

The difficult targets - PIK3CA

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Potential conflicts of interest

Employment and Leadership

- Universität Duisburg-Essen, Universitätsklinikum Essen, Ruhrlandklinik

Scientific Advice

- AstraZeneca, Boehringer Ingelheim, Bristol-Myers Squibb, Celgene, Institut für Qualität und Wirtschaftlichkeit im Gesundheitswesen (IQWiG), Lilly, Novartis

Stock Ownership

- none

Honoraries for Educational Lectures

- Alexion, Boehringer Ingelheim, Bristol-Myers Squibb, Celgene, Lilly, Novartis

Research Grants to Institution

- Boehringer Ingelheim, Bristol-Myers Squibb, Novartis

Expert Testimony

- Medical Boards, Courts, Public Funding Agencies, Universities

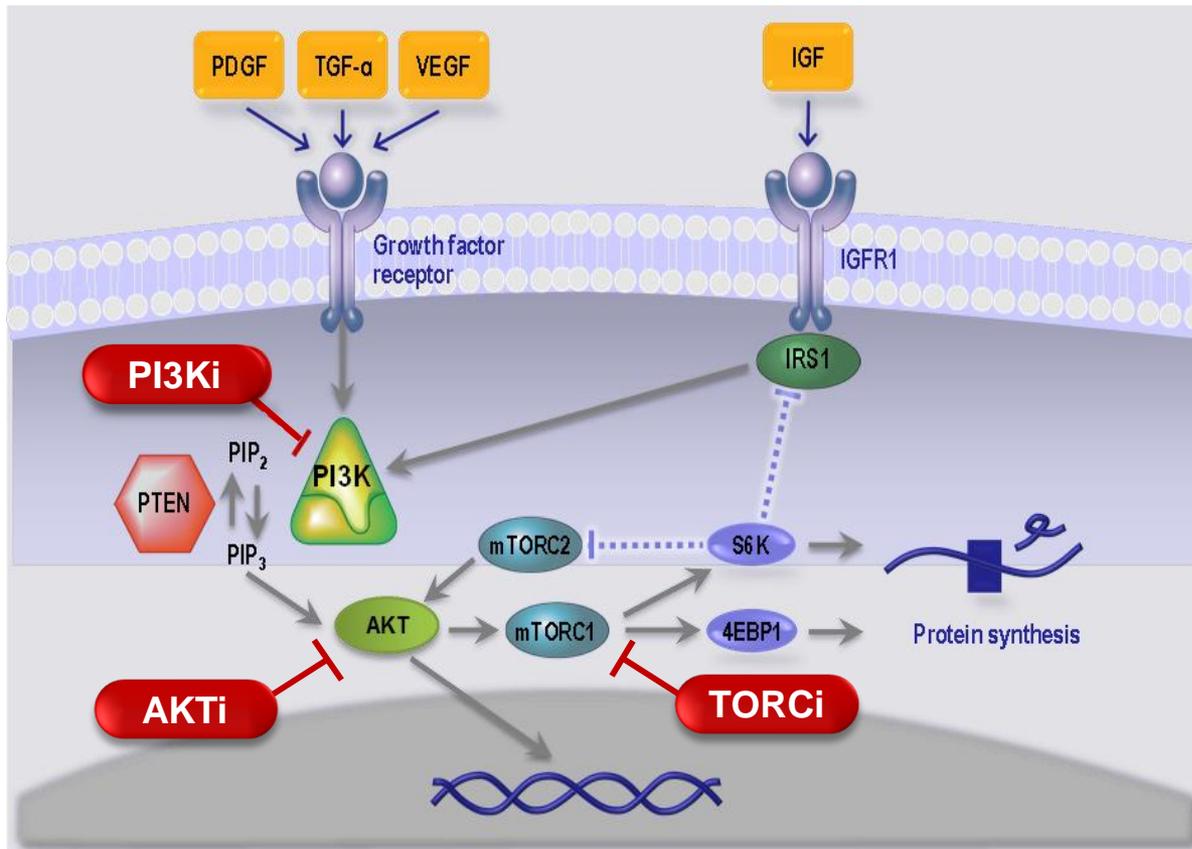
Others

- Universität Duisburg-Essen (Patents)



