

HOW DO I TREAT PATIENTS WITH HER2 MUTATIONS, BRAF MUTATIONS, MET MUTATIONS

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DECLARATION OF INTERESTS

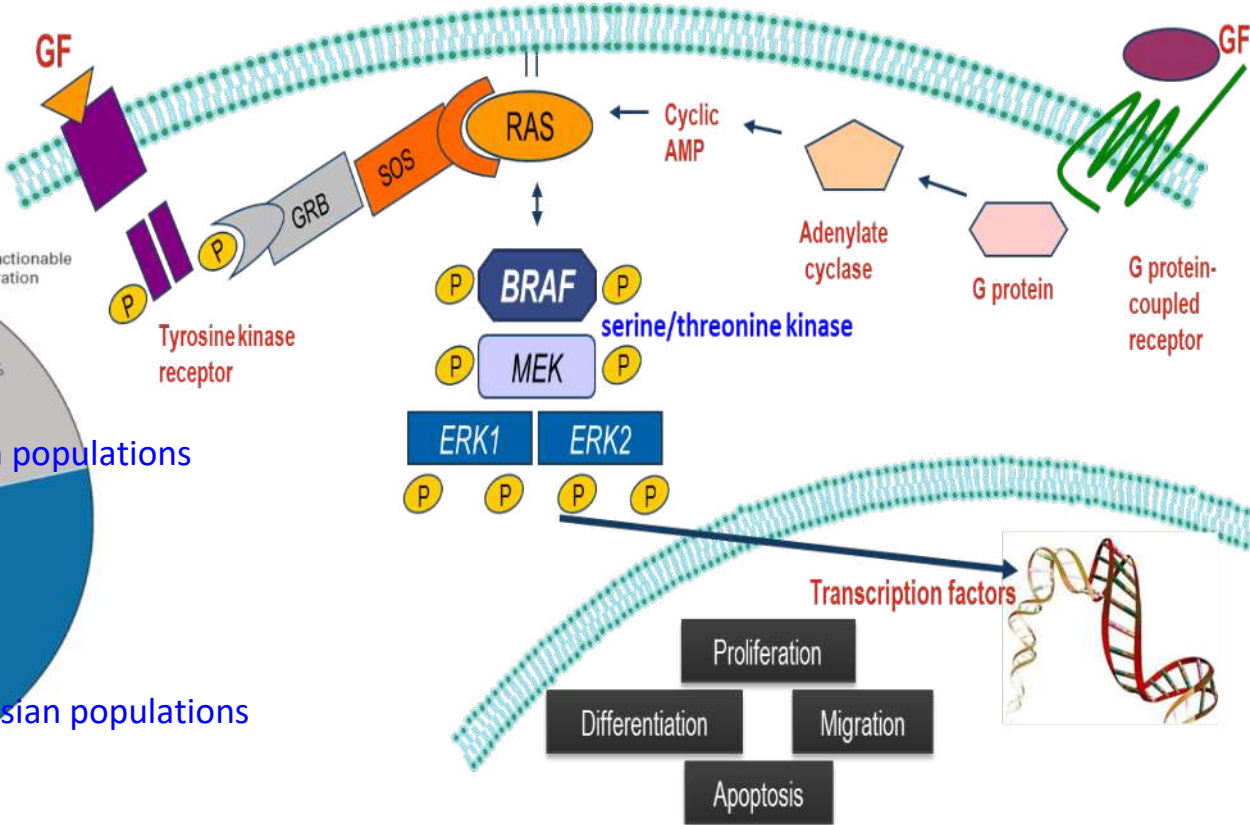
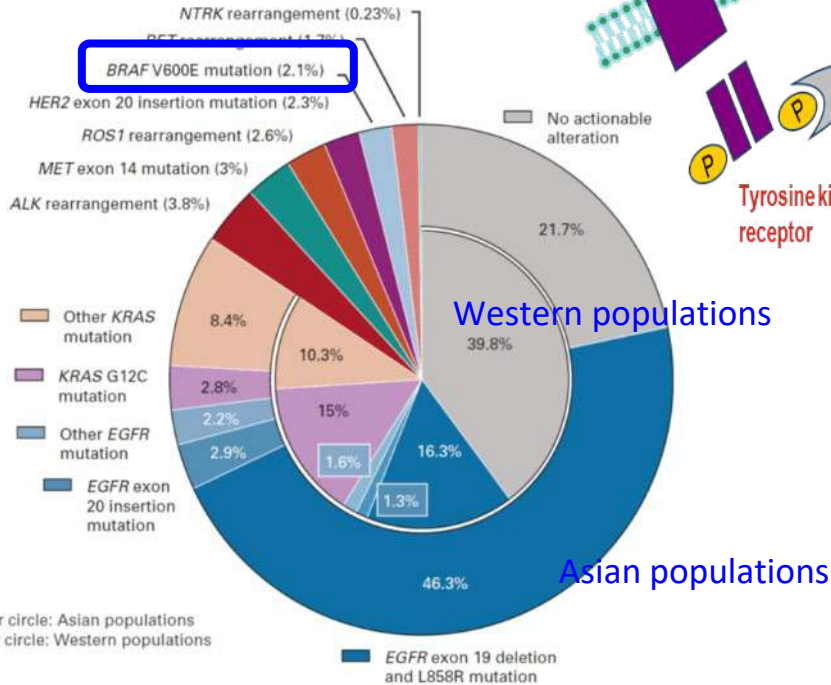
Consulting, advisory role or lectures: AstraZeneca, Bristol-Myers Squibb, Boehringer Ingelheim, Celgene, Daiichi Sankyo, Eli Lilly, Merck, Novartis, Pfizer, prIME Oncology, Peer CME, Roche, Janssen, Abbvie

Honoraria: AstraZeneca, Bristol-Myers Squibb, Boehringer Ingelheim, Celgene, Eli Lilly, Merck, Novartis, Pfizer, prIME Oncology, Peer CME, Roche, Janssen, Abbvie

Clinical trials research as principal or co-investigator (Institutional financial interests): AstraZeneca, Bristol-Myers Squibb, Boehringer Ingelheim, Eli Lilly, Merck, Novartis, Pfizer, Roche, Medimmun, Sanofi-Aventis, Taiho Pharma, Novocure, Daiichi Sankyo, Janssen, Abbvie

Travel, Accommodations, Expenses: AstraZeneca, Roche, Novartis, prIME Oncology, Pfizer

BRAF MUTATIONS IN NSCLC



1. Barlesi F et al. Lancet 2016;387:1415-1426;

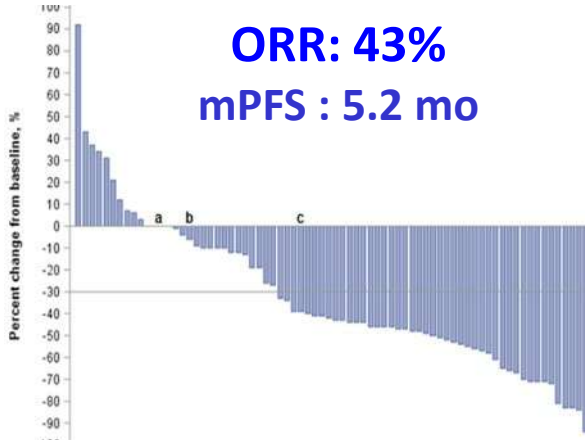
Vemurafenib and Dabrafenib in *BRAF* mutant NSCLC

AcSé trial

Vemurafenib

79 *BRAF*^{V600} NSCLC

ORR: 43%
mPFS : 5.2 mo

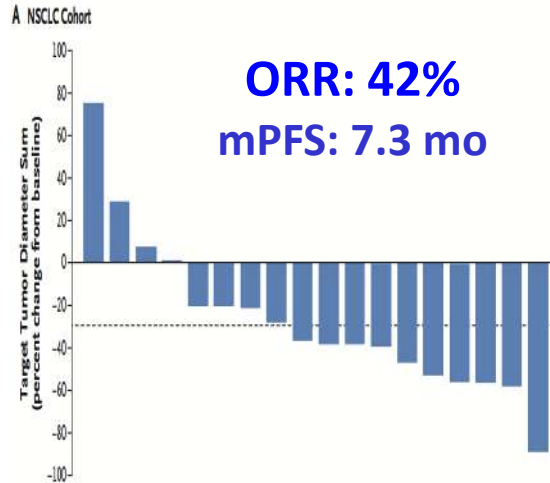


VE-Basket trial

Vemurafenib

20 *BRAF*^{V600} NSCLC

ORR: 42%
mPFS: 7.3 mo

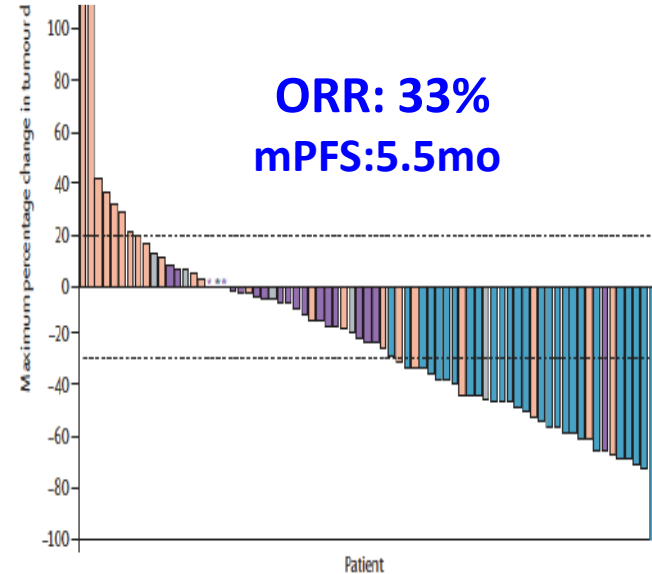


BRF113928 Study

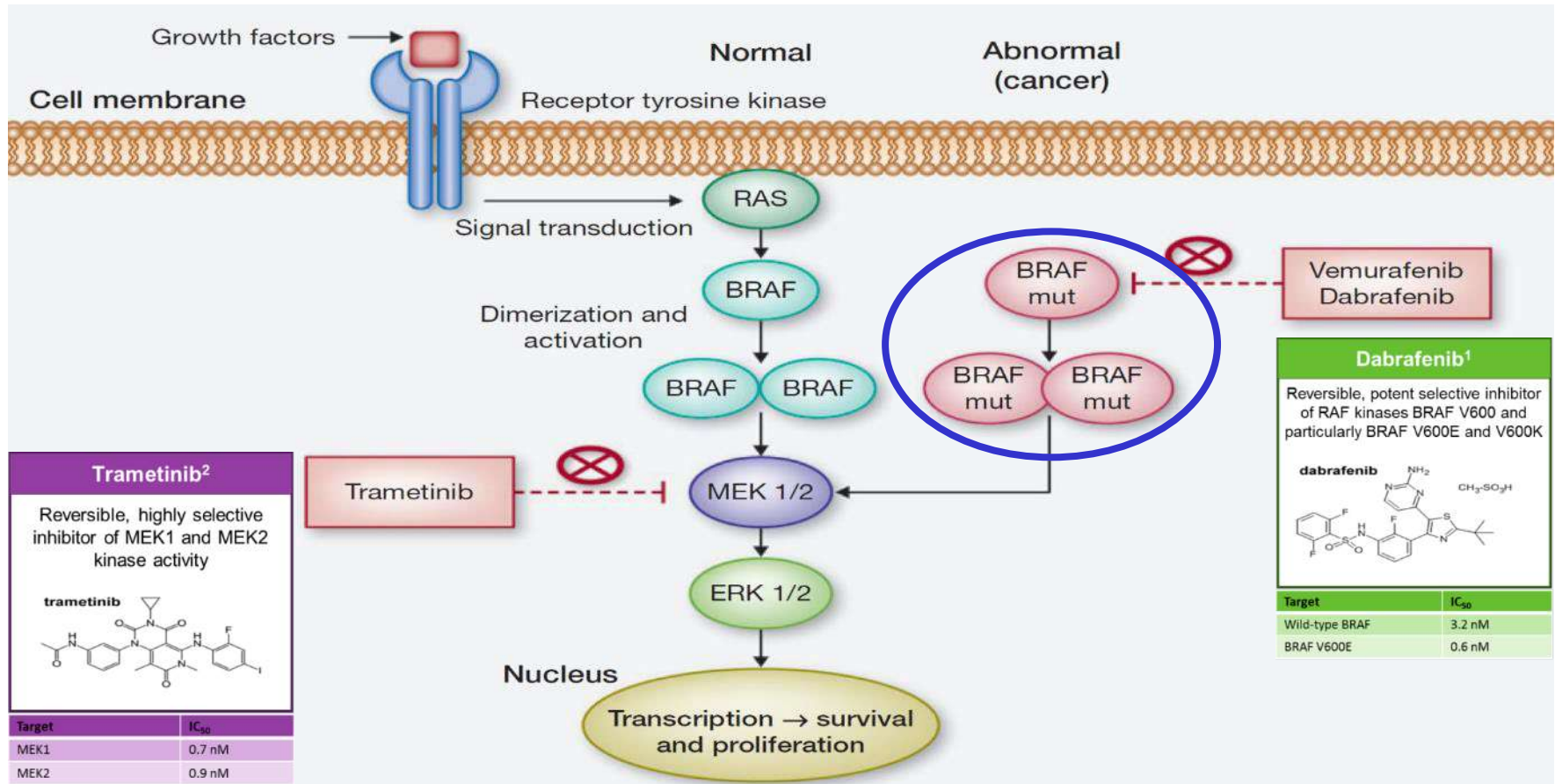
Dabrafenib

84 *BRAF*^{V600E} NSCLC

ORR: 33%
mPFS: 5.5mo



MECHANISM OF ACTION FOR DUAL MAPK PATHWAY INHIBITION WITH DABRAFENIB + TRAMETINIB TO OVERCOME ERK ESCAPE MECHANISM



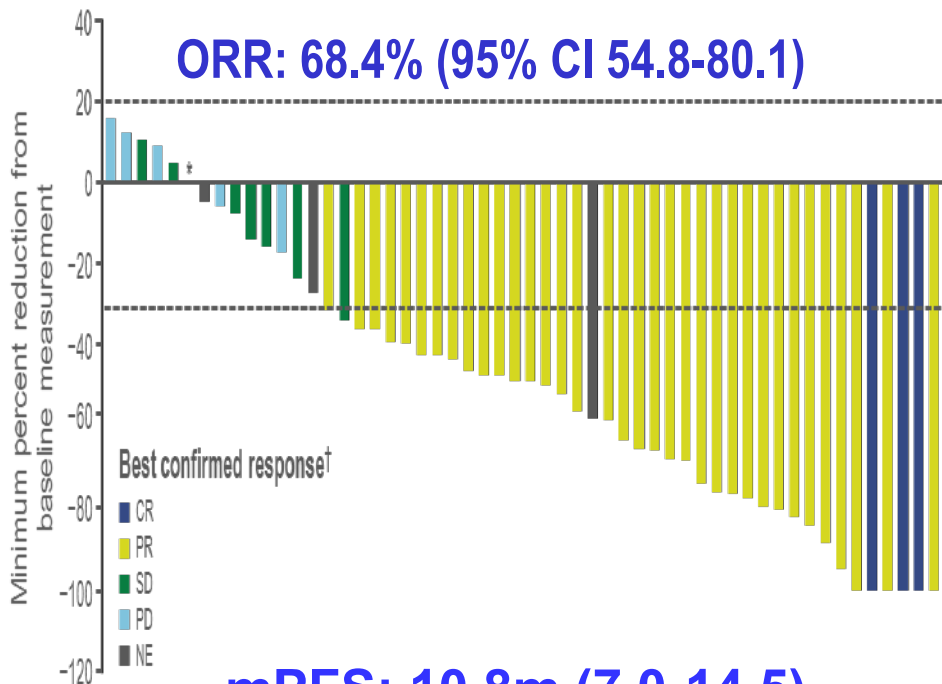
BRF113928 STUDY : MAXIMUM CHANGE IN TARGET LESION BY BEST CONFIRMED RESPONSE WITH DABRAFENIB + TRAMETINIB

Cohort B (N=57 NSCLC BRAF V600E)
2ND LINE

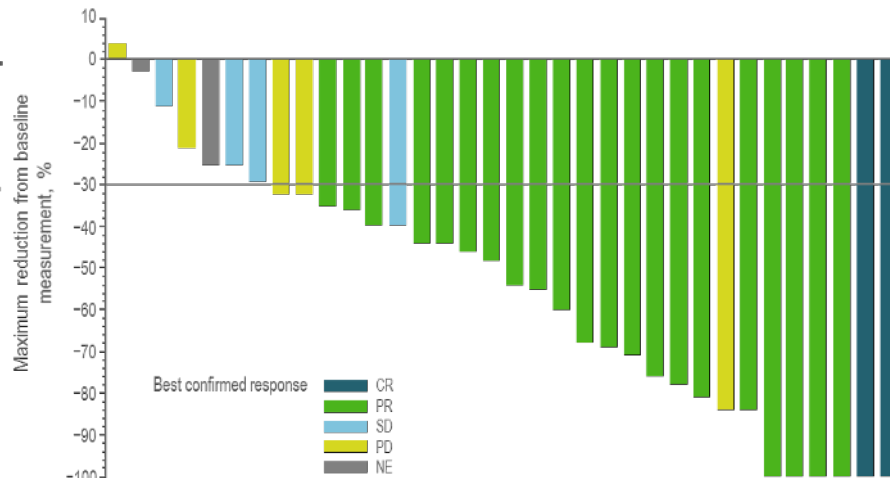
Cohort C (N=36 NSCLC BRAFV600E)
1ST LINE

ORR: 68.4% (95% CI 54.8-80.1)

ORR: 63.9% (95% CI 46.2- 79.2)



mPFS: 10.8m (7.0-14.5)



mPFS: 10.2m (6.9-16.7)

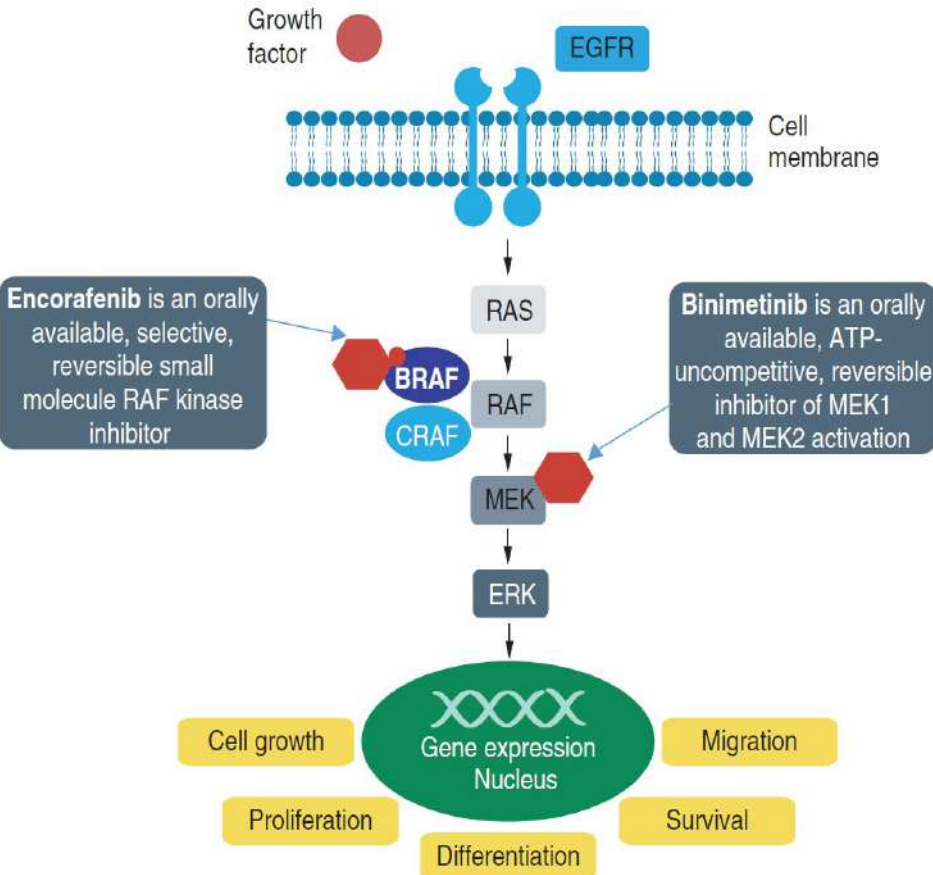
THE SAFETY PROFILE FOR DABRAFENIB, OR DABRAFENIB + TRAMETINIB

Category	AEs, n (%)	Dabrafenib + trametinib 1st line ¹		Dabrafenib + trametinib 2nd line ²		Dabrafenib monotherapy 2nd line ²	
		All grades	Grade 3	All grades	Grade 3	All grades	Grade 3
General	Pyrexia	23 (64)	4 (11)	26 (46)	1 (2)	30 (36)	2 (2)
	Asthenia	4 (11)	1 (3)	18 (32)	2 (4)	25 (30)	3 (4)
	Decreased appetite	12 (33)	0	17 (30)	0	24 (28)	1 (1)
	Chills	9 (25)	0	13 (23)	1 (2)	13 (15)	1 (1)
	Peripheral edema	13 (36)	0	13 (23)	0	–	–
	Arthralgia	5 (14)	1 (3)	11 (19)	0	14 (17)	1 (1)
Skin	Dry skin	12 (33)	0	15 (26)	1 (2)	19 (23)	0
	Rash	8 (22)	1 (3)	12 (21)	1 (2)	17 (20)	1 (1)
	Hyperkeratosis	–	–	6 (10)	1 (2)	25 (30)	1 (1)
	Basal-cell carcinoma	–	–	2 (2)	1 (2)	4 (5)	4 (5)
	Squamous-cell carcinoma	–	–	2 (4)	2 (4)	10 (12)	10 (12)
	Skin papilloma	–	–	–	–	22 (26)	0
Digestive	Nausea	20 (56)	0	23 (40)	0	23 (27)	1 (1)
	Vomiting	12 (33)	3 (8)	20 (35)	0	17 (20)	1 (1)
	Diarrhea	13 (36)	1(3)	19 (33)	1 (2)	14 (17)	1 (1)

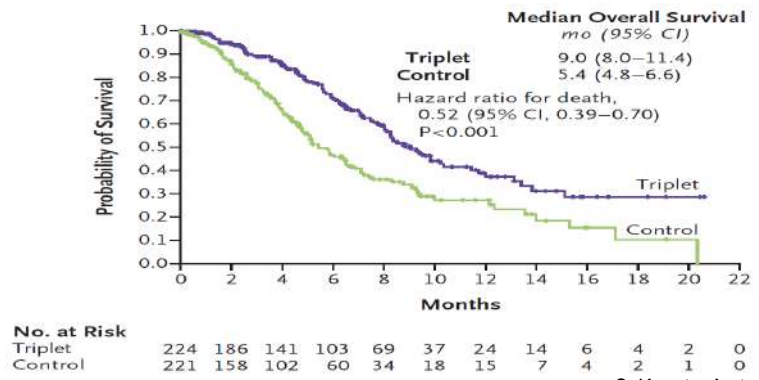
1. Planchard D *et al. Lancet Oncol* 2017;18:1307–1316;

2. Planchard D *et al. Lancet Oncol* 2016;17:984–993

The combination of encorafenib plus binimetinib

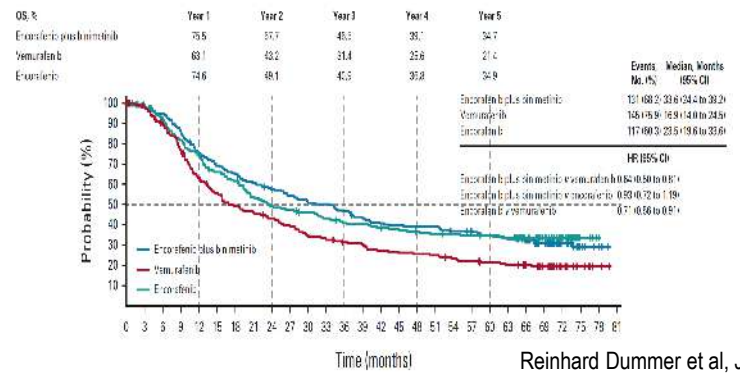


Encorafenib, Binimetinib, and Cetuximab in BRAF V600E-Mutated Colorectal Cancer



S. Kopetz, A et al, NEJM 2019

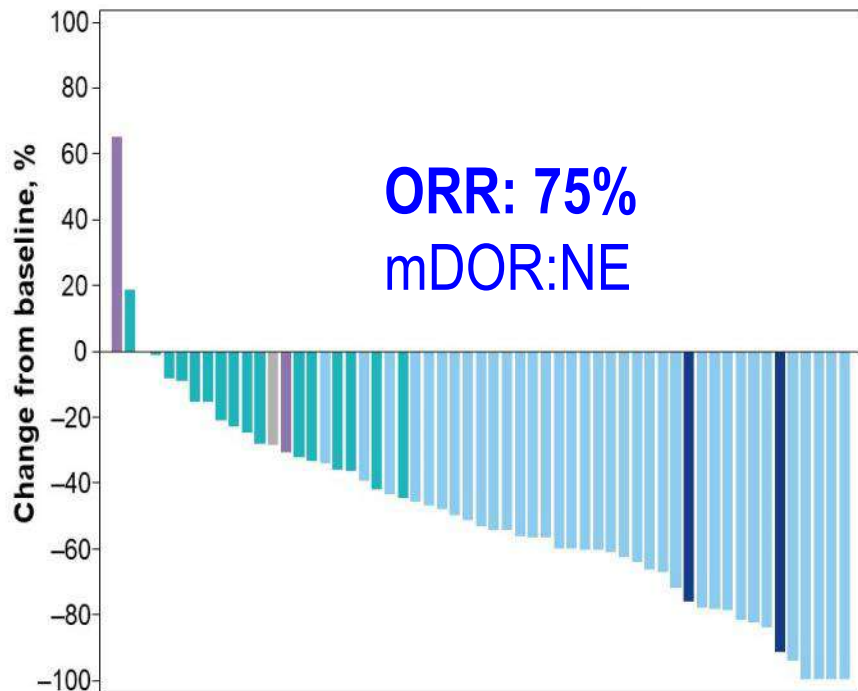
COLUMBUS: Encorafenib Plus Binimetinib Versus Vemurafenib or Encorafenib in Patients With BRAF V600-Mutant Melanoma



