

# Other Oncogenic Drivers (BRAF, MET, RET, HER2, NTRK)

**David Planchard (MD, PhD)**

Head of the Thoracic Group

Department of Cancer Medicine

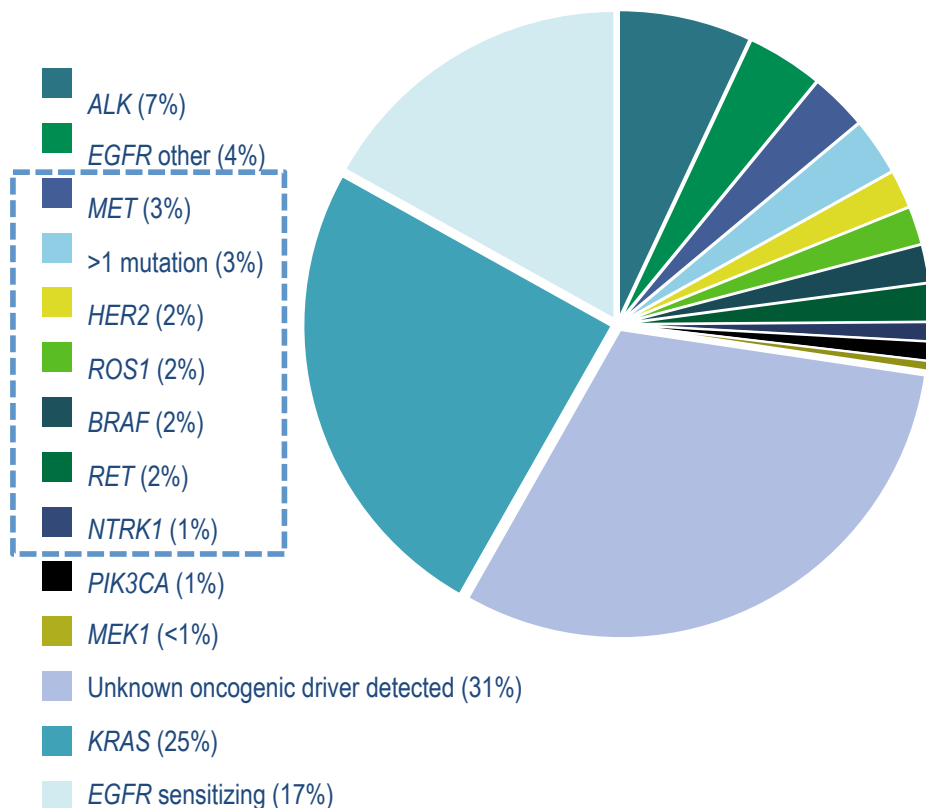
Gustave Roussy – Villejuif (France)



# Disclosure Slide

- **Honoraria:** AstraZeneca, Bristol-Myers Squibb, Boehringer Ingelheim, Celgene, Eli Lilly, Merck, MSD Oncology, Novartis, Pfizer, prIME Oncology, Roche
- **Consulting, advisory role or lectures:** AstraZeneca, Bristol-Myers Squibb, Boehringer Ingelheim, Celgene, Eli Lilly, MSD Oncology, Novartis, Pfizer, prIME Oncology, Roche
- **Travel, Accommodations, Expenses:** AstraZeneca, Roche, Novartis, prIME Oncology, Pfizer

# Great advances have been made in lung cancer therapy: targeting of oncogenic drivers



## EGFR sensitizing

Gefitinib; Erlotinib; Afatinib; Osimertinib; Dacomitinib

## ALK

Crizotinib; Alectinib; Ceritinib; Lorlatinib; Brigatinib

## ROS1

Crizotinib; Cabozantinib; Ceritinib; Lorlatinib; Entrectinib; Roprotrectinib, DS-6051b

## BRAF

Vemurafenib; Dabrafenib; Dabrafenib + Trametinib

## MET

Crizotinib; Cabozantinib; Capmatinib; Savolitinib; Tepotinib; Merestinib; Glesatinib

## HER2

Trastuzumab emtansine; Afatinib; Neratinib-temsirolimus; Dacomitinib; Poziotinib; XMT-1522; TAK-788; DS-8201a,

## RET

Cabozantinib; Alectinib; Apatinib; Vandetanib; sunitinib; Ponatinib; Lenvatinib; BLU-667; LOXO-292

## NTRK1

Entrectinib; LOXO-101 (larotrectinib); loxo-195; DS-6051b; ropotrectinib

## PIK3CA

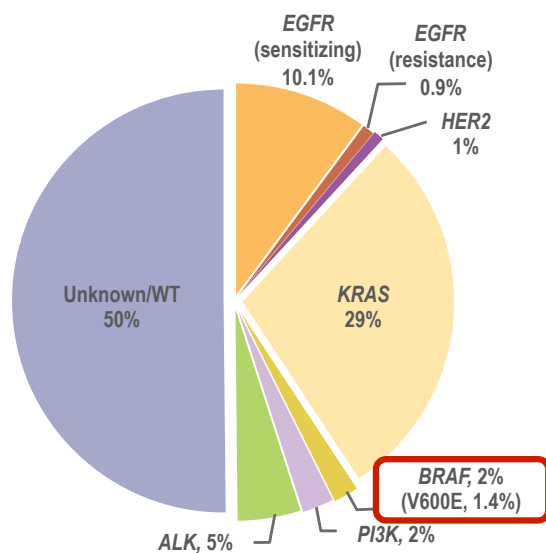
LY3023414; PQR 309

## MEK1

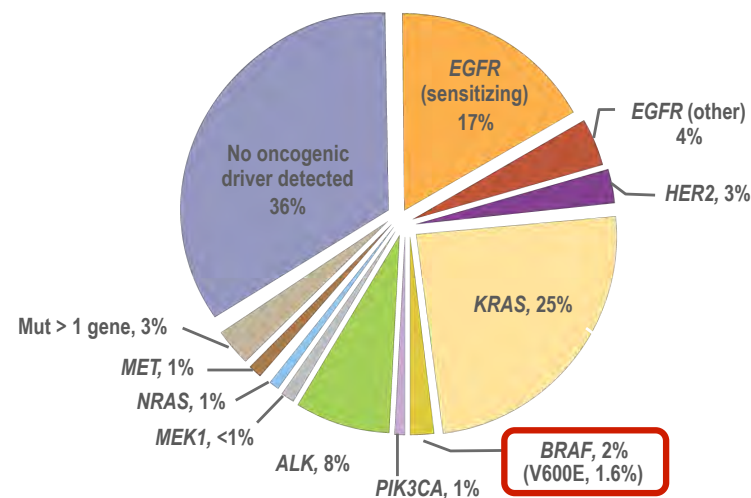
Trametinib; Selumetinib; Cobimetinib

# BRAF MUTATIONS IN NSCLC

**France<sup>1</sup>**  
NSCLC  
(Biomarkers France [IFCT]; N=17,664)



**US<sup>2</sup>**  
Adenocarcinoma  
(Lung Cancer Mutation Consortium; N=733)

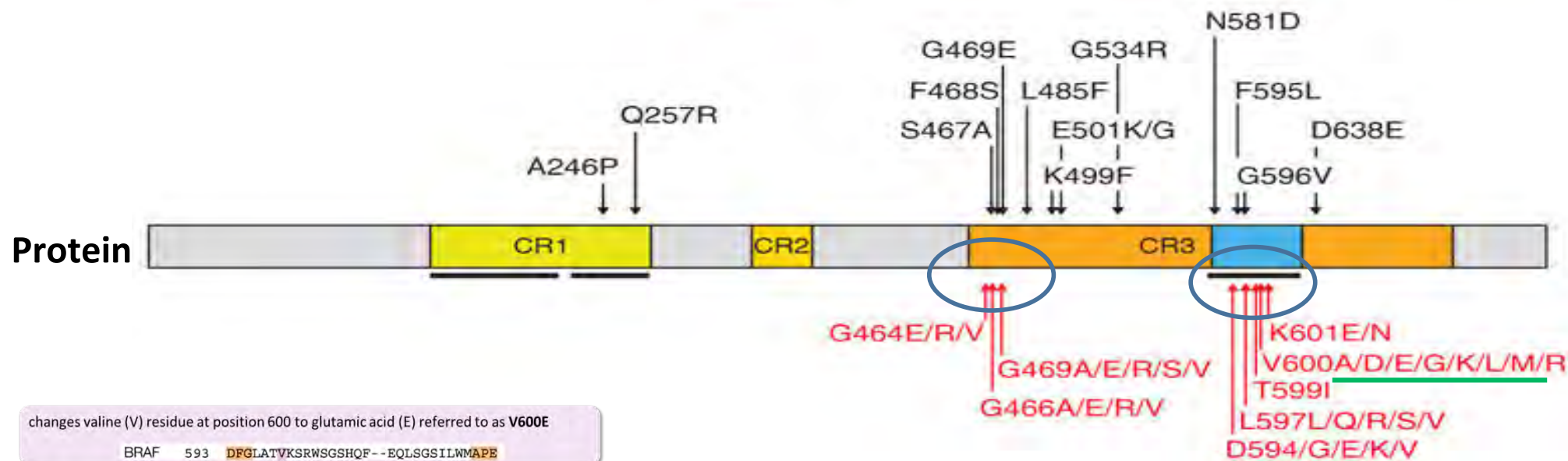


- NSCLC with *BRAF* V600E mutations has histological features suggestive of an aggressive tumor<sup>3</sup>
- Patients with *BRAF* V600E–mutant NSCLC demonstrated less-favorable outcomes with platinum-based chemotherapy<sup>3,4</sup>

1. Barlesi F *et al. Lancet* 2016;387:1415–1426; 2. Kris MG *et al. JAMA* 2014;311:1998–2006;  
3. Marchetti A *et al. J Clin Oncol* 2011;29:3574–3579; 4. Cardarella S *et al. Clin Cancer Res* 2013;19:4532–4540

# BRAF gene and protein structures with related biological aspects

18 exons and 17 introns spanning 200 Kb on the long arm of chromosome (7q34)



changes valine (V) residue at position 600 to glutamic acid (E) referred to as **V600E**  
 BRAF 593 **DFGLATVKS**RWSGSHQF--EQLSGSILWMA**PE**

Domain	RBD	Regulatory	Kinase domain
Function	RAS binding	AKT/SGK binding	P-loop and activation segment

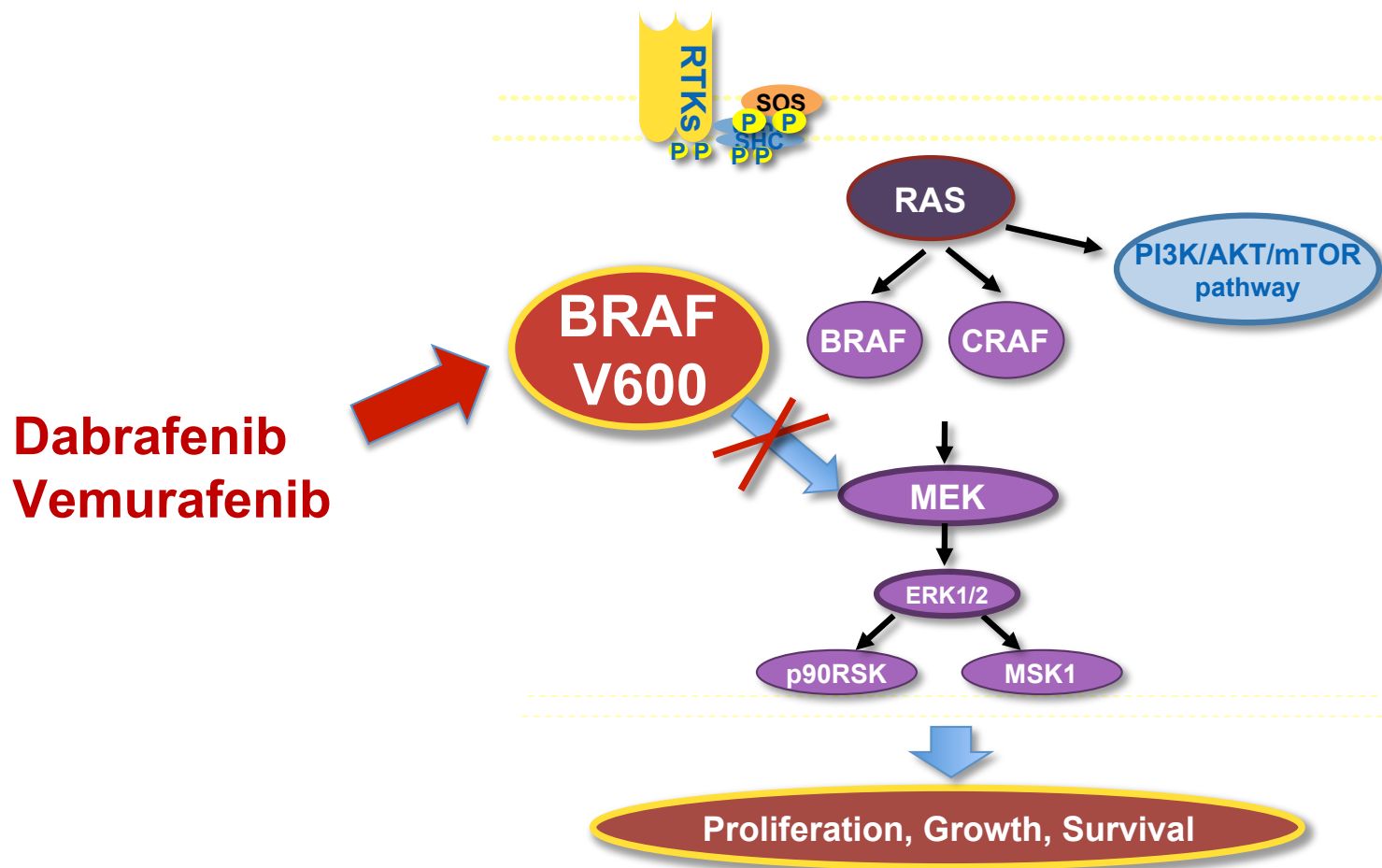
# BRAF-ASSOCIATED PATIENT CHARACTERISTICS

	<b>ALK<sup>1-4</sup></b>	<b>EGFR<sup>1,3,4,7</sup></b>	<b>KRAS<sup>4,7</sup></b>	<b>BRAF<sup>5-8</sup></b>
Age	Younger (~50)	Older (~60)	Older (~60)	Older (~65)
Male or female	None	Female predominant	Female predominant	None
Smoker or non-smoker	Never or light	Never or light	Heavy	Smoker and non-smoker
Histology	Adenocarcinoma	Adenocarcinoma	Adenocarcinoma	Adenocarcinoma
Pattern of spread	Pericardial,* pleural metastases,* liver,* intra- or extrathoracic lymph nodes,* CNS	Liver,* CNS	CNS	?

1. Shaw AT *et al. J Clin Oncol* 2009;27:4247–4253; 2. Wang Y *et al. PLoS One* 2014;9:e116017;  
 3. Tsao A *et al. J Thorac Oncol* 2006;1:231–239; 4. Doebele RC *et al. Cancer* 2012;118:4502–4511;  
 5. Kinno T *et al. Ann Oncol* 2014;25:138–142; 6. Cardarella S *et al. Clin Cancer Res* 2013;19:4532–4540;  
 7. Barlesi F *et al. Lancet* 2016;387:1415–1426; 8. Nguyen-Ngoc T *et al. J Thorac Oncol* 2015;10:1396–1403

\*Compared with triple-negative, wild-type patients

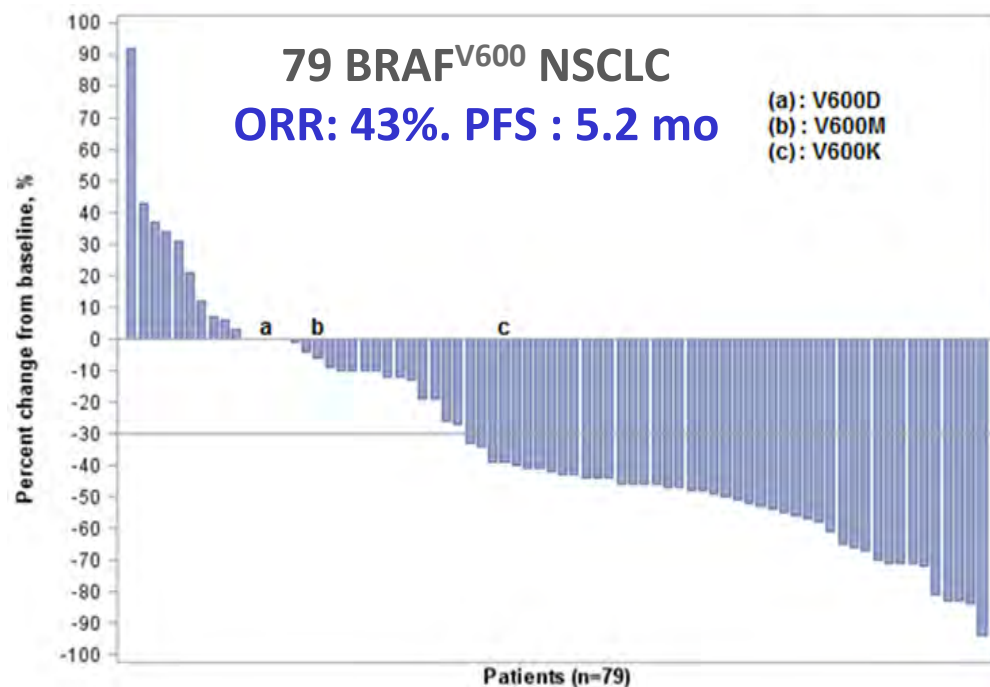
# Inhibition of BRAF V600 Kinase



# Vemurafenib in *BRAF* mutant NSCLC

## AcSé trial

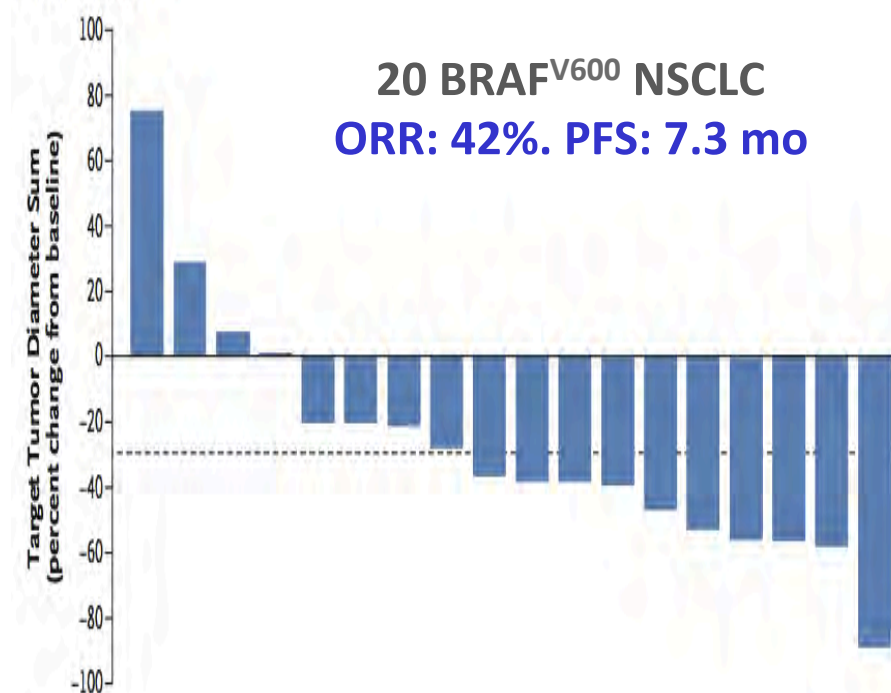
Vemurafenib



Mazières – WCLC 2018

## VE-Basket trial

Vemurafenib

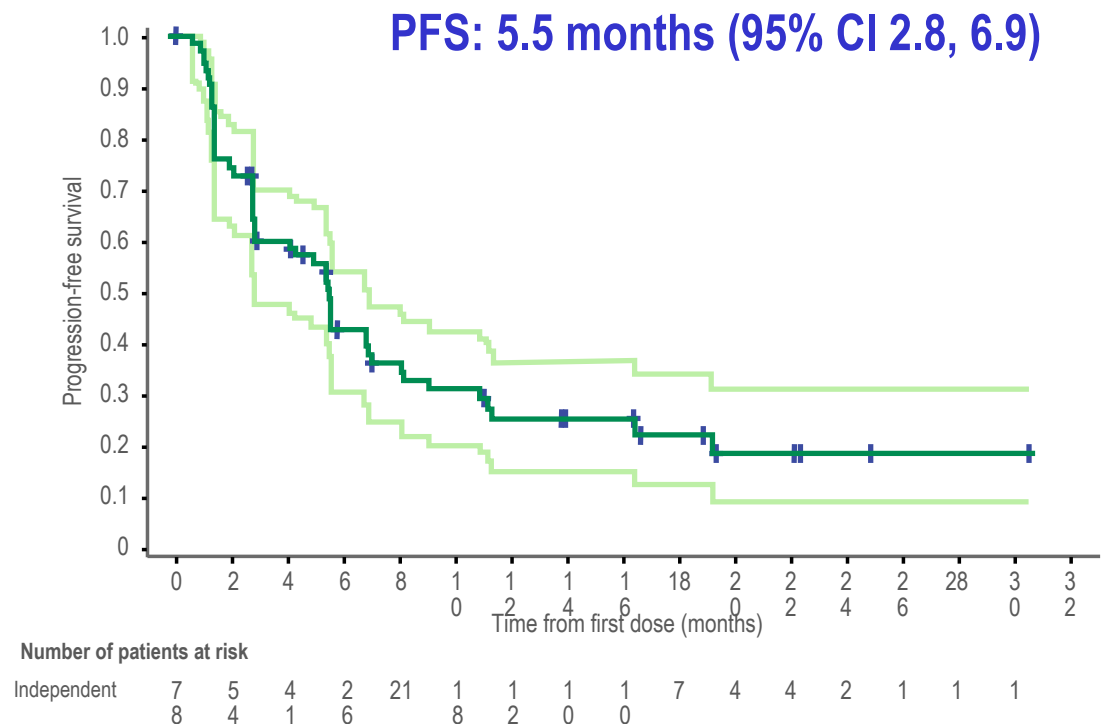
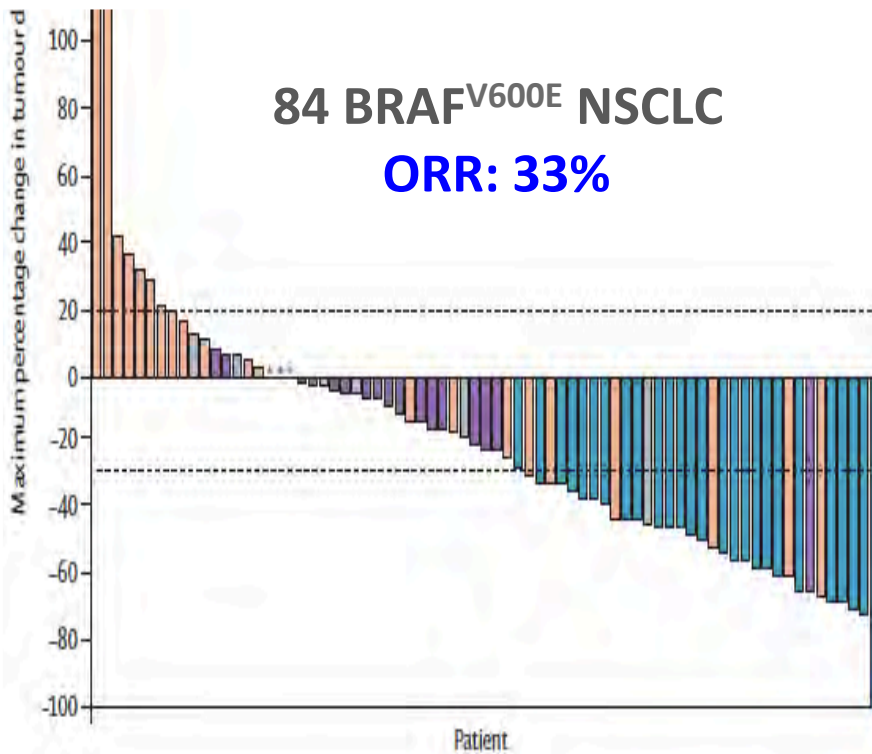