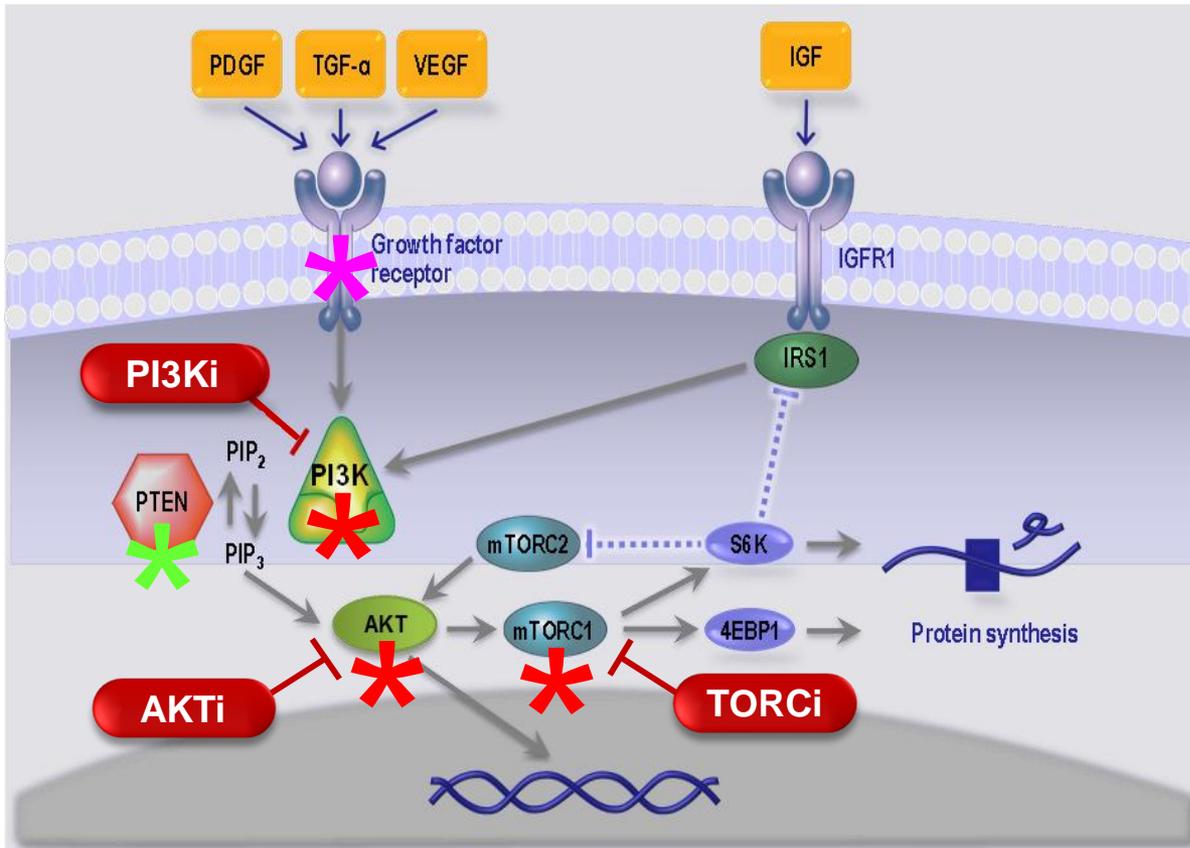
Phosphoinositide-3-kinase signaling pathway

Targeted agents in clinical development

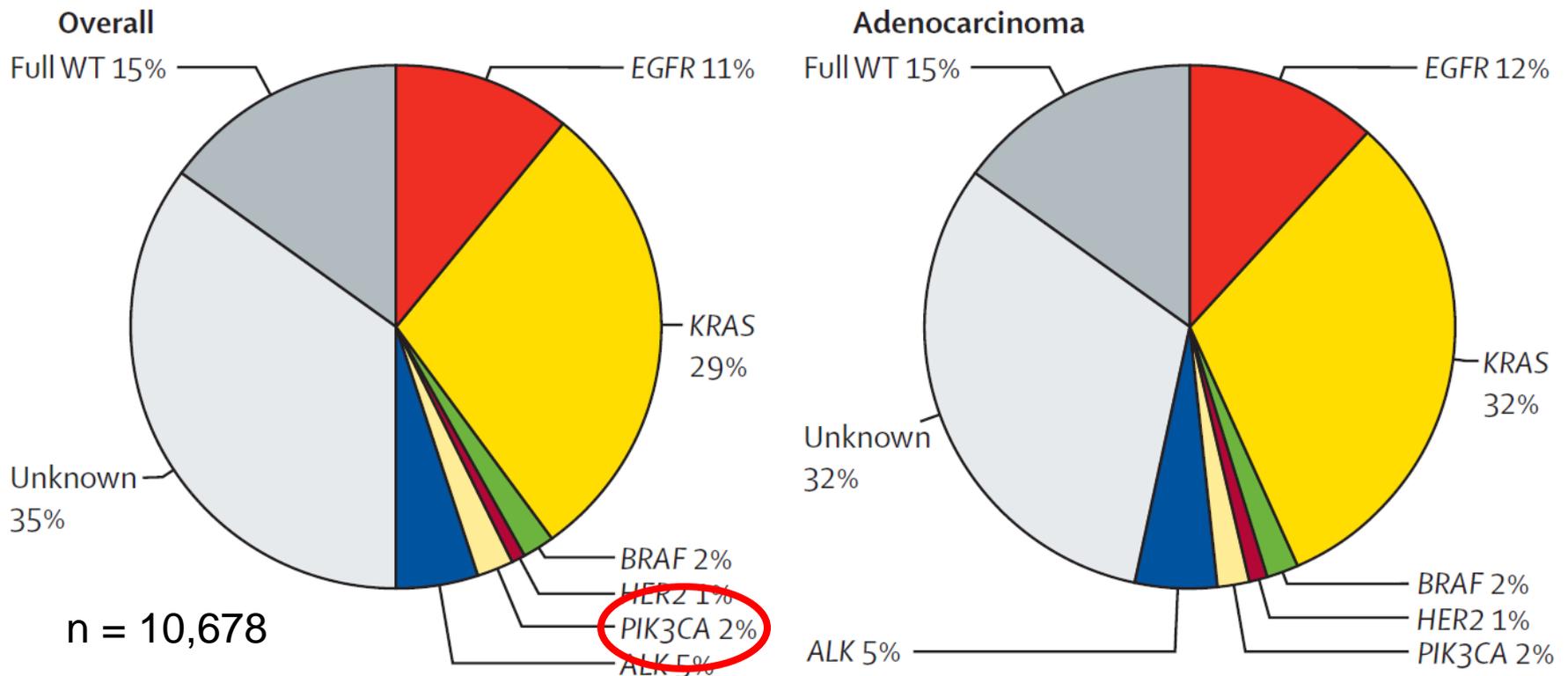


Phosphoinositide-3-kinase signaling pathway

Recurring genomic aberrations in lung cancer



Prospective screening for PI3K pathway aberrations French Cooperative Thoracic Intergroup (IFCT)



