VIRTUAL MOLECULAR TUMOR BOARD Optimizing Biomarker-Based Decision-Making for Patients with Non-Small Cell Lung Cancer with EGFR Mutations or with Other Oncogene-Addicted Lung Cancers

A 2-Part CME/MOC-Accredited Webinar Series

Tuesday, October 26, 2021 5:00 PM - 6:00 PM ET

Faculty

Marc Ladanyi, MD Andrew J McKenzie, PhD Joel W Neal, MD, PhD



Faculty



Marc Ladanyi, MD
Chief, Molecular Diagnostics Service
William J Ruane Chair in Molecular Oncology
Memorial Sloan Kettering Cancer Center
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Research To Practice
Miami, Florida



Commercial Support

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Dr Love — Disclosures

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Dr McKenzie — Disclosures

No relevant conflicts of interest to disclose.



Dr Neal — Disclosures

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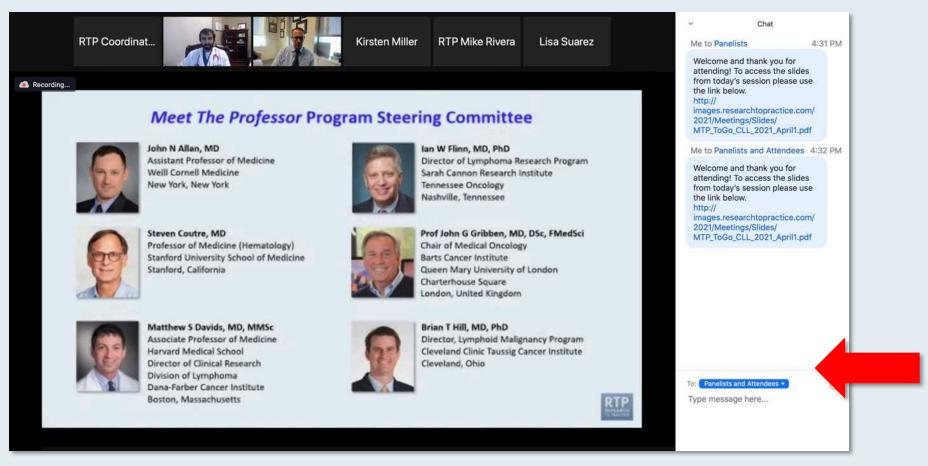


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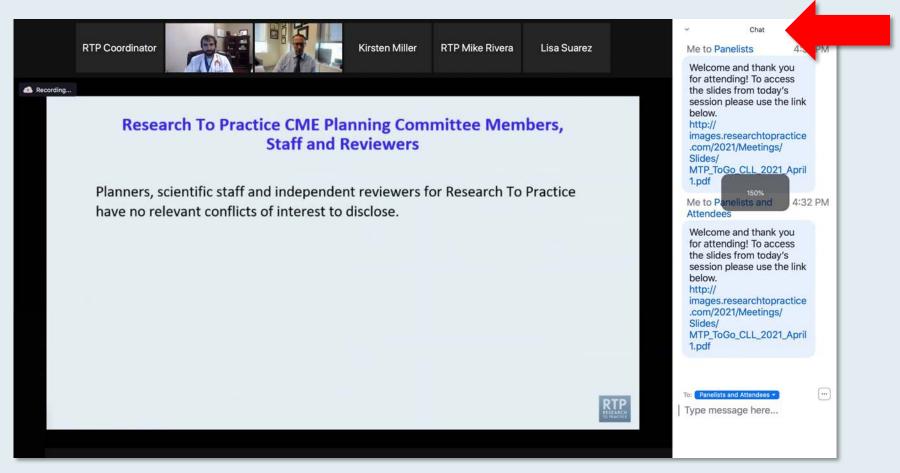


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Press Command (for Mac) or Control (for PC) and the + symbol. You may do this as many times as you need for readability.



ONCOLOGY TODAY

WITH DR NEIL LOVE

Key Presentations on Lung Cancer from the 2021 ASCO Annual Meeting



DR JOEL NEAL STANFORD UNIVERSITY









Meet The Professor Optimizing the Clinical Management of Hodgkin and Non-Hodgkin Lymphomas

Wednesday, October 27, 2021 5:00 PM - 6:00 PM ET

Faculty

Jonathan W Friedberg, MD, MMSc



Optimizing the Selection and Sequencing of Therapy for Patients with ER-Positive Breast Cancer

Thursday, October 28, 2021 5:00 PM - 6:00 PM ET

Faculty
Matthew P Goetz, MD



Meet The ProfessorManagement of BRAF-Mutant Melanoma

Monday, November 1, 2021 5:00 PM – 6:00 PM ET

Faculty
Prof Georgina Long, AO, BSc, PhD, MBBS



Optimizing the Selection and Sequencing of Therapy for Patients with Urothelial Bladder Carcinoma

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Faculty
Adam M Brufsky, MD, PhD



Key Considerations in the Optimal Clinical Care of Patients with Small Cell Lung Cancer

Thursday, November 4, 2021 5:00 PM - 6:00 PM ET

Faculty

Anne Chiang, MD, PhD
David R Spigel, MD



Meet The ProfessorOptimizing the Management of Acute Myeloid Leukemia

Monday, November 8, 2021 5:00 PM - 6:00 PM ET

Faculty
Keith W Pratz, MD



Optimizing the Management of Metastatic Castration-Resistant Prostate Cancer

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Simon Chowdhury, MD, PhD



Optimizing Biomarker-Based Decision-Making for Patients with Non-Small Cell Lung Cancer with EGFR Mutations or with Other Oncogene-Addicted Lung Cancers

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Thank you for joining us!

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Agenda

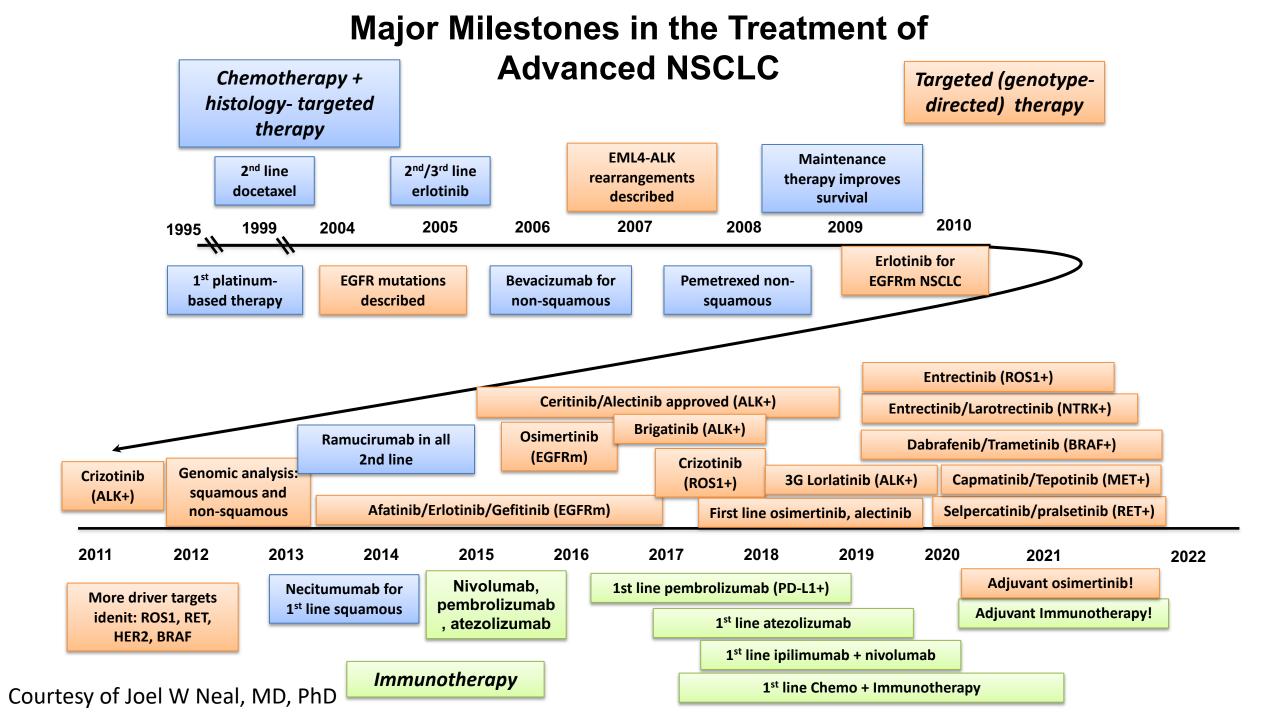
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 14 skipping and MET amplification mutations PD-L1 TPS 95%
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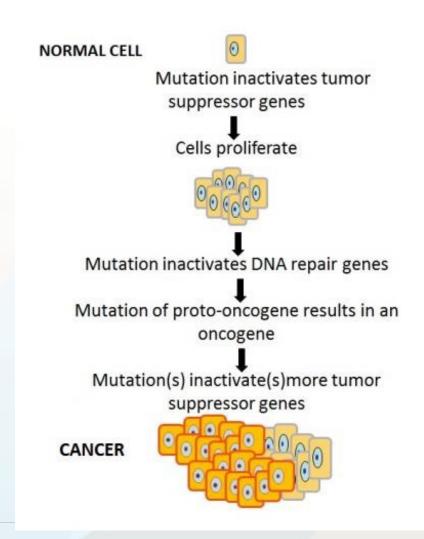




Cancer is a Disease of the Genome

Tumorigenesis (aka carcinogenesis)

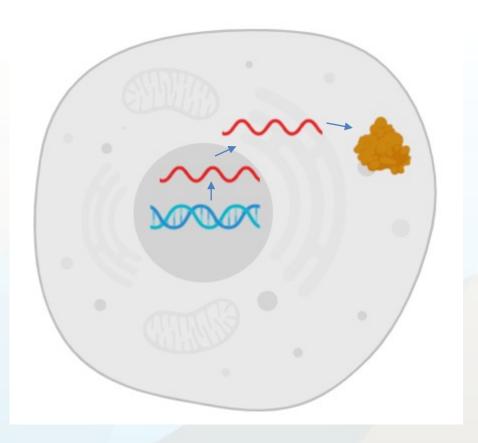
- Every type of cancer starts from a single cell (clone) with genetic mutations that confer a growth advantage over other cells (mutated cells divide and grow rapidly).
- Cancer causing mutations can occur in various classes of genes involved in cell growth and DNA repair including:
 - Oncogenes
 - Tumor Suppressor Genes
 - DNA Repair Genes
 - Cell cycle checkpoint genes
 - Cell death genes
 - Cell growth genes
 - Cellular differentiation genes
 - Cellular senescence genes
 - Metastasis/invasion genes



