



CARCINOMA POLMONARE: QUALI NOVITÀ NEL 2022?

**Il trattamento della malattia con alterazioni di
BRAF, *ROS1* o *RET* e altri target emergenti**

Sara Pilotto

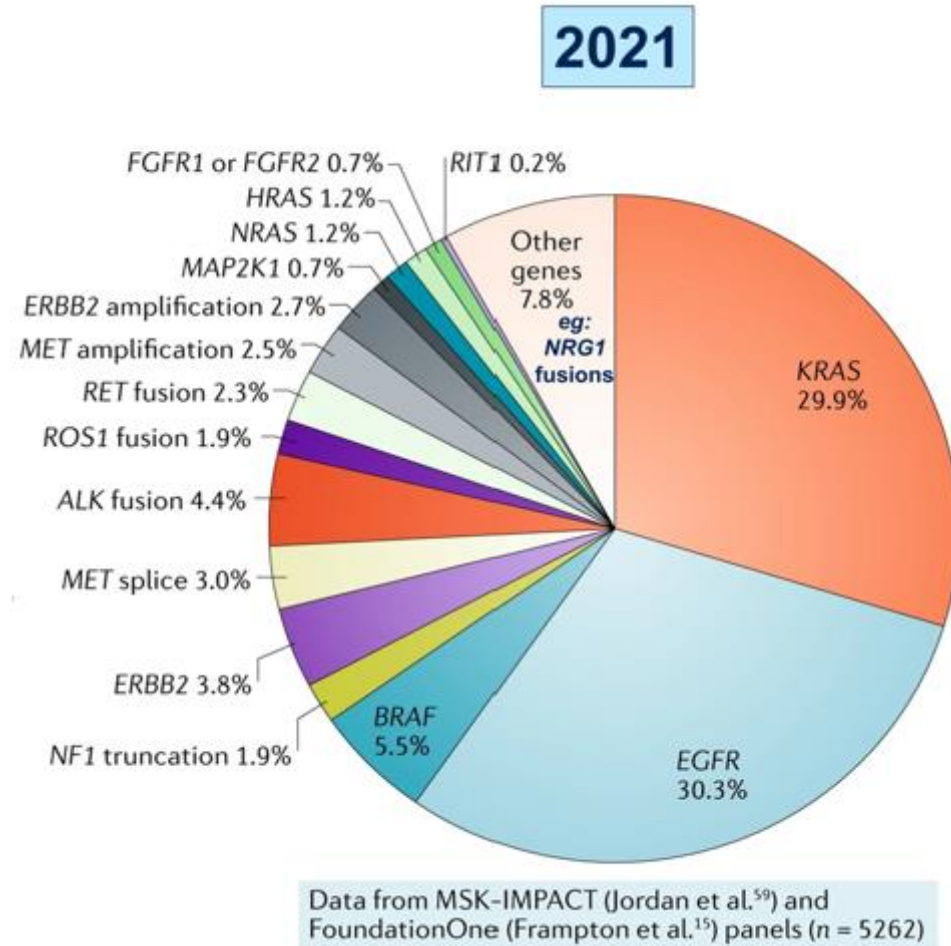
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Disclosures

Commercial Interest	Relationship(s)
AstraZeneca	Speaker's Bureau, Consultant, Research Funding
Merck&Co	Advisory Board, Speaker's Bureau, Consultant
Boehringer Ingelheim	Speaker's Bureau, Consultant
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Roche	Advisory Board, Speaker's Bureau, Consultant
AMGEN	Advisory Board, Consultant
Novartis	Advisory Board, Consultant

Precision Therapy for non-squamous (*for now*) NSCLC in 2021/2022



What we absolutely need TODAY*

- 15% EGFR mutations
- 4% ALK rearrangements
- 1% ROS1 rearrangements
- 4% BRAF p.V600E mutations
- <1% NTRK 1,2,3 rearrangements

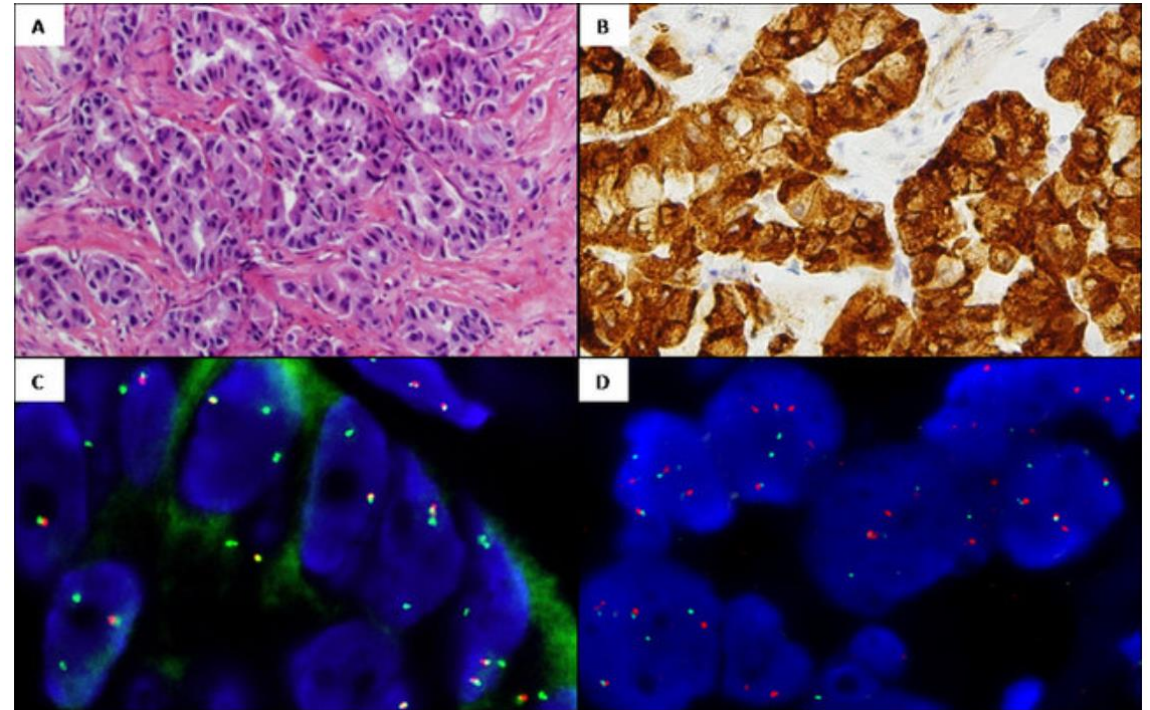
* In metastatic non-squamous NSCLC (before 1° line initiation) - in Italy



proposito, in tutti i pazienti con NSCLC in stadio IIIB-IIIC (non candidati a trattamenti loco-regionali), e IV, risulta raccomandato completare la diagnosi morfologica (di cui sopra) con la caratterizzazione delle mutazioni in *EGFR* (*Epidermal Growth Factor Receptor*) e *BRAF* (*B-Raf proto-oncogene*), la definizione delle traslocazioni a carico di *ALK* (*Anaplastic Lymphoma Kinase*), *ROS-1* (*Proto-oncogene tyrosine-protein kinase ROS*) e *NTRK 1,2 e 3* (*Neurotrophic Tyrosine Receptor Kinase*) e la valutazione dei livelli di espressione del PD-L1 (*Programmed-death ligand 1*) (secondo i cut – off validati dagli studi

***ROS1* rearrangements: do not forget this rare target!**

- Seen in 0.9% - 2% of lung adenocarcinoma
- Typically associated with younger age (average 50y), never/light smoking history, adenocarcinoma subtype



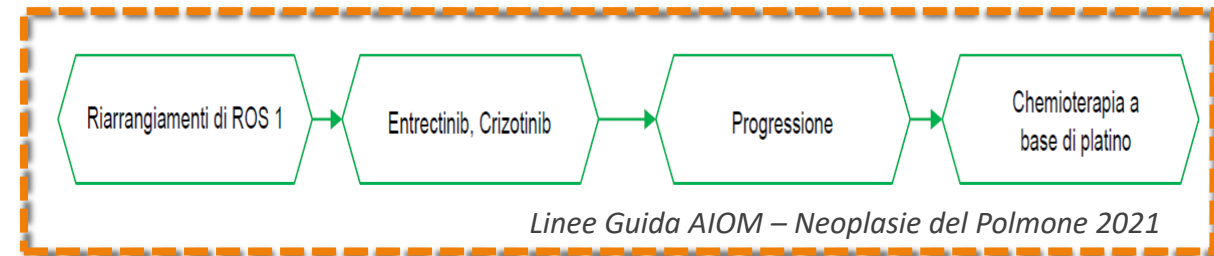
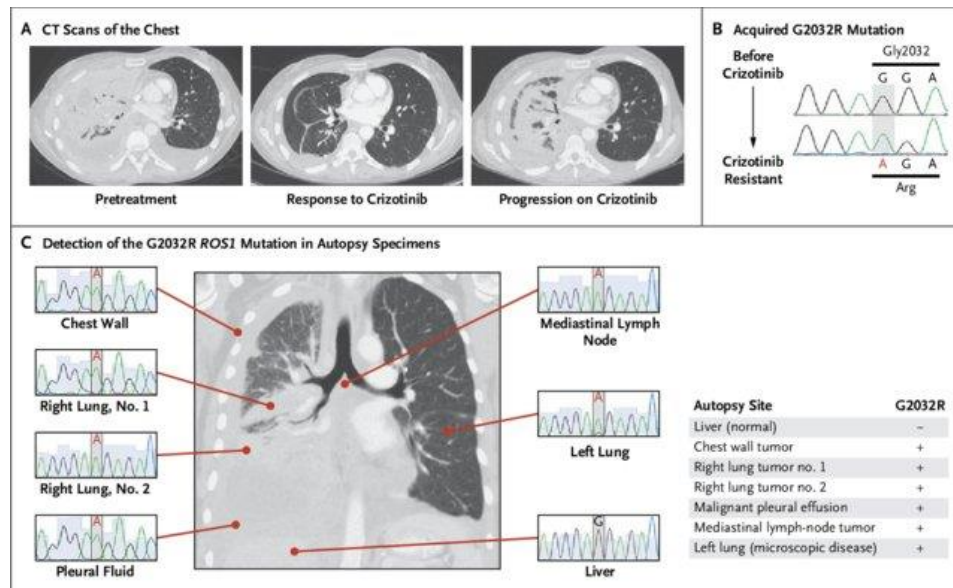
Clave S et al, Oncotarget 2016

ROS1 rearrangements: crizotinib, a (EX?) standard of care

Study	Phase	Patients, No.	ORR, %	DCR, %	mPFS, Months	mOS, Months
PROFILE 1001 ^{12,30}	I/II	53	72	90	19.3	51.4
Phase II East Asian Study ³¹	II	127	71.7	88.2	15.9	12-month rate, 83.1%
EUCROSS ³²	II	30	70	90	20	NR (12-month rate, 83%)
AcSé ³³	II	37	69.4	69.4	5.5	17.2
METROS ³⁴	II	26	65	85	22.8	NR
EUROS1 ¹¹	Retrospective	30	80	86.6	9.1	NR
ROS1 Fusion Partners ³⁵	Retrospective	49	83.3	97.2	12.6	32.7
Pemetrexed study ³⁶	Retrospective	15	80	90	9.6	NA

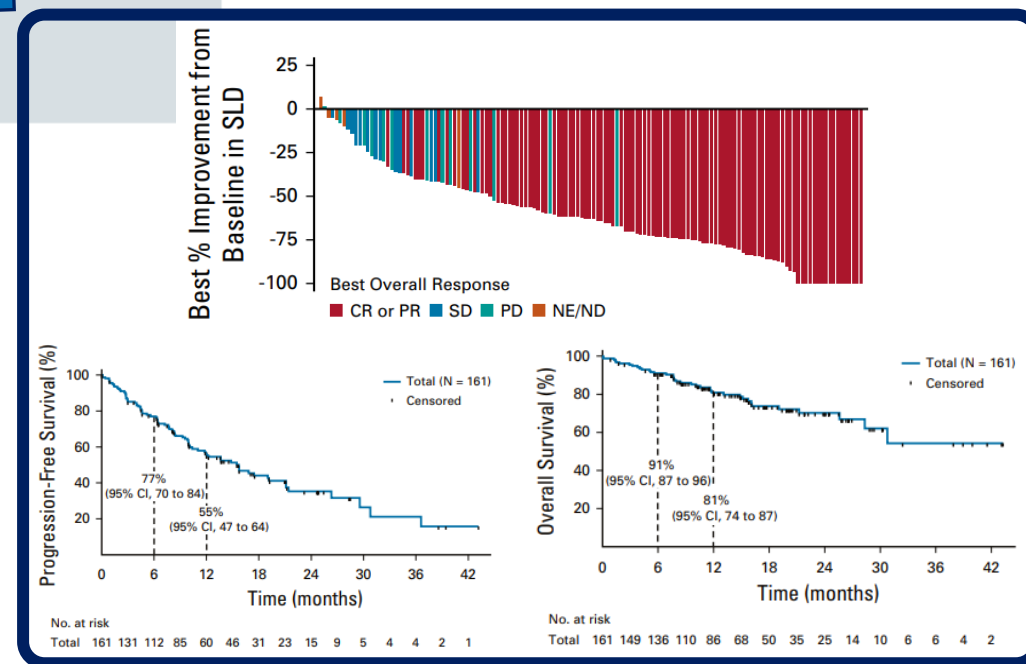
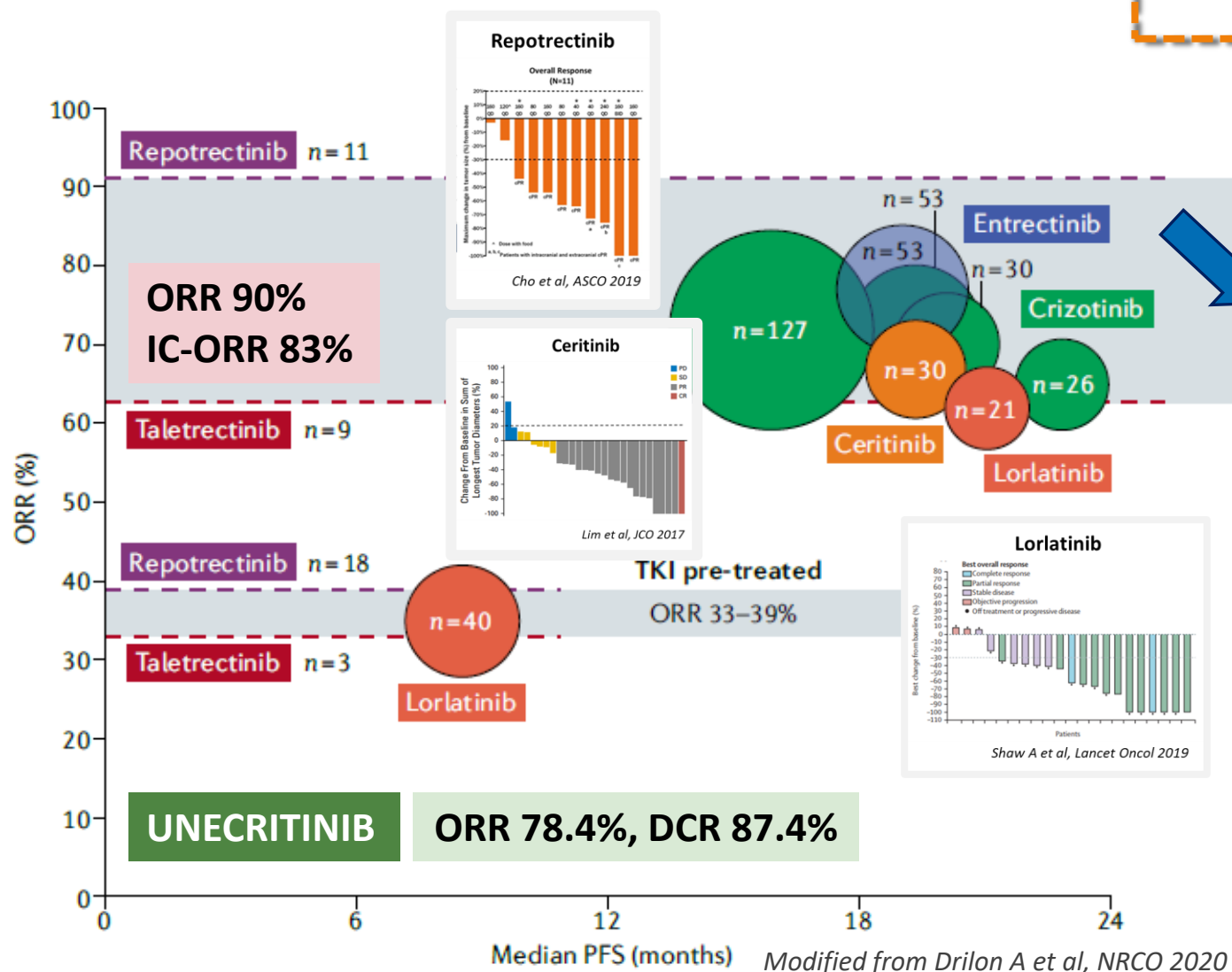
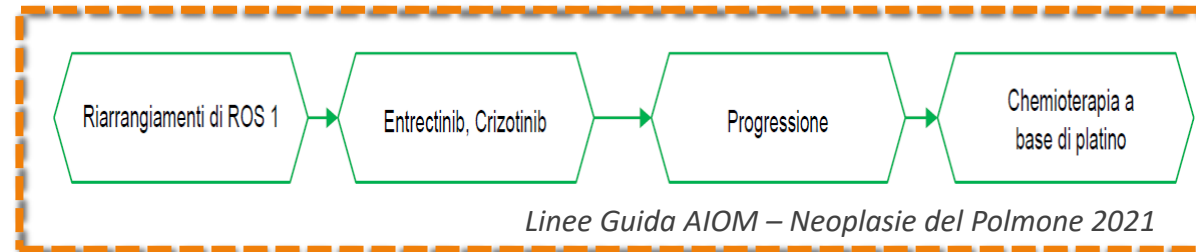
- On-target resistance mutations (*G2032R*)