Hyman – NEJM 2015



# BRF113928 STUDY : DABRAFENIB IN BRAF MUTANT NSCLC IN 2<sup>ND</sup> LINE

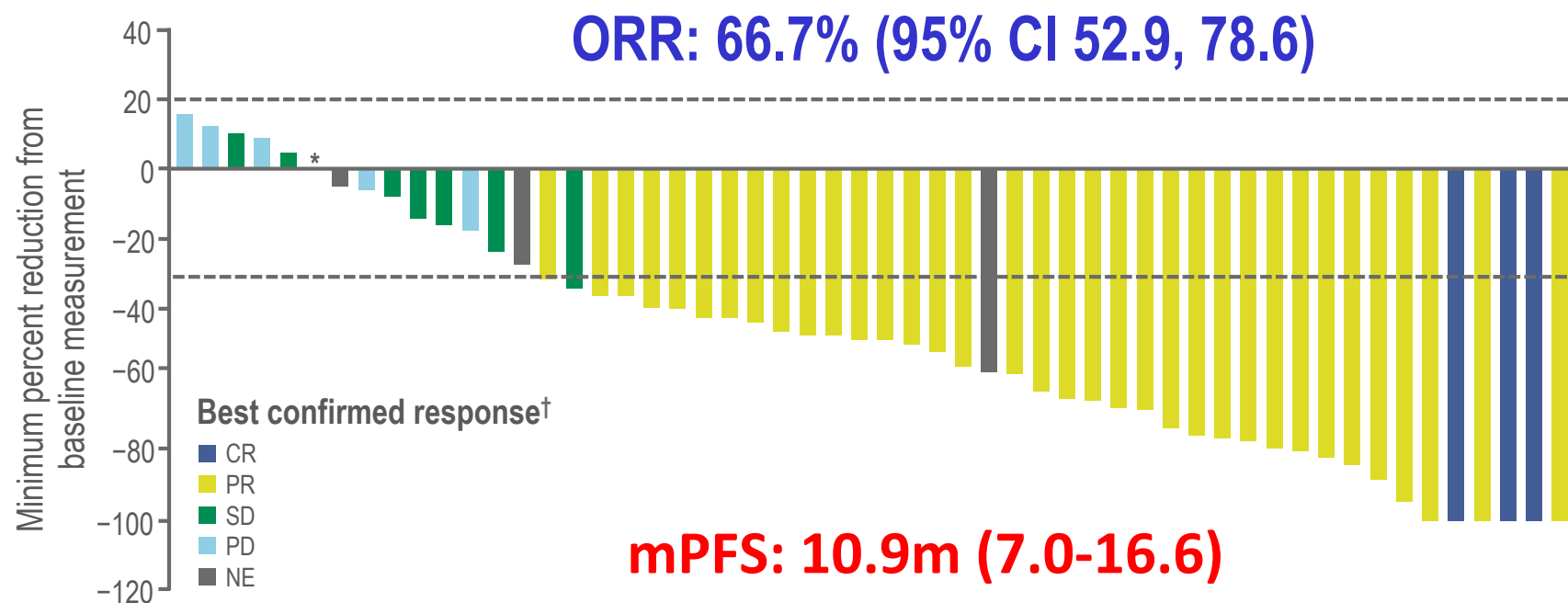
## Cohort A





# BRF113928 STUDY : MAXIMUM CHANGE IN TARGET LESION BY BEST CONFIRMED RESPONSE WITH **DABRAFENIB + TRAMETINIB** IN 2<sup>ND</sup> LINE

Cohort B

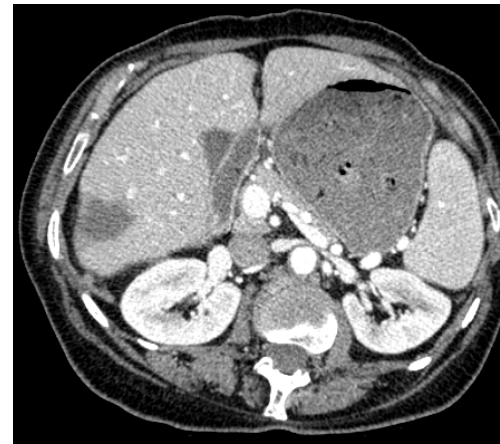
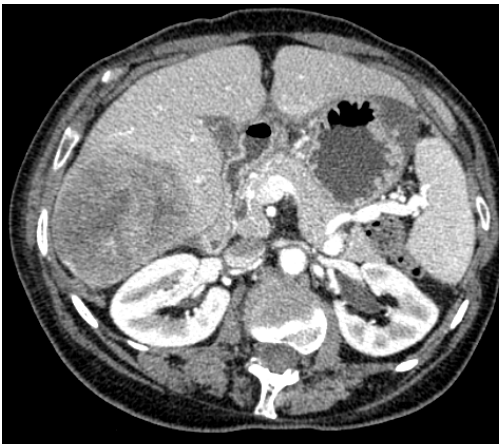


Planchard D *et al. Lancet Oncol* 2016;17:984–993;  
Planchard D *et al. J Clin Oncol* 2017;35(Suppl):Abst 9075

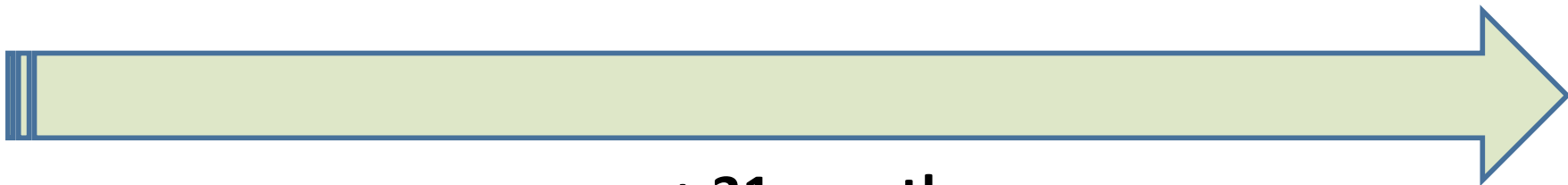


**The patient received the association:  
Dabrafenib (150mg twice a day) + Trametinib (2mg/day)**

July 2014



February 2017



**+ 31 months**

# BRAF mutated patients

Author	n	Drug	ORR	PFS (months)	OS (months)
Hyman (BASKET-trial)	20	Vemurafenib	42%	7.3	NR
Gautschi (EU-RAF, retrospective)	35	Vemurafenib	53%	5	10.8
Mazières (AcSé Vemu)	100	Vemurafenib	44.9%	5.2	9.3
Planchard (BRF cohort A)	78	Dabrafenib	33%	5.5	12.7
<b>Planchard (BRF cohort B)</b>	<b>57</b>	<b>Dabrafenib + trametinib</b>	<b>66.7%</b>	<b>10.9</b>	<b>12.7</b>
<b>Planchard (BRF cohort C)</b>	<b>36</b>	<b>Dabrafenib + trametinib 1L</b>	<b>64%</b>	<b>10.2</b>	<b>24.6</b>

## EMA and FDA approvals 2017

# BRAF non V600 cohort (AcSé Vemu)

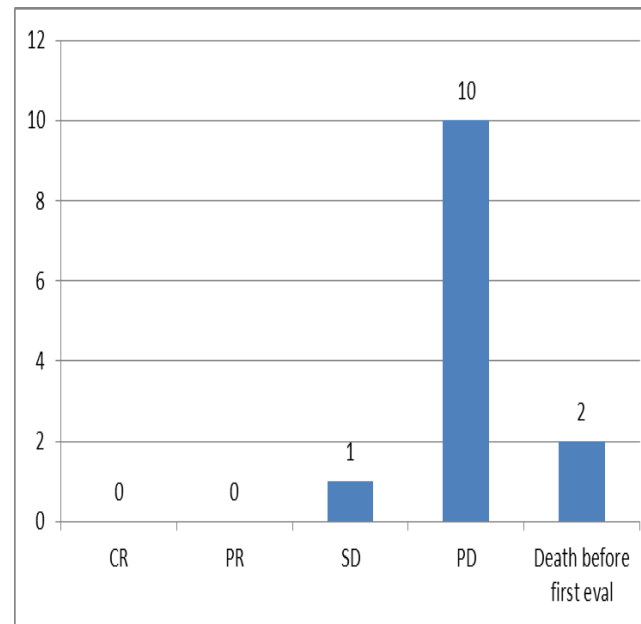
- Mean Bayesian Estimated Success rate : **5.9%** ; credibility 95%CI : [0.2%; 20.6%]
- Prob ORR < futility bound (10%): 81.5% - **study stopped**

## Non V600 mutations

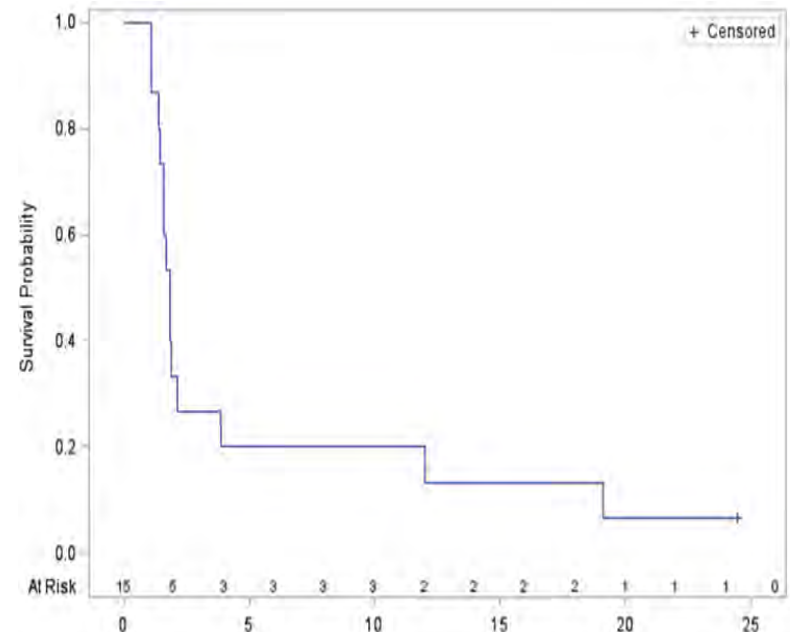
n = 17

G466A : n=1  
 G466V : n=3  
 G469A : n=3  
 G469V : n=1  
 N581S : n=3  
 G596R : n=1  
 K601E : n=3  
 K601N : n=2

**Response rate: 0%**



**PFS: 1.8 m. [1.4;2.1]**



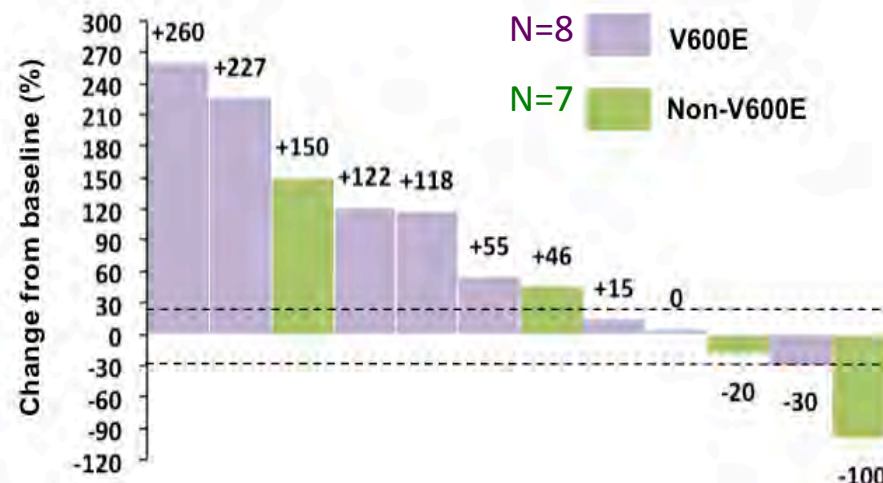
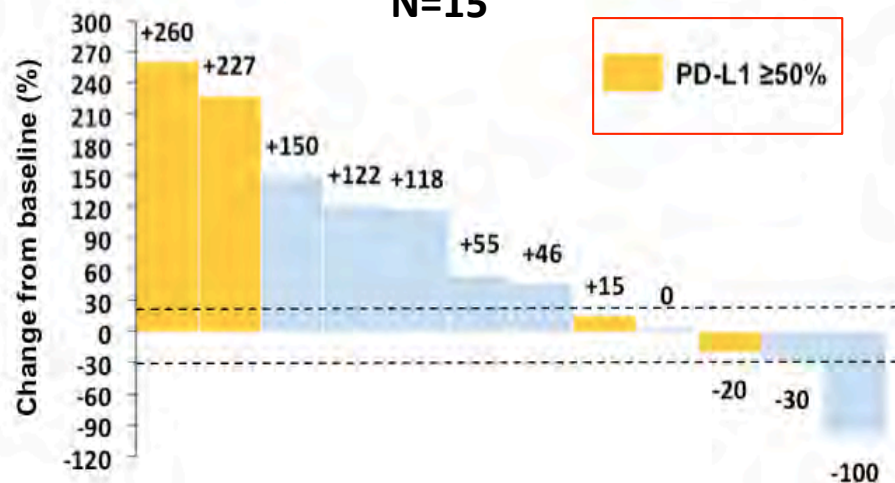
# BRAF and immunotherapy

45% of *BRAF*-mutant & high PD-L1 expression levels ( $\geq 50\%$  by 22C3 IHQ)

10% of cases associated with high tumor mutational burden ( $\geq 20$  Mb)

ORR: 17%

N=15



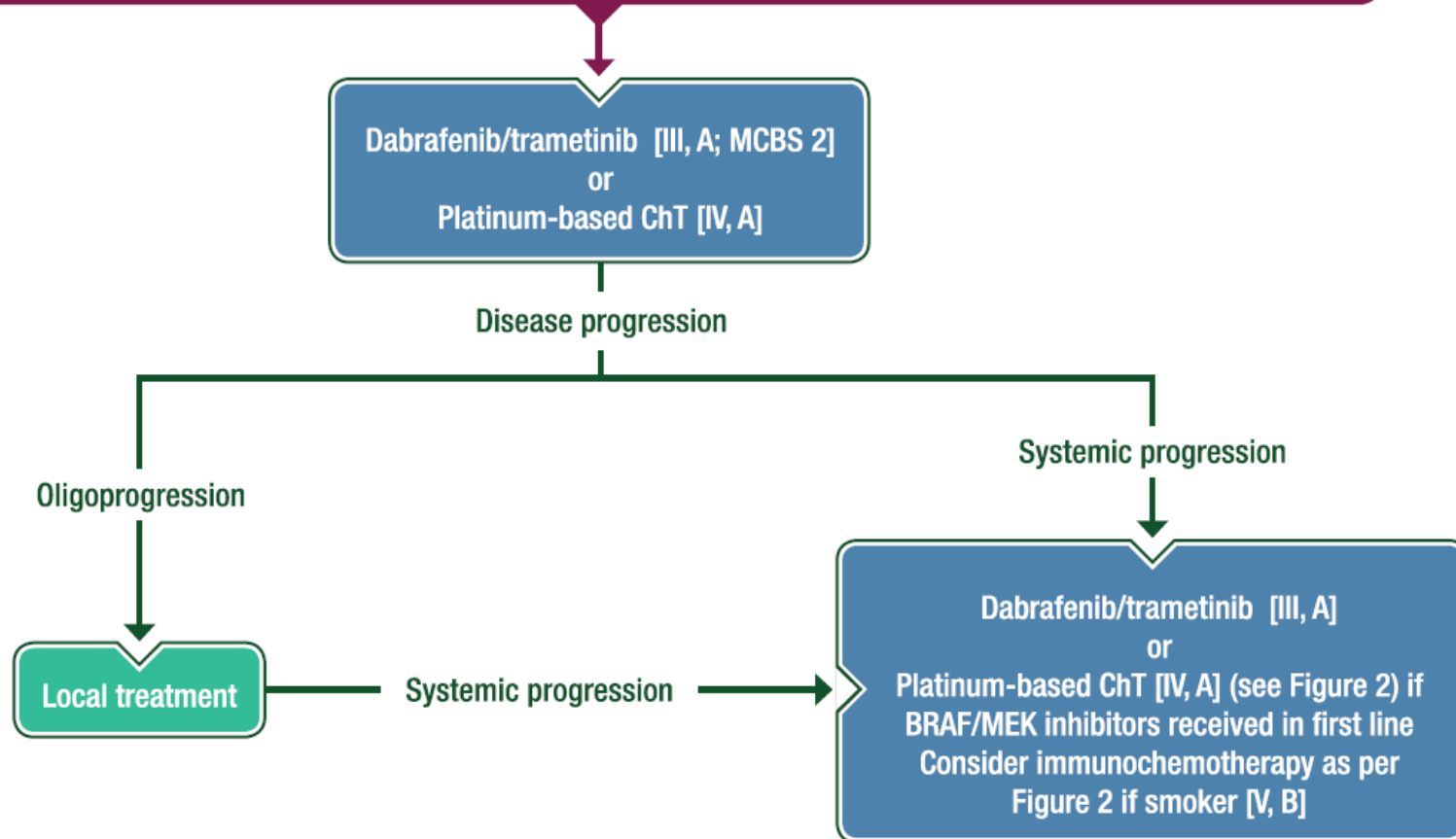
PFS / OS V600E vs. Non-V600E: 6.1 mo. vs. 2.6 mo. ( $p=0.67$ ) / NR vs. 33.9 ( $p=0.47$ )



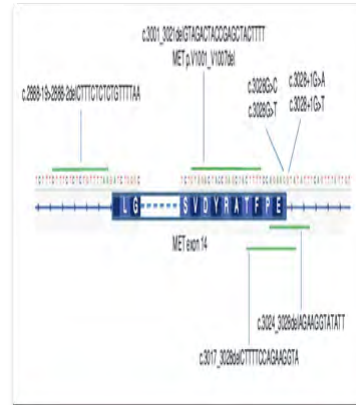
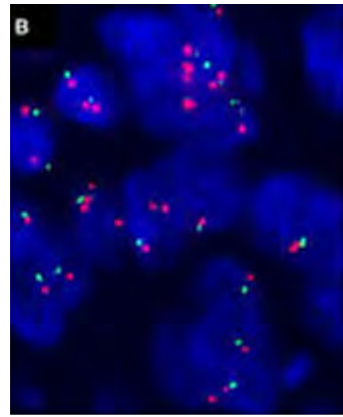
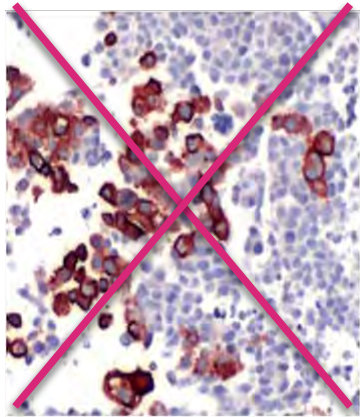
# Immunotarget- Low benefit of immunotherapy in case of molecular alteration...need for specific studies

Driver	n	RR	PFS	OS	Impact (+/X) on PFS of				Comments
					PDL1	Smoking	Nb line	Subtype	
Total		19%	2.8	13.3					Outcome consistent with registration trials for ICI
KRAS	271	26%	3.2	13.5	+	X	X	X	Clear benefit across all subgroups
EGFR	125	12%	2.1	10	+	X	X	X	Could be considered in PDL1 + after TKIs exhaustion
<b>BRAF</b>	<b>43</b>	<b>24%</b>	<b>3.1</b>	<b>13.6</b>	X	+	X	NA	Could be considered in smokers
MET	36	16%	3.4	18.4	NA	X	NA	X	Could be considered after conventionnal treatment
HER2	29	7%	2.5	20.3	NA	+	X	NA	
ALK	23	0	2.5	17	X	X	X	NA	Poor outcome. New biomarker needed.
RET	16	6%	2.1	21.3					
ROS1	7	17%	-	-					

Stage IV lung carcinoma with *BRAF V600* mutation



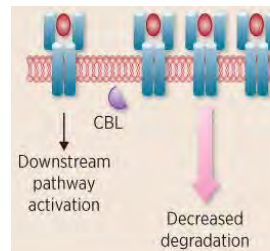
# MET aberrations in NSCLC



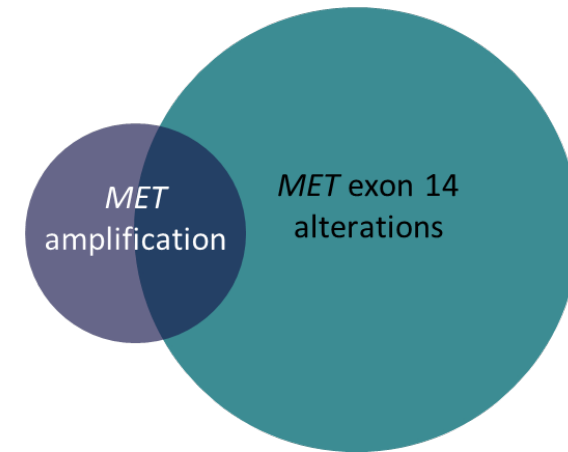
**MET Overexpression**  
25-75%

**MET Amplification**  
3-7%

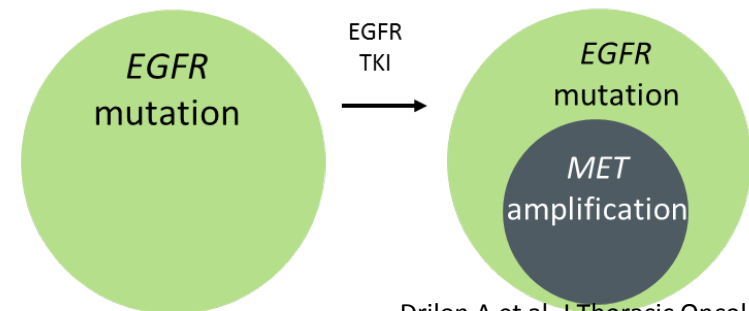
**MET exon14 Skipping**  
3%



## MET as a primary driver



## MET as a secondary/co-driver



# Type 1 MET Inhibitors

## Good Drug

- High ORR
- Potent
- Selective
- Tolerable
- CNS Activity

## Criz Tep

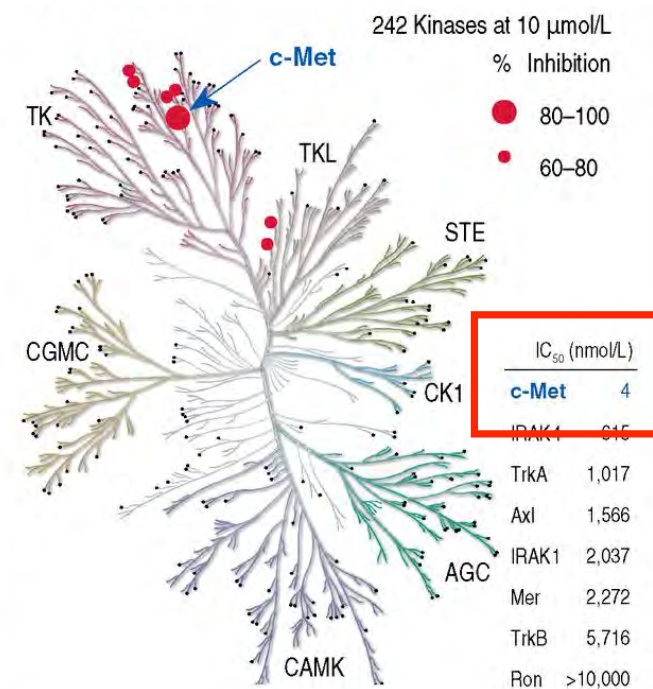
**Modest**



## Crizotinib

Kinase	IC <sub>50</sub> (nM) mean*	Selectivity ratio
c-MET	8	—
ALK	20	2X
RON	298	34X
	189	22X
Axl	294	34X
	322	37X
Tie-2	448	52X
Trk A	580	67X
Trk B	399	46X
Abl	1,159	166X

## Tepotinib

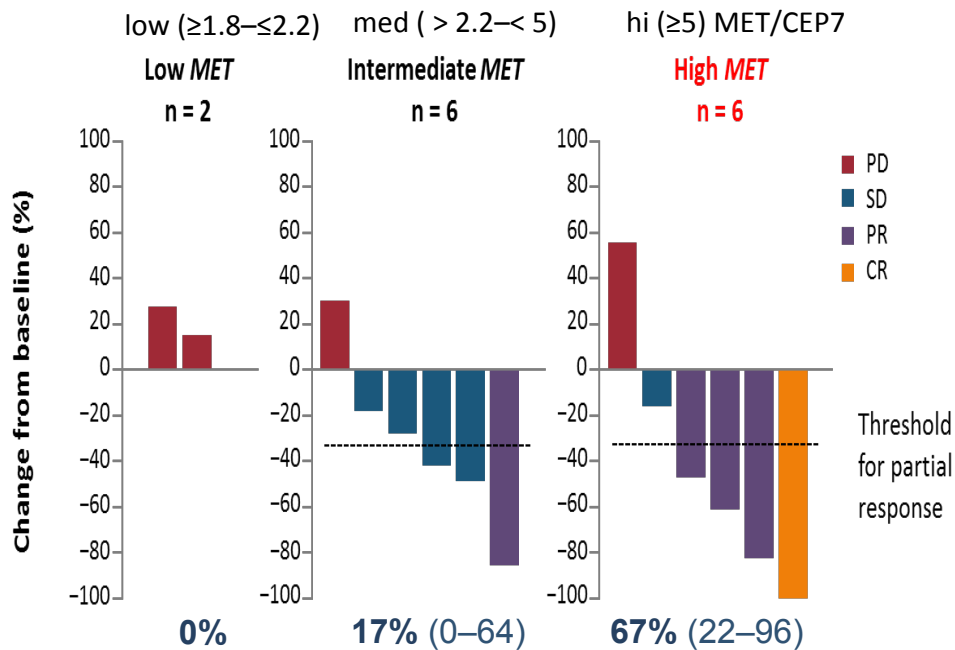


Capmatinib

Savolitinib

# Tumour shrinkage seen with crizotinib or capmatinib treatment in intermediate and high MET amplified

## Crizotinib

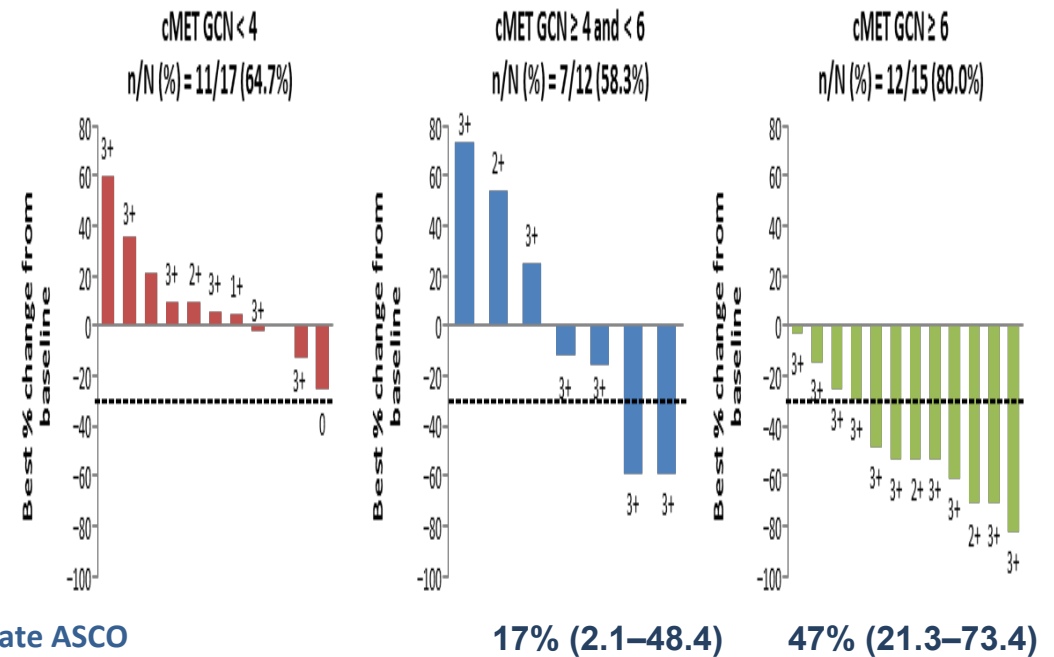


ORR **33.3% (0.8-90.6)** **14.3% (1.8-42.8)** **40.0% (19.1-63.9)**  
 PFS **1.8 (0.8-14.0)** **6.7 (3.4-7.4)** **6.7 (3.4-7.4)**  
 n = 3 n = 14 n = 20

UpDate ASCO  
D. Ross Camidge  
(Abst 9062)

Camidge DR, et al. ASCO 2014. J Clin Oncol. 2014;32:5s (suppl; abstract 8001).

