

Progetto CANOA

CARCINOMA MAMMARIO:

QUALI NOVITA' PER IL 2025?

"Saper leggere" uno studio clinico per migliorare la pratica clinica



Coordinatori Scientifici:
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Giovanni L. Pappagallo

Verona, 28 - 29 Marzo 2025
Hotel Crowne Plaza

AIGOM
ASSOCIAZIONE ITALIANA
GRUPPI ONCOLOGICI MULTIDISCIPLINARI



UPO
UNIVERSITÀ DEL PIEMONTE ORIENTALE

Alterazioni del pathway PIK3CA/AKT/PTEN: significato biologico, incidenza e caratteristiche delle pazienti

Benedetta Conte, MD

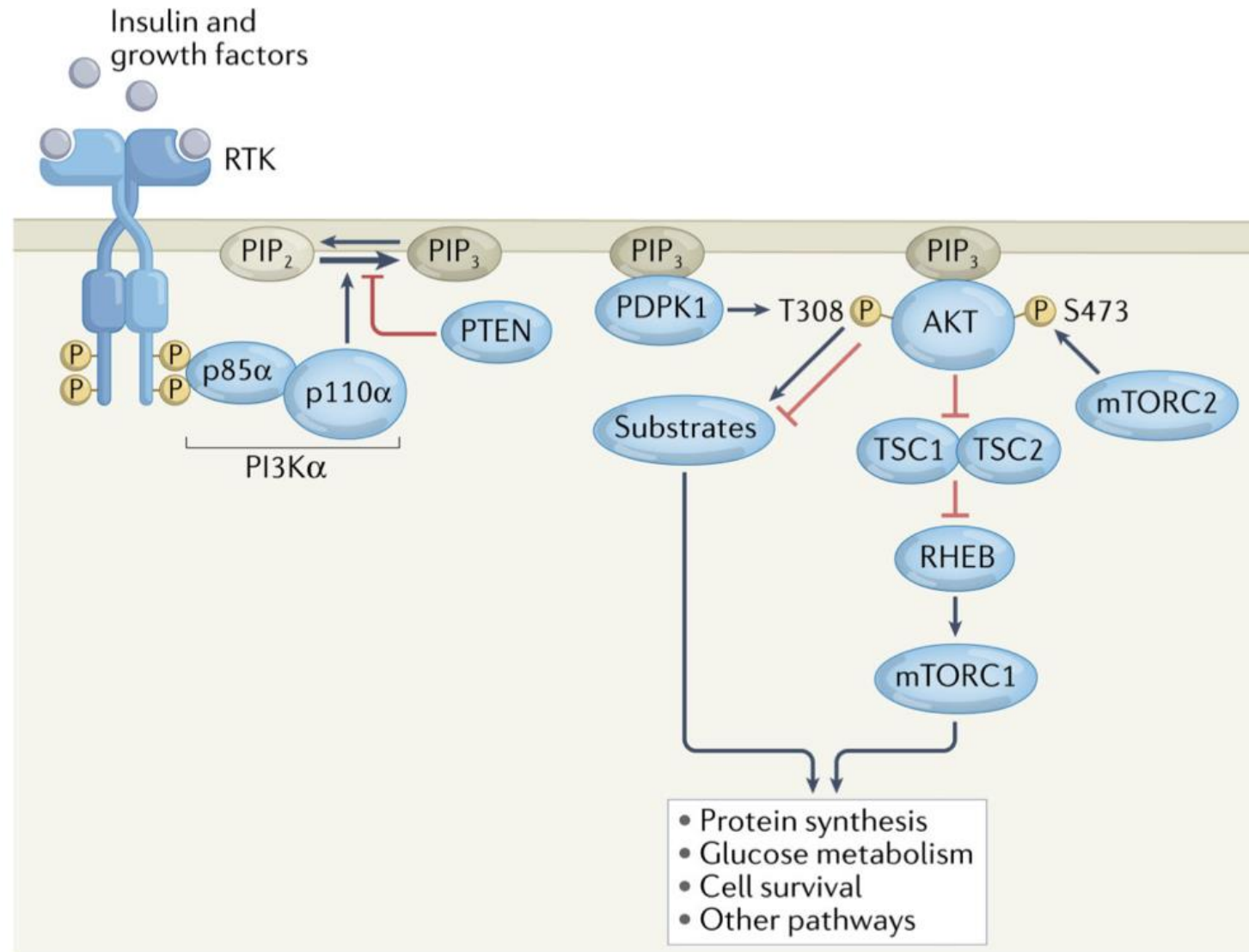
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Novara*

Declaration of interests

Benedetta Contew, MD

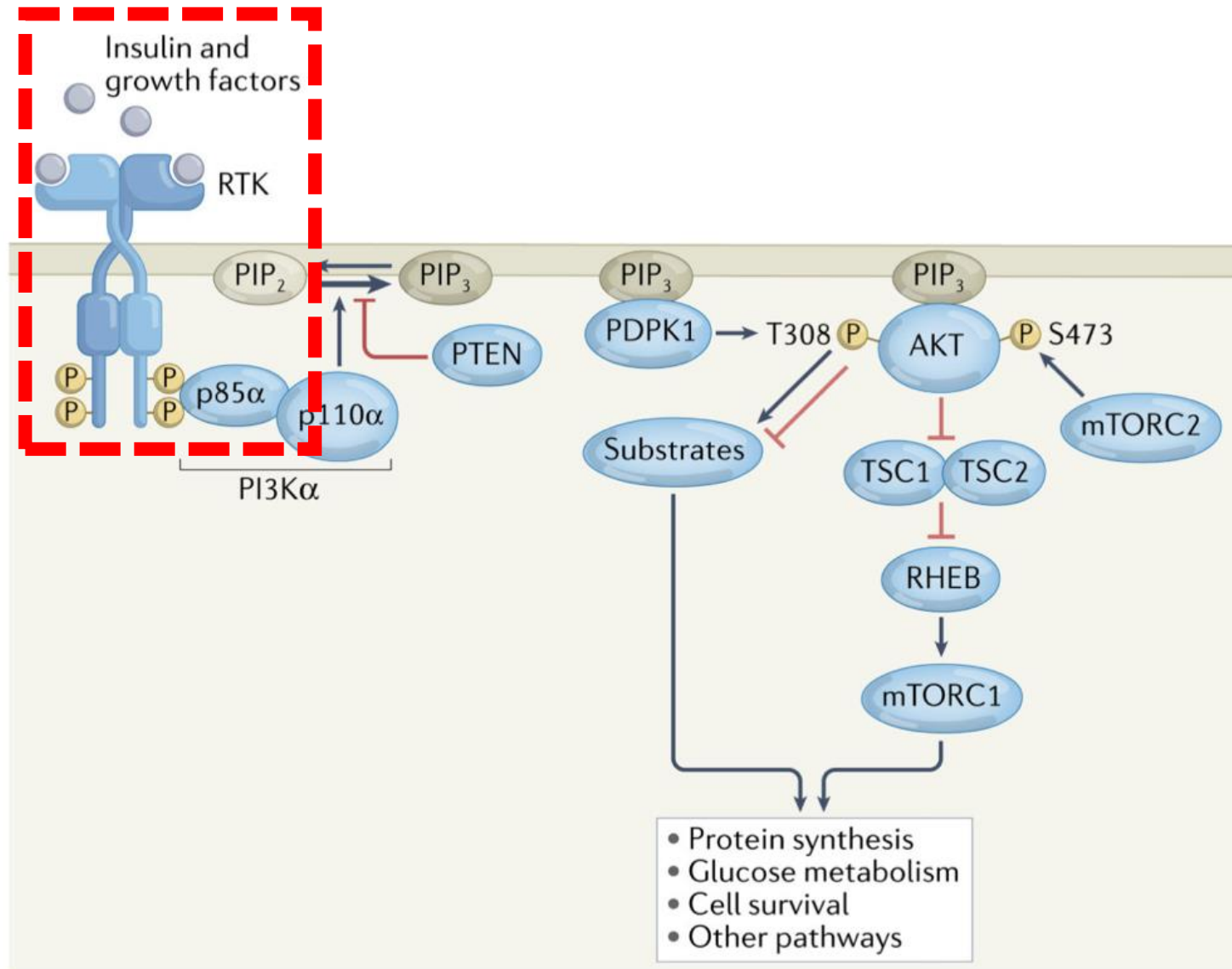
Non profit research support: GILEAD

PI3K pathway in non-malignant cells

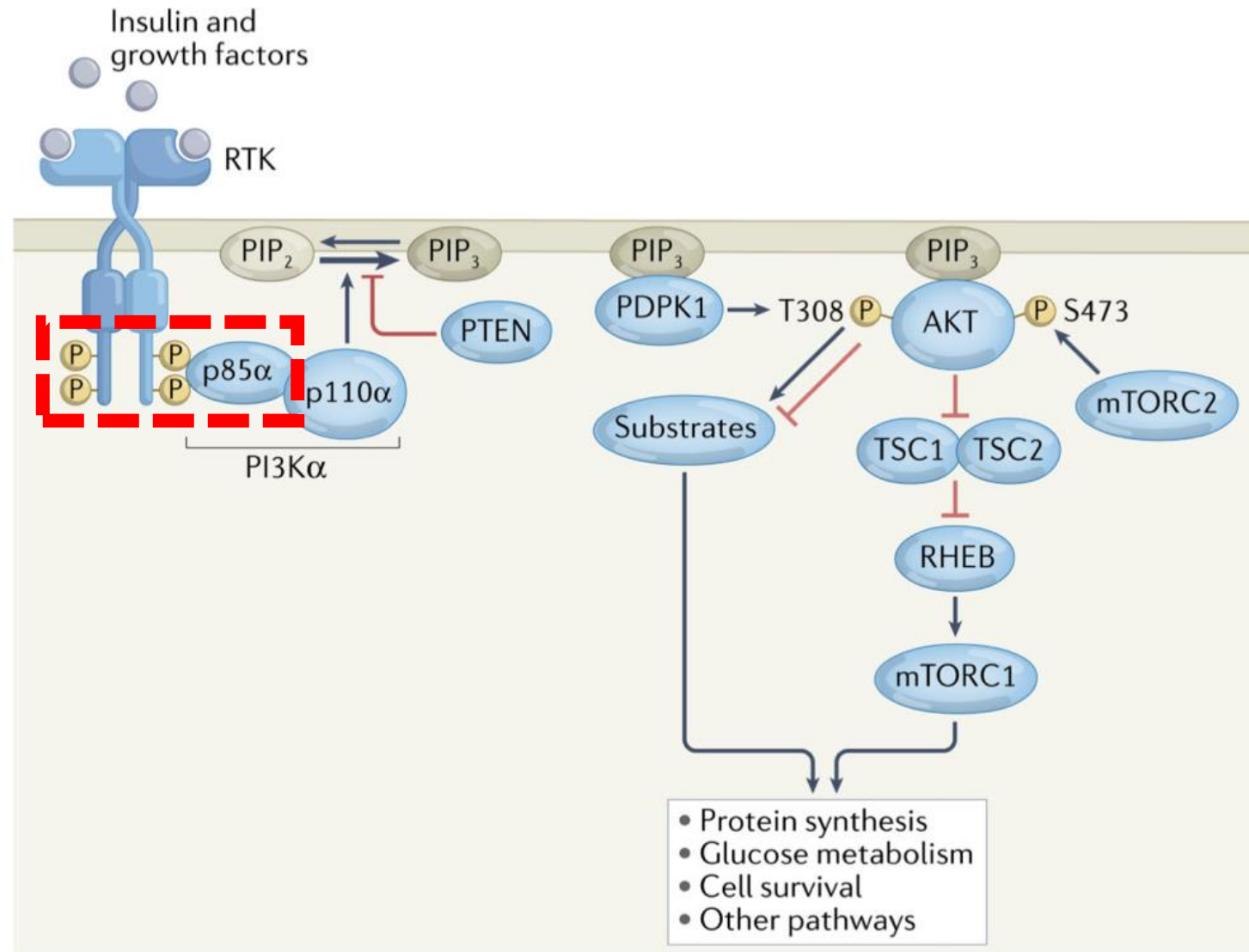


PI3K pathway in non-malignant cells

Insulin and other growth factors stimulate receptor tyrosine kinases (RTKs), leading to phosphorylation at C-terminal domain tyrosine residues.