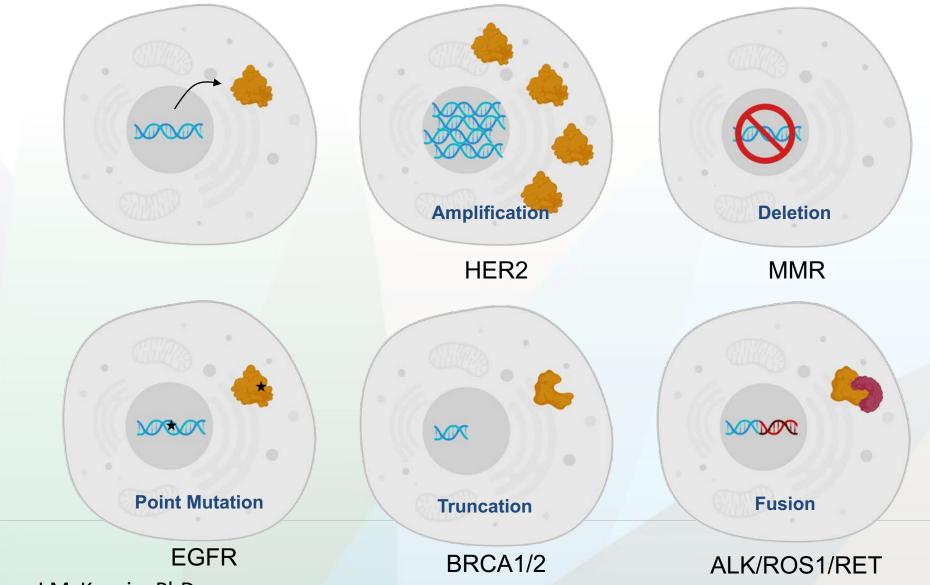
DNA, RNA, and Protein

THE CENTRAL DOGMA

• DNA Carries instructions for how to make Protein , the molecular workhorse of the cell, by way of an RNA intermediate

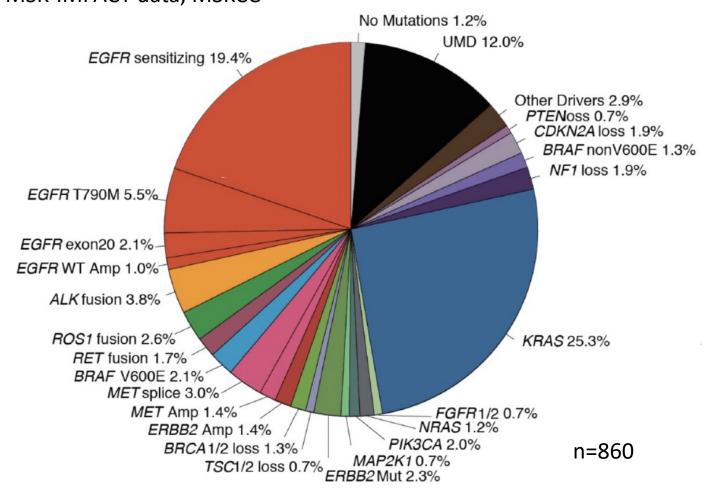


Types of Genetic Alterations (aka: mutations)



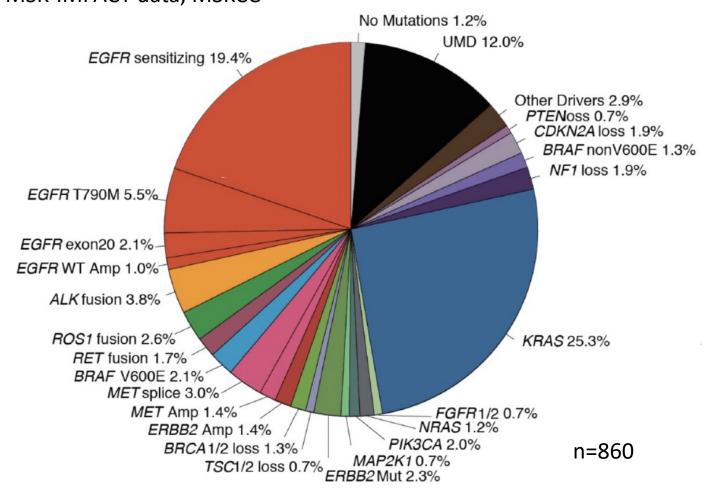
Courtesy of Andrew J McKenzie, PhD

MSK-IMPACT data, MSKCC



EGFR sensitizing 19.4% EGFR T790M 5.5% EGFR exon20 2.1% EGFR WT Amp 1.0%

MSK-IMPACT data, MSKCC

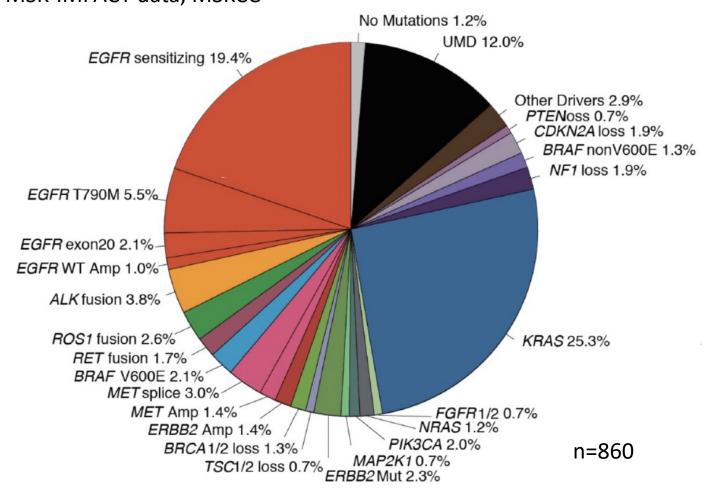


KRAS 25.3%

UMD 12.0%

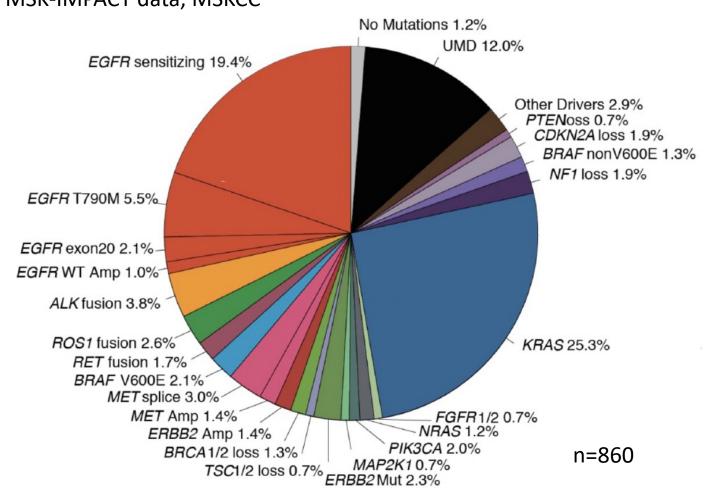
No Mutations 1.2%

MSK-IMPACT data, MSKCC



ALK fusion 3.8%
ROS1 Fusion 2.6%
RET Fusion 1.7%
BRAF V600E 2.1%
MET Splice 3.0%
MET Amp 1.4%
ERBB2 Amp 1.4%
ERBB2 Mut 2.3%

MSK-IMPACT data, MSKCC



BRCA 1/2 loss 1.3%

TSC 1/2 loss 0.7%

MAP2K1 0.7%

PIK3CA 2.0%

NRAS 1.2%

FGFR 1/2 0.7%

NF1 loss 1.9%

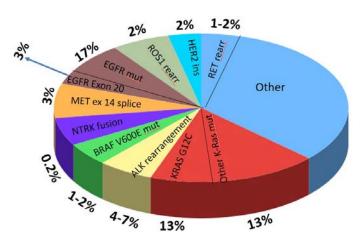
BRAF nonV600E 1.3%

CDKN2A loss 1.9%

PTEN loss 0.7%

Other drivers 2.9%

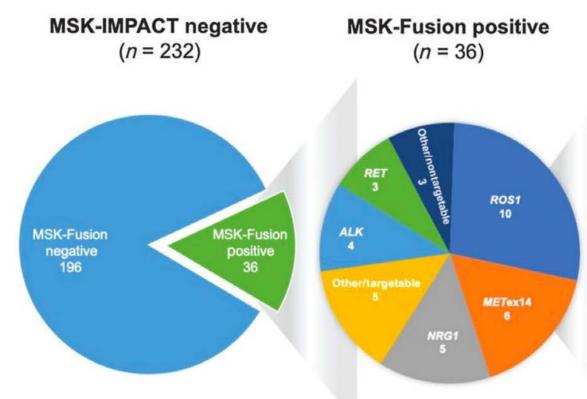
NSCLC: How to test for targets



Target	Direct/ hotspot seq	FISH testing	IHC staining	NGS DNA tumor seq	NGS plasma seq	NGS RNA tumor seq
KRAS mutations	++	=	-	++	+	++
EGFR mutations (including exon 20 insertions)	++	-	+/-	++	+	++
ALK rearrangements	-	++	+	+	+	++
ROS1 rearrangements	-	++	-	+	+	++
BRAF mutations (including V600E)	++	-	-	++	+	++
MET exon 14 mutation	+/-	-	-	++	+	++
HER2 mutations	++	_	-	++	+	++
RET rearrangements	-	++	-	+	+	++
MET amplification	-	++	+/-	+	+/-	+
NTRK rearrangements	-	++	-	+	+	++
PD-L1 Protein Expression	-	-	++	-	-	-
Turnaround time (optimistically)	1-3 days	1-3 days	1 day	7-14 days	7-10 days	7-14 days

- ++ Most sensitive
- + Less sensitive
- +/- Least sensitive (or technical limitations)
- Not appropriate

What is the incremental value of targeted RNAseq in Lung Adenocarcinomas studied by MSK-IMPACT targeted DNAseq?



Clinical relevance of complementary RNA sequencing

High Yield of RNA Sequencing for Targetable Kinase Fusions in Lung Adenocarcinomas with No Mitogenic Driver Alteration

Detected by DNA Sequencing and Low Tumor Mutation Burden

Benayed R. et al. Clin Cancer Res May 2019

Clinical benefit of matched targeted therapy (n = 10)

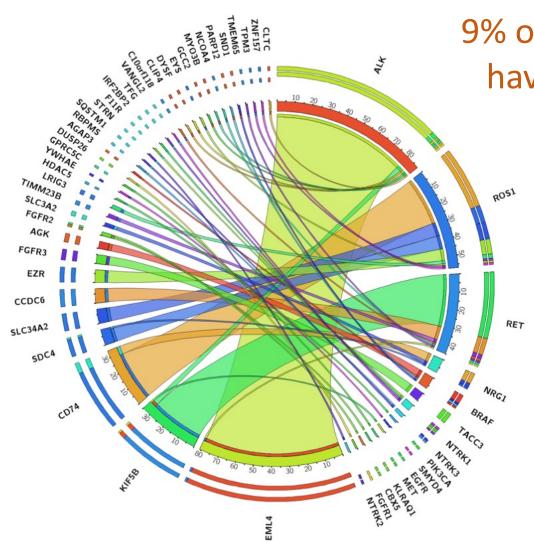
Rearrangement	Matched therapy	Best response*
EML4-ALK	Alectinib	SD
CD74-ROS1	Entrectinib	SD
SQSTM1-NTRK3	Larotrectinib	PR**
STRN-NTRK2	Larotrectinib	SD
CD74-ROS1	Entrectinib	PR**
CD74-NRG1	Afatinib	SD
MET Exon14 Skipping	Crizotinib	SD
SLC34A2-ROS1	Crizotinib	PD
SLC34A2-ROS1	Crizotinib	SD
SDC4-NRG1	Afatinib	PD

^{*} Response assessment by RECIST version 1.1. **, Confirmed PR.

8 of the 10 treated pts had clinical benefit from matched therapy

Gene fusions landscape in 2,522 lung adenocarcinomas

Detected by comprehensive DNASeq and RNASeq

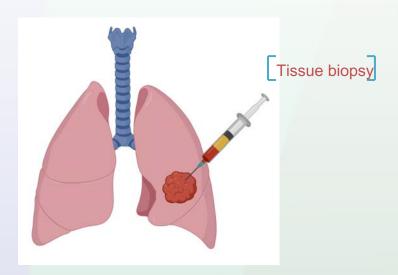


9% of lung adenoCa patients have in-frame, targetable gene fusions.