Class I PI3K inhibition in PI3K-activated lung cancer

Two-stage phase II study of buparlisib (BASALT-1)

Open-label, Phase II study in patients with previously treated metastatic NSCLC with PI3K pathway activation (Stage 1)*

Pre-screening

- All patients pre-screened for PI3K pathway activation prior to enrollment
- PI3K pathway activation defined as: *PIK3CA* mutation, *PTEN* mutation, and/or PTEN negative (<10% protein expression by IHC)

Study entry if patient has known PI3K pathway-activated tumor

Squamous NSCLC

Previously treated with one prior platinum-based chemotherapy line

Buparlisib
100 mg/day[†]

Non-squamous NSCLC

Previously treated with one or two prior systemic antineoplastic therapy lines

Buparlisib
100 mg/day[†]

Primary endpoint: PFS rate at 12 weeks
Secondary endpoints: PFS, ORR, DCR, safety



Class I PI3K inhibition in PI3K-activated lung cancer

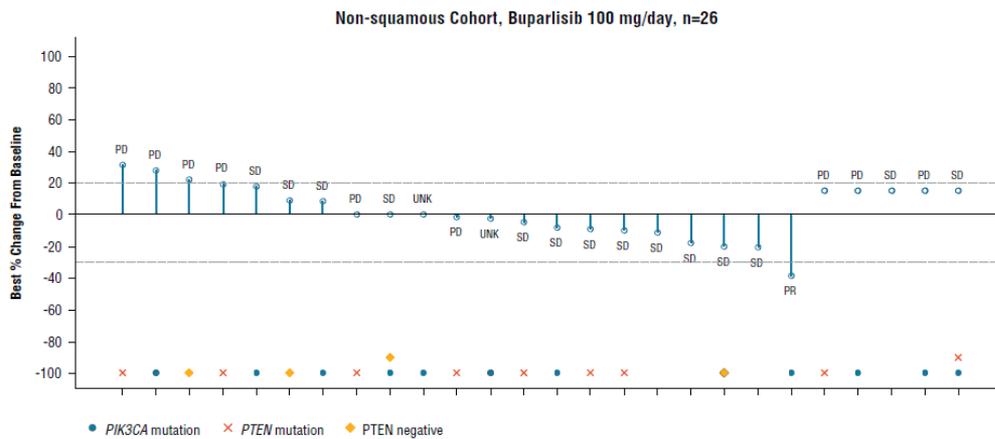
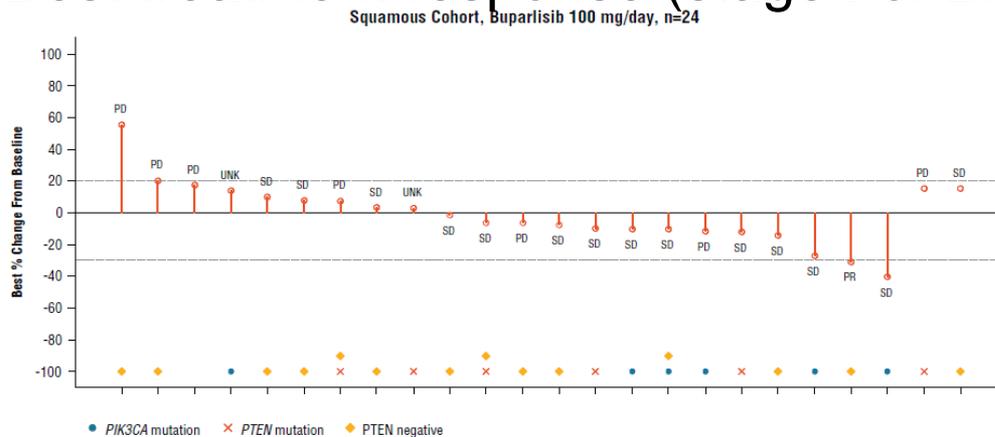
Biomarker prescreening results from BASALT-1

PI3K pathway status	Squamous (N=668) n (%)	Non-squamous (N=574) n (%)	Total (N=1242)* n (%)
PI3K pathway activation†	103 (15.4)	65 (11.3)	168 (13.5)
<i>PIK3CA</i> mutation	n‡=388 (58.1)	n‡=349 (60.8)	n‡=737 (59.3)
Yes	26 (6.7)	22 (6.3)	48 (6.5)
No	362 (93.3)	327 (93.7)	689 (93.5)
Unknown	69	95	164
Missing	211	130	341
<i>PIK3CA</i> mutation only	21 (5.4)	15 (4.3)	36 (4.9)
<i>PTEN</i> Mutation	n‡=394	n‡=342	n‡=736
Yes	25 (6.3)	36 (10.5)	61 (8.3)
No	369 (93.6)	306 (89.5)	675 (91.7)
Unknown	61	90	151
Missing	213	142	355
<i>PTEN</i> mutation only	18 (4.6)	29 (8.5)	47 (6.4)
<i>PTEN</i> Negative	n‡=444	n‡=358	n‡=802
Yes	64 (14.4)	16 (4.5)	80 (10)
No	380 (85.6)	342 (95.5)	722 (90.0)
Unknown	17	32	49
Missing	207	184	395
<i>PTEN</i> negative only	53 (11.9)	12 (3.4)	65 (8.1)



