



Clinical Considerations in *EGFR* Mutation–Positive NSCLC: Does Treatment Sequence Matter?

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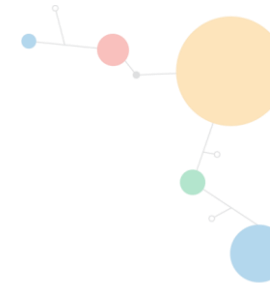




Faculty Disclosure

- Honoraria: AstraZeneca, Boehringer Ingelheim, Bristol-Myers Squibb, Eli Lilly, Merck, Novartis, Pfizer





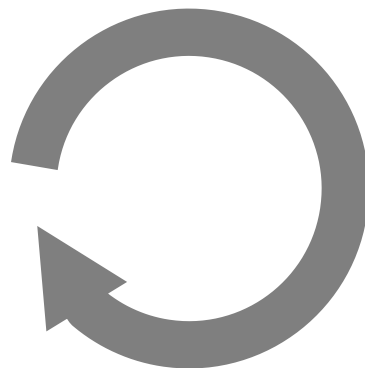
Choosing the Sequence in *EGFR*-Mutant NSCLC

Evidence #1

**TKIs are standard up front
but they are not equal**

Evidence #3

Sequence affects survival



Evidence #2

**Mutational subgroup/
resistance pattern determines
treatment choice**



Choosing the Sequence in *EGFR*-Mutant NSCLC

Evidence #1

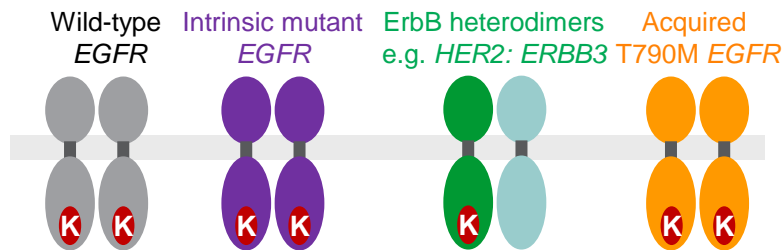
**TKIs are standard up front
but they are not equal**



First-, Second-, and Third-Generation *EGFR* TKIs Are Not Equal: Activity Against *EGFR* Mutations



K Kinase domain



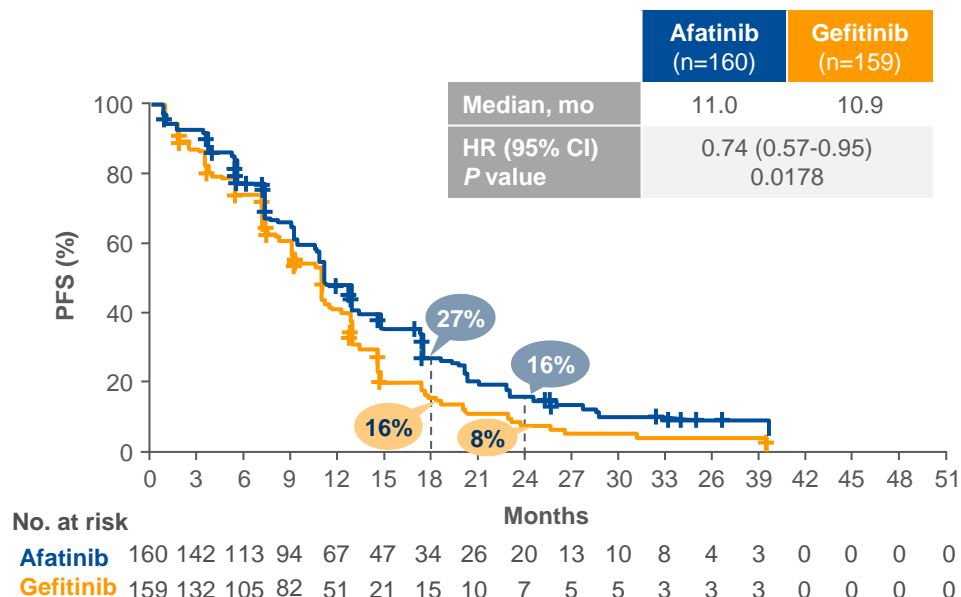
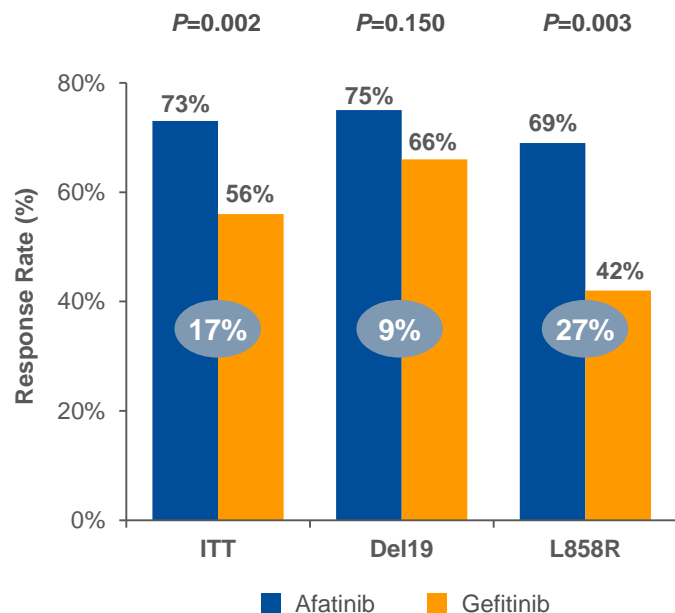
Drug	Metabolised by CYP Enzymes	Possible Drug-Drug Interactions
Erlotinib	✓	✓
Gefitinib	✓	✓
Afatinib	X	X
Dacomitinib	✓	✓
Osimertinib	✓	✓

Drug	Activity range
Erlotinib Gefitinib	<p>1st-generation TKI EGFR inhibition</p> <ul style="list-style-type: none"> Reversible binding to wild-type and mutant EGFR Inactive on T790M mutant
Afatinib Dacomitinib	<p>2nd-generation TKI ErbB family blockade</p> <ul style="list-style-type: none"> Irreversible covalent binding to EGFR, ErbB2 and ErbB4 to inhibit all ErbB family signalling Broader activity to overcome EGFR TKI-resistant mutations
Osimertinib	<p>3rd-generation TKI EGFR mutant-specific inhibitor</p> <ul style="list-style-type: none"> Irreversible covalent binding to mutant EGFR Specificity for EGFR T790M mutant; EGFR wild-type sparing



First- and Second-Generation EGFR TKIs Are Not Equal: LUX-Lung 7

LUX-Lung 7: Afatinib vs gefitinib as first-line treatment of patients with *EGFR* mutation–positive NSCLC: a phase 2B, open-label, randomised controlled trial



No. at risk	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45	48	51
Afatinib	160	142	113	94	67	47	34	26	20	13	10	8	4	3	0	0	0	0
Gefitinib	159	132	105	82	51	21	15	10	7	5	5	3	3	3	0	0	0	0



PFS = progression-free survival; HR = hazard ratio; CI = confidence interval.
Corral et al. *Ann Oncol.* 2017;28 (suppl 2):ii28.