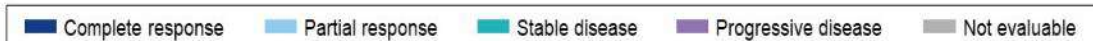
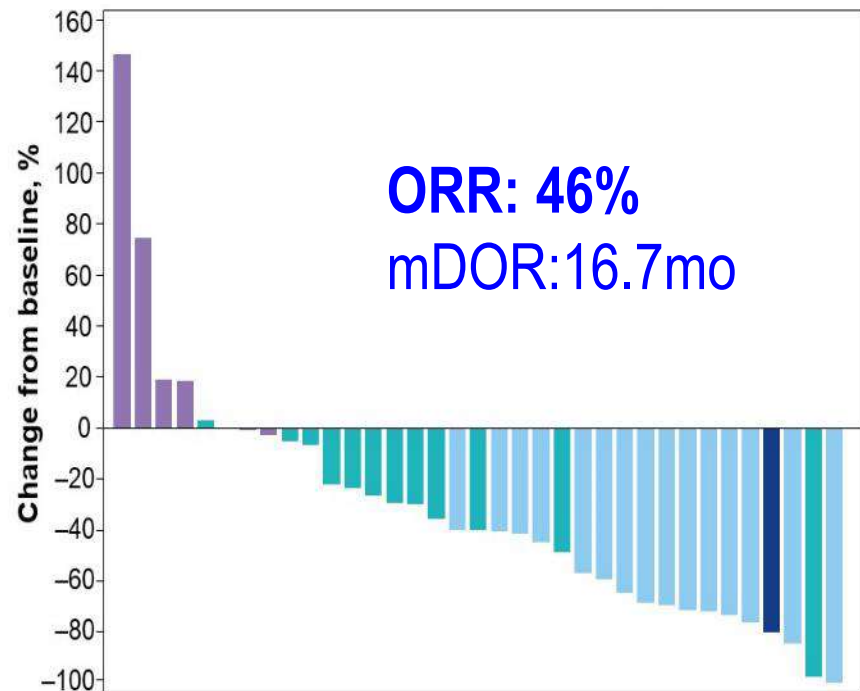
Encorafenib + Binimetinib in metastatic BRAF-V600E NSCLC

PHAROS Trial (Phase 2 Study)

Treatment naive (n=59)

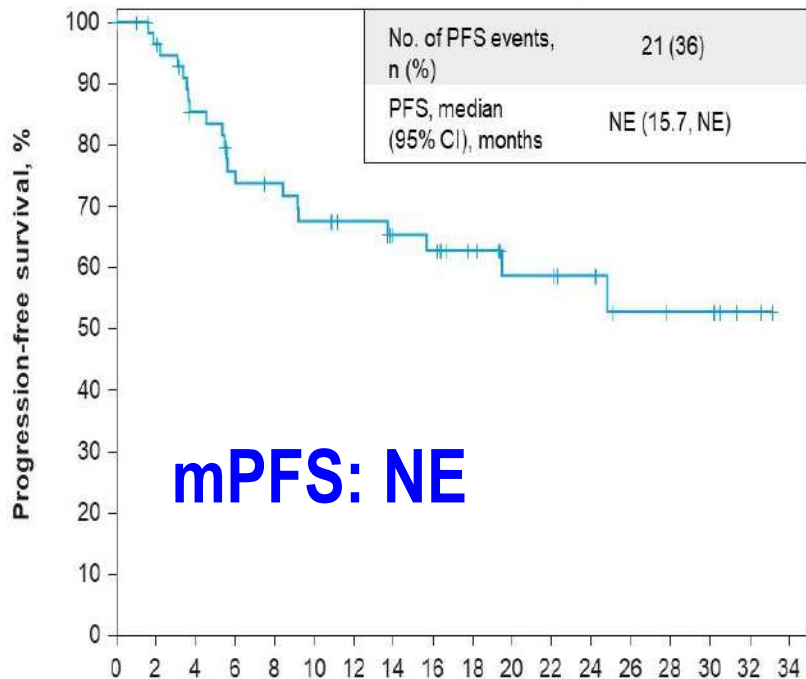


Previously treated (n=39)



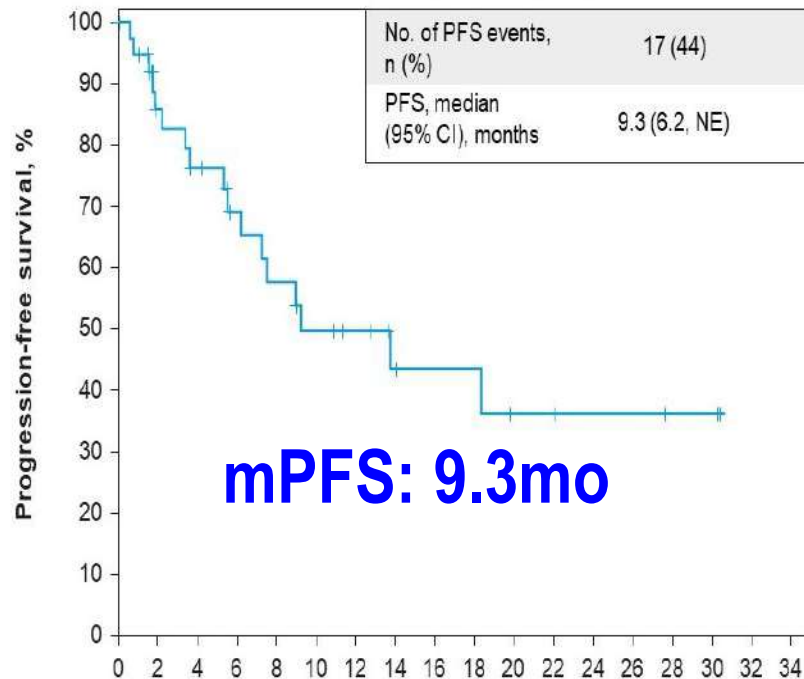
PFS by IRR

Treatment naïve (n=59)



No. at risk	0	2	4	6	8	10	12	14	16	18	20	22	24	26	28	30	32	34
Treatment naïve	59	54	45	38	36	33	30	26	25	19	14	14	12	8	7	7	2	0

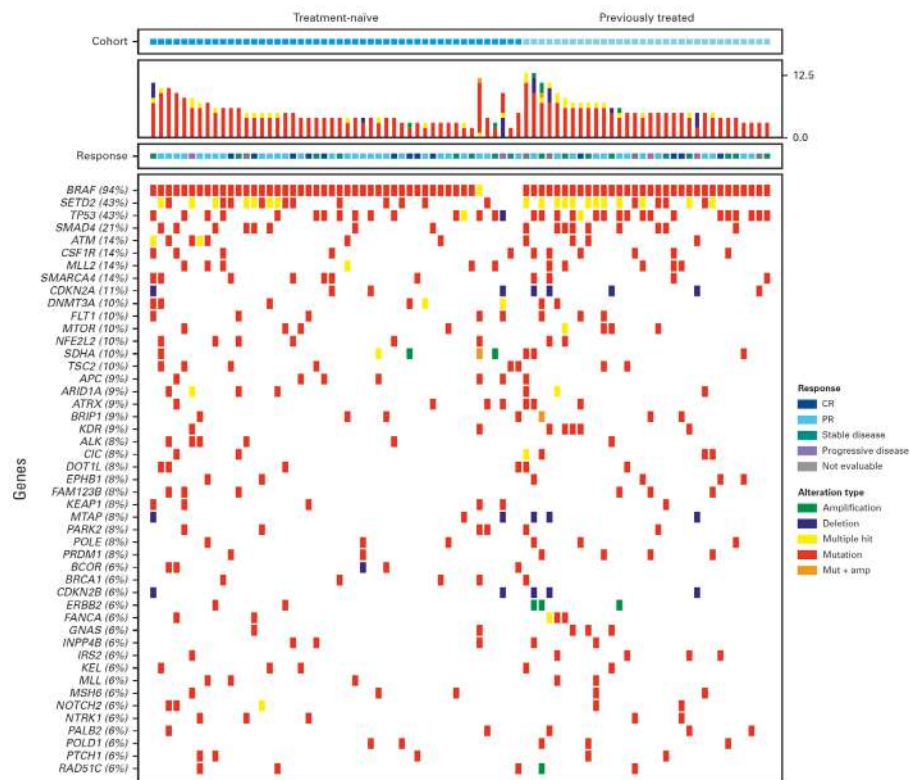
Previously treated (n=39)



No. at risk	0	2	4	6	8	10	12	14	16	18	20	22	24	26	28	30	32	34
Previously treated	39	27	23	18	15	12	10	7	6	6	4	4	3	3	2	2	0	0

Incidence of TRAEs of Any Grade $\geq 10\%$ Tumor molecular alterations in baseline biopsy

AE Preferred Term	Overall (N = 98)		
	Any Grade	Grade 3	Grade 4
Any TRAEs No. (%)	92 (94)	37 (38)	3 (3) ^a
Nausea	49 (50)	3 (3)	0
Diarrhea	42 (43)	4 (4)	0
Fatigue	31 (32)	2 (2)	0
Vomiting	28 (29)	1 (1)	0
Anemia	18 (18)	3 (3)	0
Vision blurred	17 (17)	1 (1)	0
Constipation	13 (13)	0	0
ALT increased	12 (12)	5 (5)	0
AST increased	12 (12)	7 (7)	0
Pruritus	12 (12)	0	0
Blood creatine phosphokinase increased	11 (11)	0	0
Peripheral edema	11 (11)	0	0
Abdominal pain	10 (10)	0	0
Alopecia	10 (10)	0	0
Asthenia	10 (10)	3 (3)	0
Dry skin	10 (10)	0	0



Most frequent genomic alterations identified at baseline:

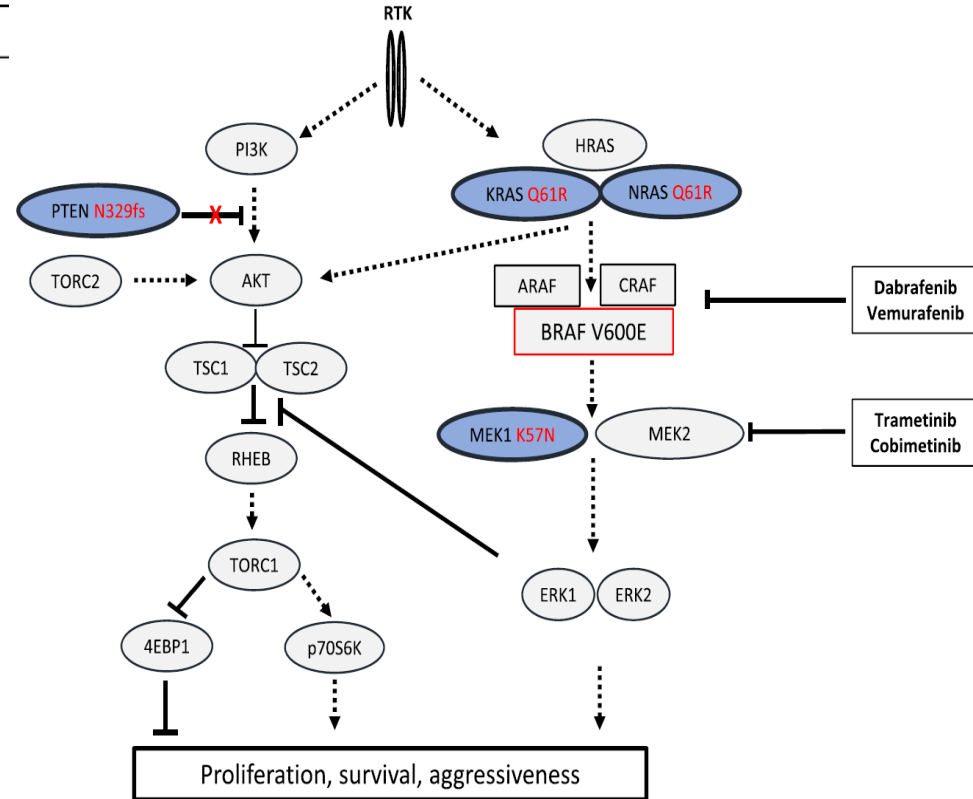
SETD2 and TP53 (43% each), SMAD4 (21%), ATM, MLL2, CSF1R, SMARCA4 (14% each), and CDKN2A (11%).

None of these alterations associated with outcome

PI3K-AKT-mTOR and RAS-RAF-MEK pathways

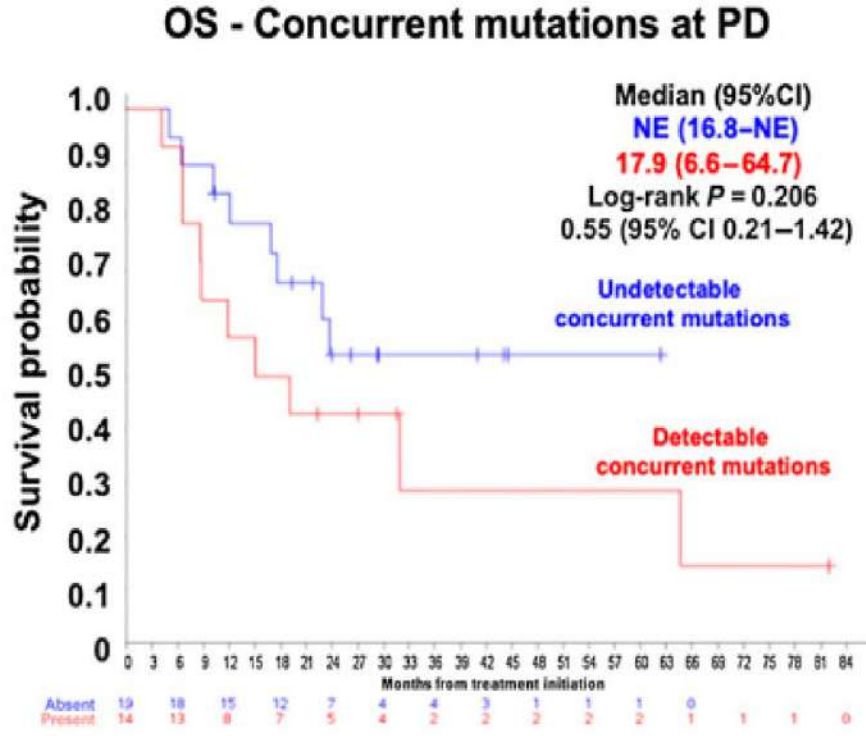
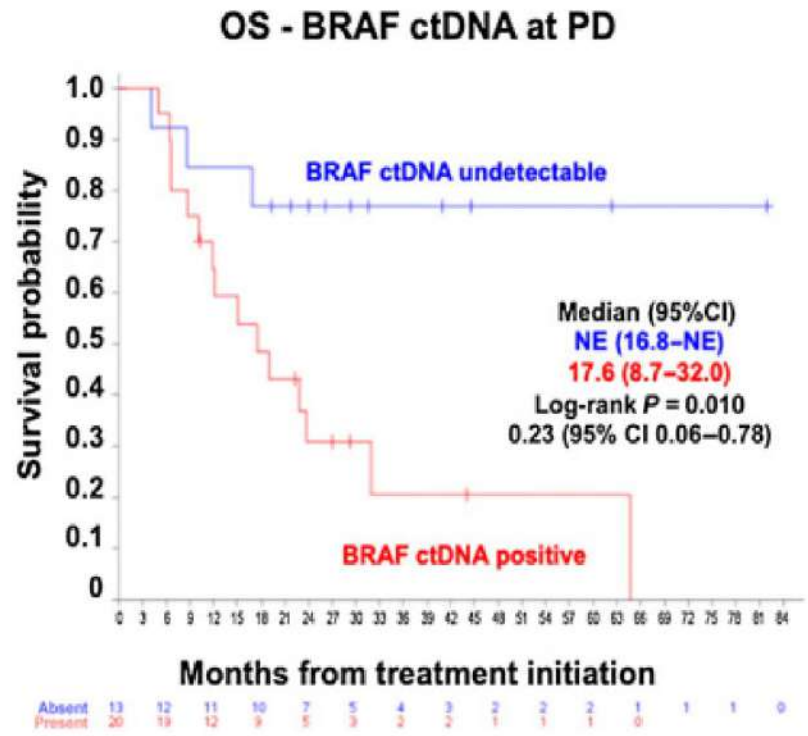
WES, RNA sequencing (RNAseq) (Illumina Integragen) and comparative genomic hybridisation (CGH) array

Patient	Tumour cell proportion	Targeted NGS	WES	RNAseq	CGH array	TMB (mut/Mb)
MR113	50%	BRAF V600E MEK1 K57N	BRAF V600E MEK1 K57N <i>FANCD2</i> Q706* <i>LRP1B</i> P203Q <i>GRIN2A</i> M960I <i>RARA</i> T285I	<i>ZNF354C-RGS7</i>	No hints	3.04
MR159	30%	BRAF V600E PTEN N329fs	BRAF V600E PTEN N329fs <i>SETD2</i> D1537Ifs <i>SETD2</i> T1171Kfs <i>MEN1</i> G230V <i>PTCD3</i> E114Rfs <i>ATXN1</i> P485A <i>AKAP6</i> D668G	No fusion transcripts	No hints	1.42
MR279	50%	BRAF V600E AKT1 E17K NRAS Q61R	BRAF V600E AKT1 E17K (<i>see baseline NGS</i>) <i>ERBB4</i> S303Y <i>SETD2</i> G1081Vfs	No fusion transcripts	No hints	1.71
MR372	50%	BRAF V600E KRAS Q61R TP53 R280I	BRAF V600E KRAS Q61R TP53 R280I <i>KMT2E</i> L1610Ffs <i>MPL</i> Q247Sfs <i>ZFH33</i> S2515* <i>MED12</i> Q2160* <i>ARID1A</i> F1809fs <i>KMT2A</i> S754F <i>KMT2A</i> S2319C <i>MYOD1</i> R281C <i>SETD2</i> C1520F <i>NCOR1</i> L866V	No fusion transcripts	No hints	3.75



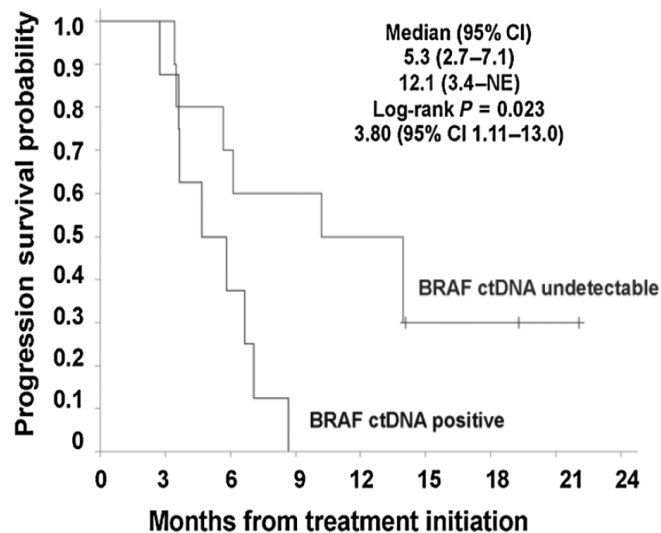
OS according to BRAF-mutant ctDNA status or concurrent genomic alterations, at PD

Prospective cohort of 78 BRAF-mutant NSCLC patients (N=208 samples)



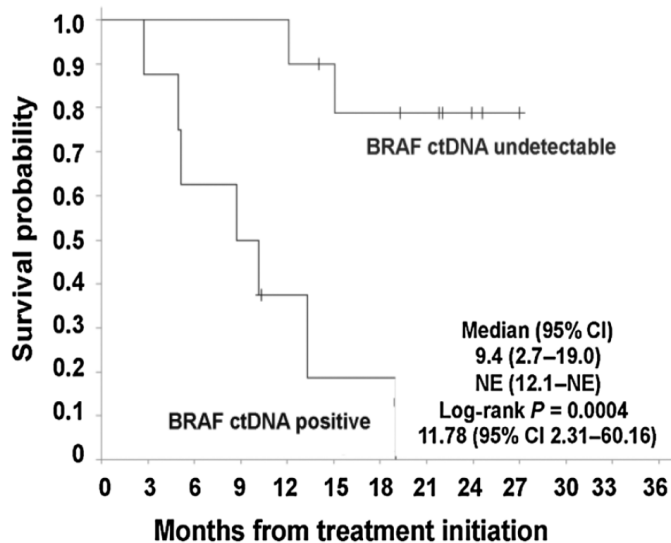
PFS and OS by ctDNA status at the first radiographic evaluation (<100 days after start of targeted therapy)

PFS - CLEARENCE TOTAL ctDNA

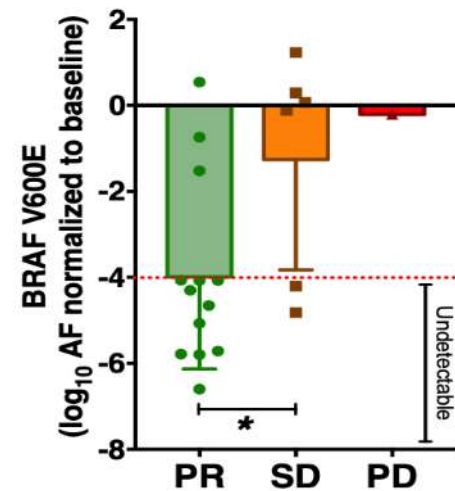


POS	8	3	0		
UND	10	7	5	2	0

OS - CLEARENCE TOTAL ctDNA



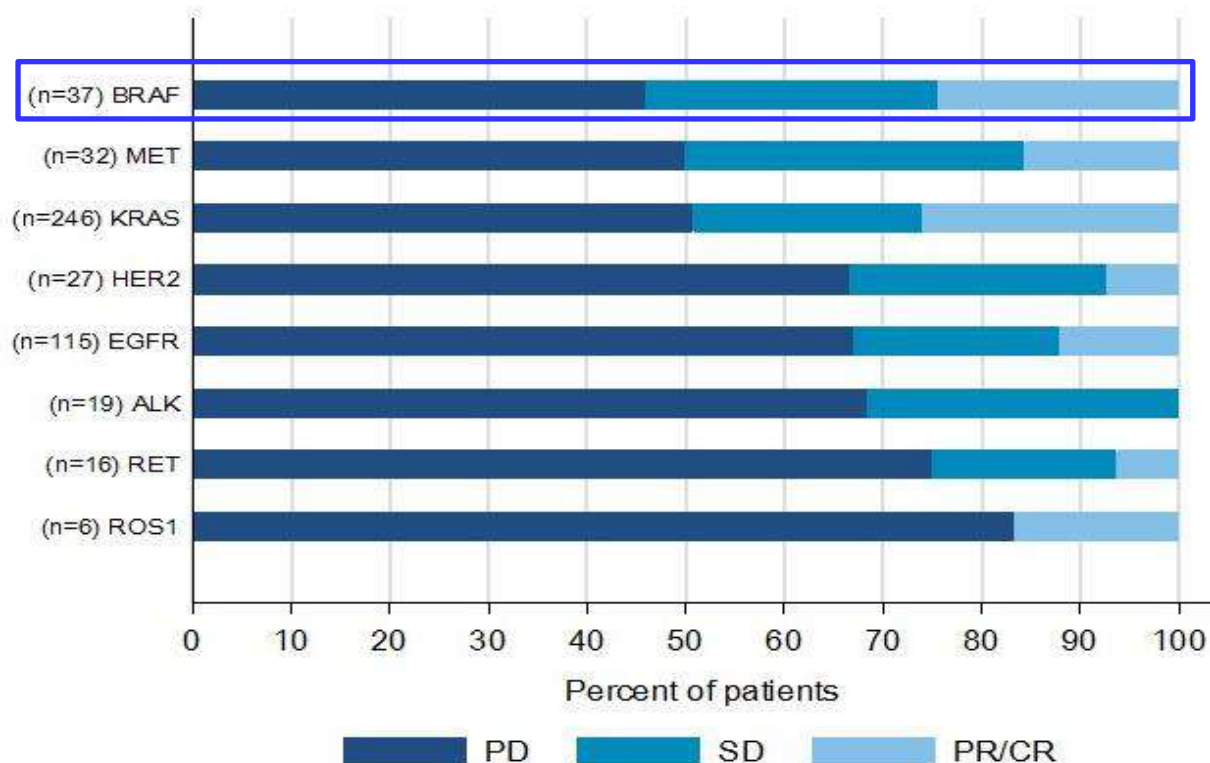
POS	8	5	2	1	0	
UND	10	10	10	7	2	0



Complete clearance of BRAF V600E at the first CT-scan evaluation* in 12/20 (60%)