- CNS progression



Almquist and Ernani, JCO Oncology Practice 2021
 Awad M et al, NEJM 2013

ROS1 rearrangements: a changing (*overcrowded*) landscape

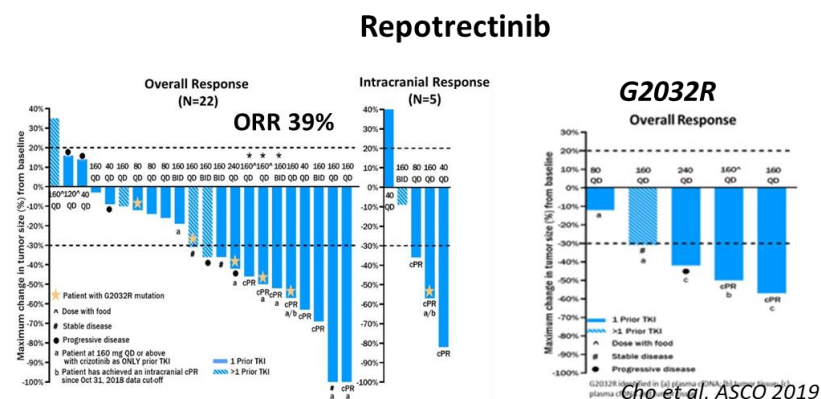


How do we best sequence?? *[yesterday, today and tomorrow]*

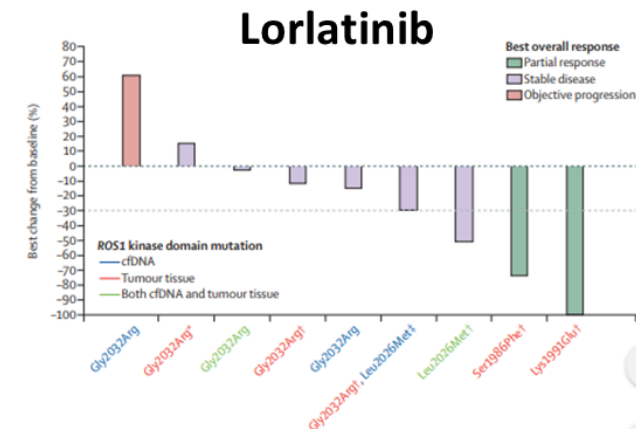
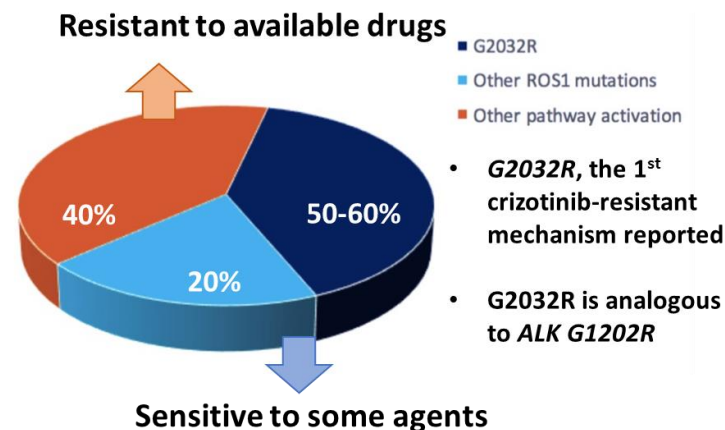


Modified from Remon J et al, CTR 2021

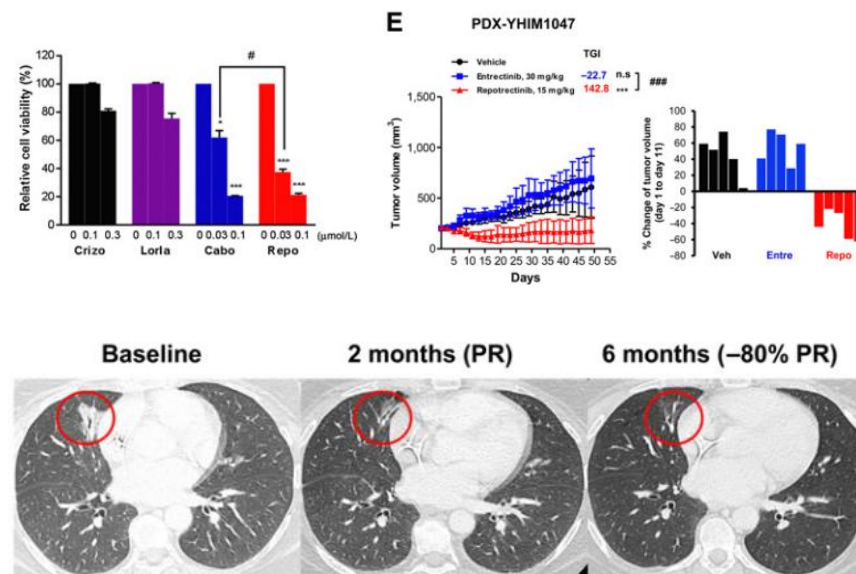
Repotrectinib showed selective and potent in vitro and in vivo activity against ROS1 G2032R



Cho et al. ASCO 2019



Shaw A et al, Lancet Oncol 2019



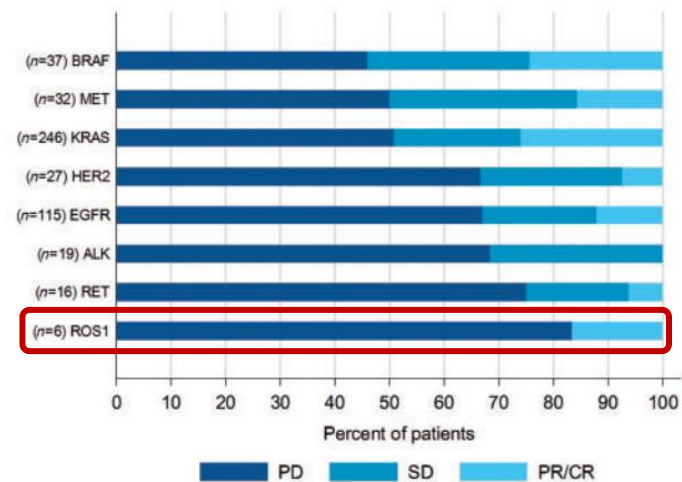
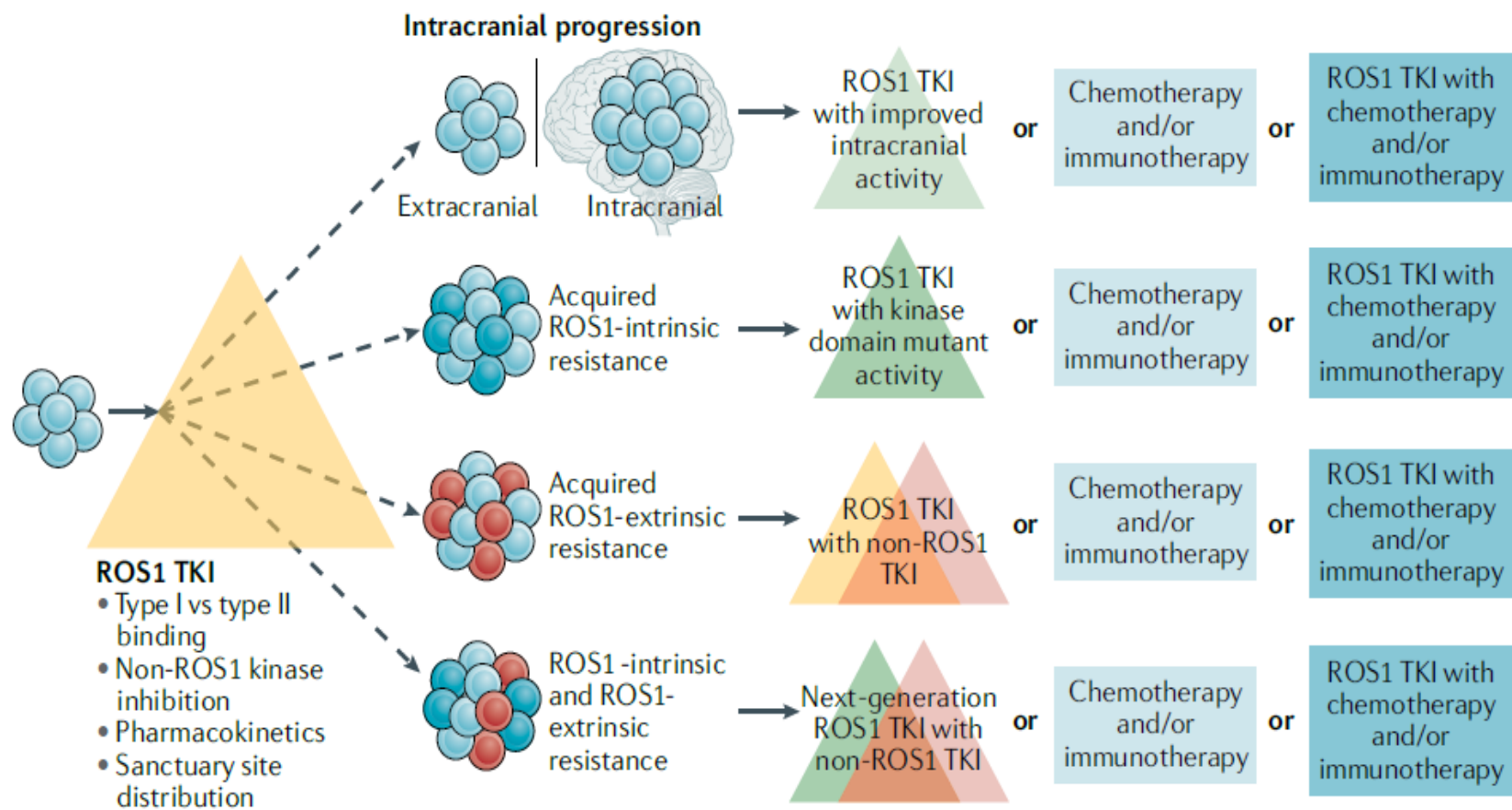
Baseline (blood/ tumor)	best response
PF-001-002 SLC34A2-ROS1, G2032R, TP53	PD
PF-001-003 EZR-ROS1, TP53	PD
PF-001-004 SDC4-ROS1, G2032R, TP53	PD
PF-001-005 SDC4-ROS1, G2032R, TP53	PD
PF-001-006 SDC4-ROS1, TP53	PD
PF-001-007 CD74-ROS1, G2032R	PD
PF-001-013 MTOR	PD
PF-001-016 none	SD
PF-001-022 Nil sample	SD
PF-001-017 TP53	SD
PF-001-021 EGFR	SD
PF-001-019 None	SD
PF-001-008 EZR-ROS1, S1861I, TP53	SD
PF-001-001 EGFR	PR
PF-001-011 SLC34A2-ROS1	PR
PF-001-018 LRI3G-ROS1, V2054A, TP53	PR
PF-001-009 intergenic-ROS1, TP53	PR
PF-001-020 TSC	PR
PF-001-010 none	PR
PF-001-012 nil sample	PR
PF-001-015 none	PR
PF-001-014 CD74-ROS1, TP53	PR

Yun MR et al, Clin Cancer Res 2020



Landi L, WCLC 2019, ESMO 2019

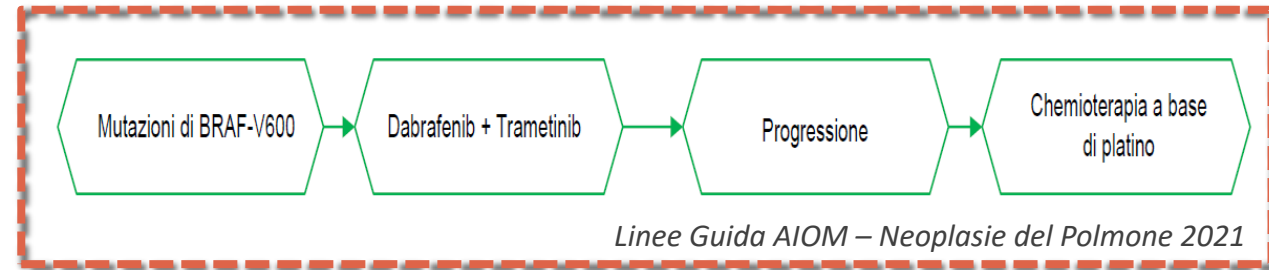
How do we best sequence?? [*yesterday, today and tomorrow*]



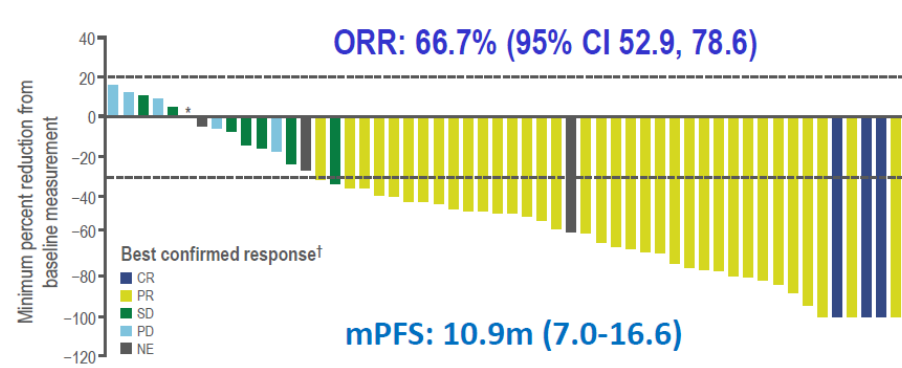
Room for IO? Not alone!
In combos? Mah!