First- and Second-Generation EGFR TKIs Are Not Equal: ARCHER 1050

ARCHER 1050: Dacomitinib vs gefitinib as first-line treatment of patients with EGFR mutation–positive NSCLC: a phase 3, open-label, randomised trial (excluding CNS metastases)

- Advanced NSCLC with *EGFR*-activating mutation(s)
- No prior systematic treatment of advanced NSCLC
- No CNS metastasis
- No prior *EGFR* TKI or other TKI
- ECOG PS 0, 1

R 1:1
(N=452)

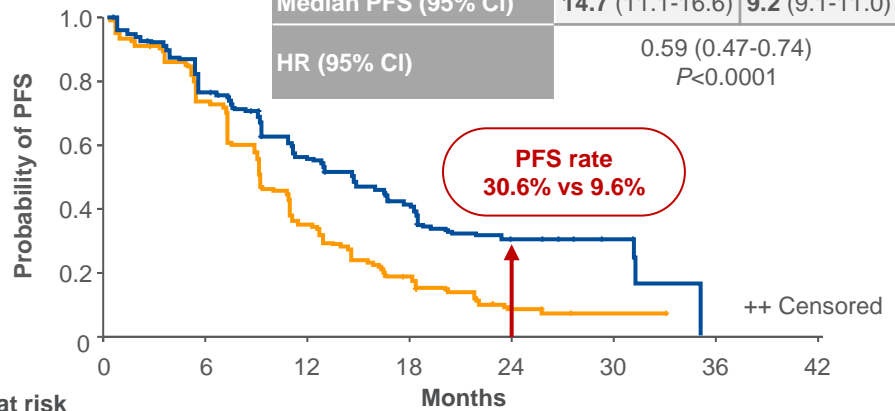
Dacomitinib
45 mg PO QD
(n=227)

Gefitinib
250 mg PO QD
(n=225)

- Primary end point**
PFS by blinded independent review
- ≥256 PFS events
 - PFS HR ≤0.667 (50%↑)
 - 90% power
 - 1-sided α=0.025
 - mPFS: 14.3 vs 9.5 months

PFS: Blinded Independent Review (ITT Population)

	Daco (n=227)	Gef (n=225)
Events, n (%)	136 (59.9%)	179 (79.6%)
Median PFS (95% CI)	14.7 (11.1-16.6)	9.2 (9.1-11.0)
HR (95% CI)	0.59 (0.47-0.74) P<0.0001	



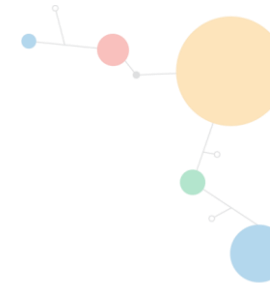
No. at risk	0	6	12	18	24	30	36	42
Dacomitinib	227	154	106	73	20	6	0	0
Gefitinib	225	155	69	34	7	1	0	0



CNS = central nervous system; ECOG PS = Eastern Cooperative Oncology Group performance status; R = randomised; PO = orally; QD = once daily; mPFS = median progression-free survival; ITT = intent-to-treat.

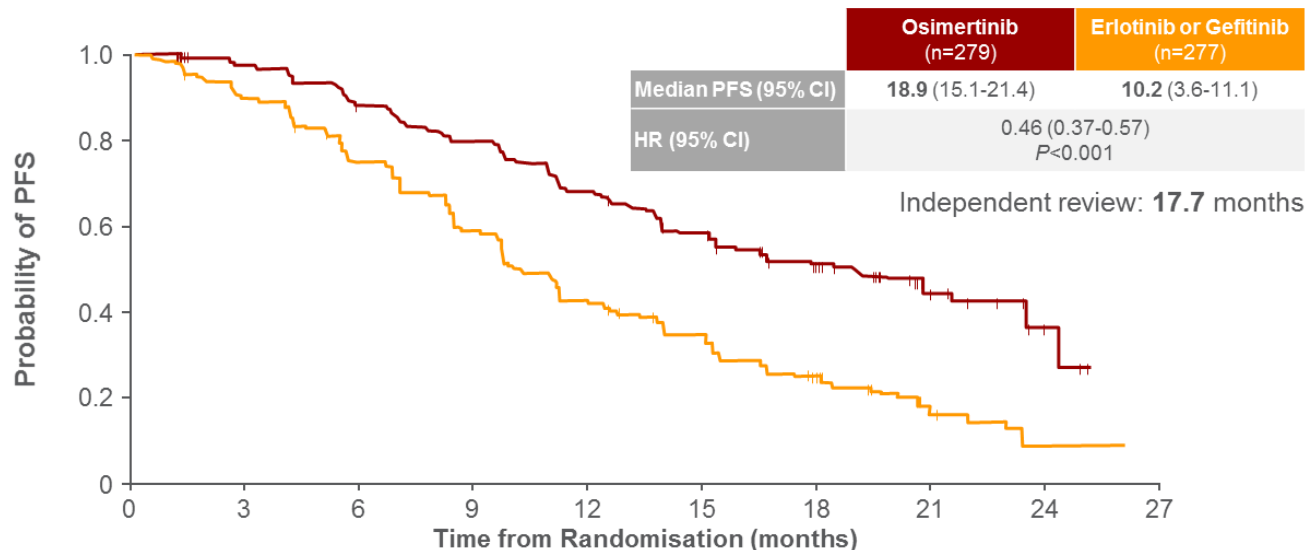
ClinicalTrials.gov. <https://clinicaltrials.gov/ct2/show/NCT01774721>. Accessed March 16, 2018; Mok et al. *J Clin Oncol.* 2017;35(suppl 18):LBA9007; Wu et al. *Lancet Oncol.* 2017;18:1454.

First- and Second-Generation EGFR TKIs Are Not Equal: FLAURA



FLAURA: Osimertinib vs erlotinib or gefitinib as first-line treatment of patients with *EGFR* mutation-positive NSCLC: a phase 3, double-blind, randomised trial

Primary endpoint: PFS (By investigator assessment)



No. at risk

Osimertinib

Erlotinib or Gefitinib

279	262	233	210	178	139	71	26	4	0
277	239	197	152	107	78	37	10	2	0



First-, Second-, and Third-Generation EGFR TKIs Are Not Equal: Safety



Second- or Third-Generation TKIs vs First-Generation TKIs

	LUX-Lung 7 ^{1,2}		ARCHER 1050 ³		FLAURA ⁴	
	Afatinib (n=160)	Gefitinib (n=159)	Dacomitinib (n=227)	Gefitinib (n=225)	Osimertinib (n=279)	First-gen TKI (n=277)
Treatment discontinuation rate	6%	6%	10%	7%	10%	14%
Most common grade ≥3 AEs	Diarrhoea, 12% Rash/acne, 9%	Liver enzyme elevation, 9% Rash/acne, 3%	Acne, 14% Diarrhoea, 8% Paronychia, 7%	Liver enzyme elevation, 12% Dyspnoea, 3%	Diarrhoea, 2% Decreased appetite, 2%	Rash/acne, 7% Liver enzyme elevation, 12%



