



# Targeted therapies in NSCLC: MET, BRAF, RET & NTRK

Janakiraman Subramanian MD, MPH

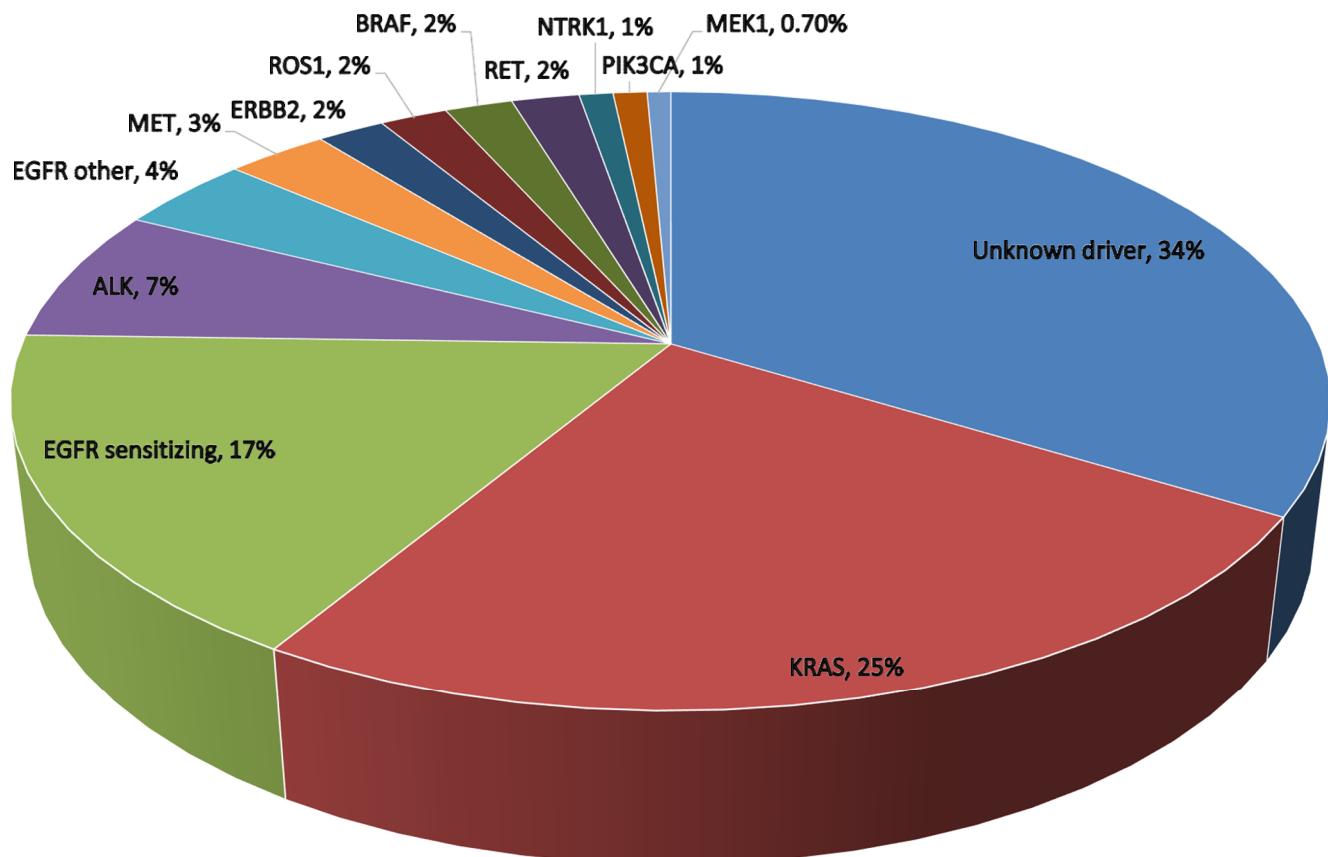


# Disclosures

- Research funding: Novartis, Merck, CanStem, Helsinn, Biocept, Incyte, Genetech & Paradigm
- Advisory role: Astra Zeneca, Boehringer Ingelheim, Novartis, Eli Lilly & Pfizer
- Speakers bureau: Astra Zeneca & Boehringer Ingelheim



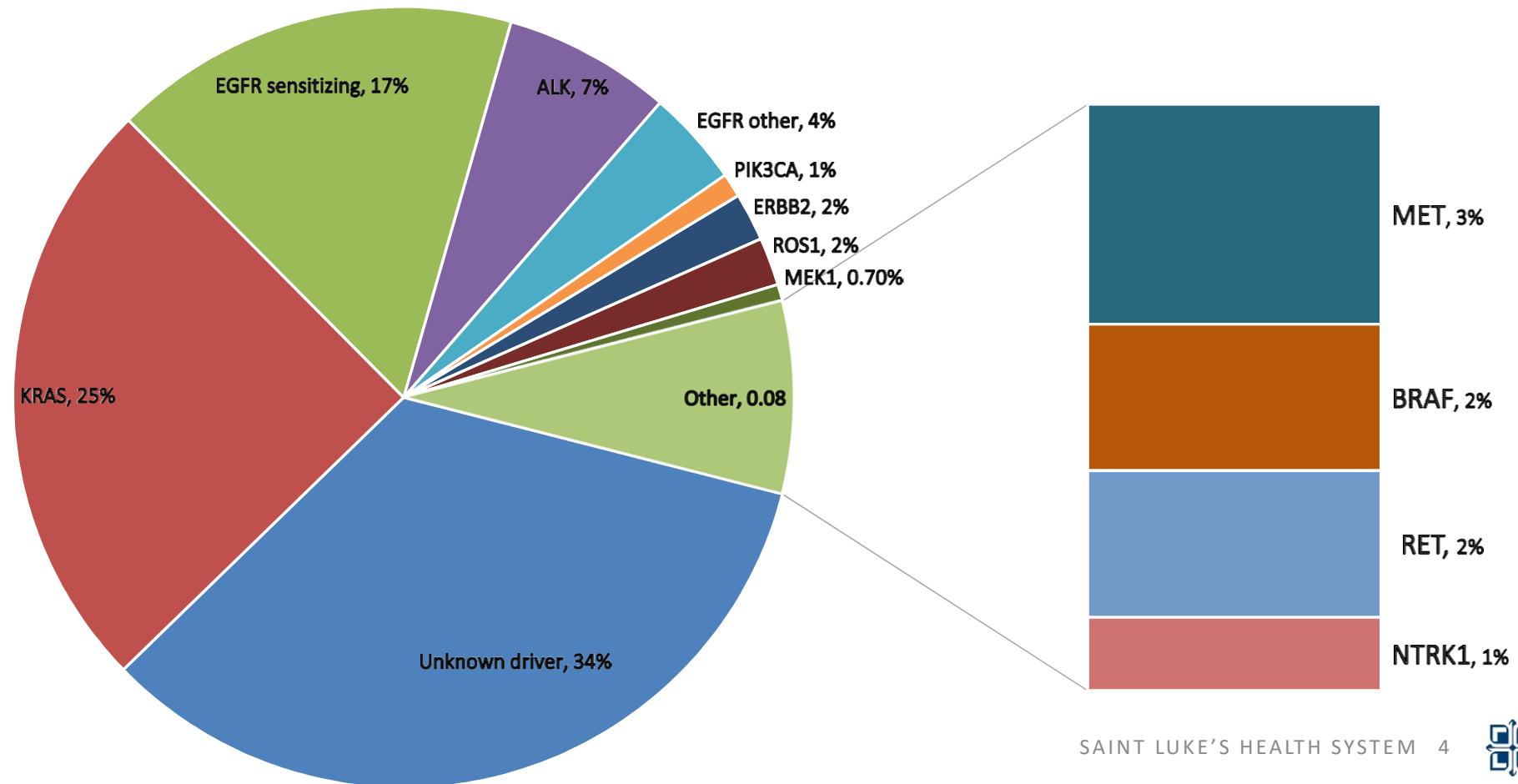
# Drivers in Lung Adenocarcinoma



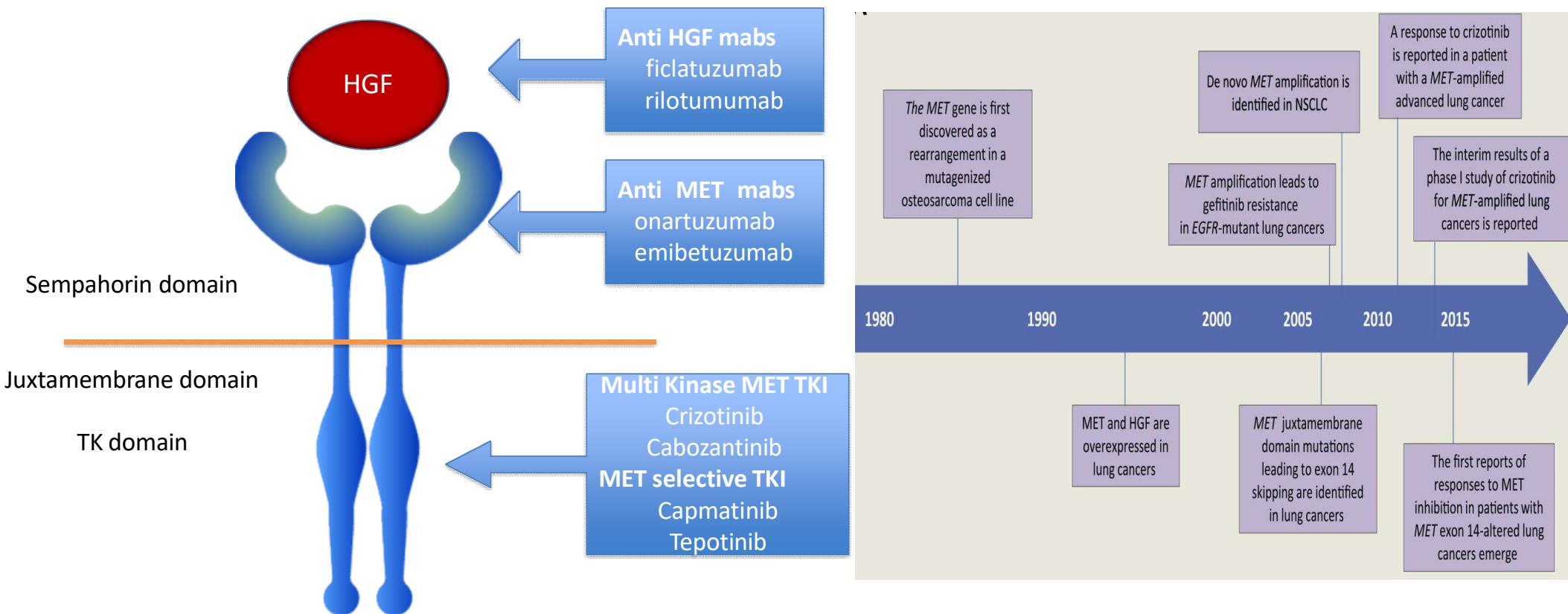
Adapted from <https://mycancergenome.org>