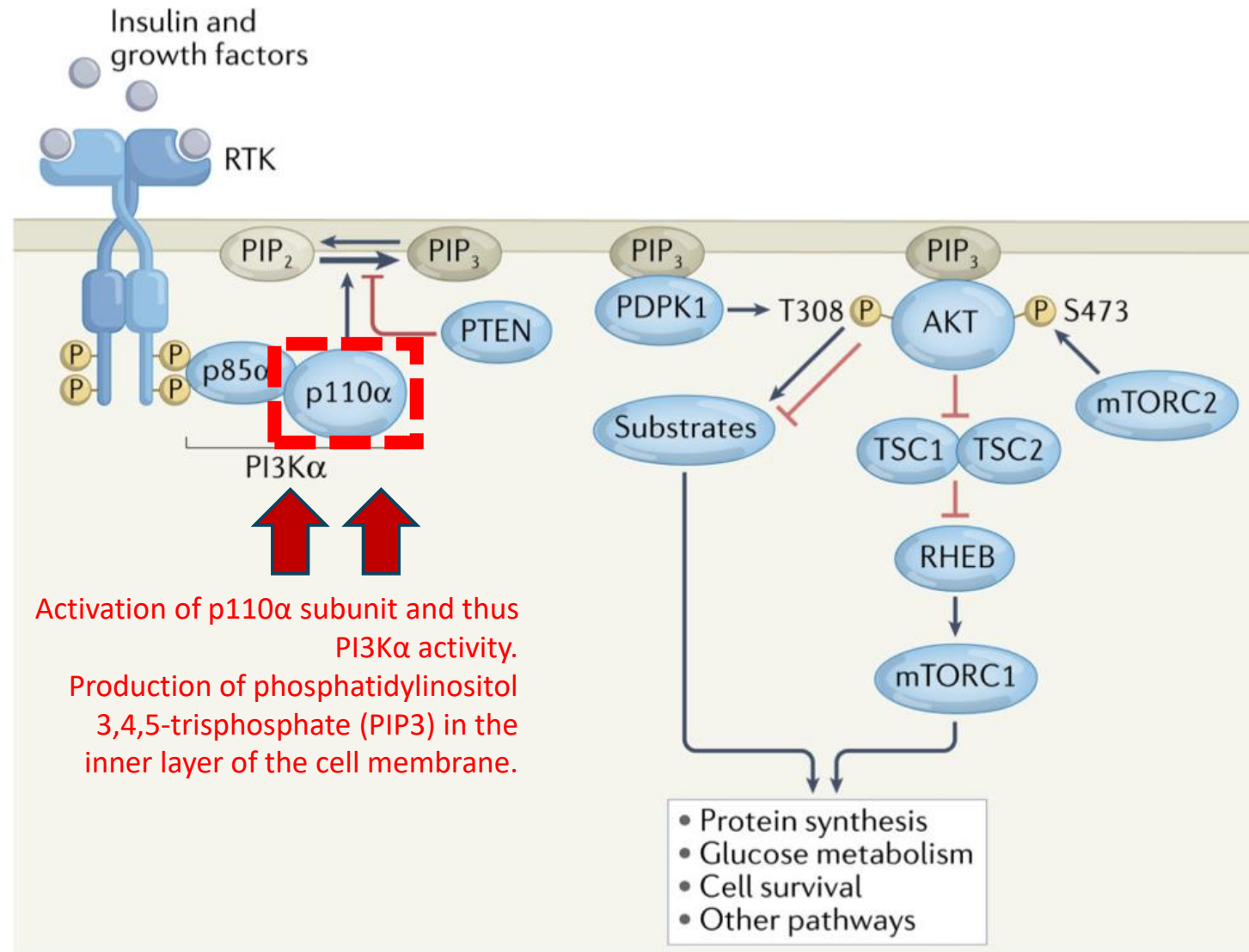


PI3K pathway in non-malignant cells

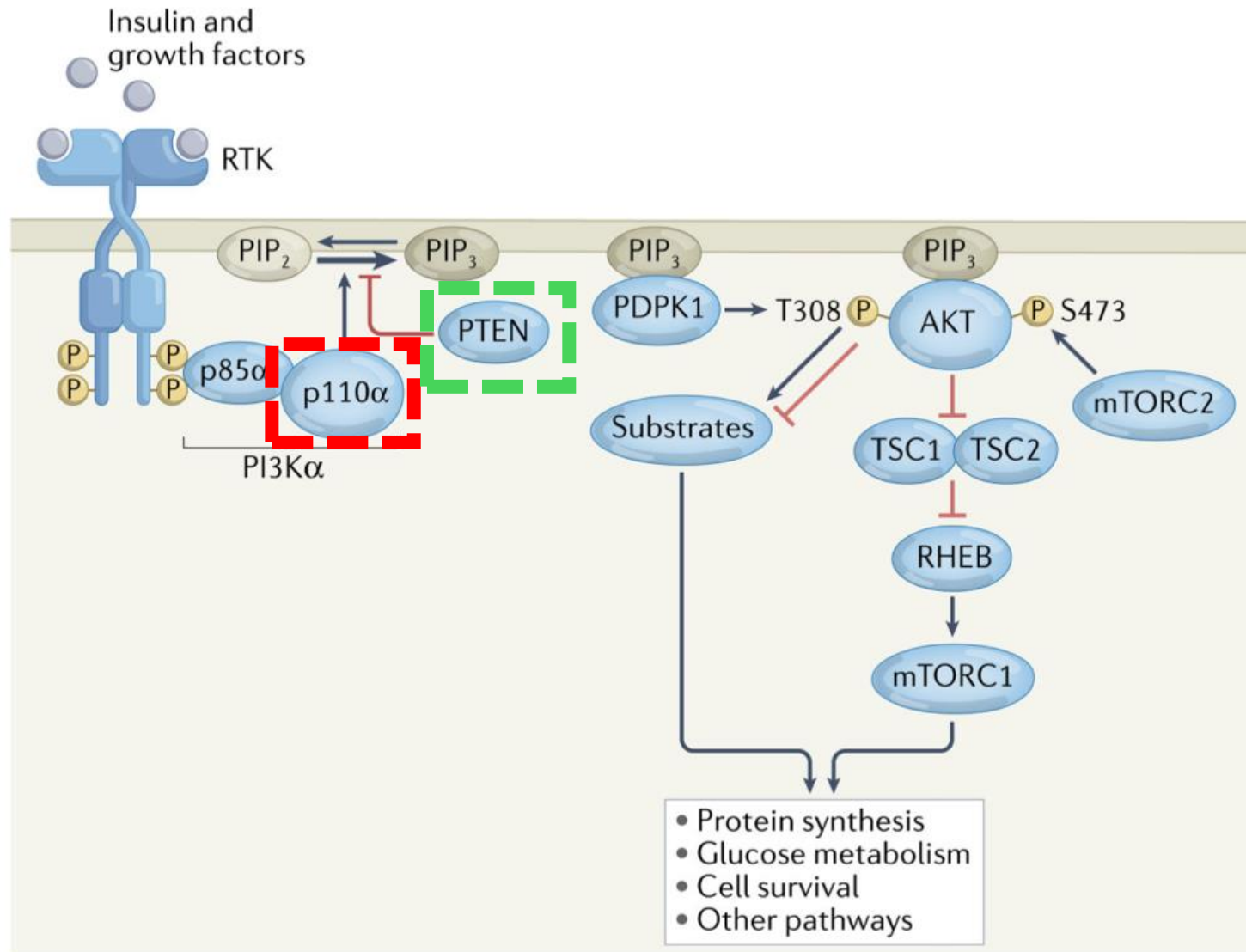


These phosphotyrosines bind to the regulatory p85α subunit of phosphatidylinositol 3-kinase-α (PI3Kα)

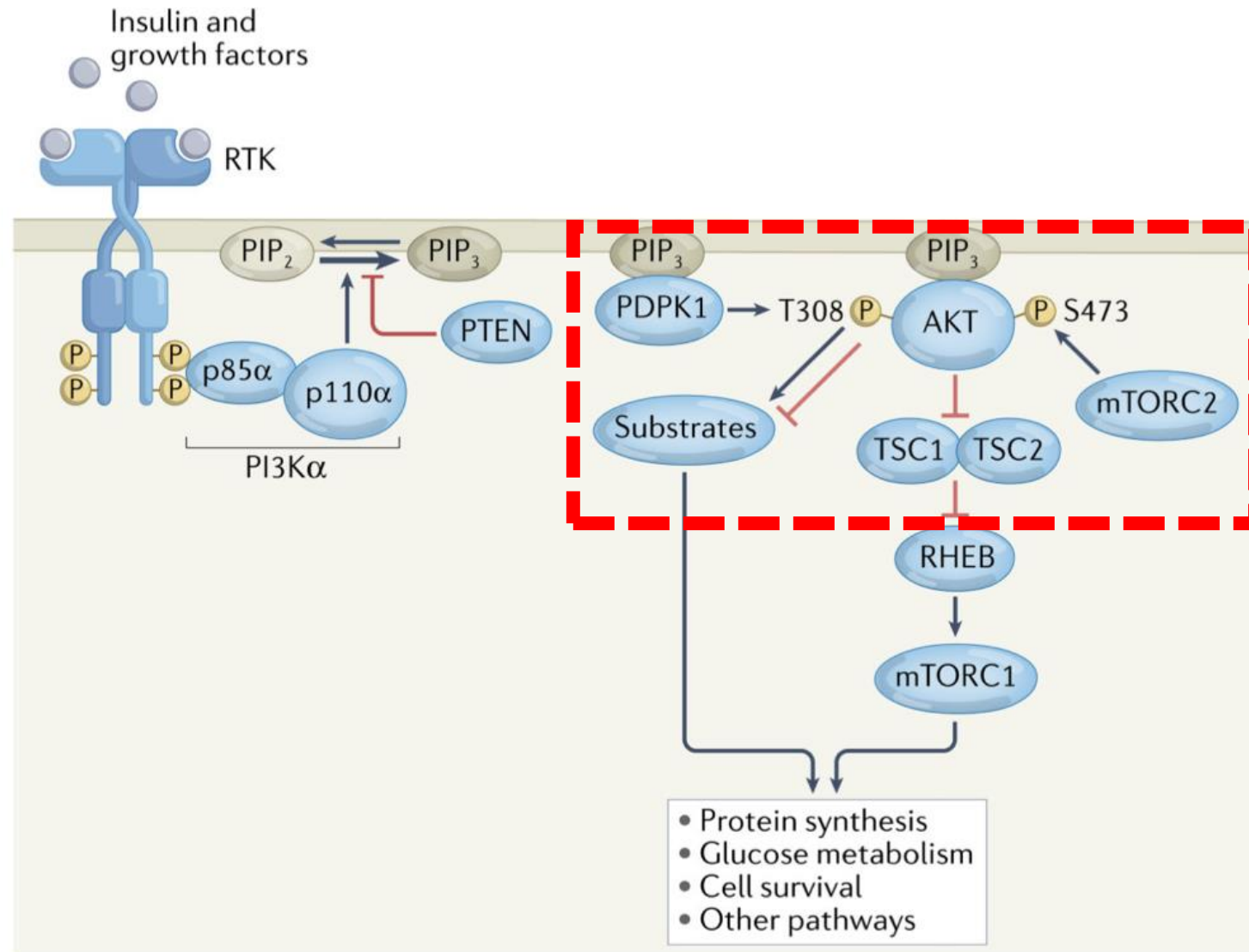
PI3K pathway in non-malignant cells



PI3K pathway in non-malignant cells



PI3K pathway in non-malignant cells



PIP₃ recruits the serine/threonine kinases PDK1 and AKT to the plasma membrane.