BRAF p.600E mutation: an established SoC

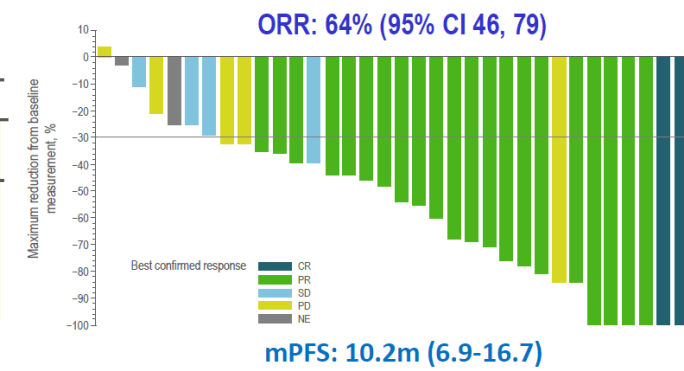
- BRAF mutations present in **2% - 4%** of NSCLC (½ BRAF V600E or class 1 BRAF mutations)
- V600E: mainly female (~ 60%), **current/former smokers** (never smokers ~ 30%)
- Non-V600E: mainly male and almost exclusively current/former smokers
- Potential **mechanism of resistance in EGFR-mutated NSCLC** (osi-treated 3-10%)



	Previously Treated				Treatment Naive
	VE-Basket trial vemurafenib (n=20)	AcSé trial vemurafenib (n=100)	BRF113928 dabrafenib (n = 78)	BRF113928 Dabrafenib Plus Trametinib (n = 57)	BRF113928 Dabrafenib Plus Trametinib (n = 36)
Male	14 (70%)	-	39 (50%)	29 (51%)	14 (39%)
Never smoker	7 (35%)	-	29 (37%)	16 (28%)	10 (28%)
ORR % (95% CI)	42 (20-67)	44.9	33 (23-45)	67 (53-79)	64 (46-79)
PFS, median (95% CI)	7.3 (3.5-10.8)	5.2	5.5 (3.4-7.3)	10.2 (6.9-16.7)	10.9 (7.0-16.6)
OS, median (95% CI)	NA	9.3	12.7 (7.3-16.3)	18.2 (14.3-NE)	24.6 (12.3-NE)



Planchard D, Lancet Oncol 2016

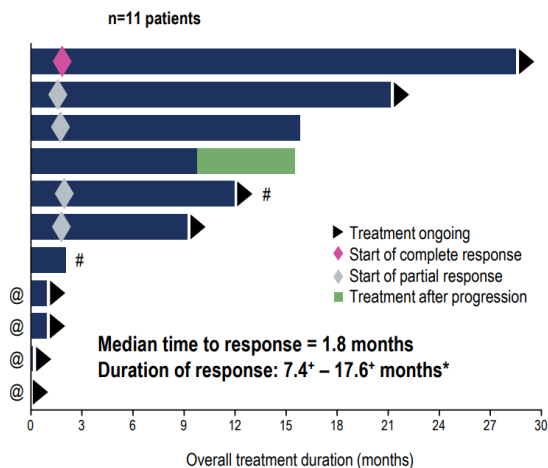
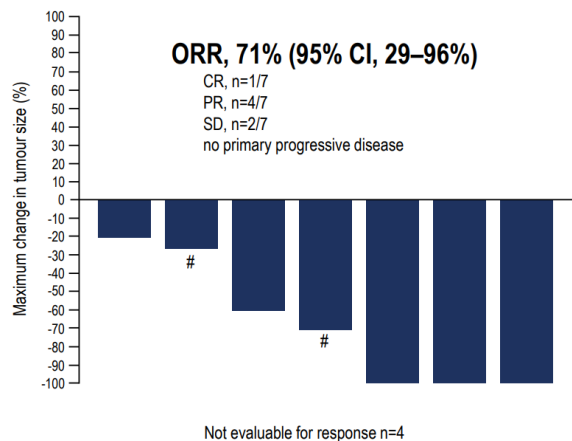
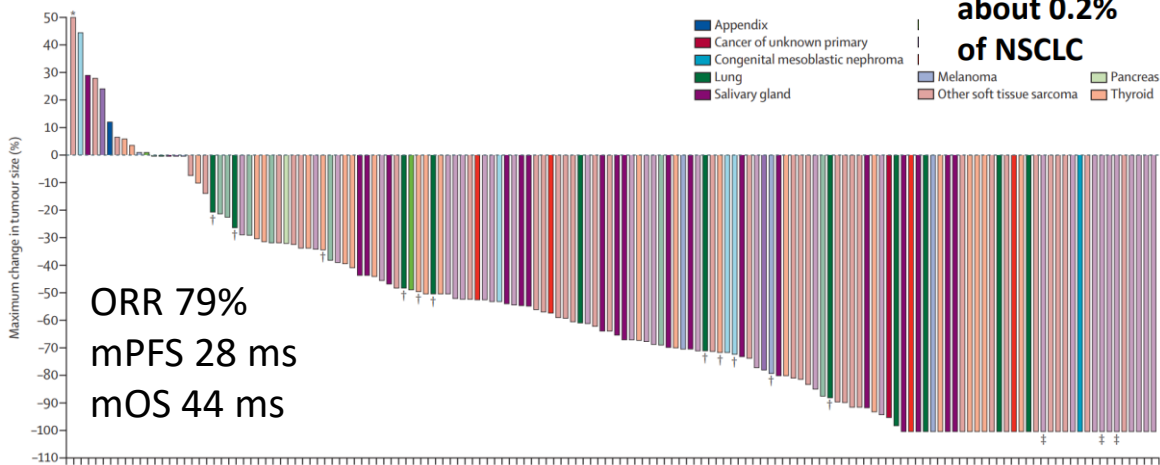


Planchard D, Lancet Oncol 2017

NTRK 1,2,3 rearrangements

NTRK fusions are rare events:
0.21% across 11,116 patients with tumors of all types

LAROTRECTINIB



Hong D et al, Lancet Oncol 2020

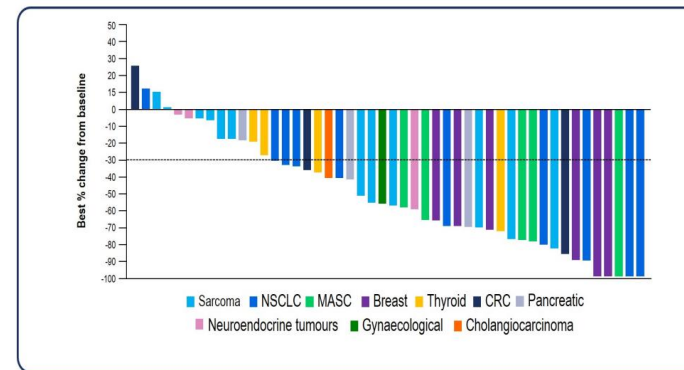
Riarrangiamenti di NTRK

Entrectinib, Larotrectinib

Progressione

Chemioterapia a base di platino

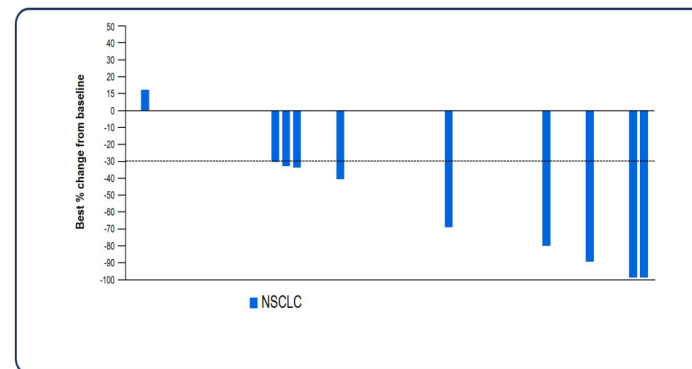
Linee Guida AIOM – Neoplasie del Polmone 2021



Data cut-off date: 31 May 2018. Note: Patients (n=6) without matched pre/post therapy scans were excluded from the plot

Efficacy outcomes	NTRK+ patients (n=54)
ORR,* %	57.4 (95% CI: 43.2-70.8)
CR,* n (%)	4 (7.4)
Median DOR,* months	10.4 (95% CI 7.1-NR)
Median PFS,* months	11.2 (95% CI 8.0-14.9)
Median OS, months	20.9 (95% CI 14.9-NR)

NTRK+ patients ORR* 57.4%



Data cut-off date: 31 May 2018. Note: Patients (n=6) without matched pre/post therapy scans were excluded from the plot. * By blinded independent central review

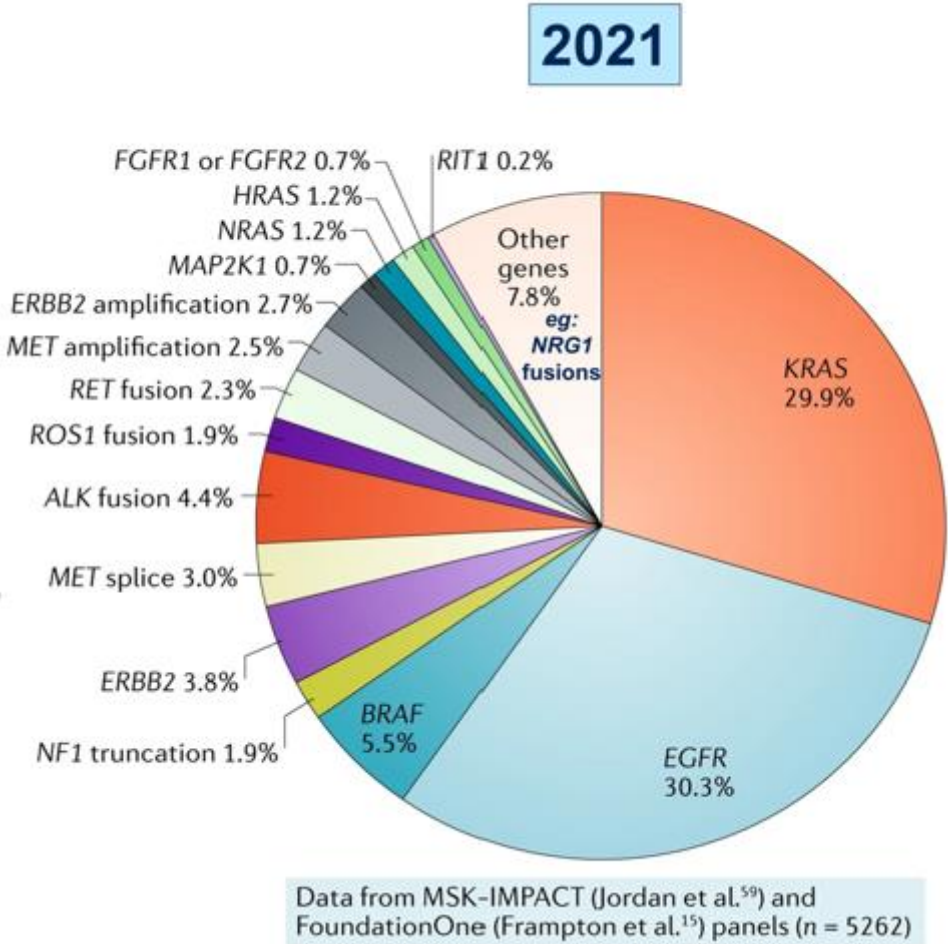
Efficacy outcomes	NTRK+ NSCLC patients (n=10)
ORR,* %	70.0 (95% CI: 34.75-93.33)
CR,* n (%)	1 (10)
PR,* n (%)	6 (60)
SD,* n (%)	1 (10)
Median DOR,* months	NE (95% CI 10.4-NE)
Median PFS,* months	14.9 (95% CI 4.7-NE)

NTRK+ NSCLC patients with CNS disease at baseline (n=6)	
Intracranial response,* n (%)	4 (66.7)
CR	2 (33.3)
PR	2 (33.3)
SD	1 (16.7)
IE	1 (16.7)

Doebele RC et al, Lancet Oncol 2020

ENTRECTINIB

Precision Therapy for non-squamous (*for now*) NSCLC in 2021/2022

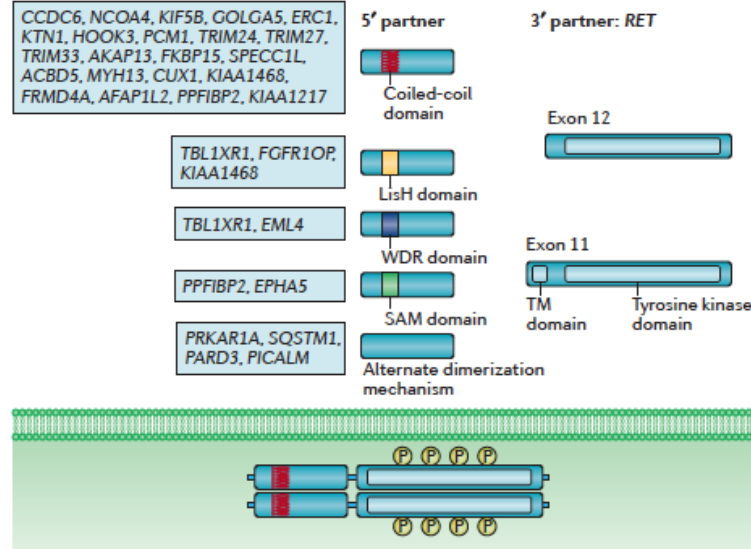


	What we absolutely need TODAY*	What we (also) need TODAY	
15%	EGFR mutations	RET rearrangements	1-2%
4%	ALK rearrangements	MET ex 14 mutations	3-4%
1%	ROS1 rearrangements	HER2 mutations	2-4%
4%	BRAF p.V600E mutations	KRAS p.G12C mutations	13%
<1%	NTRK 1,2,3 rearrangements		