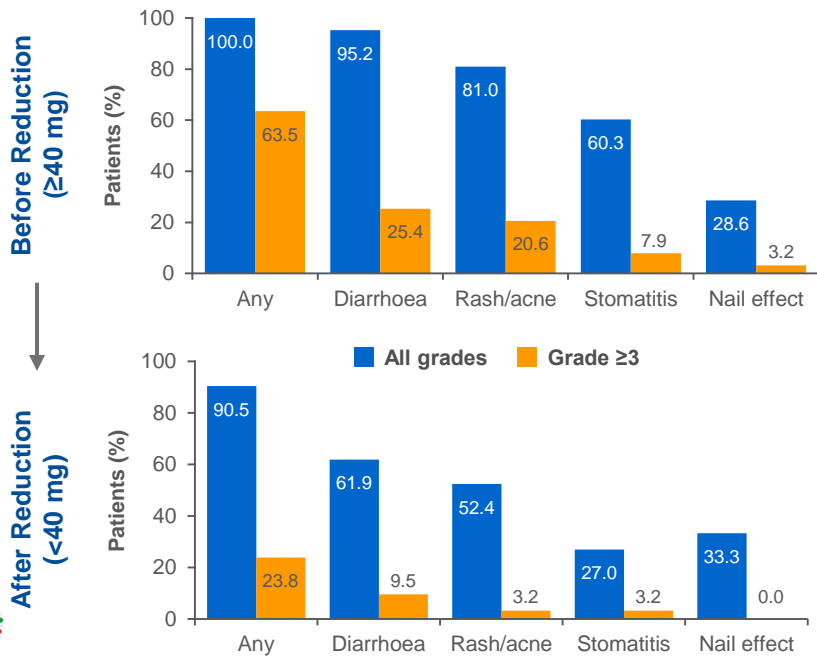
AE = adverse event.

1. Park et al. *Lancet Oncol.* 2016;17:577; 2. Paz-Ares et al. *Ann Oncol.* 2017;28:270; 3. Wu et al. *Lancet Oncol.* 2017;18:1454; 4. Soria et al. *N Engl J Med.* 2018;378:113.

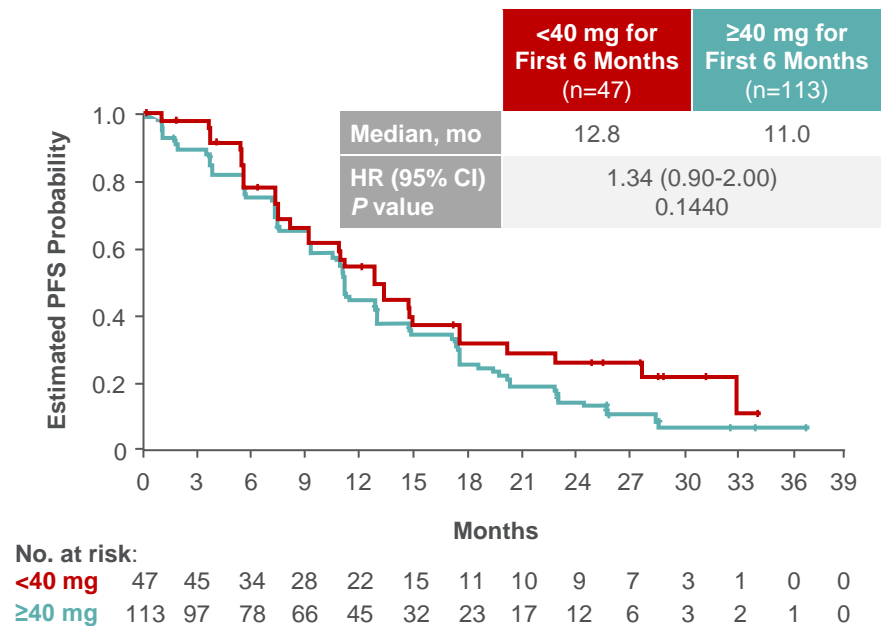
Dose Reduction of Afatinib Reduced Drug-Related AEs Without Compromising Efficacy

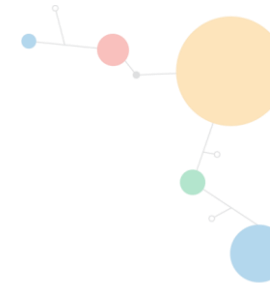


Treatment-Related AEs in Patients Who Had a Dose Reduction From 40 mg (n=63)



PFS in Patients Who Received a Dose Reduction Within the First 6 Months of Treatment

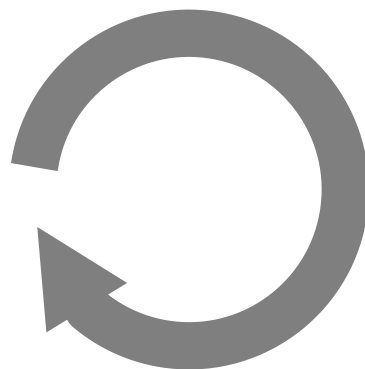




Choosing the Sequence in *EGFR*-Mutant NSCLC

Evidence #1

TKIs are standard up front
but they are not equal



Evidence #2

**Mutational subgroup/
resistance pattern determines
treatment choice**



First-Line: PFS Efficacy of TKIs Is Different in *EGFR* del19/L858R Mutations

- Impact of specific *EGFR* mutations and clinical characteristics on outcomes after treatment with *EGFR* TKIs vs chemotherapy in *EGFR* M+ lung cancer: a meta-analysis

Trial	HR	95% CI		HR	95% CI	
	Exon 19 deletion			Exon 21 L858R substitution		
ENSURE	0.20	0.12-0.33	■	0.54	0.32-0.91	■
EURTAC	0.27	0.17-0.43	■	0.53	0.29-0.97	■
LUX-Lung 3	0.28	0.18-0.44	■	0.73	0.46-1.16	■
LUX-Lung 6	0.20	0.13-0.32	■	0.32	0.19-0.54	■
NEJ002	0.24	0.15-0.38	■	0.33	0.20-0.54	■
OPTIMAL	0.13	0.07-0.24	■	0.26	0.14-0.48	■
WJTOG 3405	0.42	0.26-0.66	■	0.69	0.44-1.07	■
All	0.24	0.20-0.29	◆	0.48	0.39-0.58	◆