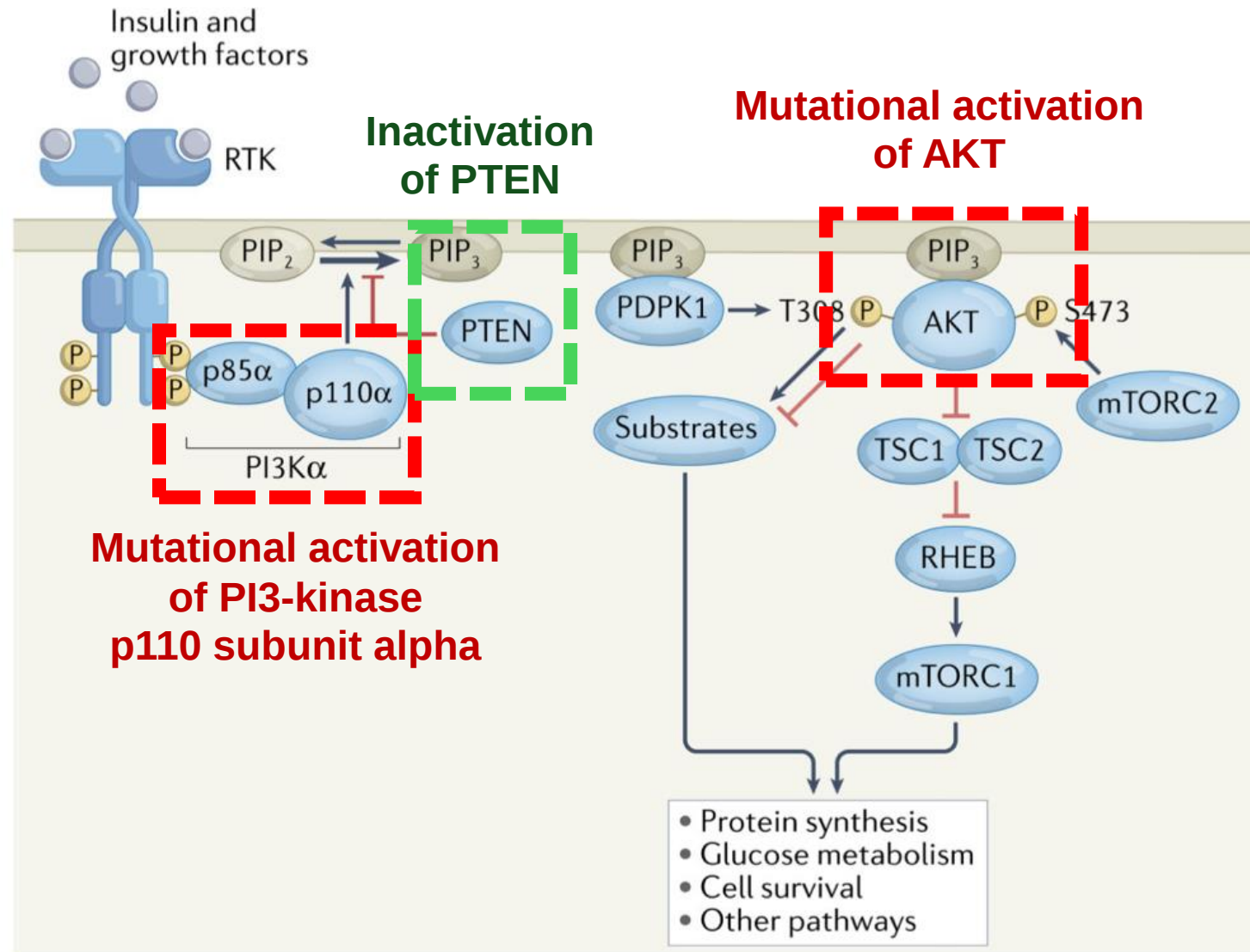
PDK1 and the mTOR complex 2 (mTORC2) phosphorylate AKT

AKT phosphorylates multiple cellular substrates, including tuberin (TSC2)

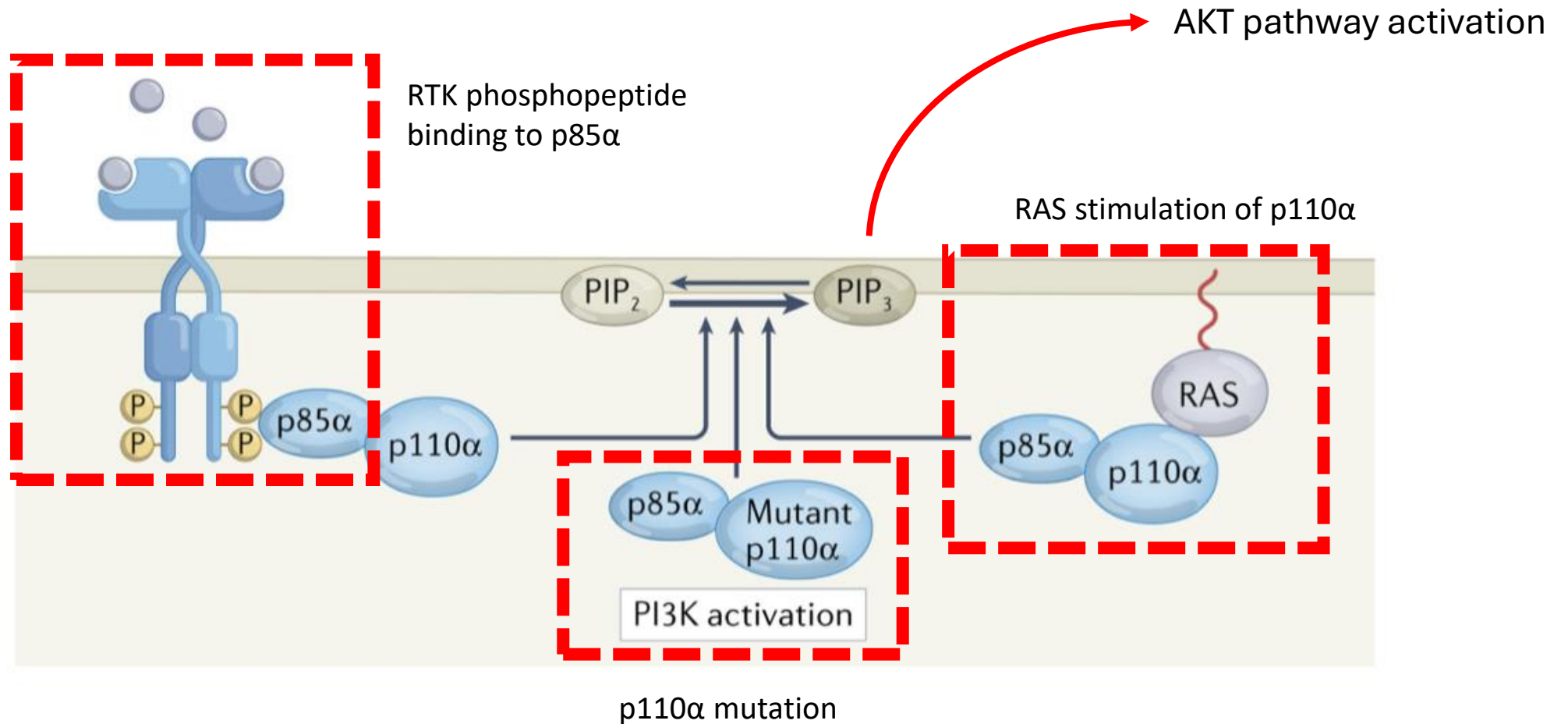
Activation of mTOR complex 1 (mTORC1)

- Protein synthesis
- Glucose metabolism
- Cell survival
- Other pathways

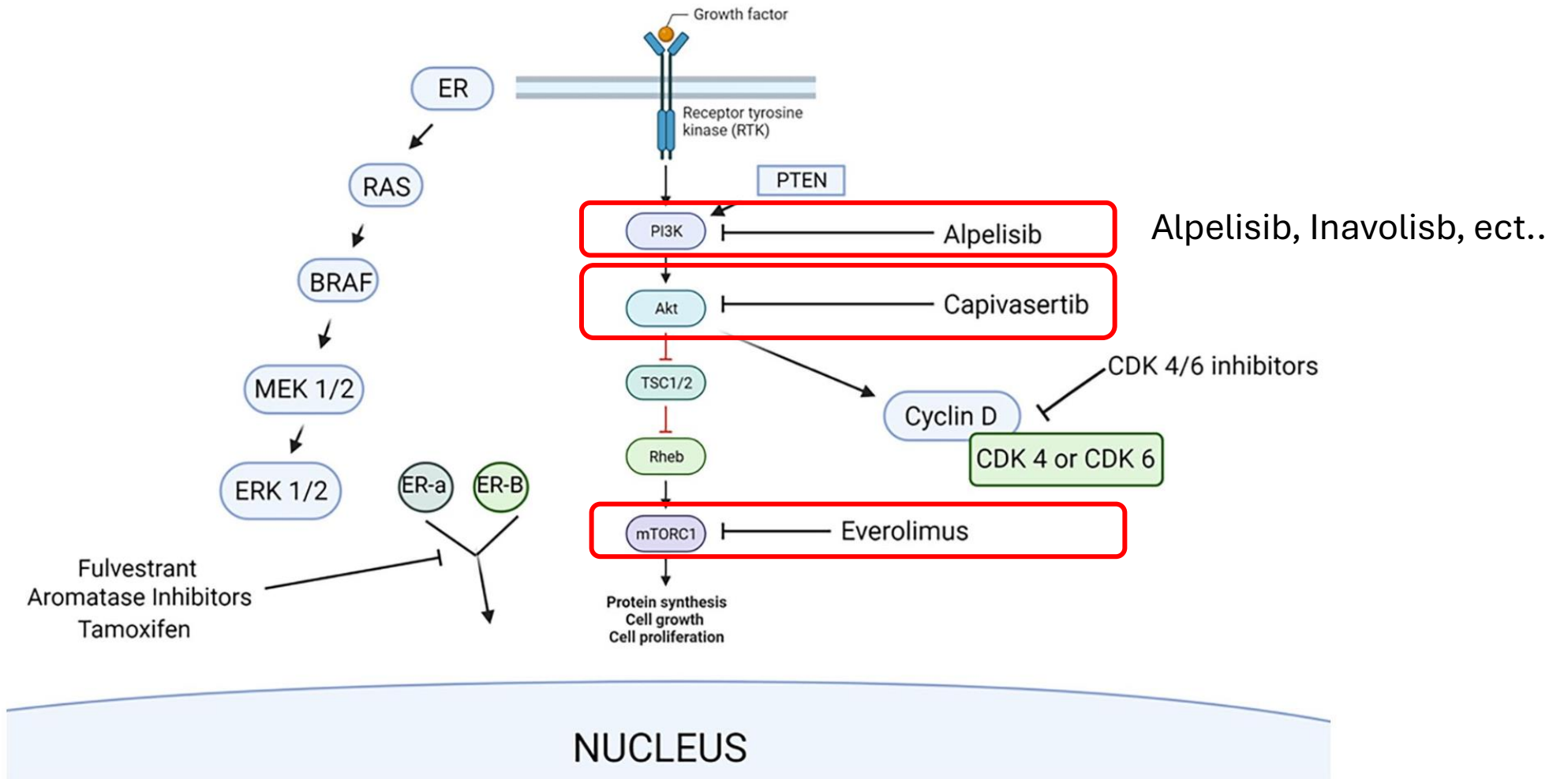
Disruption of the PI3K pathway in cancer