Tissue and Plasma Genomic Profiling

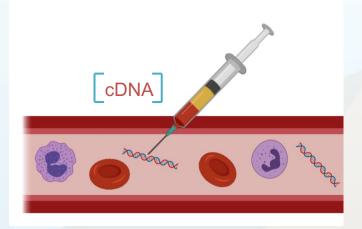
Tissue Testing

- Adequate tissue available
- High specificity and sensitivity
- Able to perform sequencing on long strands of DNA (entire coding regions) and WGS/WES
- DNA and RNA analysis capabilities
- Only tests sample that was biopsied

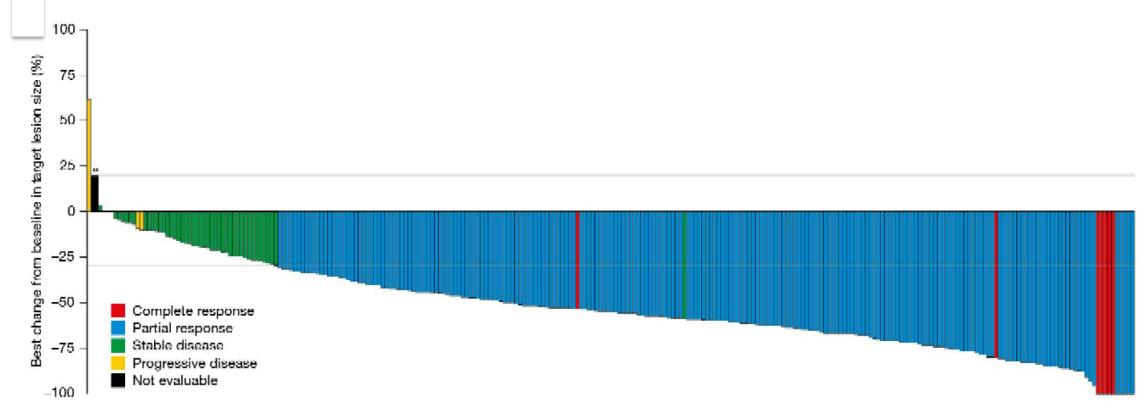


Plasma Testing

- Difficult to biopsy disease (or bone only)
- Less invasive
- Disease monitoring/resistance mechanisms
- Sequential testing
- Usually smaller panels
- DNA only (currently)
- Limited fusion detection (especially large genes i.e. NTRK)
- Monitors mutations in multiple lesions



Osimertinib (EGFR tyrosine kinase inhibitor) frontline for metastatic *EGFR* mutated NSCLC



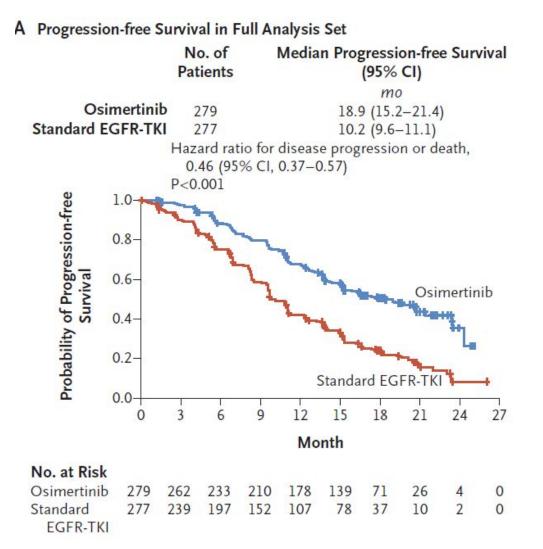
Objective response rate 80% (95% CI, 75 to 85)

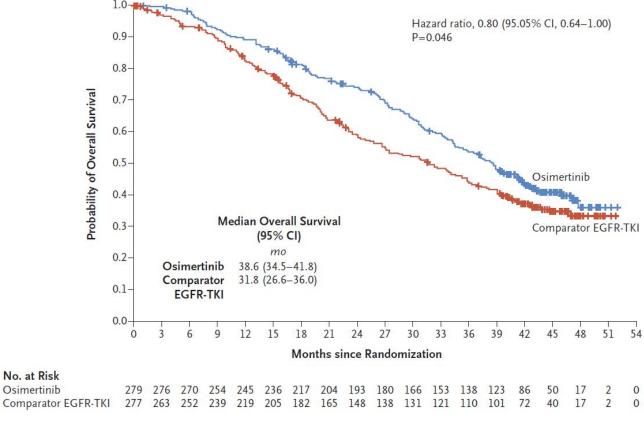
Disease-control rate was 97% (95% CI, 94 to 99)

Median best percentage shange in target lesion size. E4

Median best percentage change in target lesion size -54.7%

Osimertinib improves PFS & OS compared to older generation EGFR Tyrosine Kinase Inhibitors (TKIs)

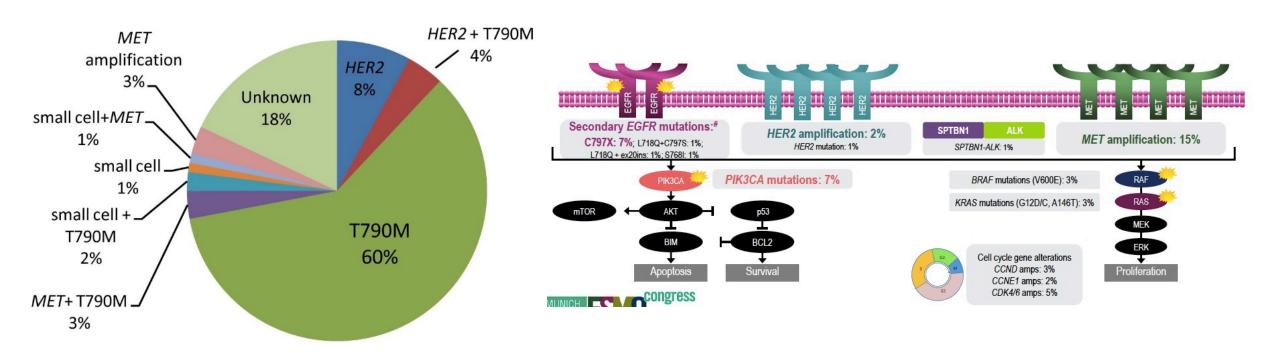




31% crossover rate

Ramalingam S et al. N Engl J Med. 2020 Jan 2;382(1):41-50.

Resistance to targeted therapy is inevitable



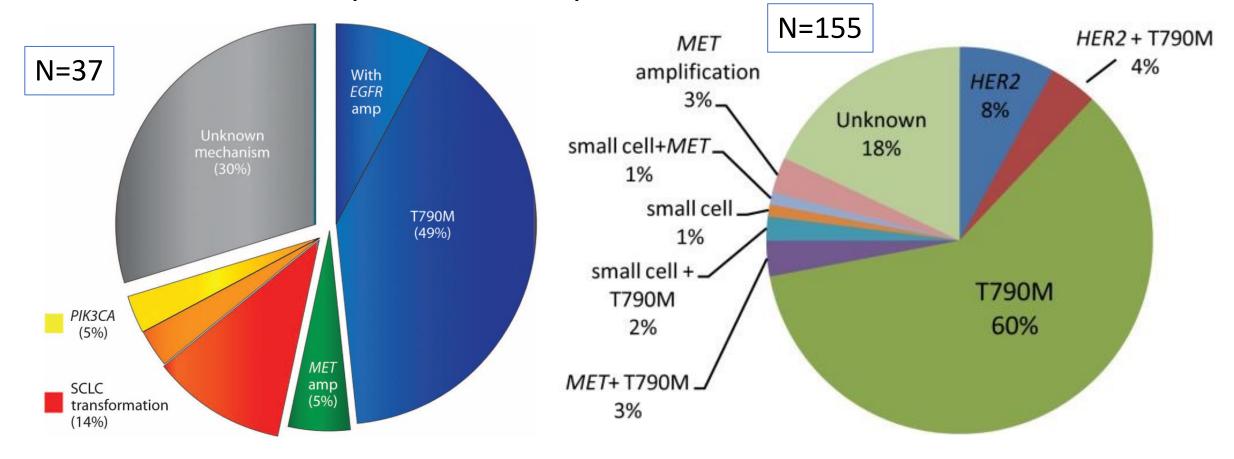
Dominant mechanism of resistance for 1st generation EGFR TKIs erlotinib or gefitinib

Yu H et al. Clin Cancer Res. 2013 Apr 15;19(8):2240-7.

Heterogeneous & multiple simultaneous mechanisms of resistance to 3rd generation EGFR TKI osimertinib

Ramanlingam SS et al. ESMO 2018, Munich

EGFR TKI AR pie charts prior to the clinical NGS era



Genotypic and Histological Evolution of Lung Cancers Acquiring Resistance to EGFR Inhibitors

Science Transl Med March 2011

Analysis of Tumor Specimens at the Time of Acquired Resistance to EGFR-TKI Therapy in 155 Patients with *EGFR*-Mutant Lung Cancers

Clinical Cancer Res March 2013

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 14 skipping and MET amplification mutations PD-L1 TPS 95%
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- 60yr Female
- Diagnosed 2017 with stage IV NSCLC adenocarcinoma
 - EGFR exon 19 deletion mutation p.E746_A751del and PDL-1 positive
- Initiated afatinib (9/2017 10/2018)
 - Developed EGFR T790M on liquid biopsy
- Osimertinib (10/2018 12/2019) until progression
 - T790M resolved; TP53 H193Y, MET/FGFR1/EGFR/CDK4/ amplifications emerge; ex19del remains
- Osimertinib + Crizotinib (2/2020 5/2020)
 - BRAF V600E, METD1228N (crizotinib resistant/cabozantinib sensitive), and KRAS G12V emerge
- Osimertinib + Pemetrexed/carboplatin (5/2020 1/2021)



Alteration	% cfDNA or Amp	Alter
EGFR E746_T751delinsA (Exon 19 deletion)	59.9%	0.06%
<i>TP53</i> H193Y	17.0%	ND
KRAS G12V	3.7%	ND_
FGFR1 I64M	0.5%	o
EGFR S220C	0.4%	o
BRAFV600E	0.3%	o
MET D1228N	0.06%	oND

CDK4 Amplification Amplifications not graphed above	High (+++)	ND_
EGFR Amplification Amplifications not graphed above	Medium (++)	ND &
BRAF Amplification Amplifications not graphed above	Medium (++)	ND &
FGFR1 Amplification Amplifications not graphed above	ND	ND.
MET Amplification Amplifications not graphed above	ND	ND_
HNF1A R229*	ND	o- ND
PDGFRA R979H	ND	o- ND
KIT Splice Site SNV	ND	o- ND
EGFR T790M	ND	o



Alteration	% cfDNA or Amp	Alteration Tre
EGFR E746_T751delinsA (Exon 19 deletion)	59.9%	0.06% 1.8%
<i>TP53</i> H193Y	17.0%	ND 0.2%
KRAS G12V	3.7%	ND ND
FGFR1 l64M	0.5%	c
EGFR S220C	0.4%	o o
BRAF V600E	0.3%	c c
<i>MET</i> D1228N	0.06%	o o

CDK4 Amplification Amplifications not graphed above	High (+++)	ND ND
		Plasn
EGFR Amplification Amplifications not graphed above	Medium (++)	ND ND
		Plasn
BRAF Amplification Amplifications not graphed above	Medium (++)	ND ND
		Plasn
FGFR1 Amplification Amplifications not graphed above	ND	ND ND
		Plasn
MET Amplification Amplifications not graphed above	ND	ND ND
		Plasn
HNF1A R229*	ND	0
		ND ND
PDGFRA R979H	ND	
		ND ND
KIT Splice Site SNV	ND	
		ND ND
EGFR T790M	ND	
		ND 0.23