# BRAF, MET, RET & NTRK in Lung Adenocarcinoma



# MET alterations in NSCLC

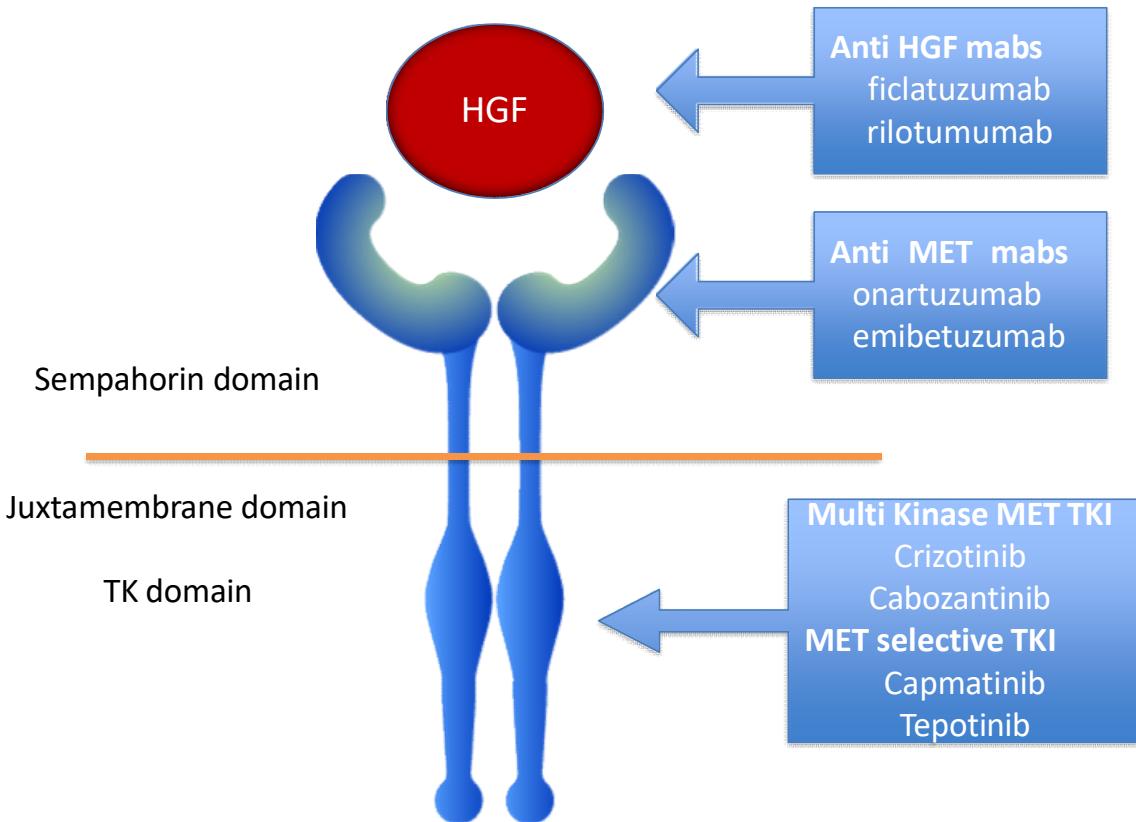


Drilon JTO 2017

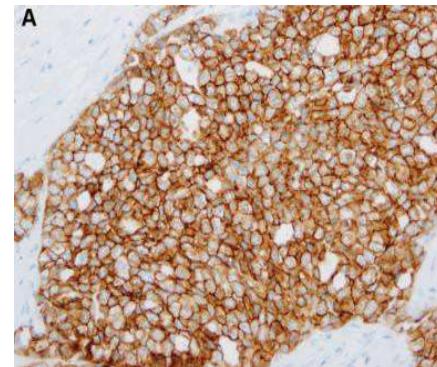
SAINT LUKE'S HEALTH SYSTEM 5



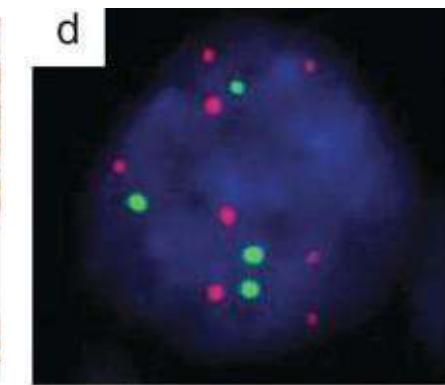
# MET alterations in NSCLC



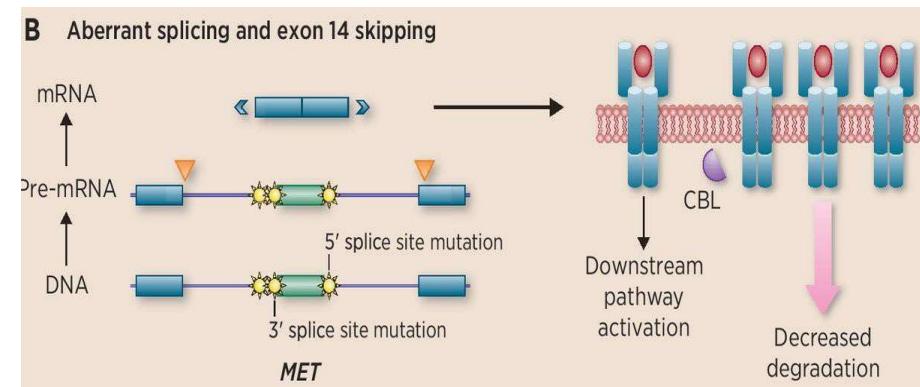
MET Over-expression  
25%-75%



MET Amplification  
5%-25%



MET Exon 14 skipping  
3%

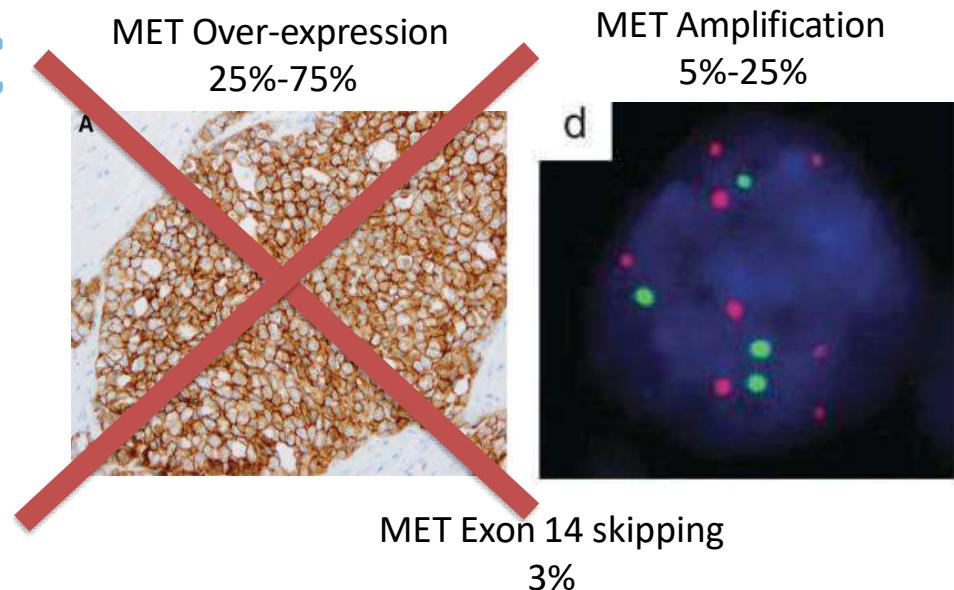
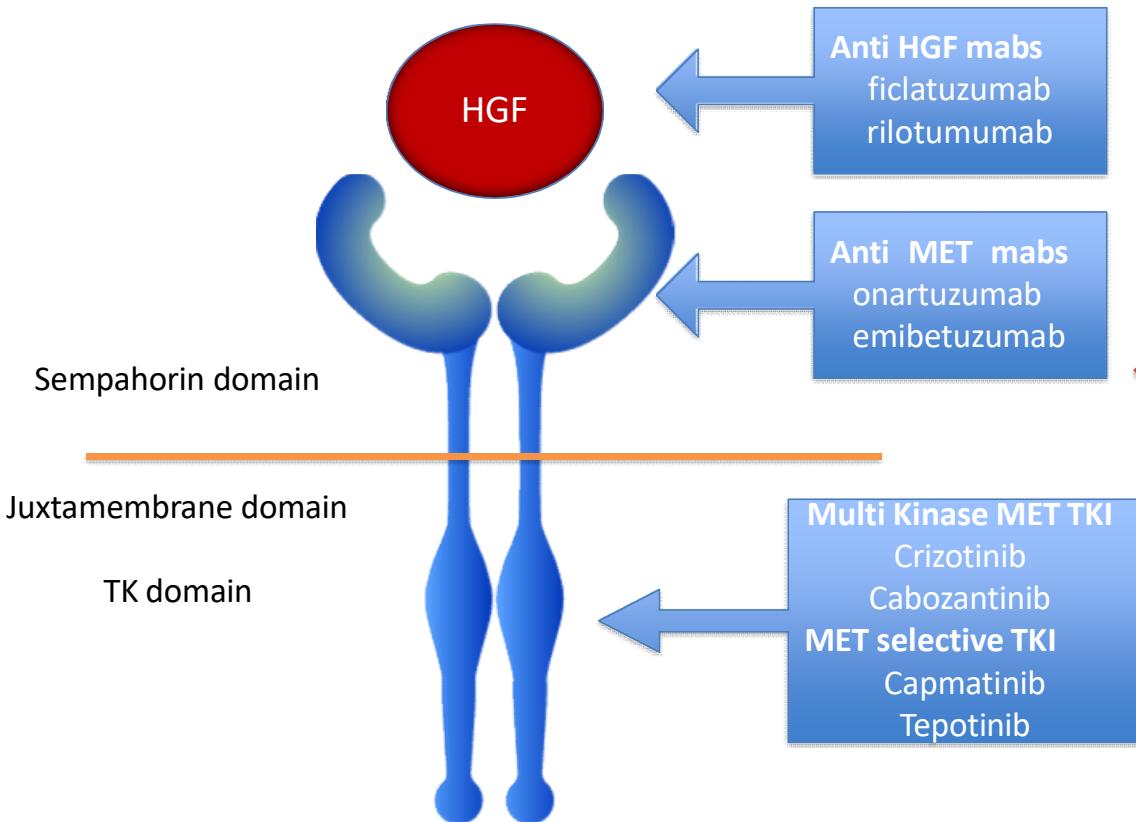


Drilon JTO 2017, Tong Clin Cancer Rsrch 2016, Drilon Clin Cancer Rsrch 2016

SAINT LUKE'S HEALTH SYSTEM 6



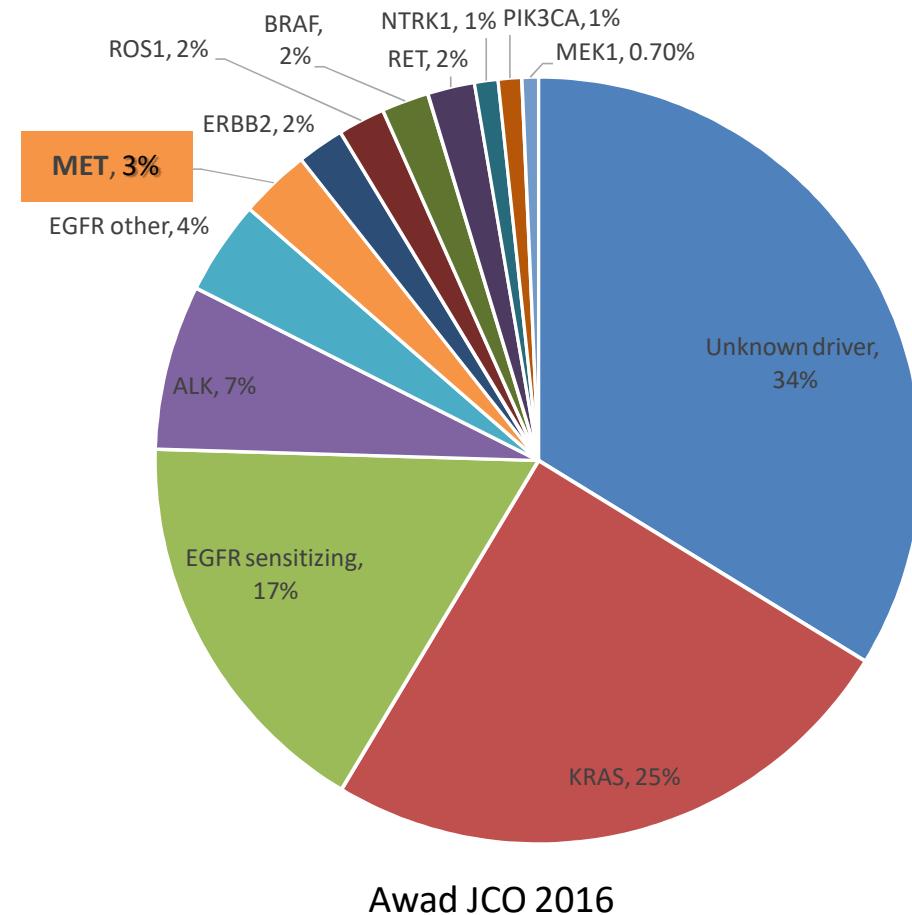
# MET alterations in NSCLC



Drilon JTO 2017, Tong Clin Cancer Rsrch 2016, Drilon Clin Cancer Rsrch 2016



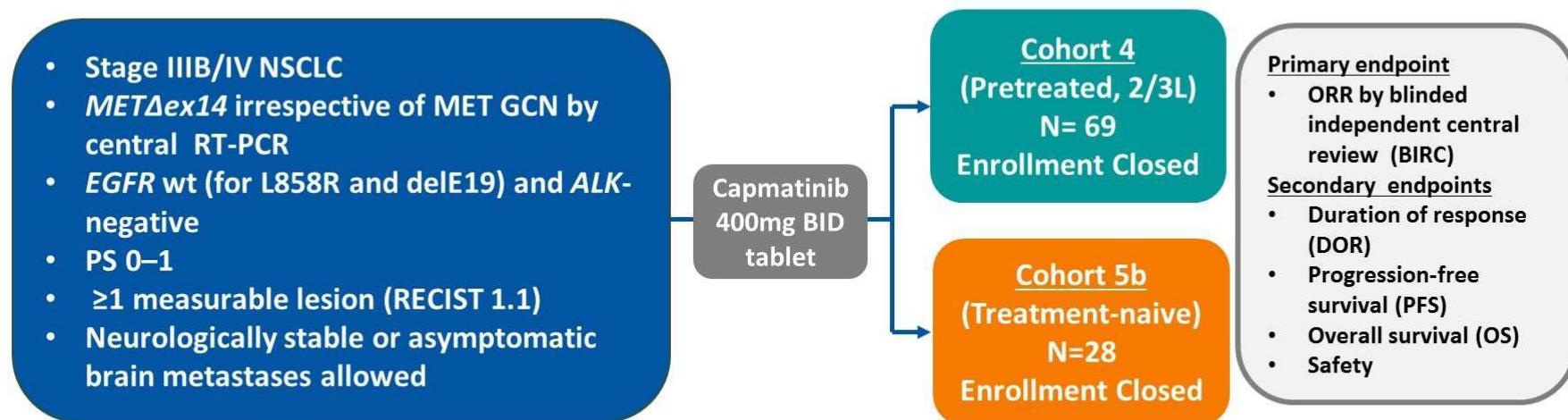
# MET Exon 14 Skipping mutation



- Older patients, median age > 70 yrs
- > 60% were tobacco smokers
- Predominantly adenocarcinoma
- Sarcomatoid pleomorphic type
- Mutually exclusive with KRAS and EGFR activating mutations
- Concurrent MET amplification in 20%.



# GEOMETRY mono-1: A phase II trial of capmatinib in patients with advanced NSCLC harboring *MET exon14 skipping* mutation



## Study methodology:

- Cohort 4 and 5b are each analyzed separately and have independent statistical hypothesis
- Primary (ORR) and key secondary (DOR) endpoints based on BIRC including 2 parallel independent radiology reviewers (+ additional one for adjudication)
- Efficacy endpoints based on BIRC and investigator assessment per RECIST 1.1

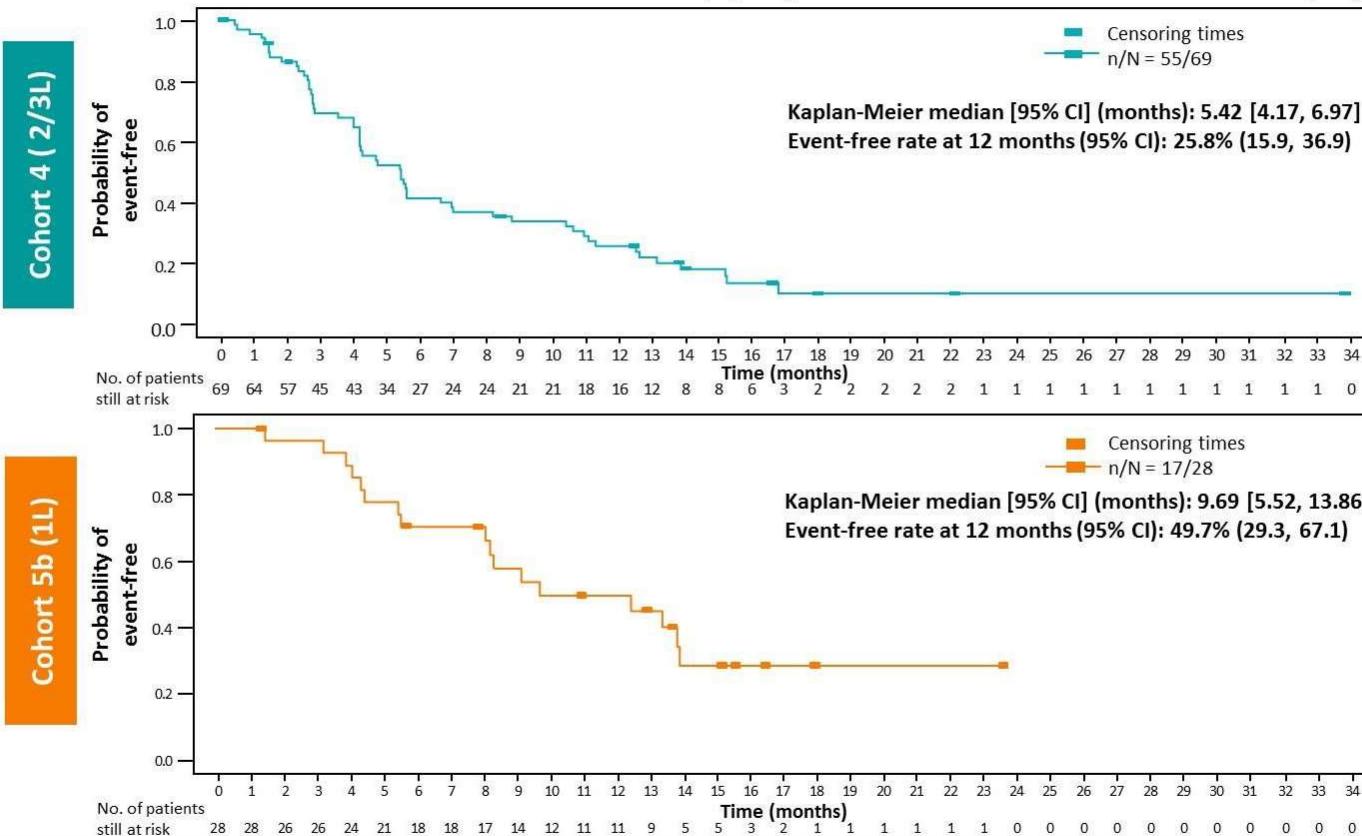
Data cut off: April 15, 2019; median duration of follow-up for DOR: 9.7 months in Cohort 4 and 9.6 months in Cohort 5b