IMMUNOTARGET registry (BRAF, n=37 pts)

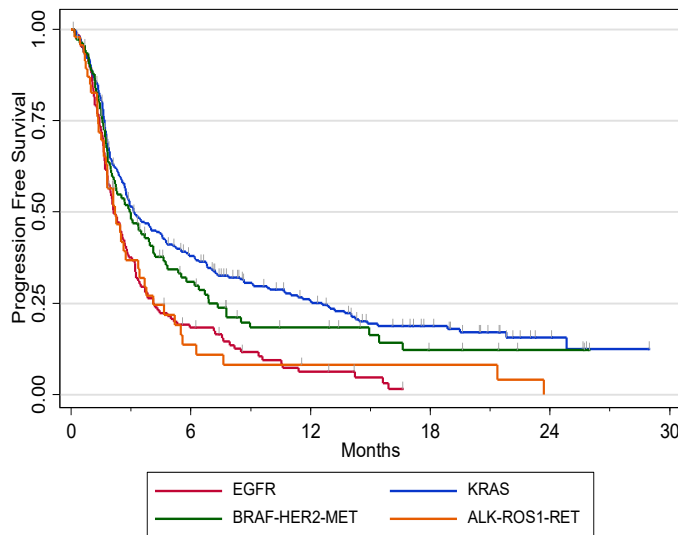
Driver	PD	SD	CR/PR
BRAF	46%	30%	24%
MET	50%	34%	16%
KRAS	51%	23%	26%
HER2	67%	26%	7%
EGFR	67%	21%	12%
ALK	68%	32%	0
RET	75%	19%	6%
ROS1	83%	0	17%
TOTAL	57%	24%	19%



IMMUNOTARGET COHORT: PFS

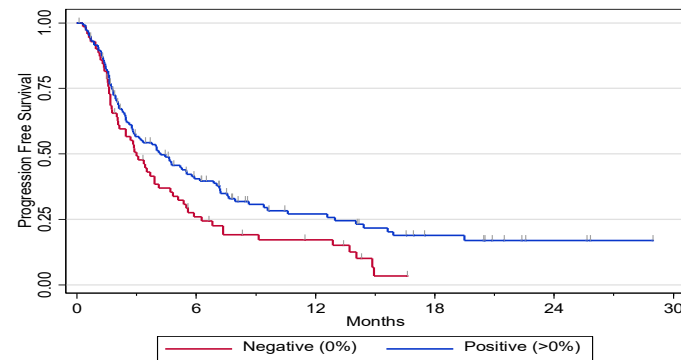
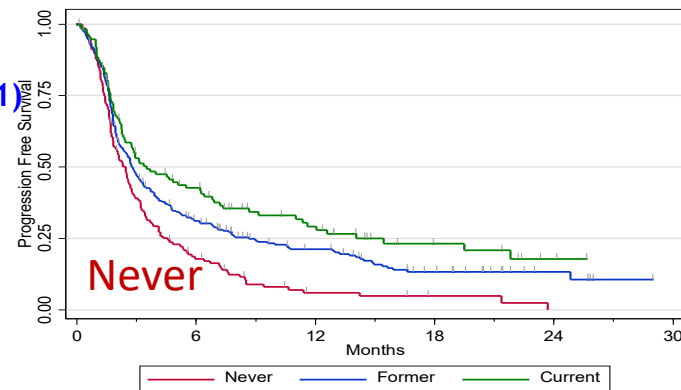
Driver	PFS (months)	
KRAS	3.2	3.2
EGFR	2.1	2.1
BRAF	3.1	
MET	3.4	2.9
HER2	2.5	
ALK	2.5	
RET	2.1	2.2
ROS1	-	
TOTAL	2.8	

PFS according to driver alteration (p < 0.001)



Median follow-up 16.1 months

PFS by smoking (p < 0.001)



PFS by PDL1 (p = 0.02)

BRAF non V600 cohort (AcSé Vemu)

Non V600 mutations

n = 17

G466A : n=1

G466V : n=3

G469A : n=3

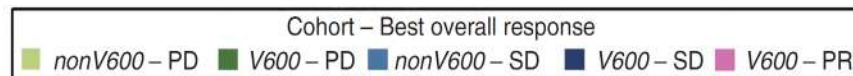
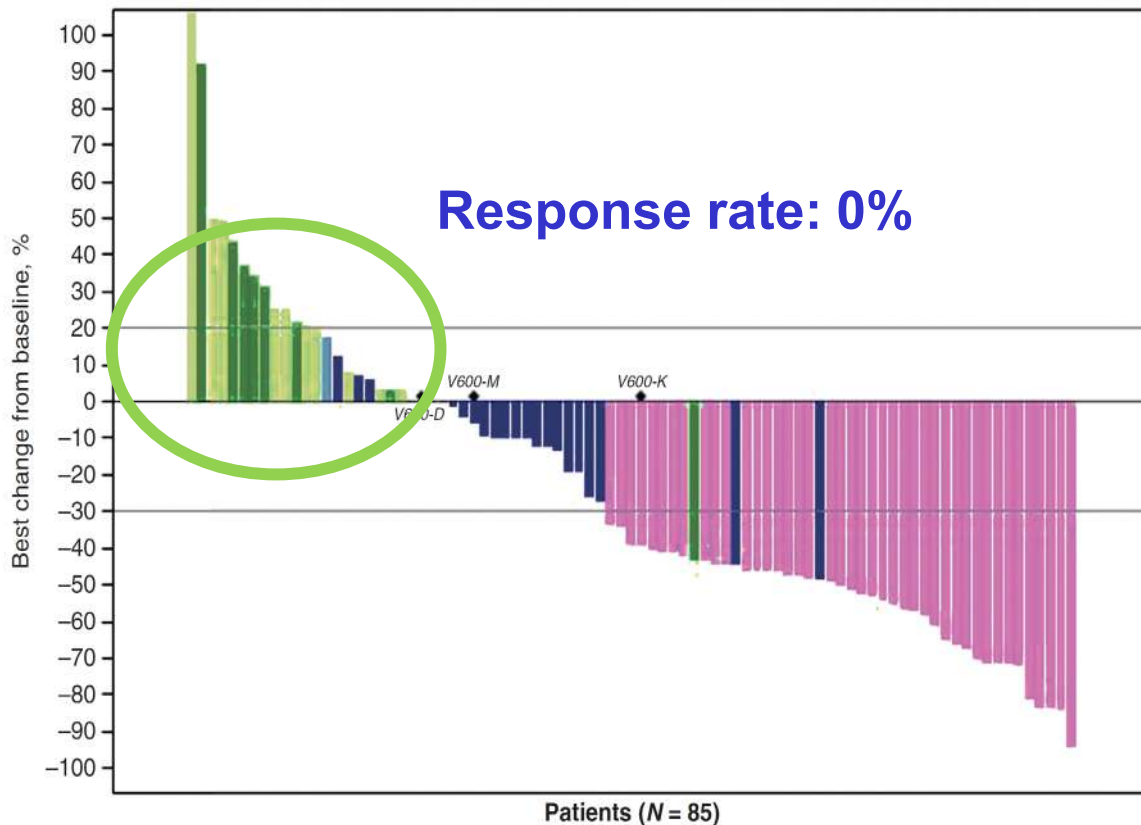
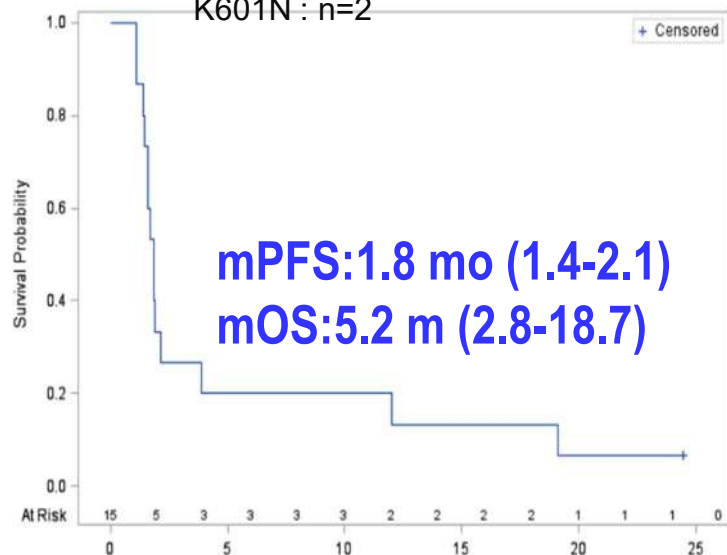
G469V : n=1

N581S : n=3

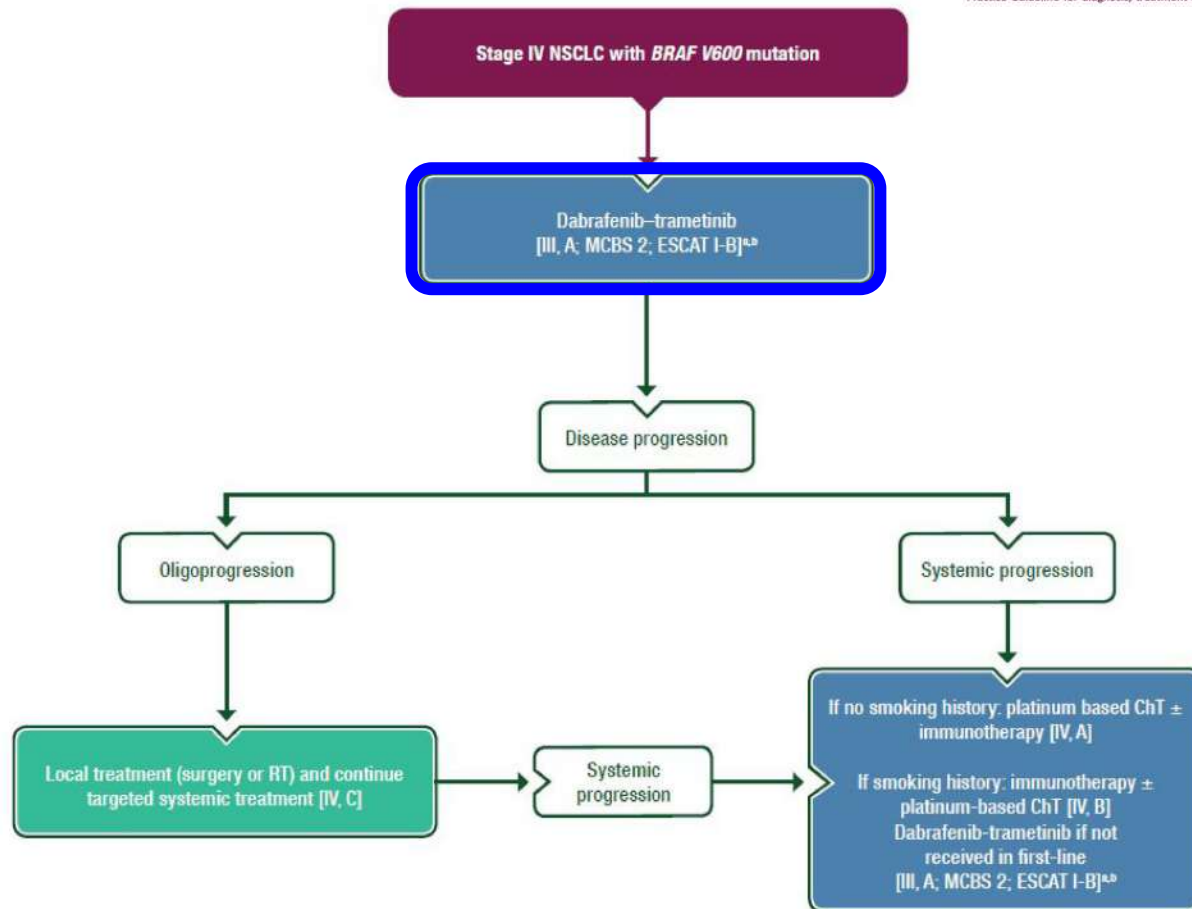
G596R : n=1

K601E : n=3

K601N : n=2

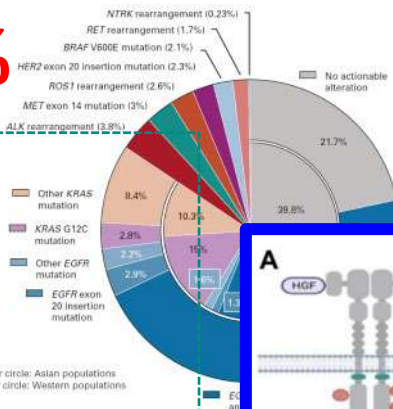


BRAFV600mut NSCLC



MET Ex14 mutant NSCLC

≈2%



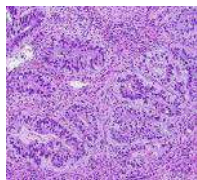
Clinical characteristics



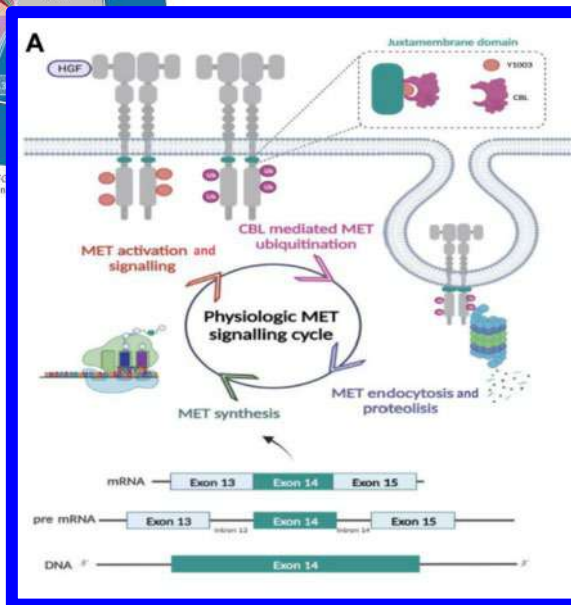
Median age 75 years



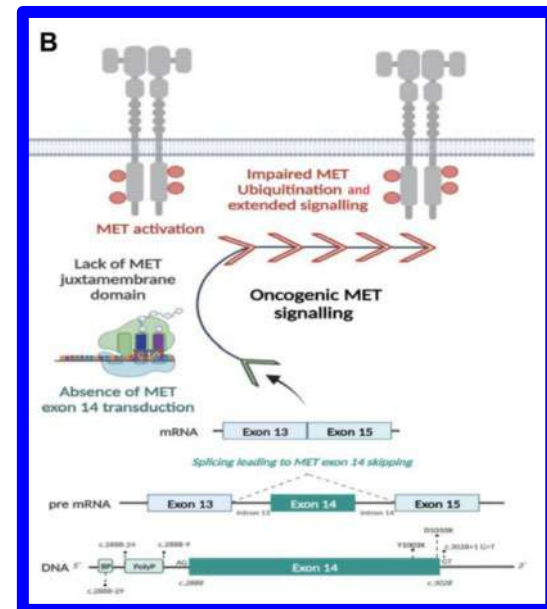
Never smokers, 50%



Non-Squamous: ~ 80%
(Sarcomatoid: 20% METex¹⁴)
Squamous: ~ 10%



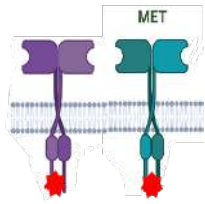
MET exon 14 skipping mutation



MET Agents – A Broad Therapeutic Landscape

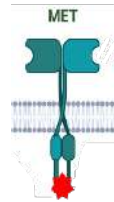
MET KINASE INHIBITORS

MULTI-TARGETED



CRIZOTINIB
CABOZANTINIB
GLESATINIB (MGCD265)
MERESTINIB (LY2801653)
SAR125844
FORETINIB (GSK1363089)
ELZOvantinib (TPX-022)

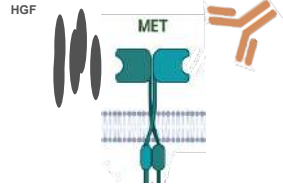
SELECTIVE



TEPOTINIB
CAPMATINIB
SAVOLITINIB
BOZITINIB (APL-101)
GLUMETINIB (SCC244)

MONOCLONAL ANTIBODIES (MET/HGF)

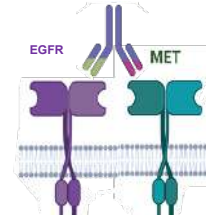
MONOSPECIFIC



HGF
FICLATUZUMAB (AV299)
RILOTUMUMAB (AMG-102)

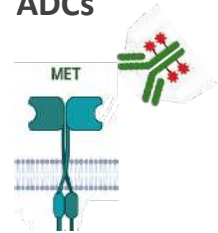
MET
HLX55

POLYSPECIFIC



EGFR-MET
AMIVANTAMAB
EMB-01
METx MET
Sym015
UBAMATAMAB (REGN5093)
EMIBETUZUMAB
(LY2875358)
EGFR-MET-MET
GB263T

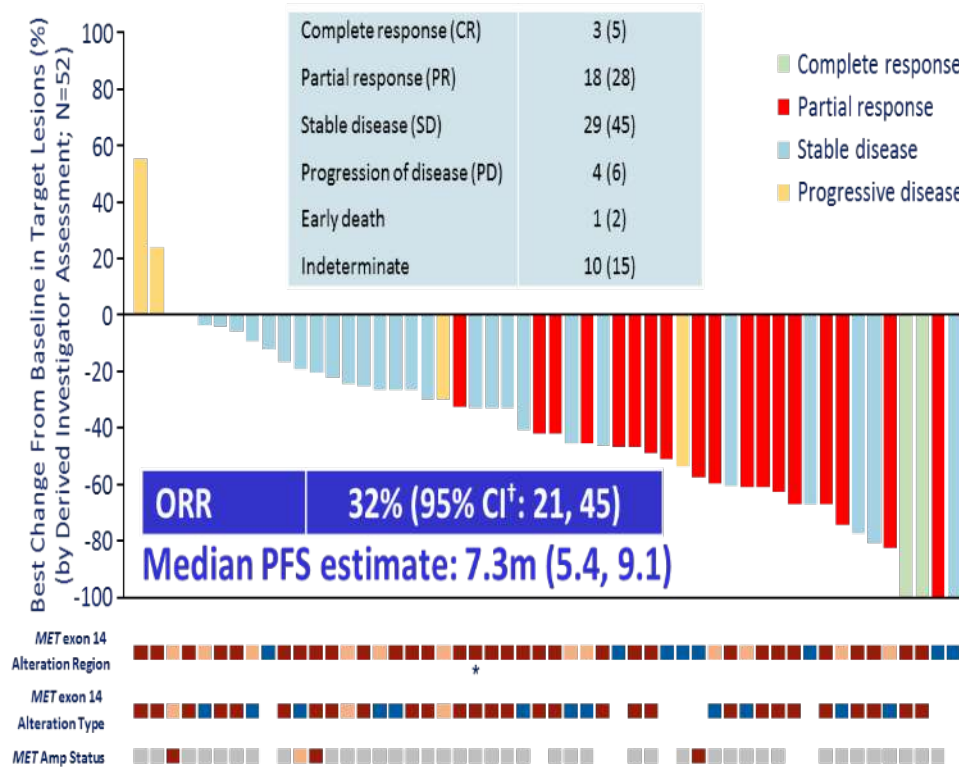
ADCs



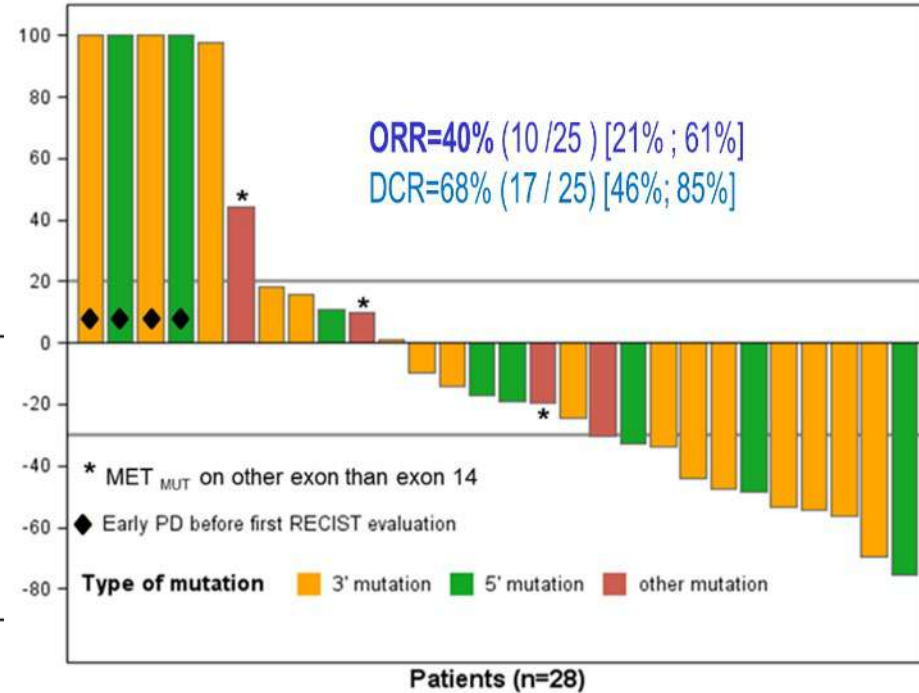
TELISO-V (ABBV-399)
SHR-A1403
TR1801-ADC

Antitumor Activity of Crizotinib MET Exon 14-Altered NSCLC

Profile 1001



AcSé trial (crizotinib), MET exon 14 mutation



VISION Study (Tepotinib)

Selective MET Kinase Inhibitor

- Tepotinib is a highly selective, ATP-competitive, reversible, potent MET tyrosine kinase inhibitor (TKI)

- IC_{50} ~1.7 nM
- At 1 μ M, only MET is inhibited out of a panel of over 300 kinases

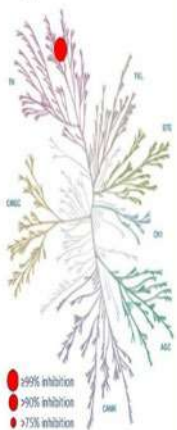
- No MTD reached at 1400 mg QD; RP2D is 500 mg QD

- Preclinical brain penetration

- High binding to rat brain tissue ($f_{u,br}$ = 0.4%)
- The $K_{p,brain}$ (ratio of free brain vs plasma concentration) in rats was 0.25, i.e. 25% of free tepotinib levels in brain, relative to levels found in plasma

- Complete brain and systemic response lasting almost 1 year in patient with NSCLC harboring MET-RB1 translocation treated with tepotinib as compassionate use (Dr Marie Florencia, MD, and Dr Raafat Alameddine at CHUM Montreal, Canada)

Tepotinib kinome¹



Key inclusion criteria

- Advanced NSCLC (EGFR/ALK wild-type, all histologies)
- Central confirmation of METex14 skipping by liquid and/or tissue biopsy
- First, second, or third line of therapy

Cohort A[‡]
METex14 skipping
(primary)

Cohort C[‡]
METex14 skipping
(confirmatory)

Tepotinib
500 mg[‡]
once
daily

Selected endpoints

Primary:

Objective response by IRC (RECIST v1.1)

Secondary:

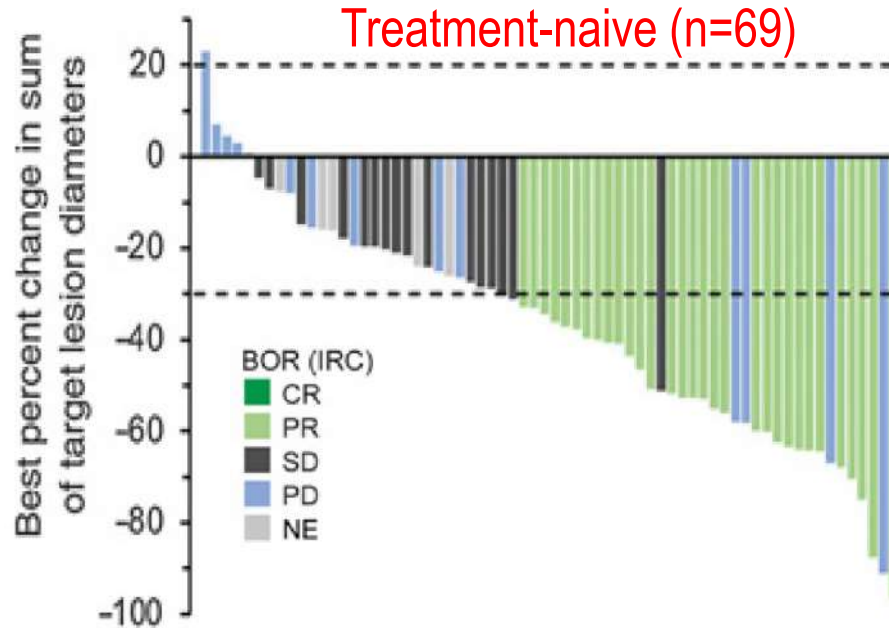
DOR, PFS, OS, Safety

Exploratory RANO-BM analysis[§]:

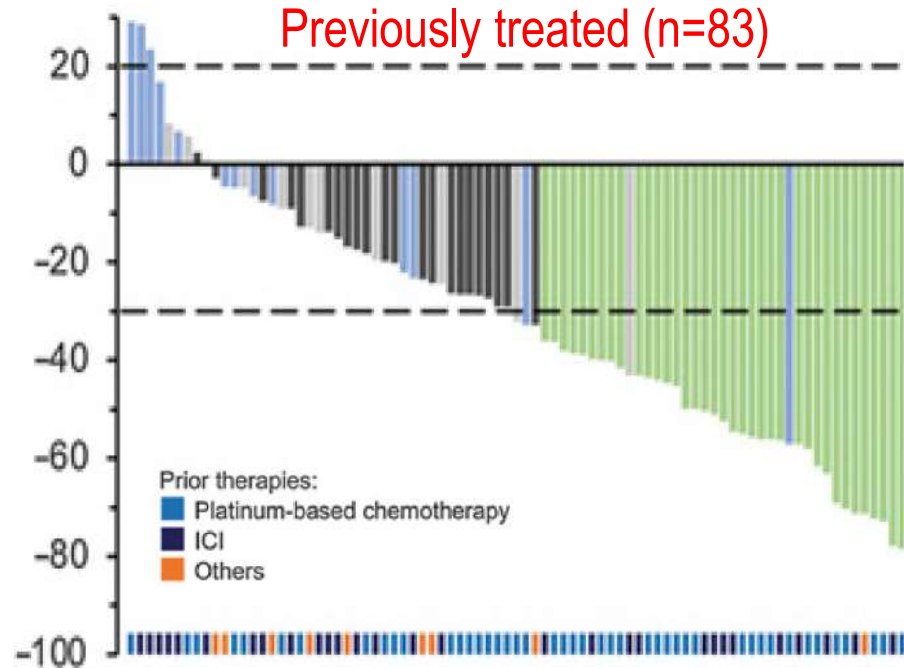
BOR per RANO-BM (patients with ≥ 1 evaluable post-baseline tumor assessment): disease control was defined as CR/PR/SD, or non-CR/non-PD

METex14 mutation: Tepotinib

VISION Study (cohort A)



ORR: 44.9%
mPFS: 8.5mo



ORR: 44.6%
mPFS: 10.9mo

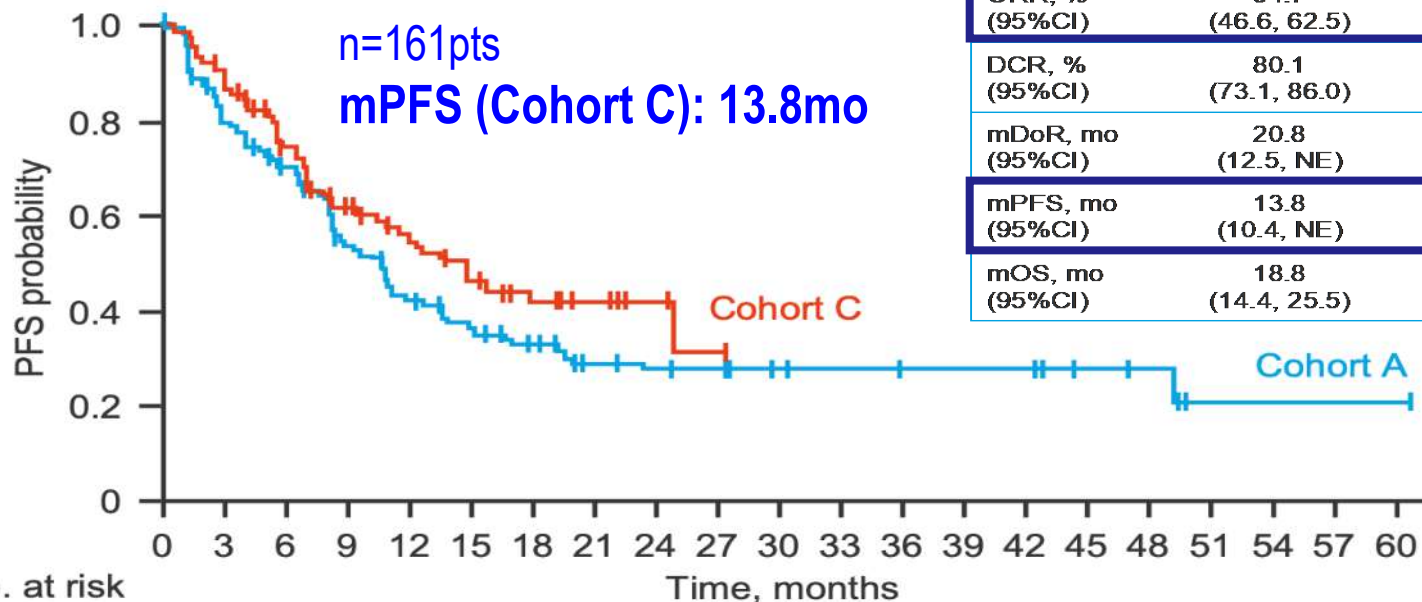
METex14 mutation: Tepotinib

VISION Study (confirmatory cohort C)

ORR: 54.7%

62.3%

51%



Response in Cohort C	All (n=161)	1L (n=69)	2L (n=51)
ORR, % (95%CI)	54.7 (46.6, 62.5)	62.3 (49.8, 73.7)	51.0 (36.6, 65.2)
DCR, % (95%CI)	80.1 (73.1, 86.0)	87.0 (76.7, 93.9)	82.4 (69.1, 91.6)
mDoR, mo (95%CI)	20.8 (12.5, NE)	NE (10.4, NE)	12.6 (4.3, NE)
mPFS, mo (95%CI)	13.8 (10.4, NE)	15.9 (10.8, NE)	13.8 (6.9, NE)
mOS, mo (95%CI)	18.8 (14.4, 25.5)	22.7 (12.7, NE)	19.6 (14.6, NE)

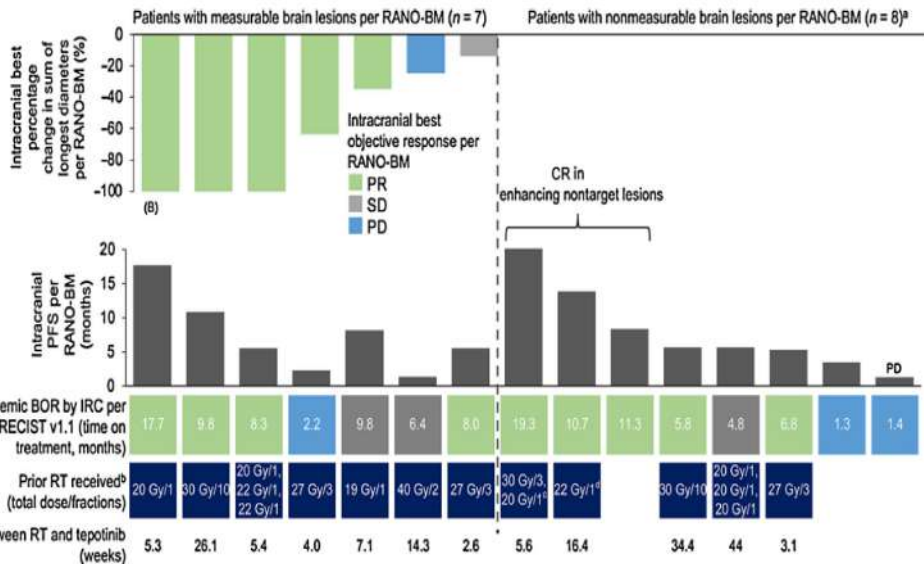
No. at risk

	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45	48	51	54	57	60
Cohort C	161	127	93	56	38	23	16	11	5	2	0	0	0	0	0	0	0	0	0	0	0
Cohort A	152	113	88	59	44	34	27	20	18	17	11	9	8	8	8	5	4	1	1	1	1

Intracranial activity of Tepotinib (METex14, VISION trial)

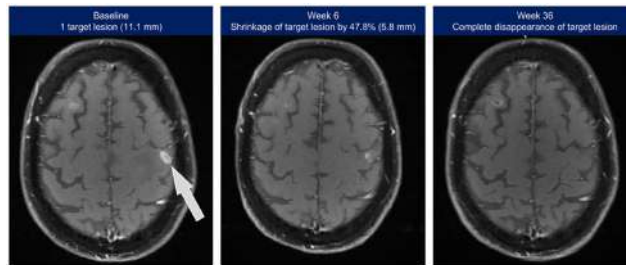
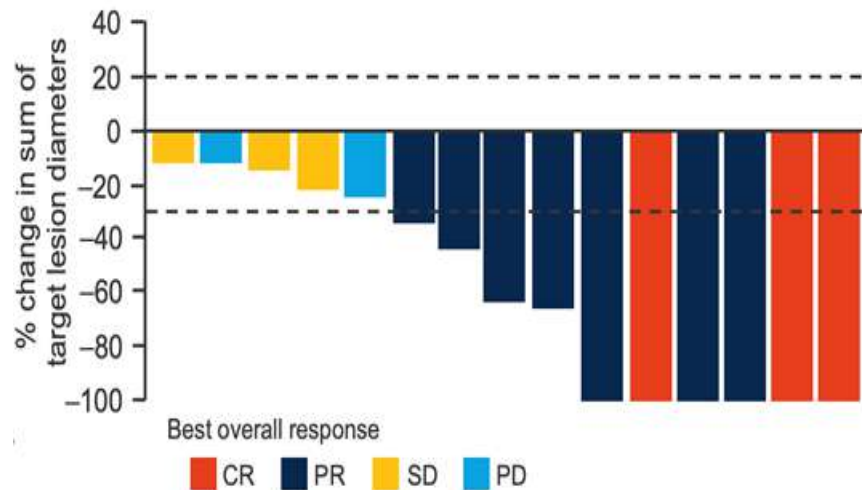
cohort A

5/7 patients

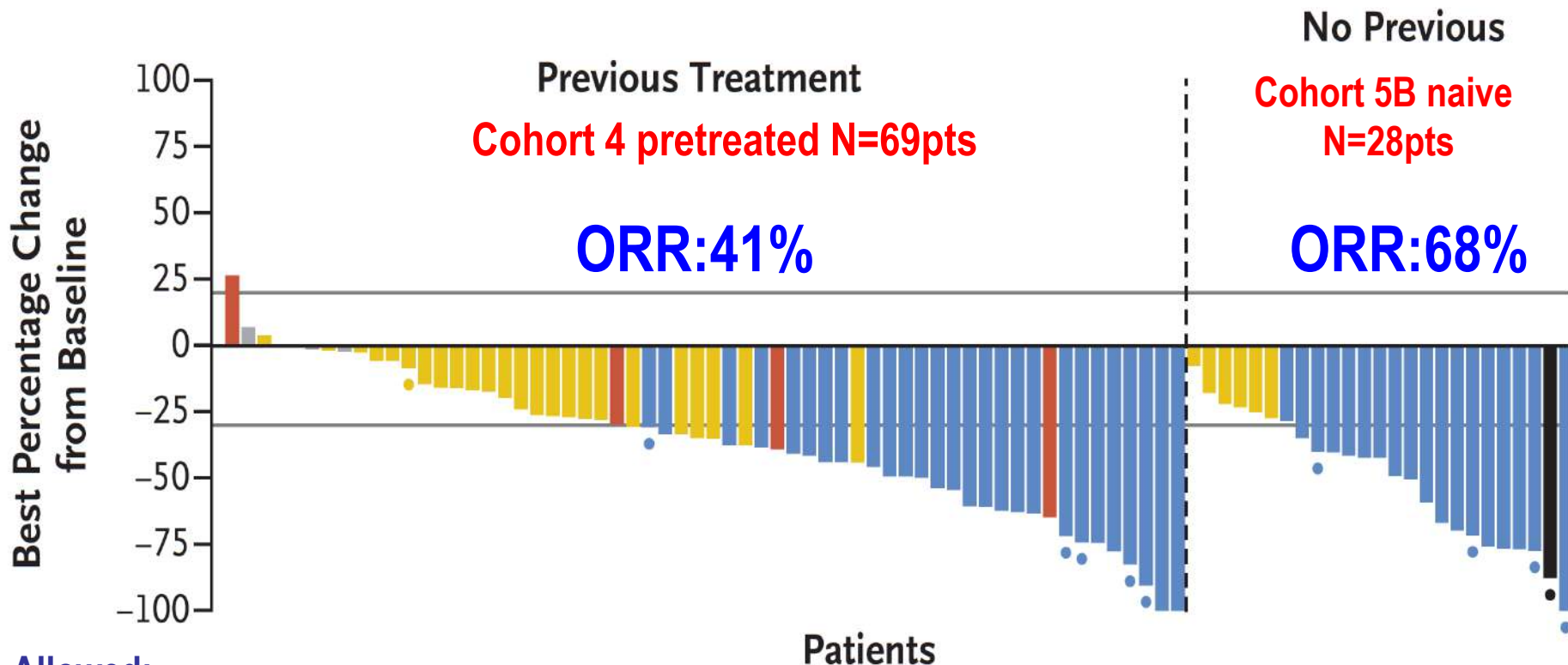


cohort C

Intracranial response in patients with target lesion (n=15)



GEOMETRY Study (Capmatinib)



Allowed:

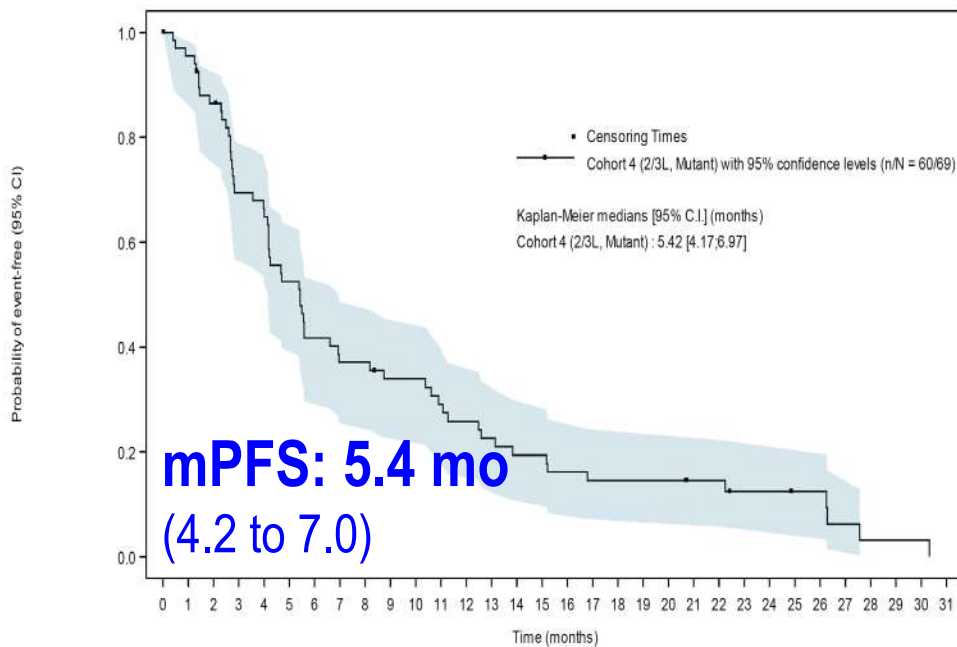
- stable CNS mets
- asymtomatic CNS mets

Primary endpoint: ORR by IRC

Capmatinib - PFS per BIRC assessment

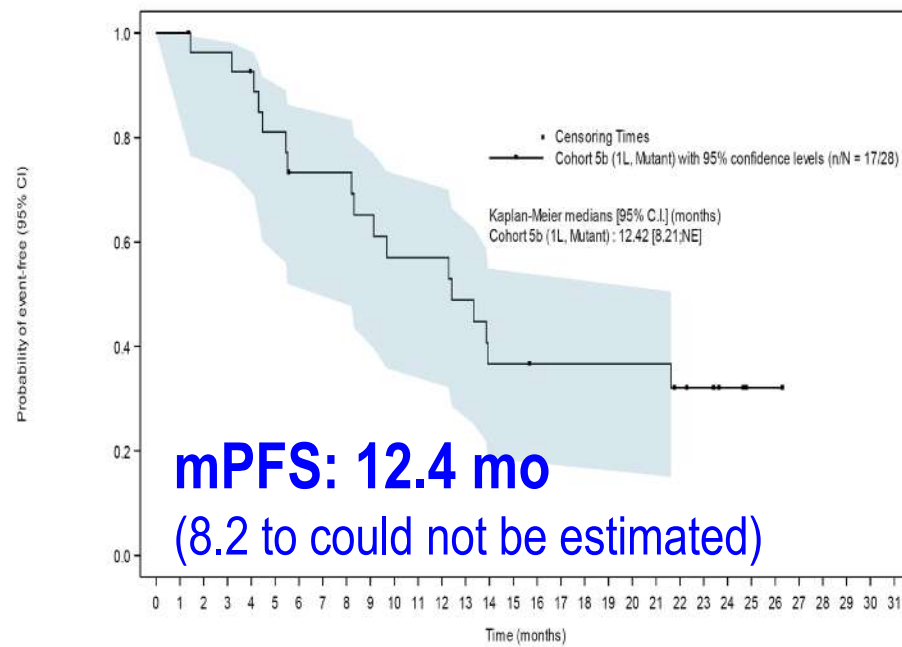
2/3 L

1L



No. of patients still at risk

Time (months)	0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31
Cohort 4 (2/3L, Mutant)	69	64	57	45	43	34	27	24	24	21	21	18	16	14	12	12	10	9	9	9	9	7	7	5	5	4	4	2	1	1	1	0



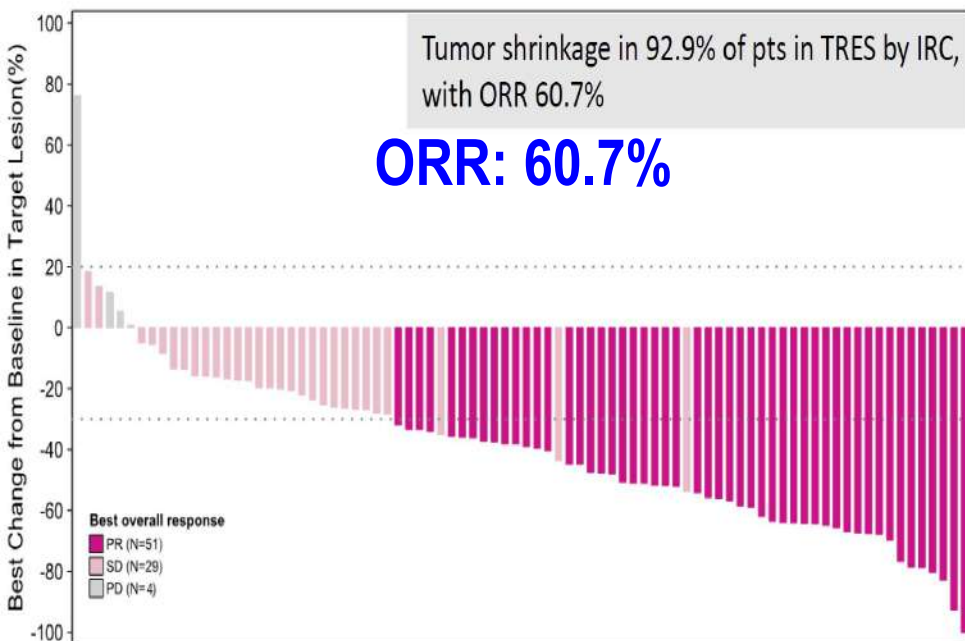
No. of patients still at risk

Time (months)	0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31
Cohort 5b (1L, Mutant)	28	28	26	26	24	21	18	18	18	16	14	14	14	12	9	9	8	8	8	8	8	8	6	5	3	1	1	0	0	0	0	0

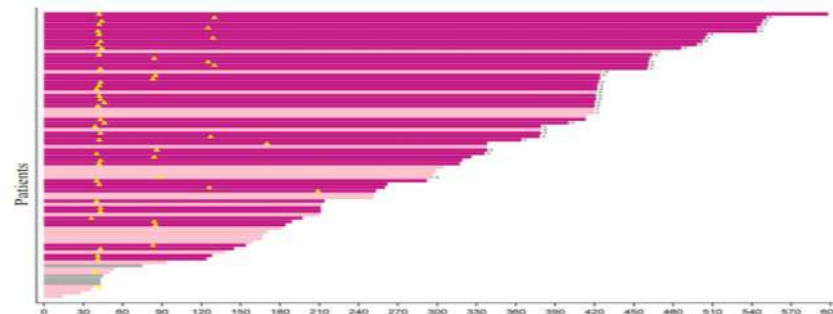
Savolitinib 1L – MET Exon14 mutation NSCLC

Deep and Durable Response (IRC assessment)

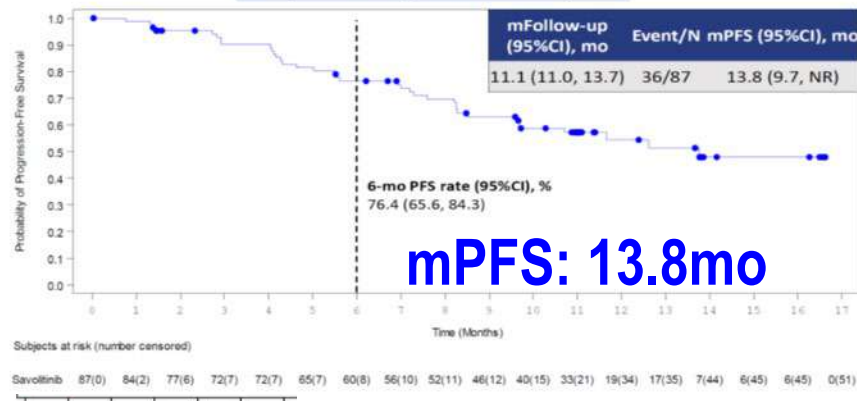
Waterfall Plot - Tumor Shrinkage of Target Lesion
(IRC Assessment) - Tumor Response Evaluable Set



Swimmer Plot - (IRC Assessment) - Tumor Response Evaluable Set



KM plot of PFS- (IRC Assessment) FAS



METex14 tyrosine kinase inhibitors

	Non-Selective	Selective TKI							
	CRIZOTINIB PROFILE 1001	CAPMATINIB GEOMETRY Mono-1		TEPOTINIB VISION (C) (TBx)		SAVOLITINIB		GLUMETINIB GLORY	
IC ₅₀ (nM)	26,5	0.6		3.0		2.1		0.42	
Dose	250 mg BID	400 mg BID		500 mg QD		400-600 mg QD		300 mg QD	
Line	≥1	1	≥2	1	≥2	1	≥2	1	≥2
N	69	60	100	161	51	87	42	42	27
RR (%)	32	68.3	44	62.3	51	60.7	40.5	66.7	51.9
DoR (mo.)	9.1	16.6	9.7	NE	12.6	NR	9.7	NE	5.1
PFS (mo.)	7.3	12.4	5.4	15.9	13.8	13.8	6.9	NE	5.7
OS (mo.)	20.5	25.5	13.6	22.7	19.6	NR	19.4	NR	NR
Comments	Shorter PFS in ctDNA positive at baseline	Higher activity in 1 st line vs. ≥2 nd line		The RR regardless Age, line & type of previous therapy		Sarcom. vs. others RR: 40% vs. 44% PFS: 5.5 vs. 6.9			



2022



2022 (2nd)



