

RELAY® THERAPEUTICS

Corporate Presentation

January 2023

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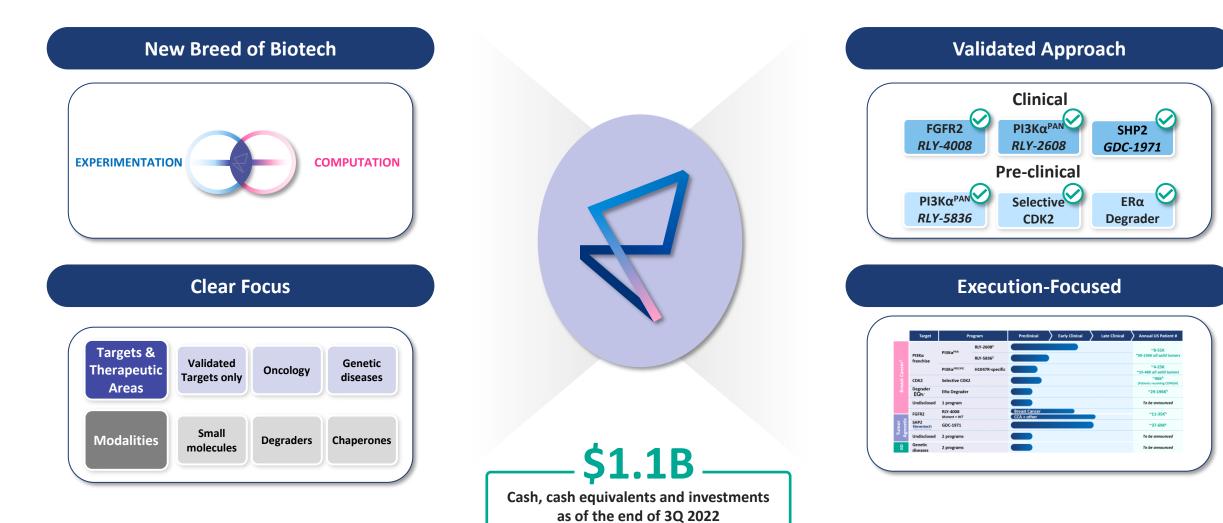
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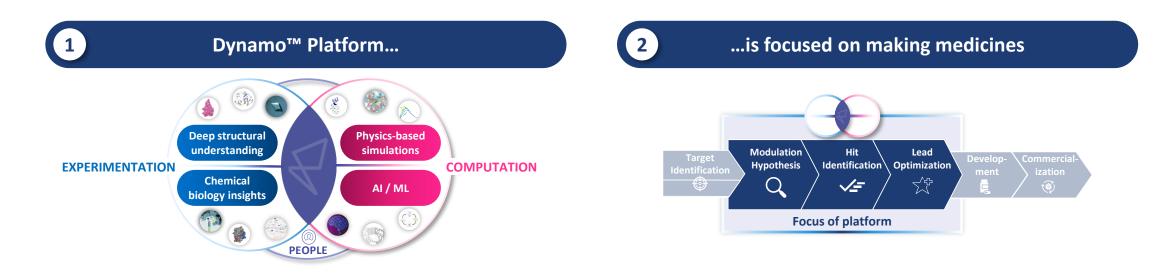
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...aims to address selectivity on validated targets



3





Company

Private

Preclinical

Over Search

Programs

2 disclosed targets

⊘ 6+ unnamed programs



ESMO & Triple Meeting

- **S**+ unnamed programs
- ✓ Platform: + ML-DEL and **Automation**



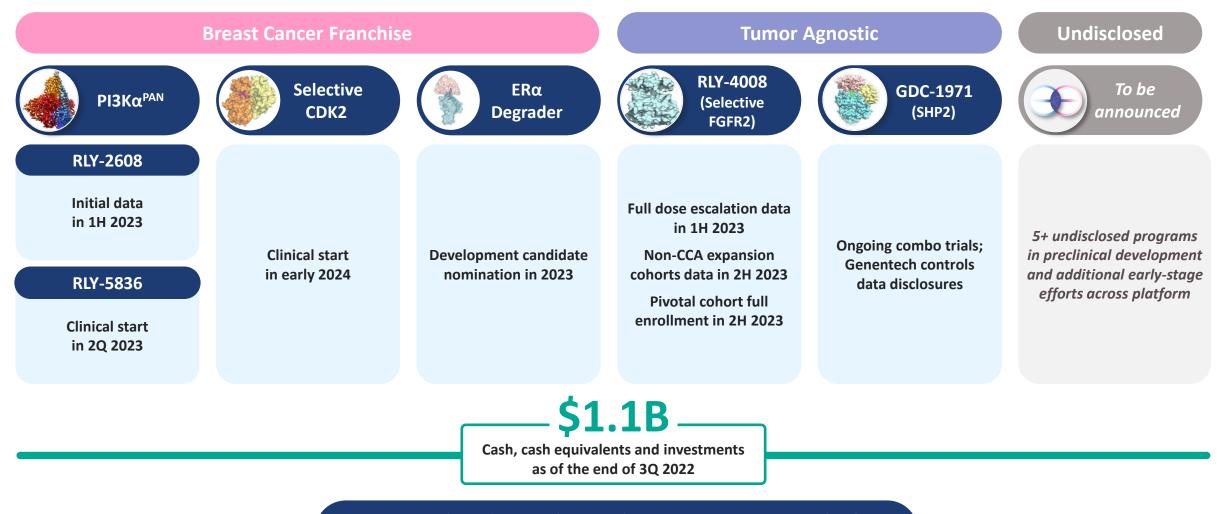
	Target	Program		Preclinical	Early Clinical	Late Clinical	Annual US Patient #
	PI3Kα franchise	ΡΙ3Κα ^{ραΝ}	RLY-2608 ²				~8-51K
		ΡΙ3Κατού	RLY-5836 ²				~50-156K all solid tumors
Cancer ¹		ΡΙ3Κα ^{specific}	H1047R-specific				~4-25K ~15-48K all solid tumors
east C	CDK2	Selective CDK2					~46K ³ (Patients receiving CDK4/6i)
Bre	Degrader EQŖ [™]	ERα Degrader					~29-196K ⁴
	Undisclosed	1 program					To be announced
	FGFR2	RLY-4008 Mutant + WT		Breast Cancer CCA + other			~11-35K⁵
Tumor Agnostic	SHP2 Genentech	GDC-1971					~37-69K ⁶
Tu Agr	Undisclosed	2 programs					To be announced
G	Genetic diseases	2 programs					To be announced

Note: Unless otherwise indicated, patient #'s refer to total annual number of US patients with late-line cancers compared to comprehensive annual incidence that may be amenable to treatment with our programs

1. Unless otherwise indicated, all breast cancer patient numbers refer to HR+/HER2- breast cancer tumors 2. RLY-2608 covers H1047X, E542X, box spots, and breast cancer patient range assumes HR+/HER2- population 3. ~46k HR+/HER2- breast cancer patients expected to receive CDK 4/6 inhibitors in adjuvant setting, first-line setting, and second-line setting in 2023, per Decision Resources Breast Cancer Market Forecast, report dated June 2022 4. HR+/HER2- US late-line breast cancer patients compared to HR+/HER2- US incident breast cancer patients 5. FGFR2 altered late-line solid tumors compared to comprehensive annual FGFR2 altered incident solid tumors including additional FGFR gene fusions and rearrangements resulting from truncation of the protein at exon 18 6. SHP2 combo only includes KRAS G12C in lung and CRC, EGFR mutations in lung, and ALK fusions in lung

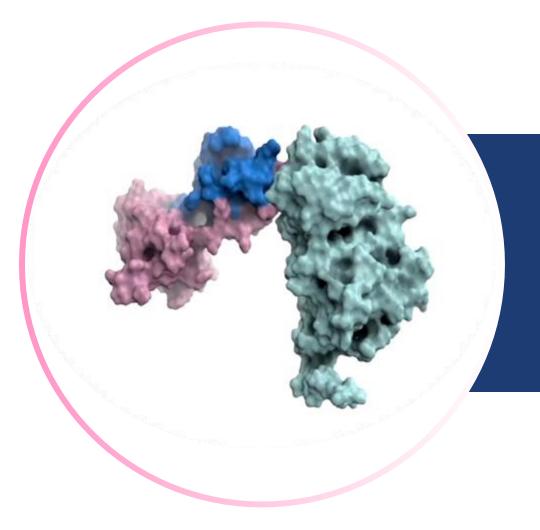
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Current cash, cash equivalents and investments are expected to be sufficient to fund current operating plan into 2025





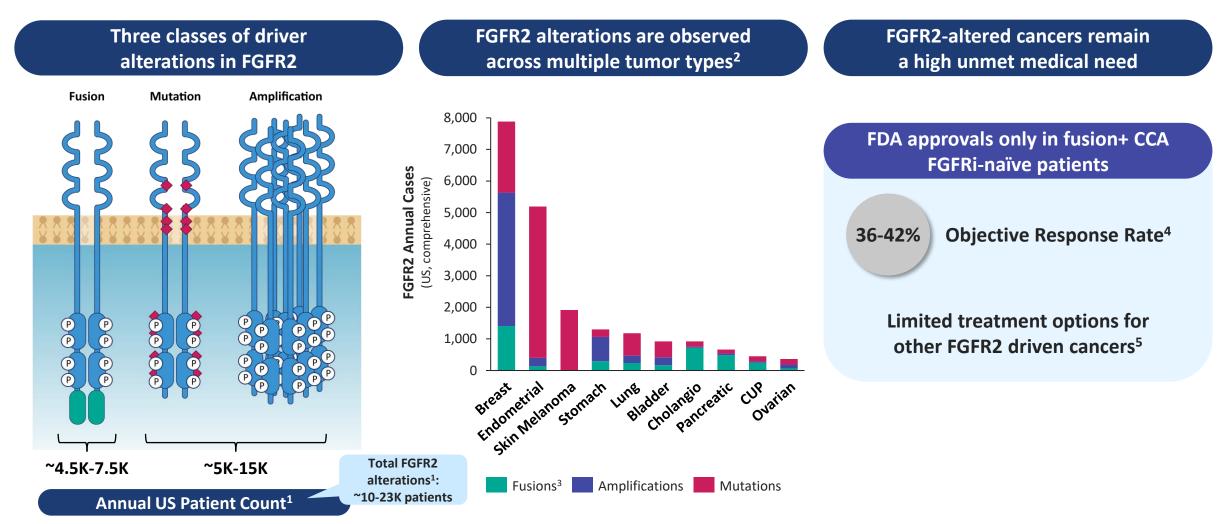
Relay Tx

Programs



	Target	Pr	ogram	Preclinical	Ear	ly Clinical	\rangle	Late Clinical
	PI3Kα franchise	ΡΙ3Κα ^{ραΝ}	RLY-2608					
5			RLY-5836					
Breast Cancer		ΡΙ3Κα ^{Specific}	H1047R-specific					
ast (CDK2	Selective CDK2						
Bre	Degrader EQ®	ERα Degrader						
	Undisclosed	1 program						
	FGFR2	RLY-4008 Mutant + WT		Breast Cancer CCA + other				
Tumor Agnostic	SHP2 Genentech	GDC-1971						
Tt Agi	Undisclosed	2 programs						
G	Genetic diseases	2 programs						