Additional data on *MET* mutated patients will be generated in Cohort 6 (2L; N~30) and Cohort 7 (1L; N~27)



## Progression-free survival per BIRC

*Median PFS was 5.42 months in Cohort 4 (2/3L) and 9.69 months in Cohort 5b (1L)*



*Median PFS per investigator was 4.80 months (95% CI: 4.11, 7.75) in Cohort 4 and 11.14 months (95% CI: 5.52, 15.24) in Cohort 5b* 11

n is the number of events, N is the number of patients



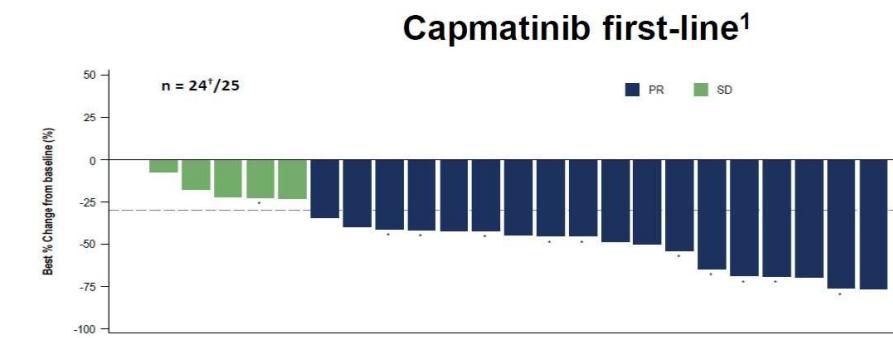
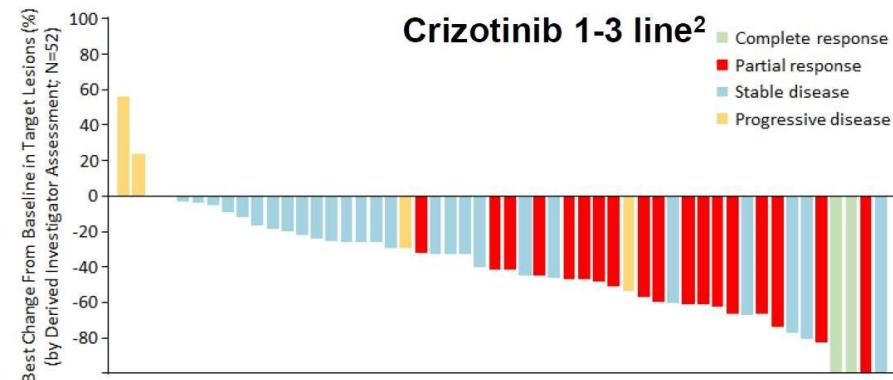
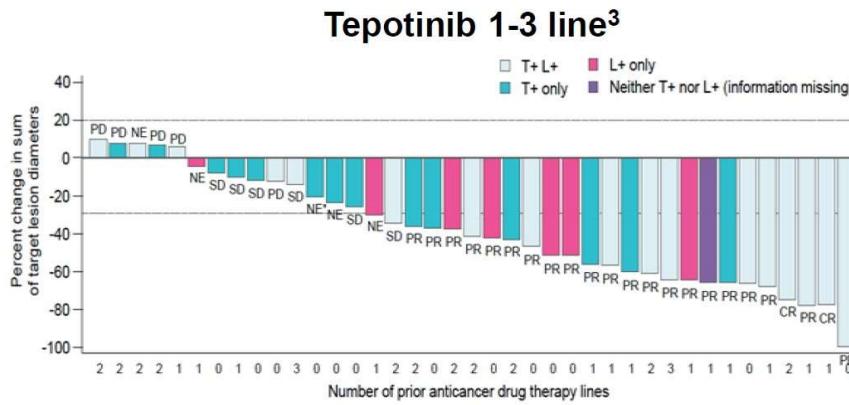
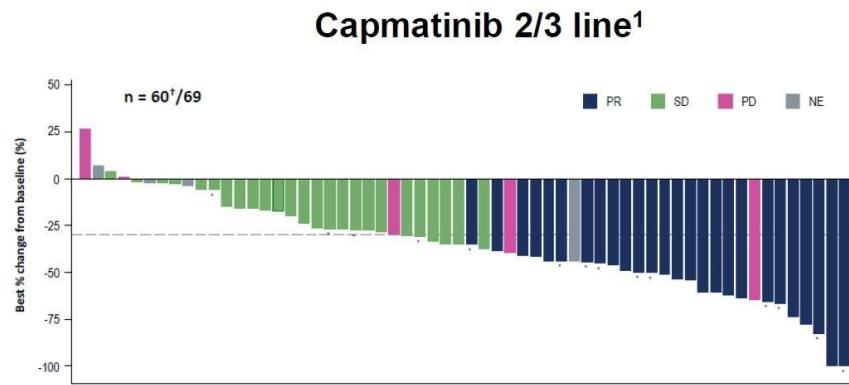
# MET TKI – Clinical Trials

Agent	Line	Testing	N	ORR (95% CI)	DOR (months)	PFS (months)
Crizotinib	1-3	Tumor/ ctDNA	65	32% (21-45)	9.1	7.3
Capmatinib	1	Tumor	28	67.9% (47.6-84.1)	11.1	9.7
	2/3	Tumor	69	40.6% (28.9-53.1)	9.7	5.4
Tepotinib	1-3	Tumor/ ctDNA	32 Tumor 24 ctDNA	45% (31.1-59.7) 50% (35.2-64.8)	15.7 12.7	10.8 9.5
Savolitinib	1-3	Tumor	31	54.8%	NA	NA

Drilon et al WCLC 2018, Wolf et al ASCO 2019, Paik et al ASCO 2019, Lu et al ASCO 2019



# Is One Better Than The Other?



Drilon et al WCLC 2018, Wolf et al ASCO 2019, Paik et al ASCO 2019, Lu et al ASCO 2019



# Adverse Events

All Grade (Grade $\geq$ 3) %	Crizotinib	Capmatinib	Tepotinib	Savolitinib
Edema	47 (3)	58.7 (4.3)	40.4 (6.3)	36.6 (4.9)
Vision disorder	40 (0)	-	-	-
Nausea	33 (0)	21.7 (0)	32.8 (1.7)	41.5 (0)
Diarrhea	30 (0)	37 (2.2)	11.6 (0)	-
Vomiting	23 (0)	-	19.2 (2.0)	24.4 (0)
Constipation	10 (0)	15.2 (0)	-	7 (0)
Bradycardia	23 (1)	-	-	-
Fatigue	13 (0)	17.4 (2.2)	13.2 (3.3)	23 (2)
Elevated Cr	-	21.7 (0)	19.2 (0)	
Elevated transaminases	10 (3)	10.9 (4.3)	-	26.8 (7.3)

Drilon et al WCLC 2018, Wolf et al ASCO 2019, Paik et al ASCO 2019, Lu et al ASCO 2019



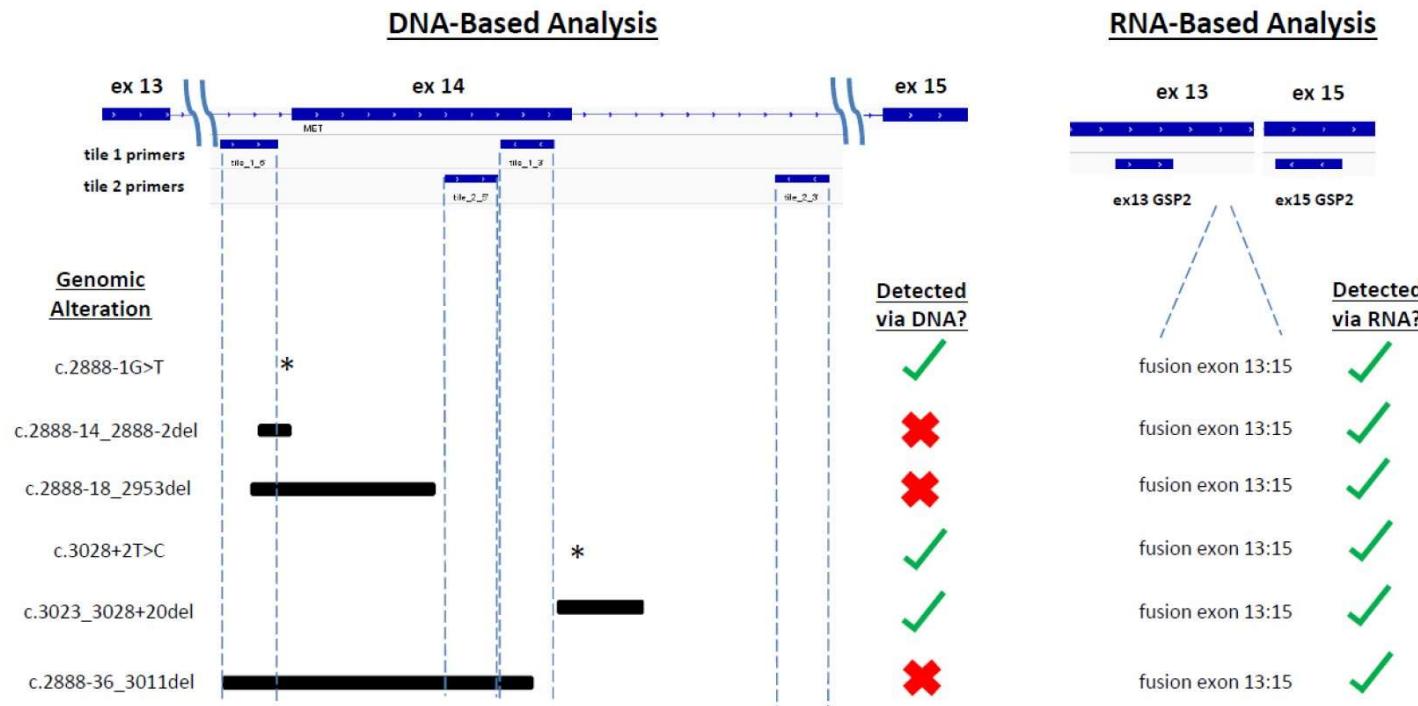
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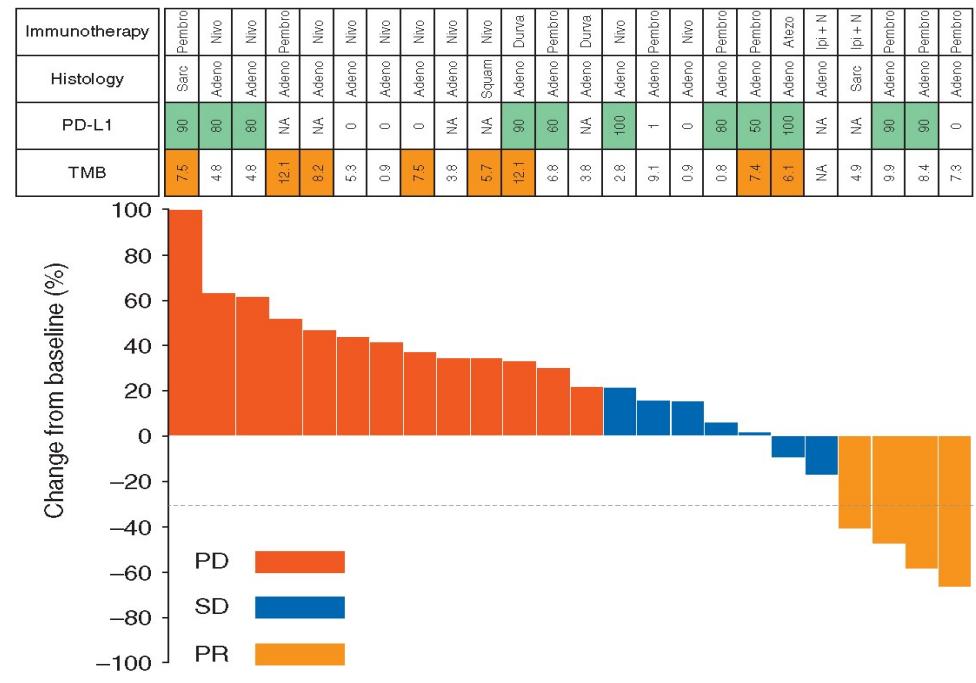
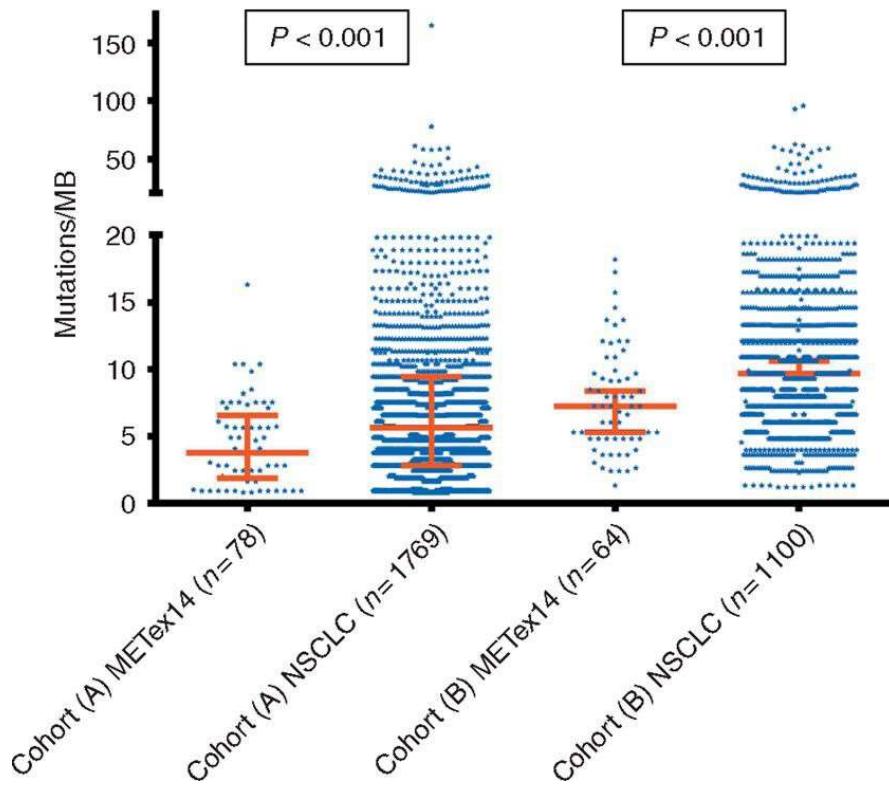
Drilon et al WCLC 2018, Wolf et al ASCO 2019, Paik et al ASCO 2019, Lu et al ASCO 2019



# NGS Testing is Key to Detection of MET Ex14 (Both DNA & RNA seq)



# IO in MET Ex 14 positive patients



ORR = 17%

Median PFS = 1.9 months

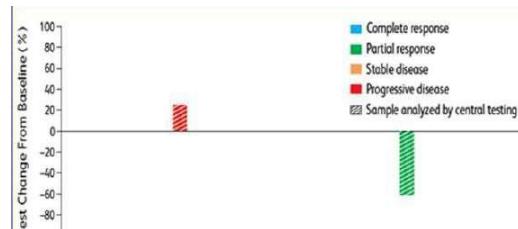
Sabari JK et al Ann Oncol 2018

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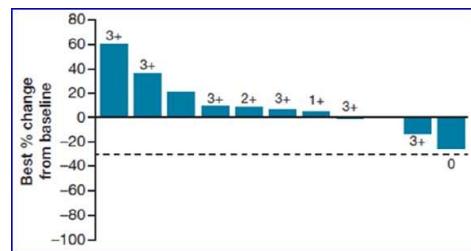
# What about *MET* copy amplified tumors?