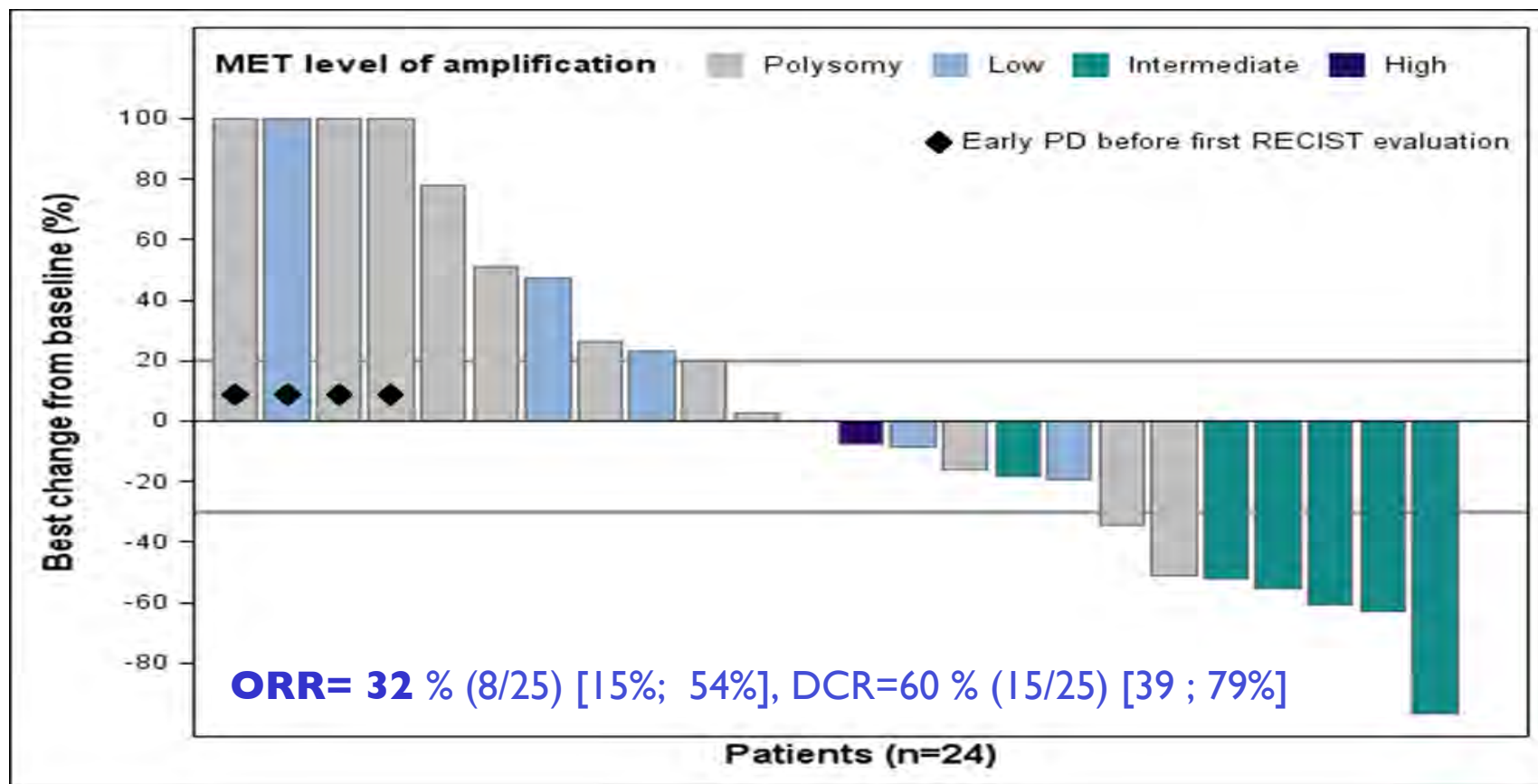
## Capmatinib



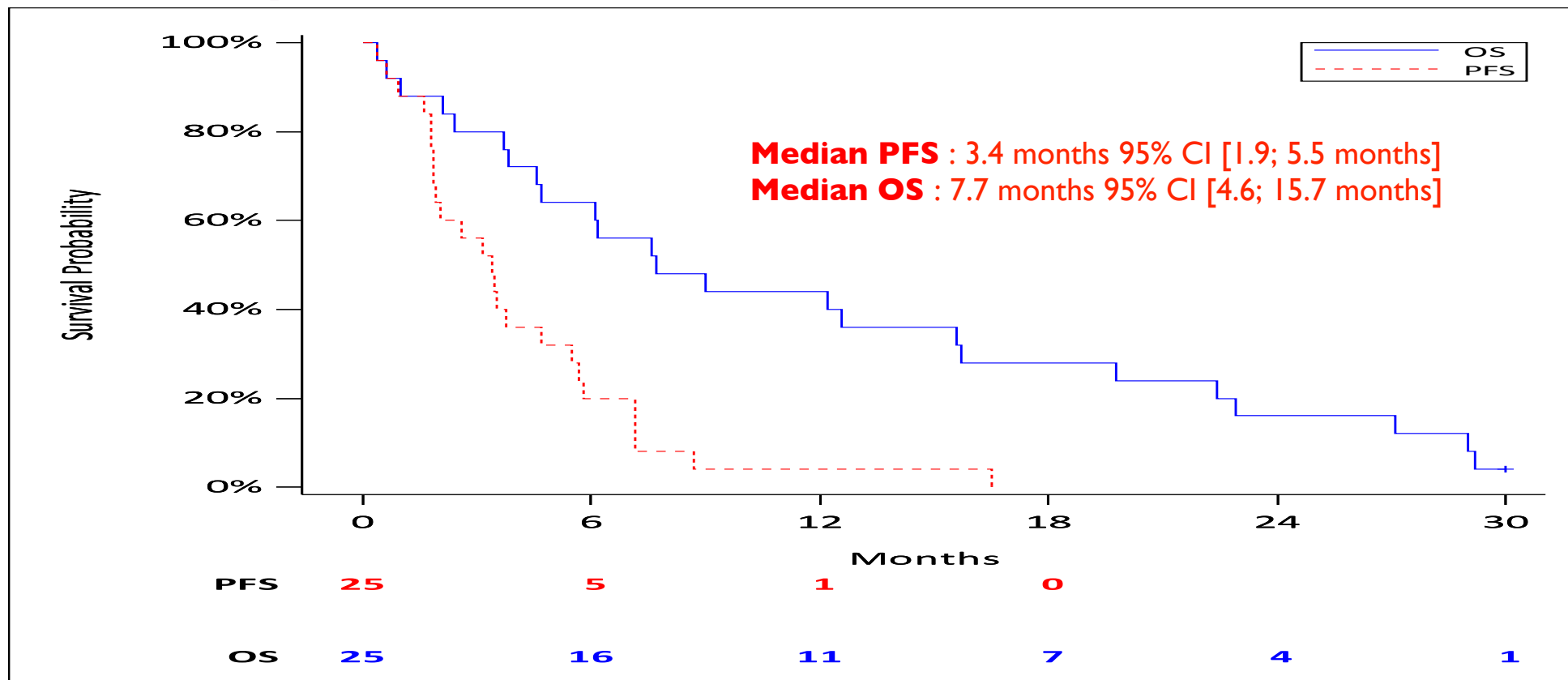
**17% (2.1-48.4)** **47% (21.3-73.4)**

Schuler M, et al. ASCO 2016. J Clin Oncol. 2016;34 Suppl:abstract 9067.

# AcSé trial, Response rate MET amplification



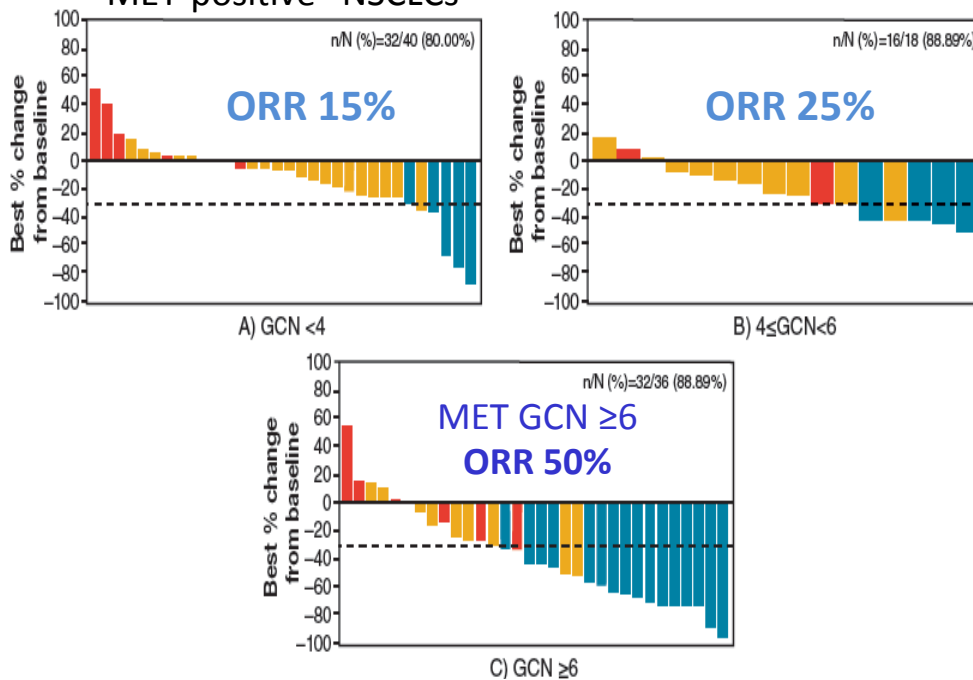
# MET amplification



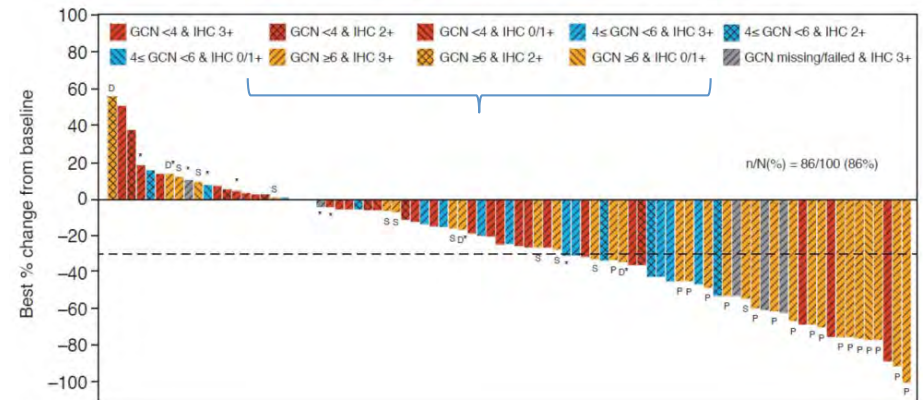
# Response to Combined EGFR- and MET-Directed Targeted Therapy (MET amplified)

## •Capmatinib + Gefitinib

- Phase 2 expansion cohort
- EGFR-mutant lung cancers with acquired resistance and "MET-positive" NSCLCs



## Phase Ib/II Study



ORR: 47% in patients with MET gene copy number ≥6

JOURNAL OF CLINICAL ONCOLOGY

EDITORIAL

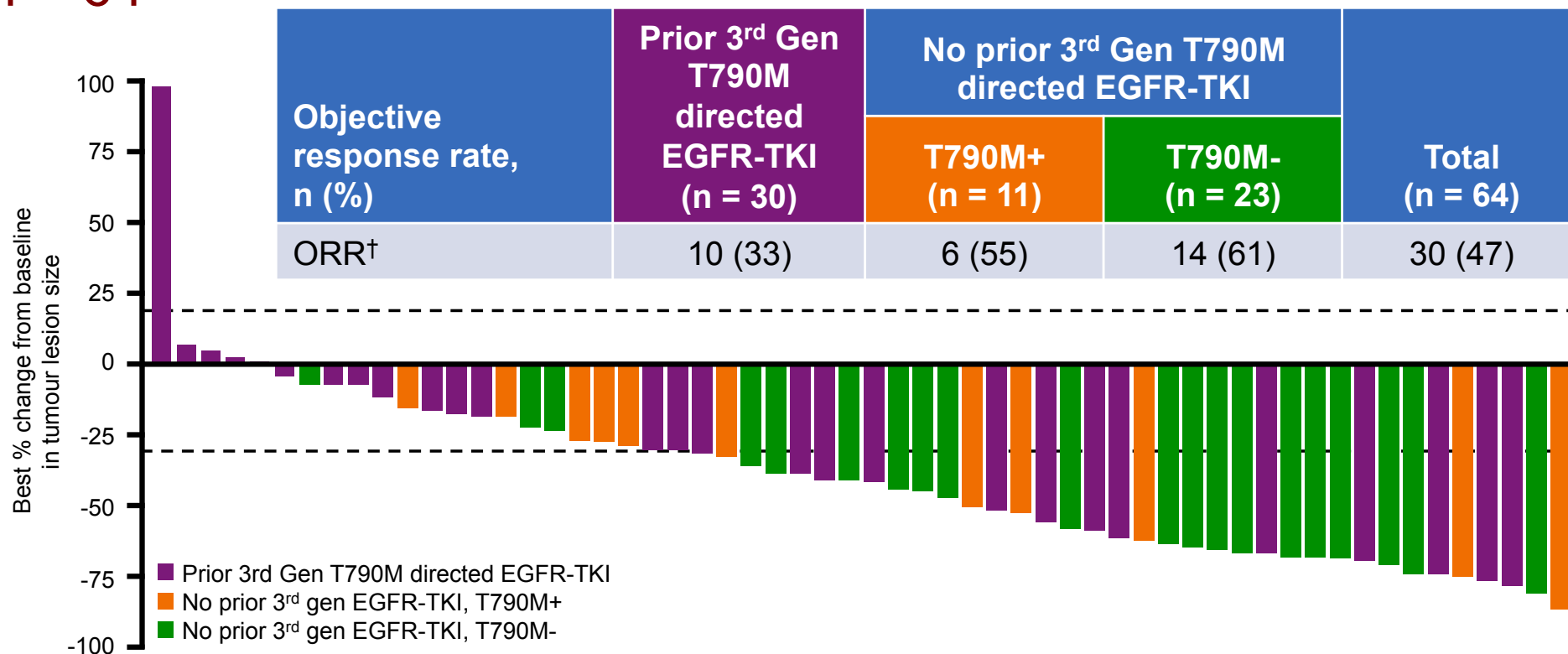
Have We Really MET a New Target?

David Planchard, Gustave Roussy, Villejuif, France



# TATTON (osimertinib+ Savolitinib)

Preliminary anti-tumour activity in all MET-positive patients\*, n = 64

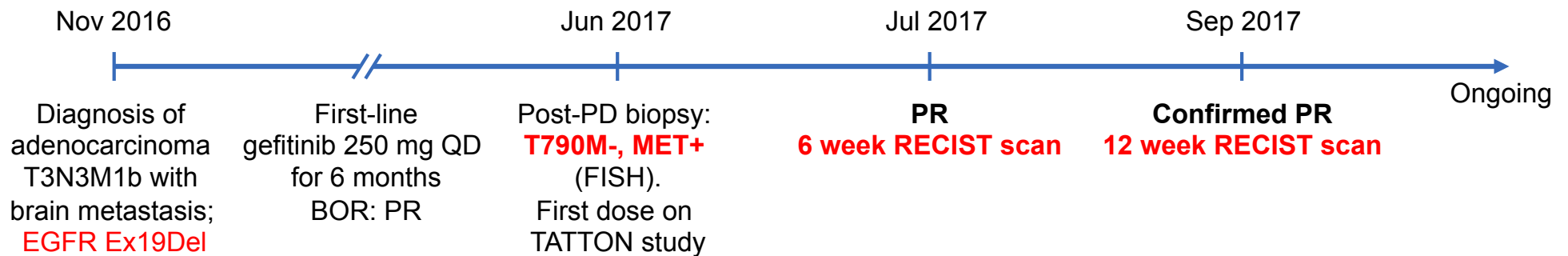
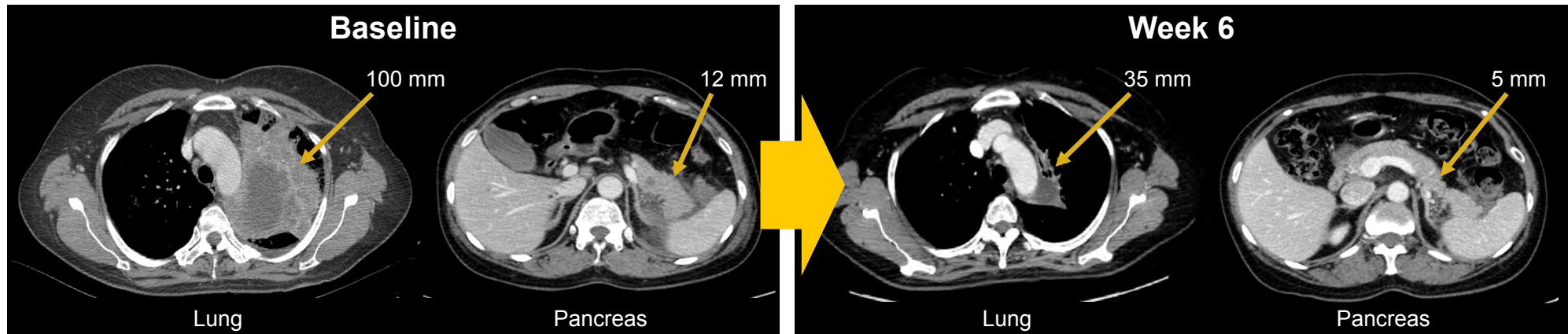


Waterfall plot based on evaluable patients (n = 64): all patients dosed and with on-treatment assessment or discontinuation prior to first tumour assessment  
Data cut-off 31 Aug 2017

\*17 patients did not have central FISH confirmation of MET-positive status (n = 6 MET-negative; n = 11 unknown by central lab); †Confirmed by a later scan performed at least 4 weeks after initial response observed