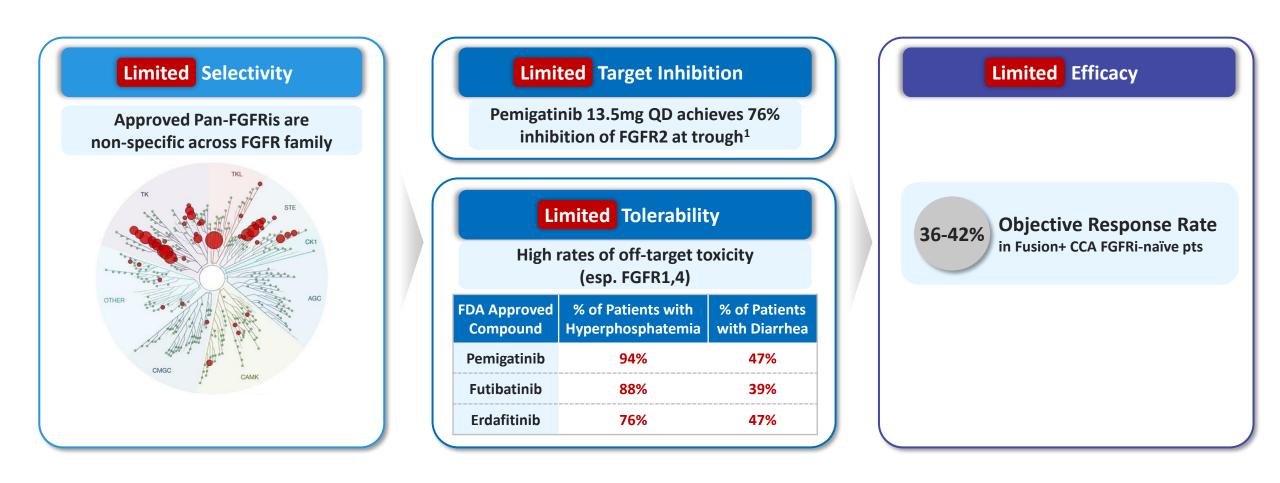


Sources: Image adapted from Babina IS, Turner NC. Nat Rev Cancer 2017;17: 318-332; Internal analysis based on third party industry data

1. All patient #'s refer to total annual number of US patients with late-line cancers vs. comprehensive annual incidence that may be amenable to treatment with our programs including additional FGFR gene fusions and rearrangements resulting from truncation of the protein at exon 18; 2. Cholangio, cholangiocarcinoma (CCA); CUP, carcinoma unknown primary; 3. FGFR2 fusion estimates include del18 truncations; 4. Based on pemigatinib, erdafitinib, and futibatinib prescribing information; 5. Erdafitinib is approved for urothelial carcinoma with FGFR2/3 alterations

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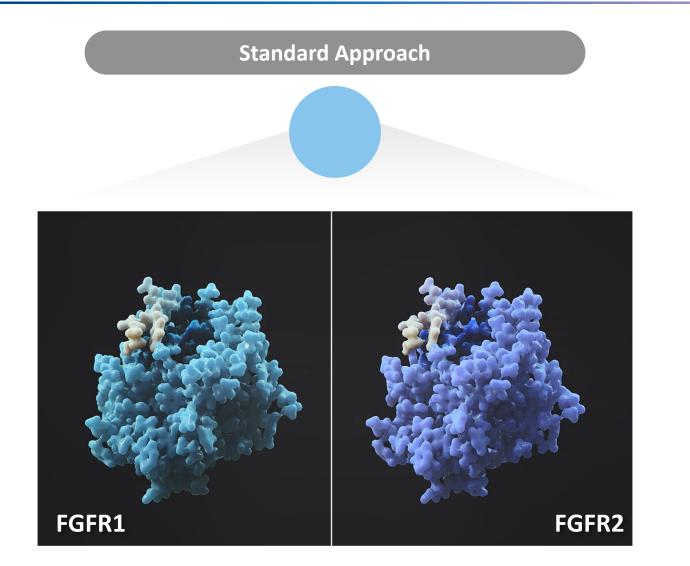


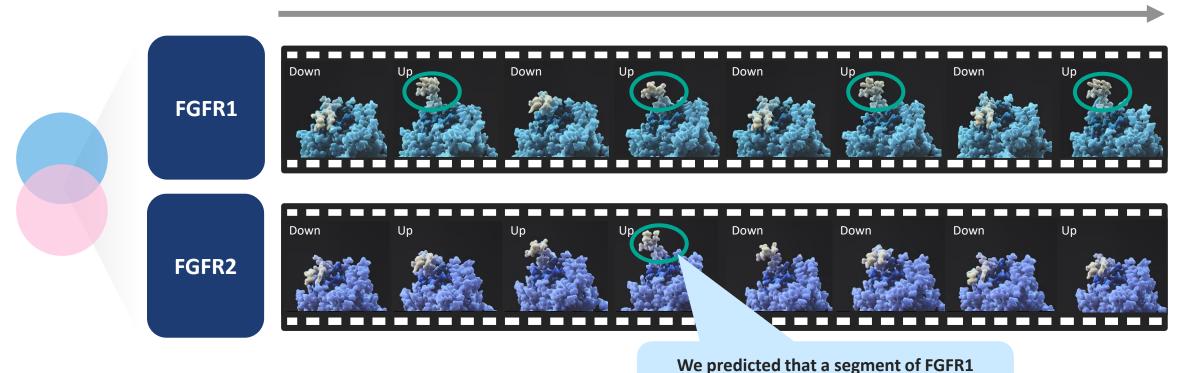


Sources: Pemigatinib – prescribing information; futibatinib – prescribing Information; erdafitinib – prescribing information 1. From pemigatinib NDA review documents: "Pemigatinib 13.5 mg daily provided 76% inhibition of ex vivo phosphorylated FGFR2α at trough" Confidential | © 2023 Relay Therapeutics

FGFR2 – Standard Approach to Discovery Has Had Limited Success



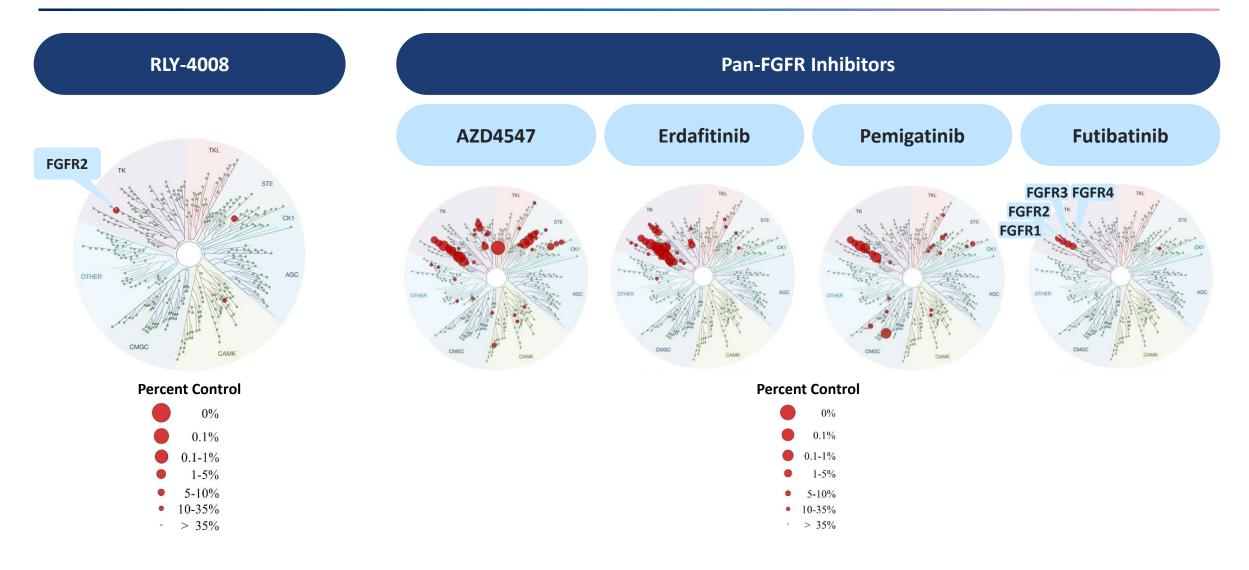




would be fully extended outwards more frequently than the same segment in FGFR2

Exploiting the dynamic difference between FGFR1 and FGFR2 enabled Relay Tx to design a selective FGFR2 inhibitor

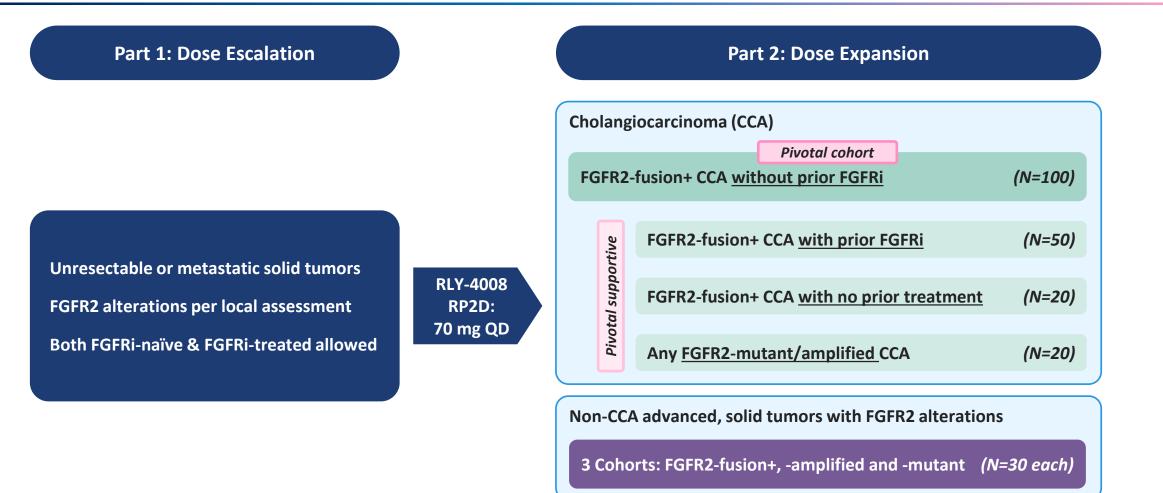




Note: Single experiment that tested each compound run at 500nM against 468 targets in the absence of ATP and without preincubation Source: KINOMEscan[™] by Eurofins DiscoverX Confidential | © 2023 Relay Therapeutics

