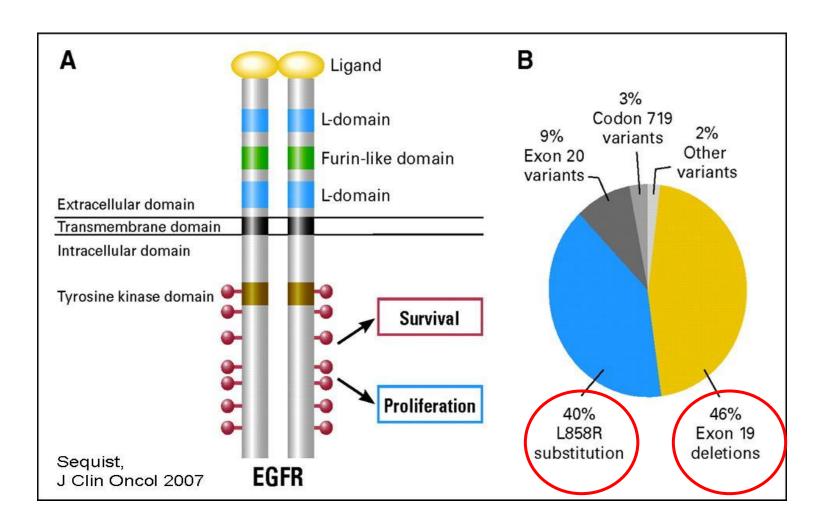
Mechanisms of resistance to HER targeting drugs Lung cancer

Fortunato Ciardiello Second University of Naples

EGFR mutations



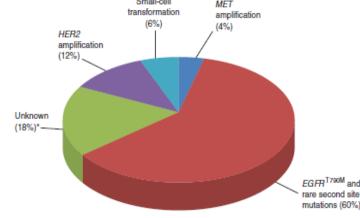
1st line: EGFR-TKIs vs CT

Author	Study	N (EGFR mut. +)	EGFR-TKI	Median PFS* (months)	PFS* HR
Mok	IPASS	261	Gefitinib	9.8 vs. 6.4	0.48
Lee	First-SIGNAL	42	Gefitinib	8.4 vs. 6.7	0.54
Mitsudomi	WJTOG 3405	177	Gefitinib	9.2 vs. 6.3	0.49
Maemondo	NEJGSG002	228	Gefitinib	10.8 vs. 5.4	0.30
Zhou	OPTIMAL	154	Erlotinib	13.1 vs. 4.6	0.16
Rosell	EURTAC	175	Erlotinib	9.7 vs. 5.2	0.37
Yang	LUX-Lung 3	345	Afatinib	11.1 vs. 6.9	0.58
Wu	LUX-Lung 6	364	Afatinib	11.0 vs 5.6	0.28
Yang	Pooled analysis LL3- LL6	1340	Afatinib		Median OS 27.3 vs 24.3 months
Shi	CONVINCE	296	Icotinib	NA	NA
Park	LUX-Lung 7	319	Afatinib vs Gefitinib	11 vs 10.9	0.73

Resistance to EGFR TKI

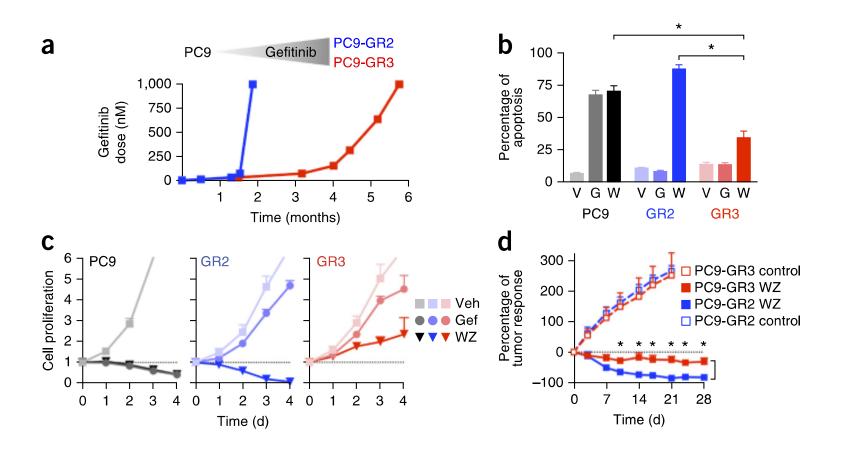
Regardless of extent of initial response, resistance to TKI develops invariably due to