TATTON Part B  
NCT02143466

# 57 year old female never smoker with NSCLC adenocarcinoma histology, ECOG PS 1

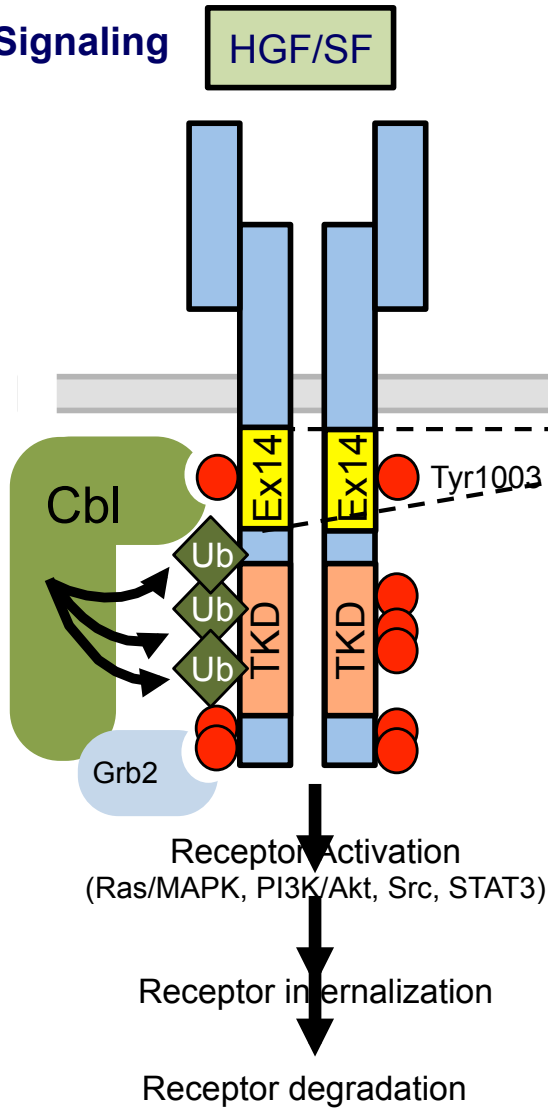


Ex19Del, exon 19 deletion

TATTON Part B  
NCT02143466

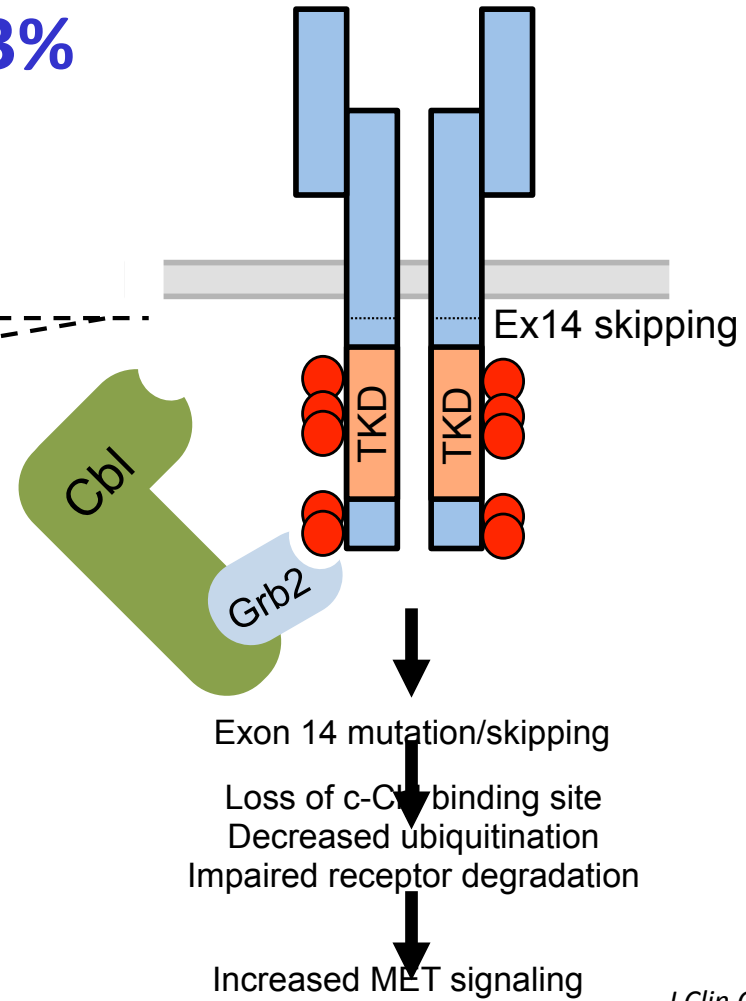
Myung-Ju Ahn et al, IASLC 2017

## Normal MET Signaling

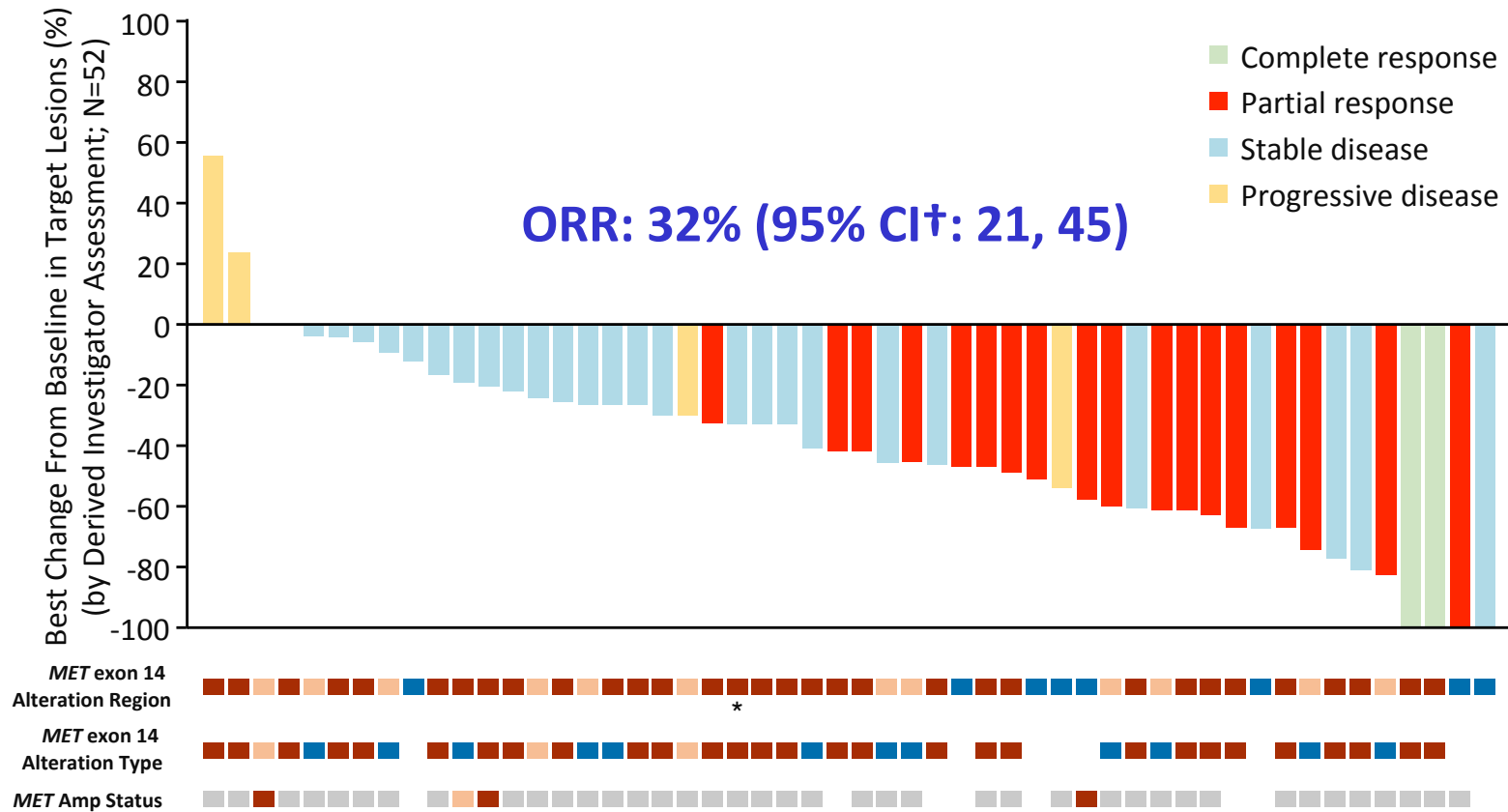


## Exon 14 Mutated/Skipped

3%



# Updated Antitumor Activity and Safety of Crizotinib in Patients With *MET* Exon 14-Altered Advanced NSCLC

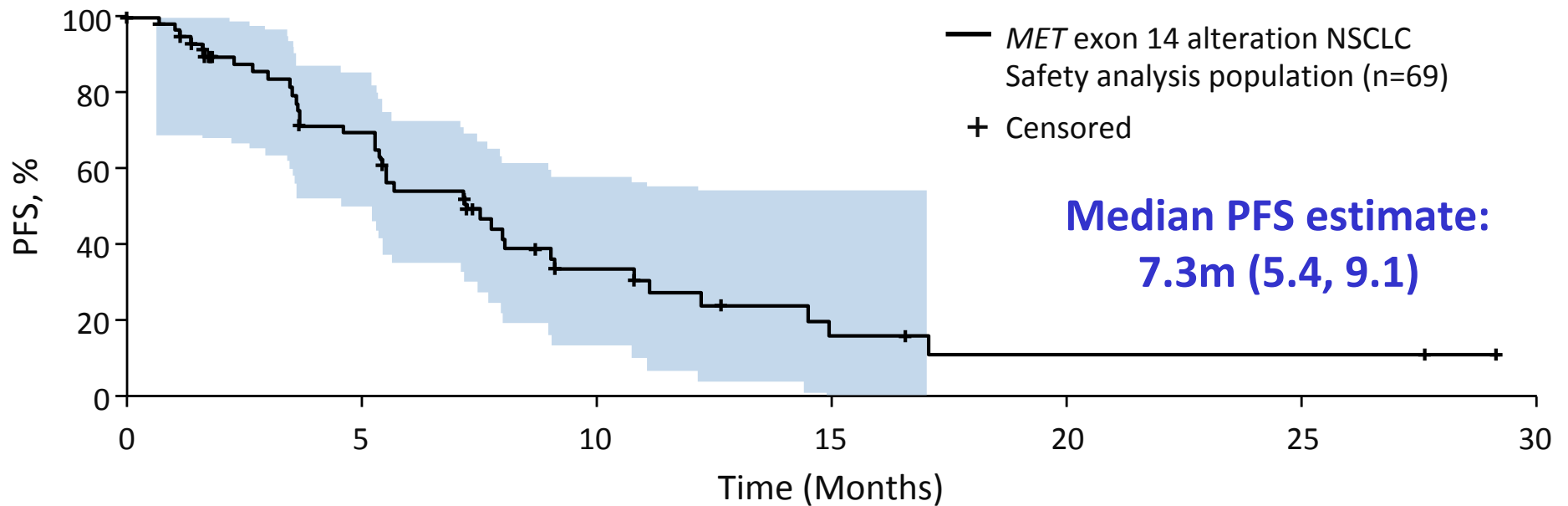


**Biomarker Data Key<sup>§</sup>**

	<i>MET</i> exon 14 alteration region	<i>MET</i> exon 14 alteration type	<i>MET</i> amp status
Red	Splice donor	Base substitution	Detected
Orange	Splice acceptor <sup>†</sup>	Large indel (>35 bp)	UIF
Blue	Canonical <sup>‡</sup>	Indel	–
Grey	Not detected	–	Not detected

\*Alterations in both splice donor and acceptor regions. †Includes alterations in the Splice Acceptor Region, Polypyrimidine Tract, and Branching Point. ‡Includes *MET* exon 14 alterations that are not associated with DNA coding region information. §White space in biomarker data rows indicates no available sample for testing, not analyzable or no results reported. bp, base pairs; UIF, uninformative.

## Progression-Free Survival (PFS) by Derived Investigator Assessment



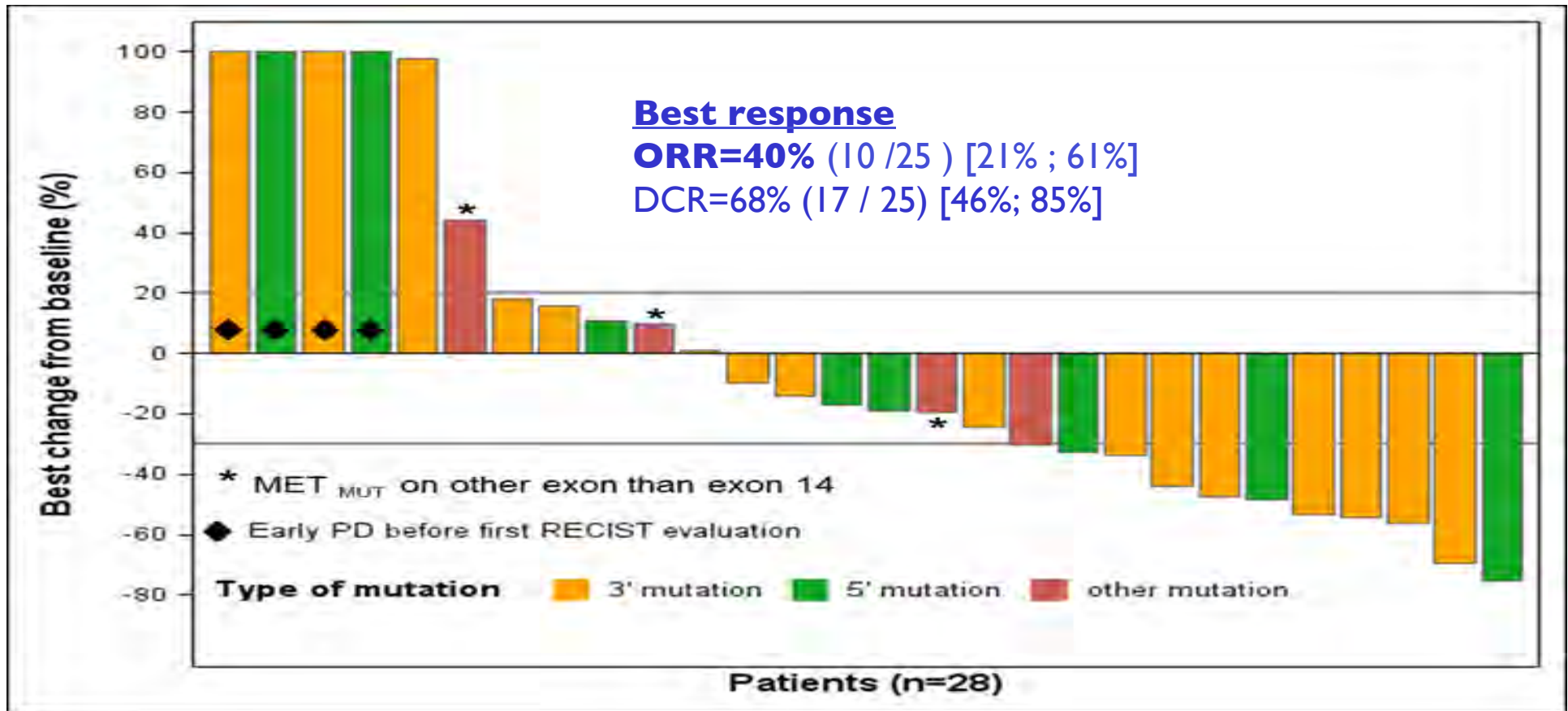
No. at risk	69	33	11	4	2	2	0
-------------	----	----	----	---	---	---	---

- OS data were not mature at time of data cutoff: 34.8% patients had died; 40.6% still in follow-up
- Median Overall Survival (OS) estimate, months (95% CI): 20.5 (14.3, 21.8)

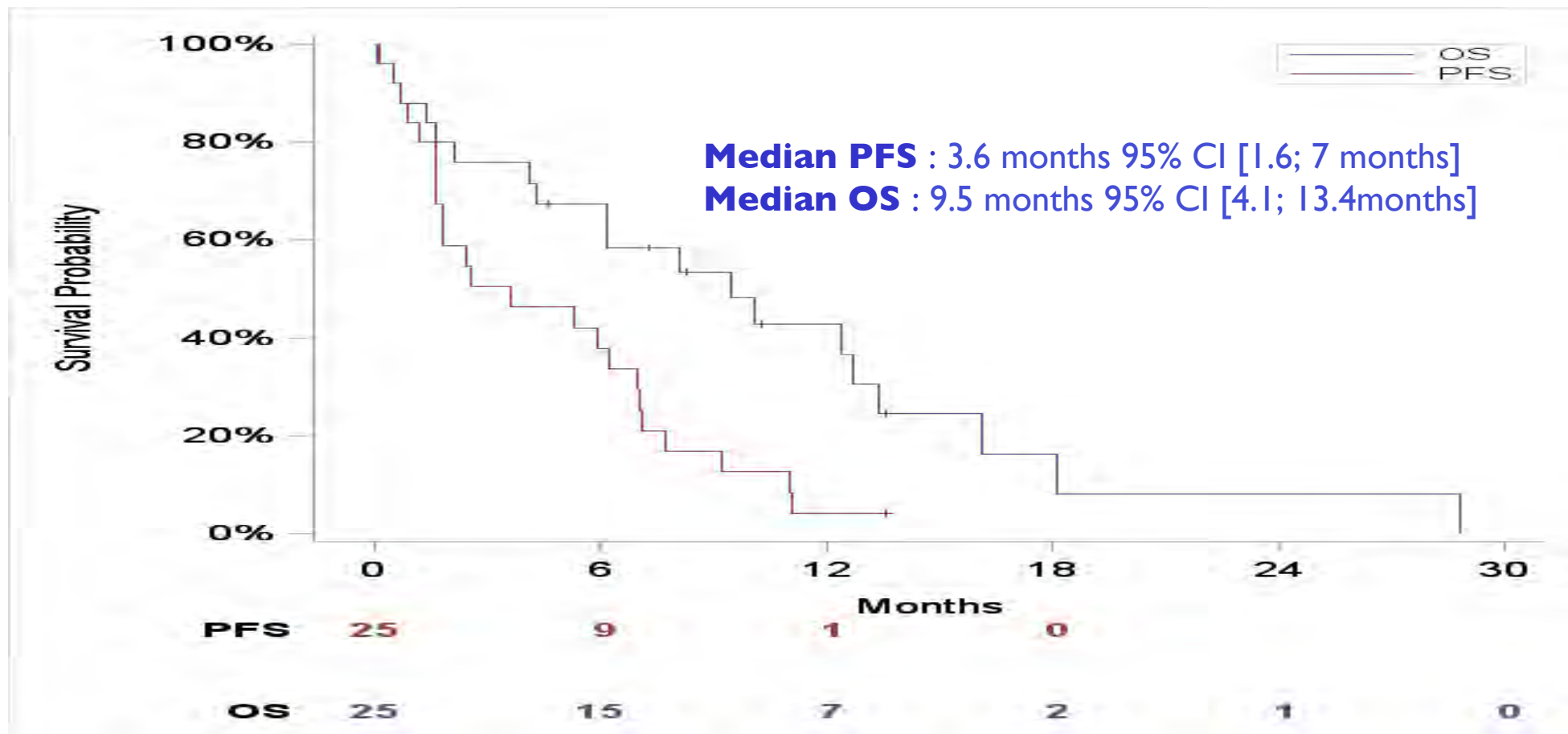
Shaded area in PFS Kaplan-Meier plot above represents 95% Hall-Wellner band. 95% CI estimates for PFS and OS based on Brookmeyer and Crowley method.

Alexander Drilon et al, WCLC 2018

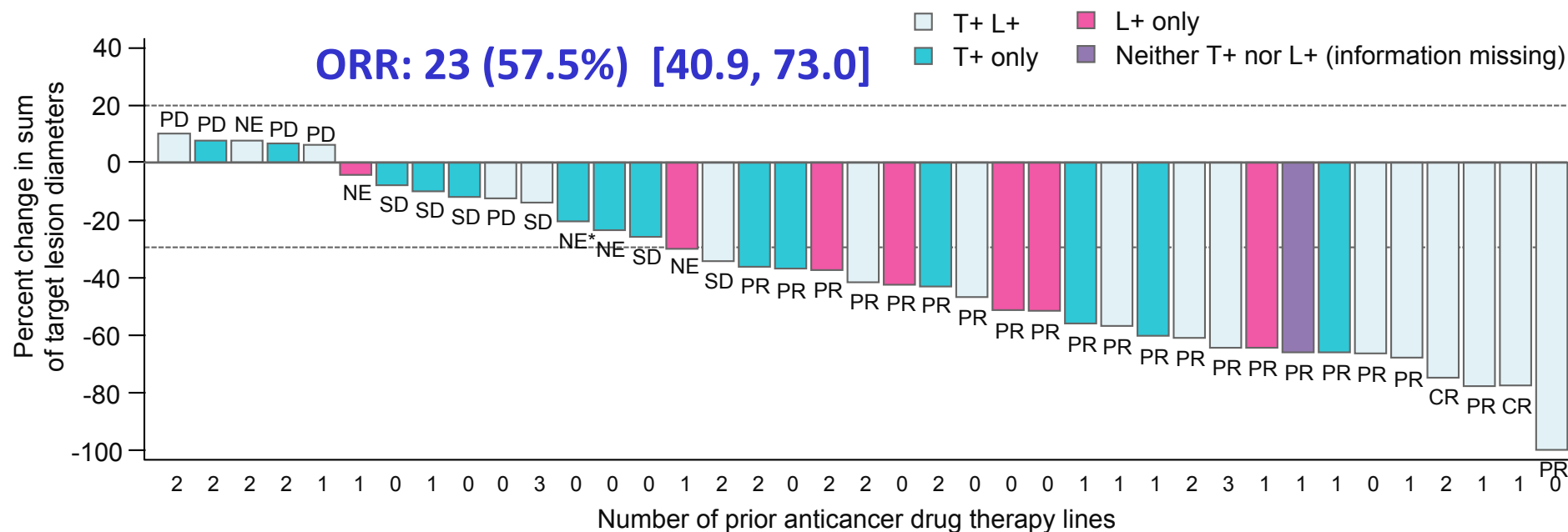
# AcSé trial, Response rate *MET* exon 14 mutation



## Response rate *MET* exon 14 mutation



# VISION: A Phase II, Single-arm Trial to Investigate Tepotinib in Advanced NSCLC with METexon14-Skipping Alterations

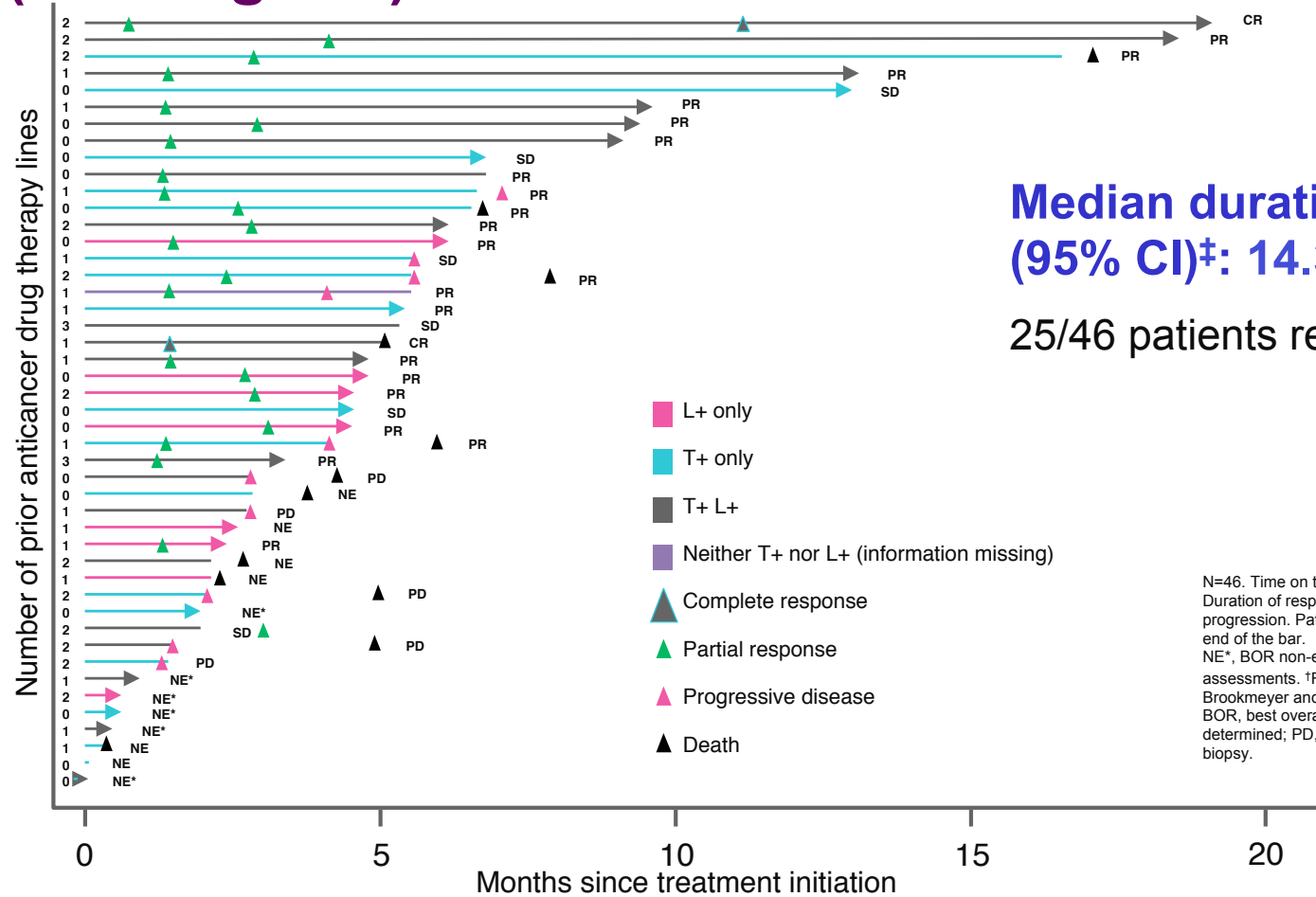


n=39. Seven patients were excluded due to baseline/on-treatment measurement not being available.  
 BOR displayed at the end of the bar. NE\*, BOR non-evaluable where ongoing patient has not had 2 post-baseline tumor assessments.  
 BOR, best overall response; CR, complete response; L, liquid biopsy; NE, non-evaluable; PD, progressive disease; PR, partial response;  
 SD, stable disease; T, tumor biopsy.

Dr Enriqueta Felip, Vall d'Hebron University Hospital, Spain



# Time on Treatment and Duration of Response (Investigator)



**Median duration of response<sup>†</sup>  
(95% CI)<sup>‡</sup>: 14.3 (3.7, nd) months**

25/46 patients remain on-treatment

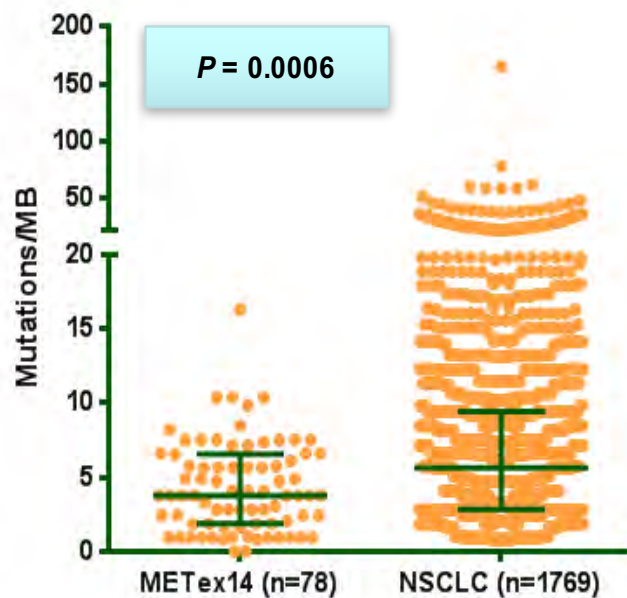
- L+ only
- T+ only
- T+ L+
- Neither T+ nor L+ (information missing)
- ▲ Complete response
- ▲ Partial response
- ▲ Progressive disease
- ▲ Death

N=46. Time on treatment is the time from treatment initiation until treatment termination. Duration of response is measured from time of initial response until documented tumor progression. Patients denoted with an arrow remain on-treatment. BOR displayed at the end of the bar.  
NE\*, BOR non-evaluable where ongoing patient has not had two post-baseline tumor assessments. <sup>†</sup>From Kaplan-Meier survival analysis. <sup>‡</sup>95% CI for the interval using the Brookmeyer and Crowley method.  
BOR, best overall response; CR, complete response; L, liquid biopsy; nd, not determined; PD, progressive disease; PR, partial response; SD, stable disease; T, tumor biopsy.

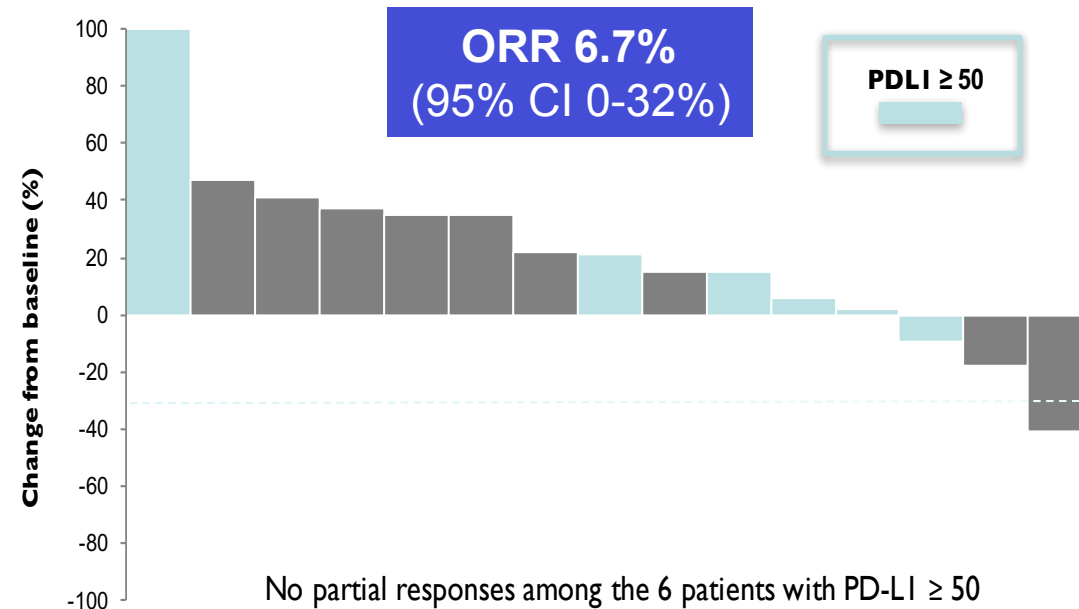
# Poor Response to Immunotherapy in MET exon 14-altered NSCLCs

*MET* exon 14-altered cancers can express high levels of PD-L1

PD-L1, Cell Signaling, Clone E1L3N assay, n=54			
	0 %	1 – 49 %	≥50 %
n, (%)	19 (35)	10 (19)	25 (46)



TMB is lower in patients with *MET* exon 14 altered NSCLCs compared to other NSCLCs



PD-LI Status    90    NA    0    0    NA    NA    NA    100    0    90    80    90    100    NA    NA

## Low benefit of immunotherapy in case of molecular alteration...need for specific studies

Driver	n	RR	PFS	OS	Impact (+/X) on PFS of				Comments
					PDL1	Smoking	Nb line	Subtype	
Total		19%	2.8	13.3					Outcome consistent with registration trials for ICI
KRAS	271	26%	3.2	13.5	+	X	X	X	Clear benefit across all subgroups
EGFR	125	12%	2.1	10	+	X	X	X	Could be considered in PDL1 + after TKIs exhaustion
BRAF	43	24%	3.1	13.6	X	+	X	NA	Could be considered in smokers
<b>MET</b>	<b>36</b>	<b>16%</b>	<b>3.4</b>	<b>18.4</b>	NA	X	NA	X	Could be considered after conventional treatment
HER2	29	7%	2.5	20.3	NA	+	X	NA	
ALK	23	0	2.5	17					
RET	16	6%	2.1	21.3	X	X	X	NA	Poor outcome. New biomarker needed.
ROS1	7	17%	-	-					