RLY-4008 – ReFocus Trial Design





RLY-4008 – Patient Characteristics



	Fusion+ CCA			
Parameter	70 mg QD (N=17)	All doses (N=38)	Total (N=195) ²	
Age (years), median (range)	57 (36-81)	58 (33-81)	59 (23-87)	
Female, %	59%	58%	62%	
Race, %				
White / Asian / Black / Unknown	41% / 24% / 0% / 35%	58% / 21% / 3% / 18%	63% / 15% / 4% / 18%	
ECOG PS ³ , %				
0	53%	50%	38%	
1	47%	50%	58%	
2	0%	0%	3%	
Prior lines of systemic therapy, %				
0	0%	0%	2%	
1	41%	47%	20%	
2	47%	32%	29%	
3+	12%	21%	49%	
Baseline sum of target lesions (RECIST 1.1, mm), median (range)	57 (10-157)	63 (10-216)	79 (10-274)	

1. Efficacy analysis includes patients with previously treated, FGFR2i-naïve CCA treated at the RP2D. Patients with measurable disease who had opportunity for ≥2 tumor assessments to confirm response or discontinued treatment with <2 tumor assessments

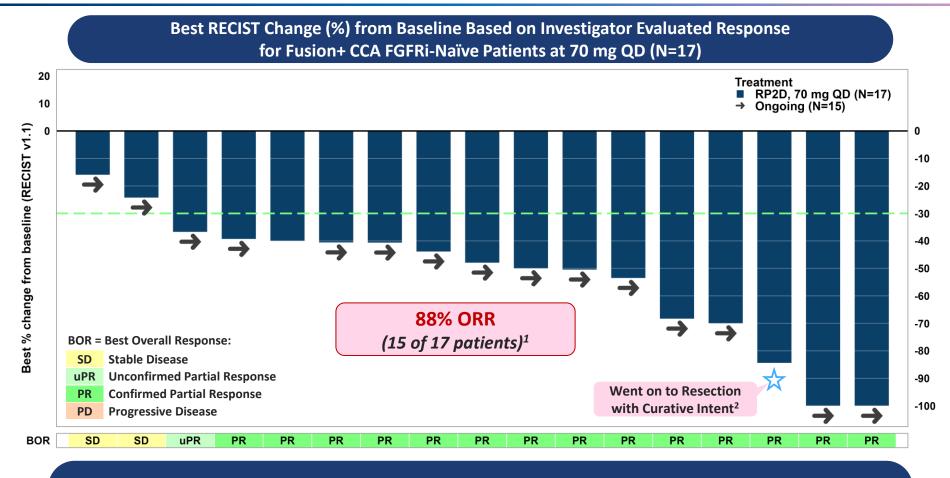
2. Patients in safety population who received ≥1 dose of RLY-4008 at any dose level

3. ECOG PS = Eastern Cooperative Oncology Group Performance Scale

RLY-4008 – Interim Response Data

FGFRi-Naïve Fusion+ CCA Patients at Pivotal Dose (70 mg QD)





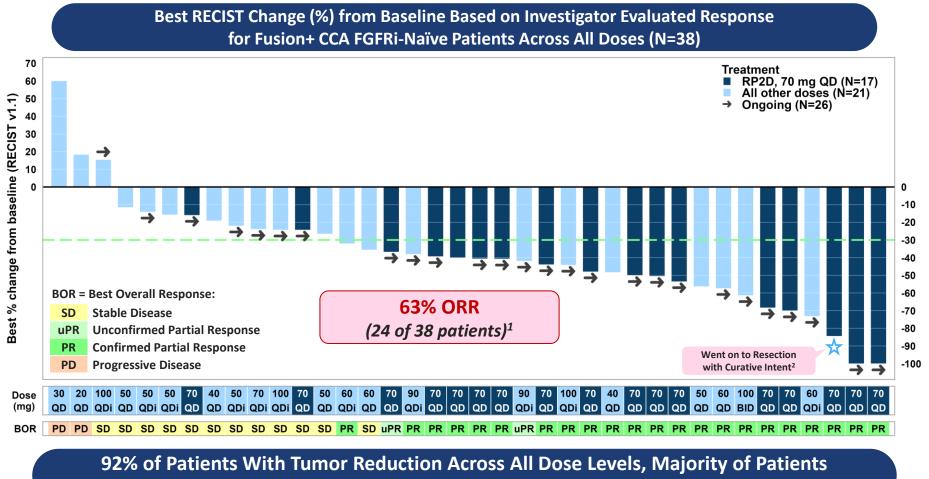
Approved Pan-FGFR Inhibitors Demonstrate 23-36% ORR in This Population³

Confirmed ORR = 82%: 14 confirmed PRs, 1 unconfirmed PR in an ongoing patient;
 Complete tumor resection: R0 resection performed in curative intent, patient remains no evidence of disease as of 31 May 2022;
 Referenced approved pan-FGFRi are Pemigatinib and Infigratinib; ORR based on prescribing information. These data are derived from different clinical trials at different points in time, with differences in trial design and patient populations. As a result, cross-trial comparisons cannot be made, and no head-to-head clinical trials have been conducted.

RLY-4008 – Interim Response Data

FGFRi-Naïve Fusion+ CCA Patients Across All Doses





With Partial Response per RECIST 1.1

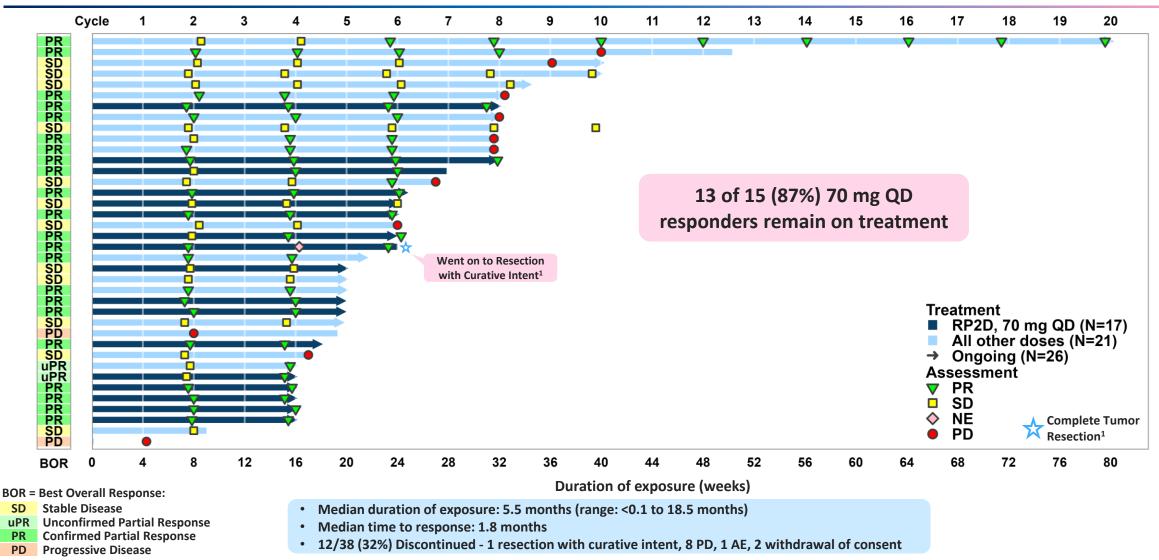
QDi = once daily dosing on an intermittent schedule (3 weeks on drug, 1 week off); BID = twice daily dosing

- 1. Confirmed ORR = 58%: 22 confirmed PRs, 2 unconfirmed PR
- 2. Complete tumor resection: R0 resection performed in curative intent, patient remains no evidence of disease as of 31 May 2022

Data from 2022 ESMO Congress (September 2022)

RLY-4008 – Time on Treatment for Fusion+ CCA FGFRi-Naïve Patients (All Doses)





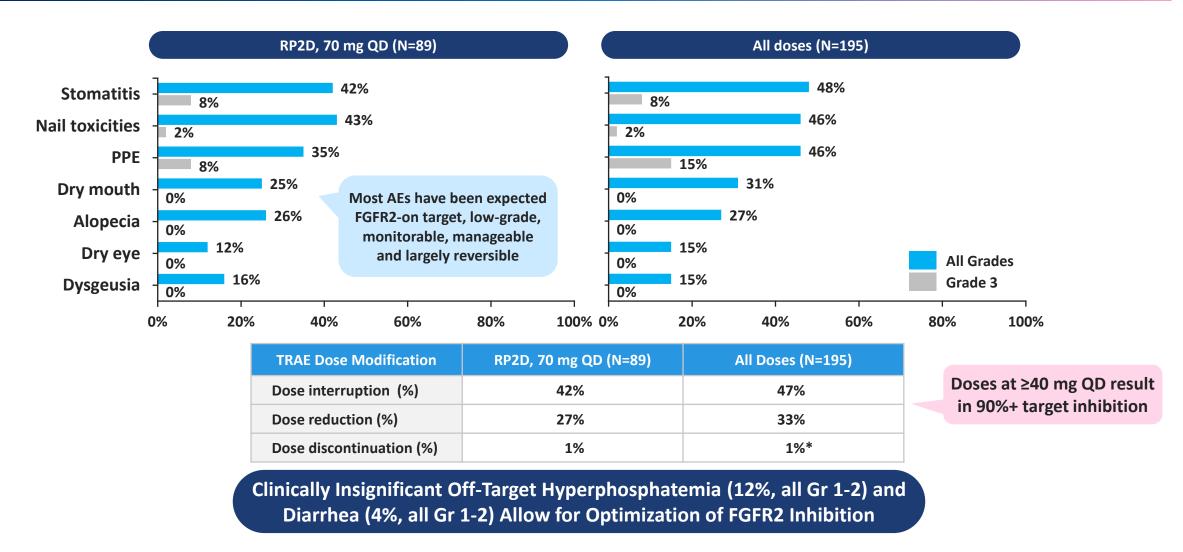
1. Complete tumor resection: R0 resection performed in curative intent, patient remains no evidence of disease as of 31 May 2022

Data cut-off: August 01, 2022; efficacy analysis includes patients with measurable disease who had opportunity for ≥ 2 tumor assessments or discontinued treatment with <2 tumor assessments or discontinued treatment with <2

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RLY-4008 – Treatment-Related Adverse Events (TRAEs) Interim Profile TRAEs > 15%