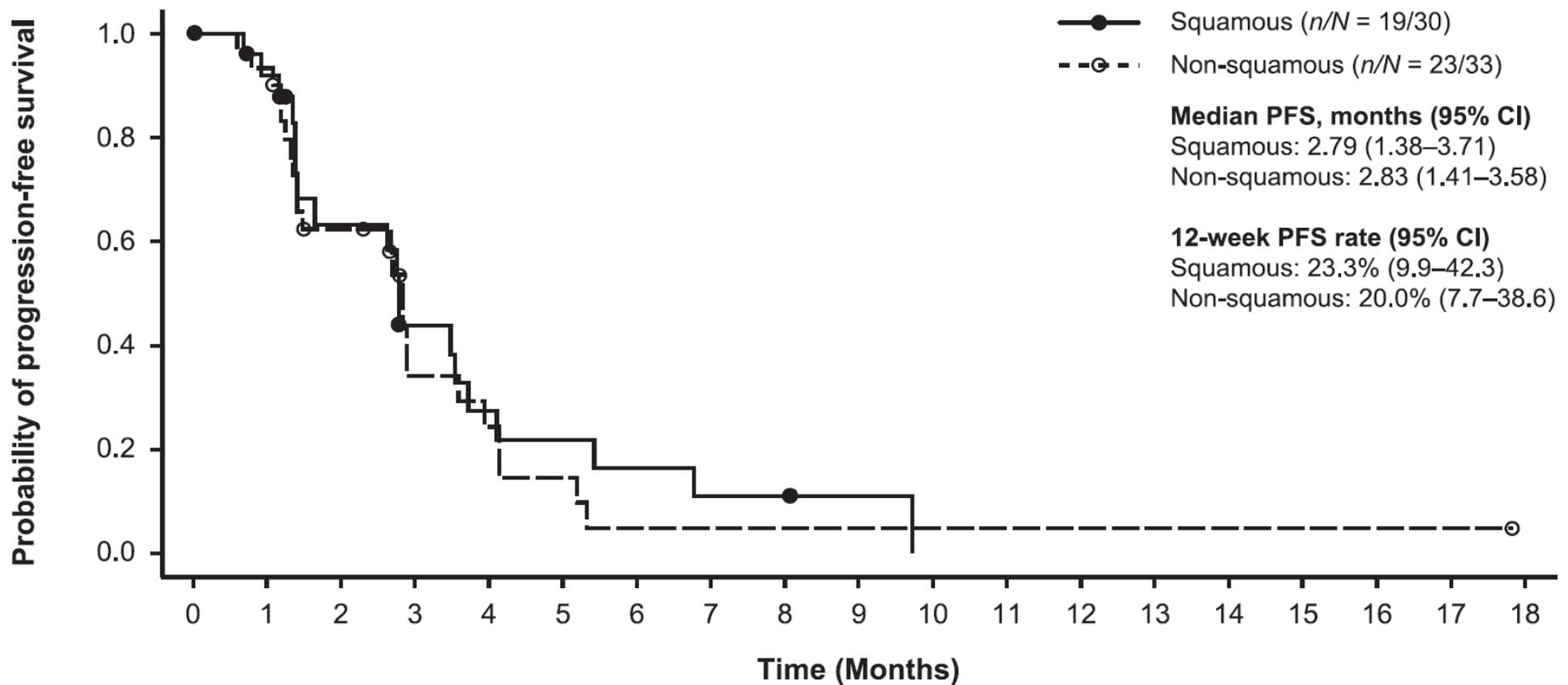
Class I PI3K inhibition in PI3K-activated lung cancer

Best treatment response (stage I of BASALT-1)