- T790 mutation (~ 50% to 60%)
- HER 2 mutation (8%-12%)
- MET amplification (~ 5% to 20%)
- Conversion to SCLC (< 5%)

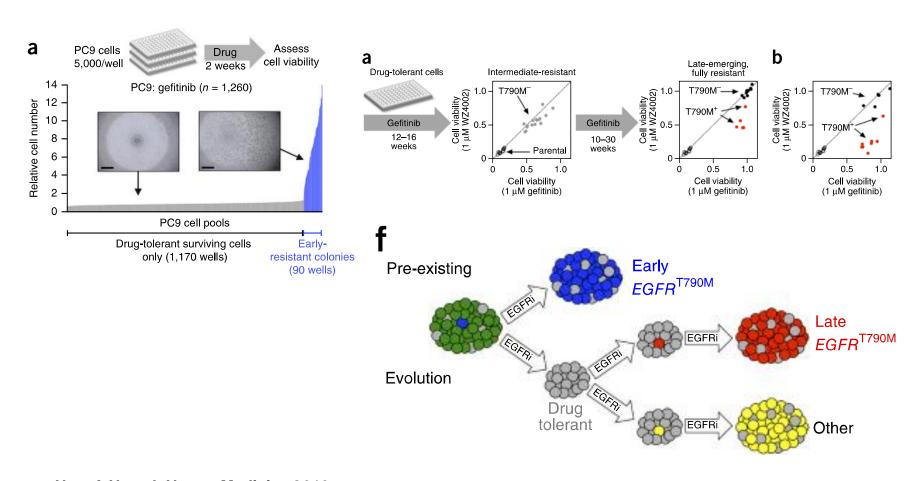


- EGFR mutated tumors might progress rapidly upon withdrawal of TKI
- Role of continuation of TKI beyond progression is under evaluation
- Afatinib, AZ9291, CO1686, HM-61713 targets T790 mutation

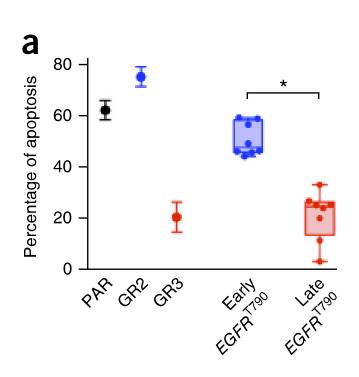
Variable sensitivity of gefitinib-resistant EGFRT790M-positive PC9 cell lines to EGFR inhibition

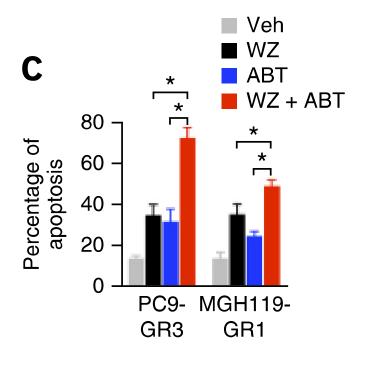


Tumor cells can follow distinct evolutionary paths to become resistant to EGFR inhibition

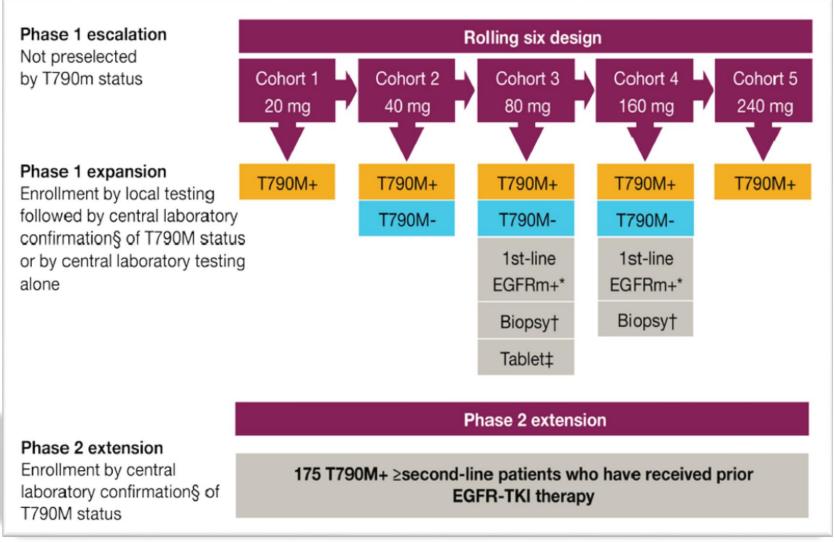


Inhibition of BCL-XL and BCL-2 restored sensitivity to third generation EGFR inhibitor