Low MET (1.8-2.2), n = 3



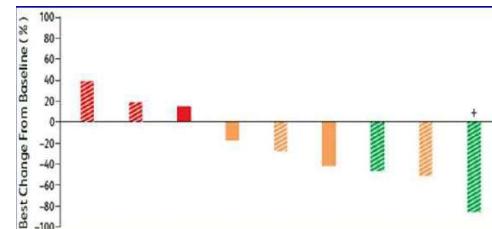
ORR 33.3% (0.8-90.6) PFS 1.8 (0.8-14)

GCN < 4, n = 17



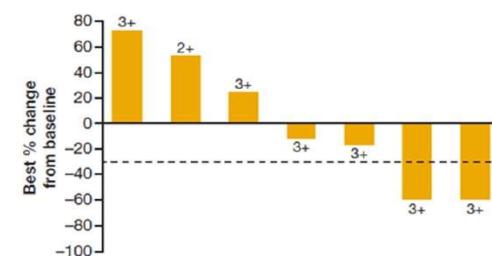
ORR 0% (0.8-90.6) PFS 1.8 (0.8-14)

Crizotinib  
Intermediate (>2.2-<4.0), n = 14



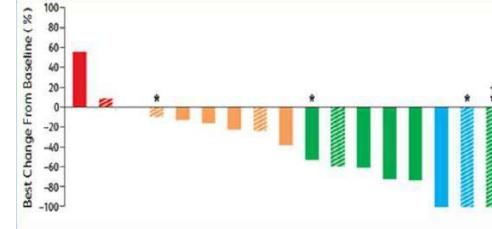
ORR 14.3% (1.8-42.8) PFS 1.9 (1.3-5.5) ORR 40% (19.1-63.9) PFS 6.7 (3.4-7.4)

Capmatinib  
GCN  $\geq$  4 & < 6. n = 12

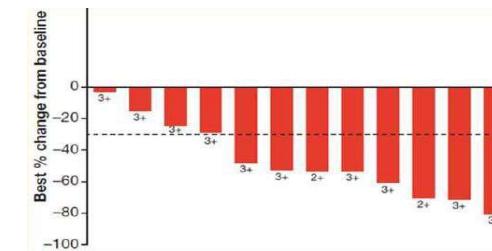


ORR 17.0% (2.1-48.4)

High  $\geq$  4.0, n = 20



GCN  $\geq$  6 n = 15

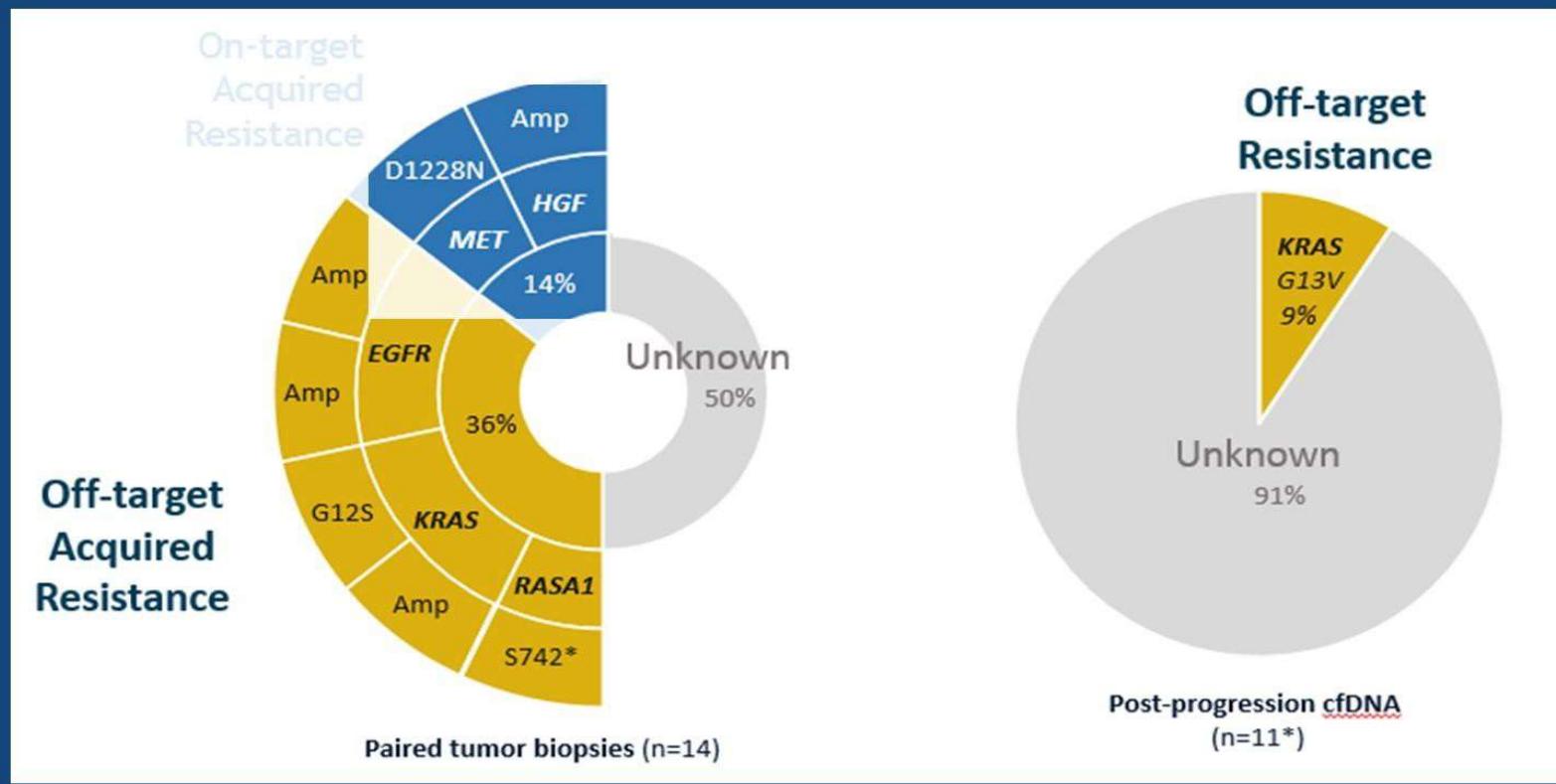


ORR 70% (21.3-73.4) PFS 1.8 (0.8-14)

Camidge ASCO 2018, Schuler ASCO 2016



# Acquired resistance involves on target and bypass pathways



Guo et al ASCO 2019

PRESENTED AT: **2019 ASCO<sup>®</sup>**  
ANNUAL MEETING

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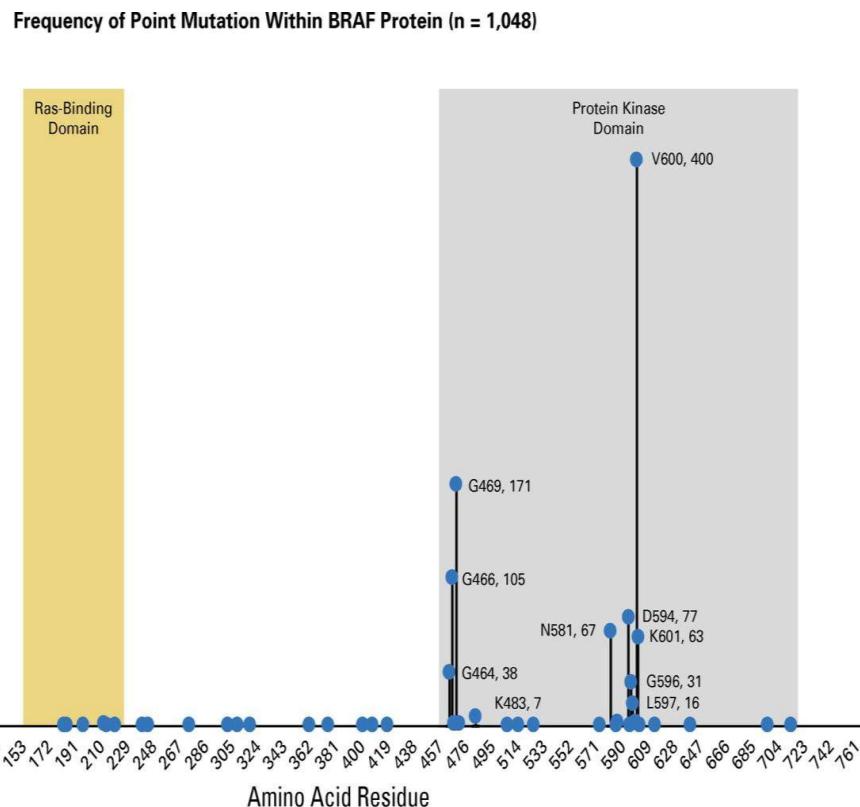
PRESENTED BY: Karen Reckamp, MD, MS

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15

# BRAF V600 mutations

A



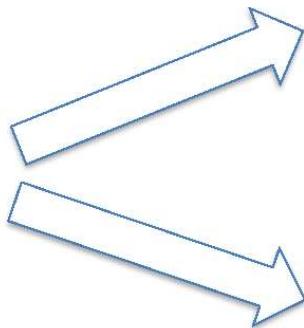
Sheikine et al J Clin Oncol 2019

- Serine-threonine kinase linking RAS GTPase to downstream MEK/ERK pathway.
- BRAF alterations incidence at 4.4%.
- BRAF V600E ~ 40% - Class I
- Median age 64 & predominantly adenocarcinoma (65%)
- Non V600\* or Class II & III generally considered to be non-actionable and ~60%.



# BRAF inhibition in Lung Cancer

Stage IV NSCLC  
BRAF V600E  
ECOG 0-2  
Line of Tx  $\geq$  2



Stage IV NSCLC  
BRAF V600E  
ECOG 0-2  
No prior Tx



Cohort A  
Dabrafenib 150mg BID  
N = 84



Dabrafenib in patients with BRAF<sup>V600E</sup>-positive advanced non-small-cell lung cancer: a single-arm, multicentre, open-label, phase 2 trial

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Cohort B  
Dabrafenib 150mg BID+  
Trametinib 2mg QD  
N = 59



Dabrafenib plus trametinib in patients with previously treated BRAF<sup>V600E</sup>-mutant metastatic non-small cell lung cancer: an open-label, multicentre phase 2 trial

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Cohort C  
Dabrafenib 150mg BID+  
Trametinib 2mg QD  
N = 36

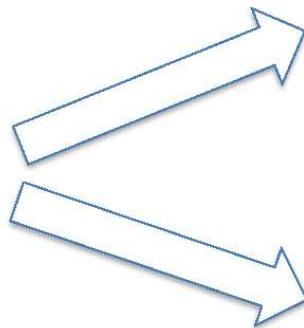
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ORR 33%, mPFS 5.5 months  
(Lancet Oncol 2016)



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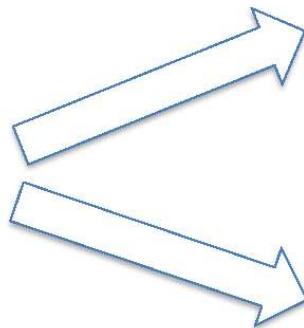
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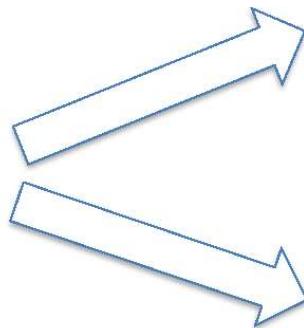
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# Non V600\* BRAF Kinase mutations