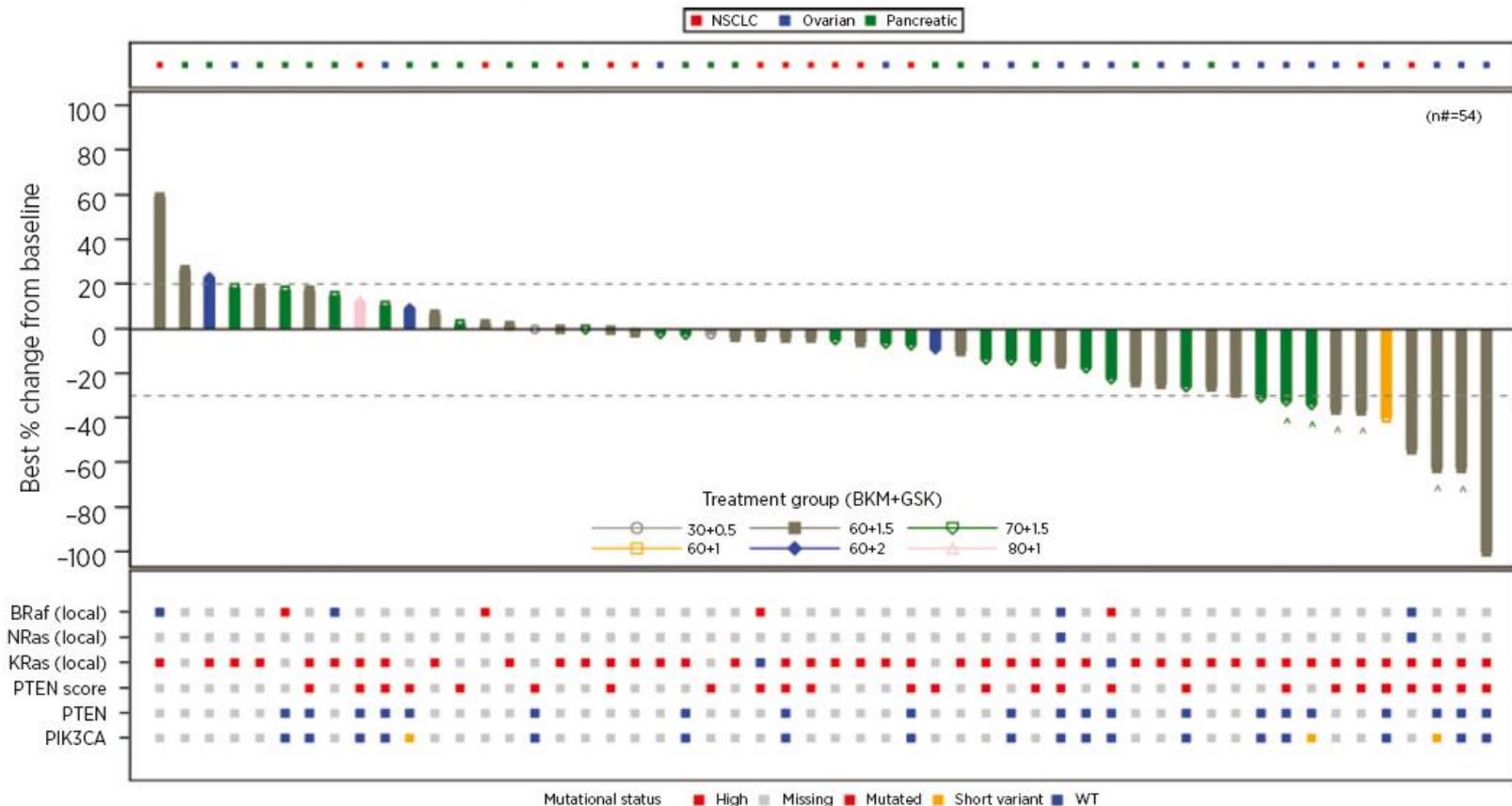
Class I PI3K inhibition in PI3K-activated lung cancer

Progression-free survival (stage I of BASALT-1)



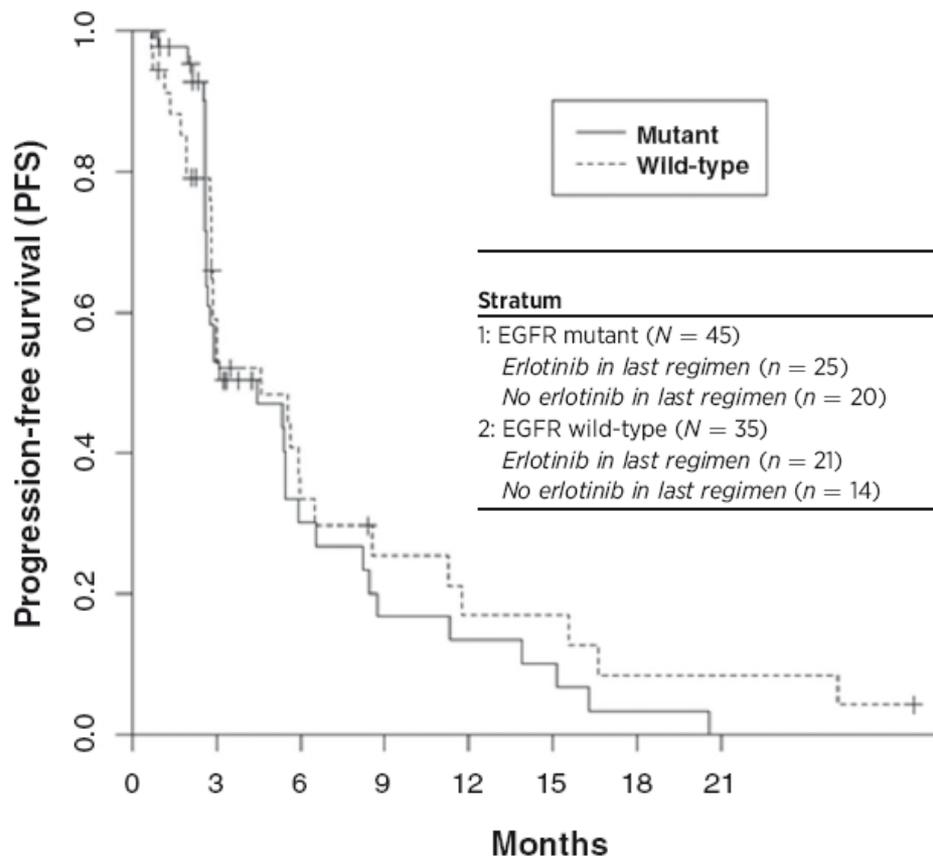
Combined PI3K and MEK inhibition in lung cancer