2021

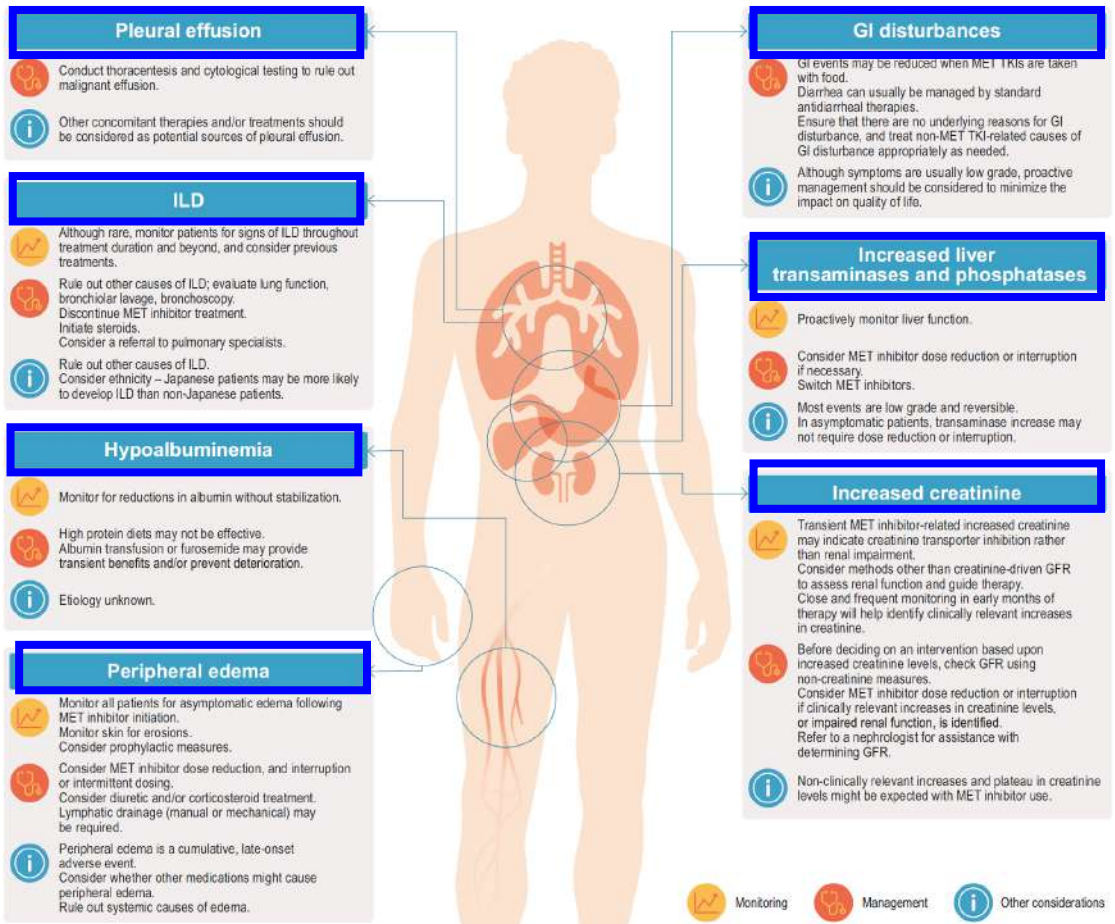


2021 (2nd)



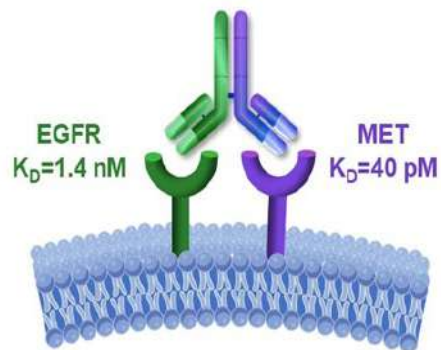
2021

Overview - key adverse events

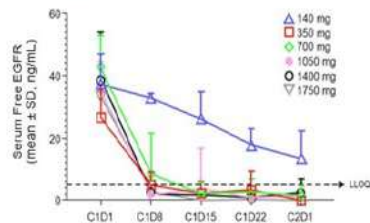


Bispecific mAb anti-EGFR & anti-MET

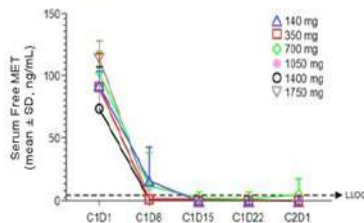
Amivantamab



EGFR



MET



CHRYSALIS Study Phase 1

Part 1: Dose Escalation

140–1750 mg

Objective: Establish RP2D

RP2D

Amivantamab
1050 mg (<80 kg)
1400 mg (≥80 kg)

Intravenous dosing
C1 QW, C2+ Q2W

Part 2: Dose Expansion

MET-2 Cohort: METex14 n=55^a
(up to 100 planned)

Objective: Safety and efficacy at the RP2D

Eligibility

- Metastatic or unresectable/advanced NSCLC
- Failed or ineligible for standard of care therapy

Eligibility for METex14 Cohort

- Measurable disease
- Primary METex14 mutation by NGS of tumor or ctDNA

Bispecific mAb anti-EGFR & anti-MET

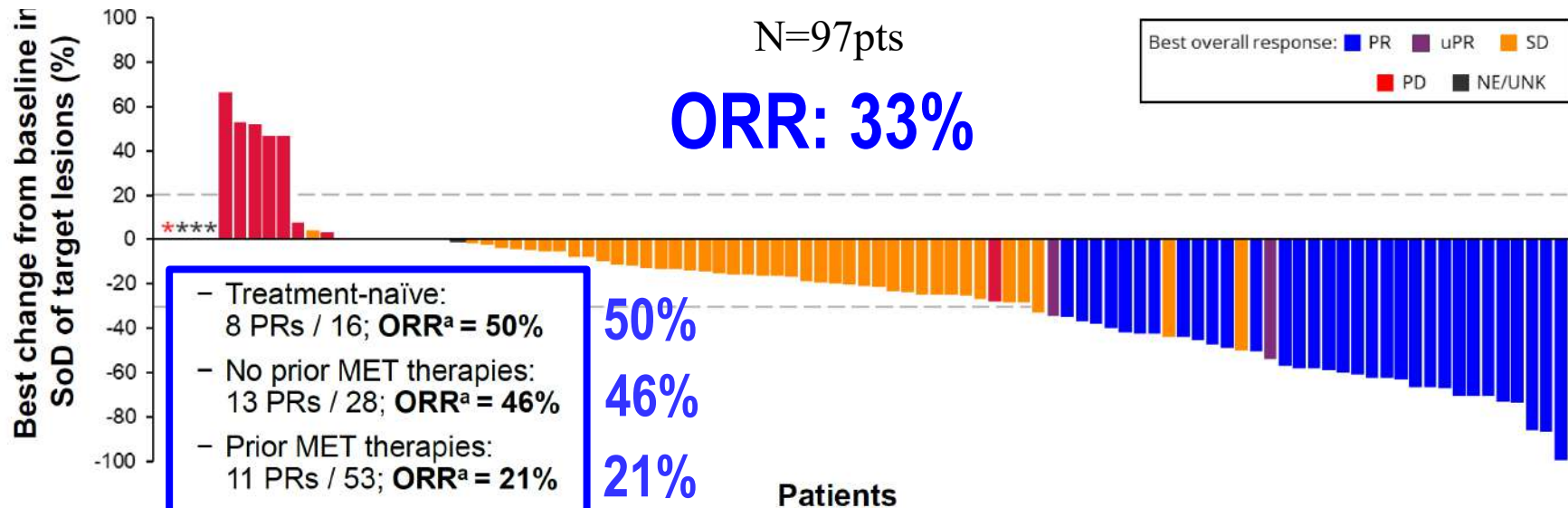
*MET*ex14 mutation: Amivantamab

CHRYSALIS Study Phase 1

N=97pts

ORR: 33%

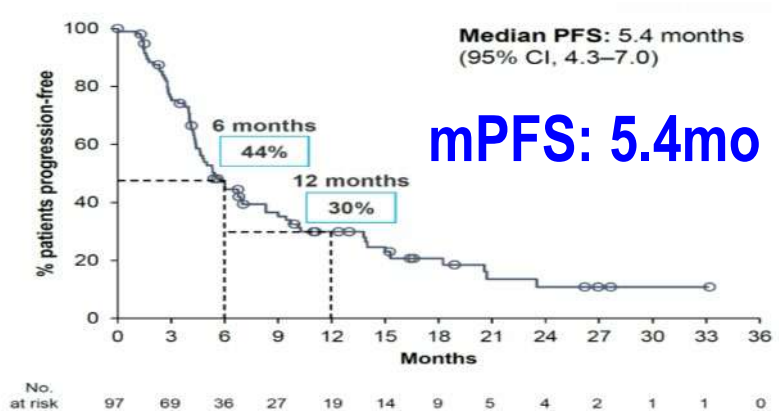
Best overall response: PR (blue), uPR (purple), SD (orange), PD (red), NE/UNK (black)



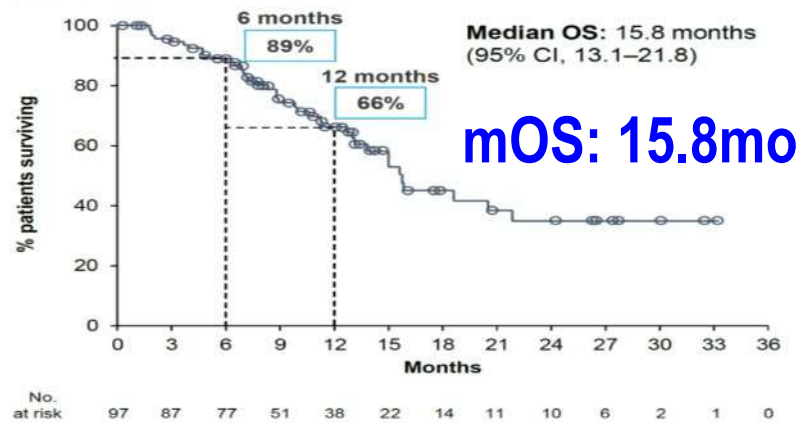
Treatment-naïve
No prior MET therapies
Prior MET therapies

PFS and OS of amivantamab monotherapy

PFS (n = 97)



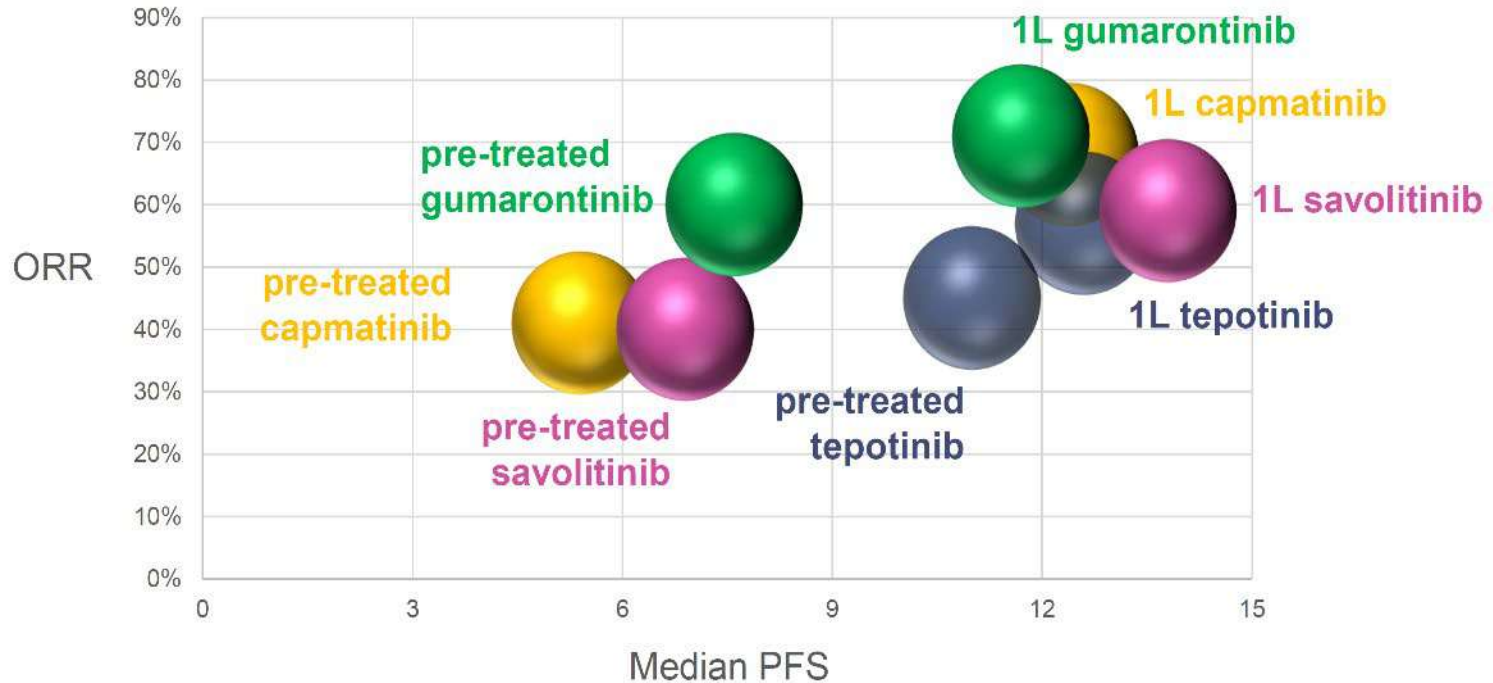
OS (n = 97)



Safety Profile

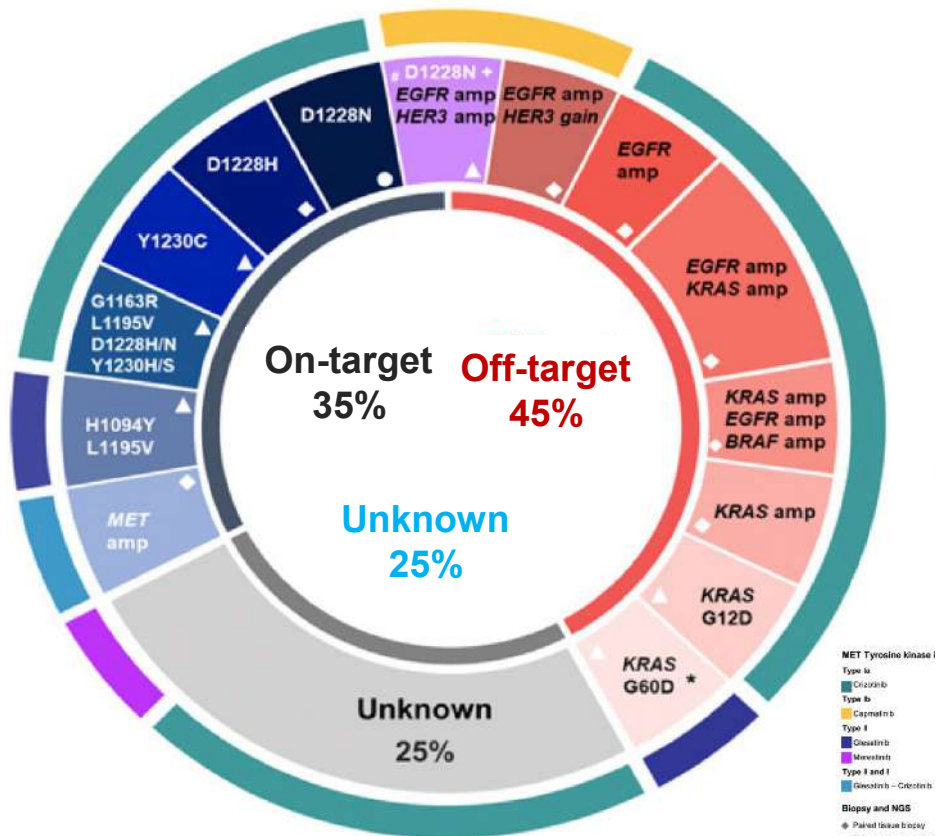
AEs (≥20%) by preferred term, n (%)	Median follow-up: 10.0 months (n = 97)	
	Total	Grade ≥3
Associated with EGFR inhibition		
Paronychia	47 (48.5)	0
Dermatitis acneiform	40 (41.2)	1 (1.0)
Rash	37 (38.1)	1 (1.0)
Stomatitis	27 (27.8)	0
Pruritus	20 (20.6)	0
Associated with MET inhibition		
Hypoalbuminemia	37 (38.1)	2 (2.1)
Peripheral edema	36 (37.1)	4 (4.1)
Other		
Infusion-related reaction	70 (72.2)	4 (4.1)
Fatigue	28 (28.9)	2 (2.1)
Dyspnea	22 (22.7)	5 (5.2)
Hypokalemia	22 (22.7)	3 (3.1)
Nausea	21 (21.6)	0
Decreased appetite	21 (21.6)	0
Alanine aminotransferase increased	20 (20.6)	2 (2.1)
AEs of special interest by grouped term, n (%)		
Rash ^a	76 (78.4)	3 (3.1)
Venous thromboembolism ^b	8 (8.2)	2 (2.1)
Interstitial lung disease ^c	4 (4.1)	1 (1.0)

First line MET TKI therapy appears to be associated with higher ORRs and PFS medians

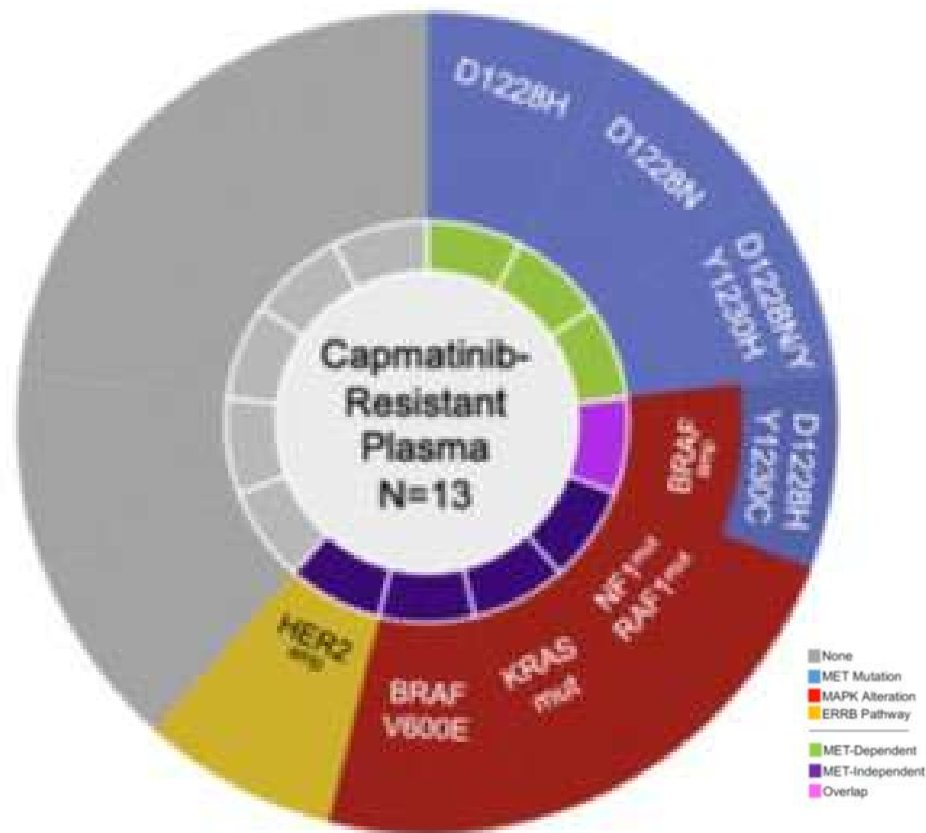


Molecular Mechanisms of Acquired Resistance to MET TKIs

Post MET TKIs plasma specimens



Postcapmatinib plasma specimens



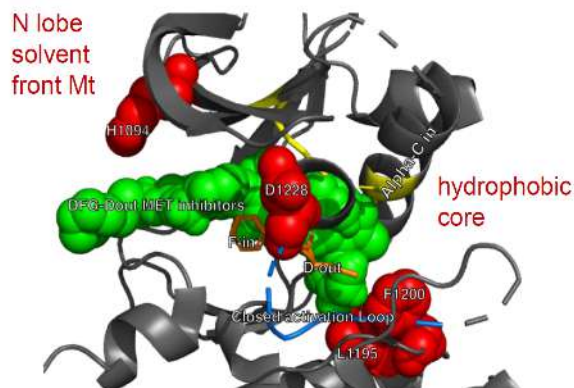
MET TKI and resistance mutations

Drug name	Inhibitor type	Selectivity	Activity against other kinases	RESISTANCE MUTATION ACTIVITY				References
				Solvent-Front	Activation-Loop	Beta-9 strand	Hydrophobic-core	
Crizotinib	DFG-Din	Multi kinase inhibitor	ALK; ROS1	G1163R	Y1230C/H/N/S; D1231Y	D1228N/H/E	L1995V	Cosmic drug resistance dataset; Recondo et al CCR 2020; Fujino et al JTO 2019
Tepotinib	DFG-Din	MET selective		G1163E	Y1230X	D1228X	L1995V	Fujino et al JTO 2019
Capmatinib	DFG-Din	MET selective		Active on cell lines	Y1230H	D1228N/H	L1995V	Cosmic drug resistance dataset; Recondo et al CCR 2020; Fujino et al JTO 2019
Savolitinib	DFG-Din	MET selective		Active on cell lines	Y1230S	D1228N/V	L1995V	Cosmic drug resistance dataset; Fujino et al JTO 2019
Cabozantinib	DFG-Dout	Multi kinase inhibitor	VEGFR; KIT; RET; AXL; FLT3; ROS1	Variably active on cell lines	Known to be active	D1228Y/A/N	L1195V; F1200L	Fujino et al JTO 2019; Fujino et al JHO 2022; Kang et al Lung Cancer 2023;
Merestinib	DFG-Dout	Multi kinase inhibitor	FLT3; AXL; ROS1	Active on cell lines	Known to be active	D1228Y	F1200I	Fujino et al JTO 2019
Glesatinib	DFG-Dout	Multi kinase inhibitor	VEGFR; TIE2; SMO	H1094Y	Known to be active	D1228Y/A	L1195V	Recondo et al CCR 2020; Fujino et al JTO 2019

- Clinical evidence of resistance
- Preclinical evidence of resistance
- Evidence of activity

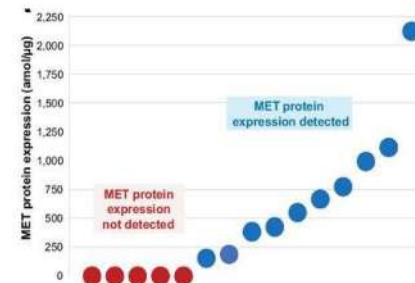
Ref/s: table courtesy of Matteo Repetto MD, Univ Milan/MSKCC

Antibody-based therapeutics may address complex ontarget and off-tatget resistance



MET Ab	Targets
Onartuzumab, telisotumab, HLX55	MET
REGN5093, Sym015	MET, MET
Amivantamab, MCLA-129, CKD702	MET, EGFR
GB263T	MET, MET, EGFR
TAVO412	MET, EGFR, VEGF

MET ADC	Warhead (if disclosed)
Telisotumab vedotin (MET)	MMAE
TR1801 (MET)	PBD
SHR-A1403 (MET)	Microtubule inhibitor
RC108 (MET)	
ABBV-400 (MET)	Topo I inhibitor
BYON3521 (MET)	Duocarmycin based
AZD9592 (MET-EGFR)	Topo I inhibitor
MYTX-011 (MET)	MMAE
REGN5093-M114 (MET-MET)	Maytansine derivative



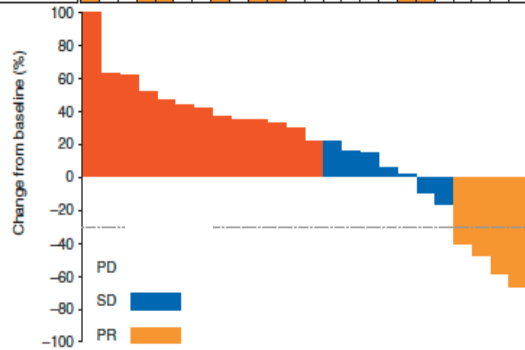
Ref/s: Bachall et al, Cancer Discov 2017; green - volumes occupied by: Merestinib (4EEV), Foretinib (6SD9) and Cabozantinib (Docked)

Attenuated efficacy of ICI in *MET*¹⁴

MSKC cohort

147MET+
ORR: 17%
PFS: 1.9 mo

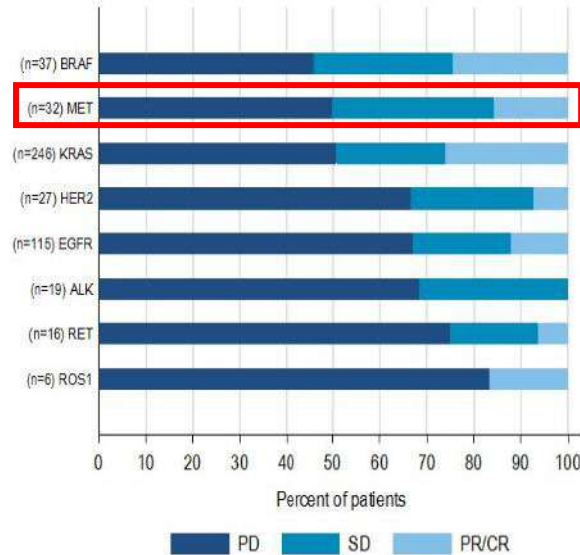
Immunotherapy	Ipilimumab	Nivolumab	Pembrolizumab	Atezolizumab	Atezolizumab + Ipilimumab	Atezolizumab + Nivolumab	Atezolizumab + Pembrolizumab	Atezolizumab + Nivolumab + Pembrolizumab
Histology	Stom	Adeno	NSCLC	NSCLC	NSCLC	NSCLC	NSCLC	NSCLC
PD-L1	90	80	80	80	80	80	80	80
TMB	75	45	45	45	45	45	45	45