Disruption of the PI3K pathway in cancer



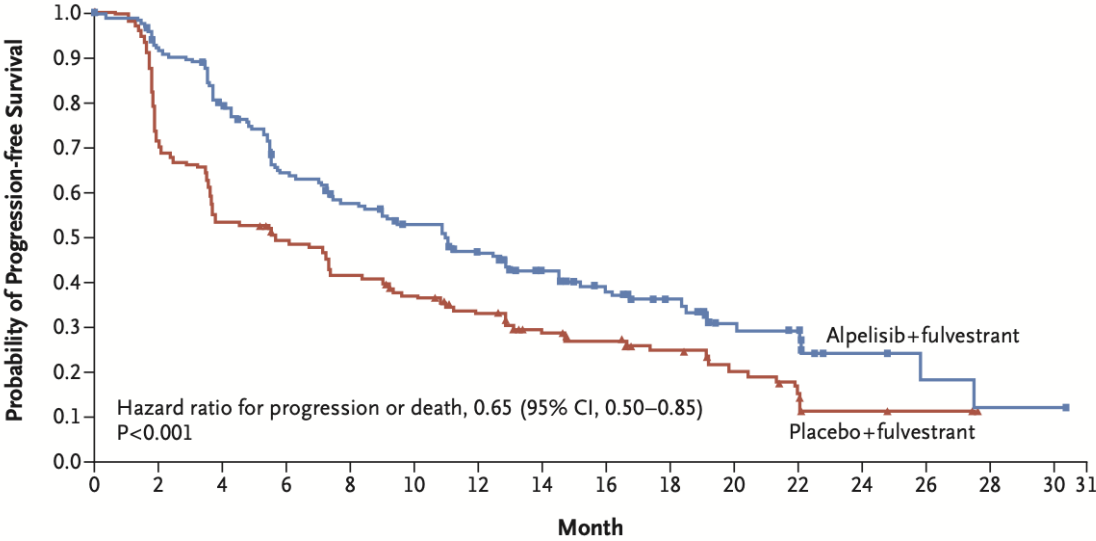
Target agents in PI3K pathway



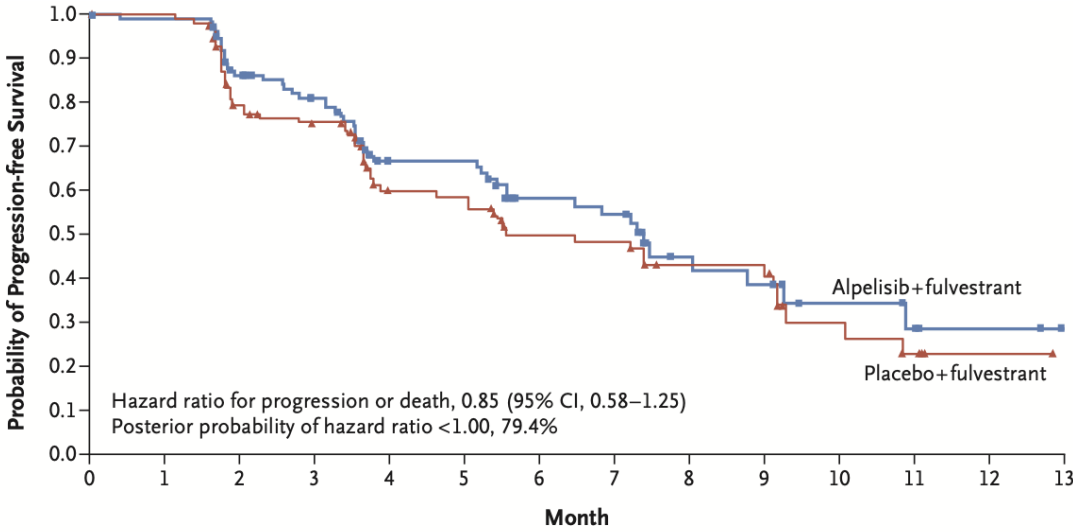
PFS with Fulvestrant + Alpelisib/placebo

SOLAR-1 trial

Cohort with *PIK3CA*-mutated cancer



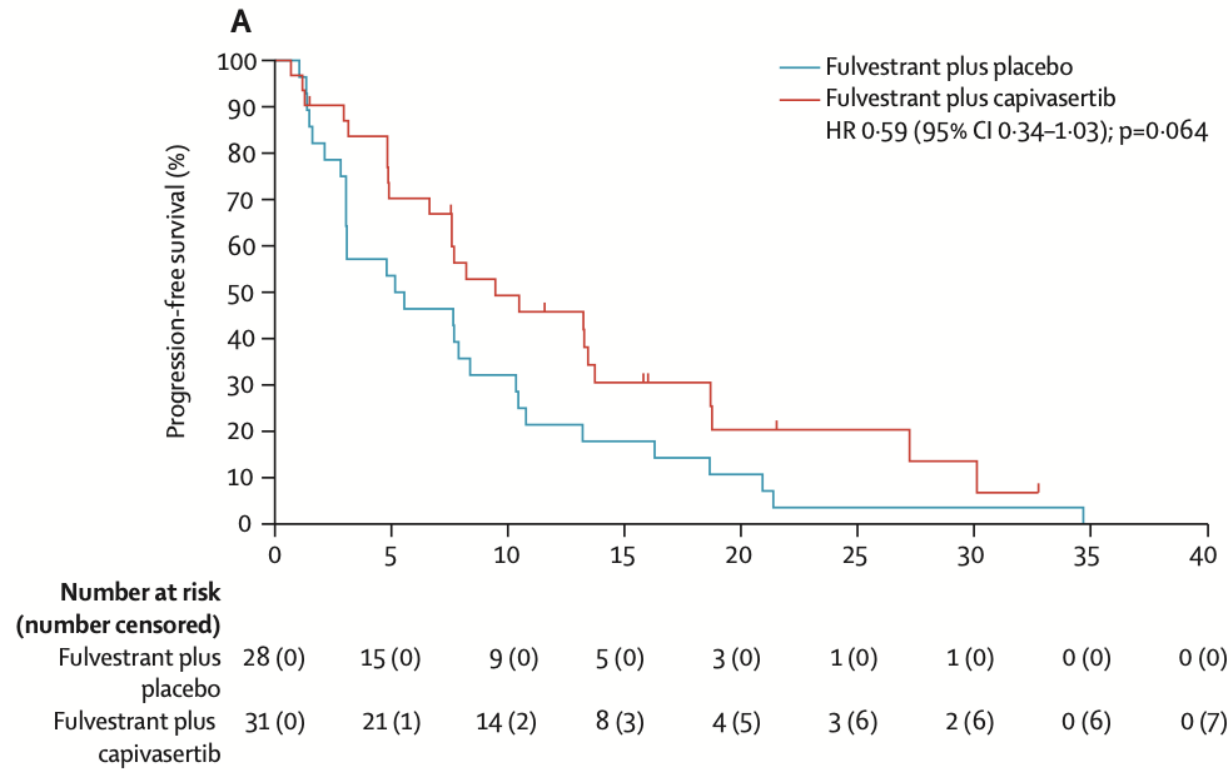
Cohort without *PIK3CA*-mutated cancer



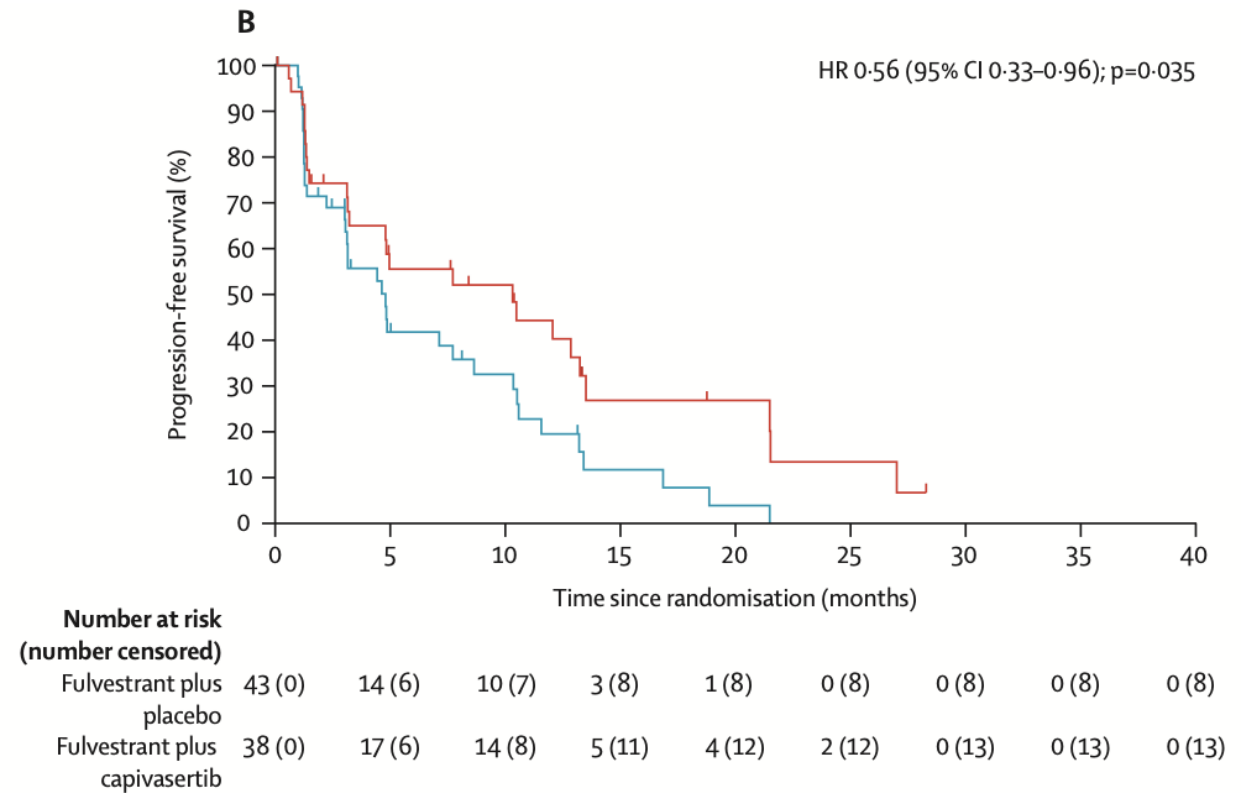
PFS with Fulvestrant + Capivasertib/placebo