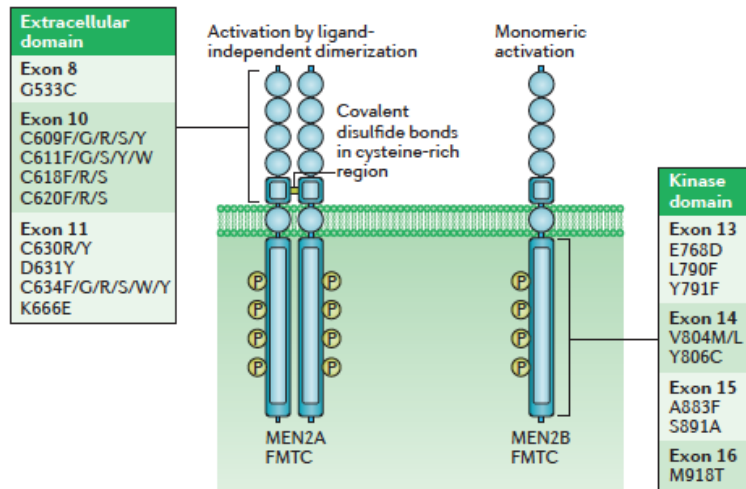
* In metastatic non-squamous NSCLC (before 1° line initiation) - in Italy

RET rearrangements: the next big target..

a RET fusion genes



b RET nonsynonymous point mutations



Younger age (≤ 60 years)¹



Never-smoker status¹



Lower likelihood of harboring other major oncogenic drivers²



Signet ring cells in $\geq 10\%$ of tumor cells¹



Smaller tumors (≤ 3 cm) with early lymph node metastases¹



Poorly differentiated tumors¹

	Alectinib	Apatinib	Cabozantinib	Dovitinib	Lenvatinib	Motesanib	Nintedanib	Ponatinib	Regorafenib	RDX-105	Sitravatinib	Sorafenib	Sunitinib	Vandetanib
RET						*								
RET C634R						*						*	*	*
RET C634W			*			*								*
RET M918T						*						*	*	*
RET V804L						*						*	*	*
RET V804M						*						*	*	*
RET Y791F														
VEGFR1														
VEGFR2									murine					
VEGFR3									murine					
EGFR														
KIT														
FGFR1														
Other targets	ALK		MET	FLT3				ABL1		BRAF**	AXL			

IC₅₀

<5 nM

5–50 nM

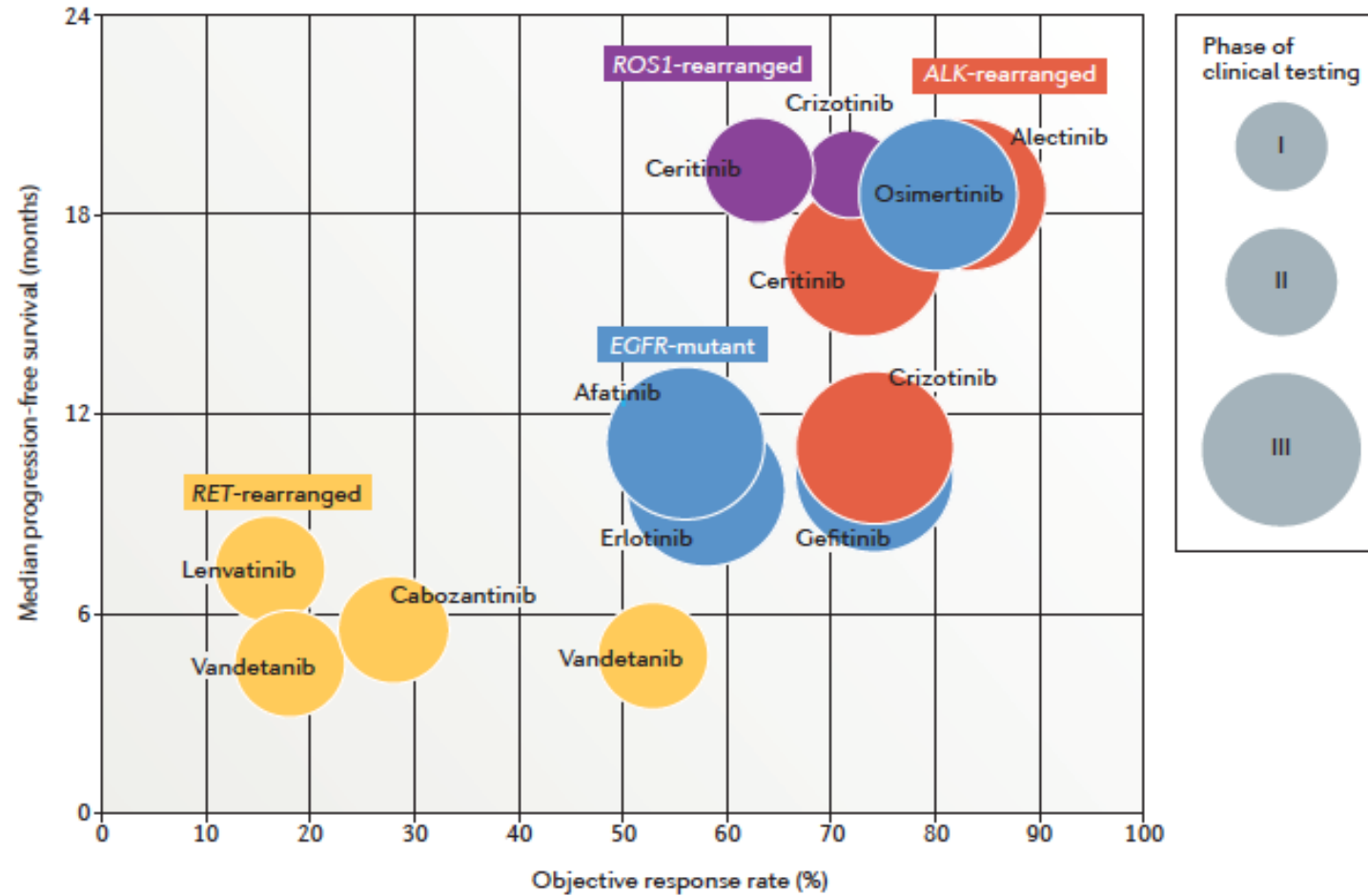
>50–200 nM

>200 nM

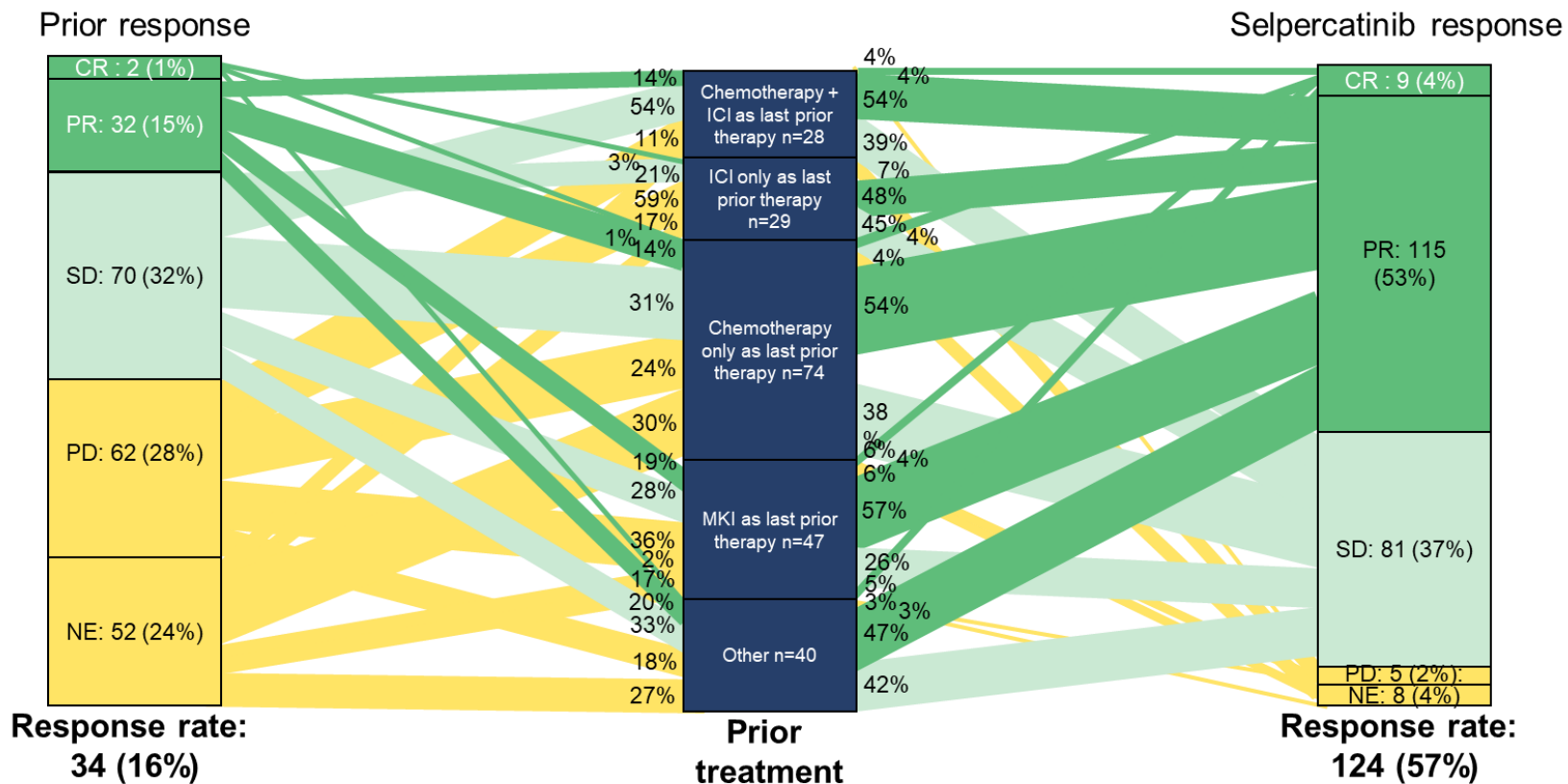
*Cell-based assay
**K_d nmol/l

Wang R, et al. JCO 2012; Drilon A, et al. Nat Rev Clin Oncol 2018

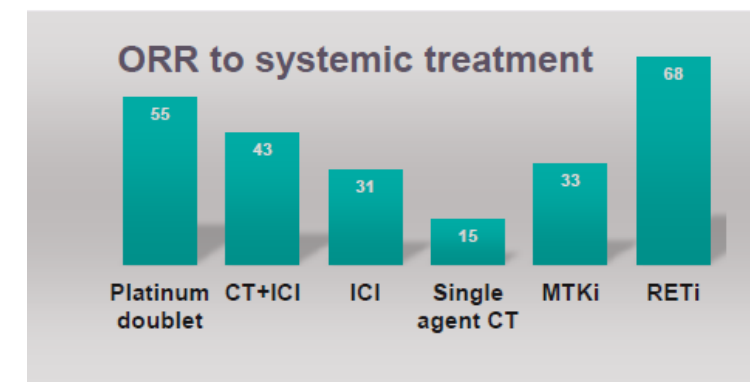
RET rearrangements as the next big target: Are we sure??



Systemic therapy in *RET* rearranged NSCLC

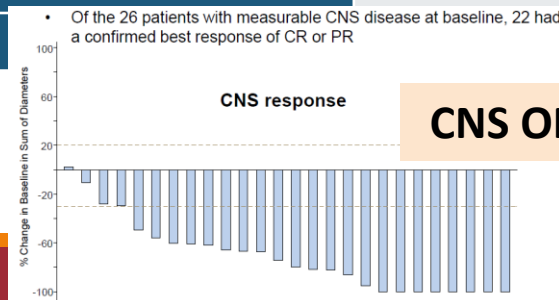


Patients' characteristics	N = 149
Median age (range)	61.9 (53-69.6)
Female sex (%)	87 (58.4)
Smoking history (%)	67 (45)
Adenocarcinoma histology (%)	138 (92.5)
Median number of lines (IQR)	2 (1-3)



RET rearrangements: two guest Stars!!

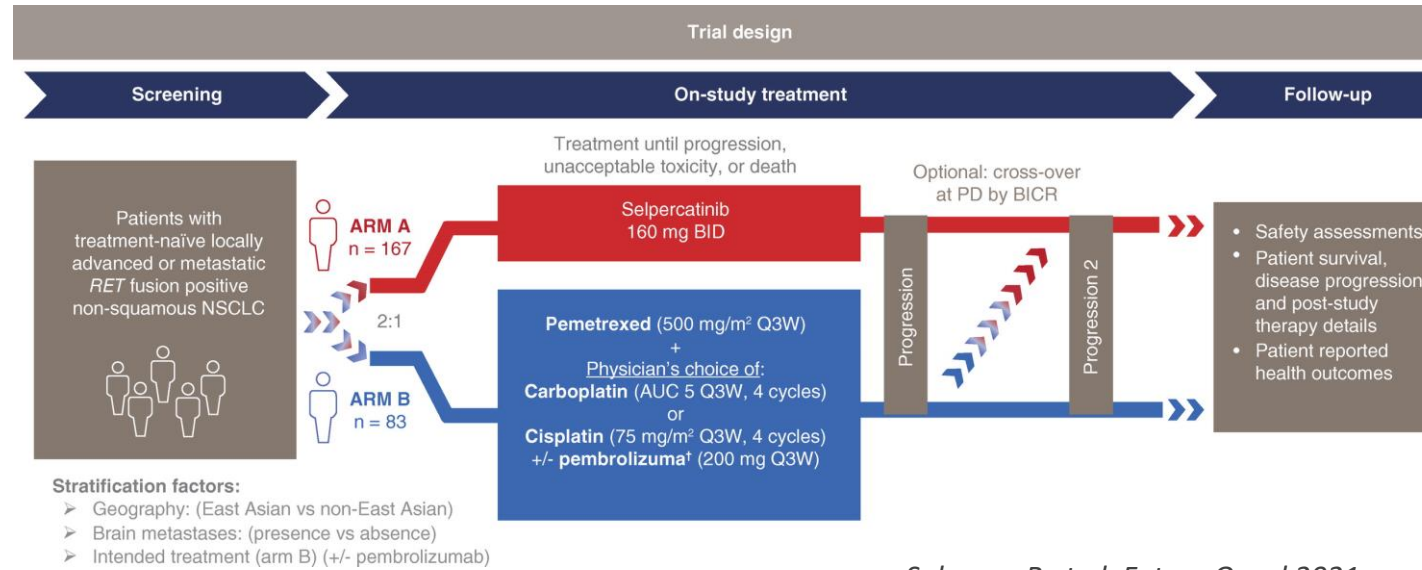
	SELPERCATINIB	PRALSETINIB
	[LIBRETTO-001]	[ARROW]
	Drilon A. et al ELCC 2022	Curigliano G. et al ASCO 2021
ORR (% , N)		
Prior platinum-based chemotherapy	61% (n = 247)	62% (n = 126)
Treatment-naïve	84% (n = 68)	88% (n = 25) ³
DCR (% , N)		
Prior platinum-based chemotherapy	94% (n = 218) ¹	91% (n = 126)
Treatment-naïve	93% (n = 48)	96% (n = 25) ³
mPFS (months, N)		
Prior platinum-based chemotherapy	24.9 (n = 247)	16.5 (n = 136)
Treatment-naïve	22.0 (n = 68)	NR
Grade ≥3 TRAEs occurring in ≥ 10% of patients (% , N)	Hypertension 12.1% (n = 746) ²	Neutropenia 19%, anemia 13%, hypertension 12% (n = 471) ⁴
Discontinuation rate (% , N)	2% (n = 746) ²	6% (n = 471) ⁴