RR not enriched in
PD-L1 ≥ 50% nor high TMB

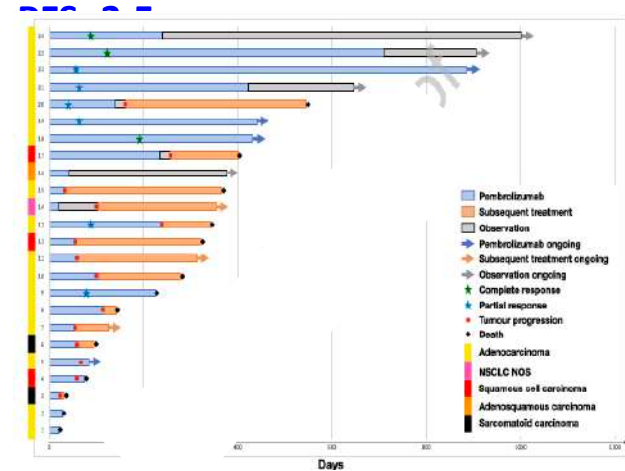
IMMUNOTARGET

36 MET+
ORR: 16%
PFS: 3.4 mo



GFPC 01-20

30 MET+ & PD-L1 ≥ 50%
ORR: 43%



Molecular tests positive (EGFR/ALK/ROS1/BRAF/RET/NTRK/MET/HER2/EGFRex20ins/KRAS G12C)

EGFR mutation
(refer to Figure 2)ALK translocation
(refer to Figure 3)ROS1
translocation
(refer to Figure 4)BRAF V600
mutation
(refer to Figure 5)

RET translocation

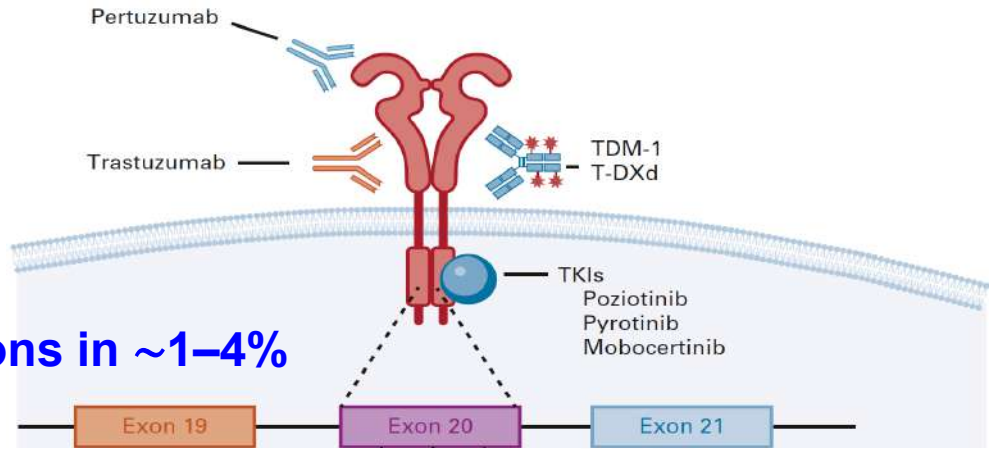
NTRK/HER2/
EGFRex20ins

METex14 skipping

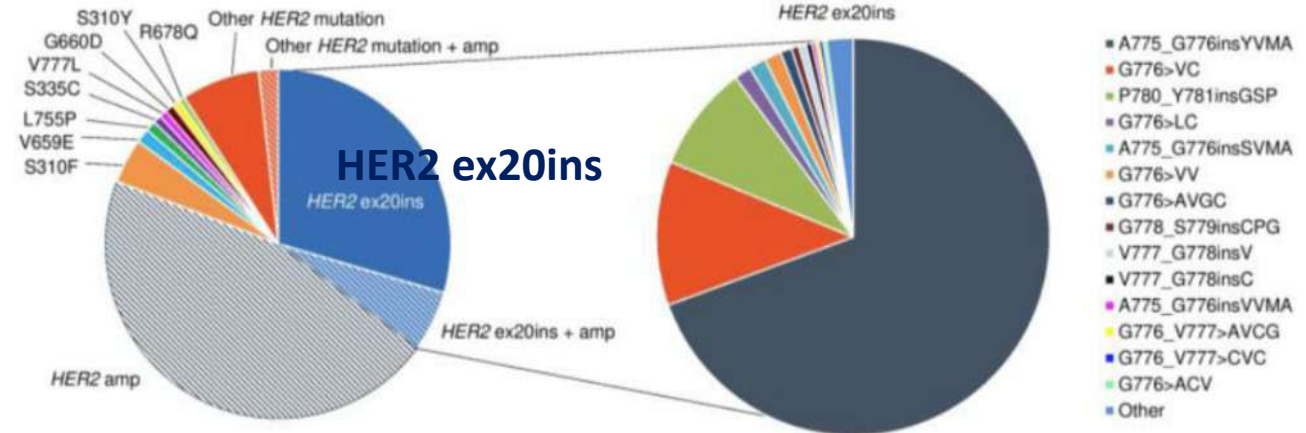
KRAS G12C
mutation2nd line

Capmatinib [III, A;
MCBS 3; ESCAT I-B]^{ALC}
Tepotinib [III, A; MCBS 3;
ESCAT I-B]^{ALC}

HER2 mutations occur mainly in the tyrosine kinase domain



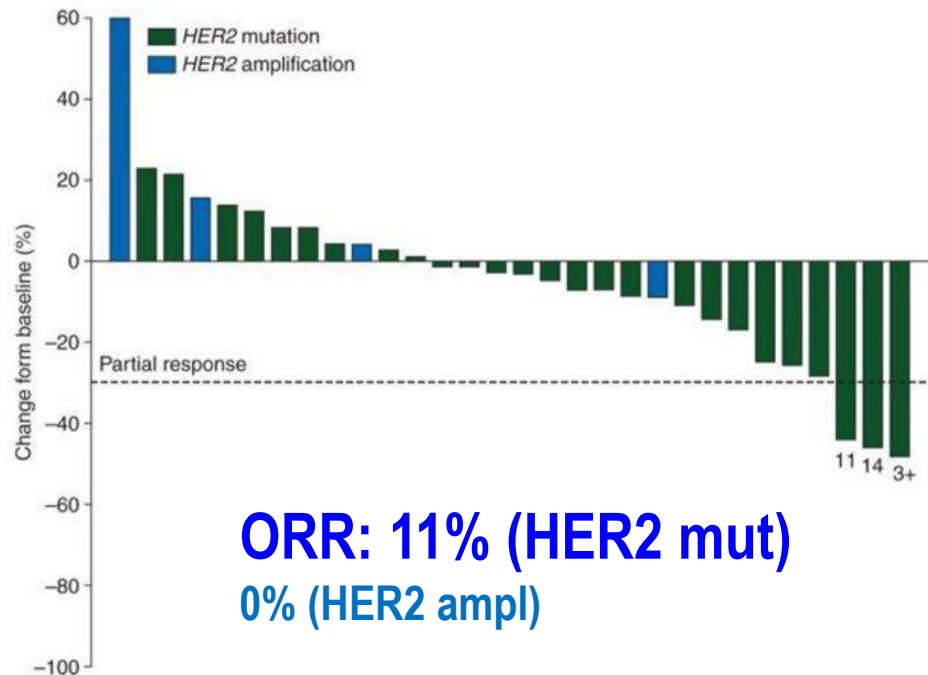
HER2 mutations in ~1–4%



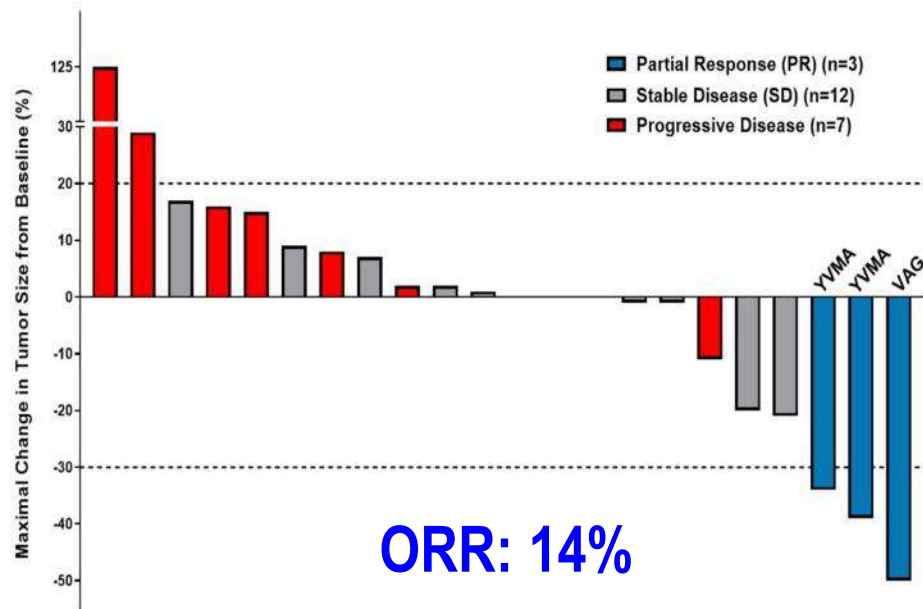
• 4.2% of NSCLCs had HER2 alterations including ex20ins (1.5%)

Dacomitinib and Afatinib for HER2 mutated NSCLC

Dacomitinib

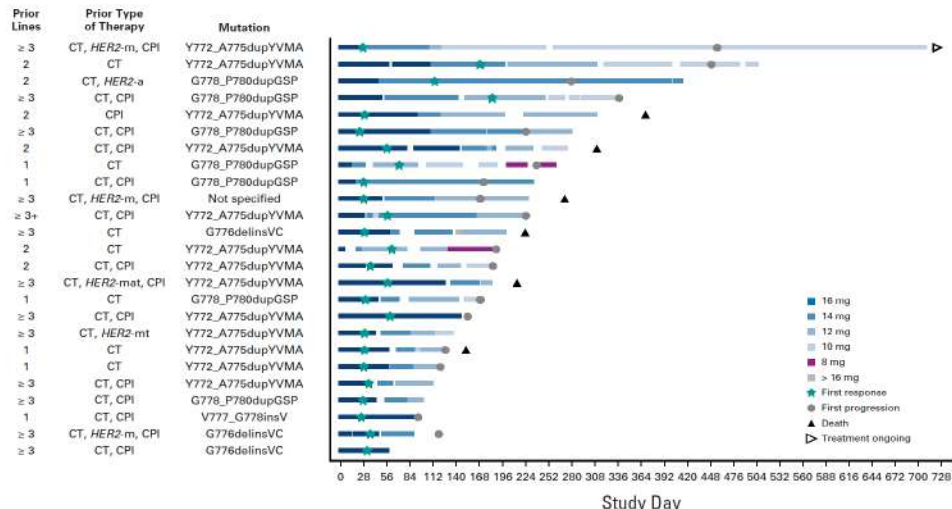
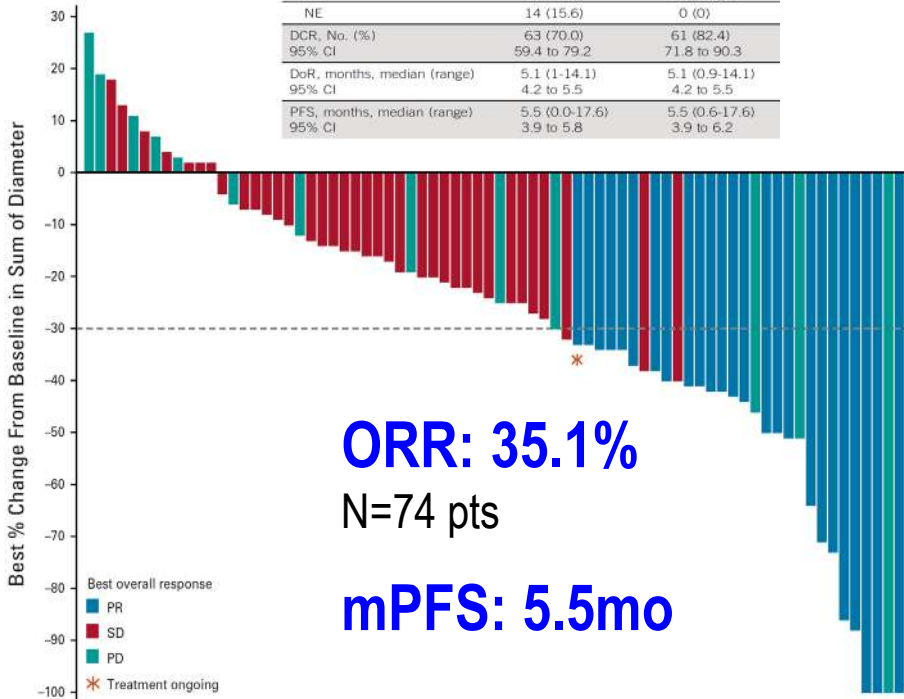


Afatinib 2)



Poziotinib - HER2 Exon 20 Insertion after Prior Therapies: ZENITH20-2 Trial

Parameter	As-Treated ^a (N = 90)	Evaluable ^b (n = 74)
ORR, No. (%)	25 (27.8) ^a	26 (35.1) ^b
95% CI	18.9 to 38.2	24.4 to 47.1
Best overall response, No. (%)		
CR	0 (0)	0 (0)
PR	25 (27.8) ^a	26 (35.1) ^b
SD	38 (42.2)	35 (47.3)
PD	13 (14.4)	13 (17.6)
NE	14 (15.6)	0 (0)
DCR, No. (%)	63 (70.0)	61 (82.4)
95% CI	59.4 to 79.2	71.8 to 90.3
DoR, months, median (range)	5.1 (1-14.1)	5.1 (0.9-14.1)
95% CI	4.2 to 5.5	4.2 to 5.5
PFS, months, median (range)	5.5 (0.0-17.6)	5.5 (0.6-17.6)
95% CI	3.9 to 5.8	3.9 to 6.2

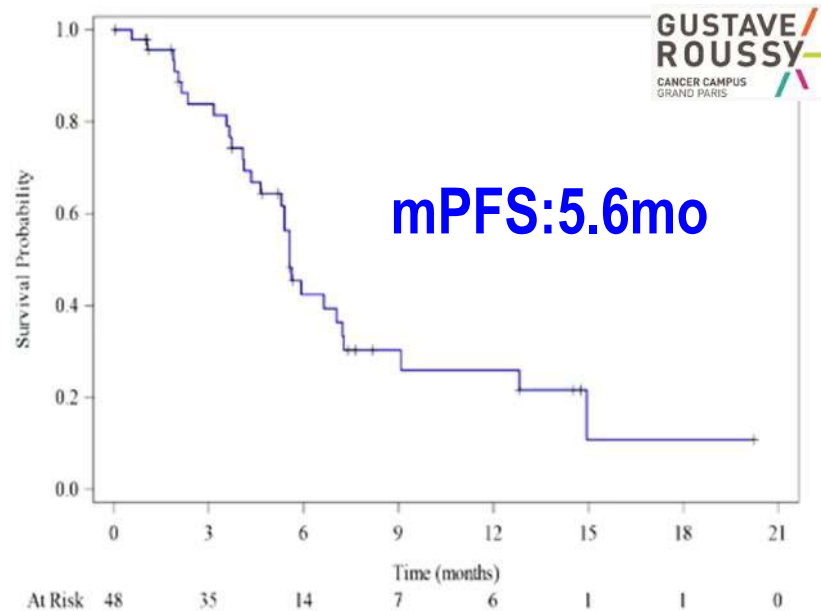
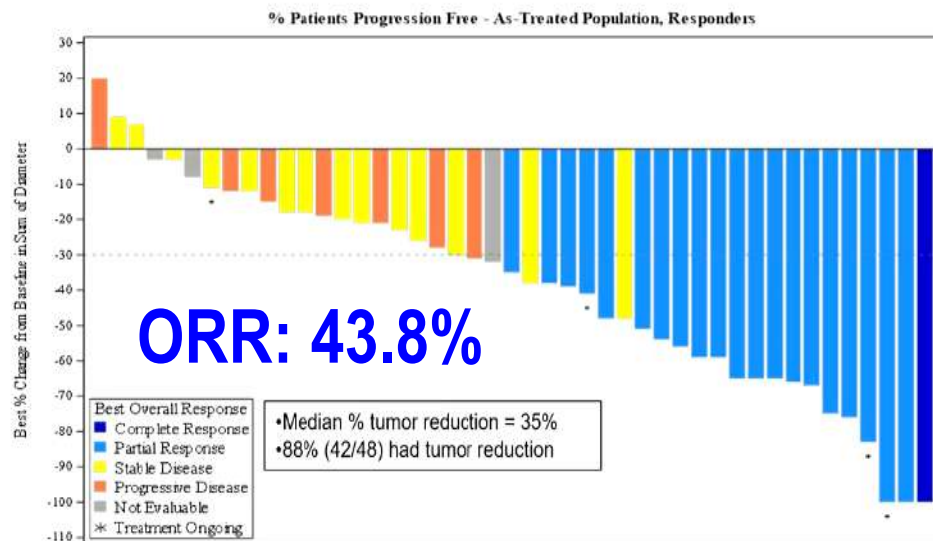


	N=90, n (%)	
Treatment-related AE	88 (98)	
Dose interruptions	78 (87)	
Dose reductions	70 (78)	

Preferred Term (PT)	N=90, n (%)		
	Any Grade	Grade 3	G4
Diarrhea	74 (82)	23 (26)	0
Rash	61 (68)	27 (30)	0
Stomatitis	59 (66)	20 (22)	1 (1)
Paronychia	34 (38)	1 (1)	0
Pneumonitis	1 (1)	0	0

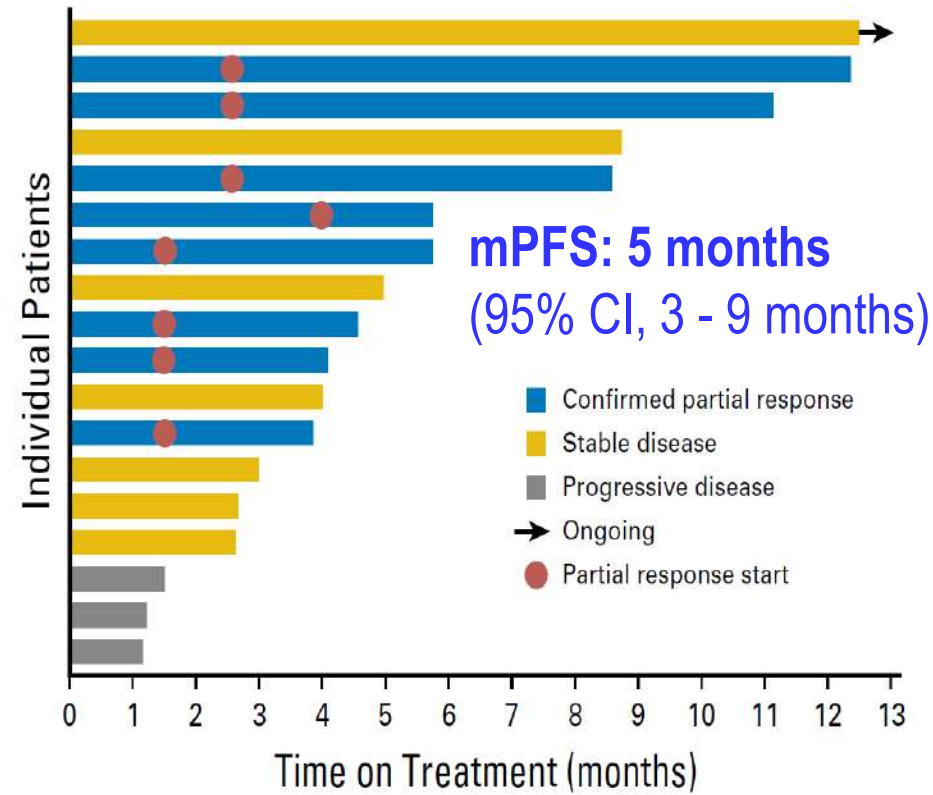
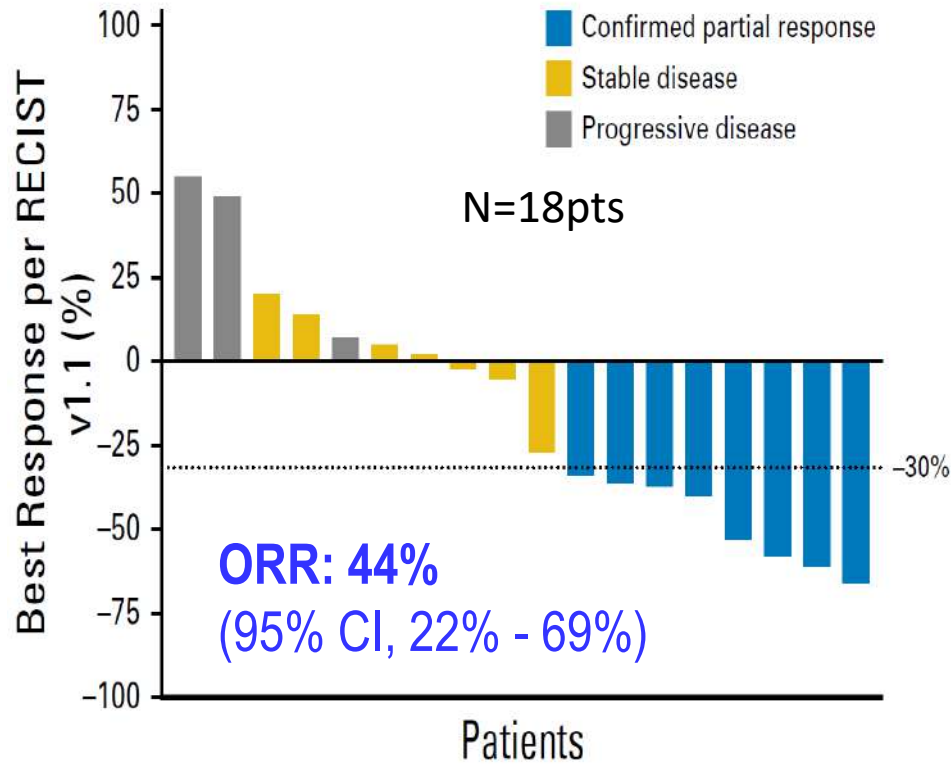
POZIOTINIB: First Line HER2 exon 20

Best %Change from Baseline in Target Tumor Size



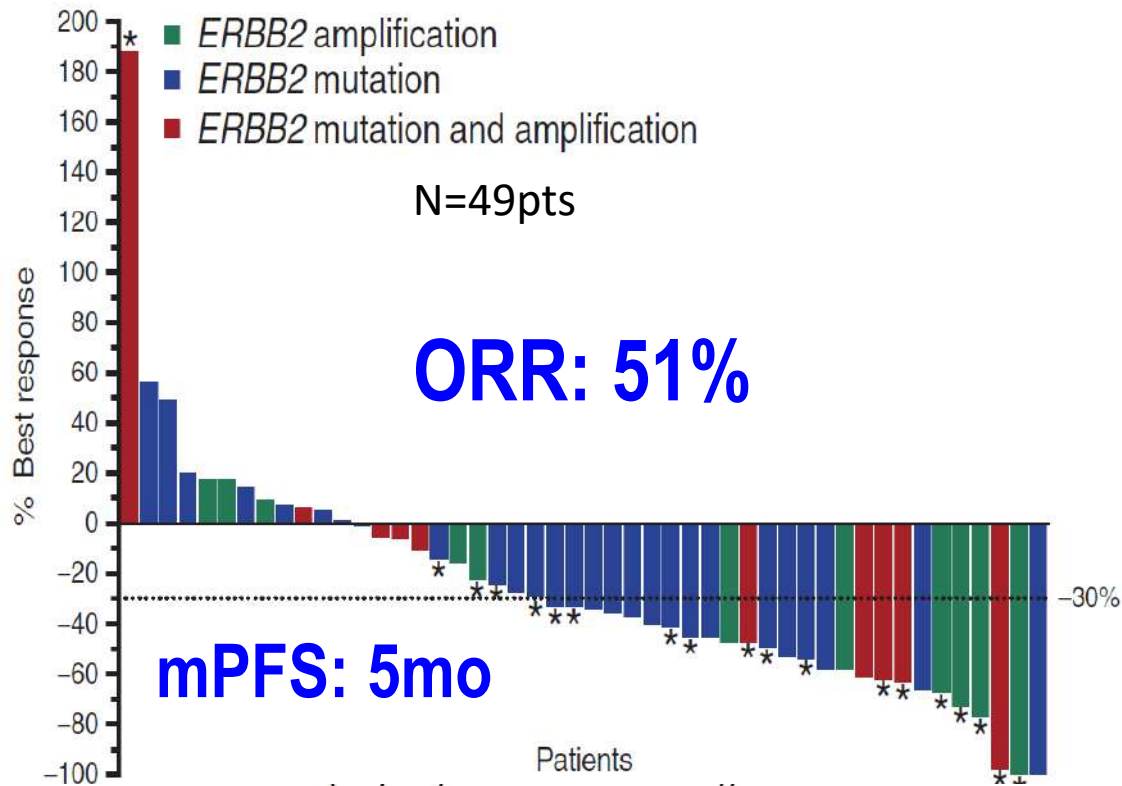
	Any Grade	Grade 3	Grade 4 / 5
Diarrhea	40 (83)	7 (15)	0
Rash	34 (69)	17 (35)	0
Stomatitis / Mucosal Inflammation	39 (81)	10 (21)	0
Paronychia	22 (46)	4 (8)	0
Pneumonitis	2 (4)	1 (2)	0