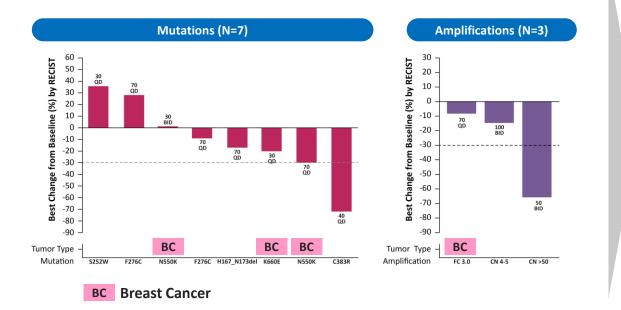


* 1 hypersensitivity, 1 retinal pigment epithelial detachment, both resolved

RLY-4008 Poised for Tumor Agnostic Validation Across FGFR2 Alterations



Tumor regression observed across FGFR2 mutations and amplifications in ReFocus Part 1 Dose Escalation Data



Continue to actively enroll tumor agnostic cohorts

Non-CCA advanced, solid tumors with FGFR2 alterations

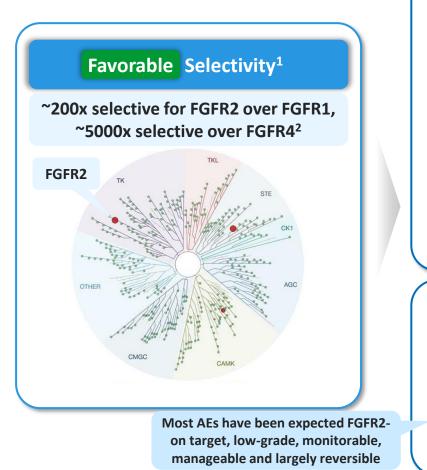
Non-CCA patients with FGFR2-fusion

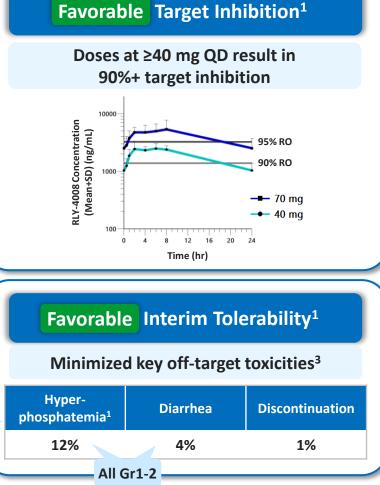
Non-CCA patients with FGFR2-amplification

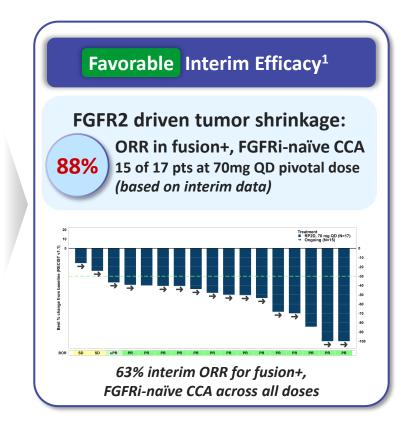
Non-CCA patients with FGFR2-mutation

Data Disclosure From Tumor Agnostic Cohorts Anticipated in 2023









Sources: KINOMEscan[™] by Eurofins DiscoverX; RLY-4008 data as presented at ESMO Congress 2022

1. Interim data as of 01 August 2022; 2. Single experiment that tested each compound run at 500nM against 468 targets in the absence of ATP and without preincubation; 3. Toxicity rates across all doses, n=195 patients



		Dose Escalation nical proof of mechanism	RP2D		Dansion and Pivotal Co se and establish efficacy	ohort
		FGFR2 alterations per local assessme e & FGFRi-treated allowed	nt;	Pivotal: Fusion+ CCA, FGFRi-naïve	FGFR2-altered, non-CCA tumors	Other FGFR2- altered CCA
	Selectivity					
ciosure		Target Inhibition				
FOCUS OT DISCIOSULE		Tolerability				
Focus		Early	Signals of Efficacy			
	40 2021:	Triple Meeting		Inter	pretable Efficacy	
	 Validated hy inhibition a Provided ea 	ypothesis with target nd acute tolerability rly efficacy signals in ous populations • Estal	3Q 2022: ESM irmed longer term safet plished stronger support on+ CCA, FGFRi-naïve eff	y and tolerability t for FGFR2 • 2H icacy • 2H	2023: Anticipated Miles I 2023: Full dose escalation da I 2023: Pivotal cohort full enro I 2023: Non-CCA expansion co	ata ollment



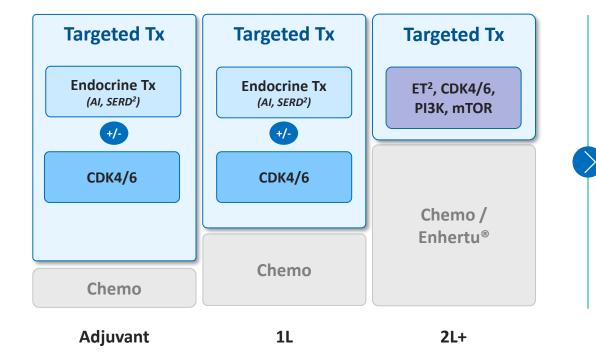
	Target	Program		Preclinical	Early Clinical	Late Clinical
			RLY-2608			
5	PI3Kα franchise	ΡΙ3Κα ^{ραΝ}	RLY-5836			
Cancer		ΡΙ3Κα ^{specific}	H1047R-specific			
Breast (CDK2	Selective CDK2				
Bre	Degrader EQ _® ™	ERα Degrader				
	Undisclosed	1 program				
	FGFR2	RLY-4008 – Mutant + WT		Breast Cancer		
U				CCA + other		
	SHP2 Genentech A Member of the Roche Group	GDC-1971				
	Undisclosed	2 programs				
G	Genetic diseases	2 programs				

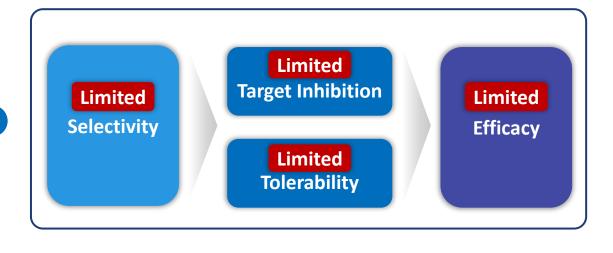


~200k annual HR+/HER2- breast cancer patients in US, of whom ~60k advance to later lines of treatment



...is limited by efficacy of available treatments

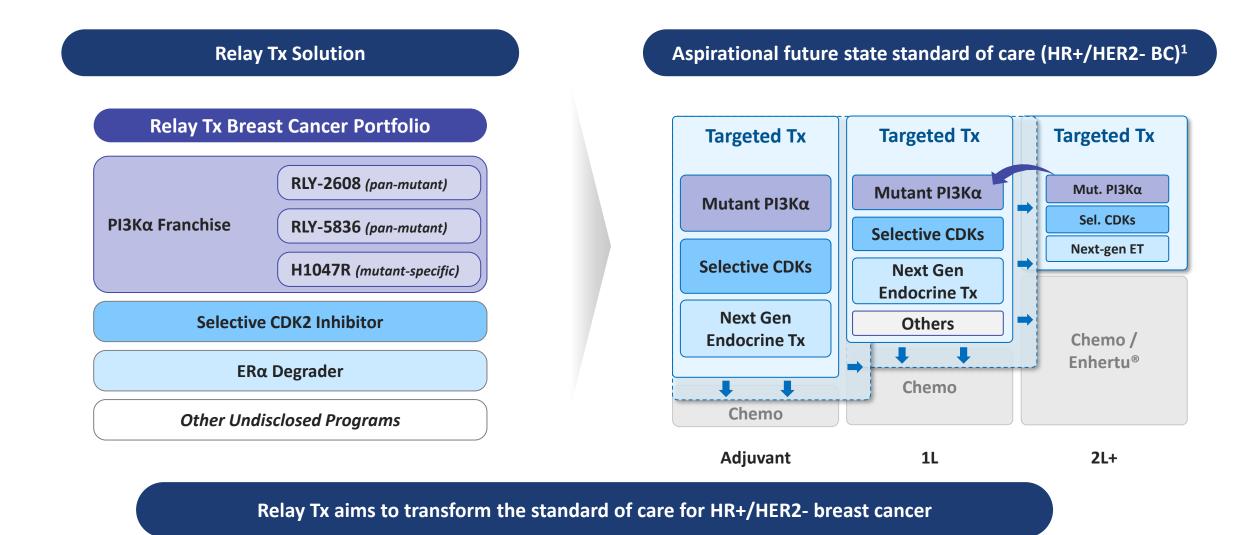




Source: Internal analysis based on third party industry data

1. Standard of care for HR+/HER2- breast cancer is illustrative; 2. AI = Aromatase Inhibitor; SERD: Selective Estrogen Receptor Degrader; ET = Endocrine Therapy





1. Aspirational future state standard of care for HR+/HER2- breast cancer is illustrative

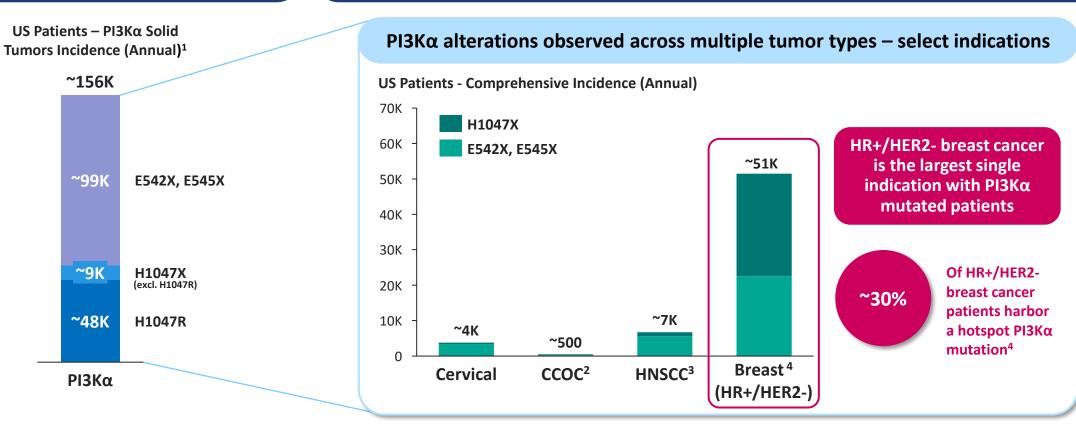


	Target	Program		Preclinical	Early Clinical	Late Clinical
	ΡΙ3Κα franchise	ΡΙ3Κα ^{ΡΑΝ}	RLY-2608			
5		ΡΙ3Κα	RLY-5836			
Breast Cancer		ΡΙ3Κα^{SPECIFIC}	H1047R-specific			
ast (CDK2	Selective CDK2				
Bre	Degrader EQŖ [™]	ERα Degrader				
	Undisclosed	1 program				
0	FGFR2	RLY-4008 Mutant + WT		Breast Cancer CCA + other		
	SHP2 Genentech A Member of the Roche Group	GDC-1971				
	Undisclosed	2 programs				
G	Genetic diseases	2 programs				





Pan-mutant selective drug is a significant clinical opportunity for solid tumors...