# Resistance to MET-Directed Targeted Therapy

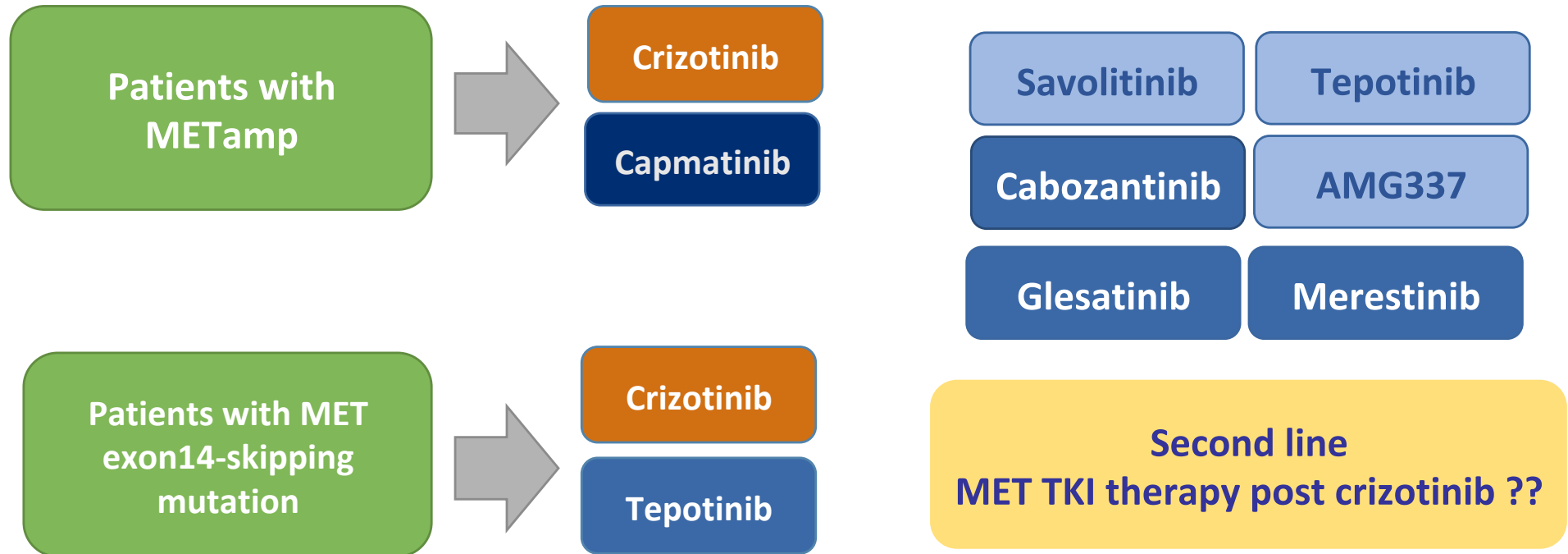
Drug Administered	MET alteration	Putative resistance mechanism	Notes
<b>Crizotinib</b> 8 mo of disease control	MET D1010H	<b>MET D1228N</b> (acquired second site mutation on tumor rebiopsy)	high total MET and phospho-MET IHC+ on post-PD biopsy
<b>Crizotinib</b> 13 mo of disease control	MET D1010H (MET Y1230C)	<b>MET Y1230C</b> (detected in ctDNA on PD)	
<b>Crizotinib</b> 8 mo of disease control	MET c.3028delG	<b>MET Y1230H</b> (acquired in tumor, MET amp + MET D1228N, Y1230H, Y1230S, and G1163R in plasma)	thereafter responded to Glesatinib
<b>Savolitinib + Osimertinib</b> 9 mo of disease control	MET amplification (+EGFR ex19 del)	<b>MET D1228V</b> (acquired second site mutation on tumor rebiopsy)	thereafter responded to Cabozantinib + Erlotinib

## Mechanisms of acquired resistance to MET TKIs in MET exon 14 mutant NSCLC

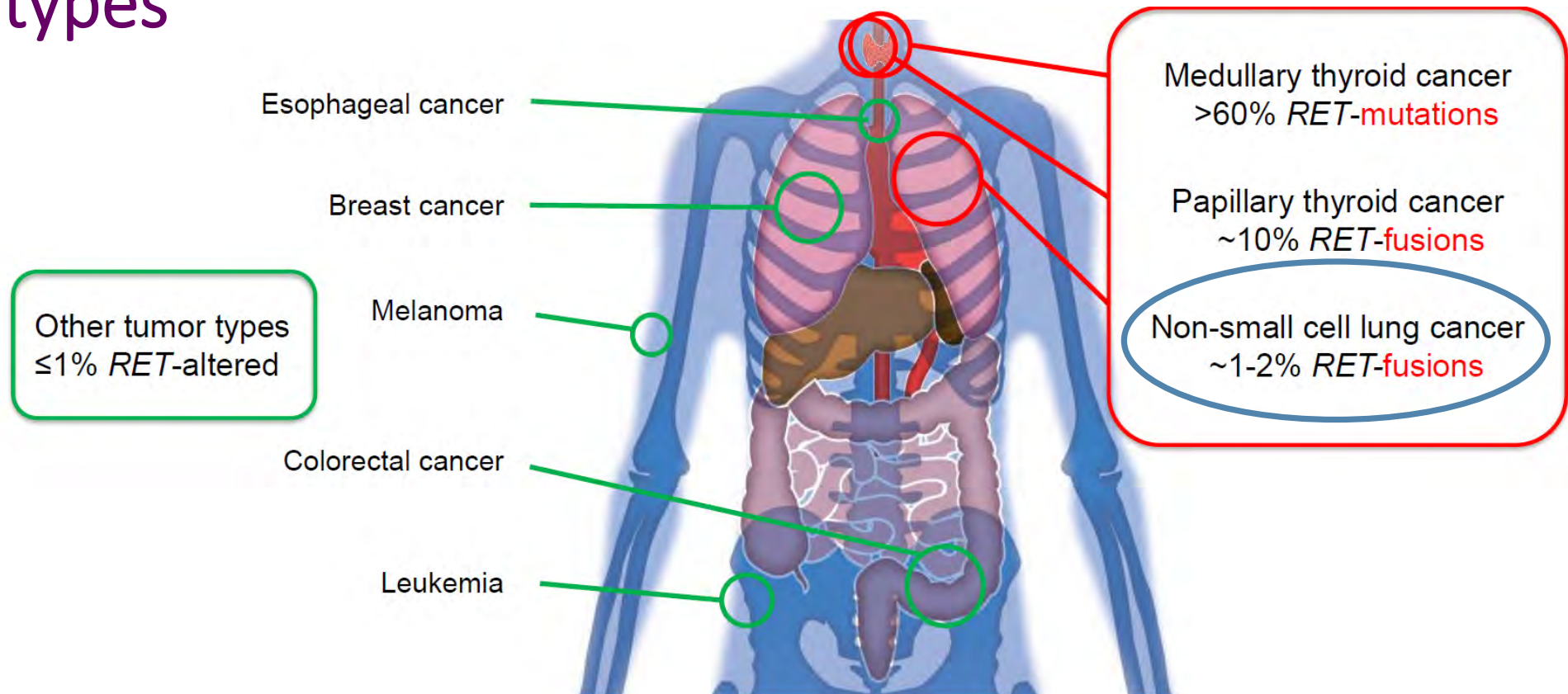
Presented Sunday, June 3, 2018. Mark M. Awad (Abst 9069)

- Secondary mutations in MET included H1094Y, G1163R, L1195F, L1195V, D1228N, Y1230H, and Y1230S.
- bypass track activation : amplification of wild-type KRAS, BRAF, and/or EGFR.
- acquired amplification of the mutated METex14 allele

# In summary for MET NSCLC

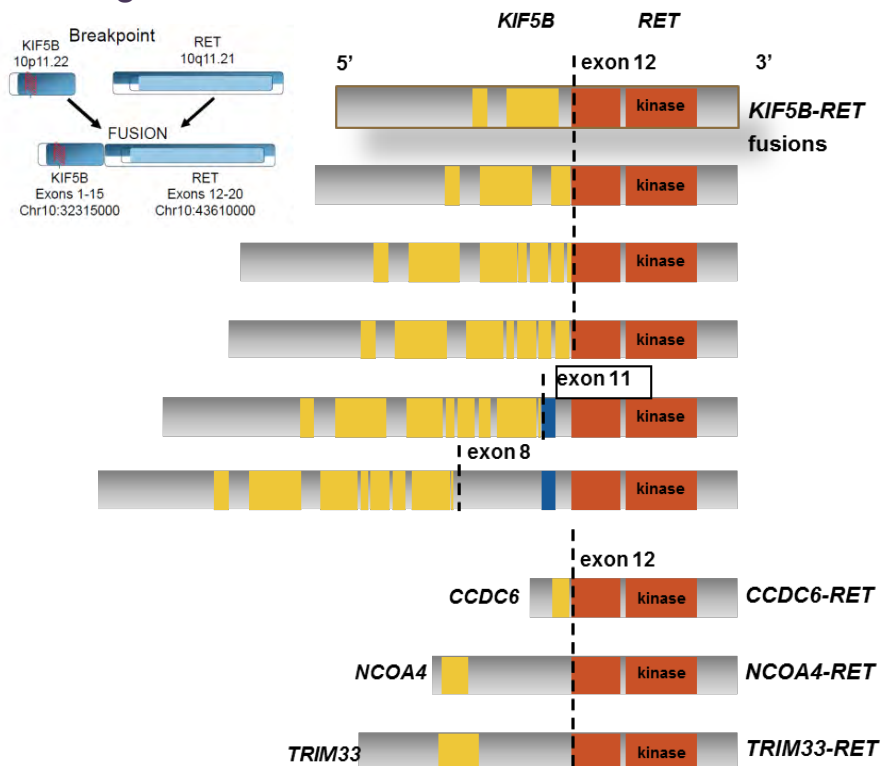


# RET is a rare driver of multiple, diverse tumor types



# RET rearrangements

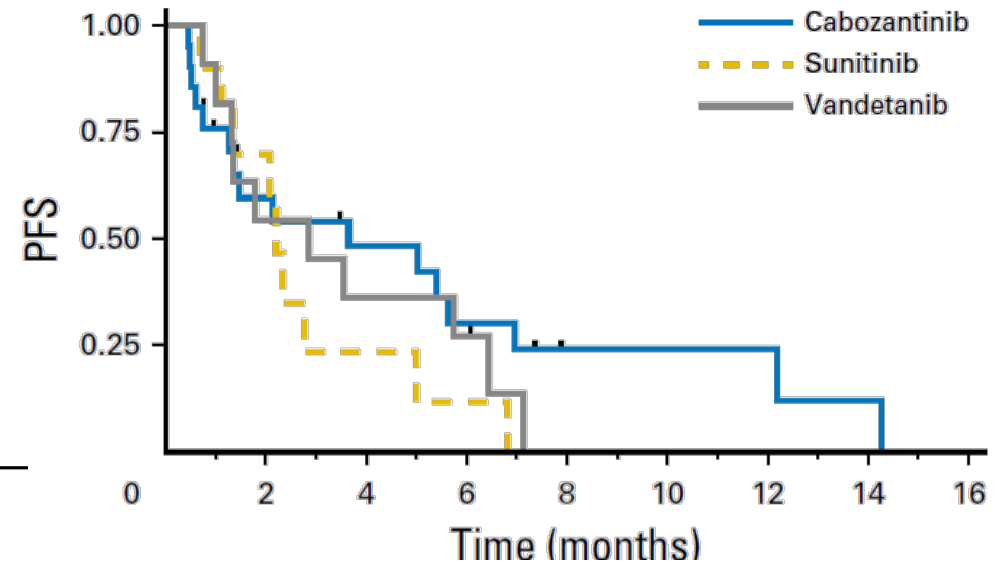
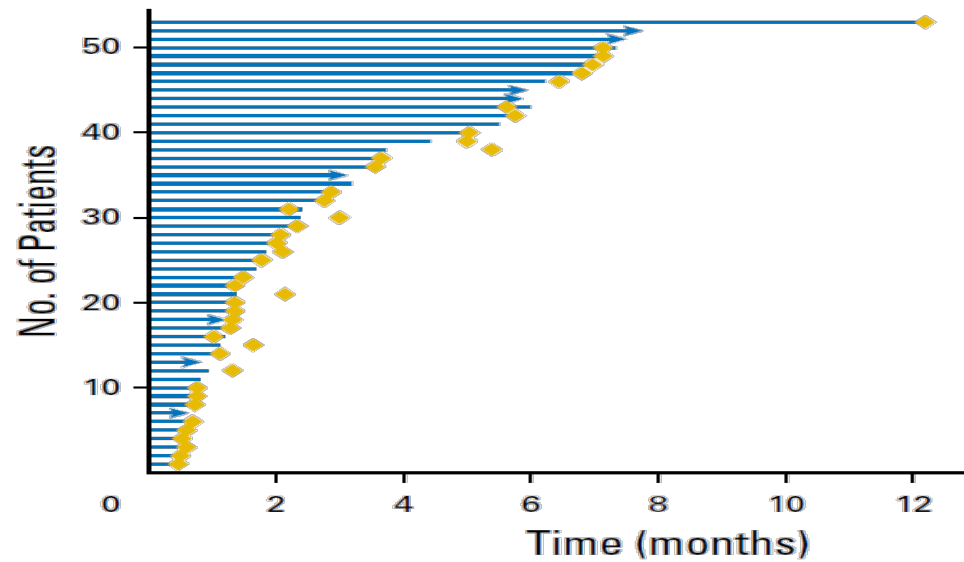
- 1–2% of unselected NSCLCs
- Clinical features: young, never or former light cigarette smokers



## Multi-tyrosine kinase inhibitors

Compound	IC <sub>50</sub> (nM) In vitro kinase	IC <sub>50</sub> (nM) Cellular kinase	IC <sub>50</sub> (nM) In vitro kinase RET V804M	Other targets
Regorafenib	1.5	~10	NR	VEGFR1-3, BRAF, c-kit, PDGF-b
Levatinib	1.5	48	NR	VEGFR1-3, FGFR1-3, c-kit, PDGFR
Alectinib	4.8	?	53 V804L (32)	ALK (1.9 nM)
Cabozantinib	5.2	27-85	4094	VEGFR2, MET
Ponatinib	7	0.7-11	12	Bcr-abl, FGFR1-4
Sunitinib	30	40-164	55	VEGFR, PDGFR, c-kit, Flt-3
Sorafenib	47	~20-50	12	RAF, VEGFR2-3, PDGFR, c-kit, Flt-3
Vandetanib	100	NR	> 10,000	VEGFR, EGFR

# Global RET Registry

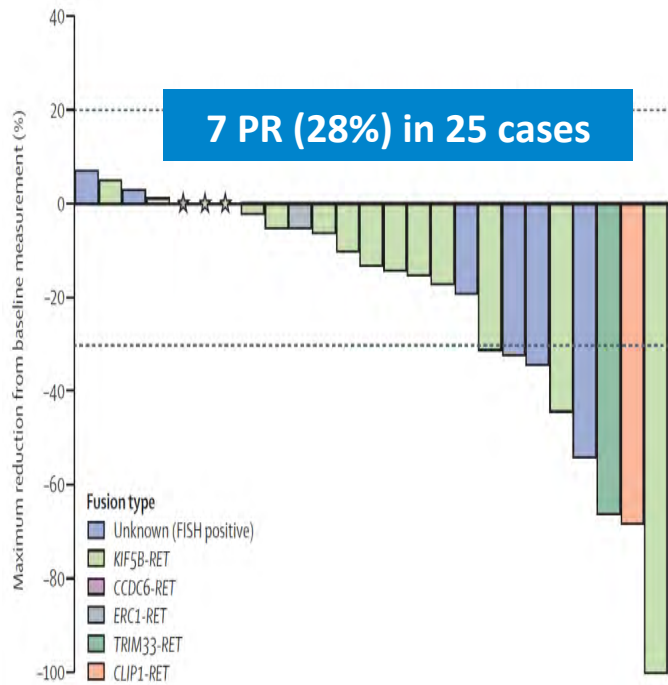


RET inhibitor	Best response (% ; 95 % CI)	Median DoT (range)	Median PFS (95% CI)	Median OS (95% CI)
Cabozantinib	37% (16.3 - 61.6)	1.6 months (0.5 -12.2)	3.6 months (1.3 - 7.0)	4.9 months (1.9 - 14.3)
Vandetanib	18% (2.3 - 51.8)	2.9 months (0.8 - 7.1)	2.9 months (1.0 - 6.4)	10.2 months (2.4 - NR)
Sunitinib	22% (2.8 - 60.0)	2.2 months (0.7 - 6.6)	2.2 months (0.7 - 5.0)	6.8 months (1.1 - NR)



# Need of more potent drugs

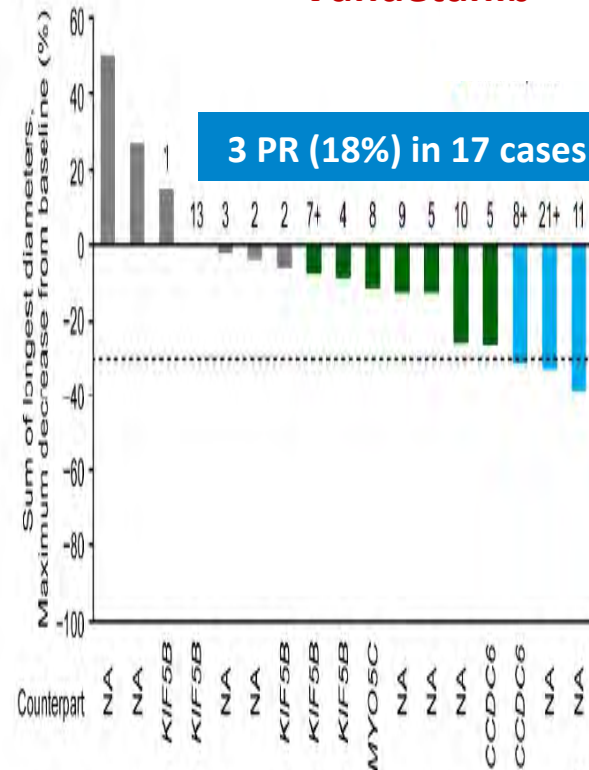
## Cabozantinib



mPFS: 5.5 months (95% CI 3.8–8.4)

Drilon et al, Lancet Oncol, 2016

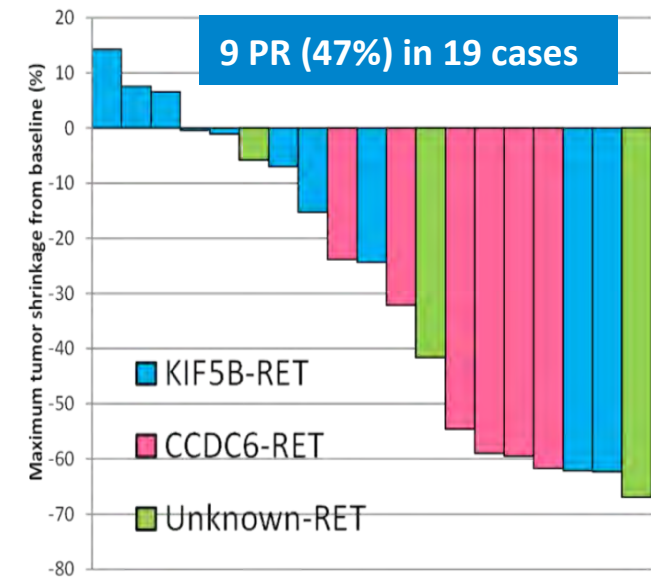
## Vandetanib



mPFS: 5.5 months

Lee et al, Ann Oncol, 2017

## Vandetanib



mPFS: 4.7 months (95% CI 2.8–8.5)

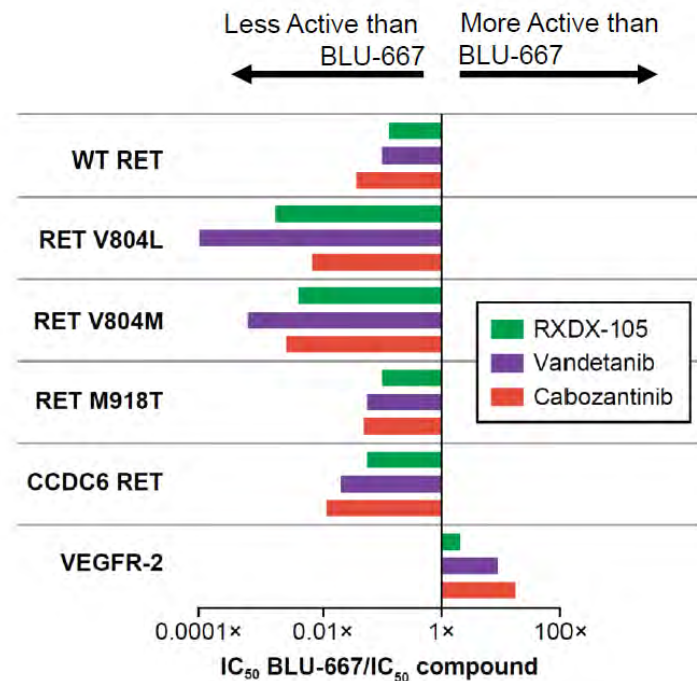
Yoh et al, Lancet Resp Med, 2017

# BLU-667 designed to treat RET-altered cancers

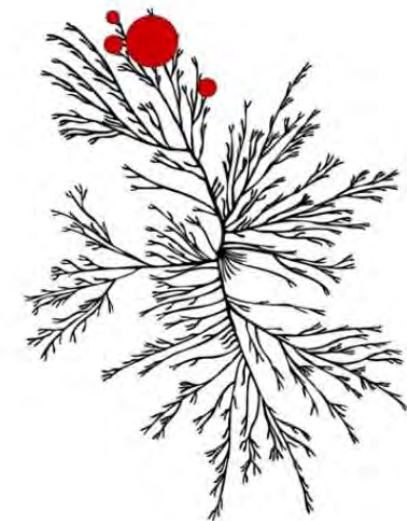
Subnanomolar potency<sup>1</sup>

Variant	Biochemical IC <sub>50</sub> (nM)
RET wildtype	0.4
RET V804L	0.3
RET V804M	0.4
RET M918T	0.4
CCDC6-RET	0.4

More Potent than MKI

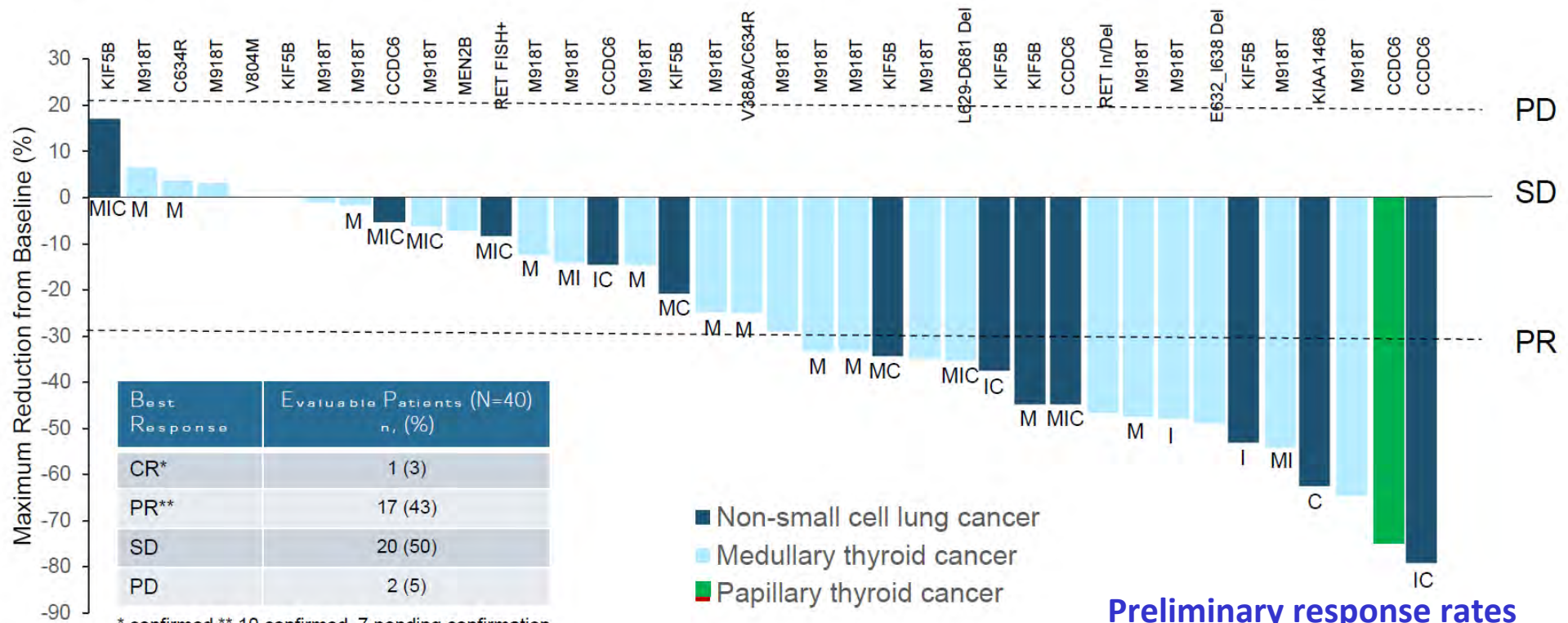


Kinome selectivity for RET



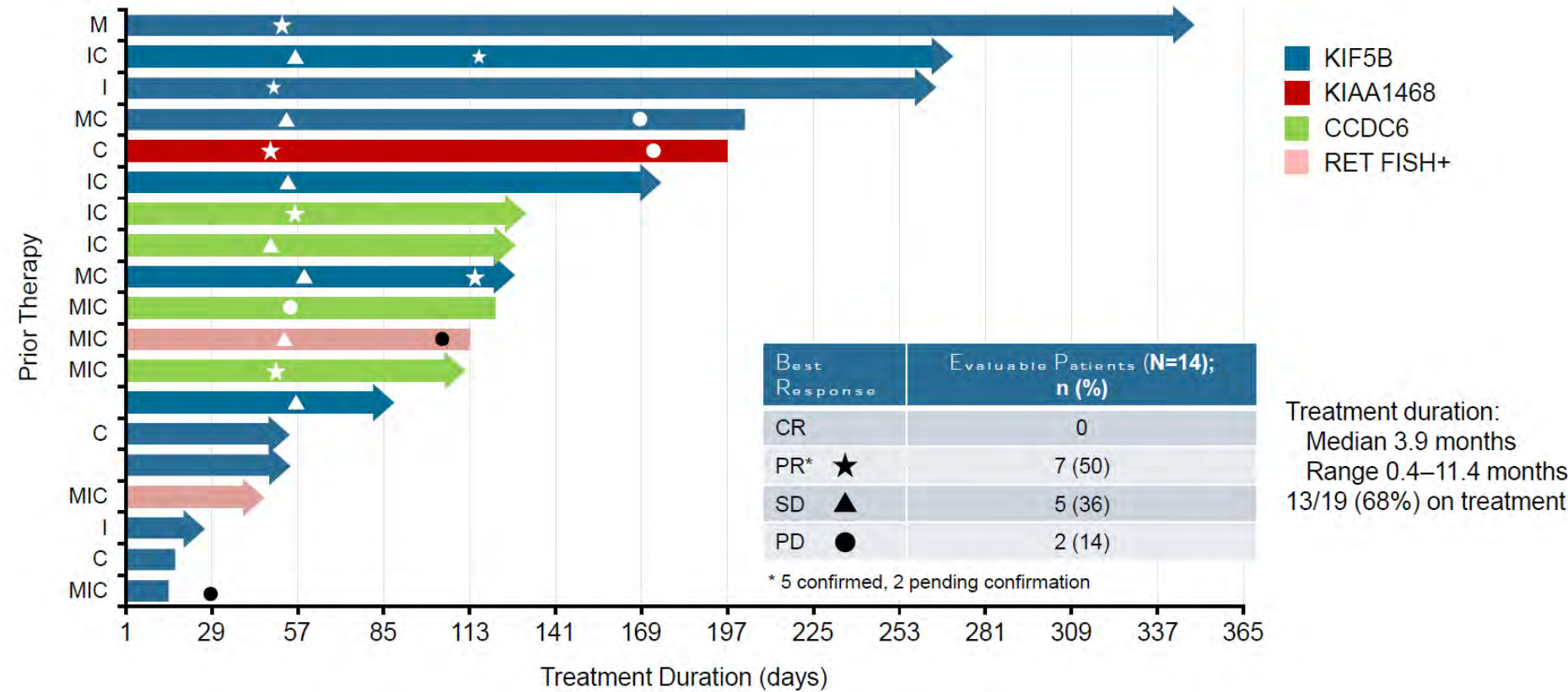
1. Subbiah V et al. *Cancer Discovery* April 15 2018