Alteration	% cfDNA or Amp	Alteration Trend		
EGFR E746_T751delinsA (Exon 19 deletion)	59.9%	0.06% 1.8% 44.9%		
TP53 H193Y	17.0%	ND 0.2% 10.5%		
KRAS G12V	3.7%	ND ND ND		
FGFR1 I64M	0.5%	o o o		
EGFR S220C	0.4%	o o o		
BRAF V600E	0.3%	o o o		
<i>MET</i> D1228N	0.06%	ND ND ND		

CDK4 Amplification Amplifications not graphed above	High (+++)	ND ND
		Plasma copy number
EGFR Amplification Amplifications not graphed above	Medium (++)	ND ND 28
		Plasma copy number
BRAF Amplification Amplifications not graphed above	Medium (++)	ND ND ND
		Plasma copy number
FGFR1 Amplification Amplifications not graphed above	ND	ND ND
		Plasma copy number
MET Amplification Amplifications not graphed above	ND	ND ND
		Plasma copy number
HNF1A R229*	ND	
		ND ND 1.7%
PDGFRA R979H	ND	
		ND ND 0.1%
	1/4.2	Т
KIT Splice Site SNV	ND	0 0
		ND ND 0.1%
EGFR T790M	ND	
		ND 0.2% ND



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EGFR E746 T751delinsA (Exon 19 deletion)	59.9%	0.06% 1.8% 44.9% 59.9%
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KRAS G12V	3.7%	ND ND ND 3.7%
FGFR1 I64M	0.5%	ND ND ND 0.5%
EGFR S220C	0.4%	ND ND ND 0.4%
BRAF V600E	0.3%	ND ND ND 0.3%
<i>MET</i> D1228N	0.06%	ND ND ND 0.08%

CDK4 Amplification Amplifications not graphed above	High (+++)	ND ND 73
		Plasma copy number
EGFR Amplification Amplifications not graphed above	Medium (++)	ND ND 2.8 3
		Plasma copy number
BRAF Amplification Amplifications not graphed above	Medium (++)	ND ND ND
		Plasma copy number
FGFR1 Amplification Amplifications not graphed above	ND	ND ND 2.3
		Plasma copy number
MET Amplification Amplifications not graphed above	ND	ND ND
		Plasma copy number
HNF1A R229*	ND	
		ND ND 1.7%
PDGFRA R979H	ND	
		ND ND 0.1% NO
KIT Splice Site SNV	ND	
		ND ND 0.1% NO
EGFR T790M	ND	
		ND 0.3% ND NO



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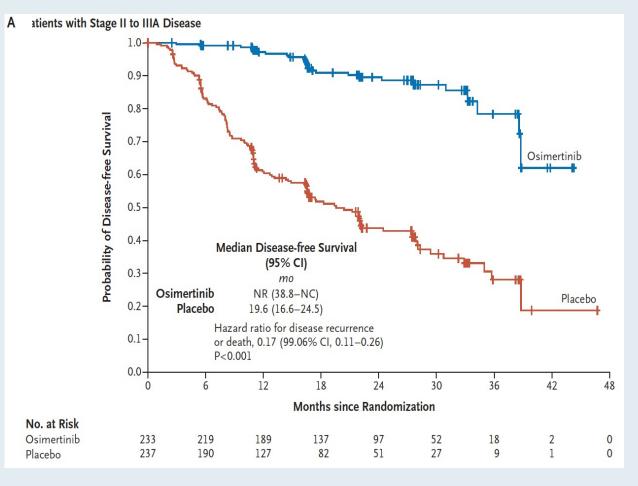
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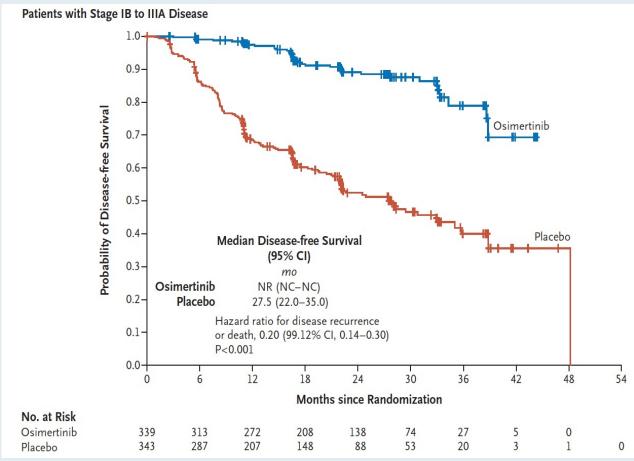
Osimertinib in Resected EGFR-Mutated Non–Small-Cell Lung Cancer

Yi-Long Wu, M.D., Masahiro Tsuboi, M.D., Jie He, M.D., Thomas John, Ph.D., Christian Grohe, M.D., Margarita Majem, M.D., Jonathan W. Goldman, M.D., Konstantin Laktionov, Ph.D., Sang-We Kim, M.D., Ph.D., Terufumi Kato, M.D., Huu-Vinh Vu, M.D., Ph.D., Shun Lu, M.D., Kye-Young Lee, M.D., Ph.D., Charuwan Akewanlop, M.D., Chong-Jen Yu, M.D., Ph.D., Filippo de Marinis, M.D., Laura Bonanno, M.D., Manuel Domine, M.D., Ph.D., Frances A. Shepherd, M.D., Lingmin Zeng, Ph.D., Rachel Hodge, M.Sc., Ajlan Atasoy, M.D., Yuri Rukazenkov, M.D., Ph.D., and Roy S. Herbst, M.D., Ph.D.,



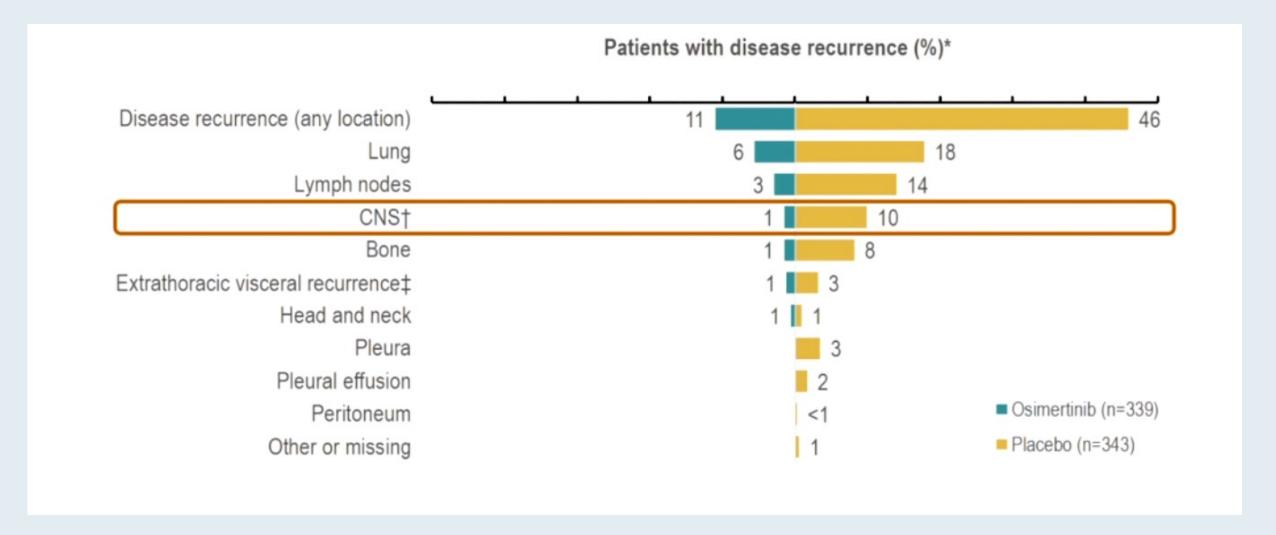
ADAURA: Disease-Free Survival by Stage





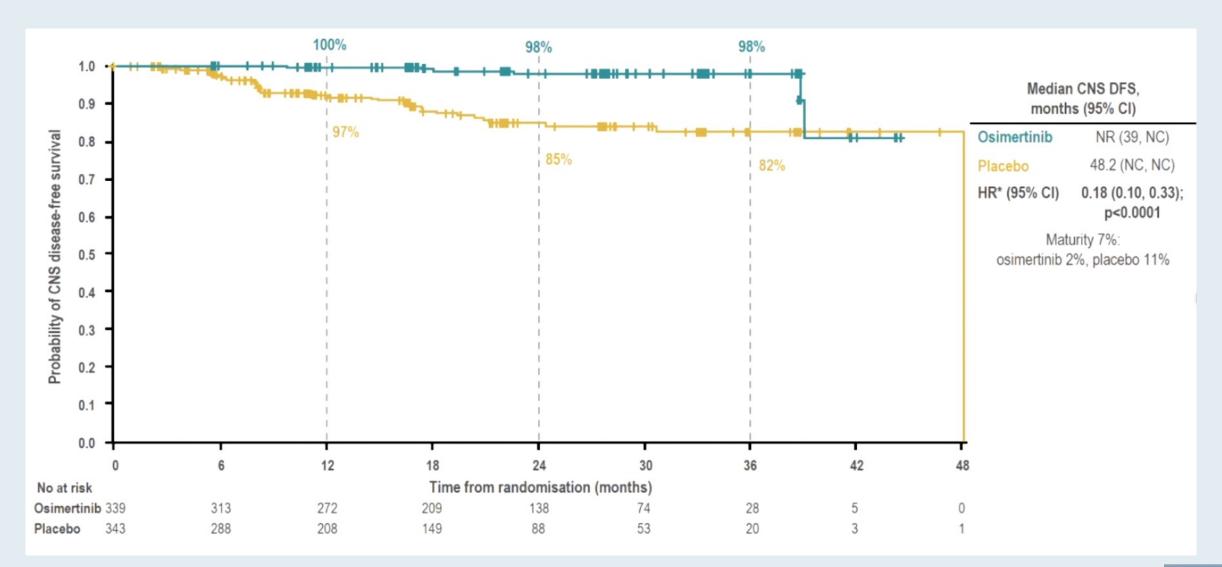


ADAURA: Sites of Disease Recurrence





ADAURA: CNS DFS in Overall Population





Agenda

Module 1: Role of Genomic Profiling; New Developments in NSCLC with an EGFR Mutation

- Dr McKenzie: A 60-year-old woman with NSCLC and an EGFR exon 19 deletion
- Dr Hussein: A 51-year-old man with metastatic lung adenocarcinoma and an EGFR exon 20 mutation

Module 2: Other Novel Targets for Patients with NSCLC

- Dr Neal: A 51-year-old man with RET-positive NSCLC
- Dr Neal: A 64-year-old woman with metastatic NSCLC and a HER2 exon 20 insertion
- Dr McKenzie: A 78-year-old man with NSCLC and a TRK fusion
- Dr Jasani: A 35-year-old woman with newly diagnosed mNSCLC ALK mutation
- Dr Mohamed: A 71-year-old woman with metastatic adenocarcinoma of the lung with MET exon
 14 skipping and MET amplification mutations PD-L1 TPS 95%
- Dr Zafar: A 64-year-old woman with discordant BRAF mutation testing results



Case Presentation – Dr Hussein: A 51-year-old man with metastatic adenocarcinoma of the lung and an EGFR exon 20 mutation