TKI Chemo

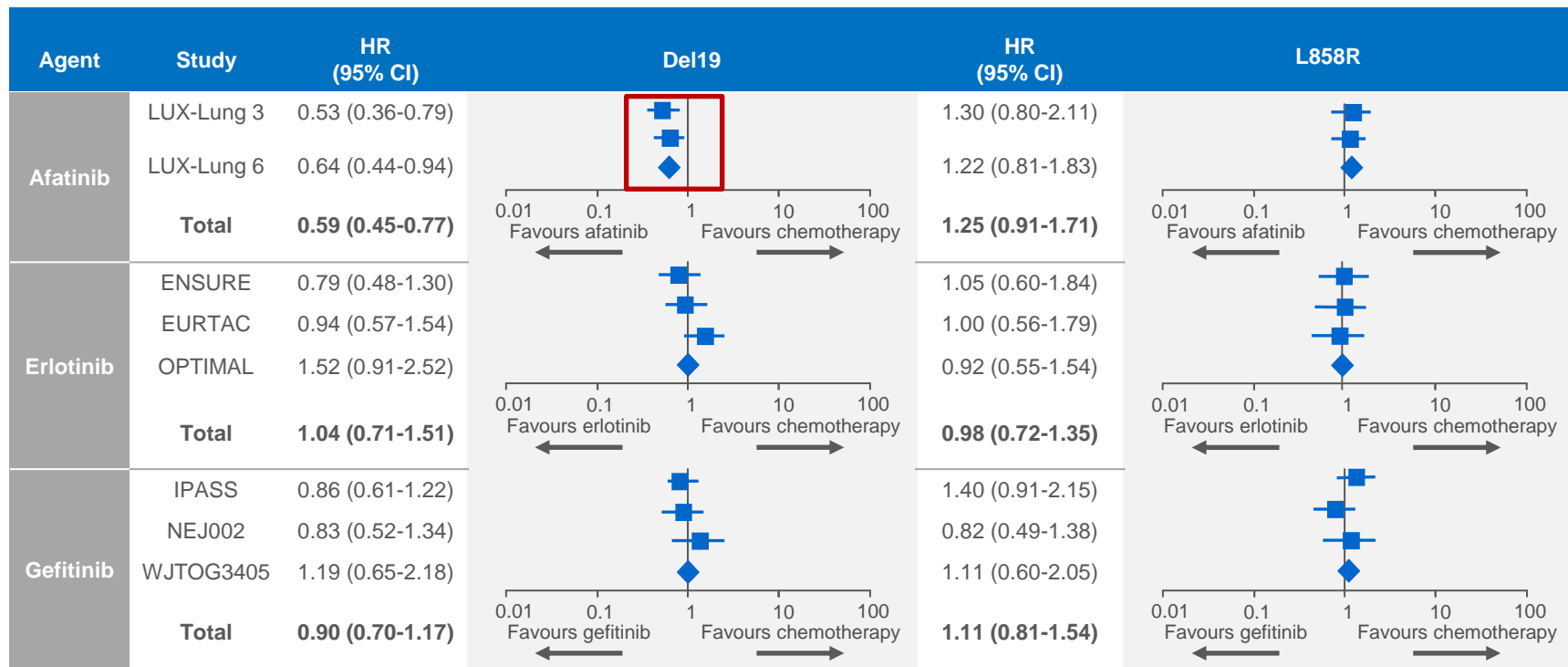
←————→

TKI Chemo

←————→



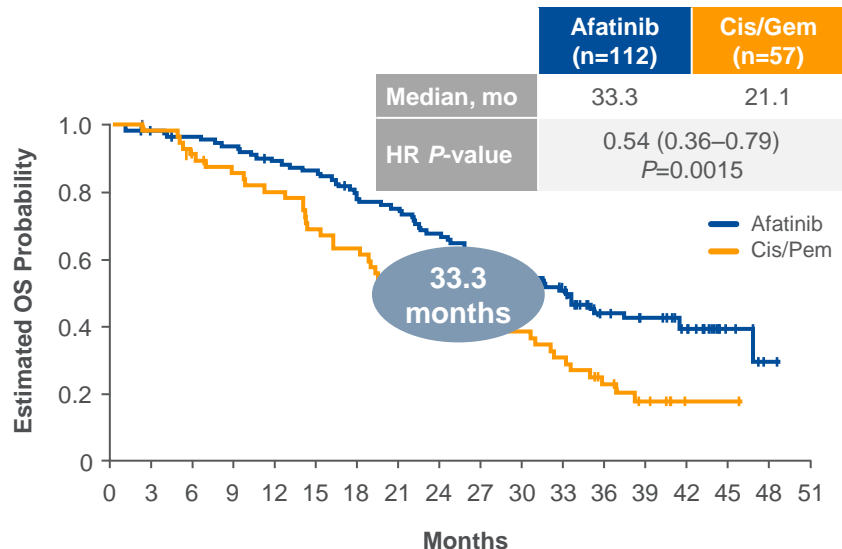
First-Line: OS Efficacy of TKIs Is Different in *EGFR* del19/L858R Mutations



LUX-Lung 3 and LUX-Lung 6: OS in Del19 Subgroup (Prespecified Endpoint)



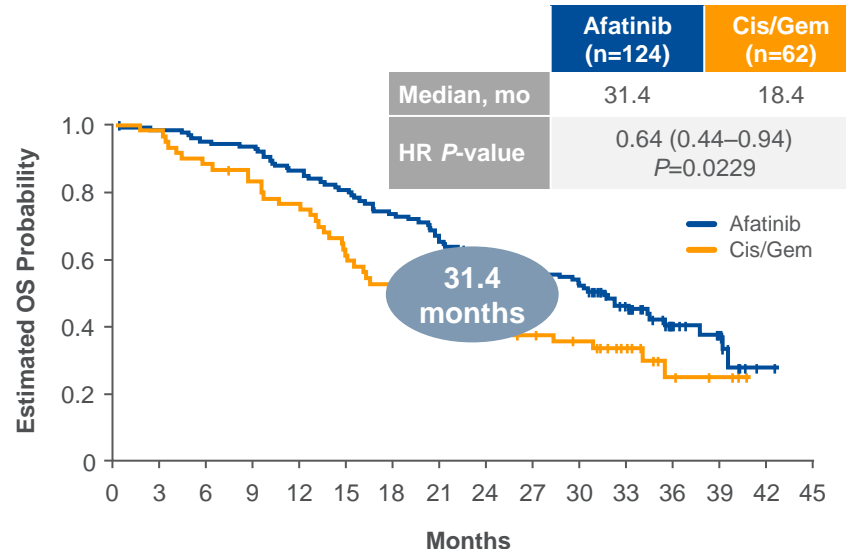
LUX-Lung 3



No. at risk:

	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45	48	51
Afatinib	112	108	105	102	96	93	83	80	72	62	58	51	34	30	21	6	1	0
Cis/Gem	57	55	50	46	43	37	33	27	25	22	20	16	10	6	1	1	0	0

LUX-Lung 6

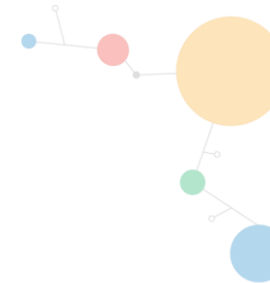


No. at risk:

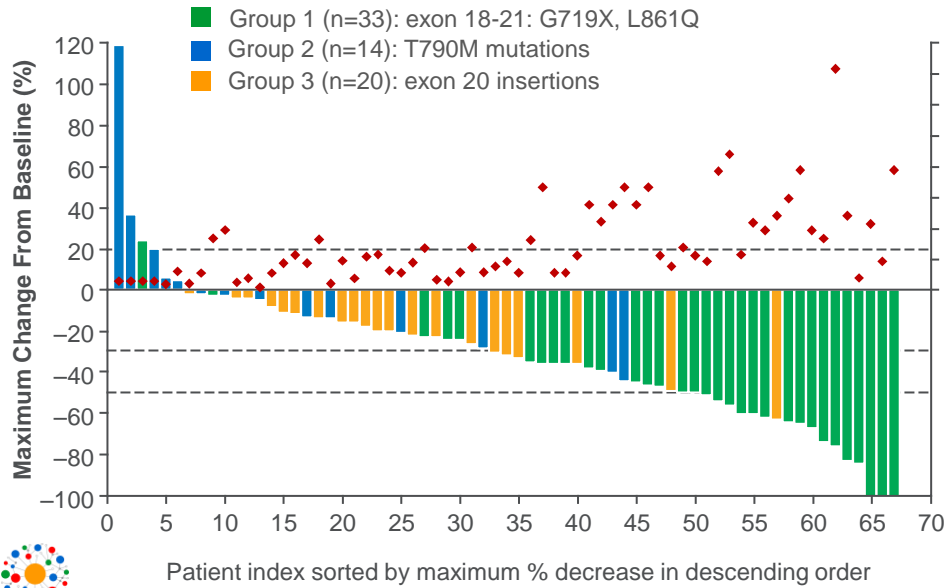
	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45
Afatinib	124	122	118	115	106	99	90	80	73	69	59	39	16	8	1	0
Cis/Gem	62	58	53	49	44	35	30	28	26	21	18	11	4	3	0	0



First-Line: Afatinib Is Effective for Uncommon *EGFR* Mutations



- Clinical activity of afatinib in patients with advanced NSCLC harbouring uncommon *EGFR* mutations: a combined post hoc analysis of LUX-Lung 2, 3, and 6



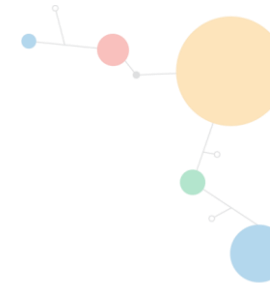
PFS (mo)

	T790M (n=14)	Exon 20 Ins (n=23)	Mut/Dup Exon 18-21 (n=38)	G719X (n=18)	L861Q (n=16)	S768I (n=8)
Response rate (%)	14.3	8.7	71.1	77.8	56.3	100.0
PFS (mo)	2.9	2.7	10.7	13.8	8.2	14.7
OS (mo)	14.9	9.2	19.4	26.9	17.1	NE

Note: A patient may be presented in more than 1 category.

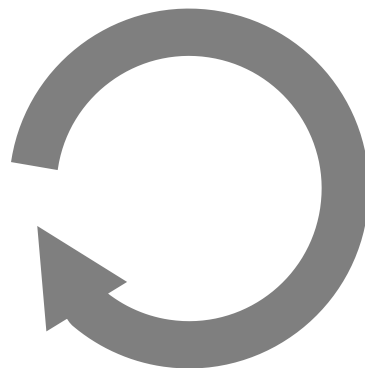


Choosing the Sequence in *EGFR*-Mutant NSCLC



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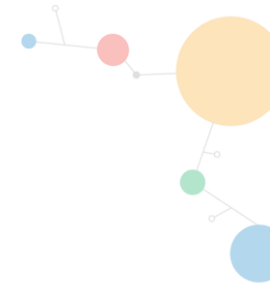


Evidence #2

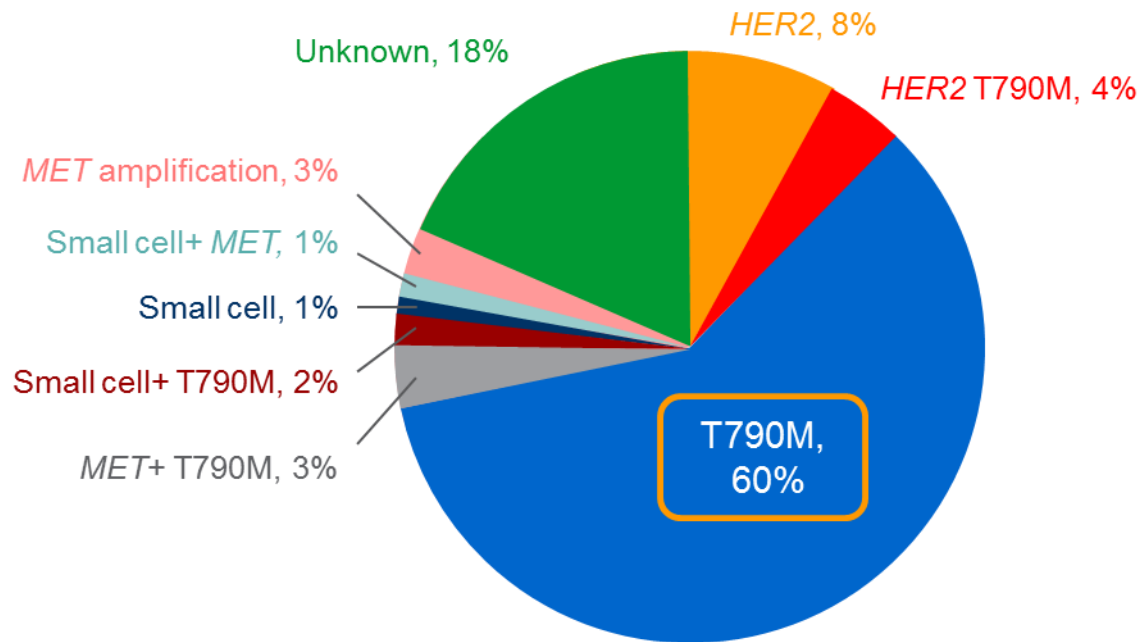
Mutational subgroup/
resistance pattern determines
treatment choice



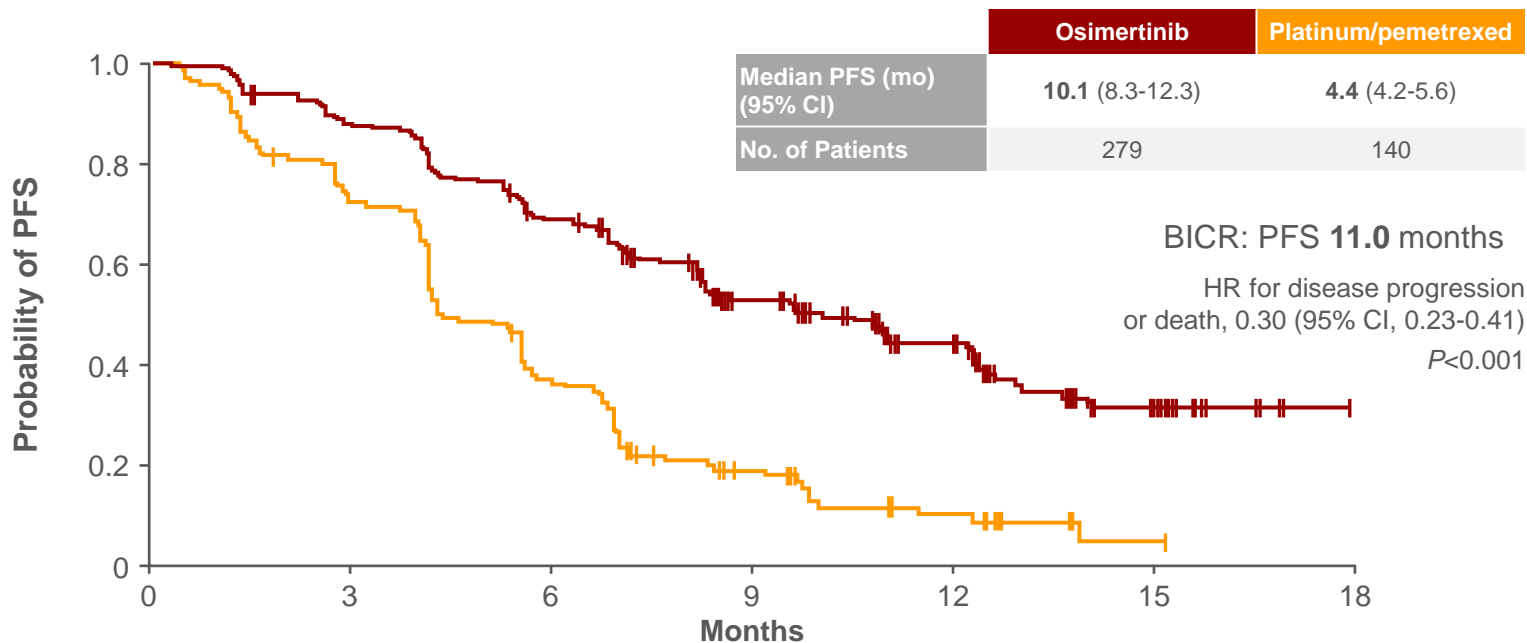
Molecular Mechanisms of Acquired Resistance to First- and Second-Generation EGFR TKIs