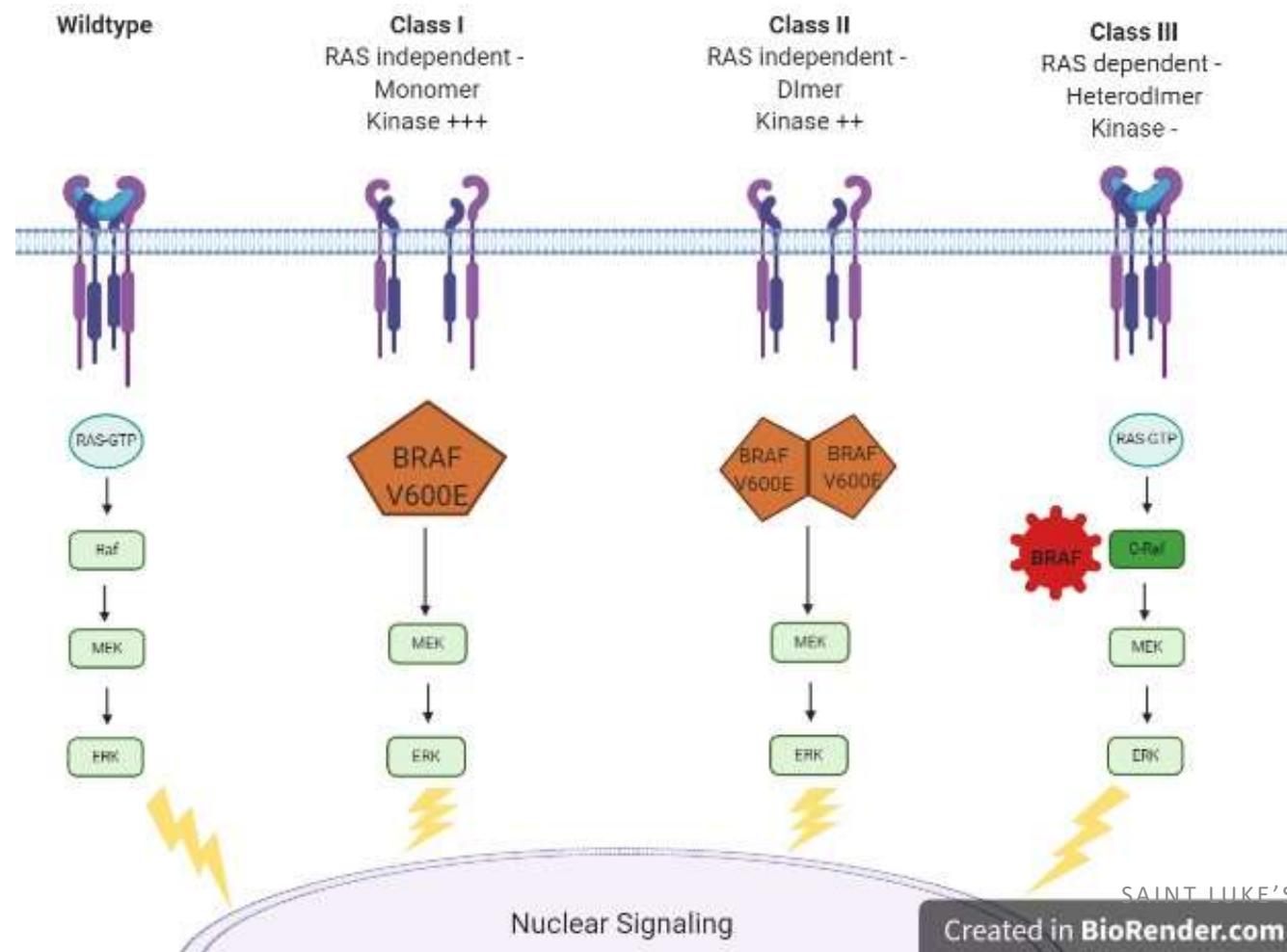
Class II Kinase Active	Class III Kinase Dead/Inactive
G464A, G464E, G464V	G466E, G466R, G466V
G466A	G469E
F468C	D594A, D594E, D594G, D594H, D594N, D594V, D594Y
G469A, G469R, G469S, G469V	G596R
N581S	T599I
E586K	
F595L	
L597Q, L597R, L597S, L597V	
K601E	

- Class II and III mutations constitute more than half of all patients with BRAF mutations.
- 2<sup>nd</sup> generation RAF kinase inhibitors are ineffective against both class II and III BRAF mutations.

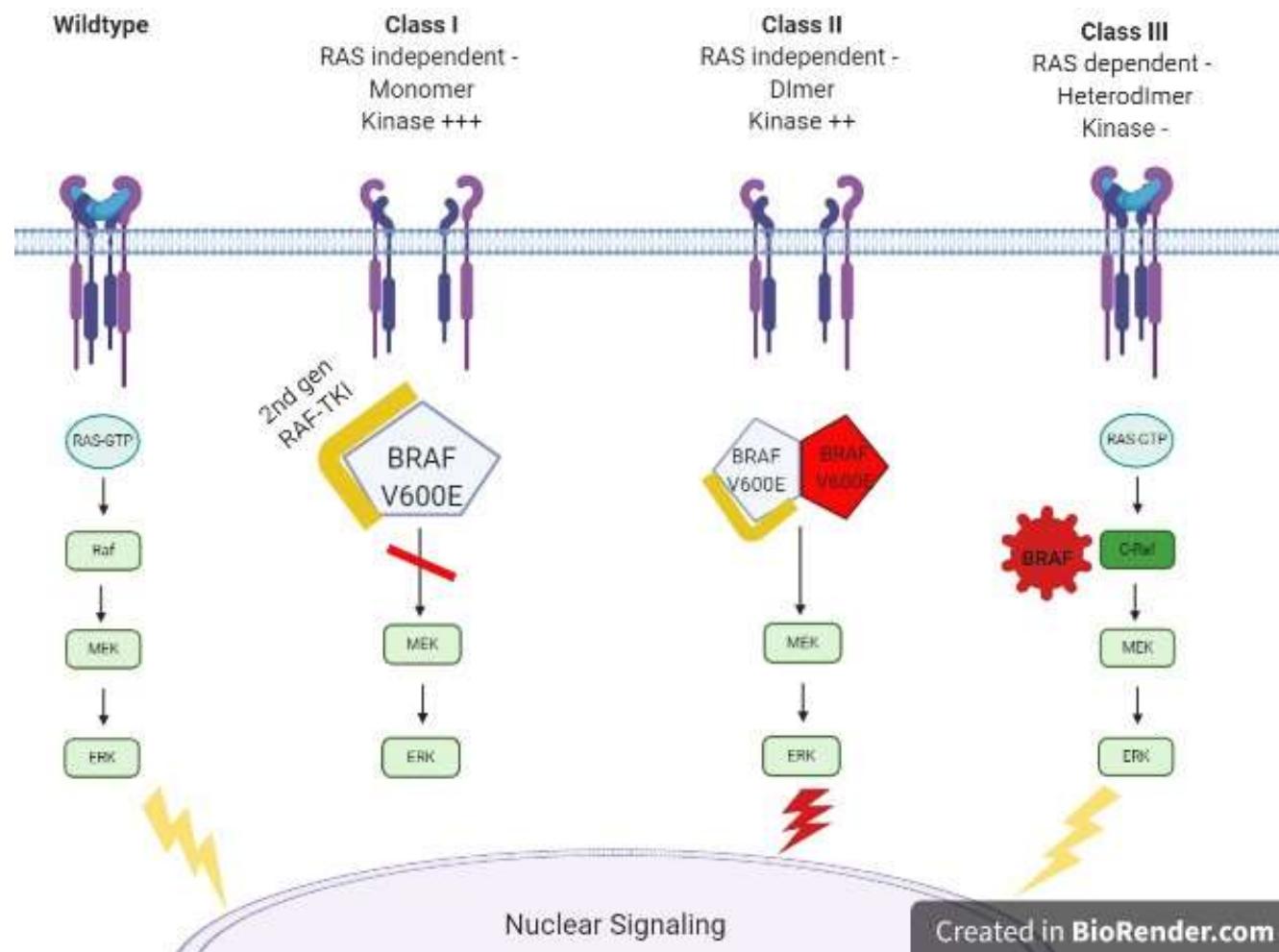
Sheikine et al J Clin Oncol 2019, Yao et al Cancer Cell 2015, Gautschi et al JTO 2015, Mazieres et al JTO 2019



# BRAF Kinase Inhibition

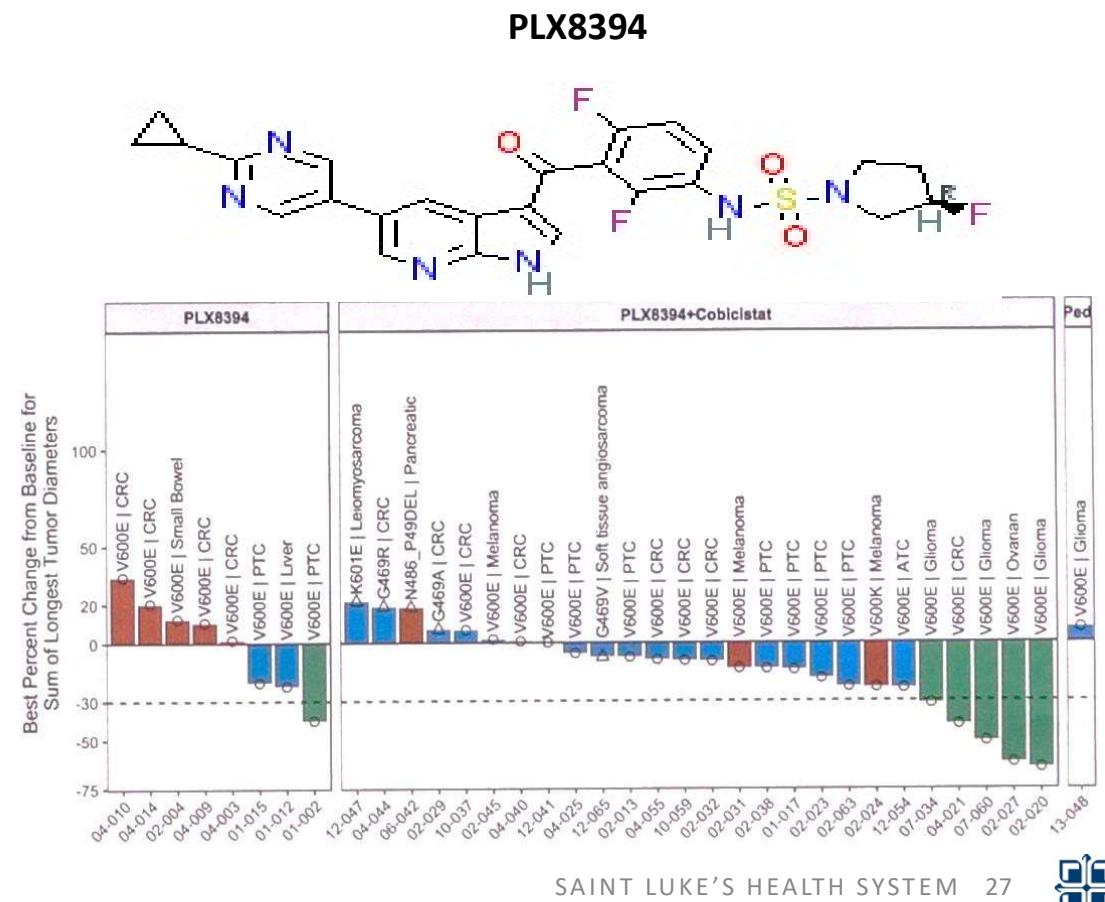


# BRAF Kinase Inhibition



# Targeting non-V600 mutations, 3<sup>rd</sup> gen RAF TKI

- 2<sup>nd</sup> gen RAF TKIs ineffective against BRAF homo & heterodimers.
- Plus risk for paradoxical ERK activation
- 3<sup>rd</sup> gen RAF TKIs avoid paradoxical activation and active against BRAF dimers.



# Way Forward in V600\* population – Triple therapy?

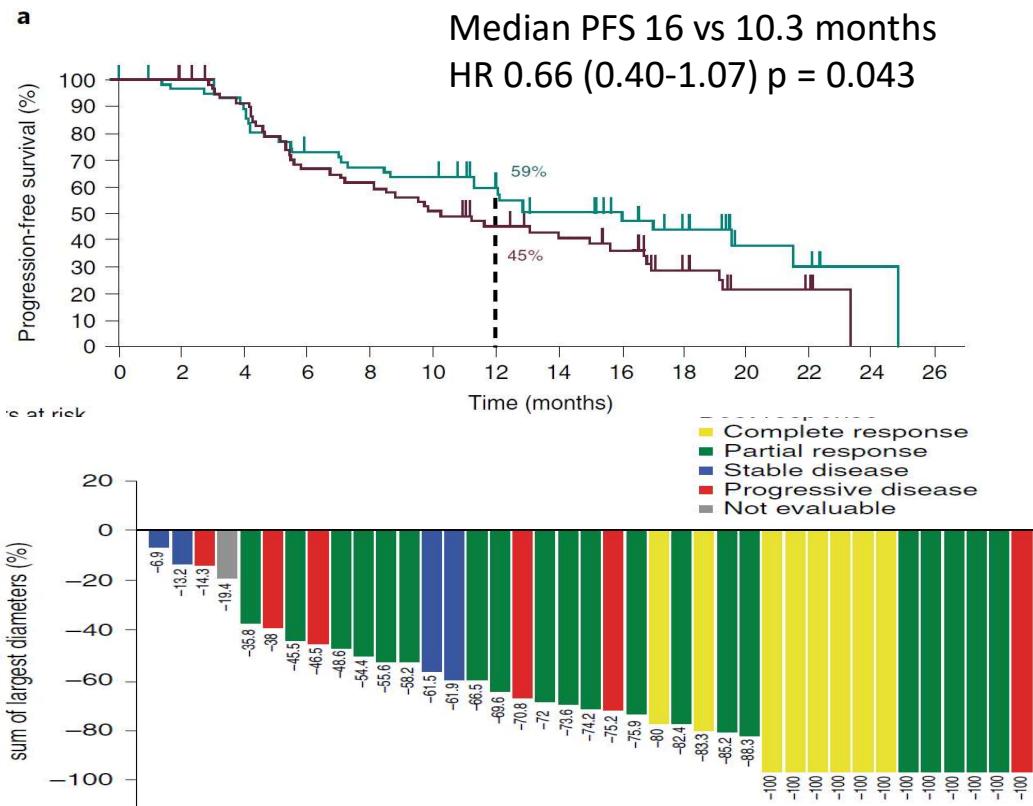
nature  
medicine

LETTERS

<https://doi.org/10.1038/s41591-019-0448-9>

## Dabrafenib, trametinib and pembrolizumab or placebo in *BRAF*-mutant melanoma

Paolo Antonio Ascierto<sup>1,18\*</sup>, Pier Francesco Ferrucci<sup>2,18\*</sup>, Rosalie Fisher<sup>3</sup>, Michele Del Vecchio<sup>4</sup>,