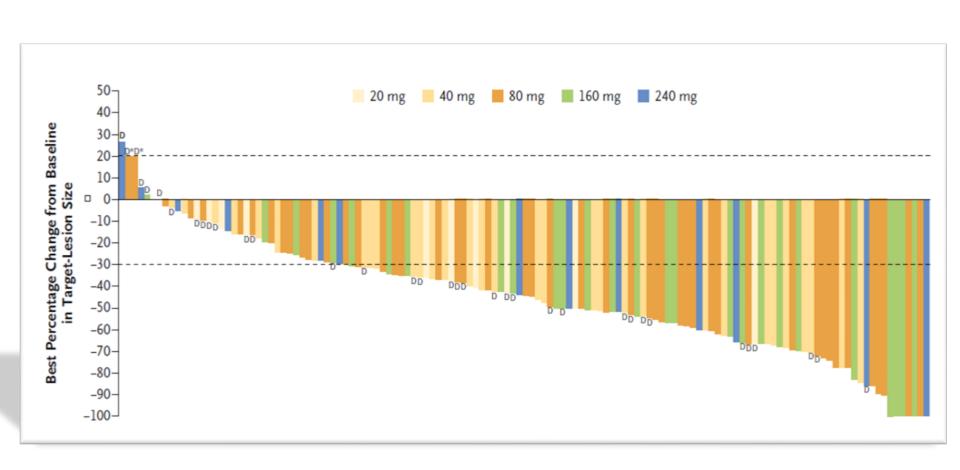


AURA (AZD9291): Overall study schema



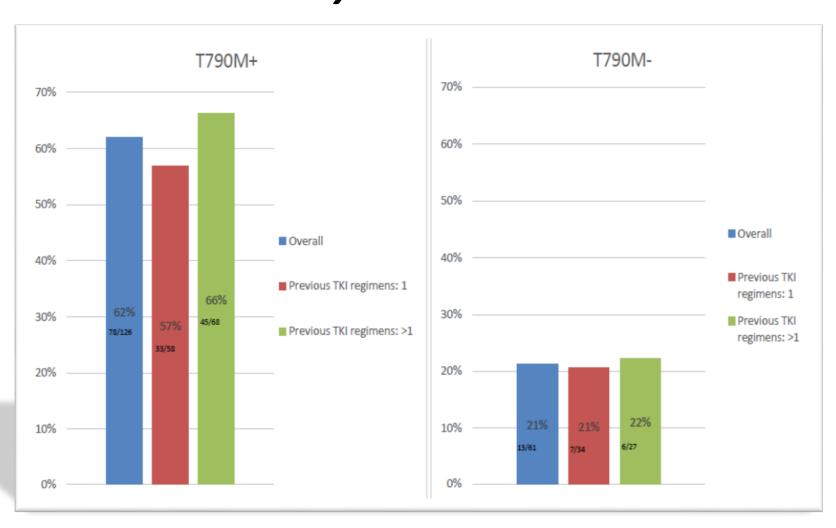
Janne PA, NEJM 2015

AURA (AZD9291): Best % change in target-lesion size



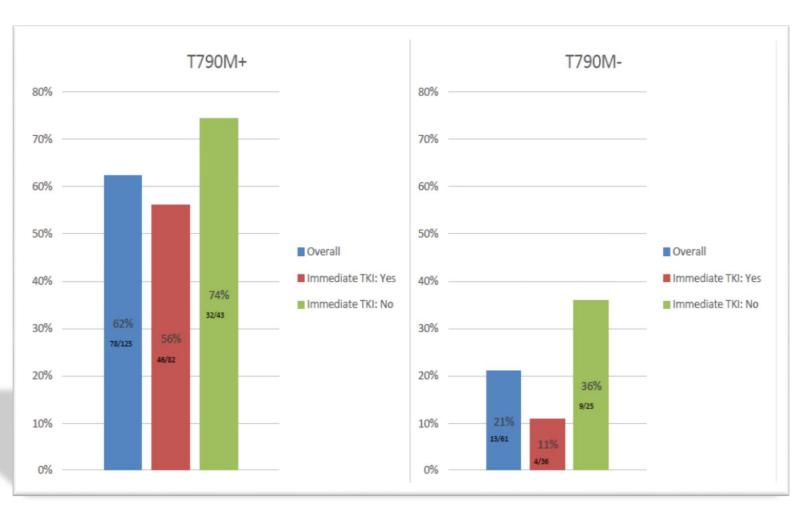
AURA (AZD9291):