- 155 *EGFR*-mutant NSCLC, acquired resistance after TKI
- Molecular analyses on rebiopsy specimen



AURA 3: Osimertinib Standard of Care for T790M+ Acquired Resistance to First- and Second-Generation *EGFR* TKIs



No. at risk

Osimertinib

Platinum/pemetrexed

279

140

240

93

162

44

88

17

50

7

13

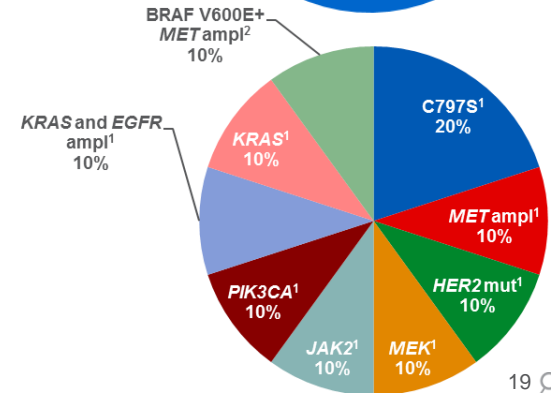
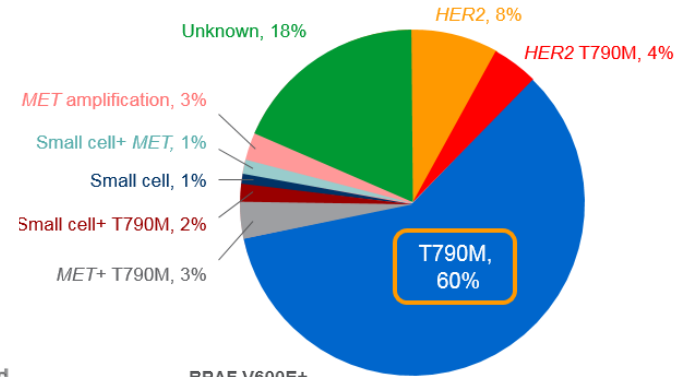
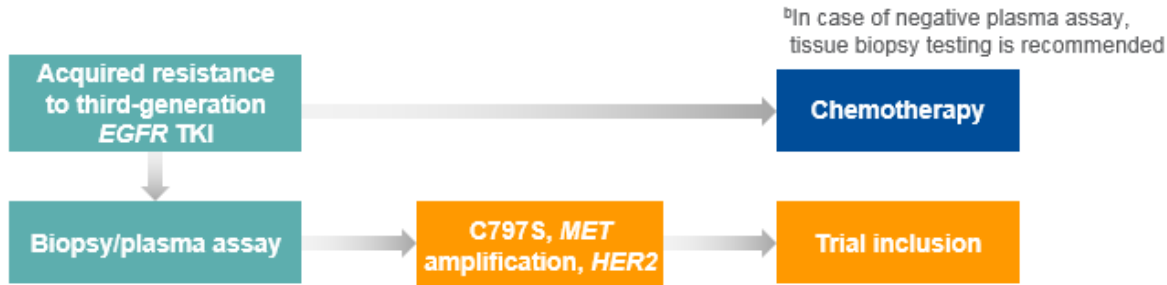
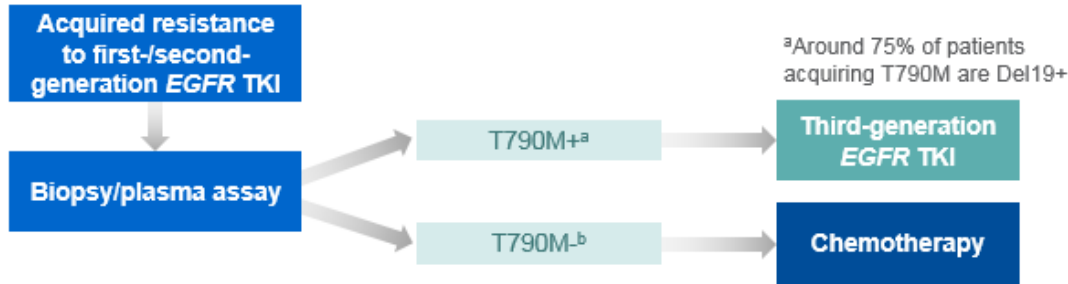
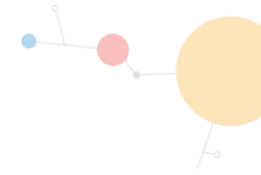
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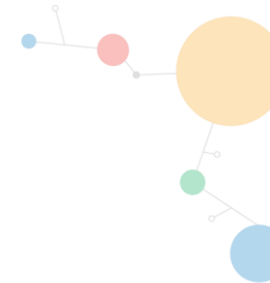
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Biopsy or Plasma May be Used to Determine EGFR T790M Status





Choosing the Sequence in *EGFR*-Mutant NSCLC

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TKIs are standard up front
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Evidence #3