FAKTION trial

PI3K pathway altered subgroup



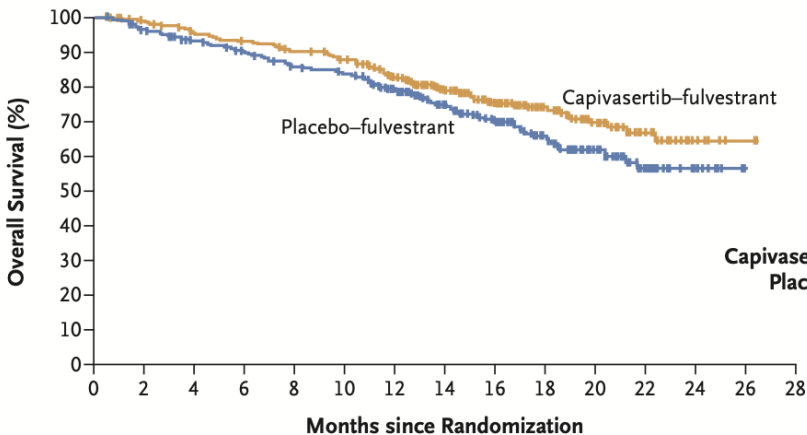
PI3K pathway non-altered subgroup



PFS with Fulvestrant + Capivasertib/placebo

CAPitello-291
trial

A Overall Population



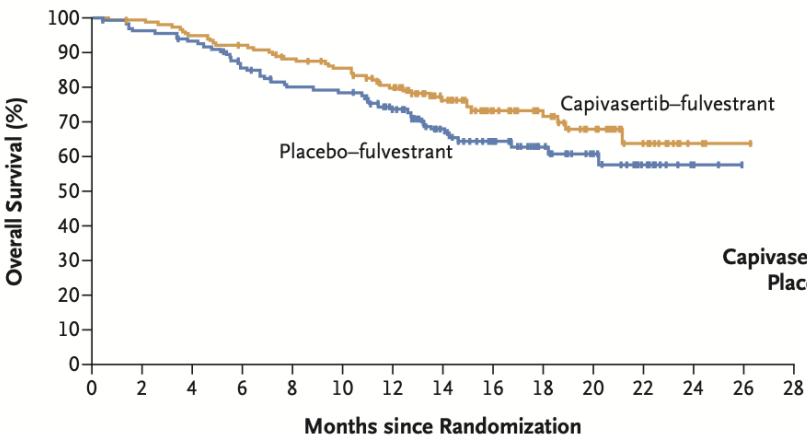
No. of Patients	No. of Deaths
355	87
353	108

Adjusted hazard ratio for death, 0.74 (95% CI, 0.56–0.98)

No. at Risk

Capivasertib-fulvestrant	355	343	327	318	306	295	258	198	144	95	63	33	9	2	0
Placebo-fulvestrant	353	334	316	301	283	274	237	181	134	90	59	30	11	0	0

B Patients with AKT Pathway-Altered Tumors



No. of Patients	No. of Deaths
155	41
134	46

Adjusted hazard ratio for death, 0.69 (95% CI, 0.45–1.05)

No. at Risk

Capivasertib-fulvestrant	155	153	144	139	131	125	111	83	60	45	30	14	3	1	0
Placebo-fulvestrant	134	127	122	112	101	99	87	62	46	31	22	13	3	0	0

PI3KCA mutation prevalence

Table 2 Description of Studies: Sample and Selected Demographics Relative to Clinical and Observational Studies of HR⁺/HER2⁻ mBC

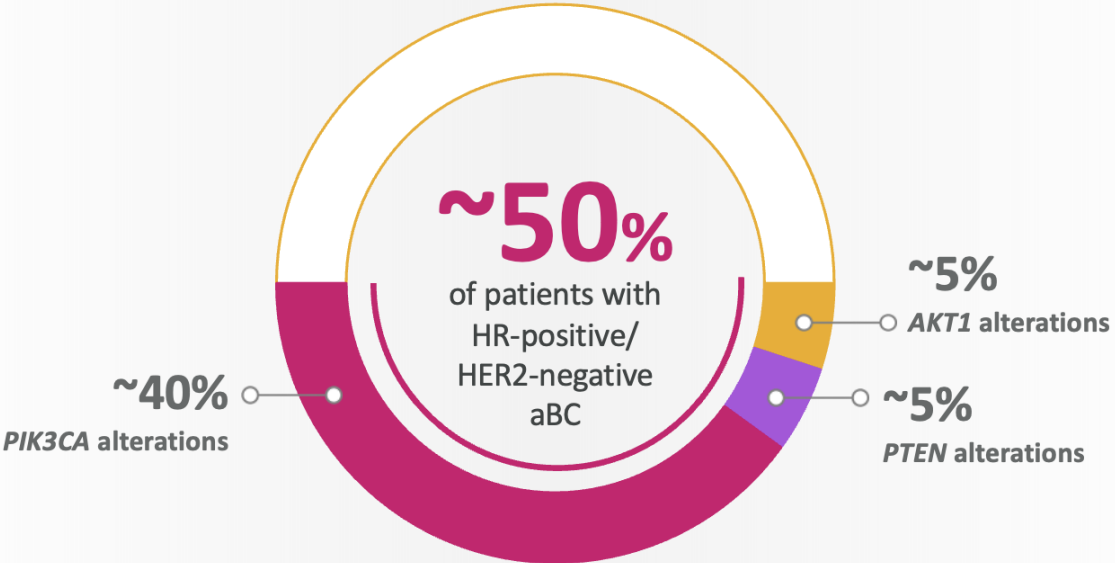
Study	Year	Study Phase	Type of Biopsy	Treatment Arms	HR ⁺ /HER2 ⁻ Sample Size Number	PIK3CA Mutant Number	PIK3CA Wild Type Number	PIK3CA Mutation Prevalence %	Median Age		ECOG PS, %				
									PIK3CA	Wild Type	0	1	2	3	Missing
Andre et al ¹⁸	2018	III	Tissue	Placebo plus fulvestrant (SOLAR-1)	288	172	116	NA ^a	64	63	192	95	NR	NR	1
Juric et al ²⁵	2018	III	Liquid	Alpelisib plus fulvestrant	284	169	115	NA ^a	63	62	196	86	NR	NR	2
Baird et al ²⁶	2016	Ib	Liquid	Taselisib plus tamoxifen	30	4	26	13.3	53	37	63				
Baselga et al ²⁷	2018	III		Taselisib plus fulvestrant (SANDPIPER)	340	340	NR	NA ^b	NR	NR	NR	NR	NR		
	2018	III		Placebo plus fulvestrant	176	176	NR	NA ^b	NR	NR	NR	NR	NR		
Baselga et al ¹⁶	2017	III	Tissue	Buparlisib plus fulvestrant (BELLE-2)	576	87 ^c	199	30.4	62	58	40	2	<1		
Campone et al ²⁸	2017	III	Tissue	Placebo plus fulvestrant	571	113 ^c	188	37.5	61	60	37	3	0		
Cristofanilli et al ²⁹	2016	III	Tissue	Palbociclib plus fulvestrant (PALOMA-3)	347	85 ^c	180	32.1	58	57	57	43			
Turner et al ³⁰	2016	III	Tissue	Placebo plus fulvestrant	174	44 ^c	86	33.8	56	55	63	35			
Di Leo et al ³¹	2018	III	Tissue	Buparlisib plus fulvestrant (BELLE-3)	289	100 ^c	132	43.1	60	60	39	1		1	
	2018	III	Tissue	Placebo plus fulvestrant	143	35 ^c	81	30.2	62	64	34	1		2	
Dickler et al ³²	2016	II	Tissue	Taselisib plus fulvestrant	60	31	29	51.7	61	62	56.7	43.3			
Fleming et al ³³	2012	II	Tissue	Temsirolimus	21	5	16	23.8	NR	NR	48	52			
Hortobagyi et al ³⁴	2017	III	Tissue	Ribociclib plus letrozole	212	69	143	32.5	NR	NR	NR	NR	NR	NR	
	2017	III	Tissue	Placebo plus letrozole	215	55	142	25.6	NR	NR	NR	NR	NR	NR	
Juric et al ²⁵	2019	Ib	Tissue	Alpelisib plus fulvestrant	87	52	33	59.8	58	NR	NR	NR	NR	NR	
Krop et al ³⁶	2016	II	Tissue	Pictilisib plus fulvestrant	89	38	45	42.7	60	68.5	31.5				
	2016	II	Tissue	Placebo plus fulvestrant	79	32	39	40.5	63	57	42				
Mayer et al ³⁷	2014	Ib	Tissue	Buparlisib plus letrozole (intermittent vs. continuous dosing)	51	16	35	31.4	55	NR	NR	NR	NR		
Mayer et al ³⁸	2016	Ib	Liquid	Alpelisib plus letrozole	26	16	10	61.5	53	NR	NR	NR	NR	NR	
Moynahan et al ³⁹	2017	III	Liquid	Everolimus plus exemestane	357	169	188	47.3	NR	NR	NR	NR	NR	NR	
	2017	III	Liquid	Placebo plus exemestane	193	69	124	35.8	NR	NR	NR	NR	NR	NR	

Abbreviations: BELLE-2 = Buparlisib Breast Cancer Clinical Evaluation-2; BELLE-3 = Buparlisib Breast Cancer Clinical Evaluation-3; ECOG PS = Eastern Cooperative Oncology Group performance status; HER2⁻ = human epidermal growth factor receptor 2-negative; HR⁺ = hormone receptor-positive; mBC = metastatic breast cancer; NR = not reported; PALOMA-3 = Palbociclib: Ongoing Trials in the Management of Breast Cancer; PIK3CA = PIK3 110α subunit, phosphatidylinositol-4,5- bisphosphate 3-kinase catalytic subunit alpha; SANDPIPER = A Study of Taselisib + Fulvestrant Versus Placebo + Fulvestrant in Participants With Advanced or Metastatic Breast Cancer Who Have Disease Recurrence or Progression During or After Aromatase Inhibitor Therapy.

^aPIK3CA mutation prevalence in SOLAR-1 is not reportable as such because patients were selected based on PIK3CA mutation status to enrich for tumors with a mutation.

^bPIK3CA mutation prevalence in SANDPIPER is not reportable because only data on the PIK3CA population was reported at time of this publication.

^cOnly a subset of the total sample of HR⁺/HER2⁻ sample was tested for the PIK3CA mutation.



Courtesy of AstraZeneca Medical
Mollon LE et al. *Clin Breast Cancer*. 2020;20(3):e232-e243

Cancer Genome Atlas Network. *Nature*. 2012;490:61–70; Martorana F et al. *Front Pharmacol*. 2021;12:662232; Miricescu D et al. *Int J Mol Sci*. 2020;22(1):173; Smyth LM et al. *Cancer Discov*. 2020;10(4):526–535; Paplomata E, O’Regan R. *Ther Adv Med Oncol*. 2014;6(4):154–166.