...with HR+/HER2- breast cancer as the single largest indication with PI3K α mutations

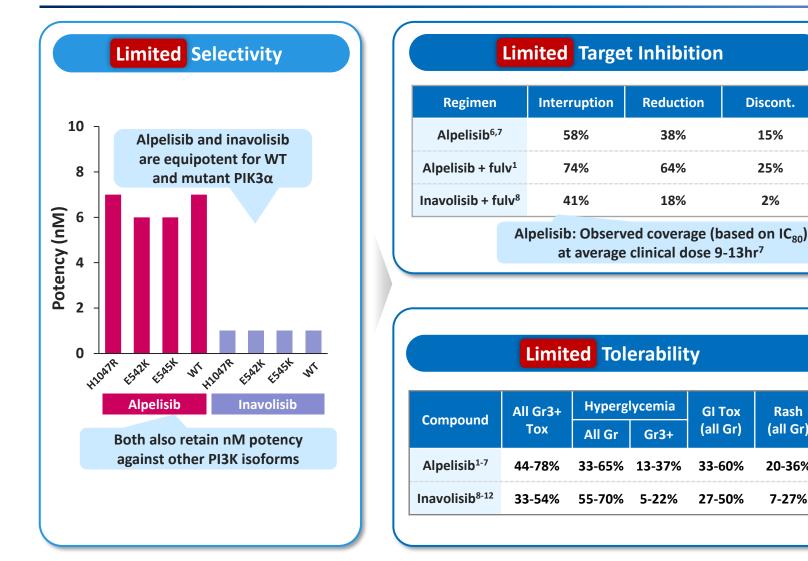
Sources: Internal analysis based on third party industry data

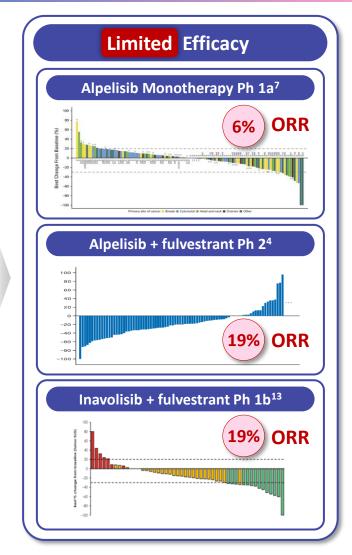
1. Annual incidence of solid tumors with PI3Ka H1047R, PI3Ka H1047X, PI3Ka E542X + E545X alterations; 2. Clear Cell Ovarian Cancer; 3. Head & Neck Squamous Cell Carcinoma;

4. HR+/HER2- breast cancer patient population with a PI3Kα hotspot alteration; alterations include: H1047X, E542X, E545X

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Note: fulv = fulvestrant; BC= breast cancer; all referenced studies are for their patient populations which are analogous to ongoing patient populations within RLY-2608 clinical trials; Alpelisib and fulvestrant are FDA approved, Inavolisib is in Phase 3 clinical trials Sources: Alpelisib – 1. SOLAR-1: Andre 2019 N Engl J Med 380:1929, 2. Ph 1b: SABCS 2013 P2-16-14, 3. Ph 1b: SABCS 2014 PD5-5, 4. Ph 2 ByLIEVE: Rugo 2021 Lancet Oncol 22:489, SABCS 2021 #P1-18-03, 5. Ph 1b mono: Annals of Oncol 25 2014 (suppl 4), 6. Ph 2 mono: Savas Cancer Discov 2022 Sep 12:2058, 7. Ph 1a mono: Juric 2018 J Clin Oncol 36:1291; Inavolisib – 8. ASCO 2022 #1052, 9. SABCS 2020 #PS11-11, 10. AACR 2020 CT109, 11. SABCS 2019 OT1-08-04; 12. SABCS 2019 P1-19-46, 13. SABCS 2021 #P5-17-05; Confidential | © 2023 Relay Therapeutics 29

Discont.

15%

25%

2%

Rash

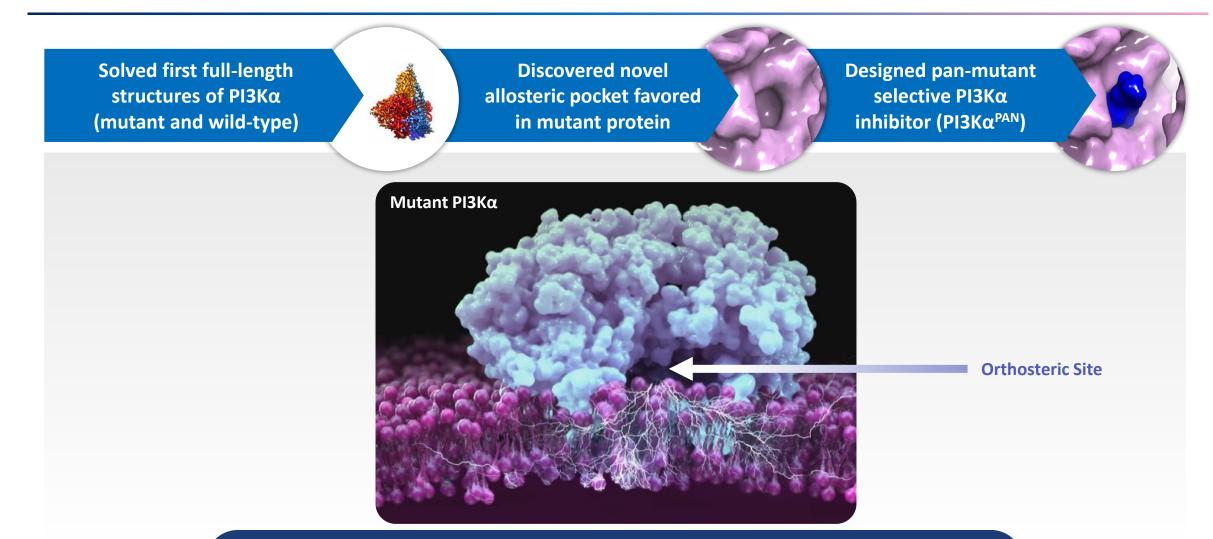
(all Gr)

20-36%

7-27%

PI3Kα – Proprietary Insights Unlock Novel Approaches

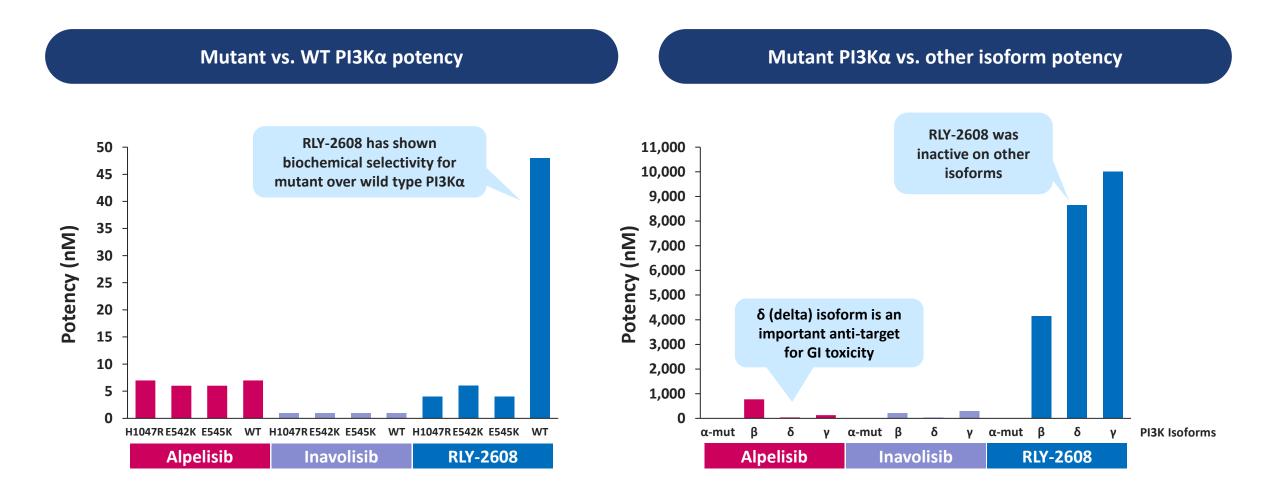




A differentiated understanding of the structure of PI3Kα and its relationship to function equips Relay Tx to design optimal mutant-selective inhibitors of PI3Kα

RLY-2608 – Mutant and Isoform Selectivity

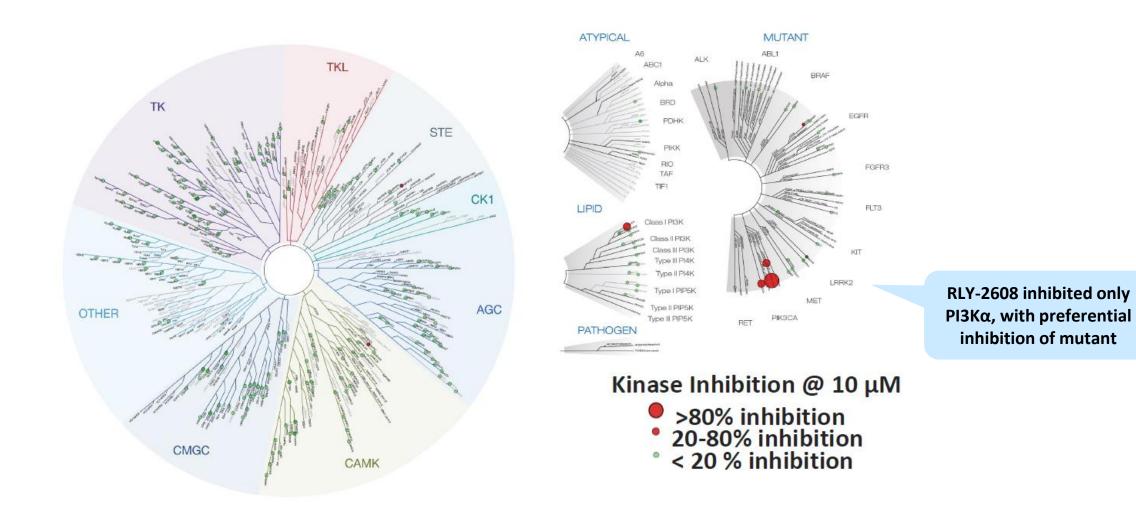




Selectivity

RLY-2608 – Selective Across the Kinome

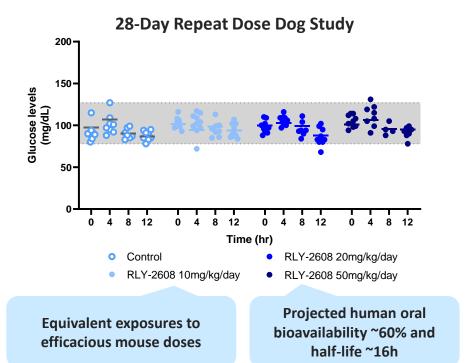




RLY-2608 – Reduced Impact on Glucose Homeostasis



Repeat dosing of RLY-2608 did not cause hyperglycemia in tox species (dog)