Sequence affects survival

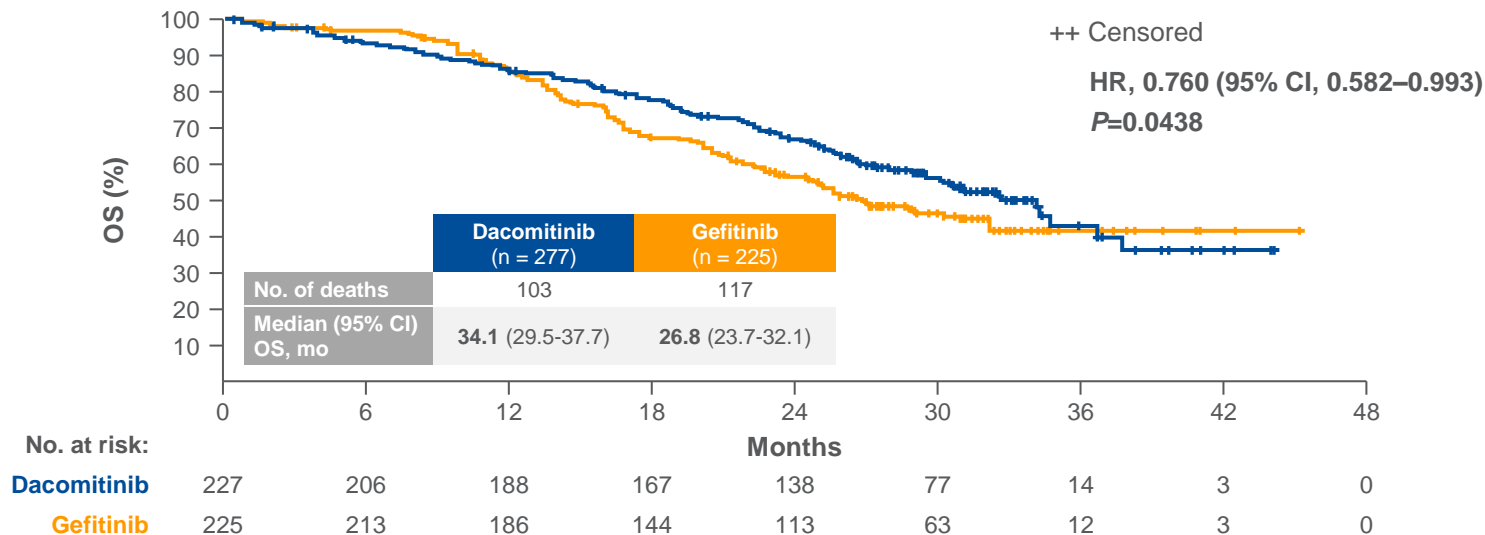


Evidence #2

Mutational subgroup/
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OS in *EGFR*-Mutant NSCLC: ARCHER 1050 Trial (Excluding Brain Metastases)

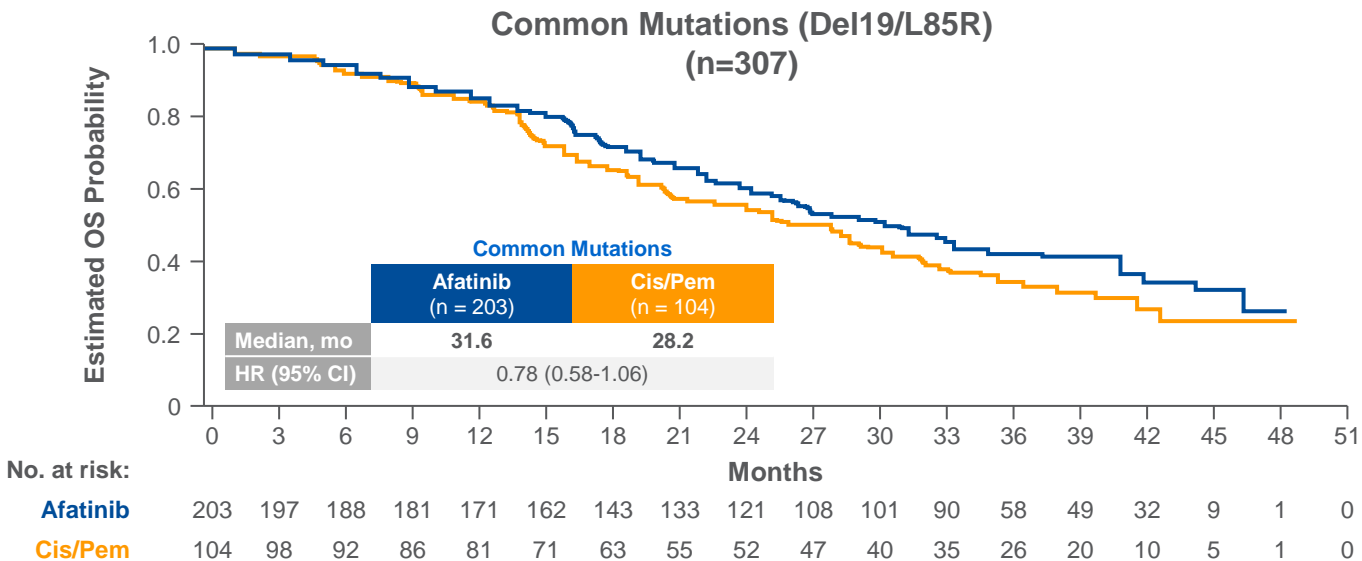


- Third-generation *EGFR* TKIs were used as a first subsequent therapy in 22 patients (9.7%) in the dacomitinib arm and in 25 patients (11.1%) in the gefitinib arm
- Median OS in patients subsequently treated with third-generation TKIs was **36.7** (95% CI: 30.1-NR) months in the dacomitinib arm





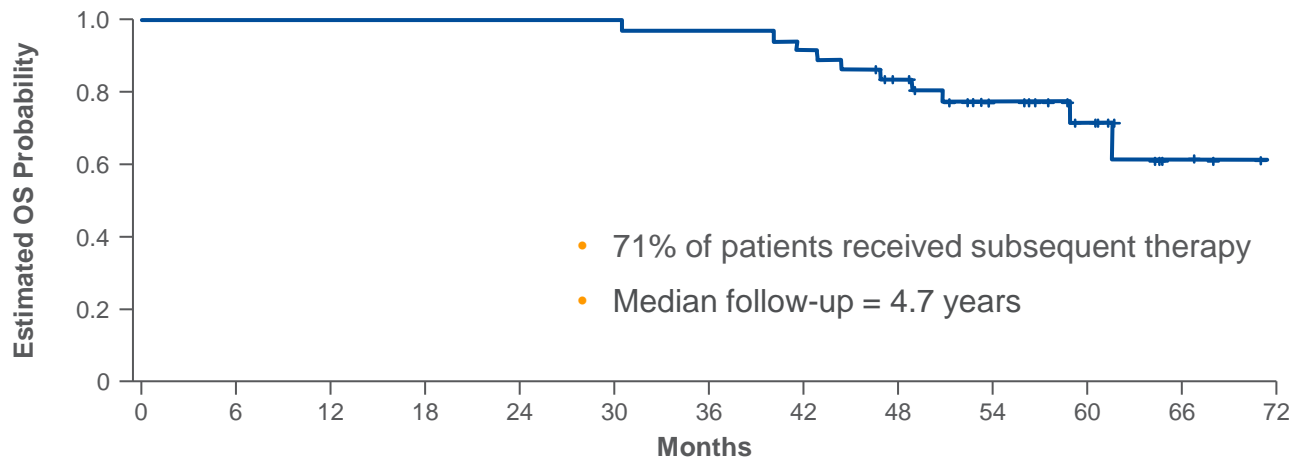
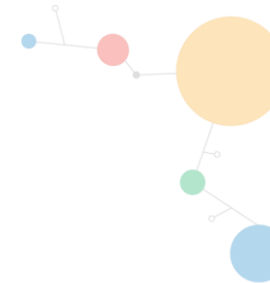
OS in *EGFR*-Mutant NSCLC: LUX-Lung 3 Trial