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LETTERS

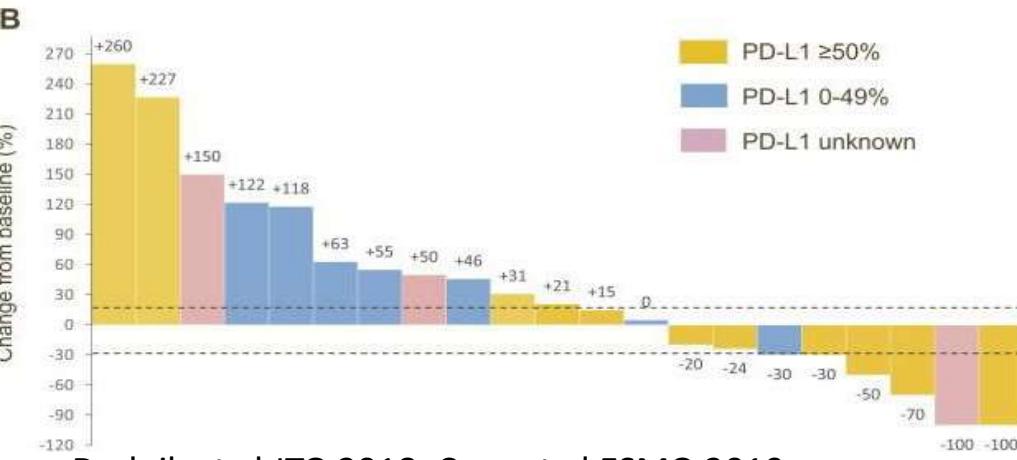
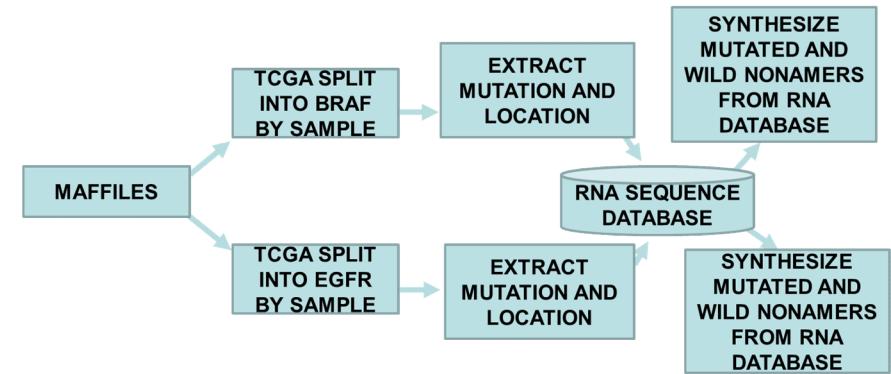
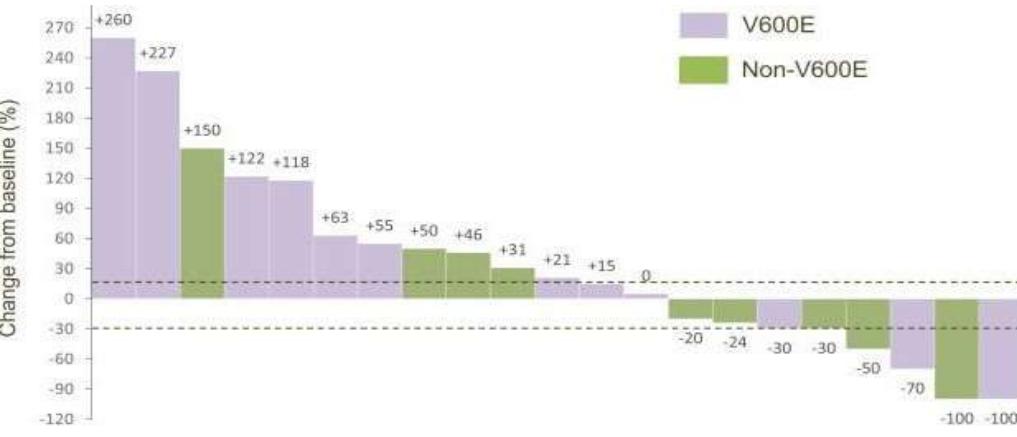
<https://doi.org/10.1038/s41591-019-0474-7>

## Atezolizumab plus cobimetinib and vemurafenib in *BRAF*-mutated melanoma patients

Ryan J. Sullivan<sup>1,2\*</sup>, Omid Hamid<sup>2</sup>, Rene Gonzalez<sup>3</sup>, Jeffrey R. Infante<sup>4</sup>,



# IO in BRAF V600\* lung cancer

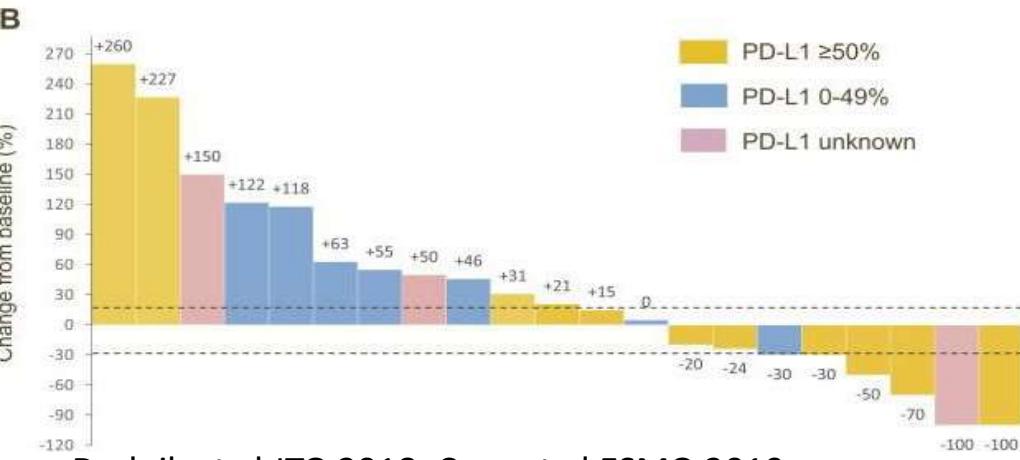
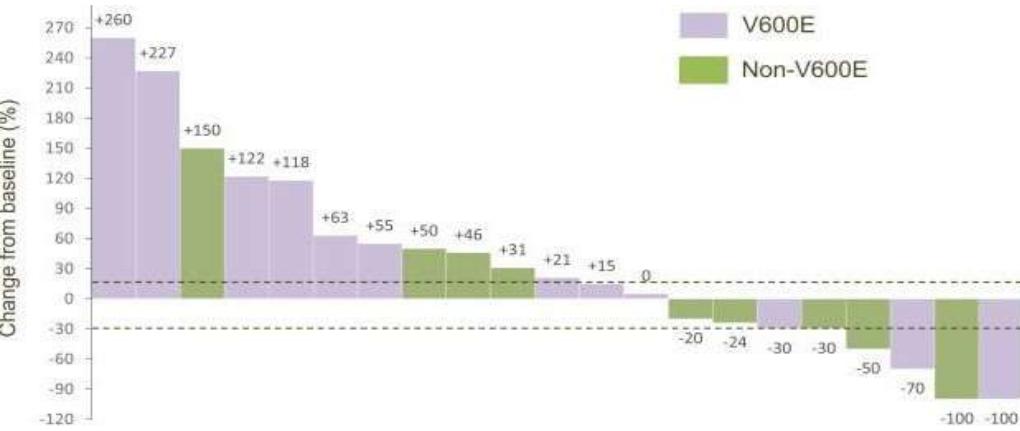


	Total	group		p-Value	Std Diff (%)
		braf n = 35	egfr n = 68		
Mutation Burden	158.0 (75.0, 488.0)	445.0 (165.0, 776.0)	90.5 (60.5, 219.5)	0.001	64.7
Number Affinities	3173.5 (1531.0, 9914.5)	9536.5 (3839.5, 14626.0)	1895.5 (1148.5, 4766.5)	< 0.001	75.3

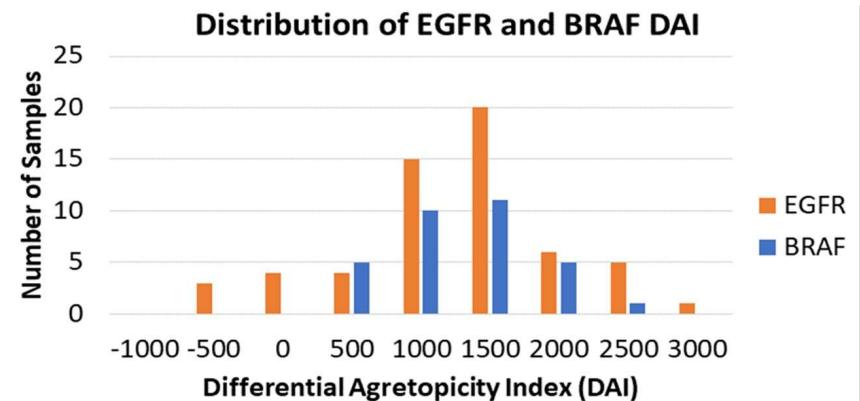
Dudnik et al JTO 2018, Case et al ESMO 2019



# IO in BRAF V600\* lung cancer



Dudnik et al JTO 2018, Case et al ESMO 2019

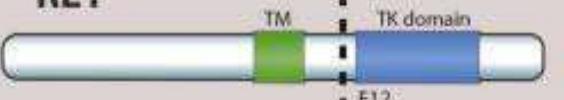


	Total n = 103	group		p-Value	Std Diff (%)
		braf n = 35	egfr n = 68		
Mutation Burden	158.0 (75.0, 488.0)	445.0 (165.0, 776.0)	90.5 (60.5, 219.5)	0.001	64.7
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# RET fusions

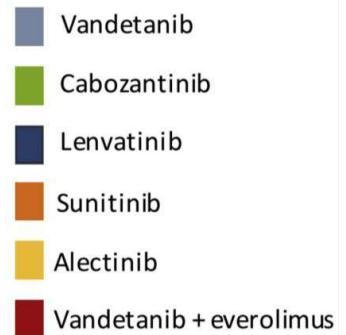
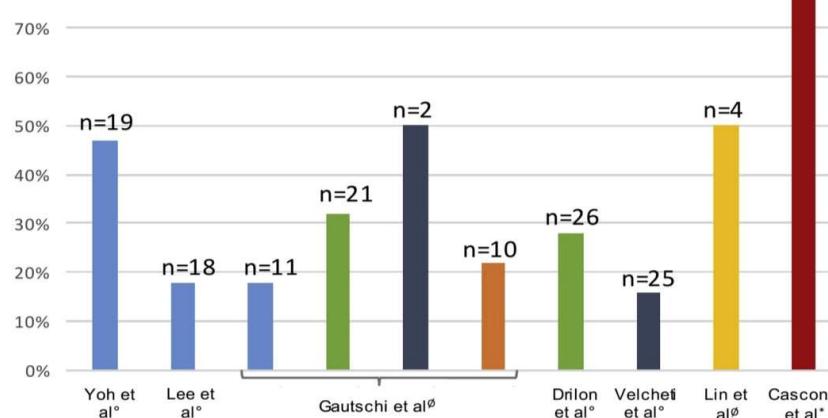
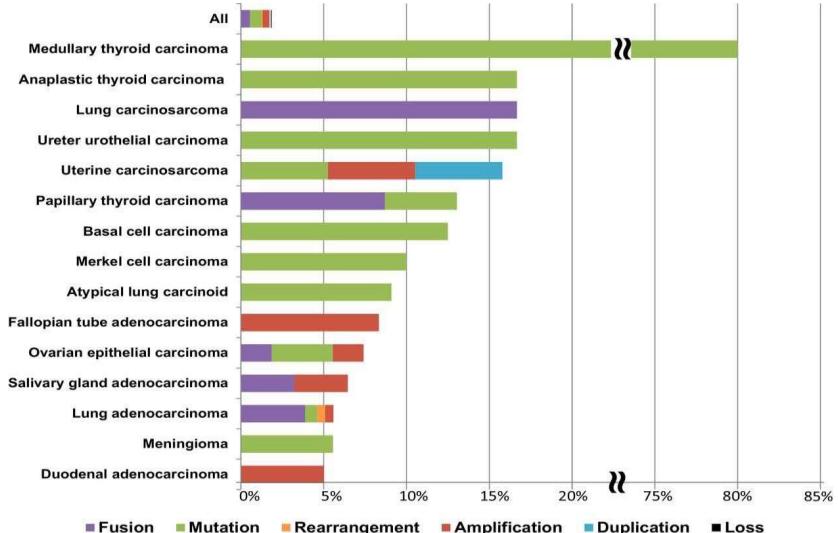
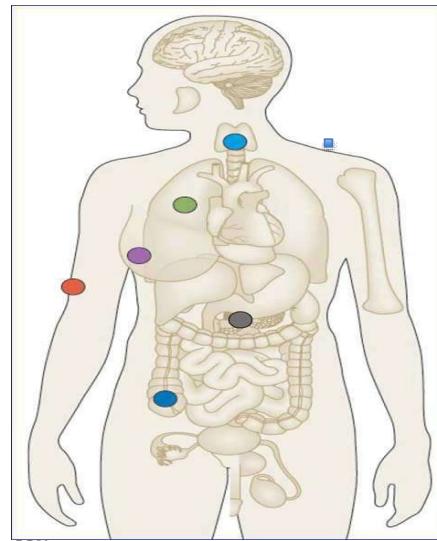
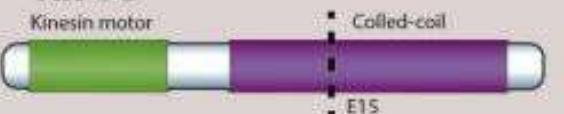
**RET**



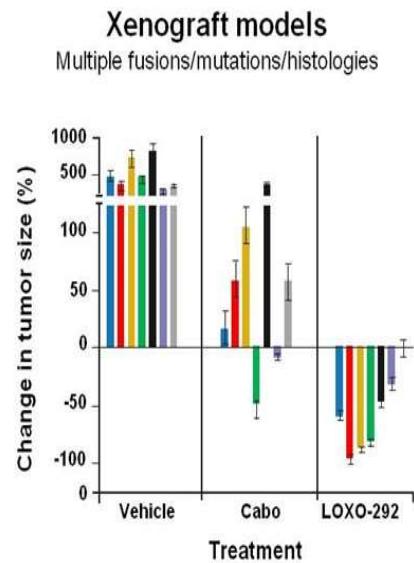
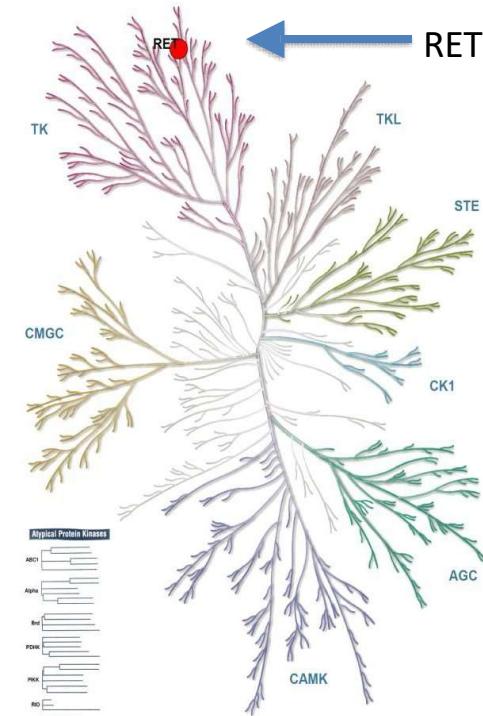
**KIF5B-RET**