Belluomini L et al, SUBMITTED

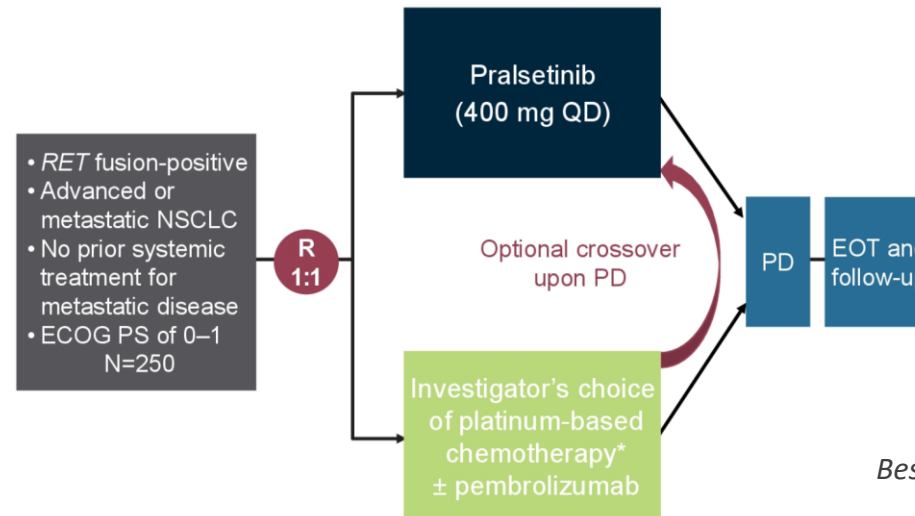
RET rearrangements: phase 3 trials are ongoing in 1st line

LIBRETTO-431



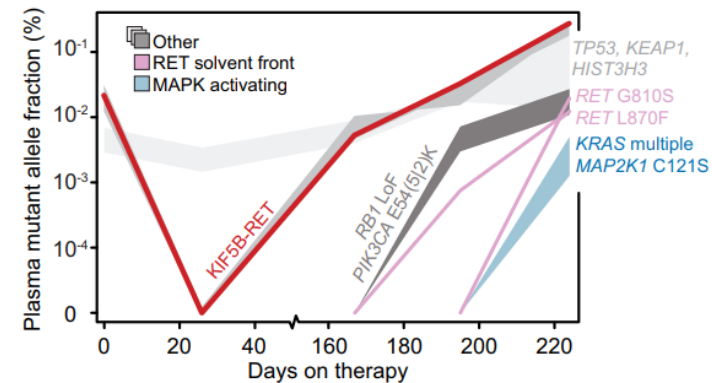
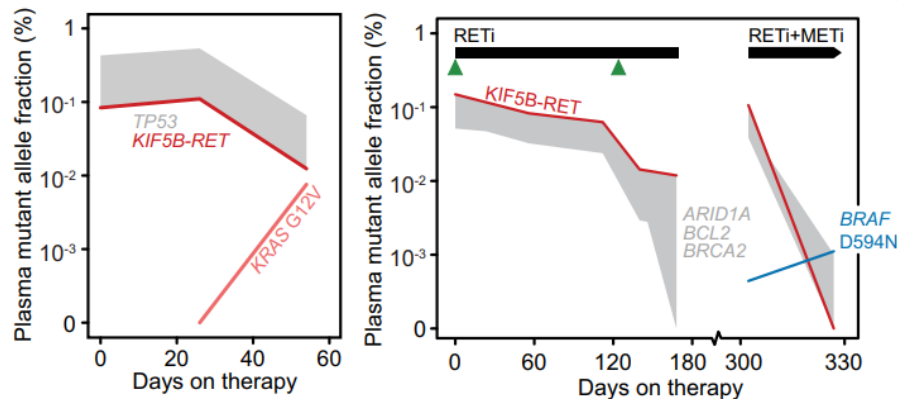
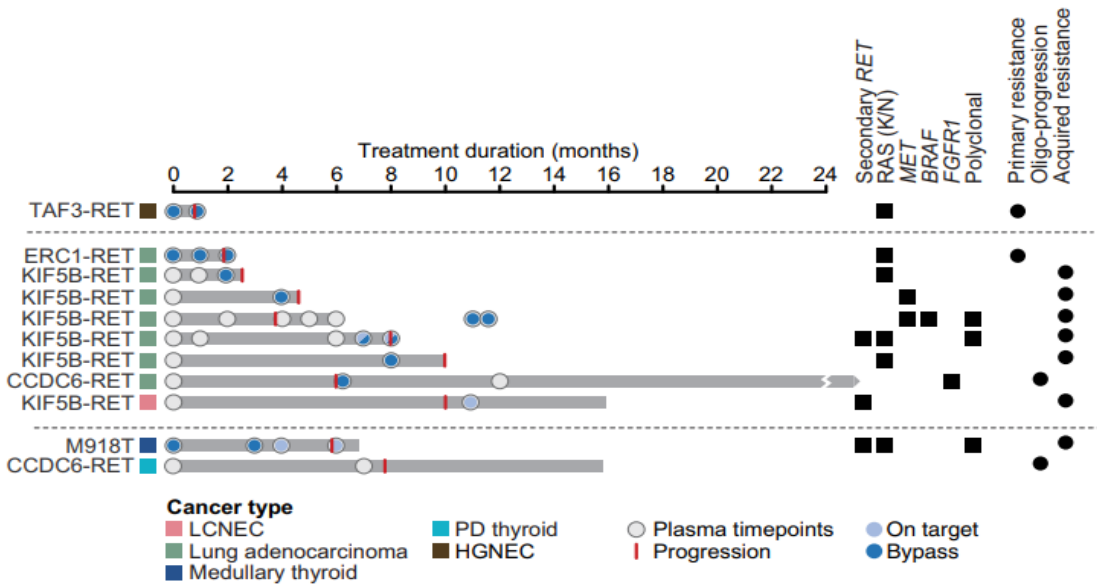
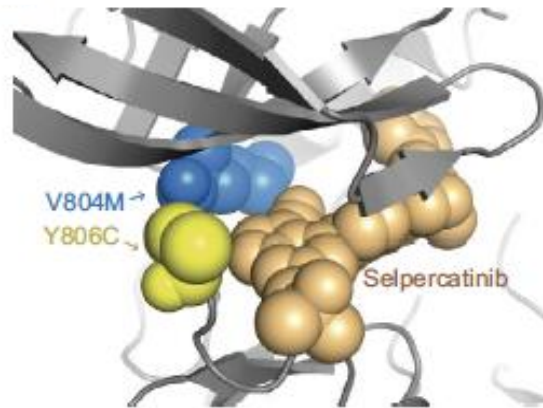
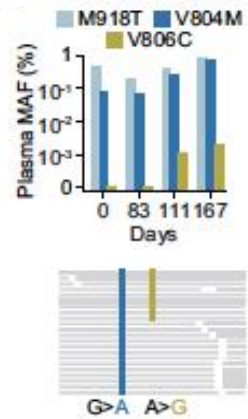
Solomon B et al, Future Oncol 2021

AcceleRET



Besse B et al, ASCO 2020

Mechanisms of acquired resistance to RETi



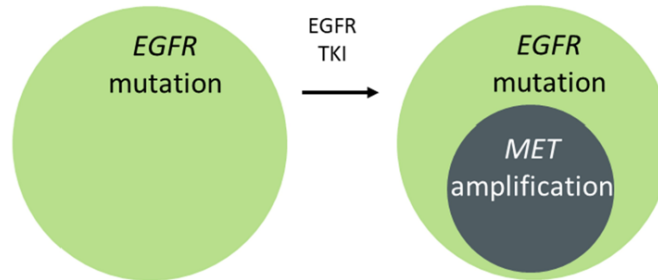
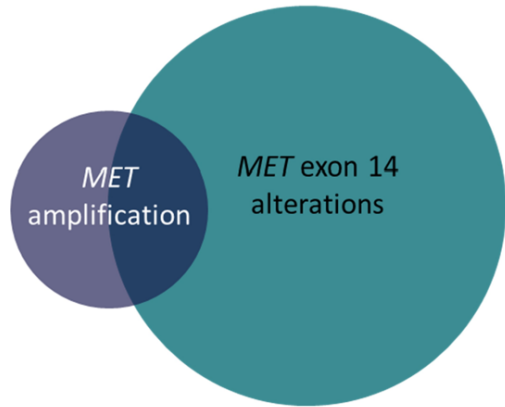
- Acquired selpercatinib resistance is driven by **MAPK pathway reactivation**, on- and off-target pathway reactivation via **secondary RET mutations** or **MET amplifications**
- Multiple distinct mechanisms are often observed in the same patient (**polyclonal resistance**)

Rosen EY et al, Nature Communications 2022

MET alterations: still a diagnostic challenge but..

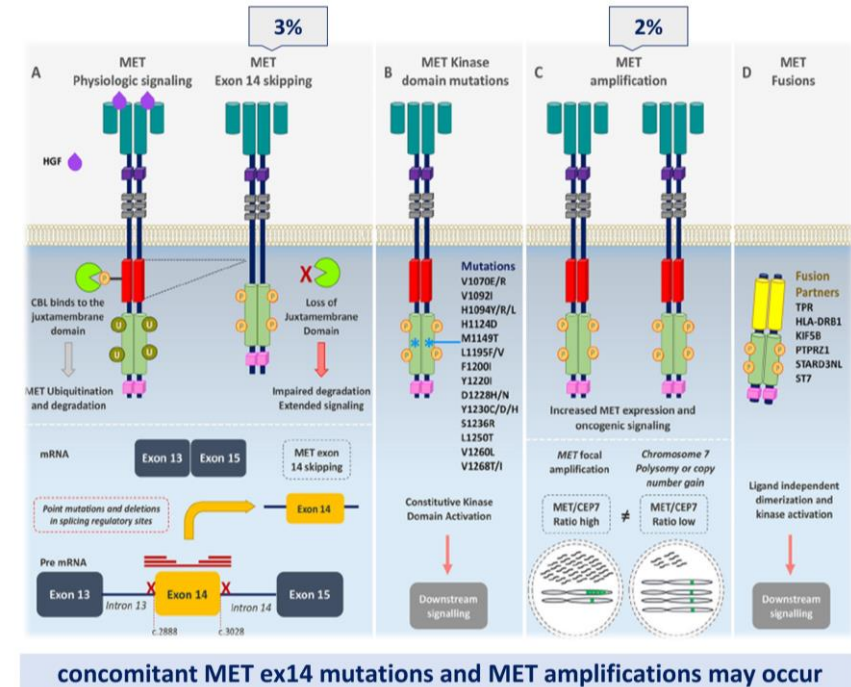
MET as a primary driver

MET as a secondary/co-driver



	Lu S, ASCO 2020	Paik PK, NEJM 2020	Drilon A, Nat Med 2020	Wolf J, ASCO 2021
METex14	-RT-PCR -NGS Geneseeq Tetradecan Panel	-NGS Guardant 360 (cfDNA) -NGS Oncomine Focus Assay (tissue)	-RT-PCR -NGS, DNA or RNA-based (different panels)	-RT-PCR -NGS FoundationONE CDx

- MET exon 14 skipping mutation on tissue, blood, both
 - DNA or **RNA** NGS
 - RT-qPCR
 - Sanger sequencing



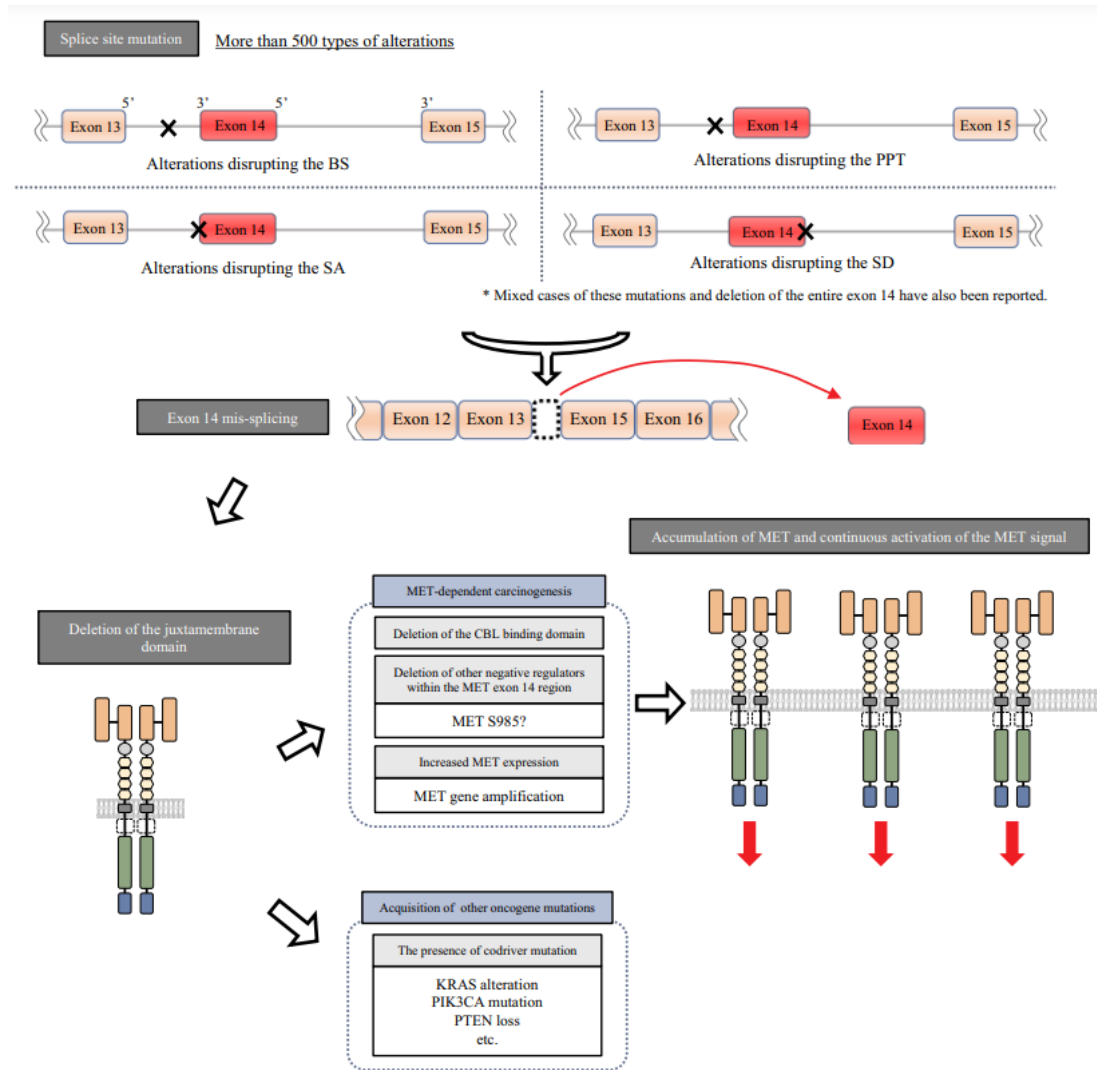
- MET amplification
 - FISH
 - NGS

Optimal copy number cut off tbd
Liquid biopsy: sensitivity, specificity

	Le X, ASCO 2021	Camidge R, JTO 2020	Wolf J, NEJM 2020
MET ampl	-NGS Guardant 360 (cfDNA): GCN ≥ 2.5	- FISH: MET/CEP7 ≥ 1.8 - NGS FoundationONE CDx (tissue): GCN ≥ 6	-NGS FoundationONE CDx (tissue): GCN ≥ 6