New ET options for ET-resistant disease (biomarker selected)

	Primary	Secondary	Sensitive
eBC	Relapse on first 2 yrs of adj ET	Relapse <1y off adjuvant ET	Relapse > 1y off adj ET/ De novo mBC
mBC	PD ≤24 wks from start of ET for mBC	PD >24 wks from start of ET for mBC	PFS >12 mo with 1L ET+ CDK4/6is*

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Trial	Design	N.	Biomarker	Prior CDK %	Primary res %	<1y of adj ET	PFS mo.	HR
EMERALD ¹	SoC ET Elacestrant	228	ESR1	100	~ 10 (ITT) 0 (ESR1mut)	-	1.9	
							3.8	0.55 (0.39-0.77)
EMBER-3 ²	SoC ET Imlunestrant	661	ESR1	~ 60	~ 9 (ITT) 0 (ESR1mut)	~ 30 (ITT) 20 (ESR1mut)	3.8	
							5.5	0.62 (0.46-0.82)
CAPItello-291 ³	Ful + Placebo Ful+Capivasertib	289	AKT path	~ 70	~ 40	-	3.1	
							7.3	0.50 (0.38-0.65)
INAVO-120 ⁴	SoC ET Elacestrant	228	PIK3CA	1	34	66	7.3	
							15	0.43 (0.32-0.59)
PALOMA-3 ⁵ MONALEESA-3 ⁶ MONARCH-2 ⁷	Fulv + Placebo Ful+CDK4/6is	-	-	0	- - ~ 25	~ 5 ~ 5 -	-	-

¹Bidard FC K et al, JCO 2022; ²Jhaveri KL et al, NEJM 2024; ³Turner NC et al, NEJM 2023; ⁴Turner NC et al, NEJM 2024 ; ⁵Cristofanilli M et al, Lancet Oncol 2016; ⁶Slamon DJ et al, JCO 2018; ⁵Sledge GW et al, JCO 2017

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¹Bidard FC K et al, JCO 2022; ²Jhaveri KL et al, NEJM 2024; ³Turner NC et al, NEJM 2023; ⁴Turner NC et al, NEJM 2024 ; ⁵Cristofanilli M et al, Lancet Oncol 2016; ⁶Slamon DJ et al, JCO 2018; ⁷Sledge GW et al, JCO 2017

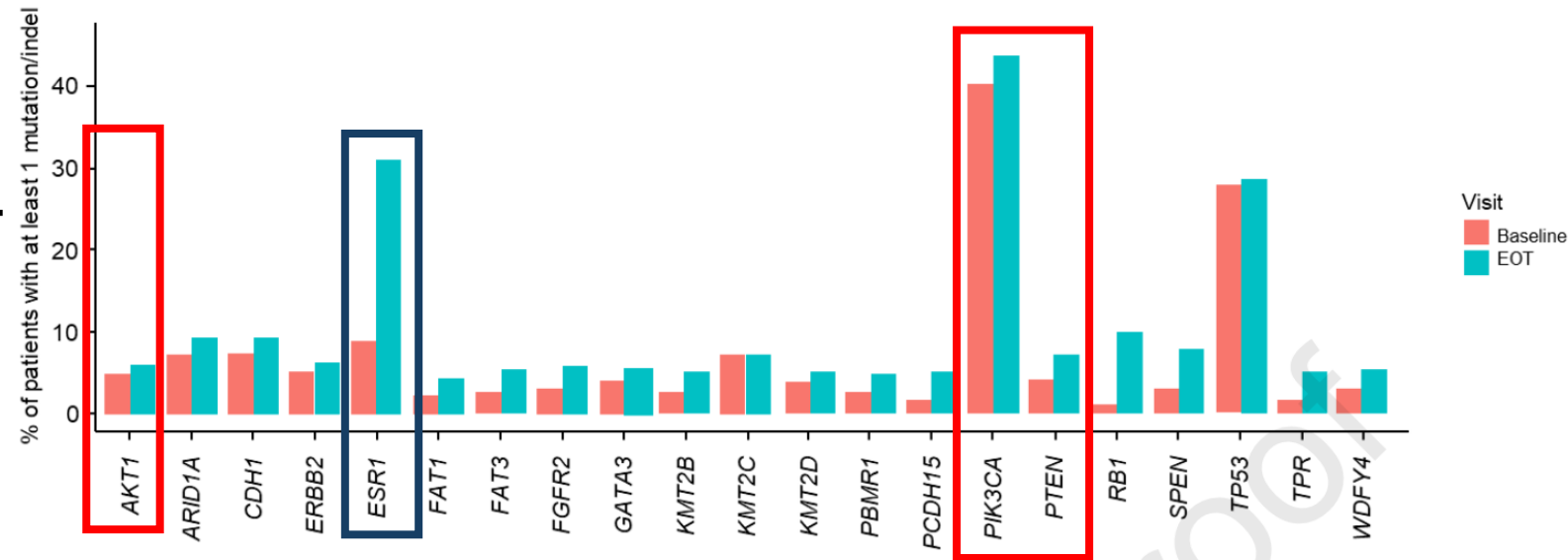
Targeting AKT pathway in ET-resistant disease: CAPItello-291

Alteration; n (%)		Capivasertib + fulvestrant (N=355)	Placebo + fulvestrant (N=353)
Any AKT pathway alteration		155 (43.7)	134 (38.0)
PIK3CA	Any	116 (32.7)	103 (29.2)
	PIK3CA only	110 (31.0)	92 (26.1)
	PIK3CA and AKT1	2 (0.6)	2 (0.6)
	PIK3CA and PTEN	4 (1.1)	9 (2.5)
AKT1 only		18 (5.1)	15 (4.2)
PTEN only		21 (5.9)	16 (4.5)
Non-altered		200 (56.3)	219 (62.0)
AKT pathway alteration not detected		142 (40.0)	171 (48.4)
Unknown		58 (16.3)	48 (13.6)
No sample available		10 (2.8)	4 (1.1)
Preanalytical failure		39 (11.0)	34 (9.6)
Post analytical failure		9 (2.5)	10 (2.8)

AKT pathway alteration status was determined centrally using next-generation sequencing **in tumor tissue** with the **FoundationOne®CDx assay** (and Burning Rock assay in China)

Prevalence of gene alterations at baseline vs EOT with ribociclib plus ET vs placebo plus ET