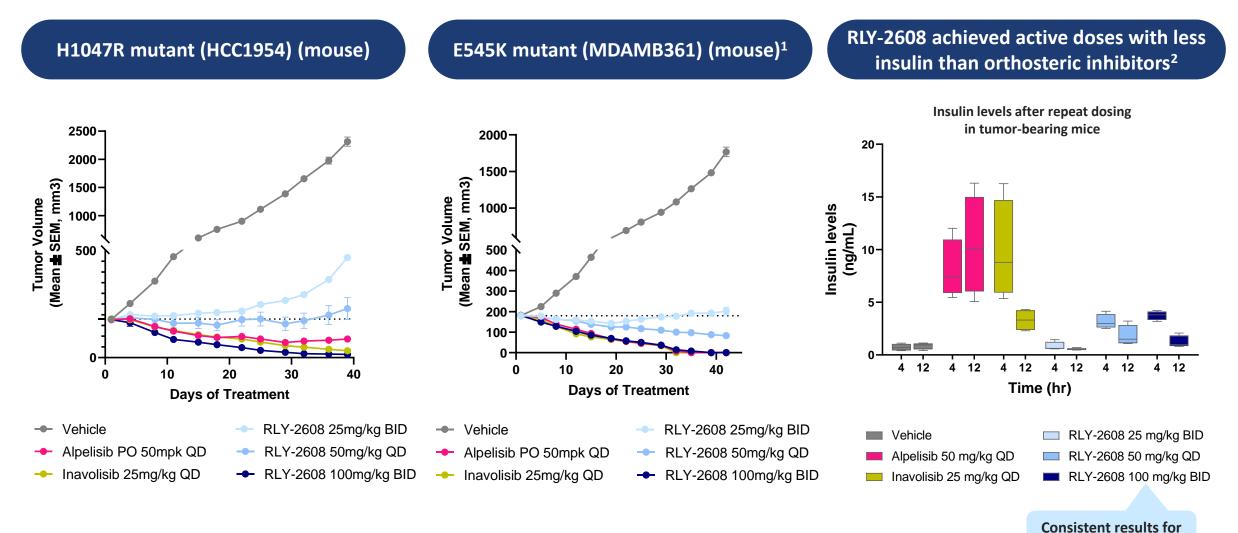
Hyperglycemia Definitions (CTCAE v5.0)

Grade	CTCAE Definition (v5.0)
Gr 1	Abnormal glucose above baseline, no medical intervention
Gr 2	Change in daily management from baseline for diabetic; oral antiglycemic agent initiated; workup for diabetes
Gr 3	Hospitilization indicated; insulin therapy initiated
Gr 4	Life-threatening consequences; urgent intervention indicated
Gr 5	Death

In higher species, dosing of RLY-2608 for 28 days showed no histopathological or ophthalmic findings associated with hyperglycemia

RLY-2608 – In Vivo Tumor Regressions



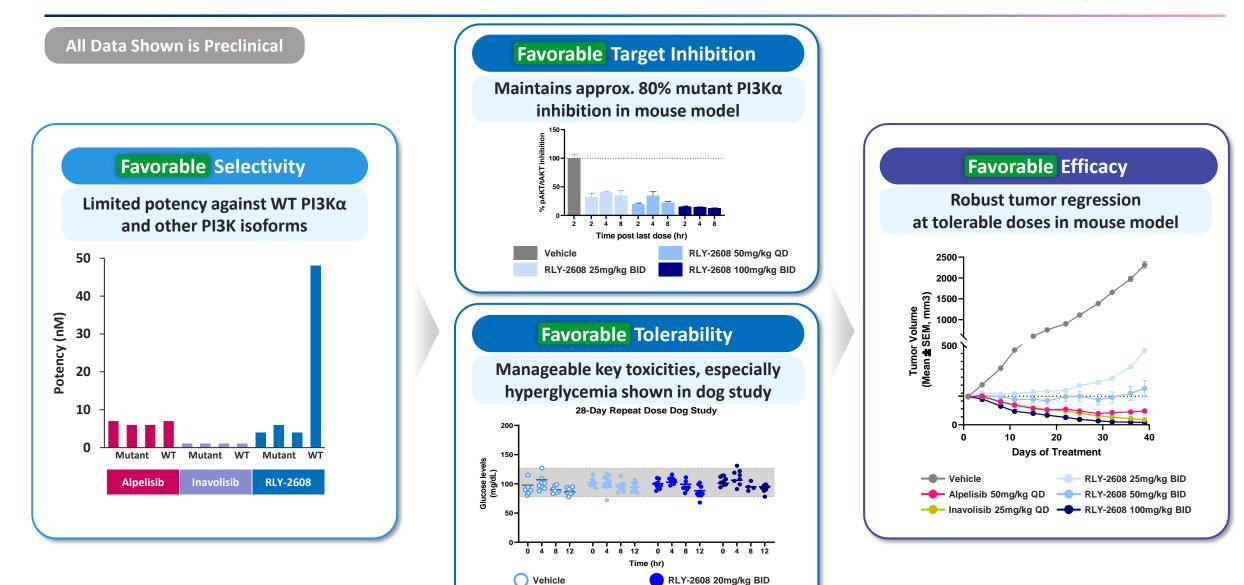


Source: RLY-2608 data as presented in 2021 AACR-NCI-EORTC Molecular Targets Conference poster presentation

1. This model also carries a second mutation at K567R; 2. HSC2 model; 3. Similar results observed in the same background strain at 1hr timepoint in the MCF7 (E545K) model

1-hour time point³

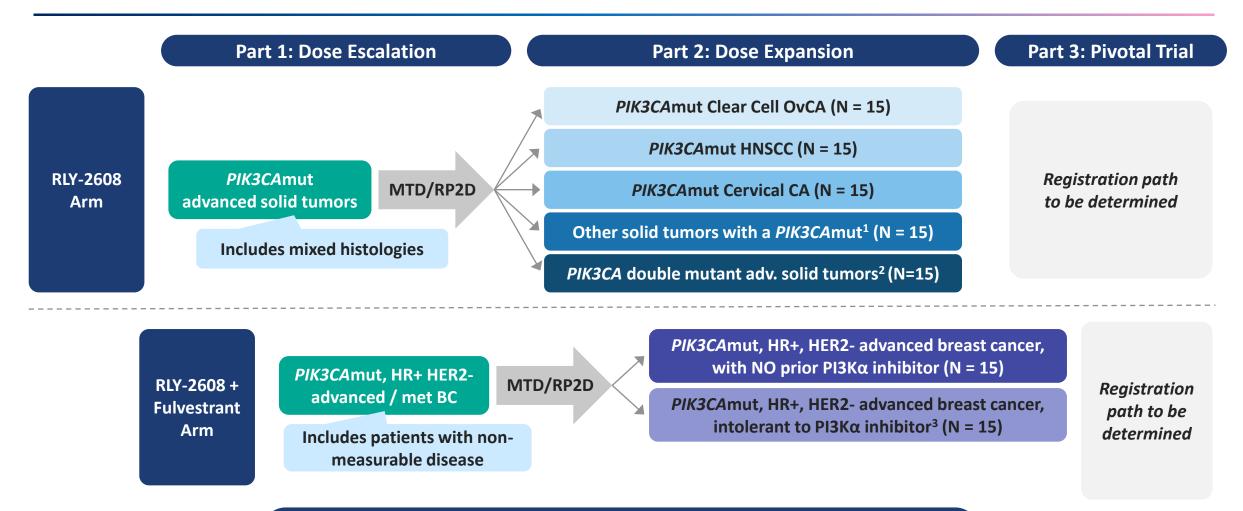




RLY-2608 10mg/kg BID

RLY-2608 50mg/kg BID

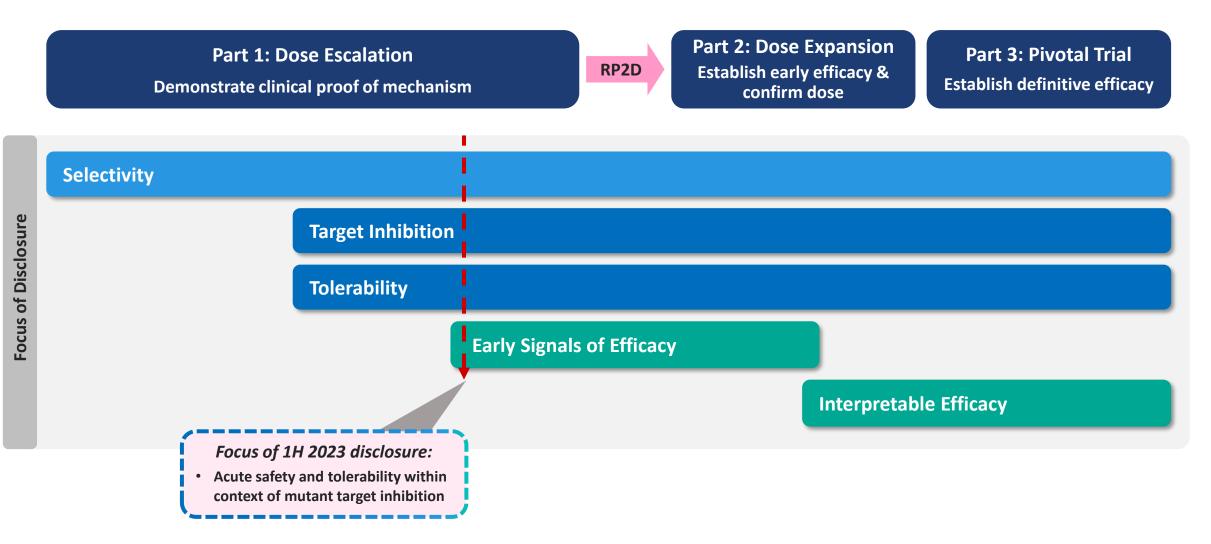




Initial clinical data update expected in 1H 2023

1. Excludes PIK3CAmut clear cell OvCA, HNSCC, and Cervical cancer patients; 2. Double mutation defined as one major PIK3CA mutation (E542X, E545X, H1047X) + \geq 1 additional PI3KCA mutation per local assessment; 3. Intolerance to PI3K α inhibitors is defined as treatment discontinuation due to treatment-related AE (e.g., hyperglycemia, rash, diarrhea, stomatitis) other than severe hypersensitivity reaction and/or life-threatening reactions, such as anaphylaxis and Stevens-Johnson syndrome.