**KIF5B**



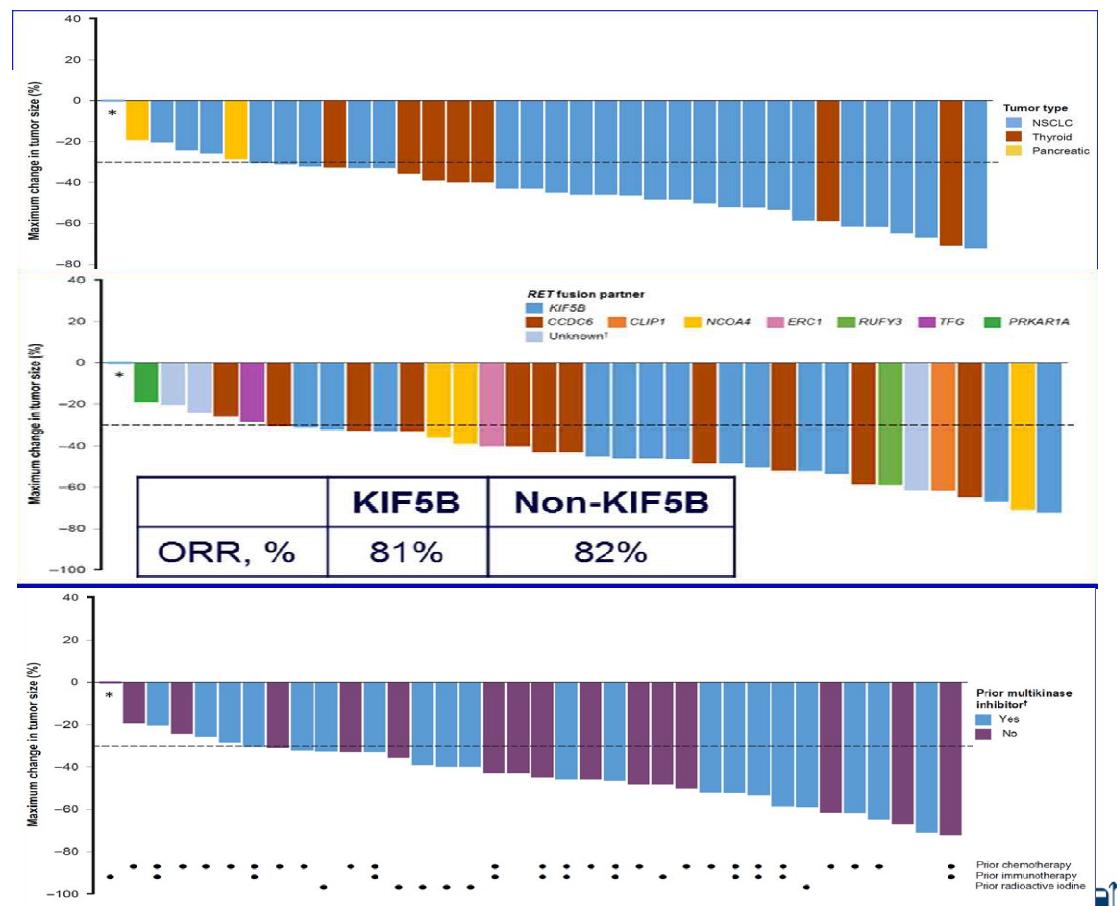
# LOXO-292 (Selpercatinib)



**Tumor models**

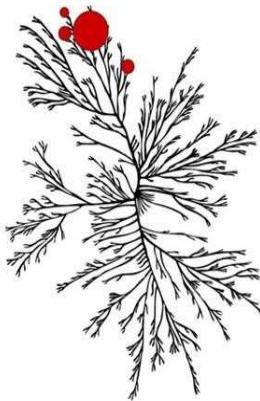
- KIF5B-RET (PDX-NSCLC)
- CCDC6-RET (PDX-CRCA)
- CCDC6-RET-V804M (PDX-CRCA)
- KIF5B-RET (NIH-3T3)
- KIF5B-RET-V804M (NIH-3T3)
- RET C634W (TT cell line-MTC)
- CCDC6-RET (LC-2/ad cell line-NSCLC)

Drilon ASCO 2018



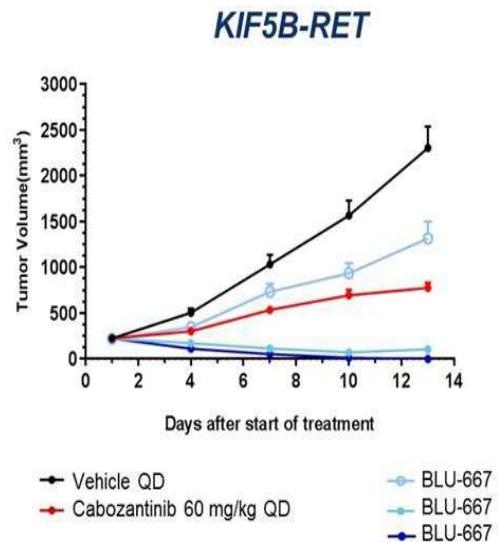
# BLU-667

## BLU-667: High kinase selectivity for RET<sup>a</sup>

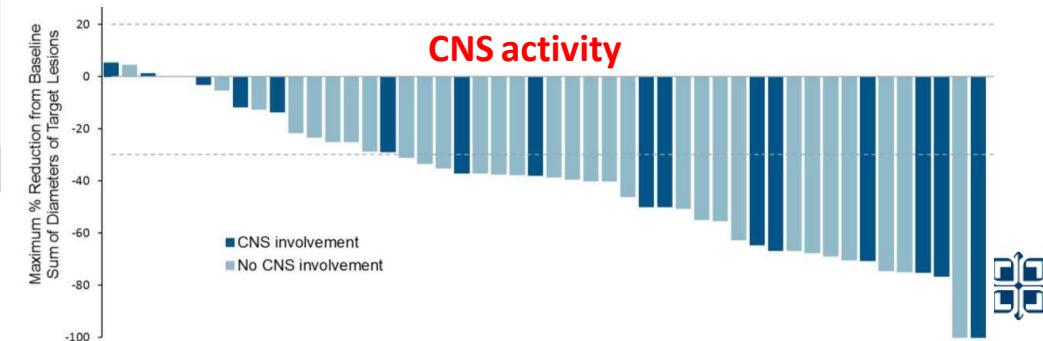
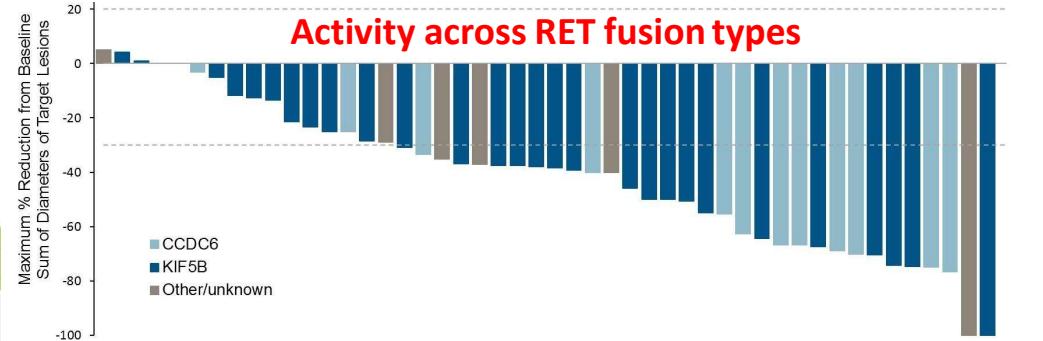


BLU-667 vs. pharmacologically relevant kinases:

- ~90-fold more selective for RET than VEGFR2
- 20-fold more selective for RET than JAK1



Best Response	All (N=48)	Prior Platinum (N=35)
ORR (95% CI)	58% (43–72)	60% (42–76)
CR*	1	1
PR*	27	20
SD	18	14
PD	2	-
DCR (95% CI)	96% (86–99)	100% (90–100)



Gainor et al ASCO 2019



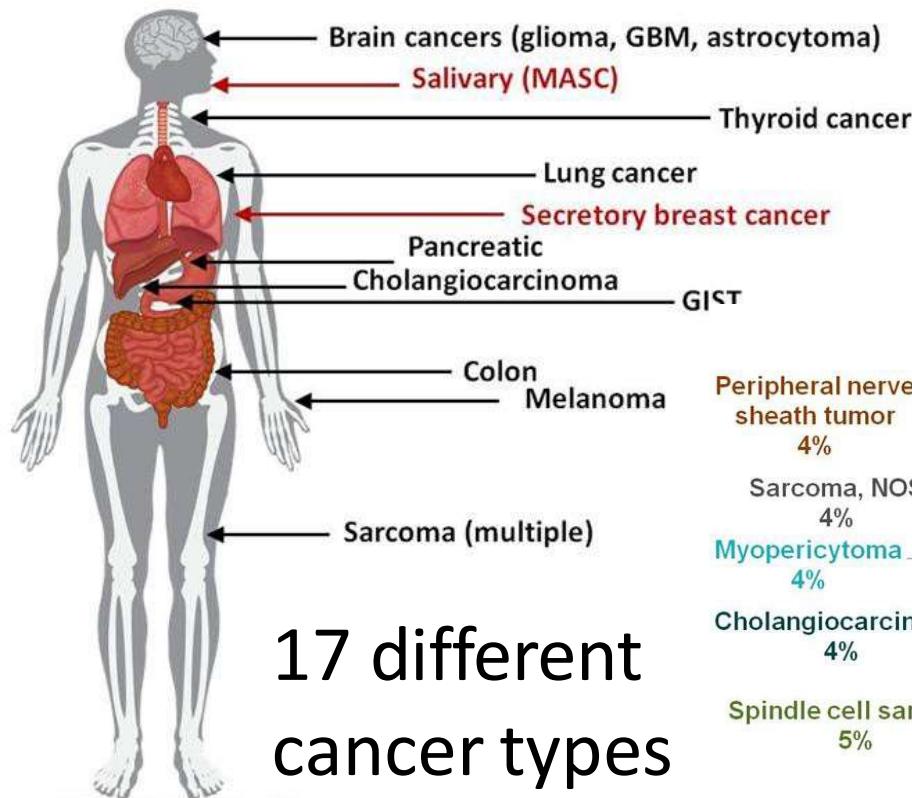
# Safety Profile LOXO-292 & BLU-667

All Grade (Grade $\geq 3$ ) %	Treatment emergent		Treatment related	
	LOXO-292 ( $\geq 10\%$ ) N = 82	BLU-667 ( $\geq 15\%$ ) N = 120	LOXO-292	BLU-667
Constipation		30 (2)	2 (-)	17 (2)
Fatigue	20 (-)	21 (3)	13 (-)	13 (3)
Diarrhea	16 (-)	18 (2)	2 (-)	9 (-)
Dry mouth	12 (-)	17 (-)	6 (-)	12 (-)
Nausea	12 (-)	-	5 (-)	-
Dyspnea	11 (1)	-	-	-
Neutropenia	-	26 (3)	-	26 (13)
Anemia	-	18 (7)	-	11 (4)
AST increased	-	24 (5)	-	20 (2)
Hypertension	-	20 (13)	-	13 (10)
ALT increased	-	17%	-	13 (2)

Gainor et al ASCO 2019, Drilon ASCO 2018

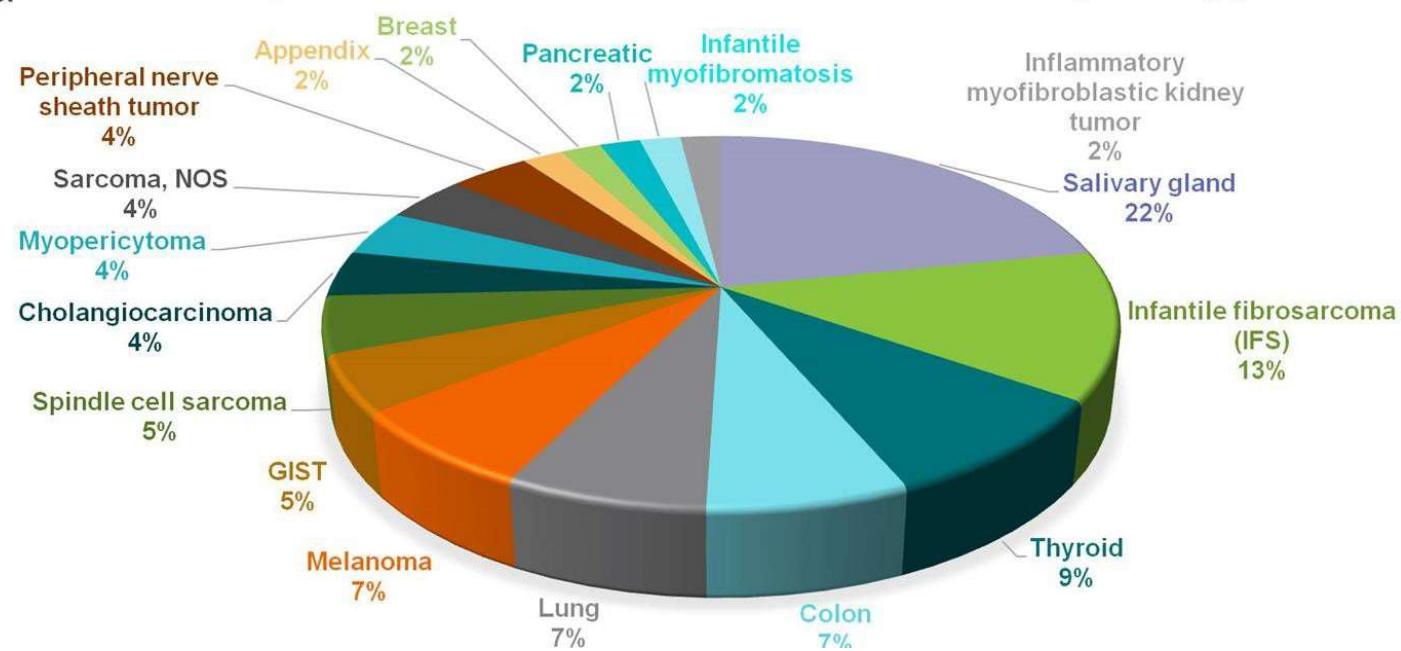


# NTRK fusions in cancer



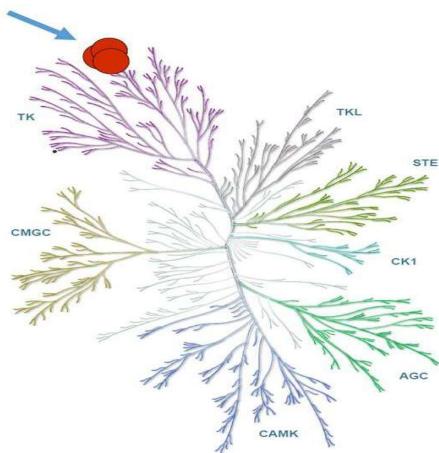
17 different cancer types

Estimated 1500 – 5000 patients with cancer harbor TRK fusions annually in the United States