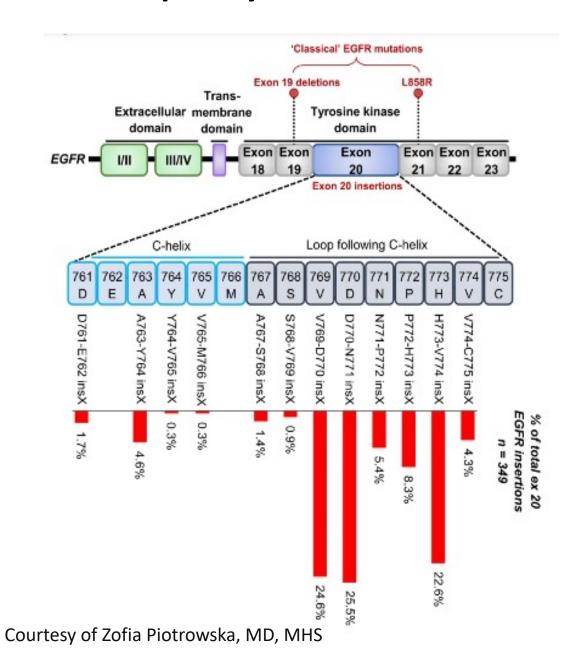


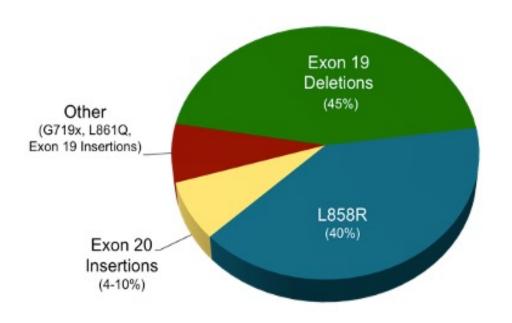
Dr Atif Hussein

- Summer 2017: Patient presented with back pain
- 02/2018: Imaging revealed T9 lytic lesion and 3 cm left upper lobe lung mass
- Biopsy of T9: Metastatic adenocarcinoma
- NGS: exon 20 EGFR mutation | MSI stable | BRAF, ROS1, and ALK no abnormalities | PD-L1 negative 0%
- Radiation therapy to T9
- Started on carboplatin/pemetrexed/bevacizumab: Good PR in the lungs → maintenance pemetrexed/bevacizumab
- 5 months later PD in the lungs, bone and brain
- Cyberknife to the brain
- Started on pembrolizumab, developed autoimmune hepatitis
 - Despite steroids LFTs continued to be abnormal; pembrolizumab discontinued due to PD
- Started on TAK788 on a clinical trial → Stable disease for 3 months then worsening disease in the lungs,
 brain with leptomeningeal disease
- Administered high dose osimertinib with no response



Frequency of EGFR Exon 20 Mutations

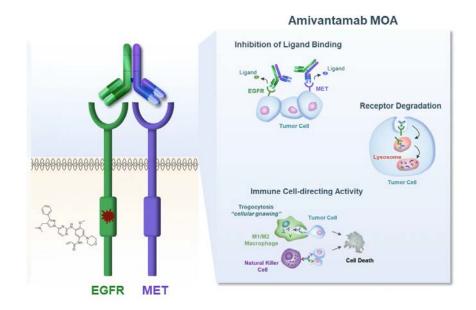




Exon 20 NSCLC: US and China				
		Exon 20 Frequency		umber of atients/year
United	EGFR	2.1%		
States	HER2	1.5%	3.6%	7700
China	EGFR	2.4%	6.3% 411	41100
Cillia	HER2	3.9%	0.576	41100

Amivantamab

Amivantamab is a bispecific antibody targeting EGFR + MET

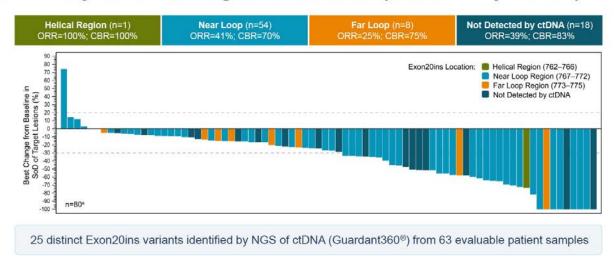


 Amivantamab recently received FDA approval for EGFR exon 20 insertion+ NSCLC

CHRYSALIS Trial

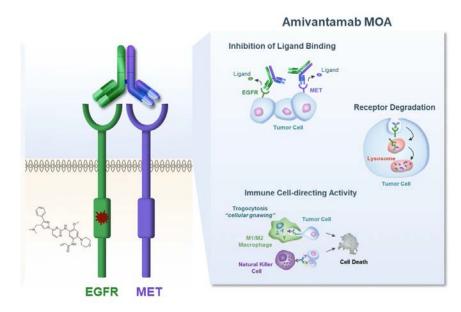
	N=81 (EGFR ins20)
ORR	40%
mDOT	11.1 mo (95% CI, 6.9-NR)
mPFS	8.3 mo (95% CI, 6.5-10.9)

Best ORR by Insertion Region of Exon 20 (detected by ctDNA)



Amivantamab + Lazertinib

Amivantamab is a bispecific antibody targeting EGFR + MET



- Amivantamab recently received FDA approval for EGFR exon 20 insertion+ NSCLC
- Amivantamab + Lazertinib (3rd gen EGFR TKI) is being evaluated in TKI-resistant EGFR+ NSCLC

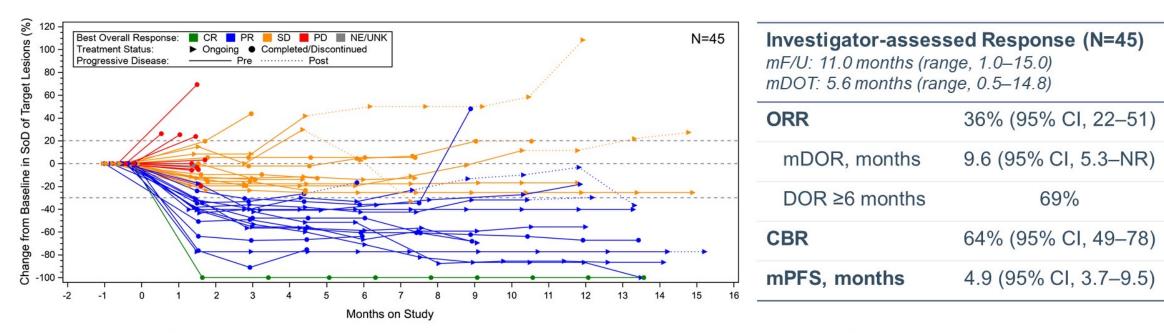
Amivantamab + Lazertinib in TKI-resistant EGFR+ NSCLC

CHRYSALIS Phase 1 Study Design: Combination Cohort (NCT02609776)

Key Eligibility Criteria Biomarker Analysis^a **Key Objectives** NGS of pretreatment tumor biopsy Metastatic/unresectable NSCLC Establish RP2CD and ctDNA collected prospectively Measurable disease (expansion cohort) · Safety and efficacy at IHC for EGFR/MET expression RP2CD EGFR Exon19del or L858R mutation 1050/1400 ma RP2CD Osimertinib-Amiyantamab relapsed, 240 mg lazertinib 1050 mg (<80 kg) chemotherapy-NGS 1400 mg (≥80 kg) IHC naïve Tumor (n=29) Intravenous dosing (n=20)EGFR Exon19del ctDNA (n=44) C1 QW, C2+ Q2W or L858R 240 mg lazertinib (N=45)240 mg lazertinib Oral daily dosing **Dose Escalation Expansion Cohort Biomarker Analysis**

1. Chul B et al. ASCO 2021

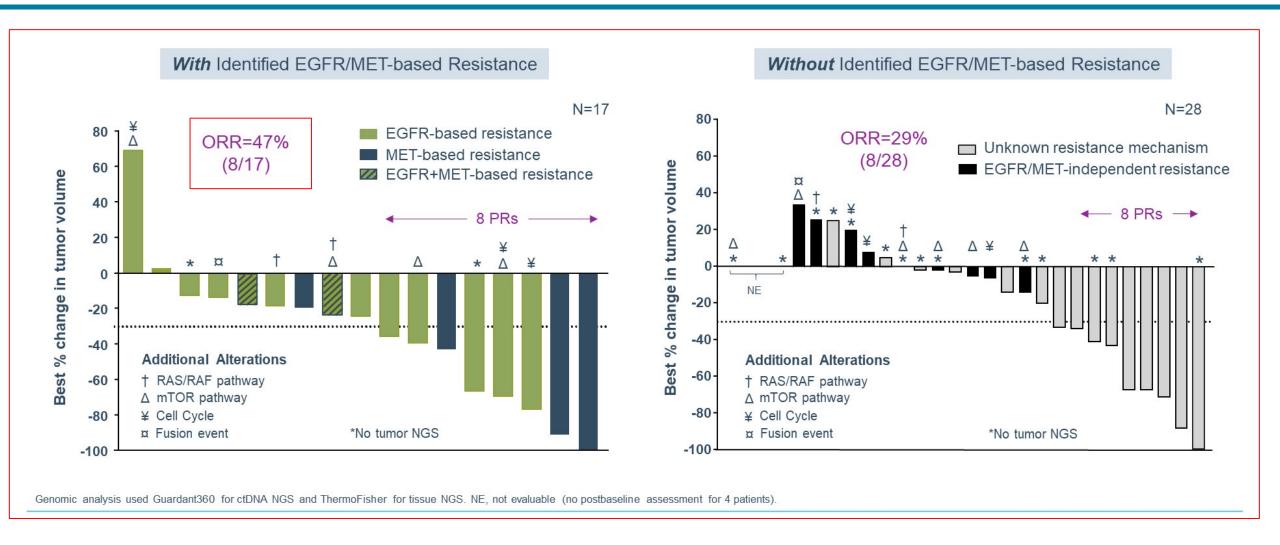
Amivantamab + Lazertinib



- Safety profile consistent with previous experience with amivantamab + lazertinib¹
- Most common AEs were IRR (78%), rash (acneiform dermatitis, 51% + rash, 27%), and paronychia (49%)
 - Majority were grade 1–2
- Treatment-related: grade ≥3 AE (16%), discontinuations (4%), dose reductions (18%)



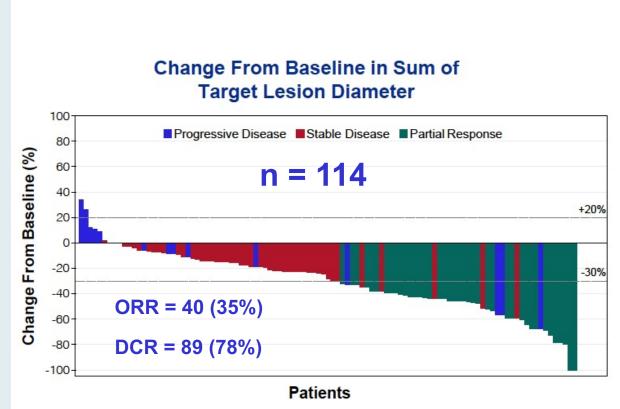
Amivantamab + Lazertinib



1. Chul B et al. ASCO 2021

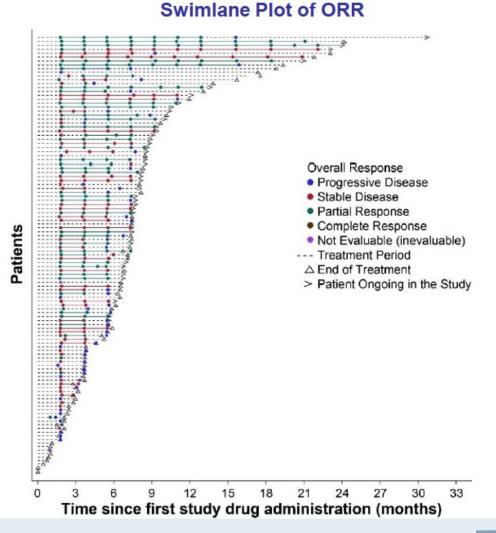
Courtesy of Zofia Piotrowska, MD, MHS

EXCLAIM: Mobocertinib in Platinum-Pretreated NSCLC with EGFR Exon 20 Insertions



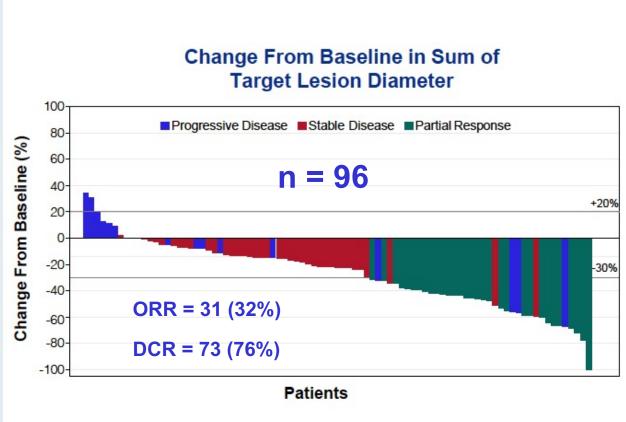
 94 patients (82%) had a reduction from baseline in the sum of target lesion diameter