# Broad anti-tumor activity against RET-altered cancers



**Preliminary response rates**  
 -ORR RET-fusion NSCLC 50%  
 -ORR RET-fusion MTC 40%

# Durable activity



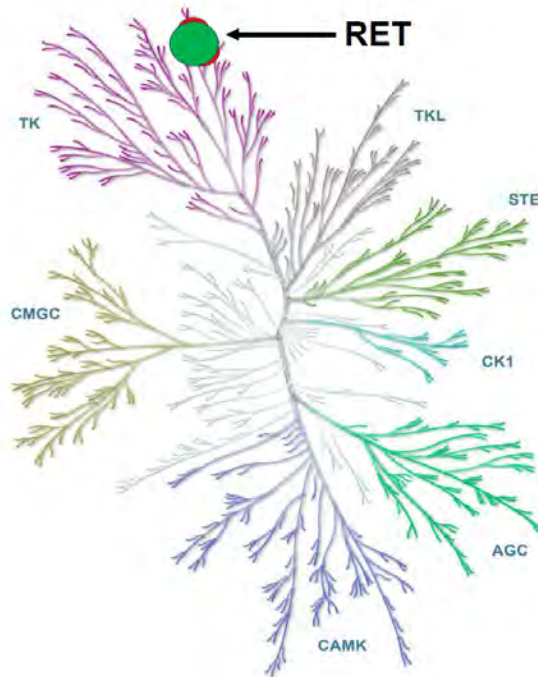
Vivek Subbiah et al, Cancer discovery 2018

Vivek Subbiah et al, AACR 2018

# LOXO-292 is a potent and selective RET inhibitor

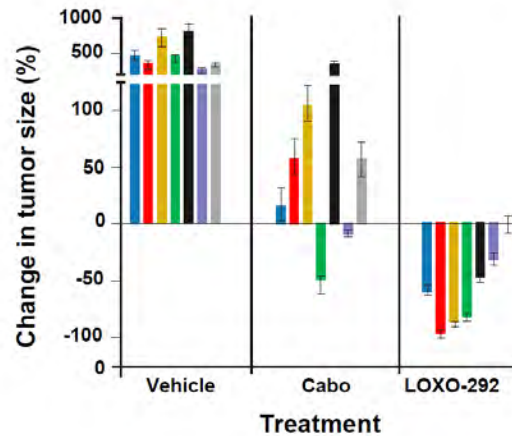
## Kinome selectivity

Highly selective for RET



## Xenograft models

Multiple fusions/mutations/histologies

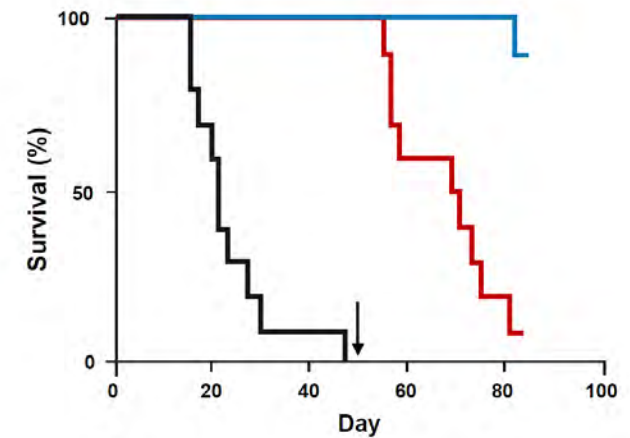


### Tumor models

- KIF5B-RET (PDX-NSCLC)
- CCDC6-RET (PDX-CRCA)
- CCDC6-RET-V804M (PDX-CRCA)
- KIF5B-RET (NIH-3T3)
- KIF5B-RET-V804M (NIH-3T3)
- RET C634W (TT cell line-MTC)
- CCDC6-RET (LC-2/ad cell line-NSCLC)

## Orthotopic brain model

CCDC6-RET orthotopic brain PDX

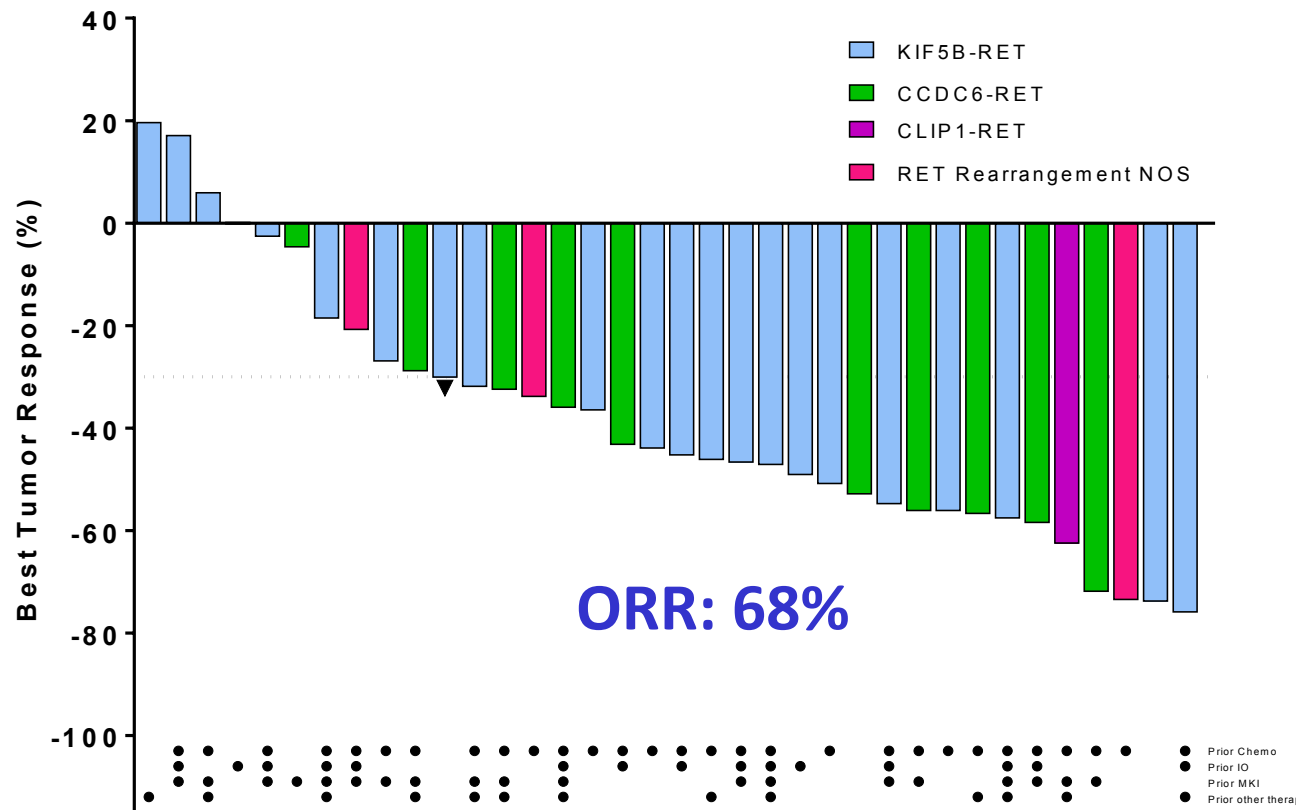


### Treatments

- Vehicle
- LOXO-292 30 mg/kg BID → Day 52 → 3 mg/kg BID
- Ponatinib 20 mg/kg QD → Day 52 → 2 mg/kg QD

Subbiah et al. Ann Oncol 2018; Cabo = cabozantinib; PDX = patient-derived xenograft; NSCLC = non-small cell lung cancer; CRC = colorectal cancer; MTC = medullary thyroid cancer; BID = twice-daily; QD = once-daily

# LOX-292: a new potent inhibitor of RET

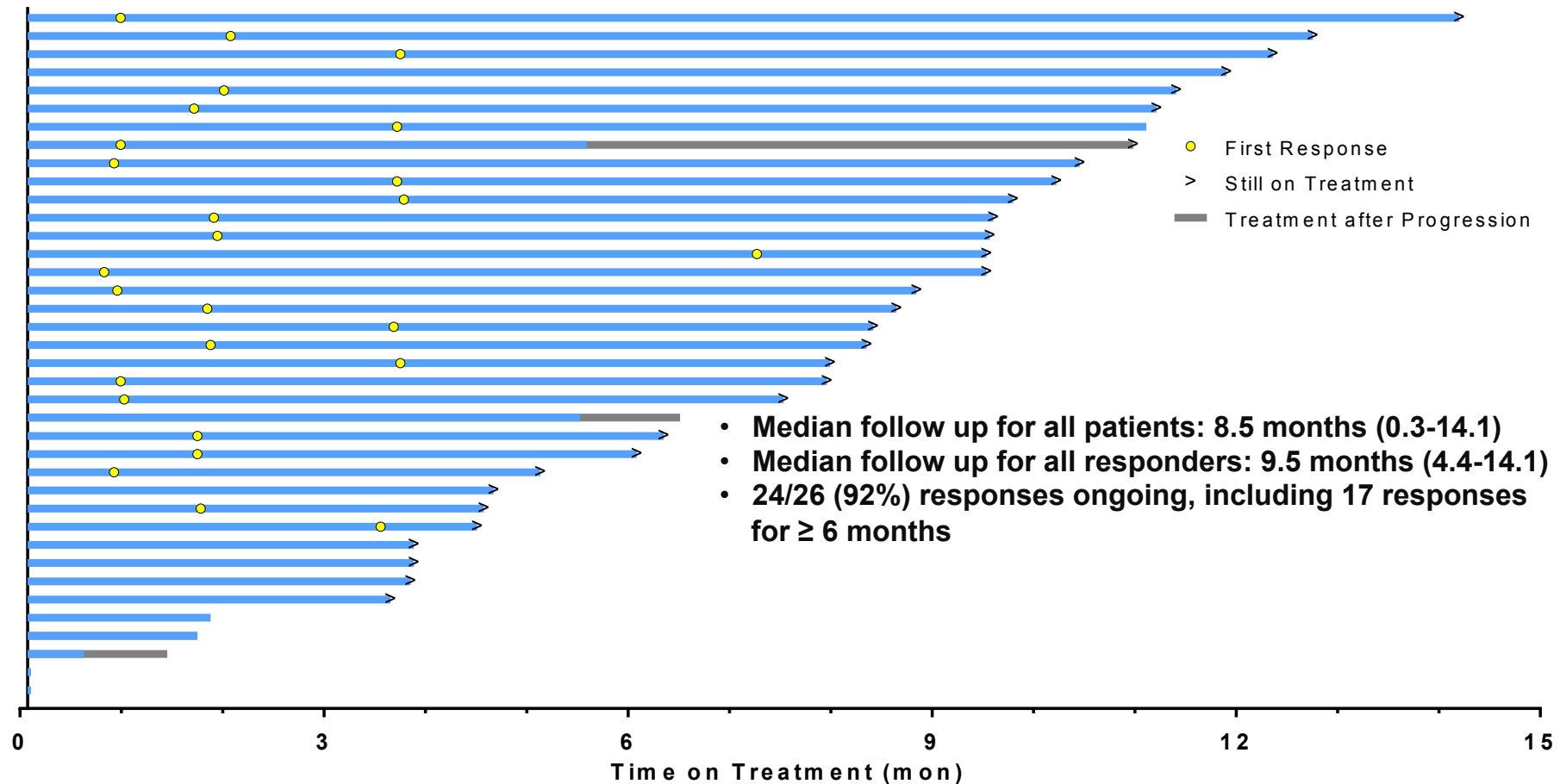


<b>RET fusion-positive NSCLC</b>	
Enrolled	38
Eligible for response evaluation	38
<b>Overall Response Rate (95% CI)</b>	<b>26/38 68% (51% - 83%)</b>
<b>Confirmed ORR*</b>	<b>25/37 68% (50% - 82%)</b>
CR	-
PR**	26
SD	8
PD	2
NE	2

**4/4 confirmed intracranial responses (1 CR, 3 PR) in patients with measurable (> 5 mm) intracranial lesions**

▼ pending confirmation; \* Excludes one patient with unconfirmed PR pending confirmation at time of data cut-off; \*\* 25 confirmed PR, 1 unconfirmed PR pending confirmation  
Follow-up as of July 19, 2018.

# Duration of LOXO-292 in *RET* fusion-positive NSCLC



NSCLC Patients enrolled as of April 2, 2018. Follow-up as of July 19, 2018.

Geoffrey R. Oxnard et al, WCLC 2018

# RET: the next big target...

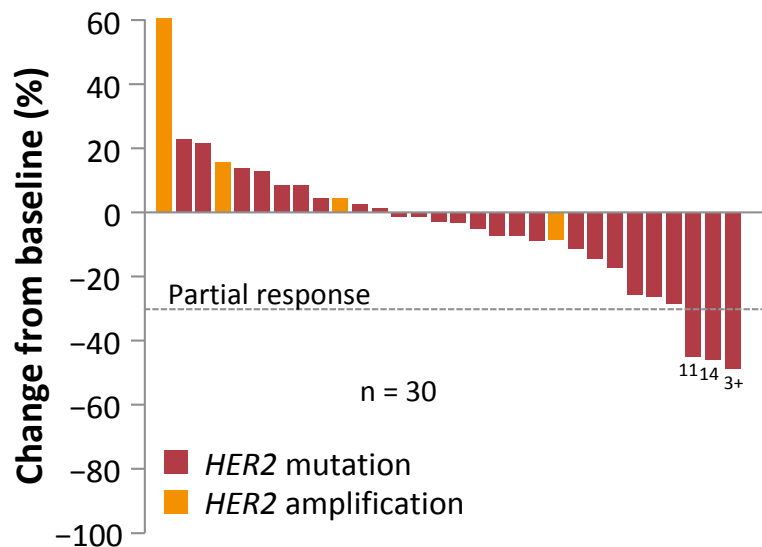
Study	Drug	n	Response rate	PFS
Drilon A, 2016	<b>Cabozantinib</b>	25	28%	NR
Lin JJ, 2016	<b>Alectinib</b>	4	50%	Duration trtt: 6m
Lee SH, 2017	<b>Vandetanib</b>	18	18%	4.5 m.
Yoh, K, 2017	<b>Vandetanib</b>	19	53%	4.7 m.
Velcheti, 2016	<b>Lenvatinib</b>	25	18%	7.3 m.
Gaustchi O, 2017	<b>Various (registry)</b>	53	18 to 37%	2.3 m.
Subbiah V, 2018 (ASCO)	<b>vandetanib + everolimus</b>	13	<b>54%</b> (7/13)	4.4m
Subbiah V, 2018 (AACR)	<b>BLU-667</b>	53 (19 NSCLC)	<b>50%</b> (NSCLC)	Duration trtt: 3.9m
Drilon A, 2018 (ASCO)	<b>LOXO-292</b>	82 (38 NSCLC)	<b>68%</b>	N.A



# Targeting *HER2* aberrations

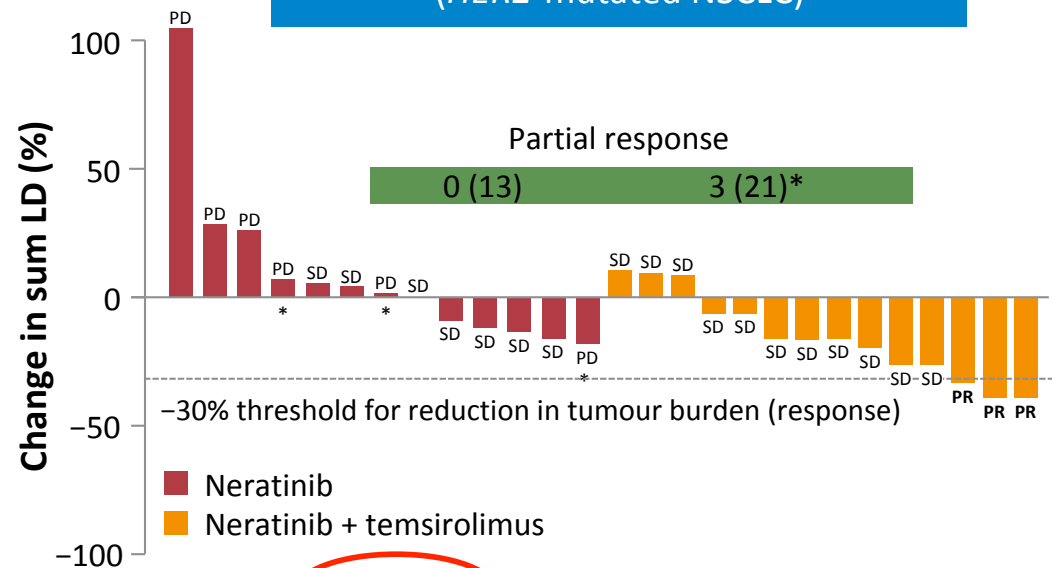
*HER2* mutations in ~1–4% and *HER2* amplifications in 2–5%

**Dacomitinib (pan-HER inhibitor)**  
(*HER2*-mutated or amplified NSCLC)



Only 3/26 of *HER2*-mutant patients had a response (**ORR 12%**)

**Neratinib (pan-HER inhibitor)**  
**± temsirolimus (mTOR inhibitor)**  
(*HER2*-mutated NSCLC)

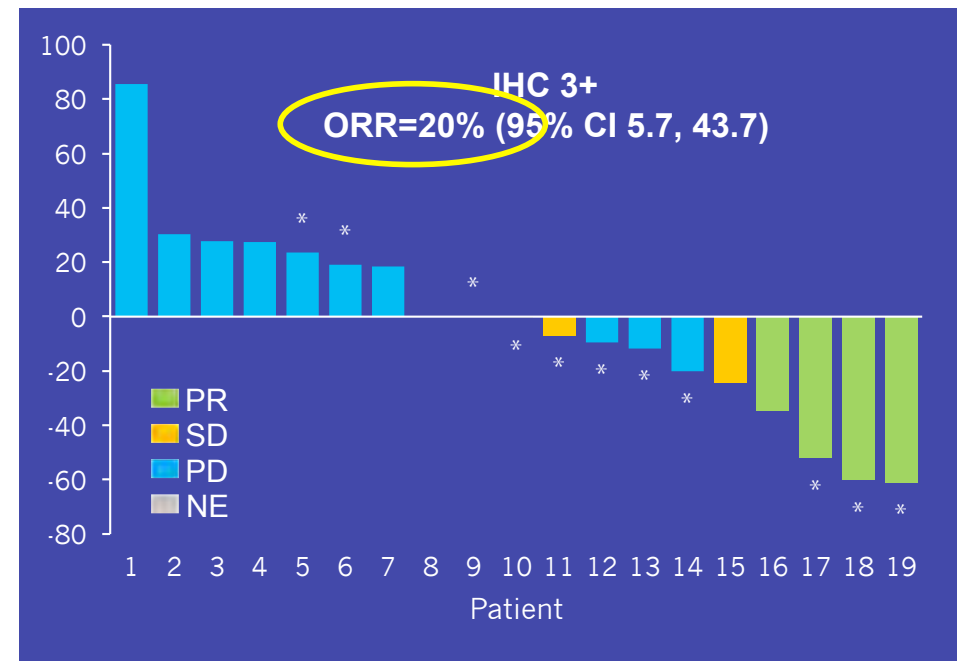
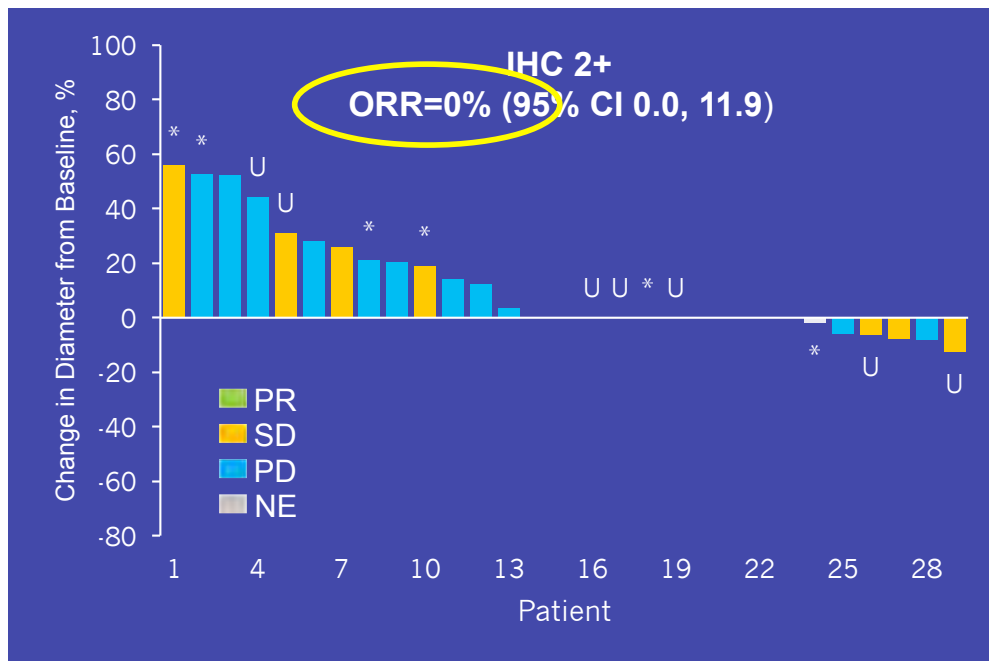


**21% ORR** and mPFS of 4 months

\* Patients had < 20% increase in tumour burden, but were considered PD due to the appearance of new lesions

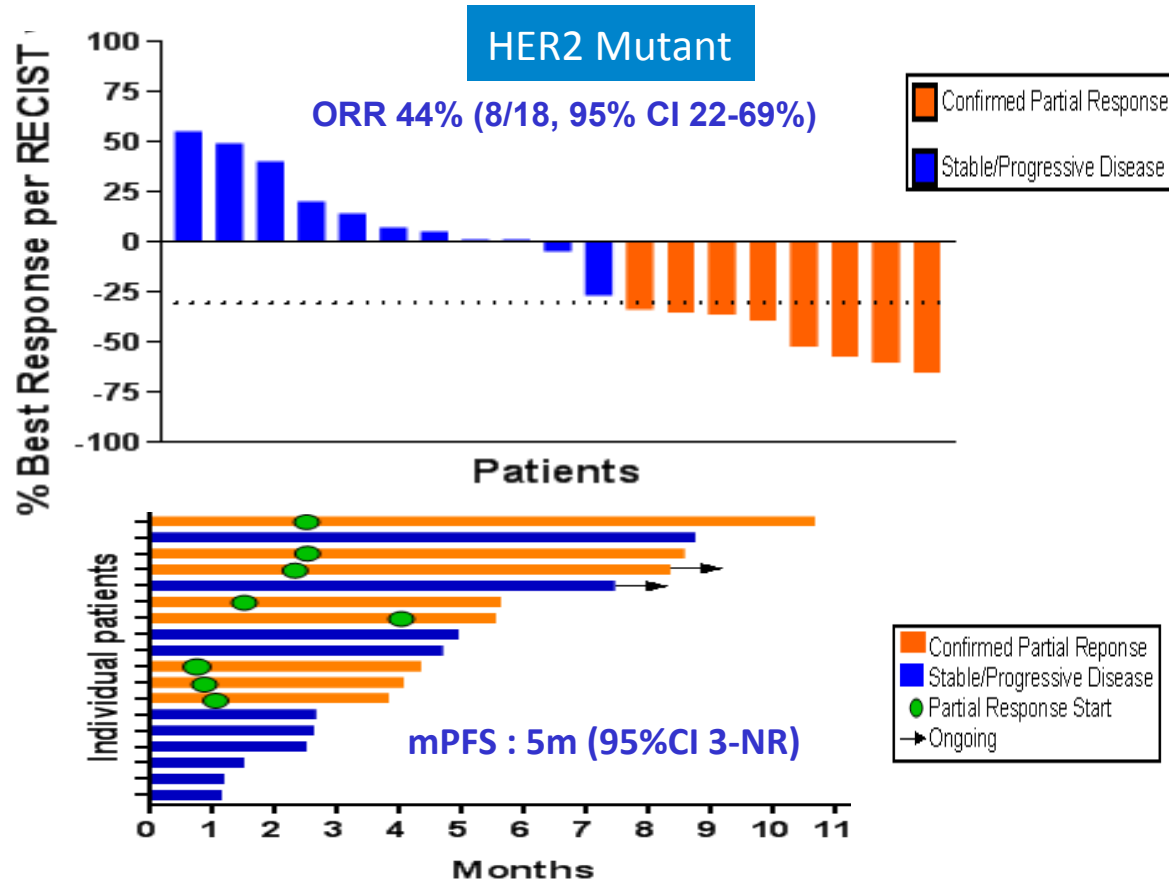
# Antibody-drug Conjugate Trastuzumab Emtansine (T-DM1) in Pts with Previously Treated HER2-Overexpressing