Phase Ib study of buparlisib plus trametinib



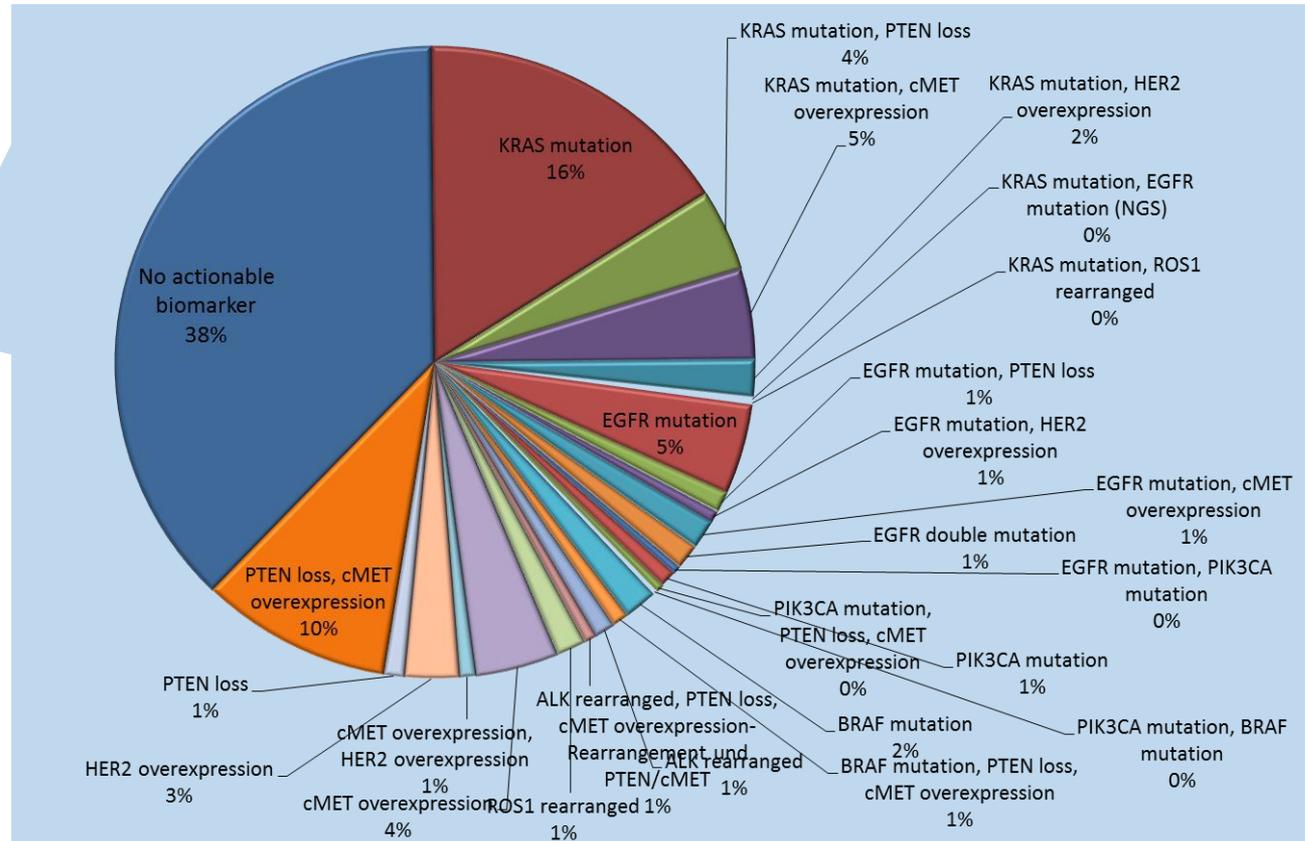
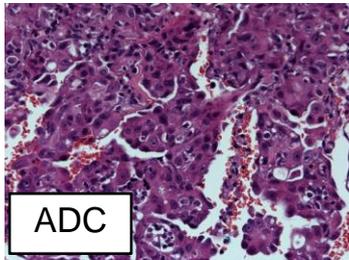
Combined PI3K and AKT inhibition in lung cancer

Phase II study of erlotinib plus MK-2206



Prospective screening for PI3K pathway aberrations

Advanced or metastatic pulmonary adenocarcinomas (WTZ-POP)



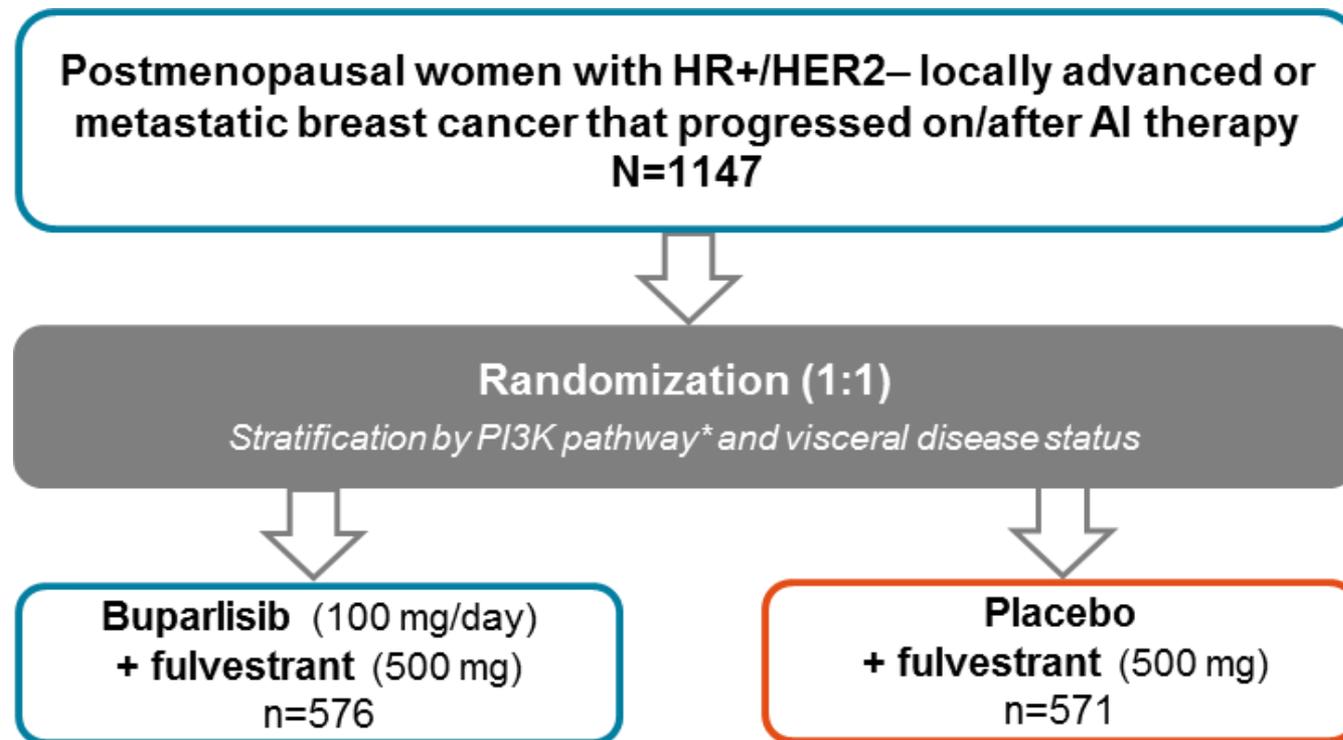
Prospective screening for PI3K pathway aberrations „Oncogenic driver“ or disease modifier in lung cancer?