ORR, objective response rate; PPP, platinum-pretreated patients



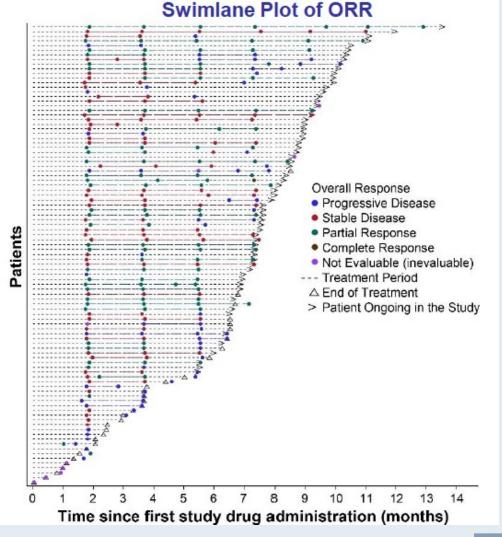


Mobocertinib Resulted in Reductions in Target Lesion Volume in EXCLAIM Cohort



 77 patients (80%) had a reduction from baseline in the sum of target lesion diameter

ORR, objective response rate





Module 1: Role of Genomic Profiling; New Developments in NSCLC with an EGFR Mutation

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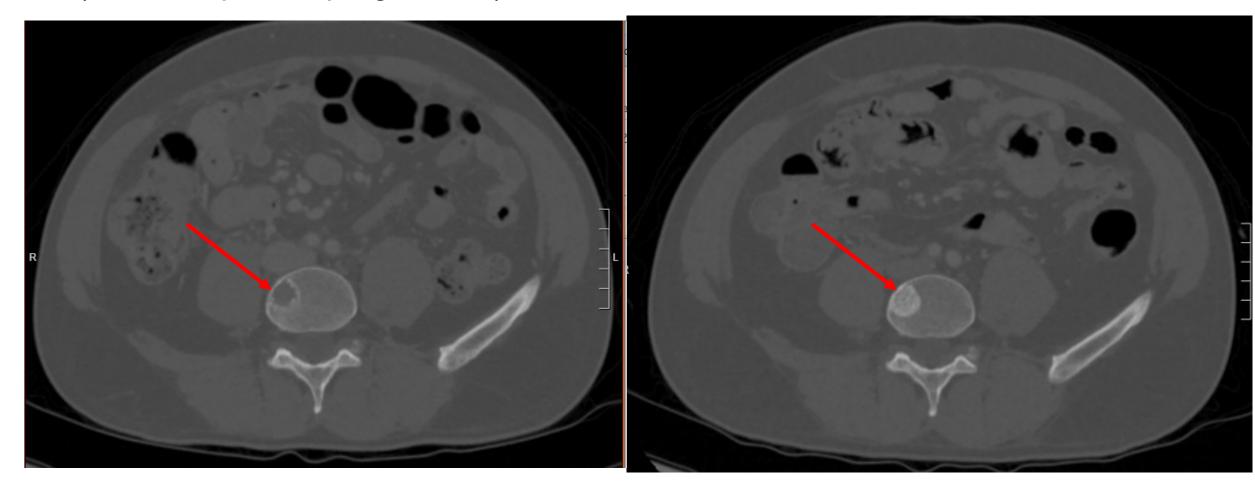


Case Presentation – Dr Neal: A 51-year-old man with RET positive NSCLC

- 51 year old Vietnamese neversmoking man presented for right sided loss of body sensation
- MRI showed right pontine hematoma with 1.3 cm underlying metastasis and other scattered mets
- CT showed 1/7 cm left lower lobe mass with hilar hode and pleural nodule, multiple bone mets
- Biospy of right radius bone adenocarcinoma, TTF1 positive, PD-L1 10%, EGFR, ALK, ROS1 negative.
- Radiosurgery to 5 brain mets
- Systemic chemotherapy with carbo/pemetrexed/pembrolizumab
- After 5 months, progression with new liver mets and progressive brain mets. Repeat biopsy with NGS sequencing: RET fusion positive
- Started selpercatinib 160 mg BID; 6 week interval response in lungs, brain. Sclerosis of bone lesions.

Case Presentation – Dr Neal: A 51-year-old man with RET positive NSCLC (continued)

Pre- and 6 week month post selpercatinib showing bone treatment response (sclerosis/pseudoprogression)

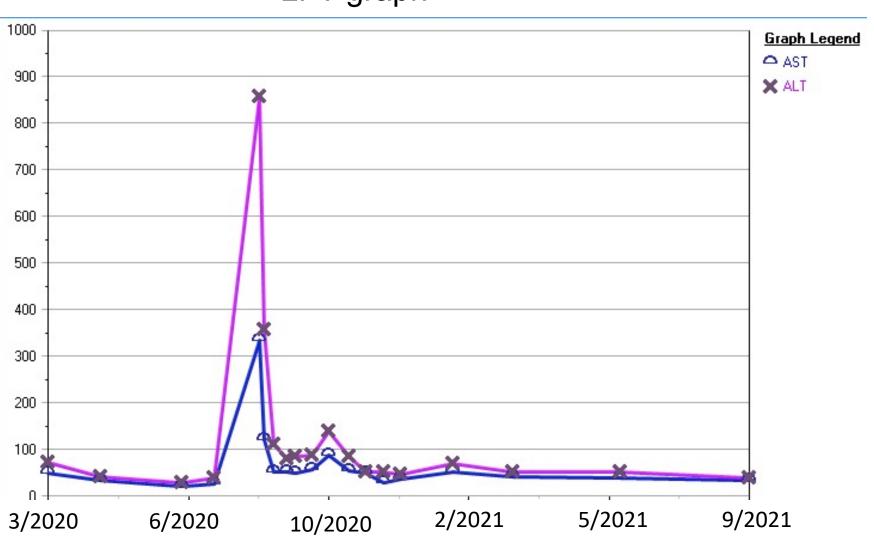


Case Presentation – Dr Neal: A 51-year-old man with RET positive NSCLC (continued)

- Aminotransferase spike 2 weeks after starting
- Held drug, reached ULN within 2 more weeks
- Restarted at 80 mg BID
- No significant LFT elevation after 4 weeks
- Escalated to 160 mg PO QAM, 80 mg PO QPM; LFTs close to ULN but stable
- Ongoing response at 14 months!

Case Presentation – Dr Neal: A 51-year-old man with RET positive NSCLC (continued)





Safety and Efficacy of Pralsetinib in Patients with Advanced *RET* Fusion-Positive Non-Small Cell Lung Cancer: Update from the ARROW Trial

Curigliano G et al.

ASCO 2021; Abstract 9089.



ARROW Study Design

Eligibility criteria

- Age ≥18 years
- Advanced or metastatic solid tumor
- RET alteration per local assessment
- Measurable disease (RECIST v1.1)
- ECOG PS 0-1

Protocol amendment
(July 19, 2019)
Eligibility criteria were
expanded to allow
treatment-naïve patients
with NSCLC who were
candidates for
platinum-based therapy

Phase 1 dose escalation (Completed)

Phase 2 dose determined: 400 mg QD

Phase 2 dose expansion (Ongoing) Treated at 400 mg QD



1º endpoints:

- ORR (BICR per RECIST v1.1)
- Safety

RET fusion-positive NSCLC

Medullary thyroid cancer^a

Other RET-altered tumors

Key 2º endpoints:

• DOR

· PFS

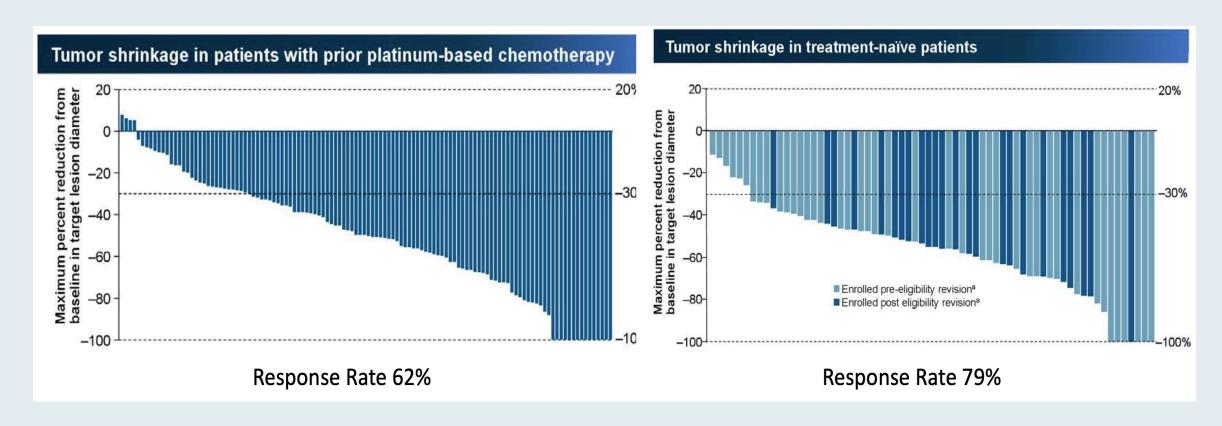
• CBR

· OS

• DCR



ARROW Primary Endpoint: Response to Pralsetinib



ORR (response-evaluable): All – 69%, Prior platinum – 62%, Treatment-naïve – 79%



ARROW: Efficacy Summary

	Measurable disease population						
	RET	Treatment-naïve			Prior treatment		
	fusion-positive NSCLC (n=216)	AII (n=68)	Pre-eligibility revision (n=43) ^a	Post eligibility revision (n=25)ª	Prior platinum (n=126)	Prior non-platinum (n=22)	
ORR, %	69	79	74	88	62	73	
(95% CI)	(62-75)	(68-88)	(59-87)	(69 - 98)	(53-70)	(50-89)	
Best overall response, n (%)							
CR	9 (4)	4 (6)	4 (9)	0	5 (4)	0	
PR	139 (64)	50 (74)	28 (65)	22 (88)	73 (58)	16 (73)	
SD	50 (23)	9 (13)	7 (16)	2 (8)	37 (29)	4 (18)	
PD	10 (5)	3 (4)	3 (7)	0	5 (4)	2 (9)	
NE	8 (4)	2 (3)	1 (2)	1 (4)	6 (5)	0	
DCR, % (95% CI)b	92 (87-95)	93 (84-98)	91 (78-97)	96 (80-100)	91 (85-96)	91 (71-99)	
CBR, % (95% CI)°	77 (71-82)	82 (71-91)	79 (64-90)	88 (69-98)	74 (65-81)	77 (55-92)	
mDOR, mo (95% CI)	22.3 (15.1-NR)	NR (9.0-NR)	11.0 (7.4-NR)	NR (NR-NR)	22.3 (15.1-NR)	NR (9.2-NR)	
mPFS, mo (95% CI)d	16.4 (11.0-24.1) n=233	13.0 (9.1–NR) n=75	10.9 (7.7–NR) n=47	NR (NR-NR) n=28	16.5 (10.5–24.1) n=136	12.8 (9.1–NR) n=22	