	IHC 2+ (n=29)	IHC 3+ (n=20)	All (N=49)
Median PFS, mo (95% CI)	2.6 (1.4, 2.8)	2.7 (1.4, 8.3)	2.6 (1.4, 2.8)



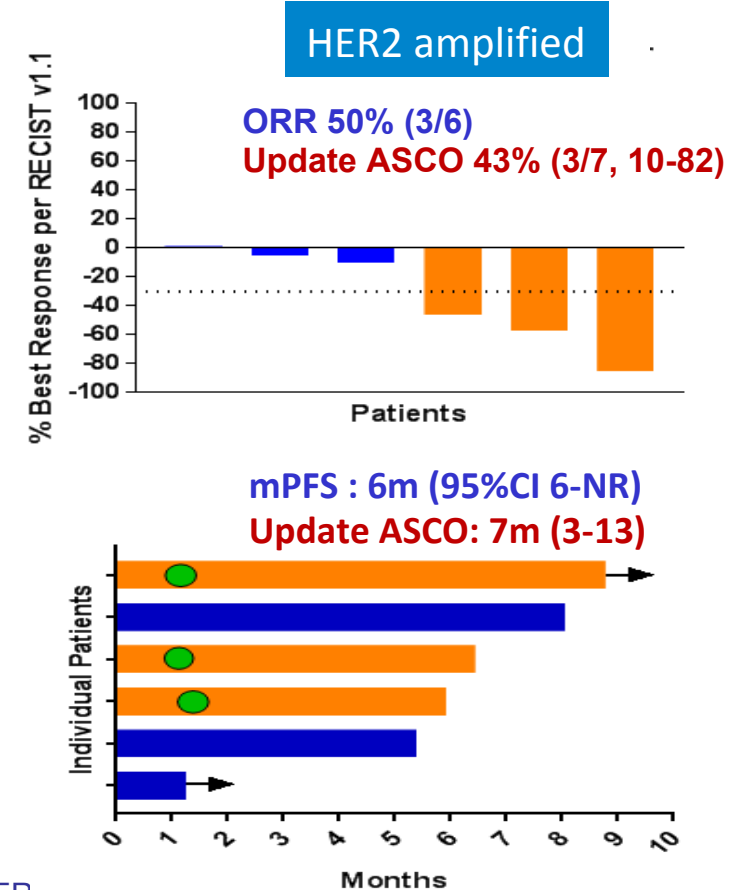
\*Indicates positive HER2 amplification; U indicates unknown HER2 amplification; All other patients' ISH status is negative

# Phase 2 trial of ado-trastuzumab emtansine for pts with *HER2* amplified or mutant cancers



6 of 8 responders were heavily pre-treated, including response to prior HER therapy neratinib, afatinib, trastuzumab

Bob T.Li et al, JCO 2018

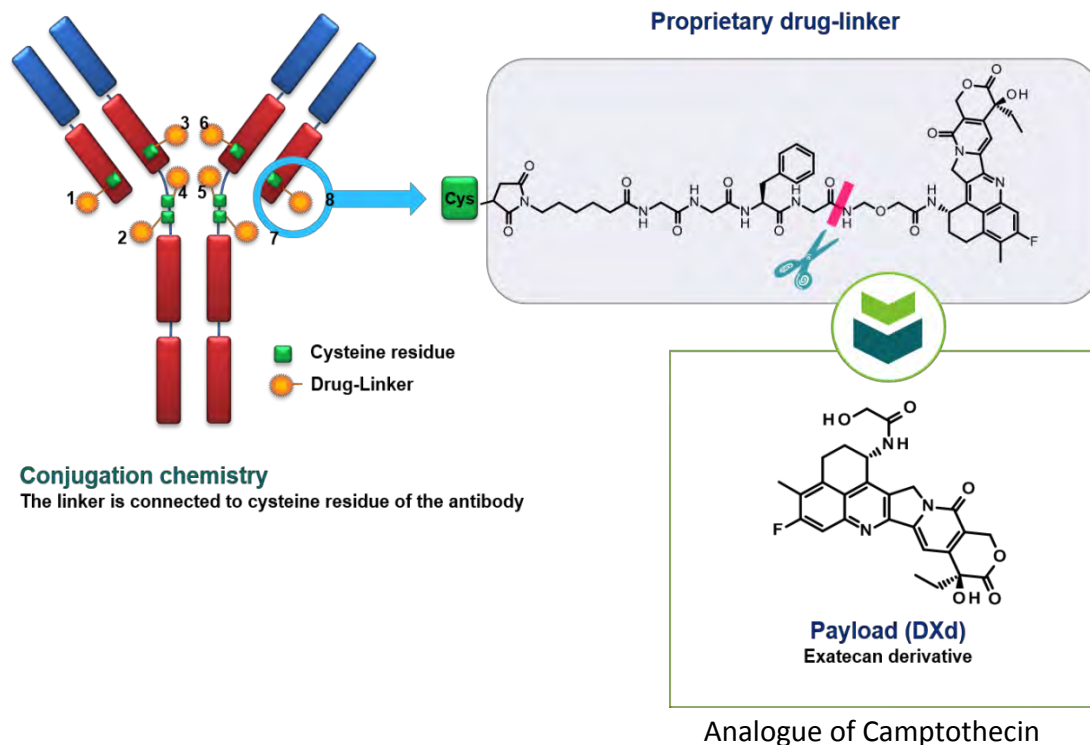


Bob T. Li, MD et al, WCLC 2017, Bob T.Li et al, ASCO 2018

# Concurrent HER2 amplification observed in 2 of 18 (11%)

NGS	FISH (HER2/CEP17)	IHC	Mass spectrometry (amol/ug)	Partial Response
Exon 20 p.A775_G776insYVMA	1.1 (2.7/2.5)	0	NA	Yes
Exon 20 p.A775_G776insYVMA	1.8 (8.1/4.5)	2+	642	
Exon 20 p.A775_G776insYVMA	NA	NA	NA	
Exon 20 p.A775_G776insYVMA	1.4 (4.5/3.3)	1+	586	Yes
Exon 20 p.A775_G776insYVMA	1.9 (5.6/2.9)	1+	548	Yes
Exon 20 p.G778_P780dup	1.6 (7.6/4.8)	1+	0	
Exon 20 p.G778_P780dup	1.8(4.6/2.5)	2+	507	Yes
Exon 20 p.G778_P780dup	1.4 (5.8/4.2)	2+	NA	
Exon 20 p.G778-779 insCPG	1.6(4.3/2.7)	0	NA	
Exon 20 p.G776_V777>VCV	NA	NA	NA	Yes
Exon 20 p.G776delinsVC	1.6 (5.7/3.6)	0	205	Yes
Exon 19 p.L755P	1.5 (3.2/2.1)	2+	434	
Exon 19 p.L755P	NA	0	NA	
Exon 17 p.V659E	1.2 (2.4/2.0)	2+	NA	
Exon 17 p.V659E	1.1 (2.3/2.0)	2+	688	Yes
Exon 8 p.S310F amplification fold change 2.8	4.1 (8.4/2.5)	2+	1495	Yes
Exon 8 p.S310F	1.8 (3.2/1.8)	0	0	
Exon 8 p.S335C	2.4 (4.8/2.0)	2+	902	

# Updated results of a phase 1 study of DS-8201a in HER2-expressing or -mutated advanced NSCLC



*Payload with a different mechanism of action*

*High potency of payload*

*Payload with short systemic half-life*

*Bystander effect*

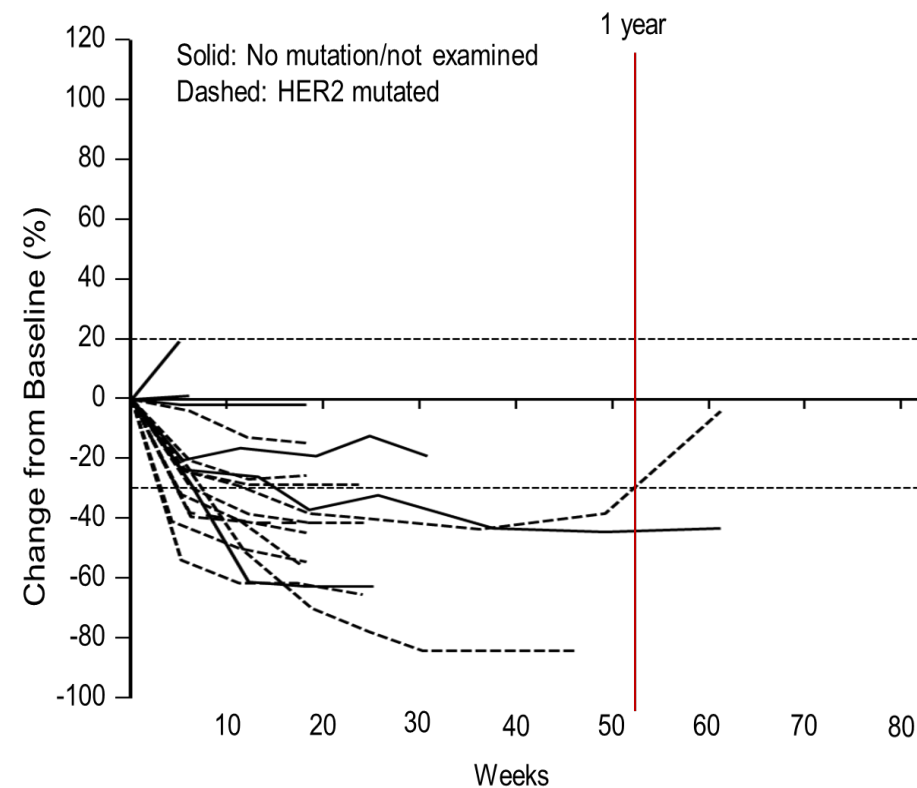
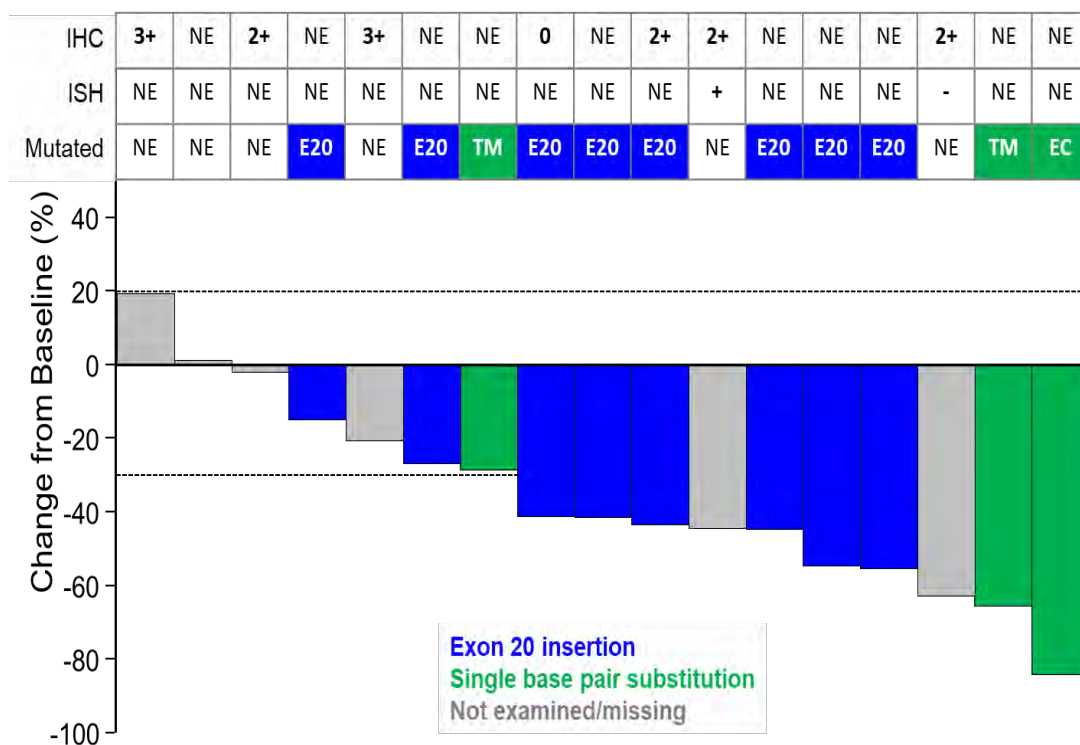
*Stable linker-payload*

*Tumor-selective cleavable linker*

*High drug-to-antibody ratio (7–8)*

- DS-8201a was designed with the goal of improving critical attributes of an ADC  
ADC, antibody drug conjugate.

# Updated results of a phase 1 study of DS-8201a in HER2-expressing or -mutated advanced NSCLC



IHC by local laboratory testing.

E20, exon 20 insertion; EC, single base pair substitution at extracellular domain; IHC, immunohistochemistry; ISH, in situ hybridization; NSCLC, non-small cell lung cancer;

NE, not examined or missing; TM, single base pair substitution in transmembrane domain.

Junji Tsurutani et al, WCLC 2018

# Efficacy Outcomes (Efficacy Evaluable Subjects)



	Confirmed ORR*, % (n/N)	DCR, % (n/N)	DOR, median (range), months	TTR, median (range), months	PFS, median (range), months
HER2-expressing or HER2-mutated NSCLC N = 18	<b>58.8% (10/17)</b>	83.3% (15/18)	9.9 (0.0+, 11.5)	1.4 (1.0, 4.2)	<b>14.1 (0.9, 14.1)</b>
HER2-mutated NSCLC n = 11	<b>72.7% (8/11)</b>	100% (11/11)	11.5 (0.03+, 11.5)	1.4 (1.0, 4.2)	<b>14.1 (4.0+, 14.1)</b>

Data cutoff, August 24, 2018.

\*Confirmed response includes subjects who had  $\geq 2$  postbaseline scans, had progressive disease, or discontinued treatment for any reason prior to second postbaseline scan.

+ after value indicates censoring.

DCR, disease control rate; DOR, duration of response; HER2, human epidermal growth factor receptor 2; NSCLC, non-small cell lung cancer; ORR, objective response rate;

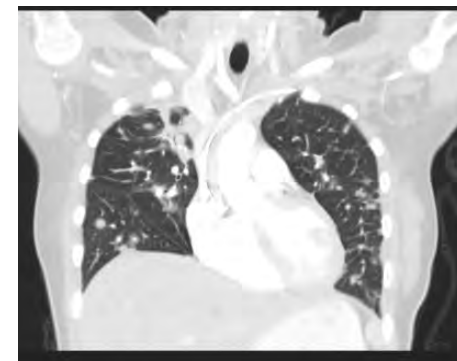
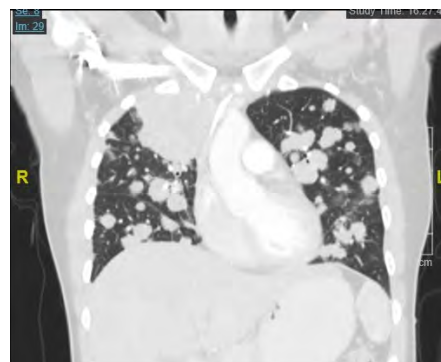
PFS, progression-free survival; TTR, time to response.

Junji Tsurutani et al, WCLC 2018

# Example CT Image from Responder to DS-8201a

- 23 years old
- Female
- Nonsmoker
- History of Type 1 Diabetes
- HER2 12 bp insertion in exon 20
- **January 2017:** presented with cough and SOB
  - Diagnosed with stage IV nonsquamous NSCLC
  - Carbo/Pem 1 cycle
- **February–June 2017:** switched to Carbo/Nab-paclitaxel due to LFT elevations
  - Best response SD
- **September–December 2017:** switched to Carbo/Pem due to progression
  - Four cycles
  - Best response SD
  - Last scan with slight increase in disease
  - Recommended HER2 targeted therapy; came to DFCI
- **February 2018:** started DS-8201a
  - Symptomatic with cough and DOE
  - Status: PR (confirmed)

HER2 insertion exon 20



February 2018 –  
baseline

May 2018 –  
C5D1

Images courtesy of Dr. Pasi Jänne. Special thanks to Dr. Pasi Jänne and Dr. Ian Krop of DFCI.

bp, base pair; Carbo, carboplatin; CT, computed tomography; DFCI, Dana-Farber Cancer Institute; DOE, dyspnea on exertion; HER2, human epidermal growth factor 2; IV, intravenous; LFT, liver function test; Nab, nab-paclitaxel; NSCLC, non-small cell lung cancer; Pem, pemetrexed; PR, partial remission; SD, stable disease; SOB, shortness of breath.

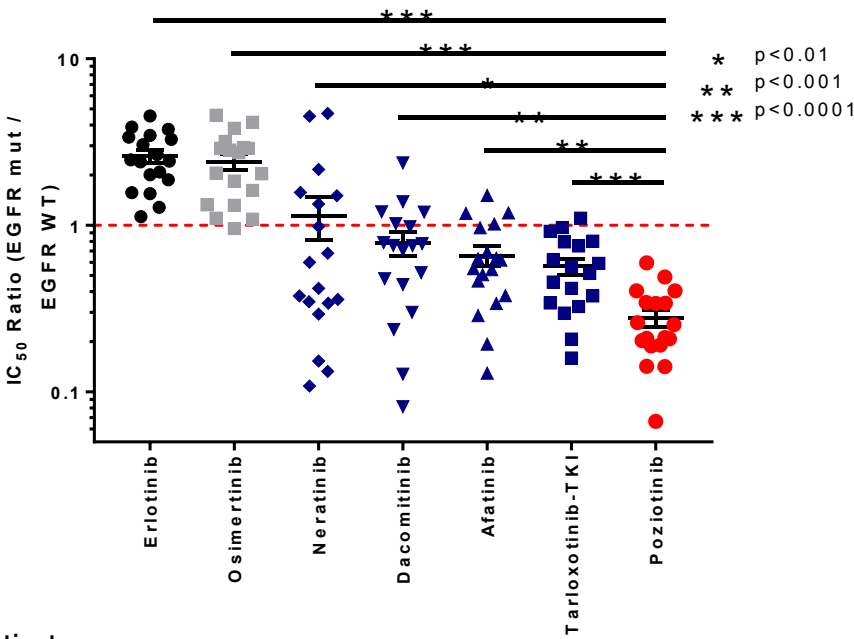
Junji Tsurutani et al, WCLC 2018



# Poziotinib is a selective (mut vs wt) inhibitor of EGFR and HER2 exon 20 mutations *in vitro*

## EGFR

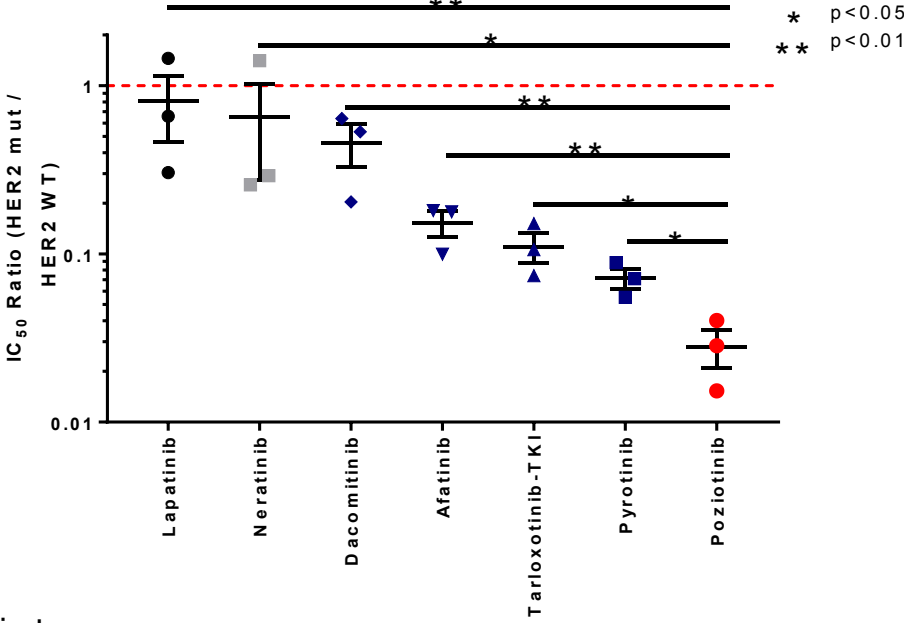
EGFR Ba/F3 Selectivity Index  
(N=20 cell lines)



Ratio to Poziotinib	9.4	8.7	4.1	2.8	2.4	2.0
---------------------	-----	-----	-----	-----	-----	-----

## HER2

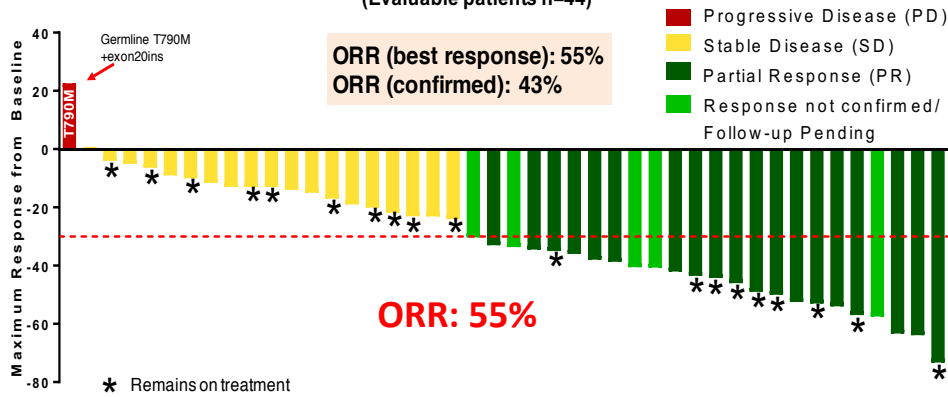
MCF10A Selectivity Index  
(N=3 Cell lines)



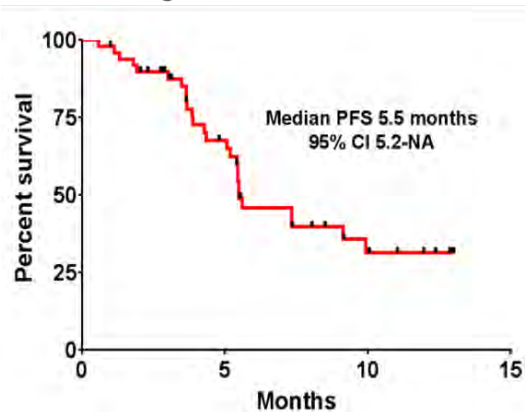
Ratio to Poziotinib	28.7	23.3	16.3	5.4	4.0	2.6
---------------------	------	------	------	-----	-----	-----

# A Phase II Trial of Poziotinib in EGFR and HER2 exon 20 Mutant Non-Small Cell Lung Cancer (NSCLC)

**Poziotinib efficacy in EGFR Exon 20 mutant NSCLC**  
(Evaluable patients n=44)

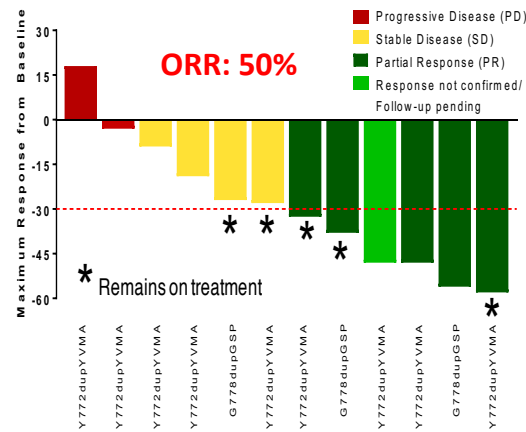


**Progression Free Survival**

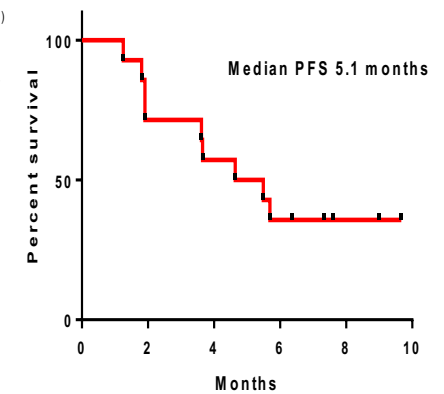


**Poziotinib efficacy in HER2 Exon 20 insertion mutant NSCLC**

**Best response HER2**  
(Evaluable patients n=12)



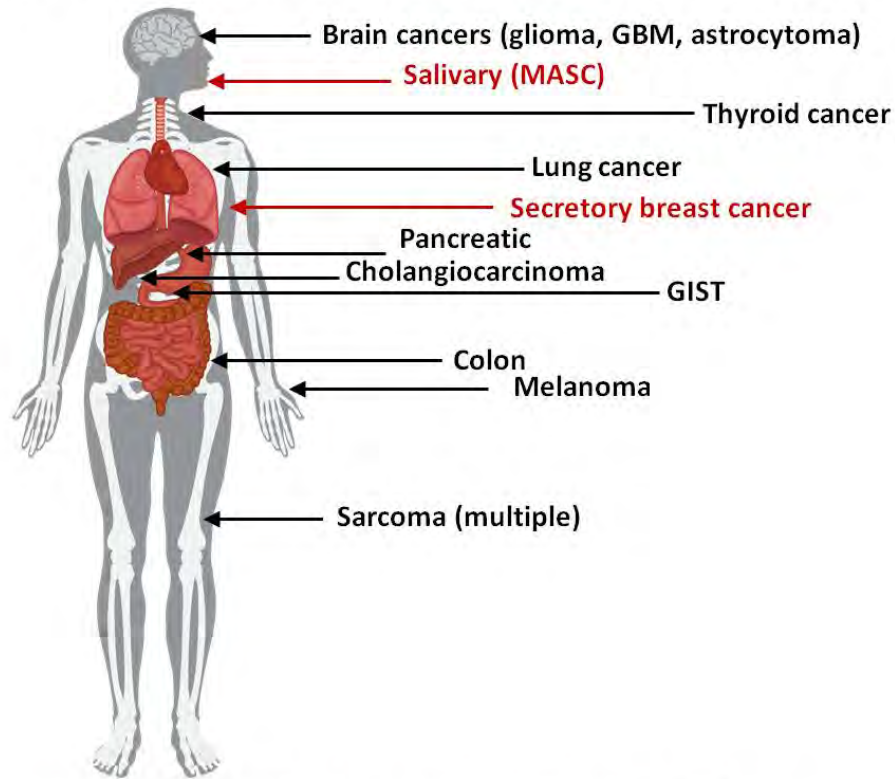
**Progression-free Survival HER2**  
(All patients n=13)