Modified from Drilon A and Pasello G, ASCO 2021

MET exon 14 skipping mutations



Up to 4% of non-squamous NSCLCs

- older patients, more smokers
- sarcomatoid tumour
- highly heterogeneous
- up to 20% with concurrent MET amplification

... we finally MET the right combo target-drugs

MET exon 14sk	N of patients	Line of treatment	RR (%)	DOR (months)	icRR (%)	PFS (months)	OS (months)	TR-AEs discontinuation
CRIZOTINIB	65	1 >1	25 36.6	9.1 (overall)	20 [§]	7.3 (overall)	20.5 (overall)	7%
CAPMATINIB	63	1 2	65.6 51.6	12.6 9.7	54	10.8 6.9	20.8* 13.6*	11%
TEPOTINIB	152	1 >1	43 43	10.8 11.1	55	8.5 (overall)	17.1 (overall)	11%
SAVOLITINIB	70	1 >1	54.4 46	6.8 NR	NA	5.6 13.8	NA	14.3%

MET <i>de novo</i> amplification	N of patients	Line of treatment	RR (%)	DOR (months)	PFS (months)	OS (months)	TR-AEs discontinuation
CRIZOTINIB	21	≥1	38.1	5.2	6.7	11.4	10.5%
CAPMATINIB	84	1 >1	40 29	7.5 8.3	4.2 4.1	NA	11%
TEPOTINIB	24	1 >1	71.4 28-30	NR	4.2 (overall)	NA	0%

*Data not mature for expansion cohorts; § retrospective data, not reported in Profile 1001

NR: not reached; NA: not available

Wolf J, 2021 ASCO meeting; Le X, 2021 ASCO meeting; Drilon A Nat Med2020; Camidge R JTO 2021; Lu S, 2020 ASCO meeting; Paik PK NEJM 2020; Wolf J NEJM 2020; Offin M, JCO Precis Oncol 2020

CAPMATINIB

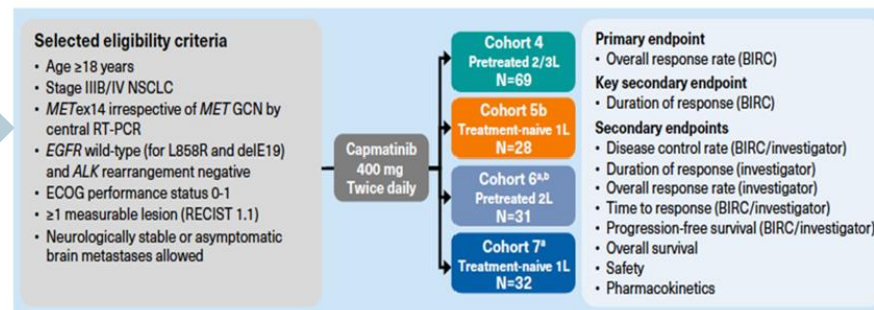
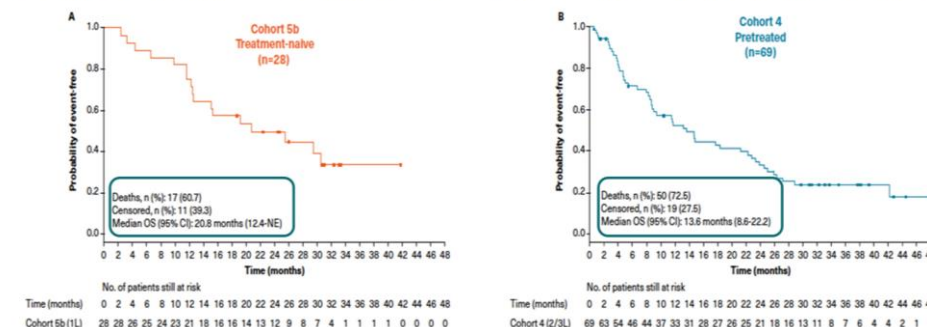


Figure 3. (A) Overall survival for treatment-naïve patients with METex14 NSCLC in Cohort 5b. (B) Overall survival for pretreated (2/3L) patients with METex14 NSCLC in Cohort 4



Wolf J et al, ASCO 2021

TEPOTINIB

- Key inclusion criteria**
- Advanced NSCLC (EGFR/ALK wild-type, all histologies)
 - Liquid or tissue biopsy MET alterations (central lab)
 - 1L, 2L or 3L treatment setting
 - Prior immunotherapy allowed
 - ECOG PS 0 or 1

Cohort A METex14 skipping
Cohort B METamp*
Cohort C METex14 skipping

Tepotinib 500 mg[†] once daily

Selected endpoints

- Primary:**
- ORR by IRC (RECIST v1.1)
- Secondary:**
- ORR by investigator, DOR, PFS, OS, safety

		Overall (n=24)	1L (n=7)	2L (n=10)	3L (n=7)
Best overall response, n (%)	PR	10 (41.7)	5 (71.4)	3 (30.0)	2 (28.6)
	SD	1 (4.2)	0	1 (10.0)	0
	PD	5 (20.8)	1 (14.3)	2 (20.0)	2 (28.6)
	NE	8 (33.3)	1 (14.3)	4 (40.0)	3 (42.9)
ORR, n (%) [95% CI]		10 (41.7) [22.1, 63.4]	5 (71.4) [29.0, 96.3]	3 (30.0) [6.7, 65.2]	2 (28.6) [3.7, 71.0]

Le X et al, ASCO 2021

MET exon 14 skipping mutations: new drug on the horizon [SCC244]

GLORY

Key patient inclusion criteria

- Locally advanced or metastatic NSCLC
- METex14 skipping mutations
- ≤2 prior systemic therapies or no prior systemic therapy
- ECOG PS 0–1

(n=73)

SCC244
300 mg/day q3w

PD/toxicity

Primary endpoint

- ORR (BIRC, RECIST v1.1.)

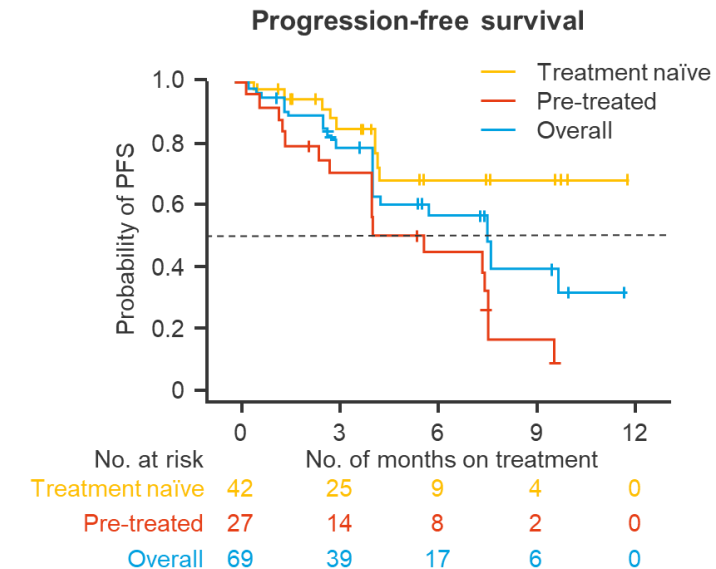
Secondary endpoints

- DoR, DCR, TTR, PFS, OS, safety

Efficacy analysis set	Treatment-naïve (n=42)	Previously treated (n=27)	All (n=69)
ORR, % (95%CI)	66.7 (50.5, 80.4)	51.9 (31.9, 71.3)	60.9 (48.4, 72.4)
DCR, % (95%CI)	88.1 (74.4, 96.0)	74.1 (53.7, 88.9)	82.6 (71.6, 90.7)
mDoR, mo (95%CI)	NE (NE, NE)	5.1 (2.8, 8.2)	8.2 (4.8, NE)
PFS events, n (%)	9 (21.4)	17 (63.0)	26 (37.7)
mPFS, mo (95%CI)	NE (4.3, NE)	5.7 (2.8, 7.6)	7.6 (4.2, NE)

TRAEs, n (%)	All (n=73)
Any	71 (97.3)
Grade ≥3	32 (43.8)
Serious	13 (17.8)
Led to dose interruption	18 (24.7)
Led to dose reduction	21 (28.8)
Led to discontinuation	5 (6.8)

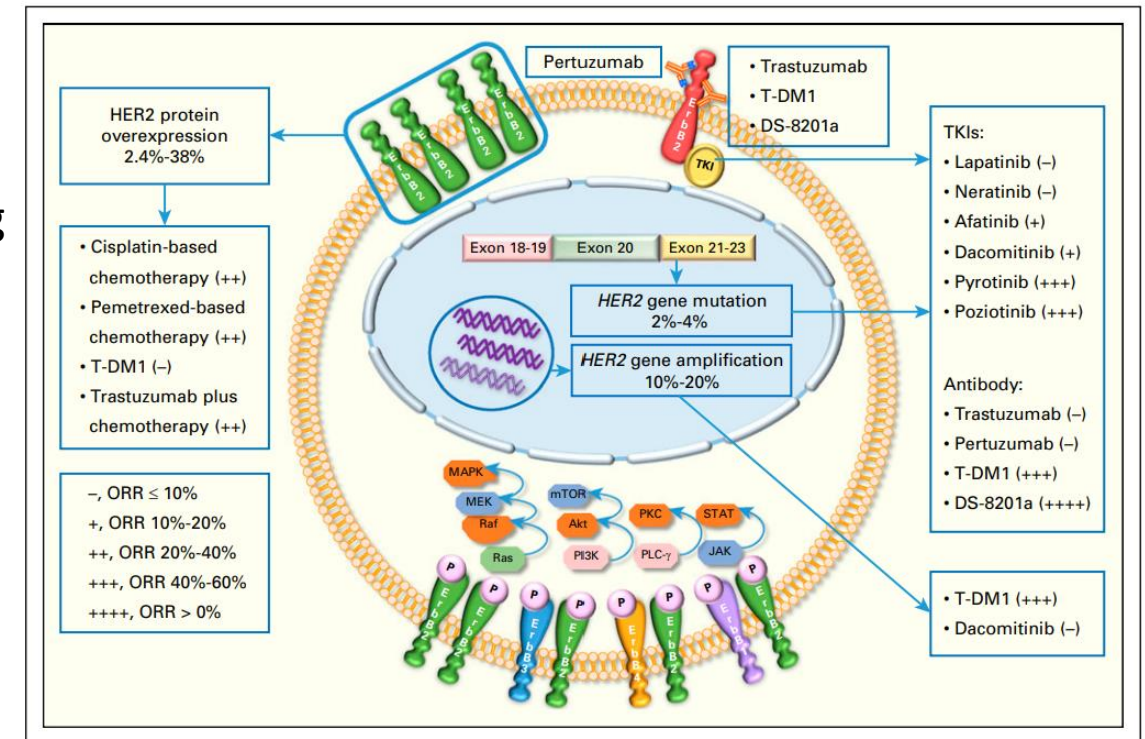
* Mainly peripheral edema, neutropenia, headache, rash



HER2 alterations: still a diagnostic challenge but...

Define “HER2 positivity” in lung cancer:

- **Overexpression** (2+/3+) in 23-35%, high (3+) in 4%
 - **Amplification** in 10-20%, high in 7% (also in EGFR)
 - **Mutations** in 2-4%, mainly insertions in exon 20-21 causing constitutive TKI activation
-
- *The association between amplification and overexpression is controversial*
 - *Lack of standardized testing techniques and cut-offs for defining HER2 positivity in lung cancer*



Zhao and Xia, JCO Precision Oncology 2020

Patients harbouring HER2 alterations more frequently **women, young, never smoker, and mainly adenocarcinomas**

HER2 alterations: should we just copy the example of breast?

- **TKIs** bind to the tyrosine domain of activated HER2 protein → several pan-HER TKIs have been tested for NSCLC with modest results (ORR 20-30%) and relevant toxicities.
- **Chemotherapy** still a standard of care (waiting for..)
- The efficacy of the **monoclonal antibodies pertuzumab and trastuzumab** has also been studied, alone or conjugated to cytotoxic drugs such as trastuzumab emtansine (T-DM1).