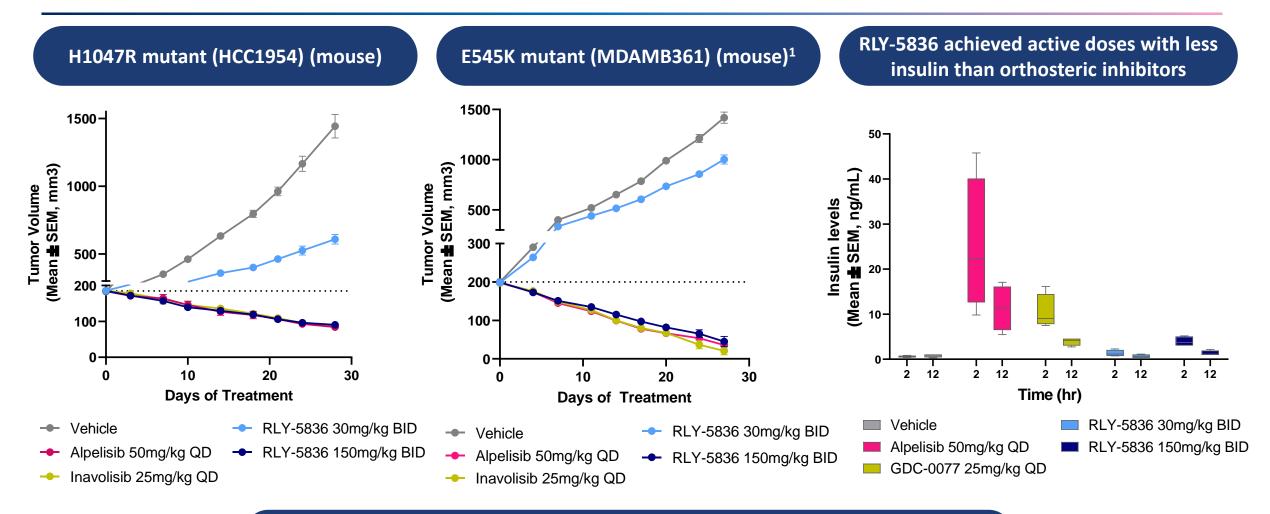




	Target	Program		Target Program Preclinical						Late Clinical
Tumor Agnostic	PI3Kα franchise	DIOK- PAN	RLY-2608							
		ΡΙ3Κα ^{ΡΑΝ}	RLY-5836							
		ΡΙ3Κα ^{Specific}	H1047R-specific							
	CDK2	Selective CDK2								
	Degrader EQŖ™	ERα Degrader								
	Undisclosed	1 program								
	FGFR2	RLY-4008 Mutant + WT		Breast Cancer CCA + other						
	SHP2 Genentech A Member of the Roche Group	GDC-1971								
	Undisclosed	2 programs								
O 5	Genetic diseases	2 programs								

RLY-5836 – Similar Pre-clinical Profile, Different Chemical Properties from RLY-2608





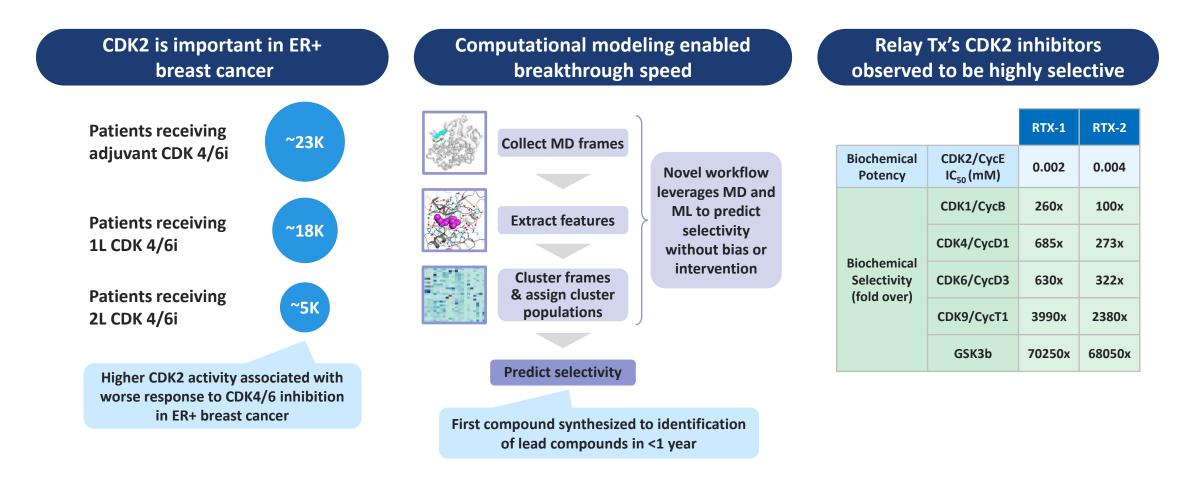
Clinical start anticipated in 2Q 2023

Source: Internal RLY-5836 data 1. This model also carries a second mutation at K567R



	Target	Program		Preclinical	Early Clinical	Late Clinical
Tumor Agnostic		ΡΙ3Κα ^{ΡΑΝ}	RLY-2608			
		ΡΙ3Κα	RLY-5836			
	PI3Kα franchise	ΡΙ3Κα ^{specific}	H1047R-specific		 	
	CDK2	Selective CDK2				
	Degrader EQ _® ™	ERα Degrader			 	
	Undisclosed	1 program				
	FGFR2	RLY-4008 Mutant + WT		Breast Cancer CCA + other		
	SHP2 Genentech A Member of the Roche Group	GDC-1971				
	Undisclosed	2 programs				
0 5	Genetic diseases	2 programs				

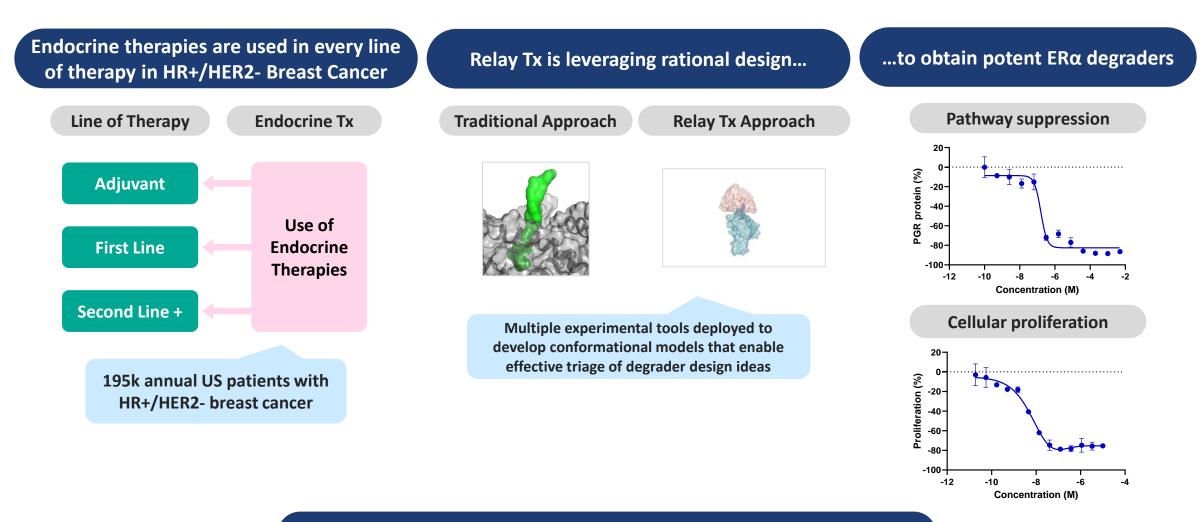




Clinical start expected in early 2024

Source: Internal analysis based on third party industry data Confidential | © 2023 Relay Therapeutics

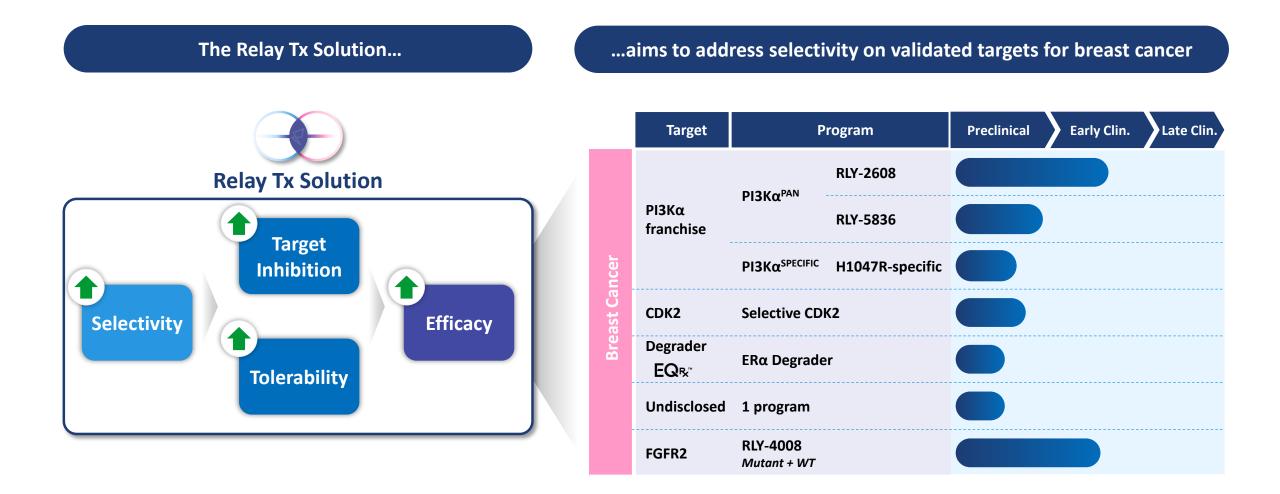




Development Candidate nomination expected in 2023

Source: Internal analysis based on third party industry data Confidential | © 2023 Relay Therapeutics