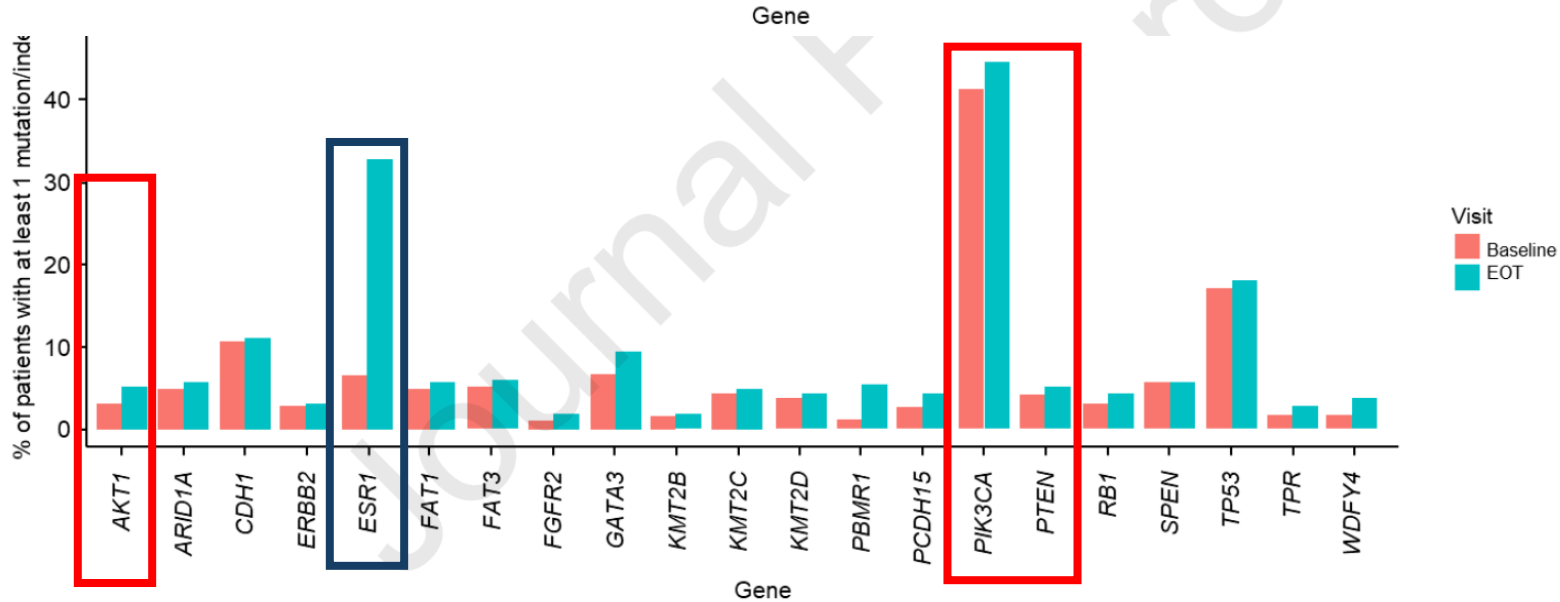
Ribo + ET



No difference in ESR1m with Ribo use

PI3KCA remains stable in time

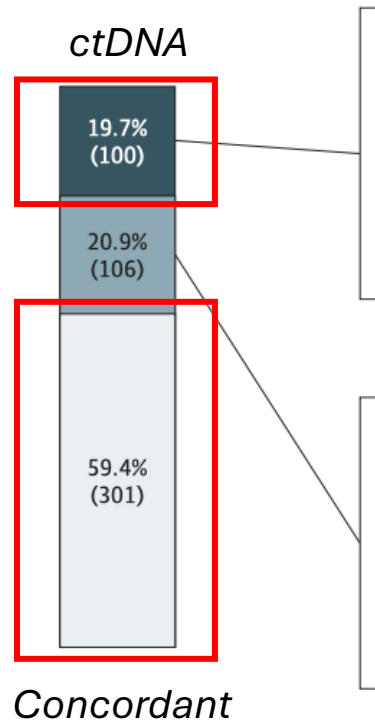
ET + Placebo



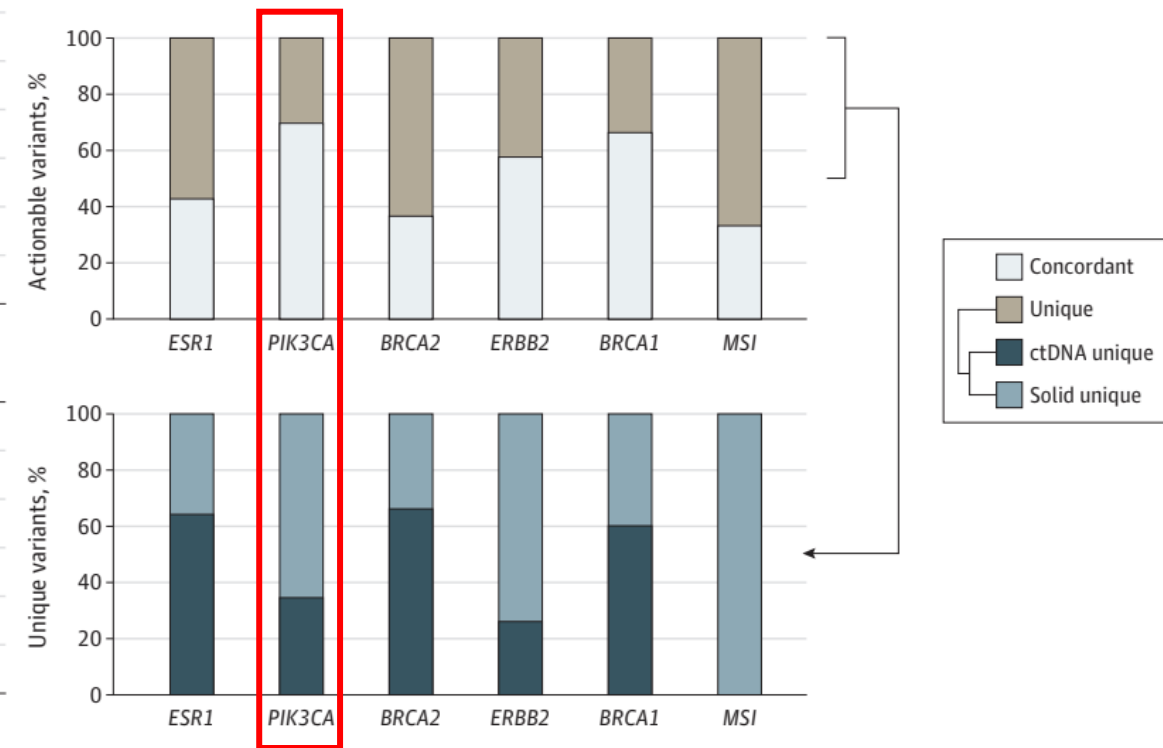
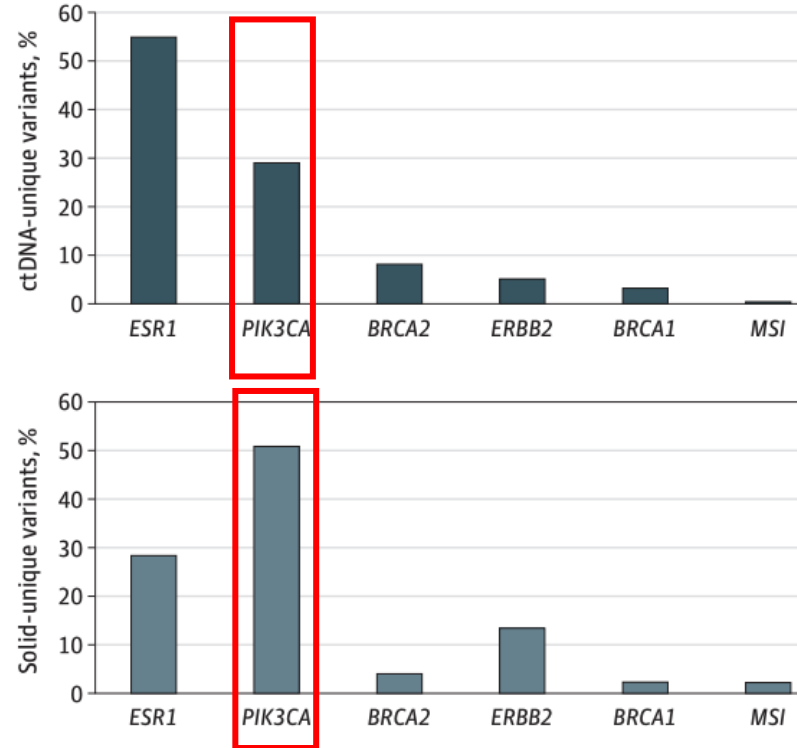
Among Ribo users only **RB1** and **SPEN** showed differences

Liquid Biopsy or tissue Biopsy?

A Distribution of actionable variants

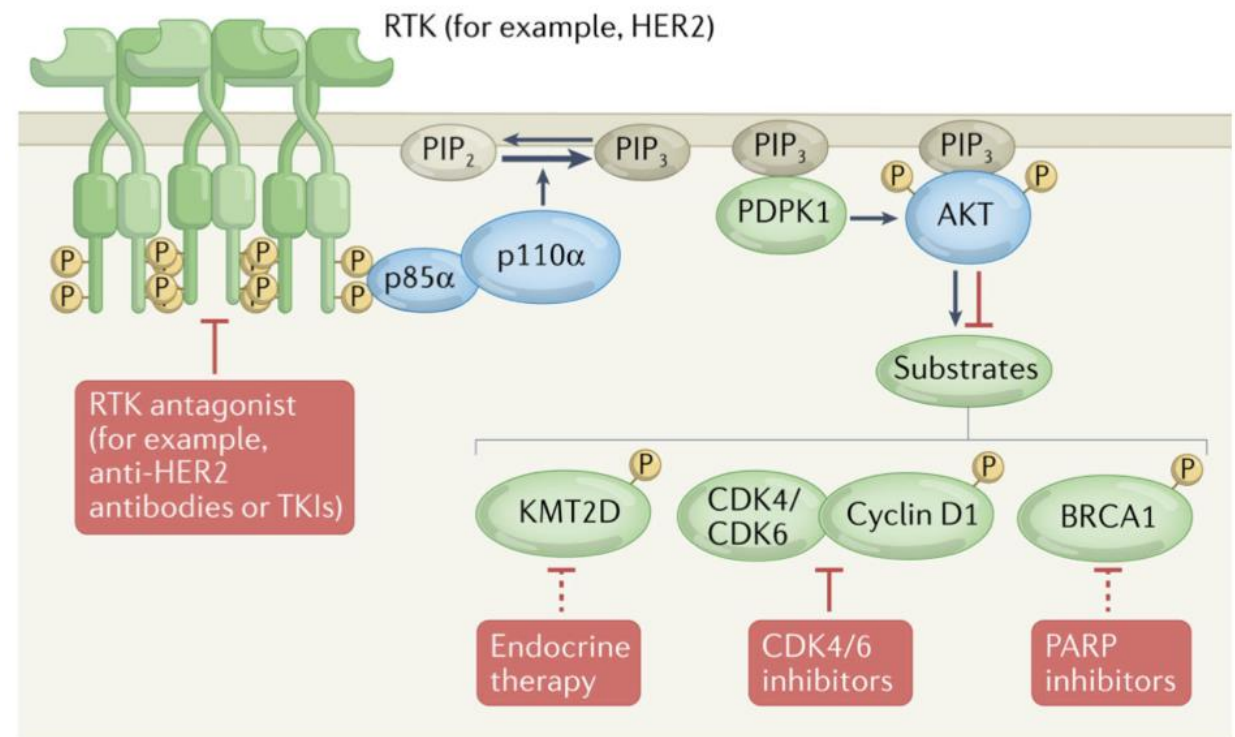
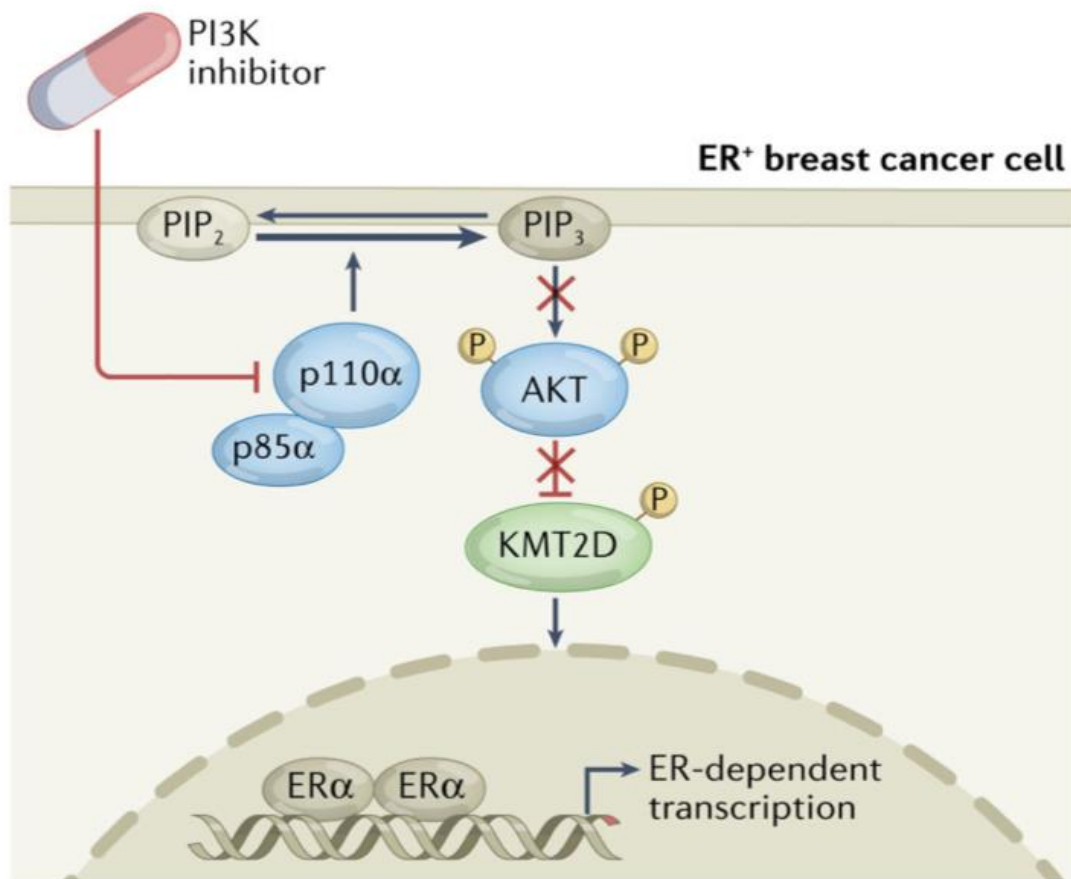


B Unique variants detected in genes

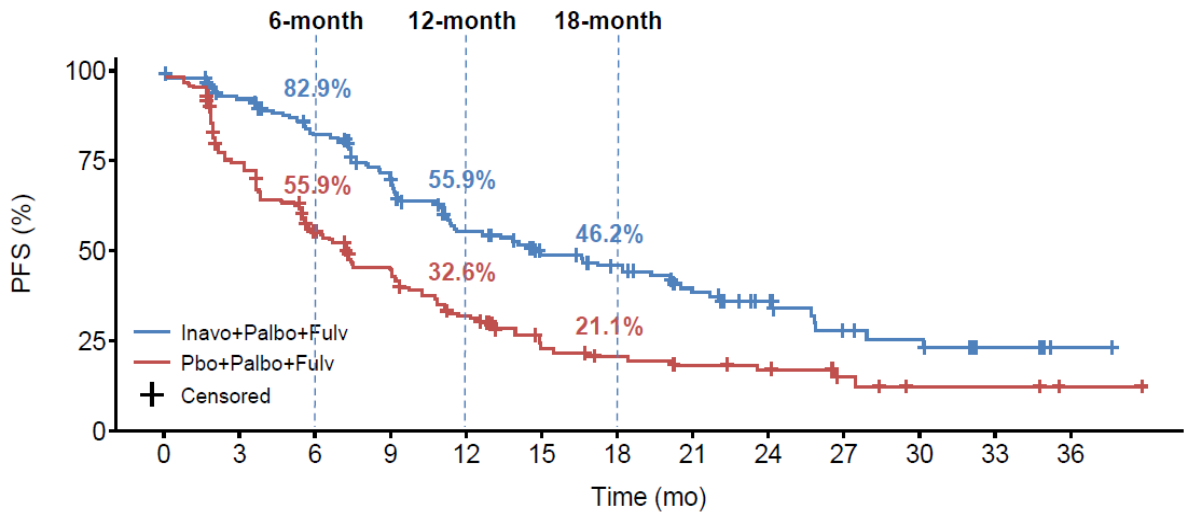


- **60 -80% concordance** between **tissue** and **ctDNA**
- **High ctDNA** identification of **actionable biomarkers**
- **19.7%** variants detected uniquely by **ctDNA** → **55% ESR1**