PIK3CA mutation	Additional oncogenic „driver“
61%	none
23%	<i>EGFR</i> mutation
8%	<i>BRAF</i> mutation
8%	<i>CMET</i> amplification



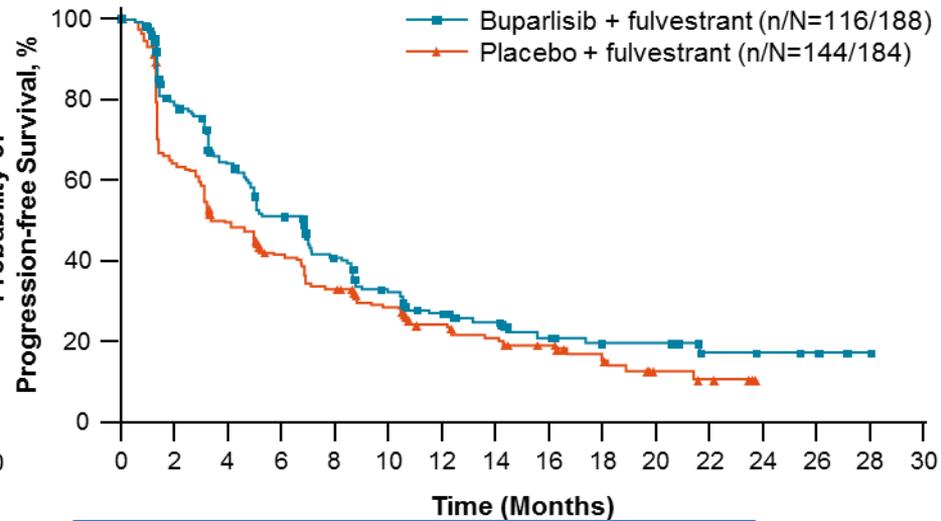
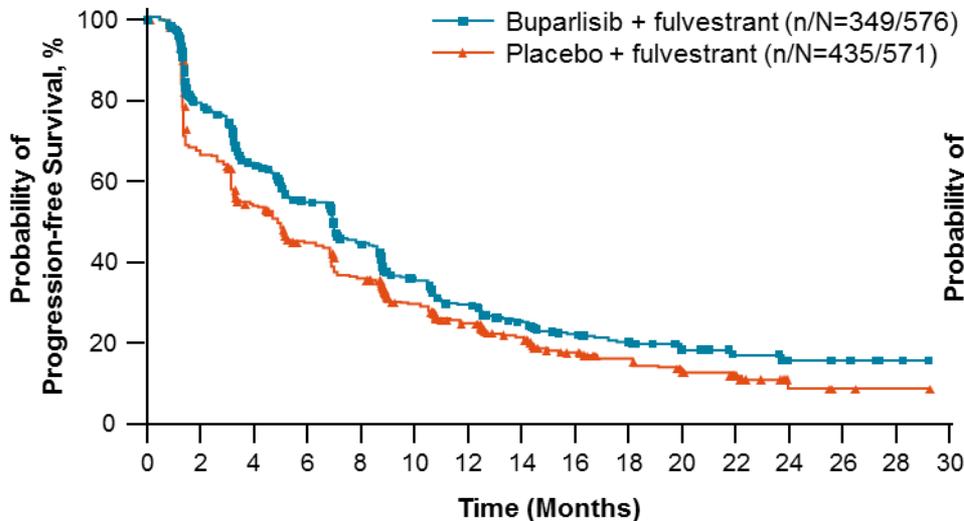
Class I PI3K inhibition in luminal breast cancer

Phase III study of fulvestrant ± buparlisib (BELLE-2)



Class I PI3K inhibition in luminal breast cancer

PFS in full and in PI3K-activated populations (archival biopsy)