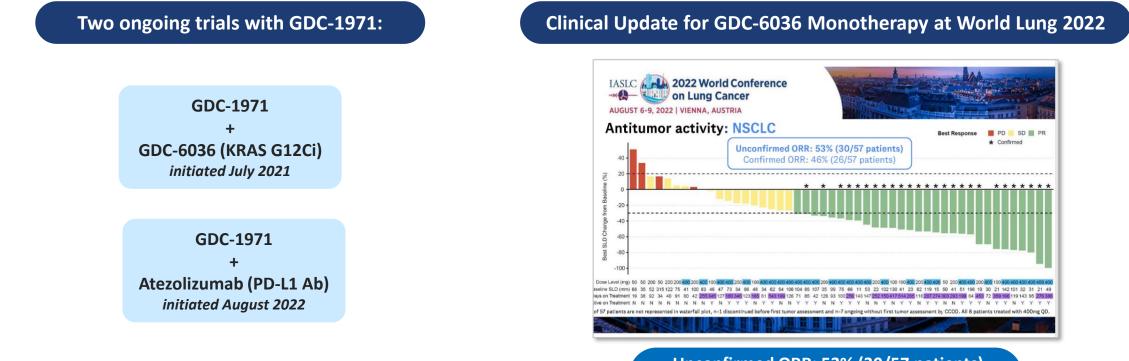






	Target	Program		Preclinical	Early Clinical	Late Clinical
	PI3Kα franchise	DIOI/ PAN	RLY-2608			
Tumor Agnostic		ΡΙ3Κα ^{ΡΑΝ}	RLY-5836)	
		ΡΙ3Κα^{SPECIFIC}	H1047R-specific			
	CDK2	Selective CDK2				
	Degrader EQR:	ERα Degrader				
	Undisclosed	1 program				
	FGFR2	RLY-4008 Mutant + WT		Breast Cancer CCA + other		
	SHP2 Genentech A Member of the Roche Group	GDC-1971				
	Undisclosed	2 programs				
QD	Genetic diseases	2 programs				





Unconfirmed ORR: 53% (30/57 patients) Confirmed ORR: 46% (26/57 patients)

Collaboration provides meaningful economics to Relay Tx¹

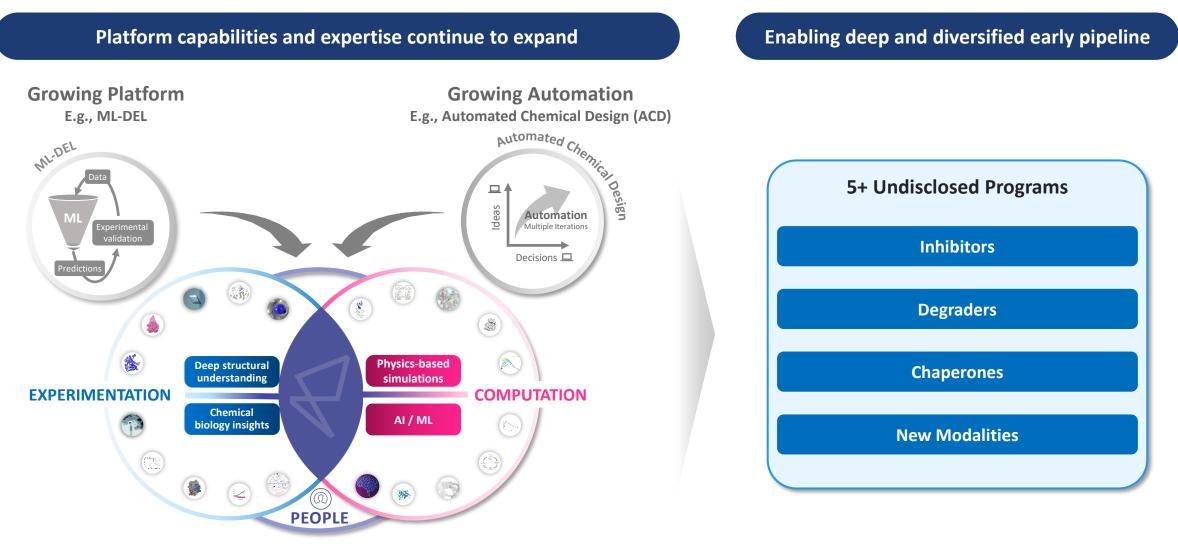
Source: World Lung 2022 #OA03.04

1. As of December 31, 2022: \$105 million in upfront & milestone payments received, plus an opt-in option for 50/50 profit share and up to \$690M in potential additional total milestones, low-to-mid teen royalties on global net sales plus eligible to receive additional royalties upon approval of GDC-1971 and GDC-6036 in combination



	Target	Program		Preclinical	Early Clinical	Late Clinical
Breast Cancer		DIOK. PAN	RLY-2608			
	PI3Kα franchise	ΡΙ3Κα^{ΡΑΝ}	RLY-5836			
		ΡΙ3Κα ^{SPECIFIC}	H1047R-specific			
	CDK2	Selective CDK2				
	Degrader EQ®	ERα Degrader				
	Undisclosed	1 program				
Tumor Agnostic	FGFR2	RLY-4008 Mutant + WT		Breast Cancer CCA + other		
	SHP2 Genentech	GDC-1971				
	Undisclosed	2 programs				
G	Genetic diseases	2 programs				







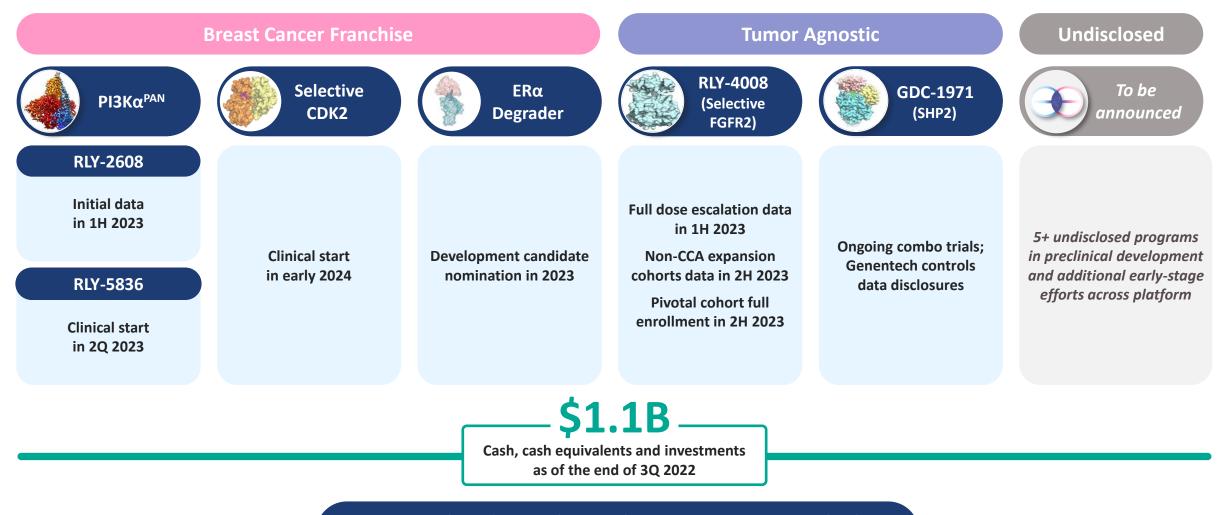
	Target	Program		Preclinical		Early Clinical	Late Clinical	Annual US Patient #	
Ę.	PI3Kα franchise	ΡΙ3Κα ^{ραΝ}	RLY-2608 ²					~8-51K	
		ΡΙ3Κα	RLY-5836 ²					~50-156K all solid tumors	
Cancer ¹		ΡΙ3Κα ^{specific}	H1047R-specific					~4-25K ~15-48K all solid tumors	
east C	CDK2	Selective CDK2						~46K³ (Patients receiving CDK4/6i)	
Tumor Agnostic	Degrader EQ®™	ERα Degrader						~29-196K⁴	
	Undisclosed	1 program						To be announced	
	FGFR2	RLY-4008 Mutant + WT		Breast Cancer CCA + other				~11-35K⁵	
	SHP2 Genentech	GDC-1971						~37-69K⁵	
	Undisclosed	2 programs						To be announced	
G	Genetic diseases	2 programs						To be announced	

Note: Unless otherwise indicated, patient #'s refer to total annual number of US patients with late-line cancers compared to comprehensive annual incidence that may be amenable to treatment with our programs

1. Unless otherwise indicated, all breast cancer patient numbers refer to HR+/HER2- breast cancer tumors 2. RLY-2608 covers H1047X, E542X, box spots, and breast cancer patient range assumes HR+/HER2- population 3. ~46k HR+/HER2- breast cancer patients expected to receive CDK 4/6 inhibitors in adjuvant setting, first-line setting, and second-line setting in 2023, per Decision Resources Breast Cancer Market Forecast, report dated June 2022 4. HR+/HER2- US late-line breast cancer patients compared to HR+/HER2- US incident breast cancer patients 5. FGFR2 altered late-line solid tumors compared to comprehensive annual FGFR2 altered incident solid tumors including additional FGFR gene fusions and rearrangements resulting from truncation of the protein at exon 18 6. SHP2 combo only includes KRAS G12C in lung and CRC, EGFR mutations in lung, and ALK fusions in lung

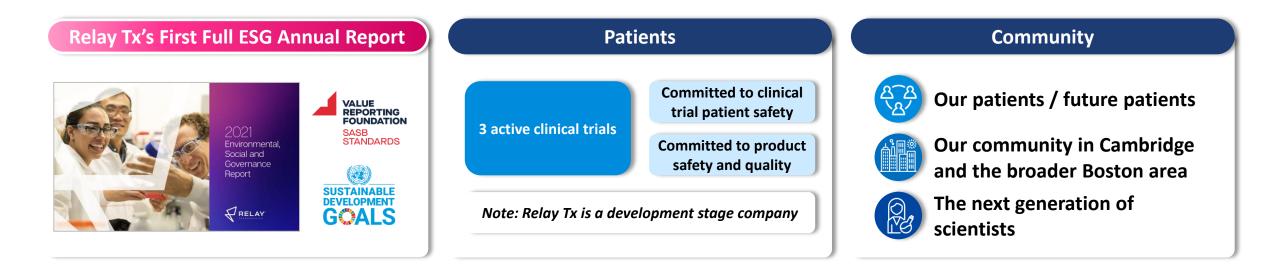
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Current cash, cash equivalents and investments are expected to be sufficient to fund current operating plan into 2025





Рео	ple		Environment		Governance			
98% employee respo "made the right decis		Ø,	Responsible energy consumption		7 Directors Total*	29%	43%	
Turnover below	Diversity & inclusion		Reducing water consumption		The Nom/Gov and Audit	Racial/Ethnic Diversity	Women	
industry average rates Training and	advisory group		Hazardous and lab waste management ef	Committees oversee ESG efforts, with the full BOD	5yrs	71%		
development opportunities	Equitable compensation	2	Non-hazardous waste management		getting ~quarterly updates	Average Tenure	Independence (Separate CEO and Chair Role) *As of December 2021	