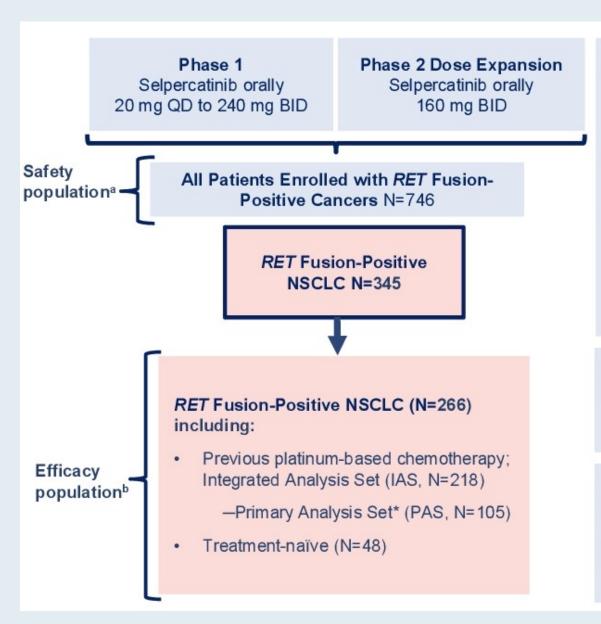
Updated Overall Efficacy and Safety of Selpercatinib in Patients with RET Fusion-Positive Non-Small-Cell Lung Cancer (NSCLC): LIBRETTO-001 Study

Besse B et al.

ASCO 2021; Abstract 9065.



LIBRETTO-001 Study Design



Study Design

- Ongoing, global, multicenter Phase 1/2 trial (NCT03157128) conducted in 16 countries and 89 sites
- Patients enrolled based on locally identified RET alterations using NGS, FISH, or PCR
- Key inclusion criteria: Diagnosis of advanced or metastatic solid tumor, ECOG PS 0 to 2, QTc of ≤470 msec, adequate organ function, asymptomatic CNS metastases permitted

Primary Endpoint

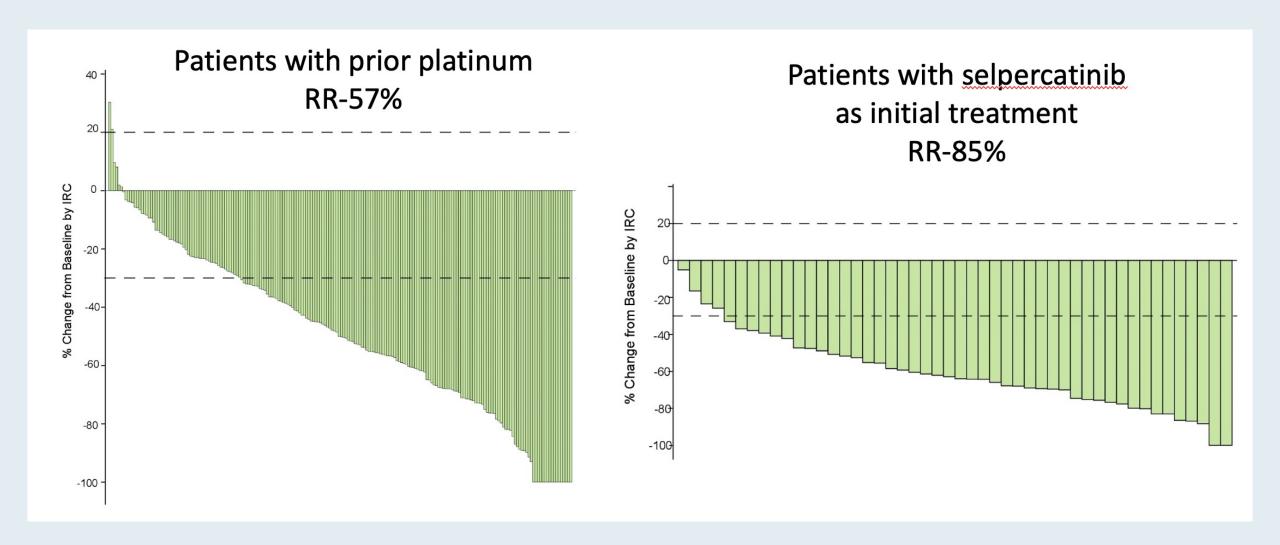
ORR (RECIST v 1.1) by Independent Review

Secondary Endpoints Included

- Duration of Response (DOR)
- Progression-Free Survival (PFS)
- Safety



LIBRETTO-001: Response to Selpercatinib





Module 1: Role of Genomic Profiling; New Developments in NSCLC with an EGFR Mutation

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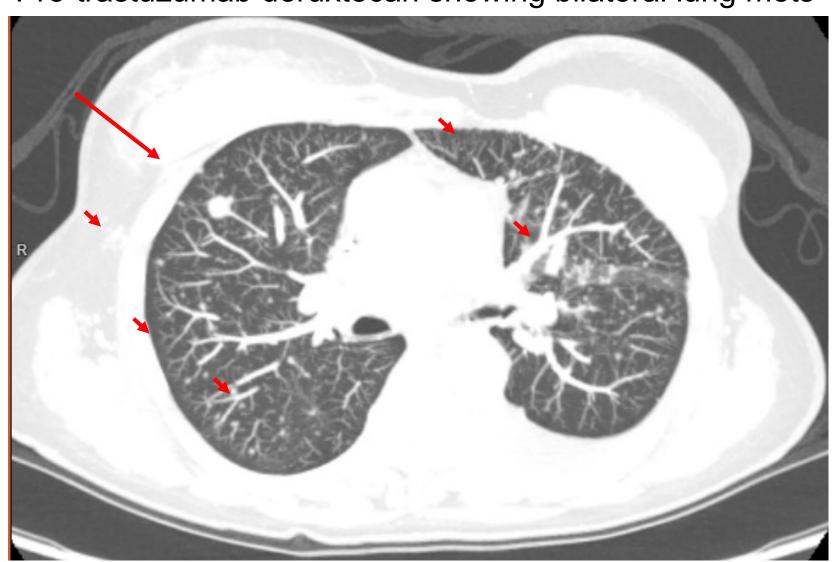


Case Presentation – Dr Neal: A 64-year-old woman with metastatic NSCLC and a HER2 exon 20 insertion

- 64 year old neversmoking Caucasian woman with h/o stage I breast cancer 3 years before and rheumatoid arthritis, no active therapy for either
- Productive cough for 3 months, CXR with opacities
- CT left upper lobe mass, bilateral lung nodules, liver mets, bone mets.
- FNA biopsy subcarinal and mediastinal nodes; adenocarcinoma. Ruled out for TB with AFB x 3
- MRI brain with 12 mets, 5-8 mm
- She has good pulmonary function and performance status (walks a mile in hills per day)
- EGFR
- 130 gene NGS panel (3 weeks after path diagnosis) ERBB2 G778_P780dup
- Interpretation: HER2 exon 20 insertion mutation
- Radiosurgery to brain mets, started carbo/pemetrexed/bevacizumab with response
- Zolendronic acid for bone mets
- After 12 months: Progression predominantly in liver mets during maintenance pemetrexed/bevacizumab
- Started trastuzumab deruxtecan off label 5.4 mg/kg IV every 3 weeks

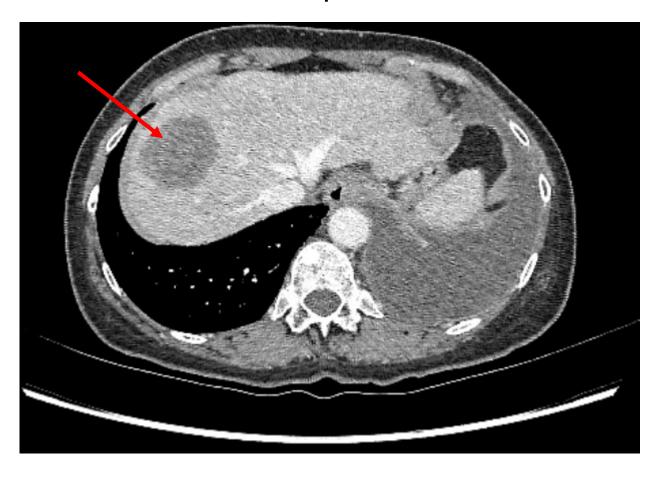
Case Presentation – Dr Neal: A 64-year-old woman with metastatic NSCLC and a HER2 exon 20 insertion (continued)

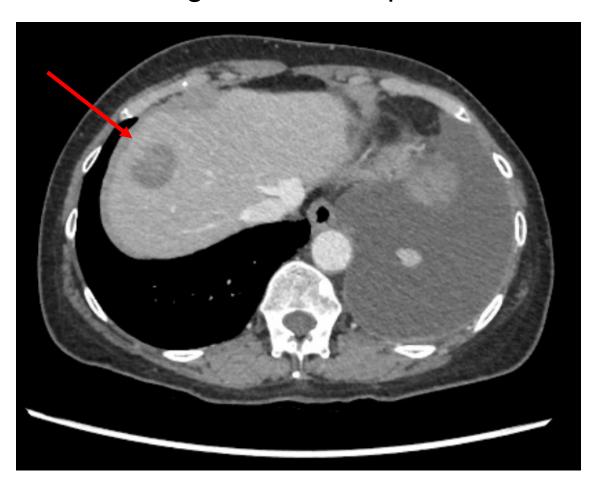
Pre-trastuzumab deruxtecan showing bilateral lung mets



Case Presentation – Dr Neal: A 64-year-old woman with metastatic NSCLC and a HER2 exon 20 insertion (continued)

Pre- and 3 month post trastuzumab deruxtecan showing liver met response





Case Presentation – Dr Neal: A 64-year-old woman with metastatic NSCLC and a HER2 exon 20 insertion (continued)

- Response after 6 weeks with continued durable response in liver mets after
- Anemia and thrombocytopenia led to dose delays for 1-2 weeks, 3-4 week cycles
- Ongoing response 10 months later

Interim Analysis of DESTINY-Lung01 in HER2-Overexpressing Cohort 1: Efficacy Summary

Parameter	IHC 3+ (n = 10)	IHC 2+ (n = 39)	Overall (n = 49)
Confirmed ORR, n (%; 95% CI)	2 (20.0; 2.5-55.6)	10 (25.6; 13.0-42.1)	12 (24.5; 13.3-38.9)
■ CR	0	1 (2.6)	1 (2.0)
■ PR	2 (20.0)	9 (23.1)	11 (22.4)
■ SD	6 (60.0)	16 (41.0)	22 (44.9)
■ PD	1 (10.0)	10 (25.6)	11 (22.4)
Not evaluable	1 (10.0)	3 (7.7)	4 (8.2)
DCR, n (%; 95% CI)	8 (80.0; 44.4-97.5)	26 (66.7; 49.8-80.9)	34 (69.4; 54.6-81.8)
Median DoR, mos (95% CI)	6.0 (NE-NE)	5.8 (3.2-NE)	6.0 (3.2-NE)

^{*}Full analysis set data.