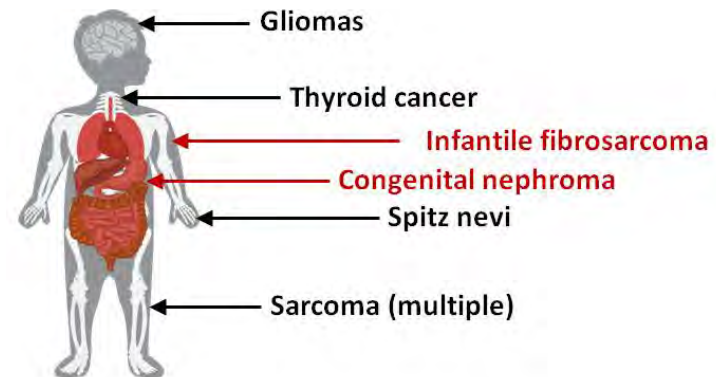
## Conclusions:

Encouraging activity has prompted a confirmatory, international, multicenter study in *EGFR* and *HER2* exon 20 mutant NSCLC patients which is currently enrolling (NCT03318939) including a first-line cohort and development of a pan-tumor basket study

# TRK fusions found in diverse cancer histologies

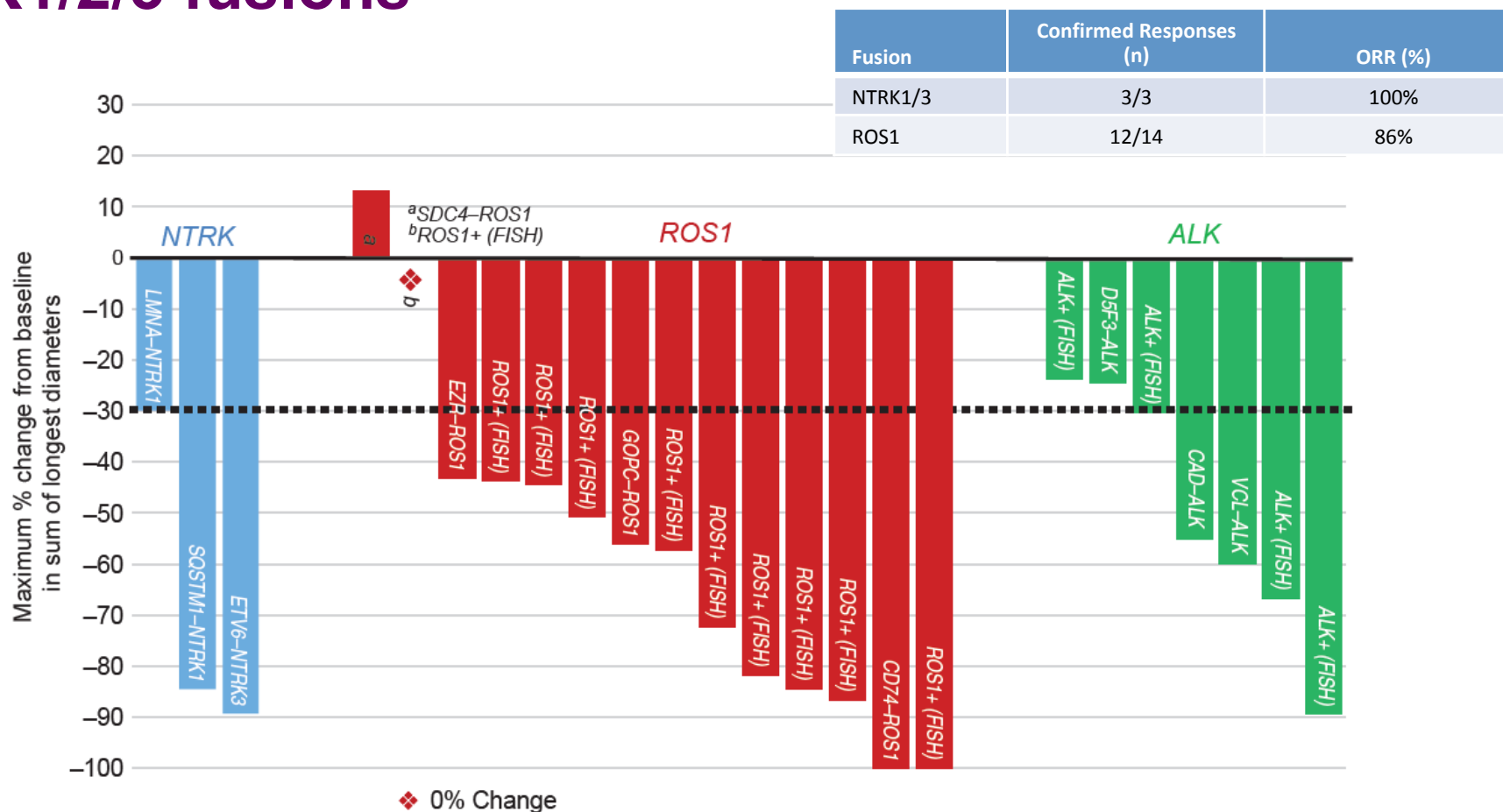


- Common cancer with low TRK fusion frequency
- Rare cancer with high TRK fusion frequency



Estimated 1,500–5,000 patients harbor TRK fusion-positive cancers in the United States annually

# Entrectinib, first in class, antitumor activity in NTRK1/2/3 fusions



# STARTRK-2 NTRK Enrollment

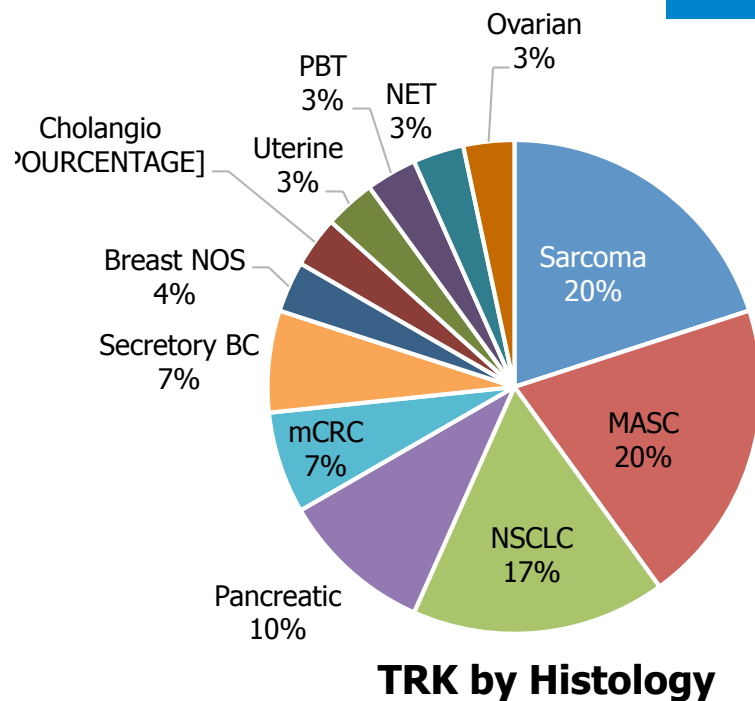
Assignment by Gene Fusion and Tumor Type

TRK

ROS1

ALK

Non  
Evaluable

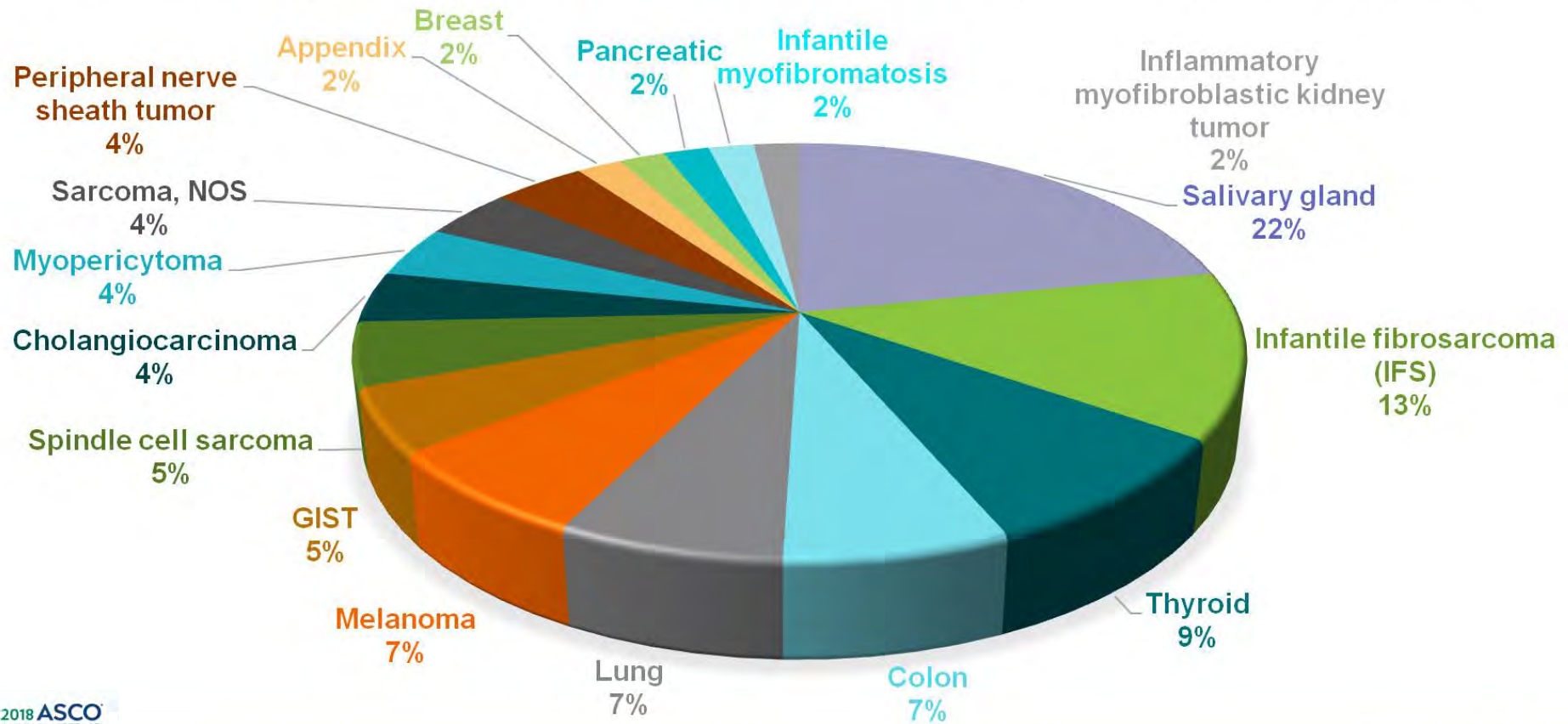


**TRK** enrollment in STARTRK-2 has been consistent with the diffuse distribution pattern described in the literature

- Patients with TRK fusions enrolled across >15 different tumor types
- All of these patients were identified using next generation sequencing (NGS)

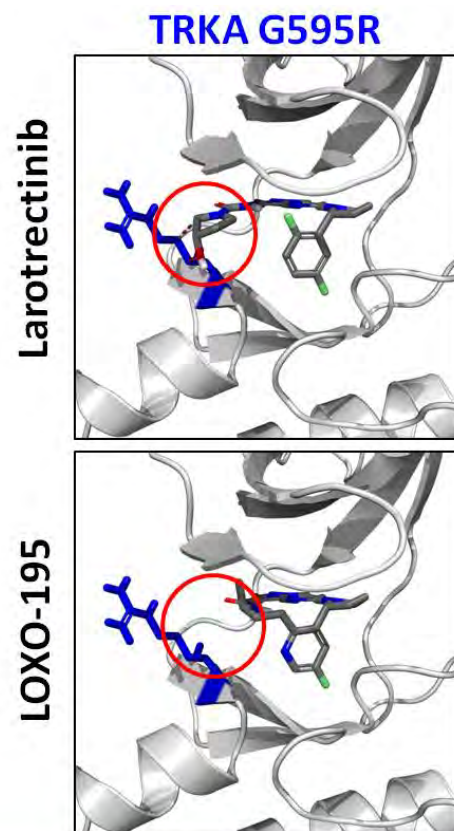
# Larotrectinib (LOXO-101), a selective TRK inhibitor

## Diversity of cancers treated - 17 unique types





# LOXO-195 to Address TRK Acquired Resistance

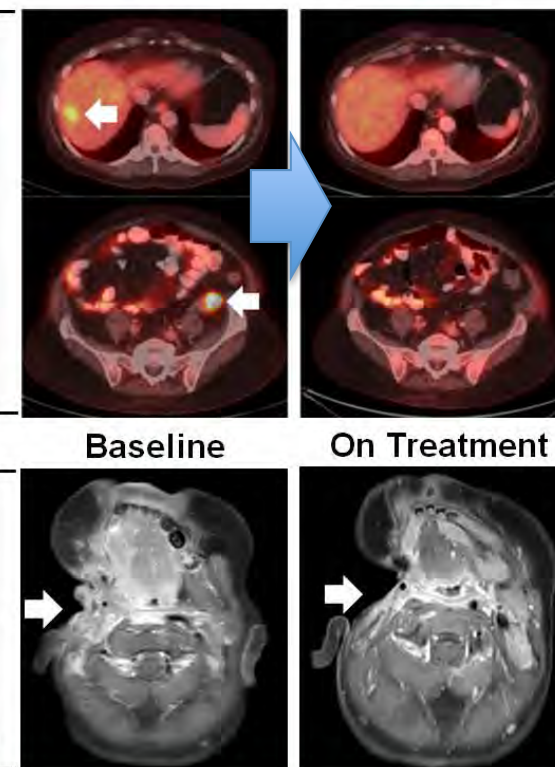


## Larotrectinib (LOXO-101)

Tumor type	Fusion	Resistance mutation
Colorectal	TPM3-NTRK1	TRKA G595R
Colorectal	LMNA-NTRK1	TRKA G595R
NSCLC	TPR-NTRK1	TRKA G595R
Sarcoma*	TPM3-NTRK1	TRKA G595R
IFS	ETV6-NTRK3	TRKC G623R
Cholangio*	LMNA-NTRK1	TRKA F589L* + GNAS Q227H

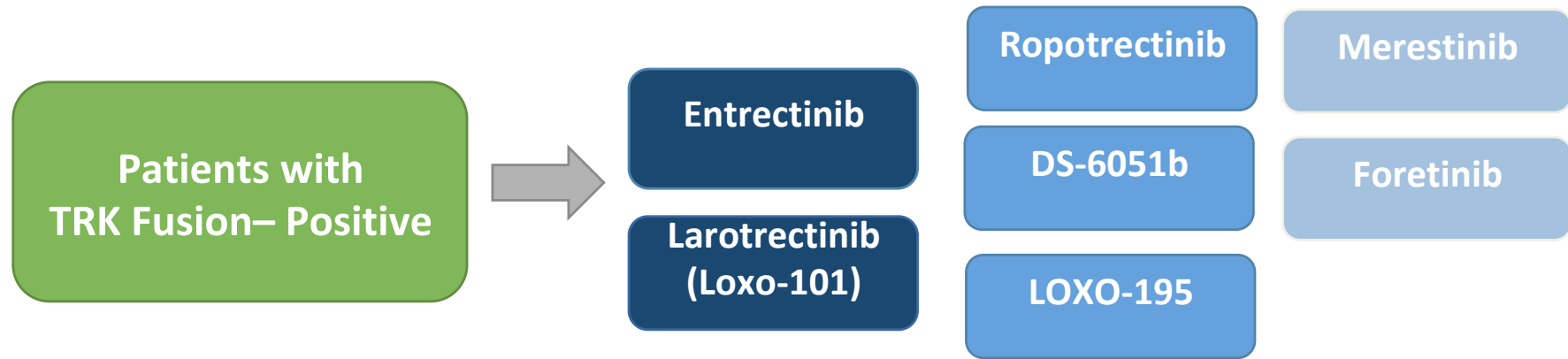
**TRK solvent front mutations detected in 5 of 6 patients with acquired resistance. First 2 patients successfully treated with LOXO-195.**

## LOXO-195 Treatment





# New data to come...



## **A phase 1 study of the next-generation ALK/ROS1/TRK inhibitor ropotrectinib (TPX-0005) in patients with advanced ALK/ROS1/NTRK+ cancers (TRIDENT-1)**

Presented Monday, June 4, 2018. Alexander E. Drilon (Abst 2513)

65 pts (28 ALK+, 29 ROS1+, and 8 NTRK+)

DS-6051b is an oral, small molecule receptor TKI with high affinity for ROS1 and NTRK kinases

## **First-in-human study of DS-6051b in patients (pts) with advanced solid tumors (AST) conducted in the US**

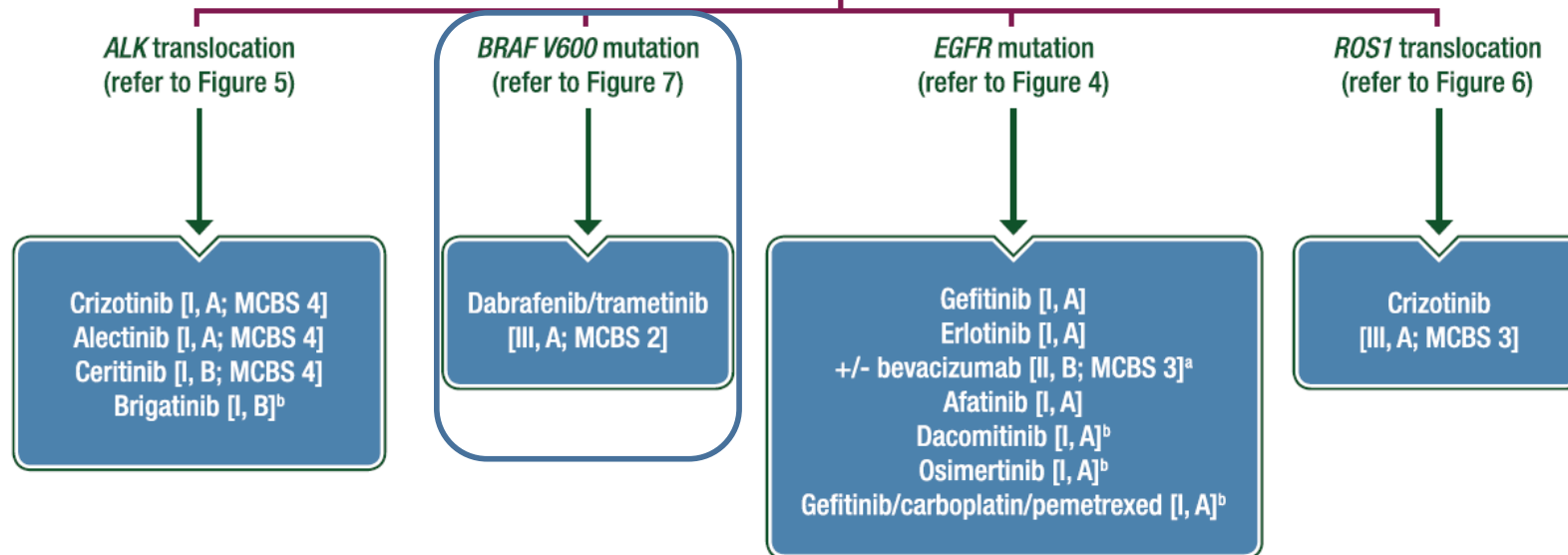
Presented Monday, June 4, 2018. Kyriakos P. Papadopoulos (Abst 2514)

CLINICAL PRACTICE GUIDELINES

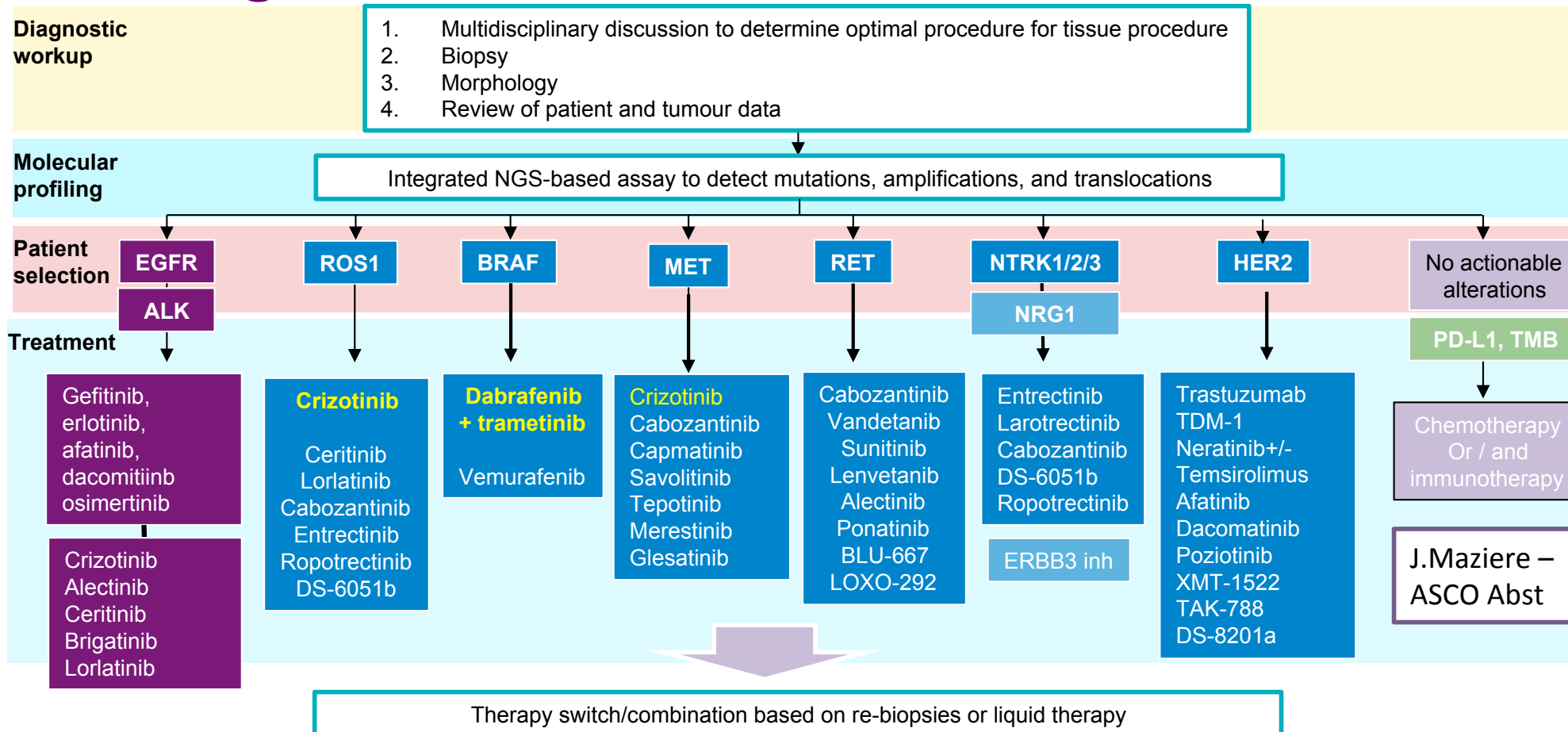
Metastatic non-small cell lung cancer: ESMO  
Clinical Practice Guidelines for diagnosis, treatment  
and follow-up<sup>†</sup>

D. Planchard<sup>1</sup>, S. Popat<sup>2</sup>, K. Kerr<sup>3</sup>, S. Novello<sup>4</sup>, E. F. Smit<sup>5</sup>, C. Faivre-Finn<sup>6</sup>, T. S. Mok<sup>7</sup>, M. Reck<sup>8</sup>,  
P. E. Van Schil<sup>9</sup>, M. D. Hellmann<sup>10</sup> & S. Peters<sup>11</sup>, on behalf of the ESMO Guidelines Committee\*

**Stage IV NSCC: Molecular tests positive (*ALK/BRAF/EGFR/ROS1*)**



# In summary, promising new drugs for old or new targets...



# THANK YOU !

## Acknowledgments

Benjamin BESSE  
Jean-Charles SORIA  
Frank Aboubakar  
Thierry LE CHEVALIER  
Laura MEZQUITA

**GUSTAVE  
ROUSSY**  
CANCER CAMPUS  
GRAND PARIS

