-care; TRAE, treatment-related adverse event.



#### Figure 5. Encouraging mPFS across PIK3CA genotypes

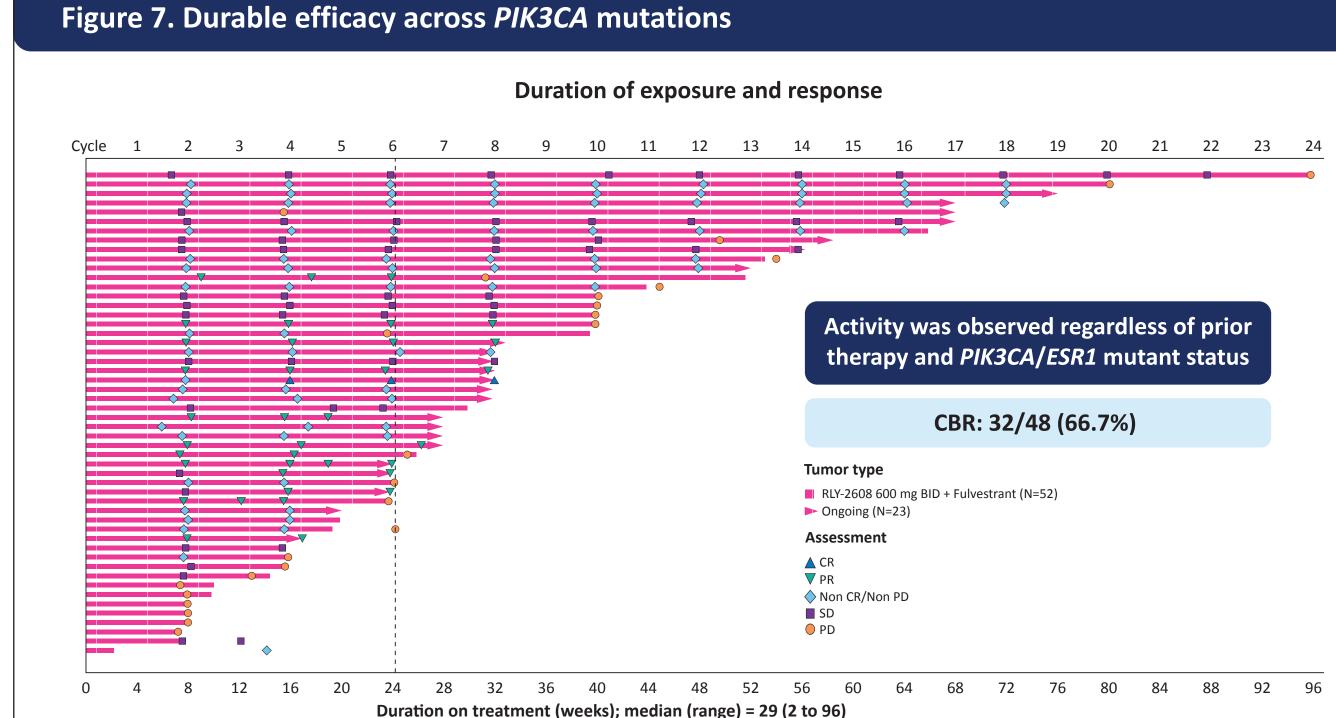


#### Figure 6. Radiographic response and PFS in patients with PIK3CA kinase mutations

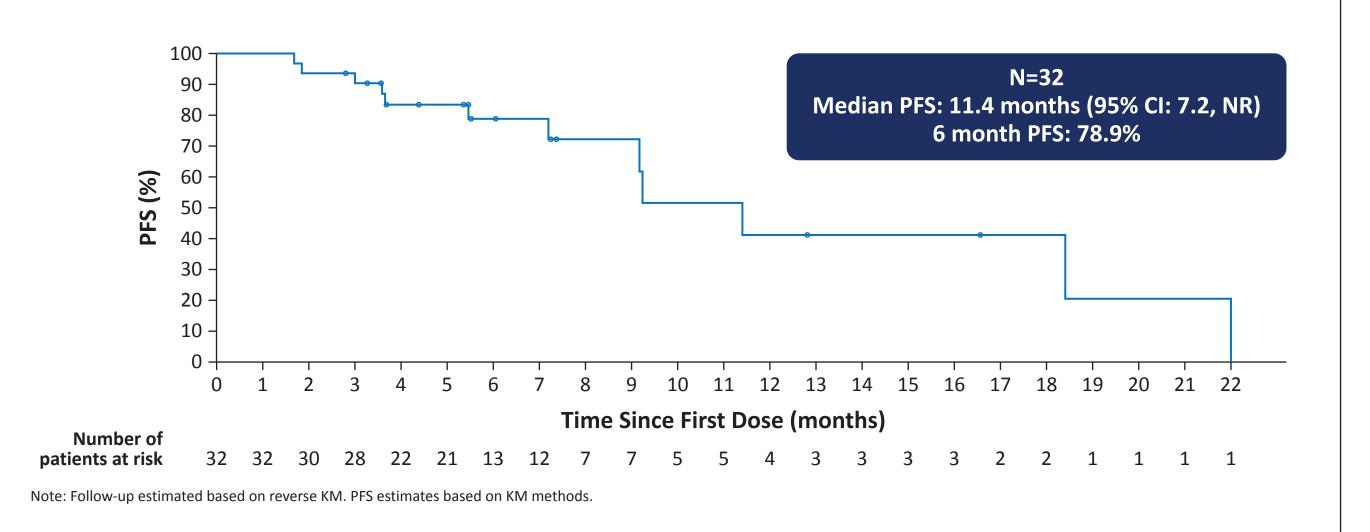


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#### Figure 8. PFS in patients treated with RLY-2608 + fulvestrant as 2L therapy



# CONCLUSIONS

Note: Median follow-up was 9.5 months.

- The ReDiscover study validates RLY-2608 as the first allosteric, pan-mutant and isoform-selective PI3Kα inhibitor
- Durable initial efficacy was demonstrated at the recommended dose (600 mg BID) across PIK3CA genotypes in patients with advanced HR+/HER2- breast cancer, confirming potent, pan-mutant targeting
- Tumor reduction observed in 74% of patients and ORR was 39% across *PIK3CA* genotypes • An ORR of 67% observed in patients with kinase domain mutations
- Highest-ever PFS observed in patients with PIK3CA-mutated HR+/HER2- advanced BC previously treated with CDK4/6i
- Median PFS 9.2 months (95% CI: 5.8–18.4) across mutation types
- Median PFS 11.4 months (95% CI: 7.2–NR) in patients receiving RLY-2608 as 2L treatment
- Forty-four percent of patients (23/52) remain ongoing with SD or response (one treated beyond progression)
- PK/PD and differentiated safety profile confirm pan-mutant selectivity
- Target inhibition was robust and continuous - The favorable safety profile eliminates or minimizes AEs commonly seen with non-selective pathway inhibitors (alpelisib, inavolisib, capivasertib, and everolimus)
- The promising efficacy and safety data observed in this FIH study suggest that RLY-2608 has broad therapeutic potential in PIK3CA-mutant, HR+/HER2- breast cancer
- A pivotal Phase 3 study will initiate in 2025: RLY-2608 + fulvestrant vs capivasertib + fulvestrant in patients with HR+/HER2- breast cancer previously treated with CDK4/6i and endocrine therapy

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