PI3K mechanism of resistance in breast cancer



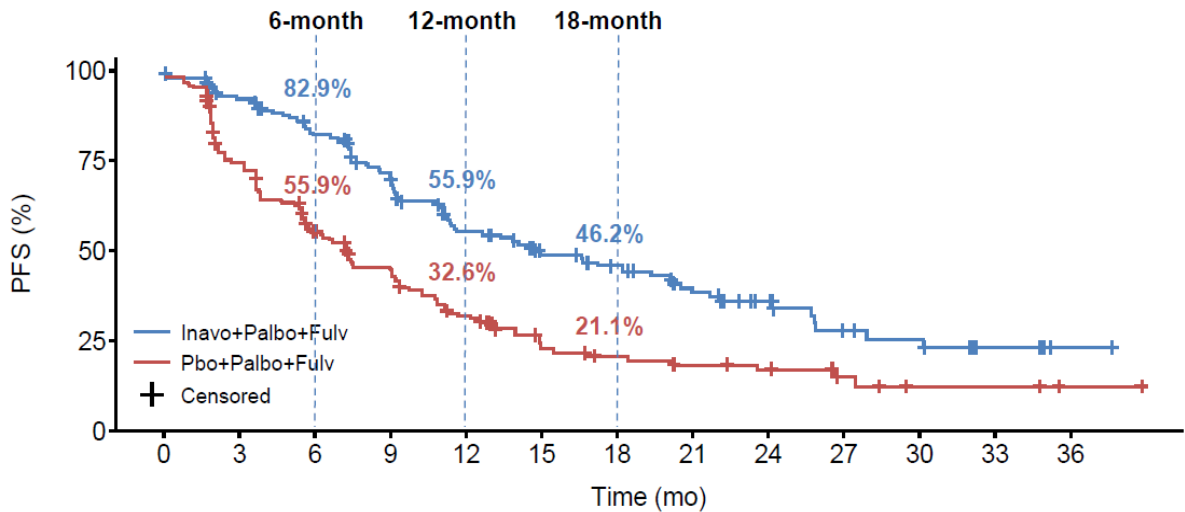
Targeting PIK3CA + CDK4/6is in ET-resistant disease: INAVO-120



161	134	111	92	66	48	41	31	22	13	11	5	1
164	113	77	59	40	23	19	16	12	6	3	3	1
Inavo+Palbo+Fulv Pbo+Palbo+Fulv												
(n=161) (n=164)												
No. of events, n (%)						82 (50.9)			113 (68.9)			
Median (95% CI), mo						15.0 (11.3, 20.5)			7.3 (5.6, 9.3)			
Stratified hazard ratio (95% CI)						0.43 (0.32, 0.59)						
						p<0.0001						

	Inavo+Palbo+Fulv		Pbo+Palbo+Fulv			Hazard ratio (95% CI)
All patients	n	Median (mo)	n	Median (mo)		
161	15.0	164	7.3			0.50* (0.38, 0.67)
Visceral disease						
No	29	25.8	36	7.4		0.43 (0.19, 0.97)
Yes	132	13.8	128	7.2		0.51 (0.38, 0.69)
Liver metastasis at enrollment						
No	84	24.2	73	11.3		0.56 (0.35, 0.90)
Yes	77	11.0	91	5.6		0.48 (0.33, 0.69)
Number of metastatic organs at enrollment						
1	21	20.2	32	7.4		0.35 (0.14, 0.87)
2	59	18.2	46	7.4		0.47 (0.29, 0.77)
>3	81	14.1	86	7.3		0.55 (0.37, 0.80)
Endocrine resistance						
Primary	53	11.4	58	3.7		0.39 (0.24, 0.61)
Secondary	108	18.2	105	9.7		0.55 (0.38, 0.80)

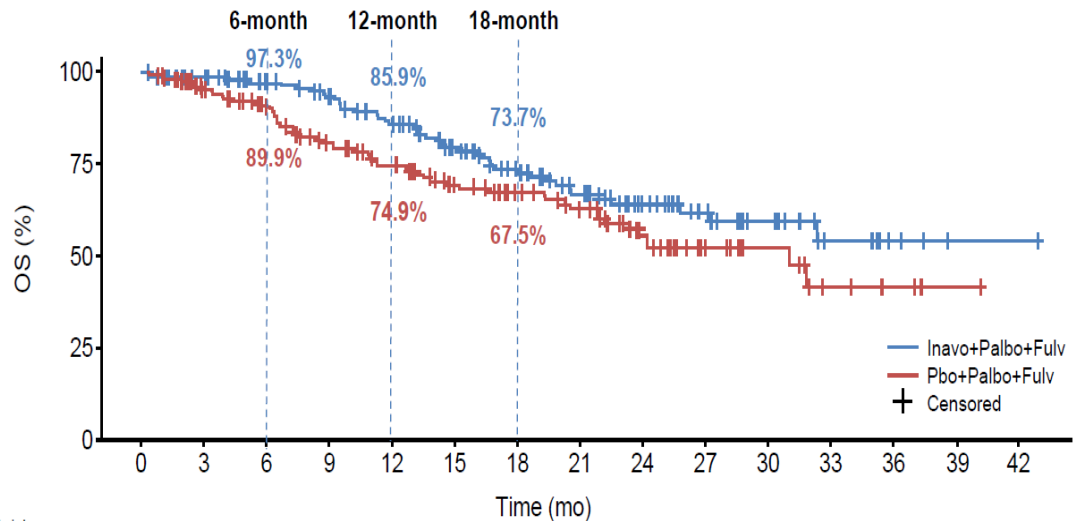
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	Inavo+Palbo+Fulv (n=161)	Pbo+Palbo+Fulv (n=164)
No. of events, n (%)	82 (50.9)	113 (68.9)
Median (95% CI), mo	15.0 (11.3, 20.5)	7.3 (5.6, 9.3)
Stratified hazard ratio (95% CI)	0.43 (0.32, 0.59) p<0.0001	

	Inavo+Palbo+Fulv		Pbo+Palbo+Fulv			Hazard ratio (95% CI)
All patients	n	Median (mo)	n	Median (mo)		
	161	15.0	164	7.3		0.50* (0.38, 0.67)
Visceral disease						
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Liver metastasis at enrollment						
No	84	24.2	73	11.3		0.56 (0.35, 0.90)
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Endocrine resistance						
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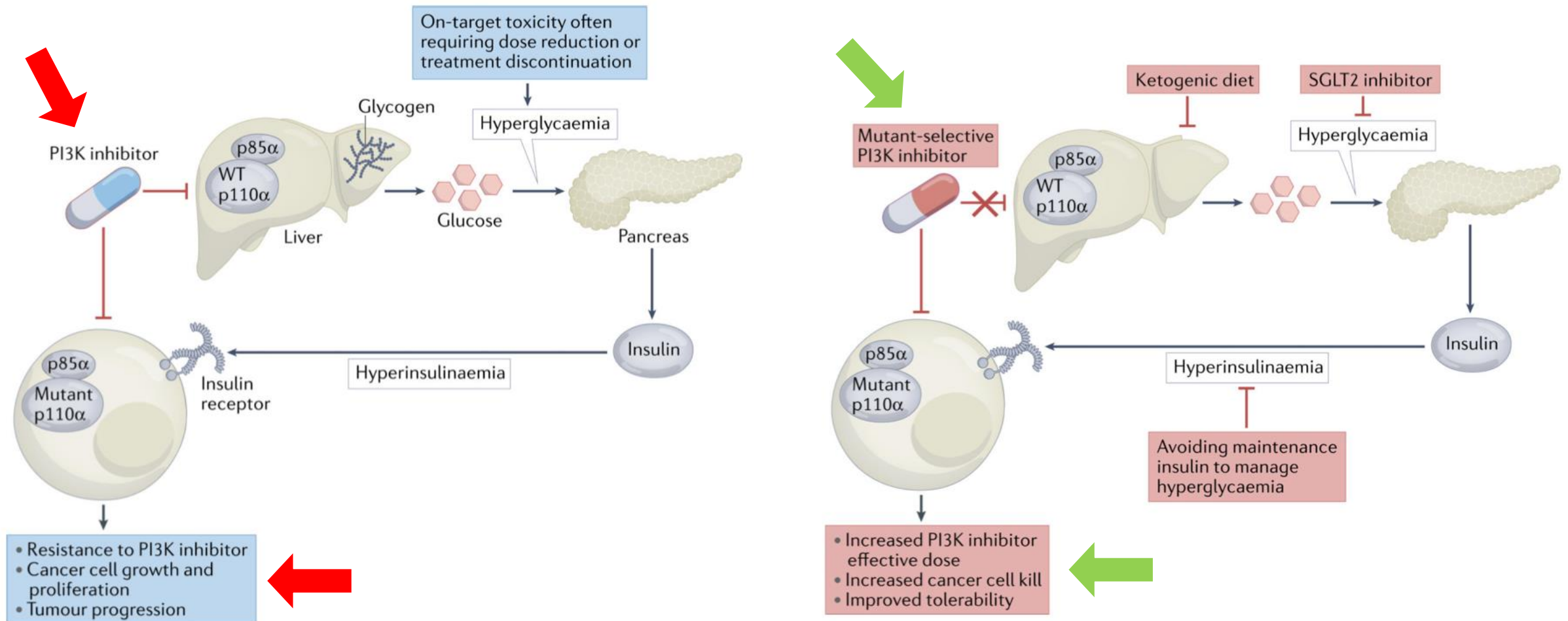


Patients at risk:	161	143	127	114	101	85	69	56	38	26	17	8	4	1	1	Median follow-up: 21.3 months
Inavo+Palbo+Fulv	164	139	120	98	87	72	61	52	33	19	11	5	3	1	0	
Pbo+Palbo+Fulv																

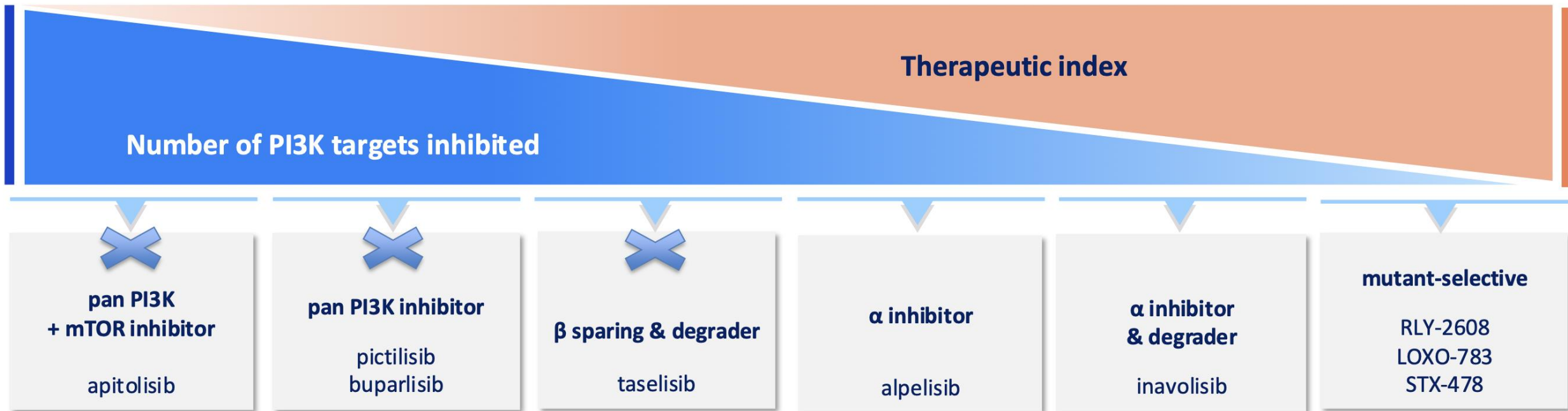
The pre-specified boundary for OS (p of 0.0098 or HR of 0.592) was not crossed at this interim analysis

	Inavo+Palbo +Fulv (n=161)	Pbo+Palbo +Fulv (n=164)
No. of events, n (%)	42 (26.1)	55 (33.5)
Median (95% CI), mo	NE (27.3, NE)	31.1 (22.3, NE)
Stratified Hazard Ratio (95% CI)	0.64 (0.43, 0.97) p=0.0338	

PI3K inhibitors effect on metabolism



PIK3 target inhibitors development





Grazie per l'attenzione

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