Ado-Trastuzumab Emtansine (T-DM1) in HER2-Mutant Lung Cancers (Phase II Basket Trial)



HER2 immunohistochemistry ranged from 0 to 2+ and did not predict response

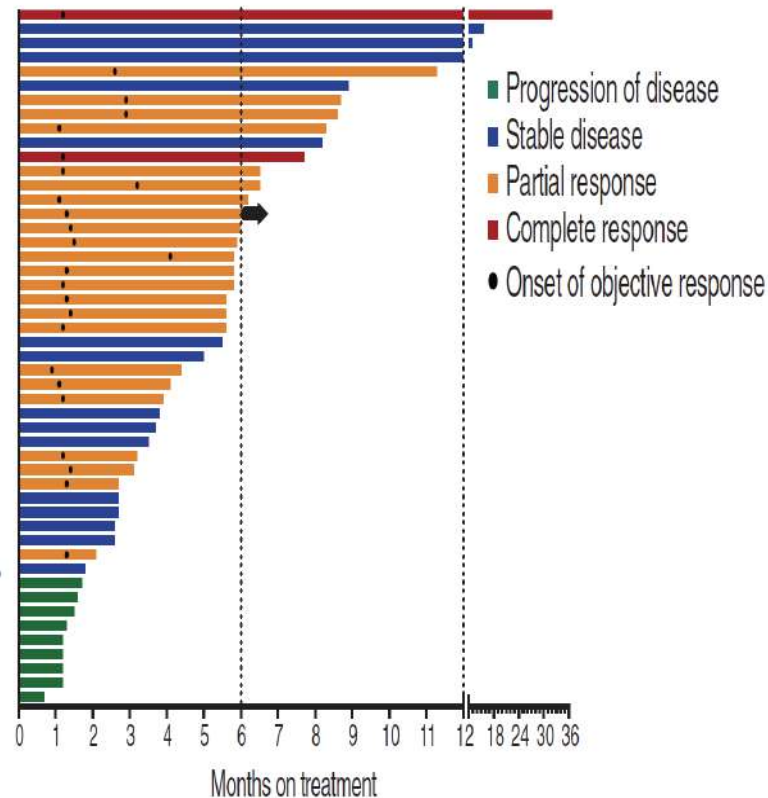
Clinical activity of T-DM1 in NSCLC



55% (6/11, 95% CI 23-83%) for *ERBB2*-amplified patients,

50% (14/28, 95% CI 31 to 69) for *ERBB2*-mutant patients

50% (5/10, 95% CI 19-81) for concurrently *ERBB2*-mutant and amplified patients



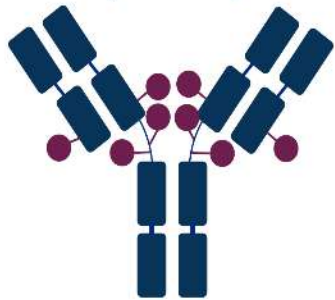
Differences between T-DXd and T-DM1

HER2-Targeting ADCs With a Similar mAB Backbone

Next-generation ADCs

Trastuzumab
deruxtecan

(T-DXd)

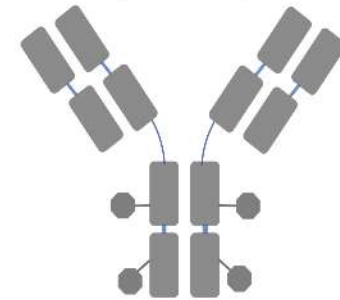


T-DXd	ADC Attributes	T-DM1
Topoisomerase I inhibitor	Payload MoA	Anti-microtubule
~8:1	Drug-to-antibody ratio	~3.5:1
Yes	Tumor-selective cleavable linker?	No
Yes	Evidence of bystander antitumor effect?	No

First generation ADCs

Trastuzumab
emtansine

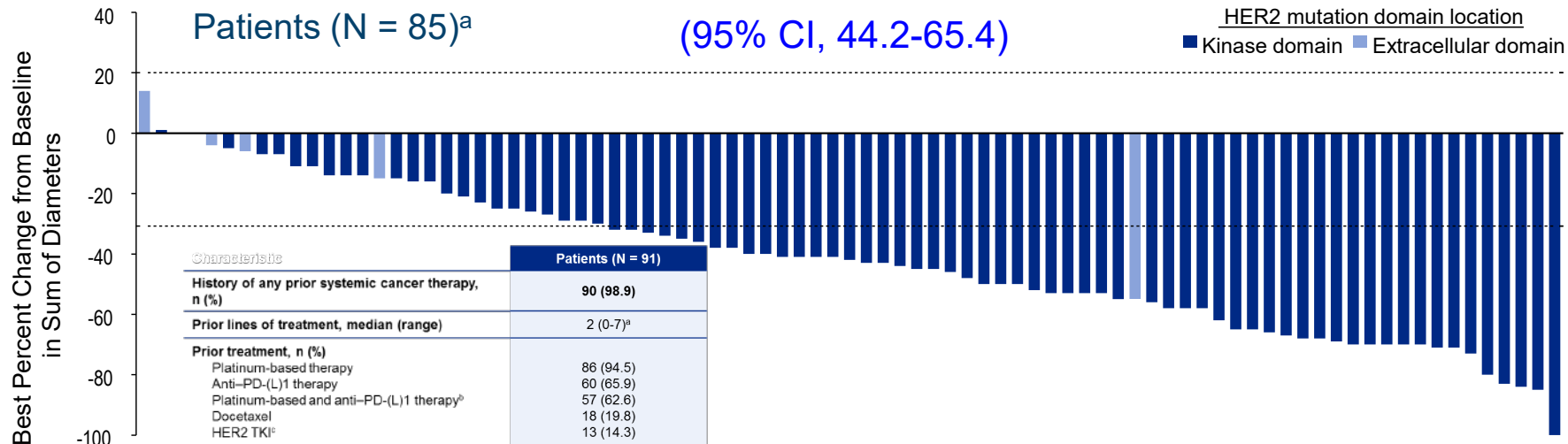
(T-DM1)



DESTINY-Lung01

Trastuzumab-Deruxtecan (T-DXd)

54.9%
(95% CI, 44.2-65.4)

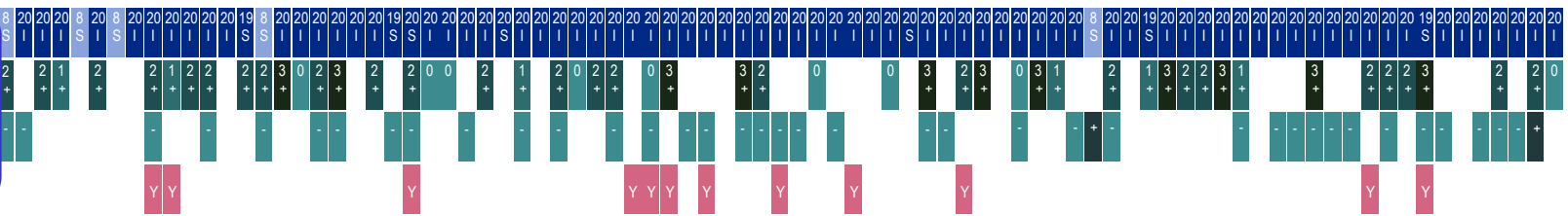


HER2 mutation (exon and subtype)

HER2 protein expression

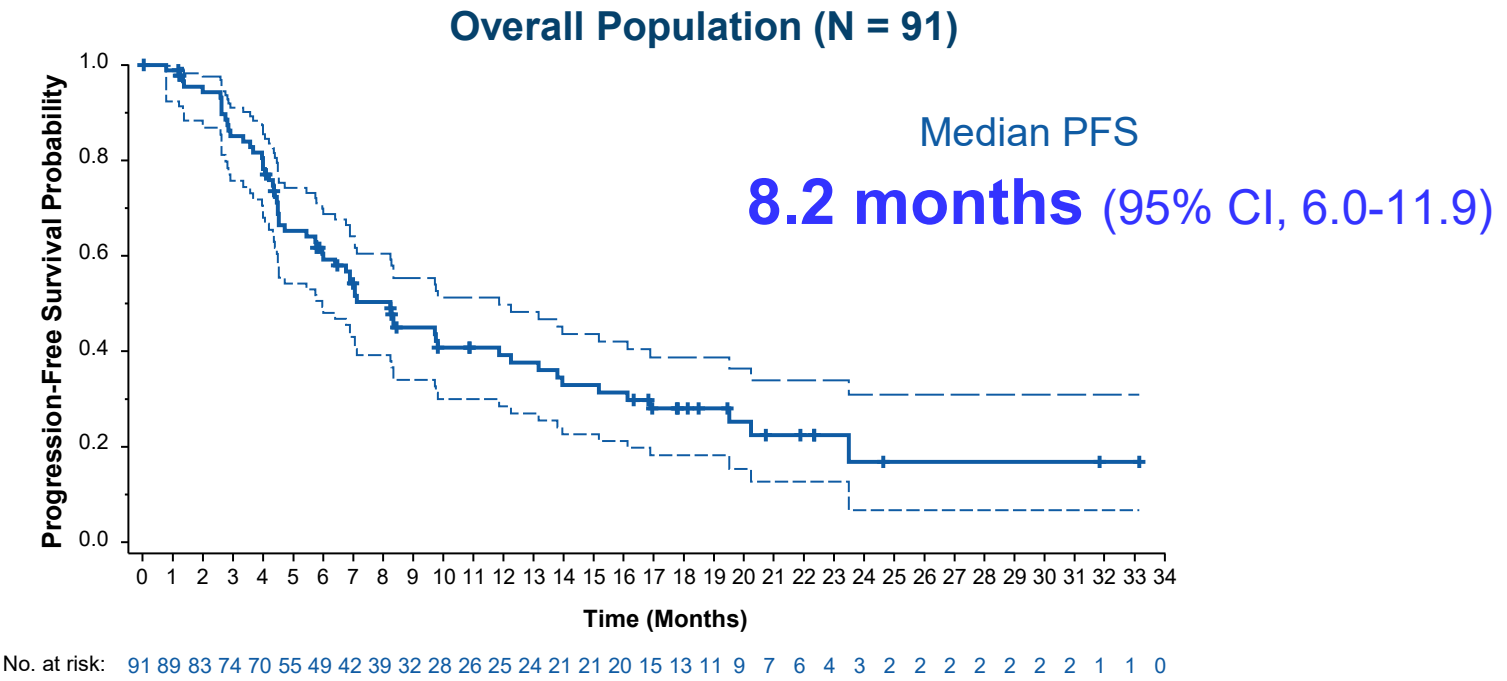
HER2 gene amplification

Prior HER2 TKI therapy



PFS in the overall *HER2m* NSCLC Population

T-DXd



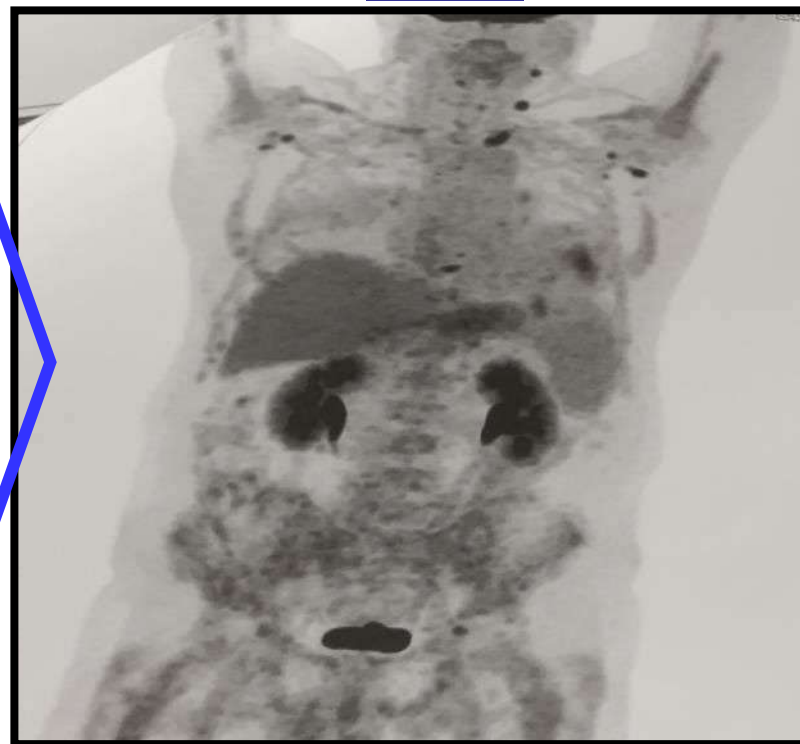
NSCLC adenocarcinoma HER2-insertion in exon 20

- Lymph node metastases, adrenal glands, hepatic, spleen, brain, bone...
- 1st line carboP-pemetrexed and 2nd line pembrolizumab

Baseline



T-DXd C7D1



Drug-related TEAEs Reported by Investigator

n (%)	Patients (N = 91)						
	Any grade	Grade ≥3					
Patients with ≥1 drug-related TEAEs	88 (96.7)	42 (46.2)					
Drug-related TEAEs with ≥20% incidence in all patients							
Nausea	66 (72.5)	8 (8.8)					
Fatigue ^a	48 (52.7)	6 (6.6)					
Alopecia	42 (46.2)	0					
Vomiting	36 (39.6)	3 (3.3)					
Neutropenia ^b	32 (35.2)	17 (18.7)					
Anemia ^c	30 (33.0)	9 (9.9)					
Diarrhea		Grade 1	Grade 2	Grade 3	Grade 4	Grade 5	Any Grade
Decreased appetite							
Leukopenia ^d	n (%)	3 (3.3)	15 (16.5%)	4 (4.4)	0	2 (2.2)	24 (26.4%)
Constipation							

Drug-Related ILD/Pneumonitis



Median time to onset of first reported drug-related **ILD/pneumonitis: 141 days** (range, 14-462 days) with a median duration of 43 days (95% CI, 24-94 days)

DESTINY-Lung02 (HER2 mut)

KEYPOINTS

- **HER2 mut** advanced NSCLC (PS 0-1)
- **≥ 1 prior therapy** (platinum based chemo)
- **Adequate washout** of prior treatment
- **CNS+ allowed** if treated/asymptomatic
- **Stratification: prior PD-(L)1 inhibitor**

Key patient inclusion criteria

- HER2+ NSCLC
 - ≥2 line therapy
 - ECOG PS 0-1
- (n=152)

R
2:1

Trastuzumab deruxtecan
5.4 mg/kg q3w
(n=102)

PD/
toxicity

Stratification

- Prior anti-PD-L1 therapy (yes vs. no)

Trastuzumab deruxtecan
6.4 mg/kg q3w
(n=50)

PD/
toxicity

Primary endpoint

- ORR (BICR)

Secondary endpoints

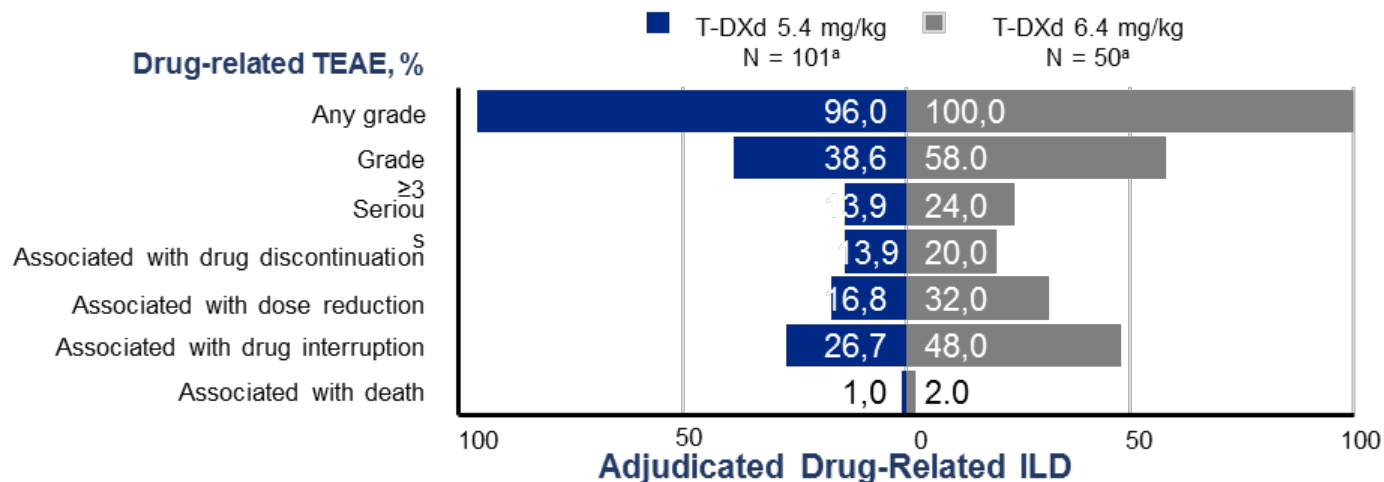
- OS, PROs, PK, safety

- **DESTINY-Lung02 is a randomized, phase 2 trial** assessing the efficacy and safety of 2 doses of **T-DXd (5.4 and 6.4 mg/kg)** in previously patients with metastatic **HER2m NSCLC**

The study was not powered to statistically compare the 2 arms

Trastuzumab Deruxtecan in Patients With HER2-Mutant Metastatic NSCLC: Primary Results of DESTINY-Lung02

Overall Safety



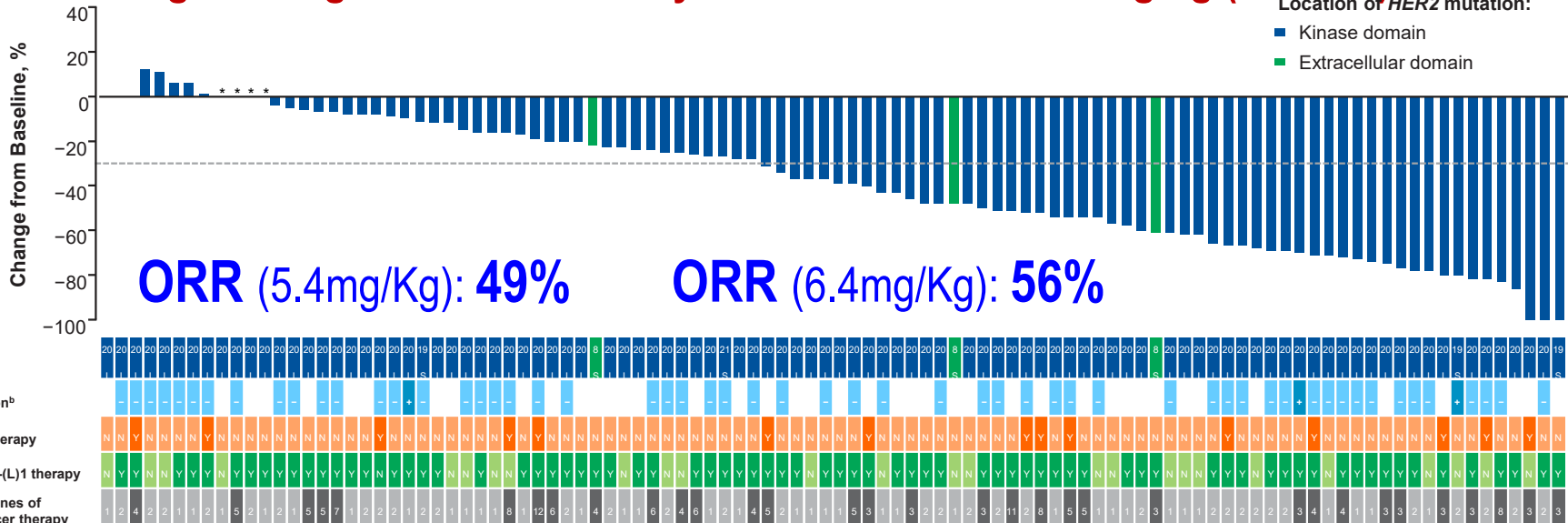
Adjudicated Drug-Related ILD

	T-DXd 5.4 mg/kg N = 101 ^a	T-DXd 6.4 mg/kg N = 50 ^a
Adjudicated as drug-related ILD		
Any grade, n (%)	13 (12.9%)	14 (28.0%)
Grade 1	4 (4.0)	4 (8.0)
Grade 2	7 (6.9)	9 (18.0)
Grade 3	1 (1.0)	0
Grade 4	0	0
Grade 5	1 (1.0)	1 (2.0)

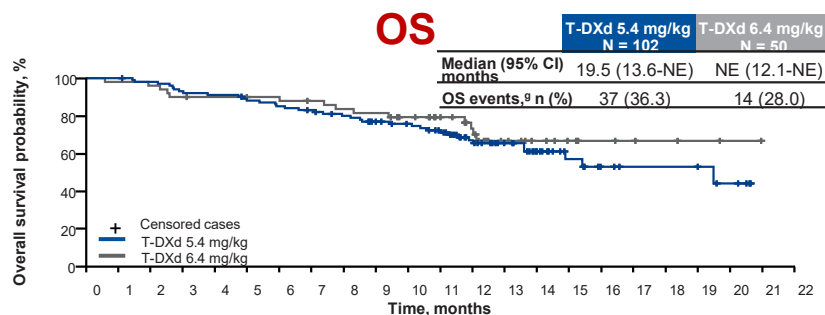
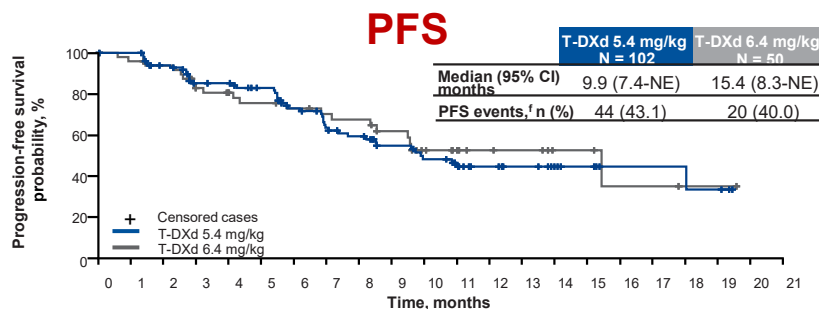
Best Percentage Change in Tumor Size by BICR with T-DXd 5.4 mg/kg (N = 102)

Location of *HER2* mutation:

- Kinase domain
- Extracellular domain



Responses observed regardless of *HER2* mutation type, *HER2* amplification status, and number or type of prior therapies



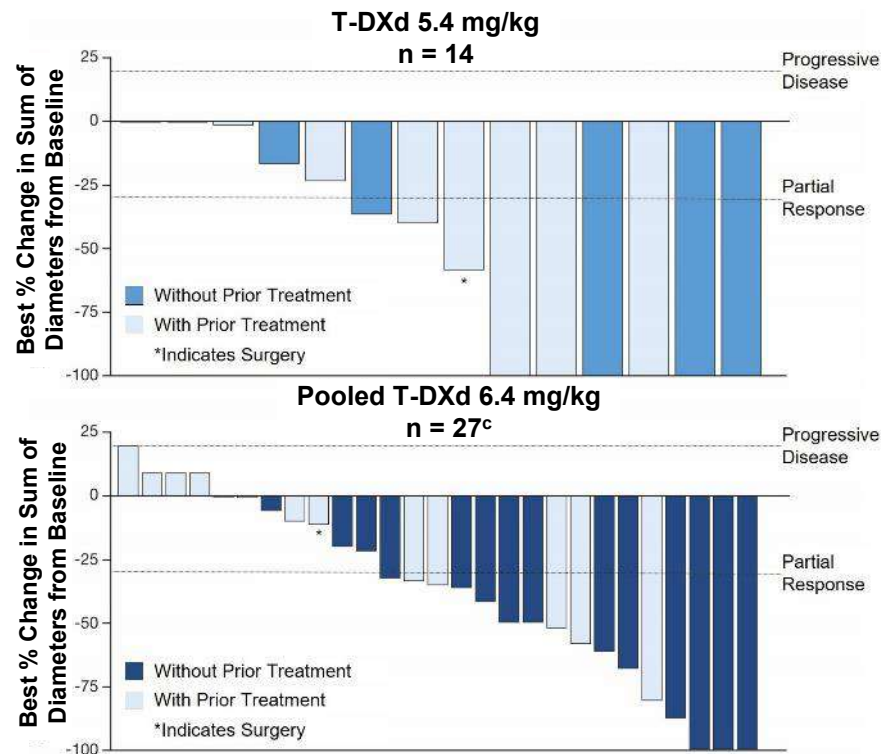


IC Objective Response Rates & Best Overall Response (BICR)