ORIGINAL ARTICLE

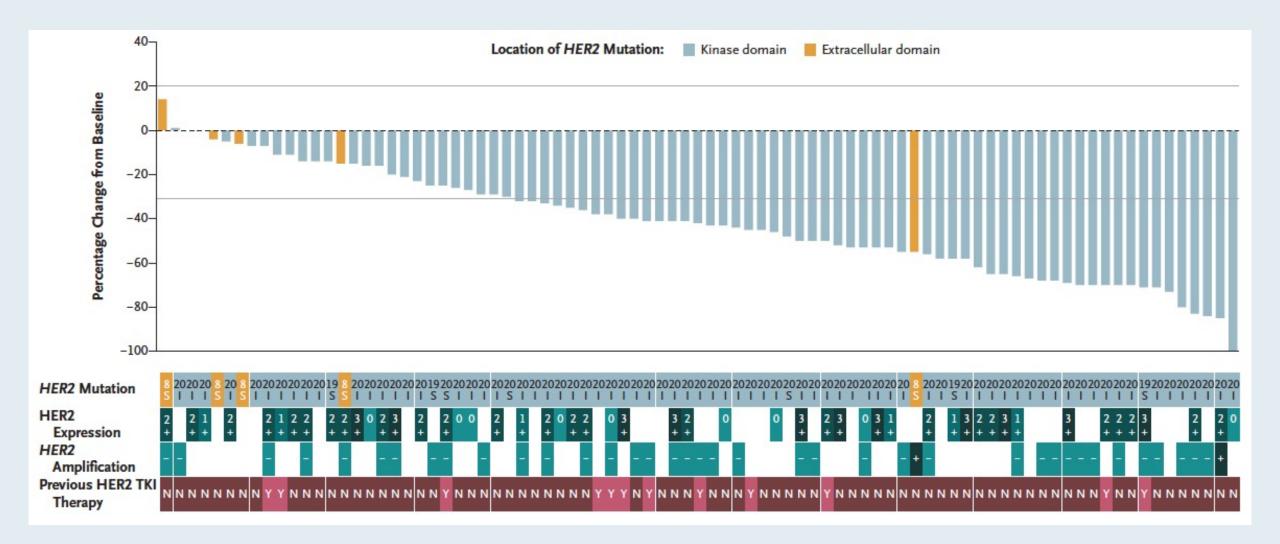
Trastuzumab Deruxtecan in HER2-Mutant Non–Small-Cell Lung Cancer

Bob T. Li, M.D., Ph.D., M.P.H., Egbert F. Smit, M.D., Ph.D.,
Yasushi Goto, M.D., Ph.D., Kazuhiko Nakagawa, M.D., Hibiki Udagawa, M.D.,
Julien Mazières, M.D., Misako Nagasaka, M.D., Ph.D., Lyudmila Bazhenova, M.D.,
Andreas N. Saltos, M.D., Enriqueta Felip, M.D., Ph.D., Jose M. Pacheco, M.D.,
Maurice Pérol, M.D., Luis Paz-Ares, M.D., Kapil Saxena, M.D., Ryota Shiga, B.Sc.,
Yingkai Cheng, M.D., Ph.D., Suddhasatta Acharyya, Ph.D., Patrik Vitazka, M.D., Ph.D.,
Javad Shahidi, M.D., David Planchard, M.D., Ph.D., and Pasi A. Jänne, M.D., Ph.D.,
for the DESTINY-Lung01 Trial Investigators*

N Engl J Med 2021; [Online ahead of print].

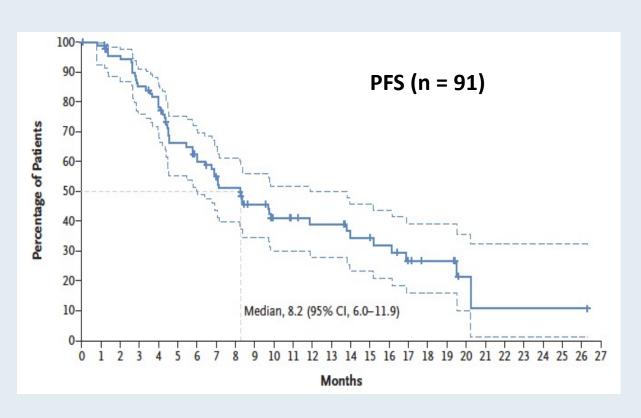


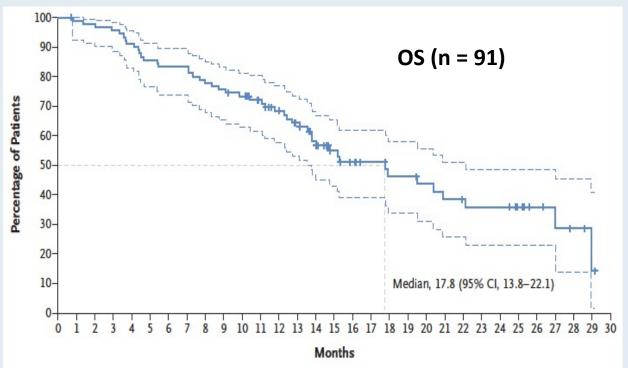
DESTINY-Lung01: Antitumor Activity





DESTINY-Lung01: Survival in the Overall Population







Module 1: Role of Genomic Profiling; New Developments in NSCLC with an EGFR Mutation

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Case Presentation – Dr McKenzie: A 78-year-old man with NSCLC and a TRK fusion

- 78yr Male never smoker
- 8/2020 Stage 3A (cT2bN2) NSCLC squamous cell carcinoma of RUL lobe endobronchial ultrasound showed mediastinal invasion into R paratracheal space
- Started XRT 9/10.
- Weekly carbo/taxol 9/11.
 - Completed chemoRT on 10/16/20.
- Started consolidative durva 11/20/20.
- Imaging 9/24/21 followed by PET showed recurrence in lungs, liver (3 nodules), and R adrenal.
- NGS sent upon metastatic diagnosis Tissue and plasma (both 10/2021)

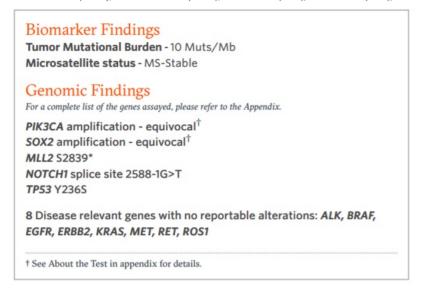


Case Presentation – Dr McKenzie: A 78-year-old man with NSCLC and a TRK fusion (continued)

Summary of Detected Somatic Alterations, Immunotherapy Biomarkers & Associated Treatment Options

KEY Approved in indication Approved in other indication (X) Lack of response						
Detected Alteration(s) / Biomarker(s)	Associated FDA-approved therapies	Clinical trial availability (see page 3)	% cfDNA or Amplification			
TTC28-NTRK3 Fusion	Entrectinib, Larotrectinib	Yes	2.5%			
TP53 Y236S	None	Yes	17.3%			
NOTCH1 Splice Site SNV	None	Yes	16.4%			

Variants of Uncertain Clinical Significance BRCA2 T1040A (21.2%), NOTCH1 N1482K (17.6%), GATA3 T78K (8.0%), ESR1 D321G (0.1%), FANCA R874_F876del (0.3%)



- Novel TRK fusion detected in plasma, but not seen in tissue testing
- Discordance between tissue and plasma is expected
 - Tumor heterogeneity & tumor evolution
 - Test differences
 - Most discrepancies are attributed to biology and not test inferiority
- Both larotrectinib and entrectinib are approved for NTRK fused cancers
 - Indicated for patients who "have no satisfactory alternative treatments or that have progressed following treatment"
- Initiating entrectinib for front-line metastatic treatment

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Case Presentation – Dr Jasani: A 35-year-old woman with newly diagnosed mNSCLC – ALK mutation



Dr Nikesh Jasani

- Presented with cough, chest pain, and fatigue
- Diagnosed with stage IV NSCLC adenocarcinoma with right hilar mass, with metastases to multiple mediastinal nodes, bone and a solitary brain mass
- Molecular studies: ALK FISH +
- Stereotactic RT to brain lesion
- 1/2021: Alectinib \rightarrow excellent response with resolution of all symptoms

Questions

- What is the optimal first-line therapy for NSCLC that is ALK mutation-positive?
- If her disease progresses, what would be the best next steps repeat biopsy and assessment of resistance mutations to determine ideal treatment?
- How do you manage CNS disease in patients with NSCLC that is ALK mutation-positive?
- What are the side effects of note with the newer ALK inhibitors such as Iorlatinib and brigatinib?



Module 1: Role of Genomic Profiling; New Developments in NSCLC with an EGFR Mutation

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- Dr Hussein: A 51-year-old man with metastatic lung adenocarcinoma and an EGFR exon 20 mutation

- Dr Neal: A 51-year-old man with RET-positive NSCLC
- Dr Neal: A 64-year-old woman with metastatic NSCLC and a HER2 exon 20 insertion
- Dr McKenzie: A 78-year-old man with NSCLC and a TRK fusion
- Dr Jasani: A 35-year-old woman with newly diagnosed mNSCLC ALK mutation
- Dr Mohamed: A 71-year-old woman with metastatic adenocarcinoma of the lung with MET exon
 14 skipping and MET amplification mutations PD-L1 TPS 95%
- Dr Zafar: A 64-year-old woman with discordant BRAF mutation testing results



Case Presentation – Dr Mohamed: A 71-year-old woman with metastatic adenocarcinoma of the lung with MET exon 14 skipping and MET amplification mutations – PD-L1 TPS 95%



Dr Mohamed Mohamed

- Never smoker presents with bilateral flank pain and 5-month history of weight loss and fatigue
- Workup reveals Stage IV poorly differentiated adenocarcinoma of the lung; lung obstructions and 2 small, subcentimeter lesions in brain detected
- Molecular analyses: MET exon 14 skipping mutation, MET amplification
- PD-L1: 95%
- Palliative RT to the lung; SRS to the brain
- Capmatinib initiated

Questions

- If she didn't need to receive RT due to the lung obstructions, should I have initiated treatment with targeted therapy alone? Does capmatinib work on brain lesions by itself?
- With a PD-L1 of 95%, should she have been considered for immunotherapy, or should targeted therapy precede that as first-line therapy?
- Are mutations in MET regarded in the same way as mutations in EGFR and ALK in terms of immune therapy? Or are they regarded like BRAF mutations in that you could use immune therapy?



Module 1: Role of Genomic Profiling; New Developments in NSCLC with an EGFR Mutation

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Case Presentation – Dr Zafar: A 64-year-old woman with discordant BRAF mutation testing results

Dr Syed Zafar

- 2020: Diagnosed with metastatic adenocarcinoma with several pulmonary lesions, mediastinal lymphadenopathy, and brain metastases
- Liquid biopsy and NGS ordered
 - Liquid biopsy reveals BRAF V600E mutation
 - NGS results do not reveal any actionable targets
- Patient is symptomatic: Cough, shortness of breath, effusion
- PD-L1-positive
- Considering symptomatology of patient, chemotherapy/IO combination initiated
- Patient's symptoms have improved on treatment
- Holding BRAF-targeted treatment in reserve as potential future therapy



Meet The Professor Optimizing the Clinical Management of Hodgkin and Non-Hodgkin Lymphomas

Wednesday, October 27, 2021 5:00 PM - 6:00 PM ET

Faculty

Jonathan W Friedberg, MD, MMSc

Moderator Neil Love, MD



Thank you for joining us!

CME credit information will be emailed to each participant within 3 business days.