Trastuzumab/pertuzumab/docetaxel

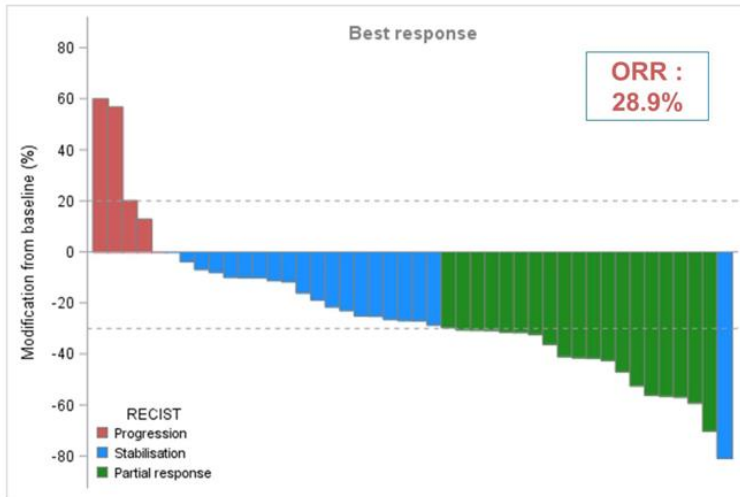
N=46 (HER2-mutant/insertions)

ORR 29%

mPFS 6.8 months

mOS 17.6 months

TRAEs G3-4 64%



Mazieres J et al, ASCO 2021

T-DM1

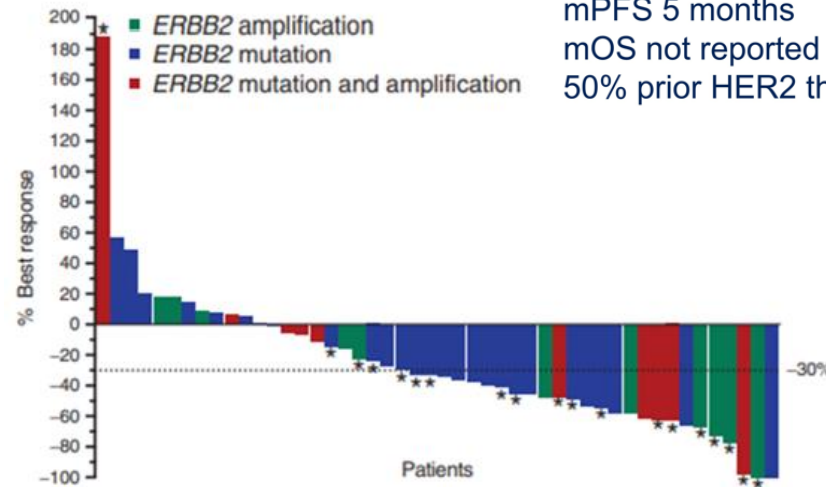
N=38 (HER2-mutant)

ORR 50% (95% CI 31-69%) RECIST/mPERCIST

mPFS 5 months

mOS not reported

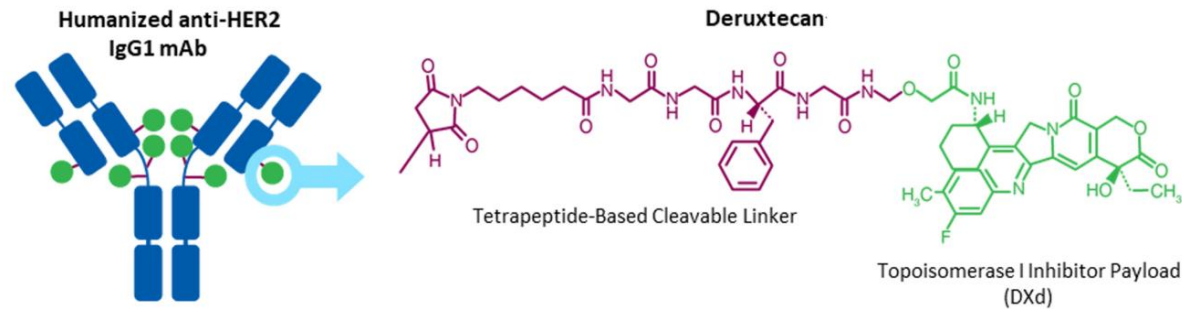
50% prior HER2 therapy



Li, BT; et al. Cancer Discovery 2020

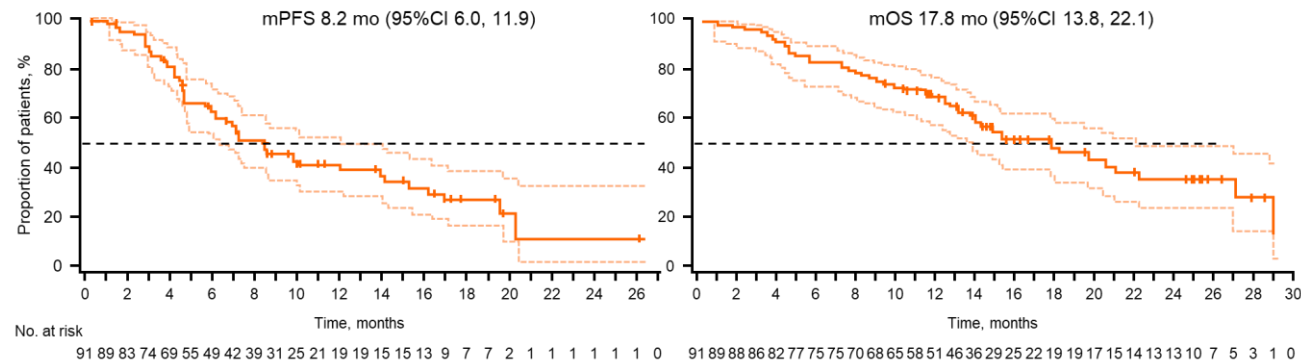
The revolution of ADC for *HER2* mutations

TRASTUZUMAB DERUXTECAN

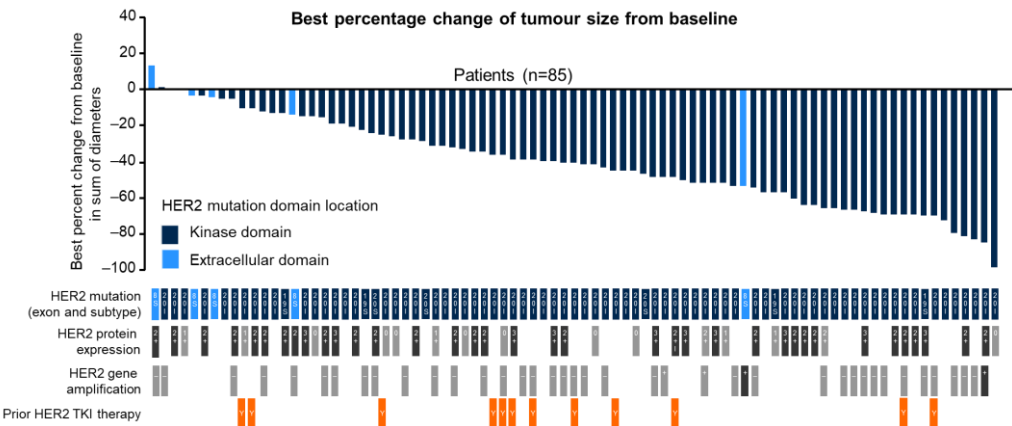


Progression-free survival

Overall survival

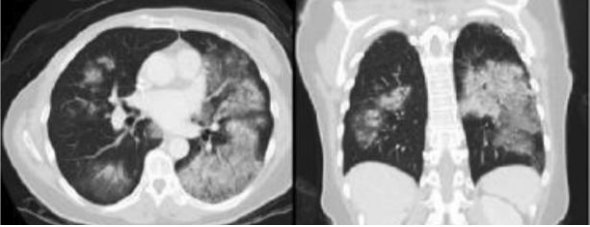


Trastuzumab deruxtecan (n=91)	
Confirmed ORR, n (%) [95%CI]	50 (54.9) [44.2, 65.4]
BOR, n (%)	
CR	1 (1.1)
PR	49 (53.8)
SD	34 (37.4)
PD	3 (3.3)
NE	4 (4.4)
DCR, n (%) [95%CI]	84 (92.3) [84.8, 96.9]
mDoR, mo (95%CI)	9.3 (5.7, 14.7)
Median follow-up, mo (range)	13.1 (0.7–29.1)



Li BT et al. NEJM 2022

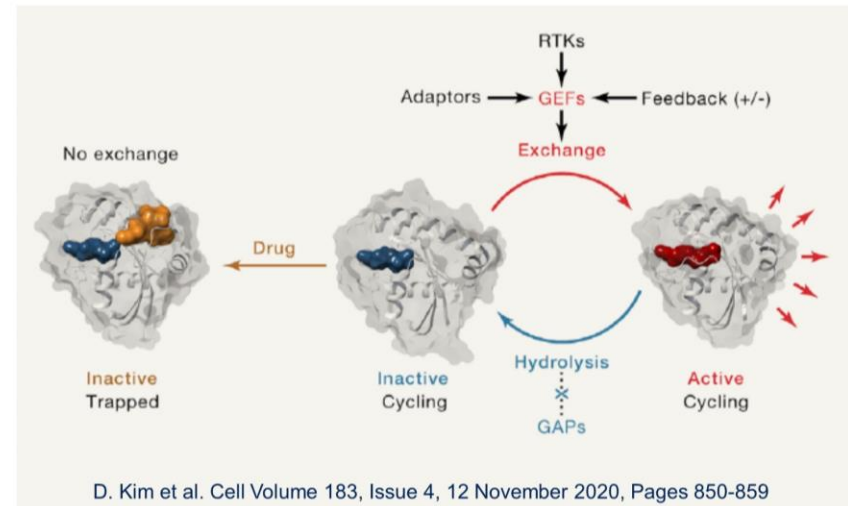
Azar I et al. Lung Cancer: Targets and Therapy 2021



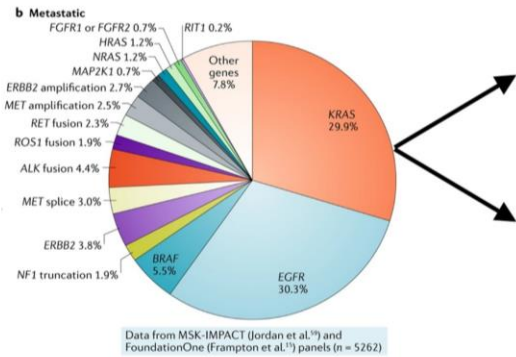
KRAS p.G12C mutations [Drugging the 'undruggable' KRAS]

KRAS G12C Inhibitors - Mechanism of Action

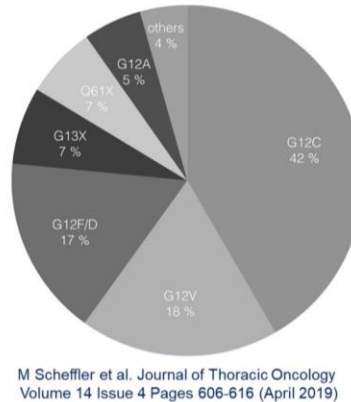
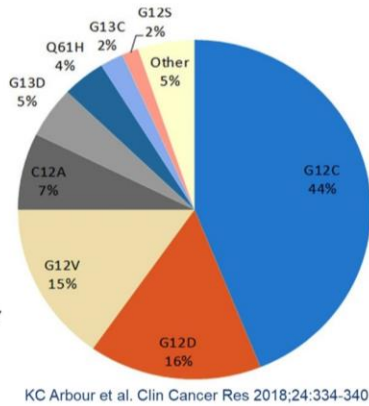
- Novel class of drugs → these are targeted therapies but they are not TKIs
- Allele-specific inhibitors targeting the Cysteine (C) residue.
- The inhibitors bind covalently to the mutant cysteine residue and occupy a pocket in the switch II region (SIIP) when KRAS G12C is in its inactive GDP-bound state (inactive-state selective drugs).



2021: Molecular Subsets of Lung Cancer



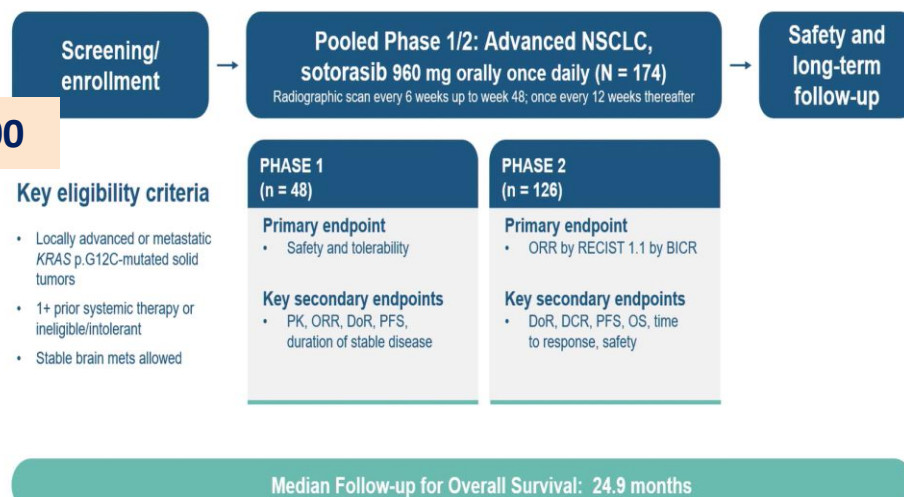
Skoulidis, F., Heymach, J.V. Nat Rev Cancer 19, 495–509 (2019).



- KRAS G12C most common KRAS variant
- 13% (1 in 8) of all lung adenocarcinomas
- Multiple KRAS G12C inhibitors being developed

The first-in-class in *KRAS p.G12C* mutations: sotorasib

CODEBREAK 100



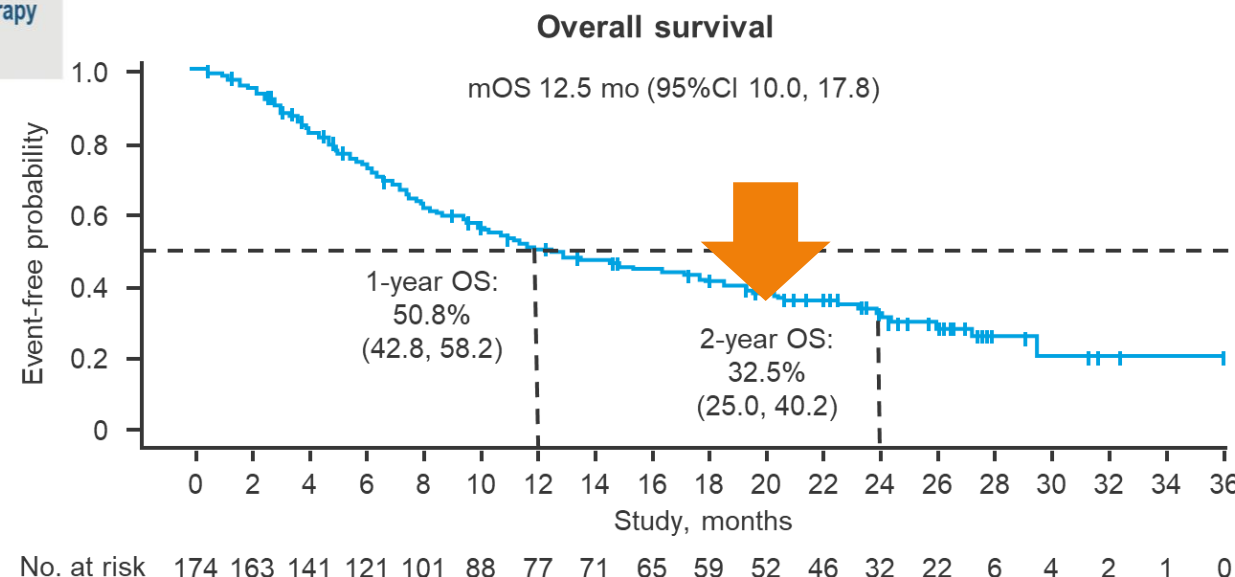
TRAEs, %	Overall (n=174)	Onset after 1 year (n=45)
Any	70	24
Grade 2	20	9
Grade 3	20	2
Grade 4	1	0
Led to treatment modification	22	2
Led to discontinuation	6	0