Measurable BM at Baseline

	T-DXd 5.4 mg/kg DL-02 BM n = 14	Pooled T-DXd 6.4 mg/kg DL-01 <i>HER2m</i> /DL-02 BM n = 30
IC-cORR, n (%)^a	7 (50.0)	9 (30.0)
95% CI ^b	23.0-77.0	14.7-49.4
CR	3 (21.4)	0
PR	4 (28.6)	9 (30.0)
SD	6 (42.9)	13 (43.3)
PD	1 (7.1)	4 (13.3)
NE ^c	0	2 (6.7)
Missing	0	2 (6.7)
IC-DCR, n (%)^a	13 (92.9)	22 (73.3)
95% CI ^b	66.1-99.8	54.1-87.7
IC-DoR, months^d		
Median, (95% CI) ^e	9.5 (3.6-NE)	4.4 (2.9-10.2)

12/14 (86%) patients with measurable BM receiving T-DXd 5.4 mg/kg and 21/27 (78%) in the pooled 6.4 mg/kg group experienced a reduction in brain lesion size from baseline as their best overall response



FDA and EMA approval

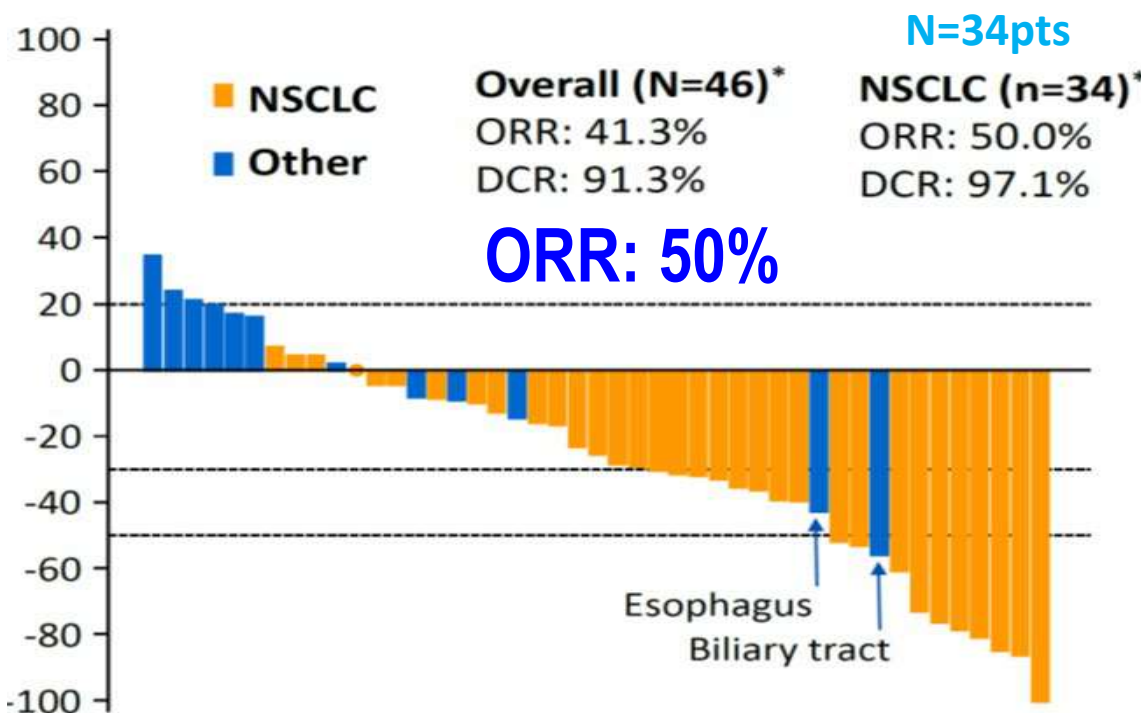
First drug approved for HER2-mutant NSCLC

On August 11, 2022, the FDA granted accelerated approval to fam-trastuzumab deruxtecan-nxki for adult patients with **unresectable or metastatic NSCLC whose tumors have activating HER2 (ERBB2) mutations**, and who have received a prior systemic therapy.

September 2023, Enhertu as monotherapy is indicated for the treatment of adult patients with advanced NSCLC whose tumours have an activating HER2 (ERBB2) mutation and who require systemic therapy following platinum-based chemotherapy with or without immunotherapy.

Drug	Study	N	Patient population	Study arm	Control arm	Endpoints
Trastuzumab-deruxtecan	DESTINY-Lung04	264	<ul style="list-style-type: none">• First-line advanced NSCLC• HER2 ex19 or ex20 mutations	Trastuzumab-deruxtecan	Pembrolizumab + pemetrexed + platinum	PFS

BEAMING Lung-1, Phase I of ZONGERTINIB (BI 1810631), in pts with HER2 aberrations

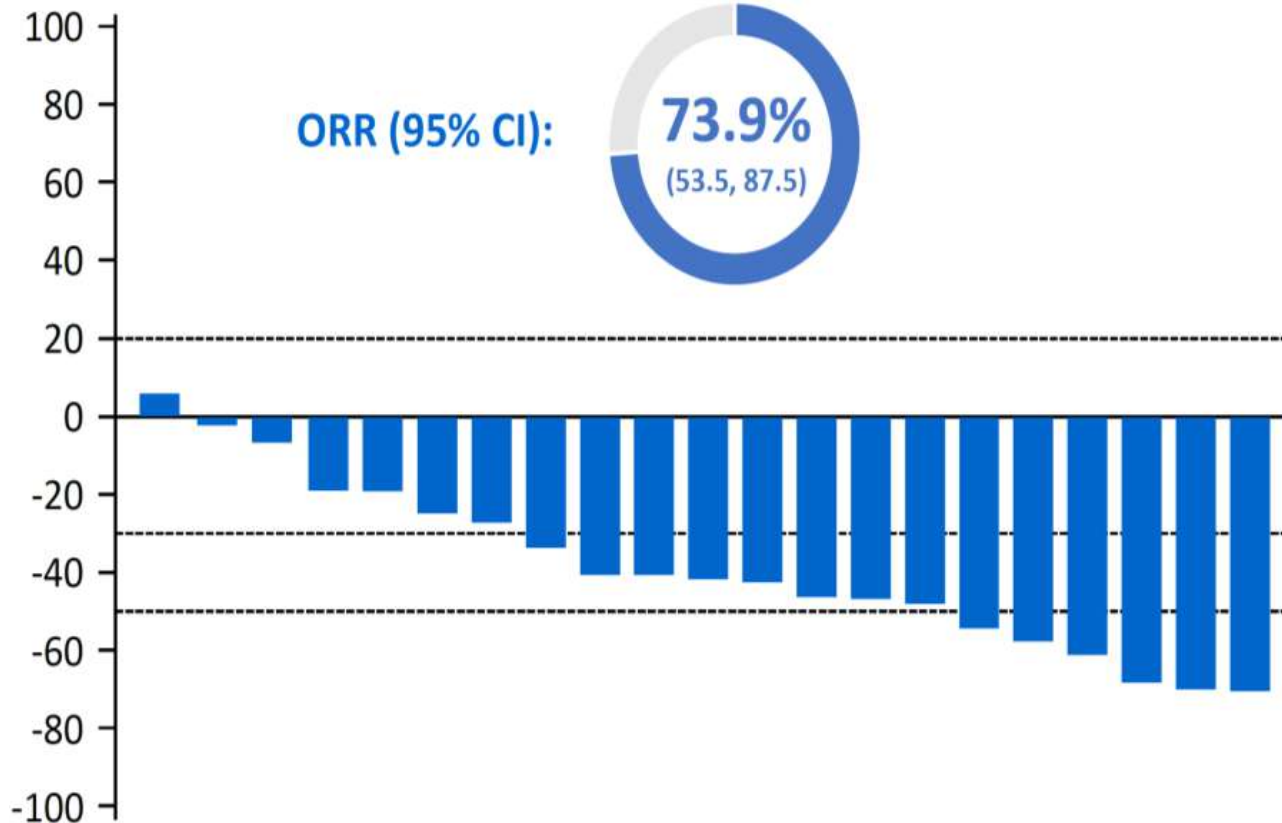
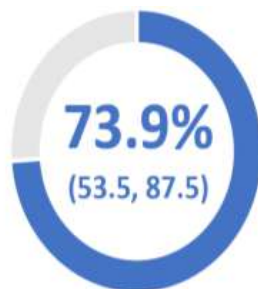


Phase Ia dose escalation and safety

Phase Ia TRAEs (%) [†]	Zongertinib BID (n=17)		Zongertinib QD (n=33)		Total (N=50)	
	Any	Grade ≥3	Any	Grade ≥3	Any	Grade ≥3
Any TRAE	76.5	5.9	84.8	12.1	82.0	10.0
Diarrhea	47.1	—	36.4	—	40.0	—
AST increased	5.9	—	18.2	3.0	14.0	2.0
Rash [†]	11.8	—	15.2	—	14.0	—
ALT increased	5.9	5.9	15.2	6.1	12.0	6.0
Paronychia	5.9	—	12.1	—	10.0	—
Dry skin	11.8	—	6.1	—	8.0	—
Anemia	11.8	—	6.1	—	8.0	—

Antitumor activity in Phase Ib - ZONGERTINIB (BI 1810631)

ORR (95% CI):



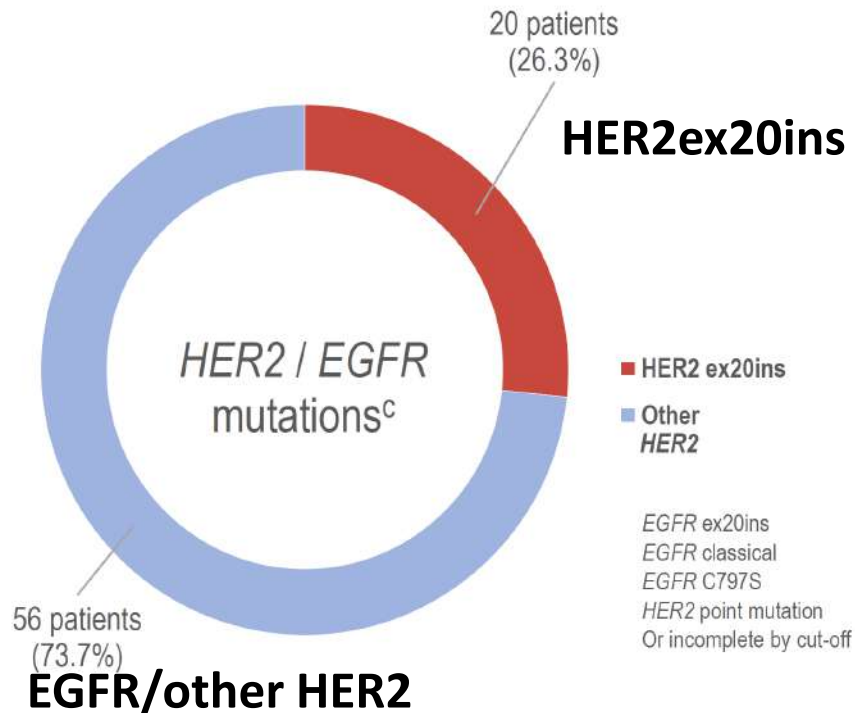
Overall (N=23)*

- The first interim analysis in Cohort 1 was passed
- Patients included in the efficacy analysis all had between 2–5 cycles of treatment at cut off
- DCR: 91.3%
- Median best percentage change from baseline in target lesions: -41.2%

Early evidence of efficacy in pts with HER2 exon20ins in phase I with BAY292088

Patient characteristics

	All treated patients (N=76)
Females, n (%)	45 (59.2)
Median age, years (range)	60.0 (35.0-81.0)
Baseline ECOG PS, n (%)	
1	49 (64.5)
Patients who have never smoked, n (%)	55 (72.4)
Number of previous systemic anti-cancer treatments, n (%)	
0	2 (2.6)
1	20 (26.3)
2	16 (21.1)
≥3	38 (50.0)
NSCLC histology, n (%)	
Squamous cell carcinoma, small cell, non-keratinizing	1 (1.3)
Adenocarcinoma, not otherwise specified	70 (92.1)
Adenocarcinoma with mixed subtypes	1 (1.3)
Papillary adenocarcinoma, not otherwise specified	1 (1.3)
Missing	3 (3.9)
Median time since initial diagnosis, months (range) ^a	28.2 (2.1-195.2)
Median time since most recent progression / relapse to first administration of study treatment, months (range) ^b	1.5 (0-30.0)



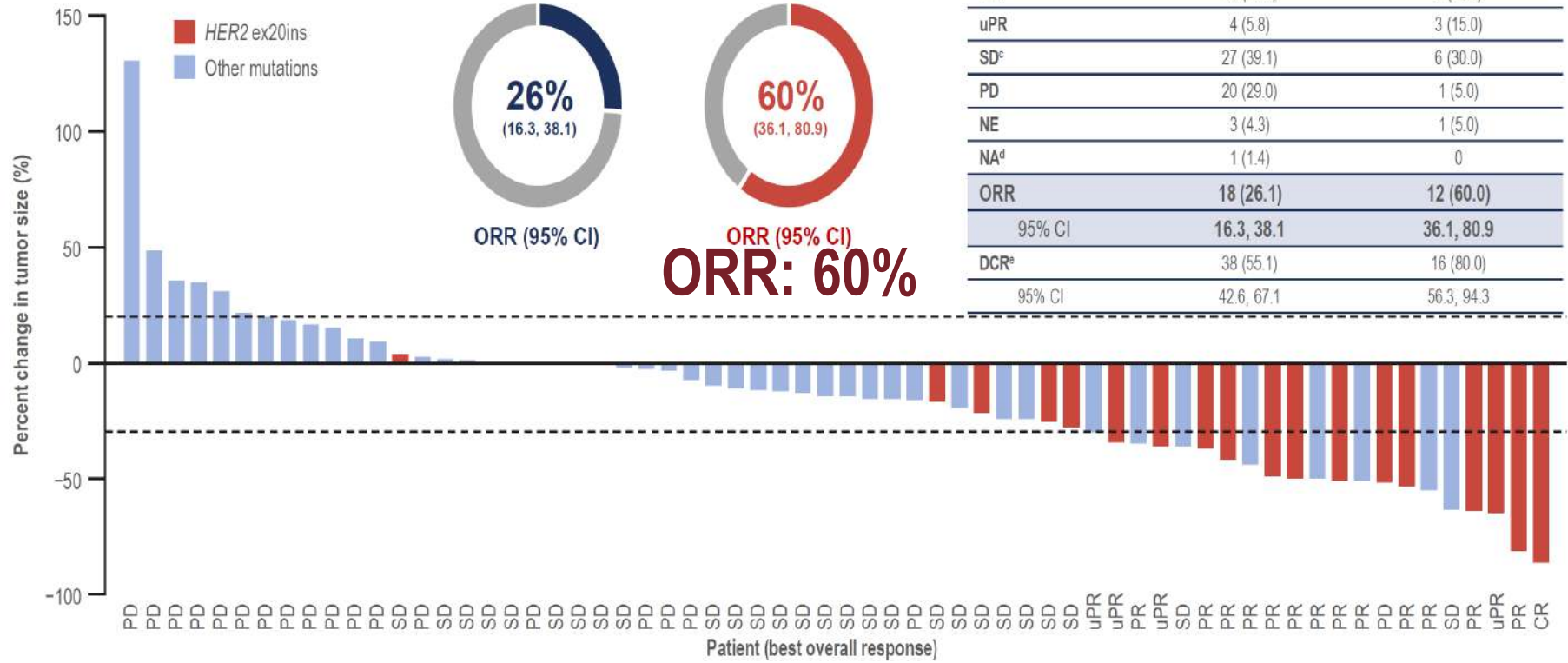
Best % change in target lesion size across all dose levels

N=69pts
All patients

N=20pts
HER2 ex20ins



ORR: 60%



n (%)	All evaluable patients* (n=69)	HER2 ex20ins subgroup ^b (n=20)
CR	1 (1.4)	1 (5.0)
PR	13 (18.8)	8 (40.0)
uPR	4 (5.8)	3 (15.0)
SD ^c	27 (39.1)	6 (30.0)
PD	20 (29.0)	1 (5.0)
NE	3 (4.3)	1 (5.0)
NA ^d	1 (1.4)	0
ORR	18 (26.1)	12 (60.0)
95% CI	16.3, 38.1	36.1, 80.9
DCR ^e	38 (55.1)	16 (80.0)
95% CI	42.6, 67.1	56.3, 94.3

Dose escalation / backfill: safety

n (%)	All grades (N=76)	Grade ≥ 3 (N=76)
Any TRAE	66 (86.8)	19 (25.0)
Most common TRAEs occurring in $\geq 10\%$ of patients		
Diarrhea	57 (75.0)	12 (15.8)
Paronychia	19 (25.0)	0
Dry skin	17 (22.4)	0
Dermatitis acneiform	16 (21.1)	0
Stomatitis	14 (18.4)	1 (1.3)
Pruritus	12 (15.8)	2 (2.6)
Vomiting	12 (15.8)	1 (1.3)
Rash	12 (15.8)	0
Decreased appetite	11 (14.5)	0
Increased amylase	11 (14.5)	2 (2.6)
Nausea	10 (13.2)	0
Hypokalemia	10 (13.2)	5 (6.6)
Increased alanine aminotransferase	8 (10.5)	2 (2.6)

Safety summary

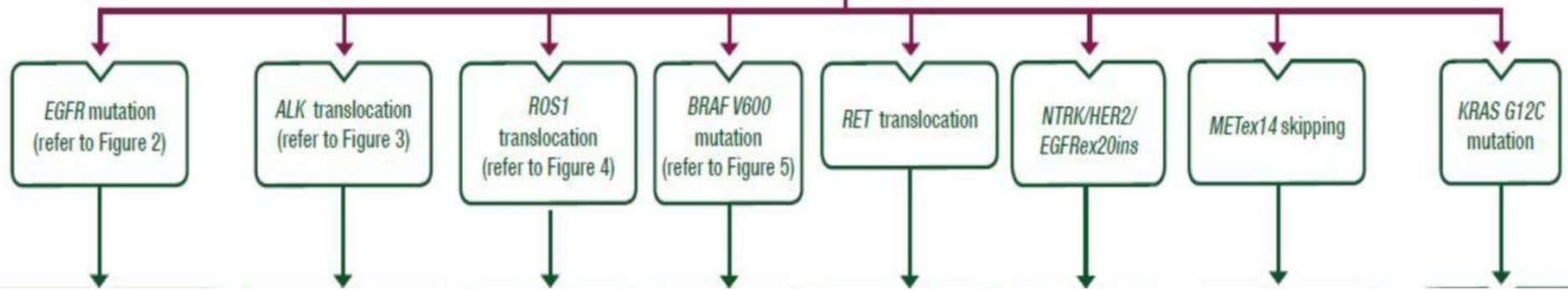
- 76 patients have been treated in dose-escalation / backfill cohorts at the cut-off
- 5 patients with dose-limiting toxicities (3/5 in 40 mg BID, 1/9 in 30 mg BID, and 1/5 in 60 mg QD)
- No discontinuation due to TRAEs
- 20 patients (26.3%) with dose reductions due to TRAEs
- 2 patients (2.6%) with serious TRAEs (diarrhea, vomiting)
- Most AEs were reversible and manageable

Next Generation of EGFRwt sparing HER2 TKIs

Investigational Compounds	IC50	NCT Number	Trial Status	Indication	Sponsor
ELVN-002 ^{2,3}	4.2 nM	NCT05650879	Phase 1 Recruiting	Advanced/metastatic <i>HER2</i> mutation–positive NSCLC	Enliven Therapeutics
JIN-A04 ^{4,5}	11.1 nM	N/A	Preclinical	N/A	J INTS BIO
NVL-330 ⁶	<20 nM	N/A	Preclinical	N/A	Nuvalent
ENT-H1 ⁷	NR	N/A	Preclinical	N/A	Entos

2. ClinicalTrials.gov. <https://classic.clinicaltrials.gov/ct2/show/NCT05650879>. Accessed July 25, 2023; 3. Aujay M, et al. Presented at the AACR Annual Meeting 2023, Orlando, USA, April 14–19, 2023. Poster 4019; 4. Yu M, et al. Presented at the AACR Annual Meeting 2023, Orlando, USA, April 14–19, 2023. Abstract 4029; 5. Chang M. Korea Biomedical Review. Accessed on July 27, 2023; 6. Andrews KL, et al. *Eur J Cancer*. 2022;174S1:S3–S128. 7. Zhao, C, et al. Presented at the AACR Annual Meeting 2023, Orlando, USA, April 14–19, 2023. Poster 4034.

Molecular tests positive (*EGFR/ALK/ROS1/BRAF/RET/NTRK/MET/HER2/EGFR_{ex20ins}/KRAS G12C*)



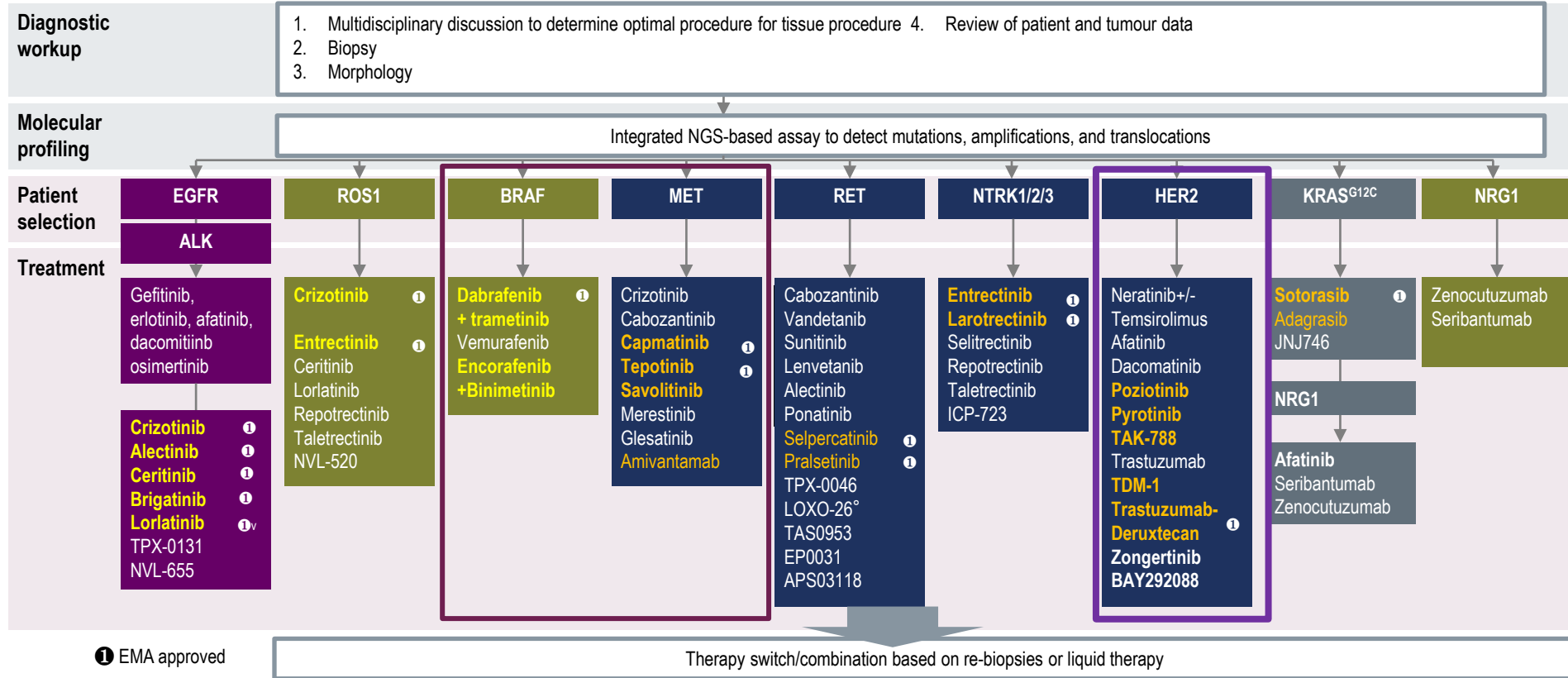
HER2 mutation

2nd line

Trastuzumab–
deruxtecan
[III, B; ESCAT II, B]^o

GREAT ADVANCES HAVE BEEN MADE IN LUNG CANCER THERAPY FOR BRAF, MET AND HER2...

Personalised therapy in advanced-stage NSCLC



THANK YOU !



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