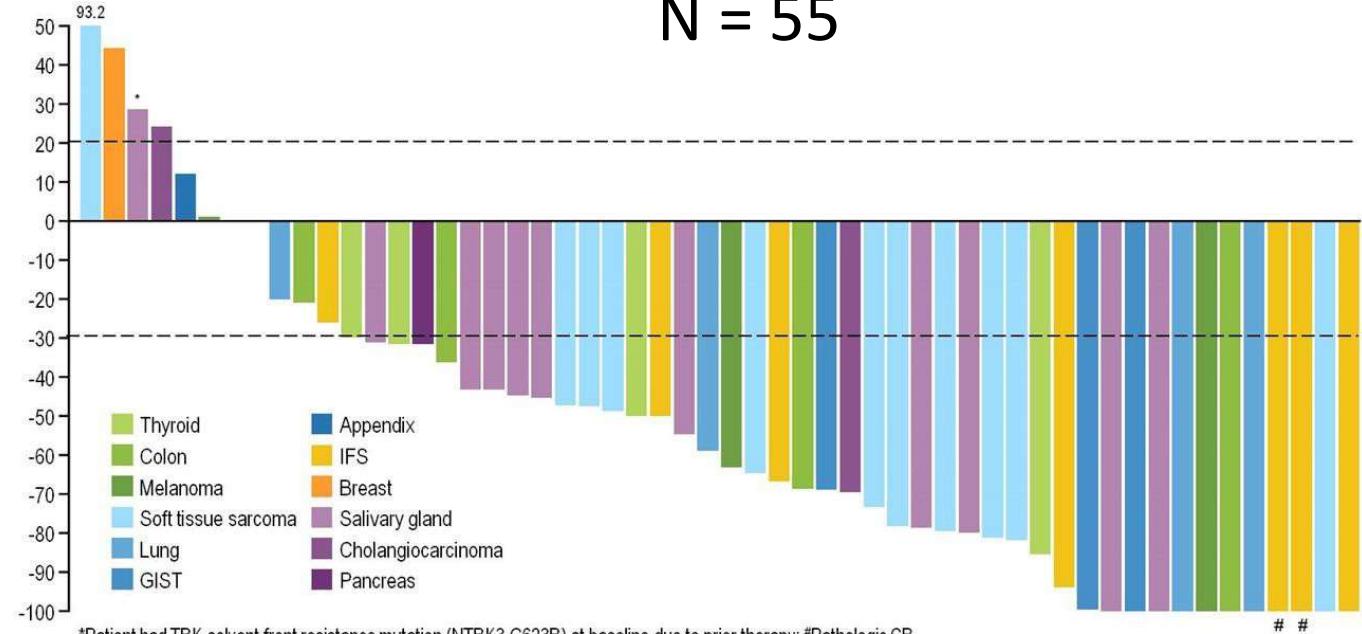
# Larotrectinib (LOXO-101), highly selective TRK inhibitor

TRKA/B/C



- Highly selective NTRK inhibitor
- Development program ( $n = 55$ )
  - Adult phase I ( $n = 8$ )
  - SCOUT pediatric phase II ( $n = 12$ )
  - NAVIGATE adult/adolescent phase II basket trial ( $n = 35$ )

Effective irrespective of tumor type  
 $N = 55$



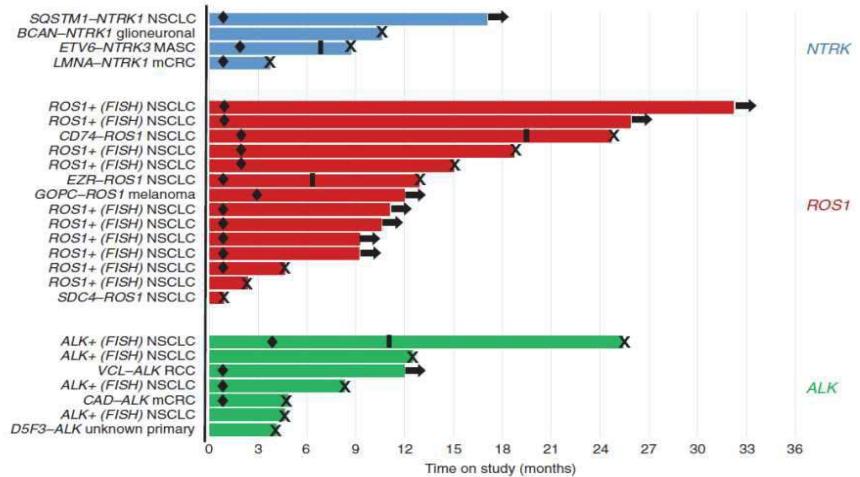
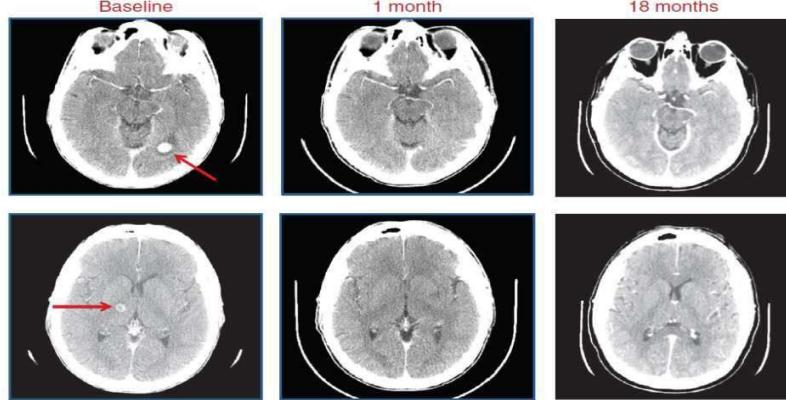
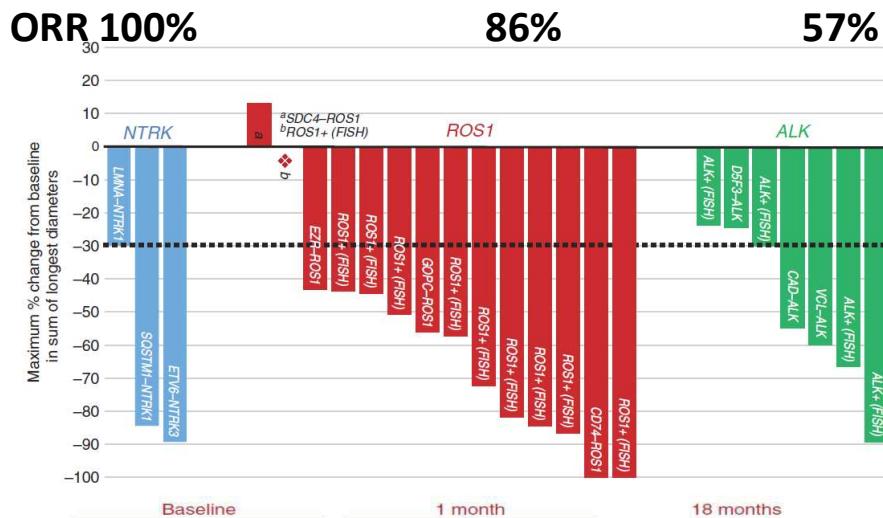
\*Patient had TRK solvent front resistance mutation (NTRK3 G623R) at baseline due to prior therapy; #Pathologic CR

Note: One patient not shown here. Patient experienced clinical progression and no post-baseline tumor measurements were recorded.

Drilon NEJM 2018



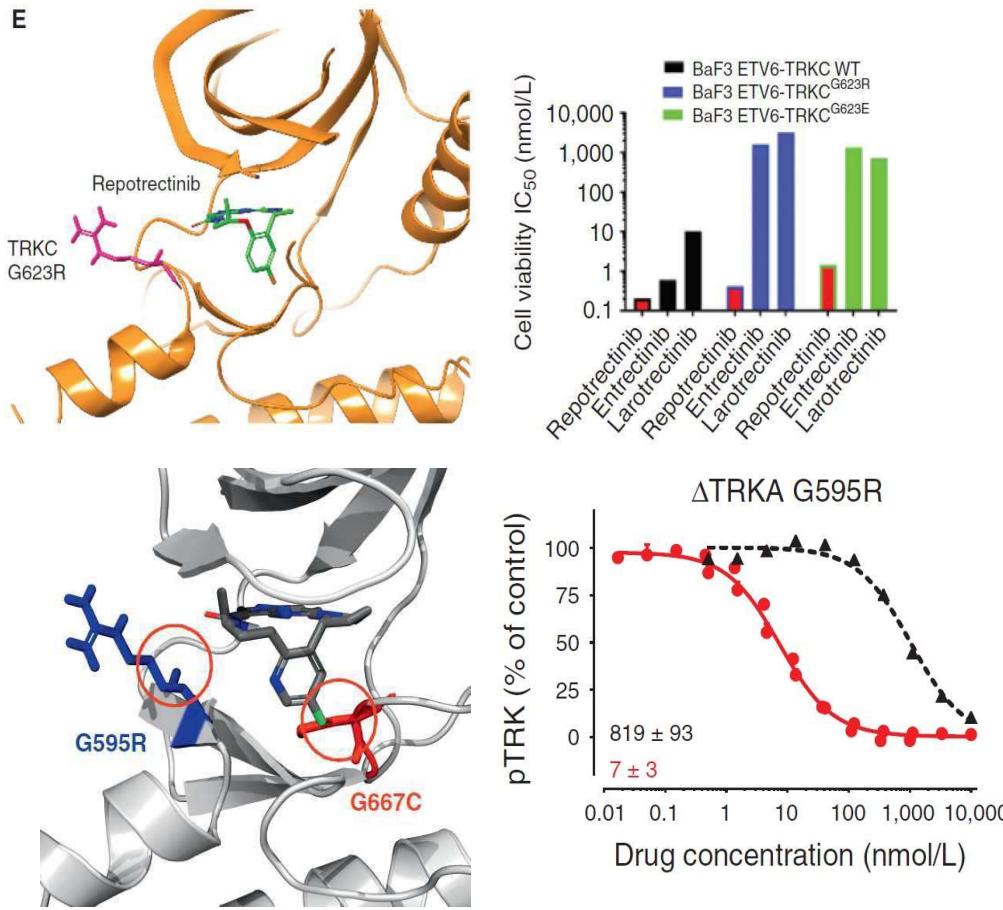
# Entrectinib (RXDX-101), Pan-TRK, ALK & ROS1 inhibitor



- N = 25, ALKA-372-001 & STARTRK-1 trials
- TKI naïve
- Intracranial response rate 63%



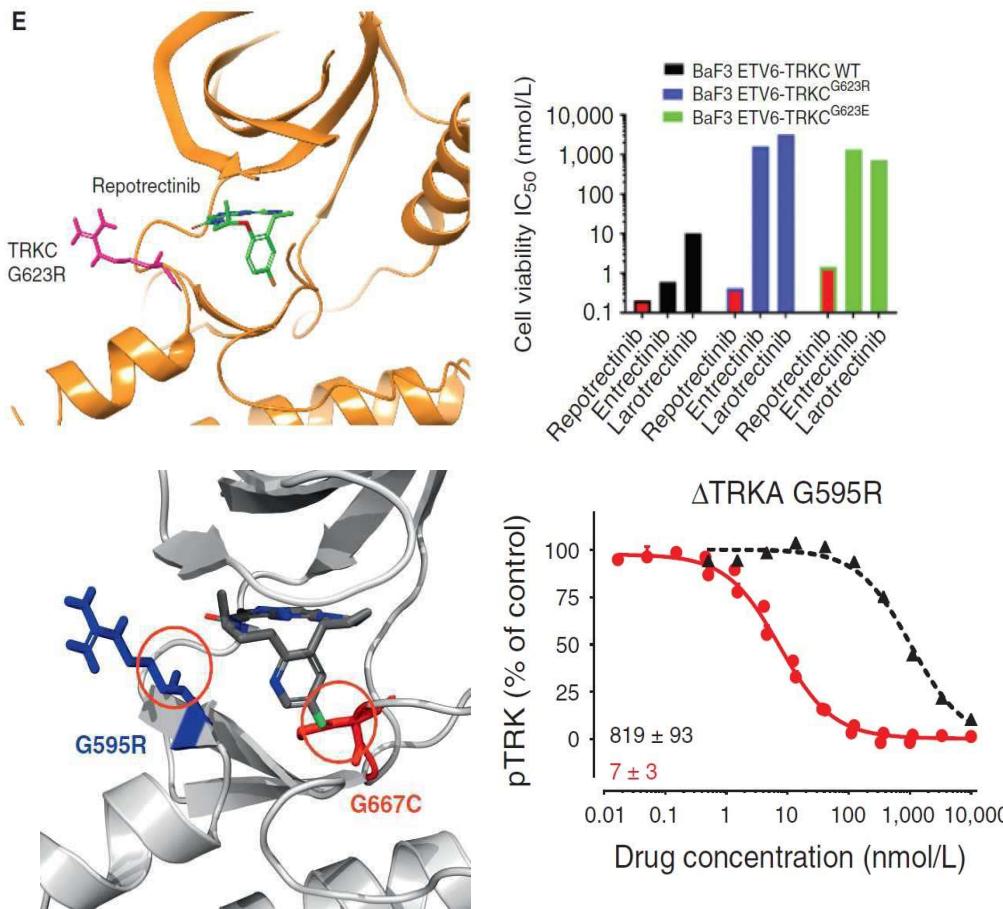
# Resistance and potential 2<sup>nd</sup> line options



- Kinase domain mutations
  - Solvent front
    - TRKA G595R, TRKC G623R
  - Gatekeeper
    - TRKA F589L
  - Activation loop
    - TRKA G667S, TRKC G696A



# Resistance and potential 2<sup>nd</sup> line options



- Kinase domain mutations
  - Solvent front
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# Current Landscape & the Way Forward

- MET
  - Specific TKIs to target MET ex14 are expected to be approved in the near future (capmatinib & tepotinib).
  - NGS testing key to detectiong MET ex14.
  - Is MET amplification a target?
  - Met protein expression by IHC not a therapeutic biomarker.
- RET
  - LOXO-292 & BLU-667 highly selective and effective RET inhibitors.
  - Is LOXO-292 (selpercatinib) better tolerated?



# Current Landscape & the Way Forward

- BRAF
  - Established target- need to build on existing treatment paradigms
    - Rational combinations with IO and other small molecule inhibitors ? ERK.
  - 3<sup>rd</sup> generation RAF TKIs needed to target non V600 BRAF and limit effects of paradoxic activation.
- NTRK
  - Tumor and age agnostic target.
  - Highly effective inhibitors available (Larotrectinib & Entrectinib).



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# Thank You

- Questions?