* Mainly diarrhea, ALT/AST increase, nausea and fatigue

Global study with 83% of enrolled patients having received prior platinum-based chemotherapy and anti-PD-(L)1 therapy

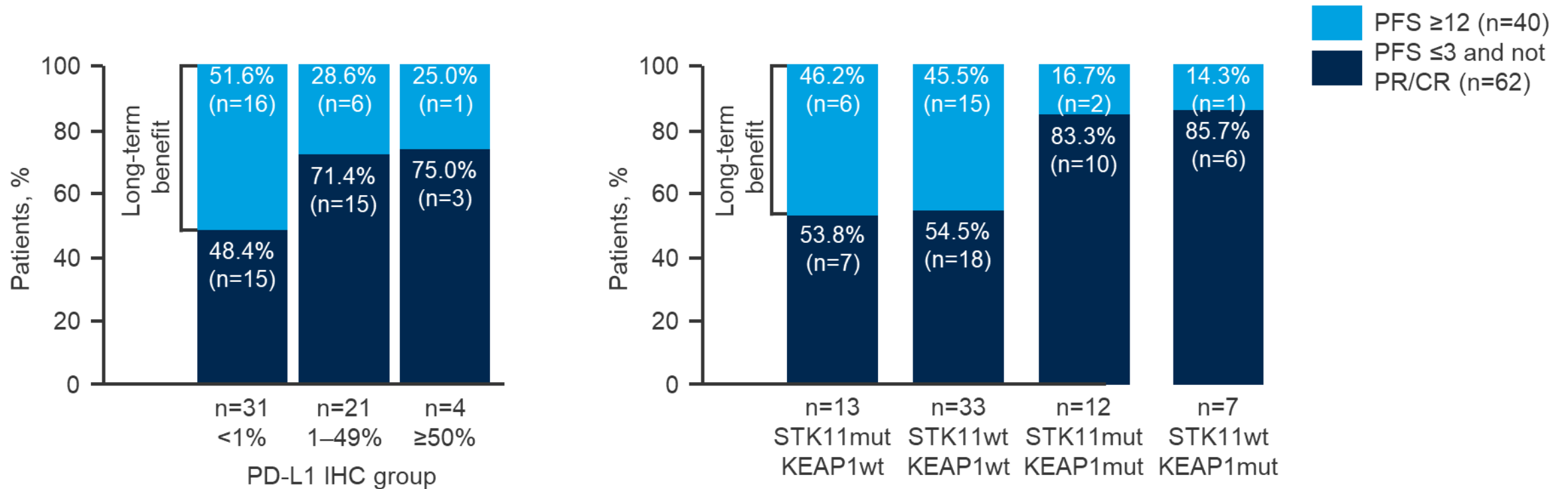
Data cutoff: February 22, 2022

Response	n=172
ORR, % (95%CI)	40.7 (33.3, 48.4)
BOR, n (%)	
CR	5 (2.9)
PR	65 (37.8)
SD	74 (43.0)
PD	23 (13.4)
NE	5 (2.9)
DCR, % (95%CI)	83.7 (77.3, 88.9)
mPFS, mo (95%CI)	6.3 (5.3, 8.2)
mDoR, mo (95%CI)	12.3 (7.1, 15.0)



Dy GK et al, AACR 2022

The first-in-class in *KRAS* p.G12C mutations: sotorasib



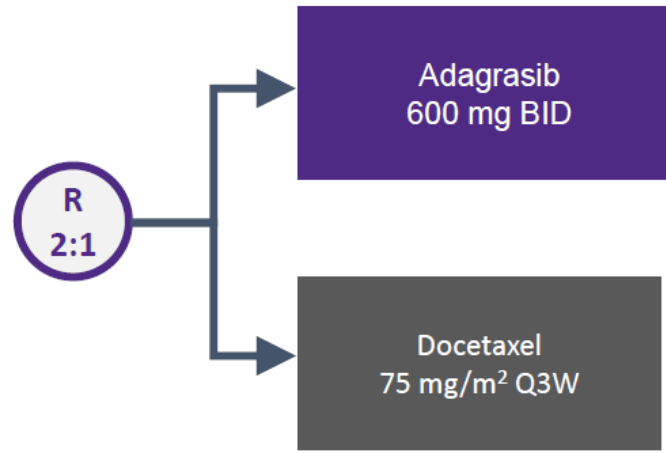
- Long-term benefit was seen regardless of PD-L1 expression, *KRAS* G12C allele frequency, co-mutations
- Patients with long-term benefit had lower plasma ctDNA at baseline and these levels correlated with tumor burden

KRAS p.G12C mutations: phase 3 trials are ongoing in 2nd line

KRYSTAL-12: Phase 3 Randomized, Open-Label Trial of 2L Adagrasib vs Docetaxel in Patients With Previously Treated NSCLC With *KRAS*^{G12C} Mutation^{1,2}

Key Eligibility Criteria (n=452)

- NSCLC with *KRAS*^{G12C} mutation based on sponsor-approved test
- ECOG PS of 0 or 1
- No active brain metastases
- Prior treatment with platinum-based regimen and a checkpoint inhibitor
- No prior treatment with a *KRAS* inhibitor



Endpoints

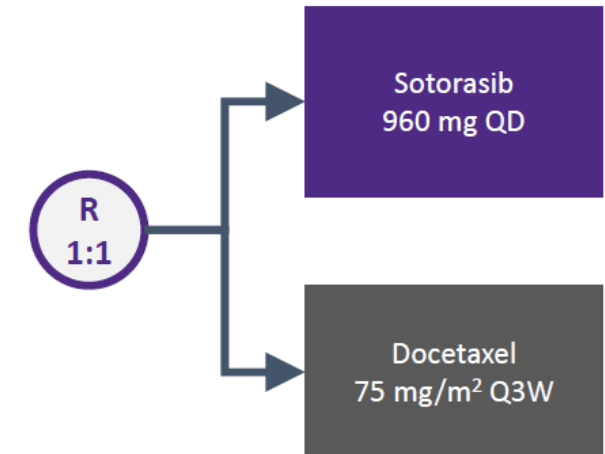
Primary: PFS, OS

Secondary: AEs, ORR, DOR, PROs, 1-year survival rate, PK

CodeBreak 200: Phase 3 Randomized, Open-Label Trial of Sotorasib vs Docetaxel in Patients with Previously Treated Metastatic NSCLC with *KRAS*^{G12C} Mutation^{3,4}

Key Eligibility Criteria (n=345)

- NSCLC with *KRAS*^{G12C} mutation confirmed through central testing
- ECOG PS of 0 or 1
- No active brain metastases
- Prior treatment with platinum-based regimen and a checkpoint inhibitor
- No prior treatment with a *KRAS*^{G12C} inhibitor

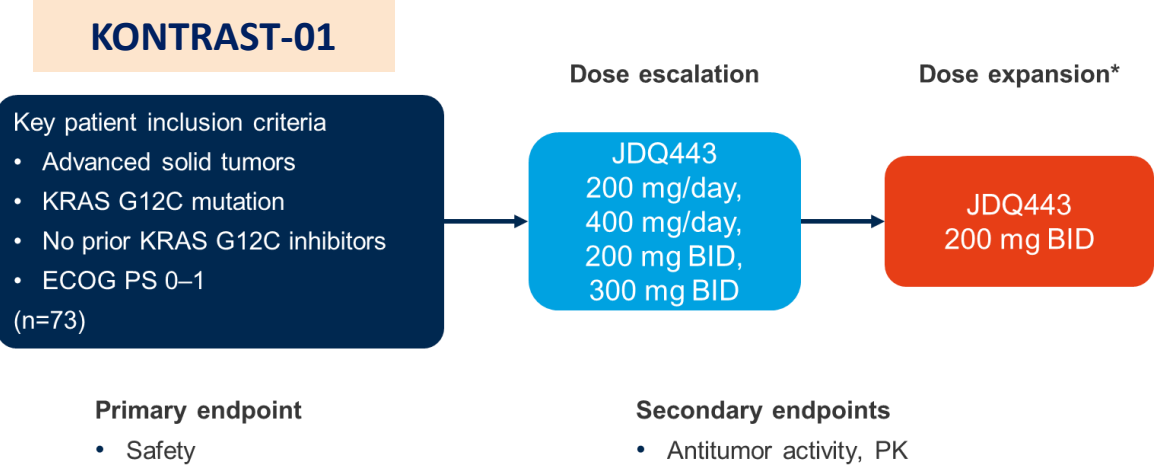


Endpoints

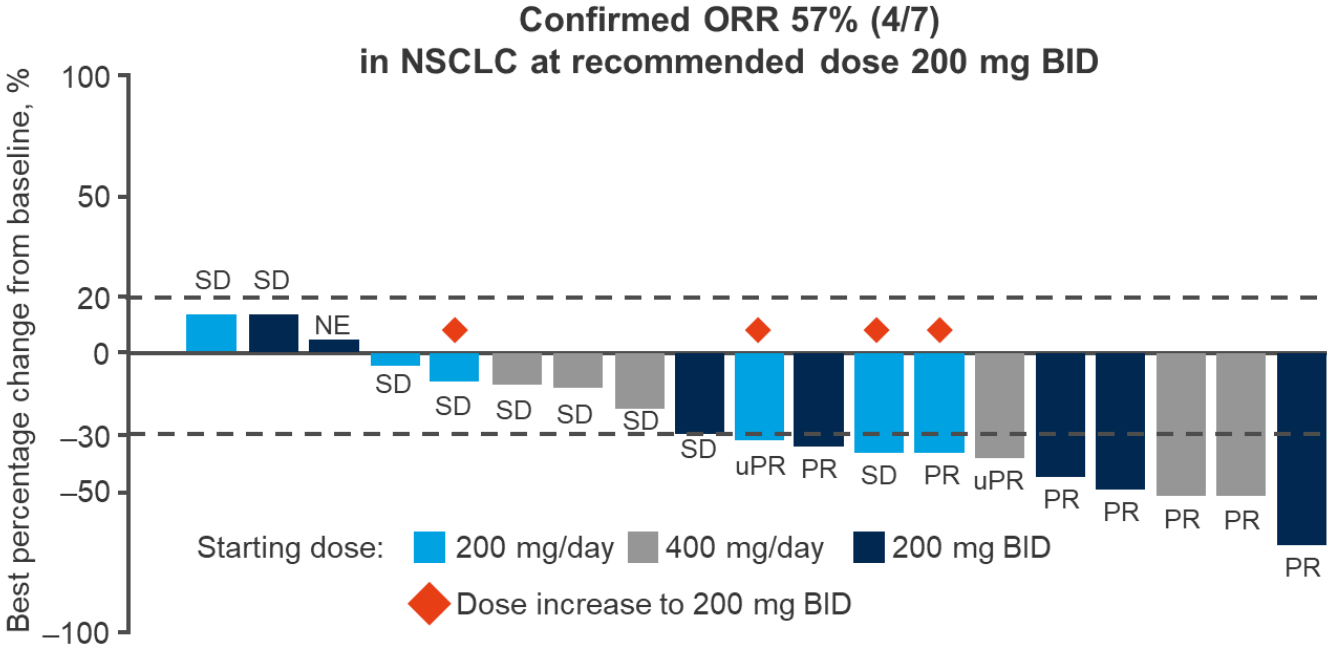
Primary: PFS

Secondary: OS, ORR, DOR, TTR, DCR, PRO, QoL, PK

KRAS p.G12C mutations: new drug on the horizon [JDQ443]



Grade ≥3 TRAEs, n (%)	JDQ443 200 mg BID (n=11)	JDQ443 all patients (n=39)
Any	0	5 (12.8)
Photosensitivity reaction	0	2 (5.1)
Fatigue	0	1 (2.6)
Neutropenia	0	1 (2.6)



BoR, n (%)	Patients with NSCLC (n=20)
PR (confirmed)	7 (35)
SD	11 (55)
PD	0
NE	2 (10)
ORR (confirmed and unconfirmed)	9 (45)
ORR (confirmed)	7 (35)

How did we arrive here & where are we going?



❖ Enlarging the *family* of oncogene-addicted

- New drugs and targets on the horizon
- Understanding the *difficult* targets
- Drugging the *undruggable(s)*

❖ Deciphering resistance & Patients' selection

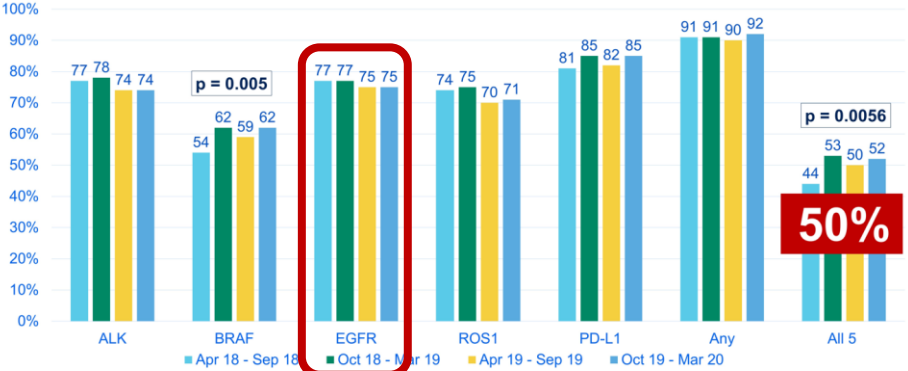
- In both oncogene-addicted and not oncogene-addicted

❖ Bring innovation as early as possible

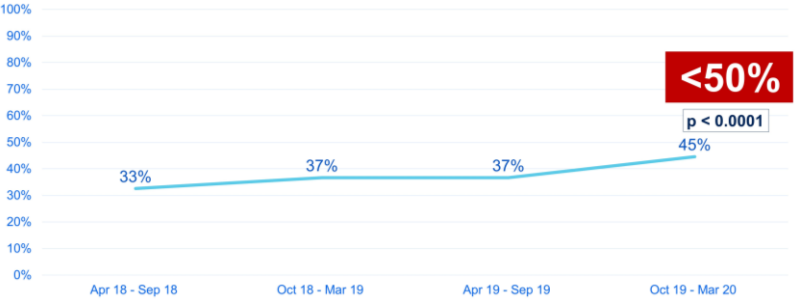
- Oncogene-addicted vs. not?
- Neoadjuvant vs/with adjuvant? For how long? Combos?

The *primum movens* is always the Test!!

Biomarkers testing rates over time



NGS testing rates over time



Robert NJ et al, ASCO 2021

EGFR mutation testing data collected in 36 Italian institutions in 2020



Malapelle U et al, CROH 2021





ATLAS

BOARD SCIENTIFICO

Il coordinamento delle attività dei due gruppi è affidato a un board scientifico.



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CARCINOMA POLMONARE: QUALI NOVITÀ NEL 2022?

Thanks!
😊