- OS was up to **41.5 months** in del19 patients in countries with a **universal healthcare reimbursement** policy



OS in Patients in LUX-Lung 3, 6, and 7 Treated Subsequently With Osimertinib



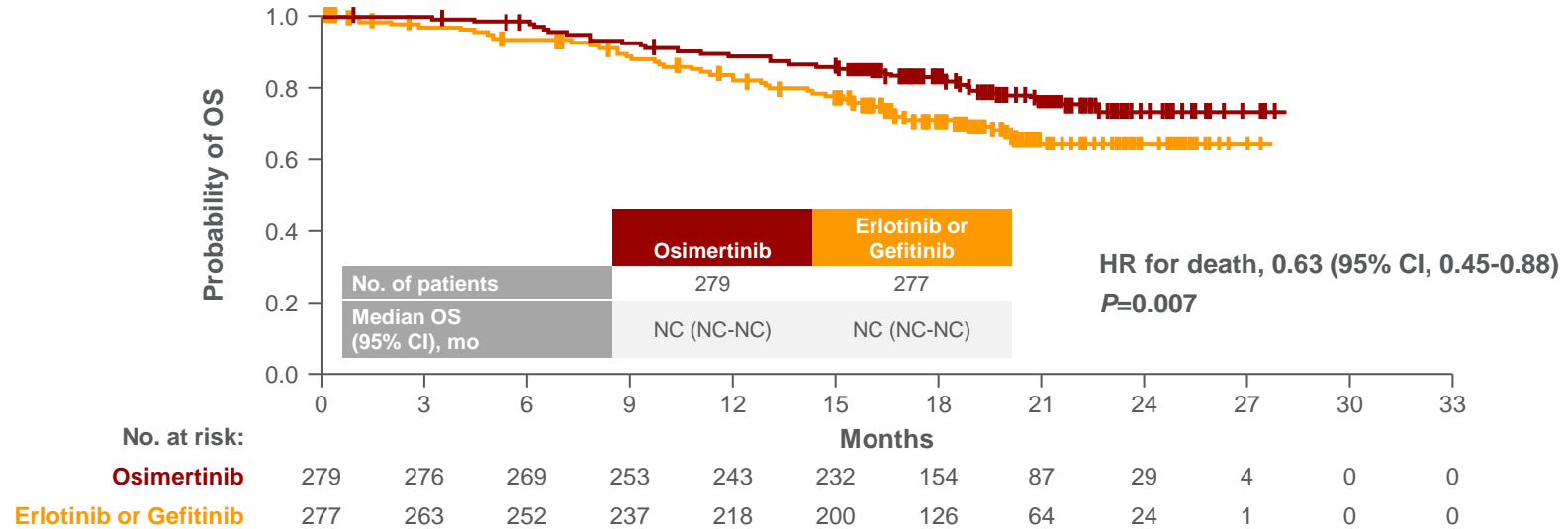
No. at risk:

Afatinib

37 37 37 37 37 37 36 35 28 20 12 3 0



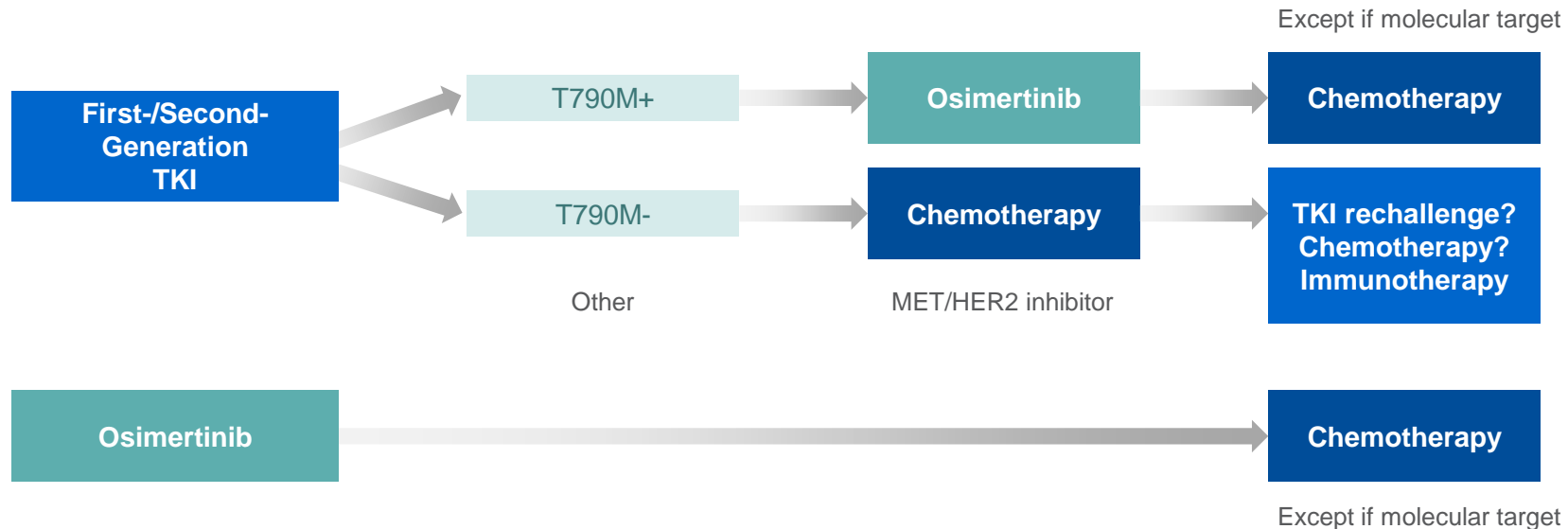
OS in Patients Treated With First-Line Osimertinib



FLAURA OS DATA ARE IMMATURE (25% MATURITY)



Treatment Sequences in *EGFR*-Mutant NSCLC After First-Line *EGFR* TKI



NEED MATURE OS AND TREATMENT SEQUENCES FROM FLAURA and AURA 3





How Do We Optimise Sequence?

Clinical factors

- CNS disease and progression
 - Data for afatinib and osimertinib show delay in onset and progression of CNS metastases
- Loss of patients from one line to another
- Treatment of oligoprogression, treatment beyond PD

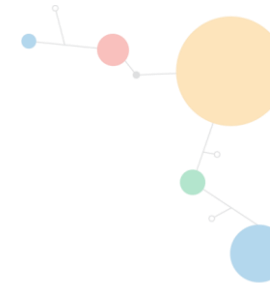
Optimisation of treatments

- Antiangiogenics
- Anti-*EGFR* antibodies
- Chemotherapy + TKI

Understanding of biology

- *EGFR* mutational subgroups (eg, del19, L858R, uncommon mutations) determine treatment choice
- Resistance mechanisms impact subsequent therapy choice





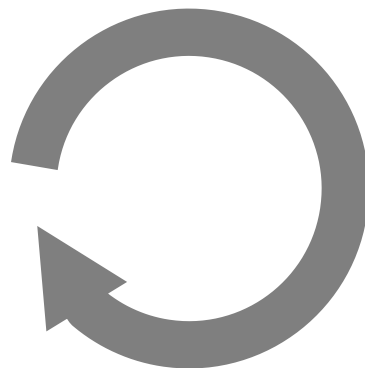
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