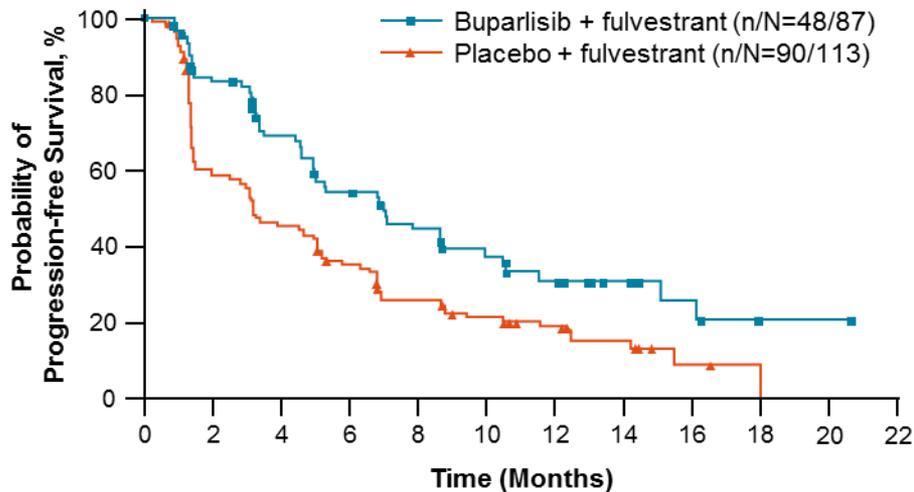
Full Population (N=1047)	Buparlisib + Fulvestrant n=576	Placebo + Fulvestrant n=571
Median PFS, months (95% CI)	6.9 (6.8–7.8)	5.0 (4.0–5.2)
HR (95% CI)	0.78 (0.67–0.89)	
One-sided P value	<0.001	

PI3K Activated Group (N=372)	Buparlisib + Fulvestrant n=188	Placebo + Fulvestrant n=184
Median PFS, months (95% CI)	6.8 (4.9–7.1)	4.0 (3.1–5.2)
HR (95% CI)	0.76 (0.60–0.97)	
One-sided P value*	0.014	

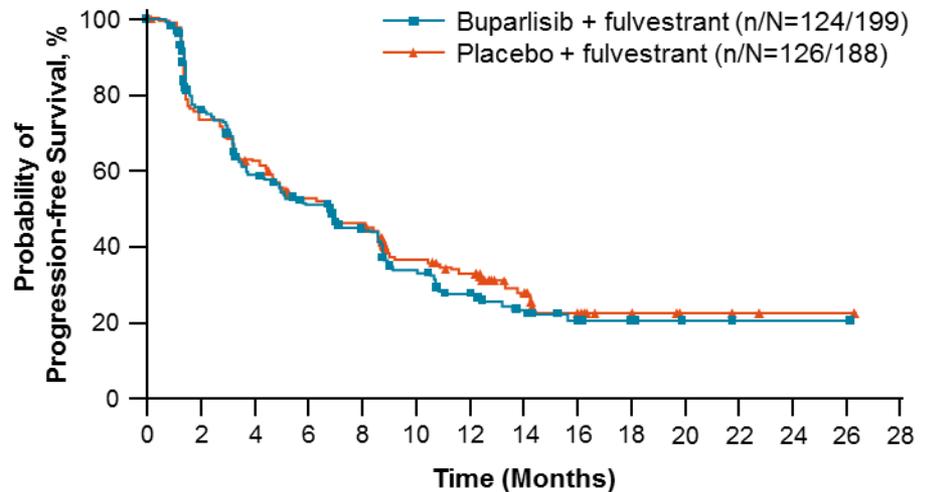


Class I PI3K inhibition in luminal breast cancer

PFS in relation to *PIK3CA* mutation detected in ctDNA



ctDNA <i>PIK3CA</i> Mutant n=200	Buparlisib + Fulvestrant n=87	Placebo + Fulvestrant n=113
Median PFS, months (95% CI)	7.0 (5.0–10.0)	3.2 (2.0–5.1)
HR (95% CI)	0.56 (0.39–0.80)	
One-sided nominal <i>P</i> value	<0.001	



ctDNA <i>PIK3CA</i> Non-mutant n=387	Buparlisib + Fulvestrant n=199	Placebo + Fulvestrant n=188
Median PFS, months (95% CI)	6.8 (4.7–8.5)	6.8 (4.7–8.6)
HR (95% CI)	1.05 (0.82–1.34)	
One-sided nominal <i>P</i> value	0.642	



Prospective screening for PI3K pathway aberrations

Discordant *PIK3CA* mutation prevalence in breast cancer

	Primary tumor		Metastasis		<i>p</i> (χ -square)
	<i>N</i>	% positive	<i>N</i>	% positive	
All	52		70		
PR	30	57.7	35	50.0	0.320
ER	34	65.4	49	70.0	0.509
HR	37	71.2	52	74.3	0.700
Her2/neu+	13	25.0	15	21.4	0.629
Her2/neu−/HR+	28	53.8	45	64.3	0.308
Her2/neu+/HR+	9	17.3	6	8.6	0.249
Her2/neu+/HR−	4	7.7	9	12.9	0.441
Triple negative	11	21.2	9	12.9	0.221
<i>FGFR1</i> amplified	6	11.5	3	4.3	0.246
<i>PI3KCA</i> mutant	7	13.5	21	30	0.034*
<i>PI3KCA</i> amplified	1	1.9	2	2.9	0.471
PTEN loss	7	13.5	9	12.9	0.138

* Denotes statistically significant



Targeting PIK3CA in lung cancer

Summary

- Clinical validation of *PIK3CA* mutations as biomarker for tractable dominant oncogenic dependency in stage IV lung cancer has failed
- Concomitant mutations of additional „oncogenic drivers“ frequently observed
- Questionable predictive value of *PIK3CA* mutation detection in archival tumor biopsy (breast cancer)
- Combination therapies involving PI3K pathway-targeting agents limited by clinical toxicities



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