RR in patients treated with 1 or >1 prior EGFR-TKI and centrally tested T790M status

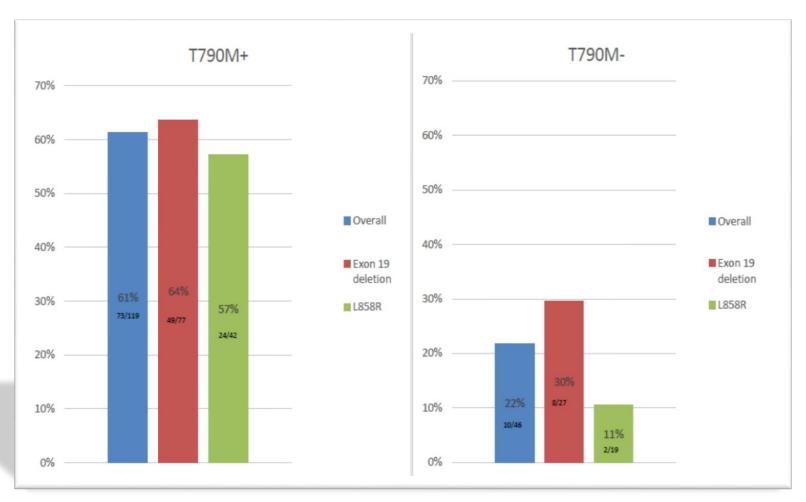


AURA (AZD9291):

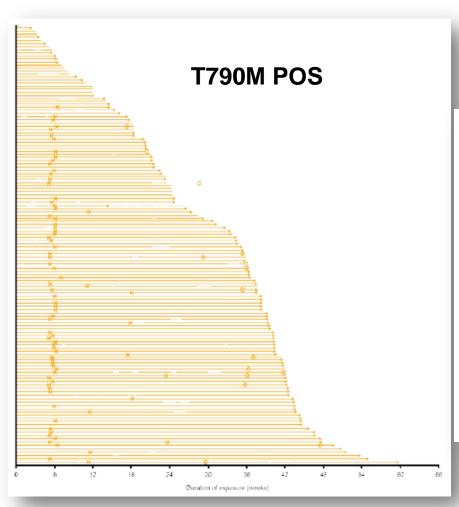
RR in treated patients immediate vs no immediate prior EGFR-TKI and centrally tested T790M status

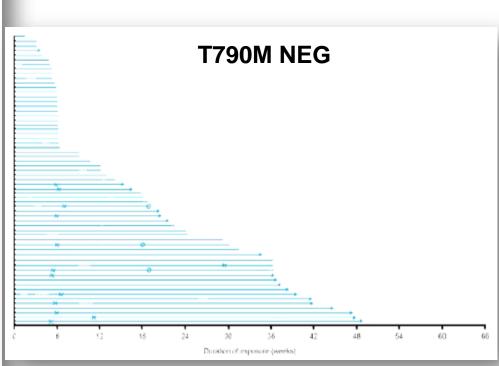


AURA (AZD9291): RR in patients with EGFR mutation type L858R versus exon 19 deletion and T790M status

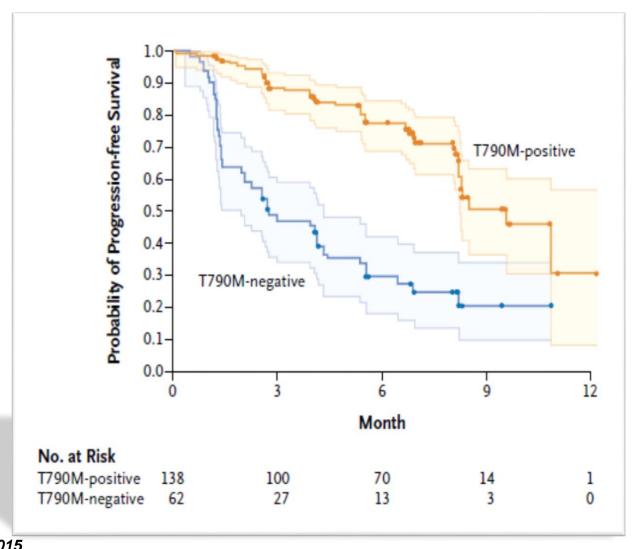


AURA (AZD9291): Duration of treatment and response by centrally tested T790M

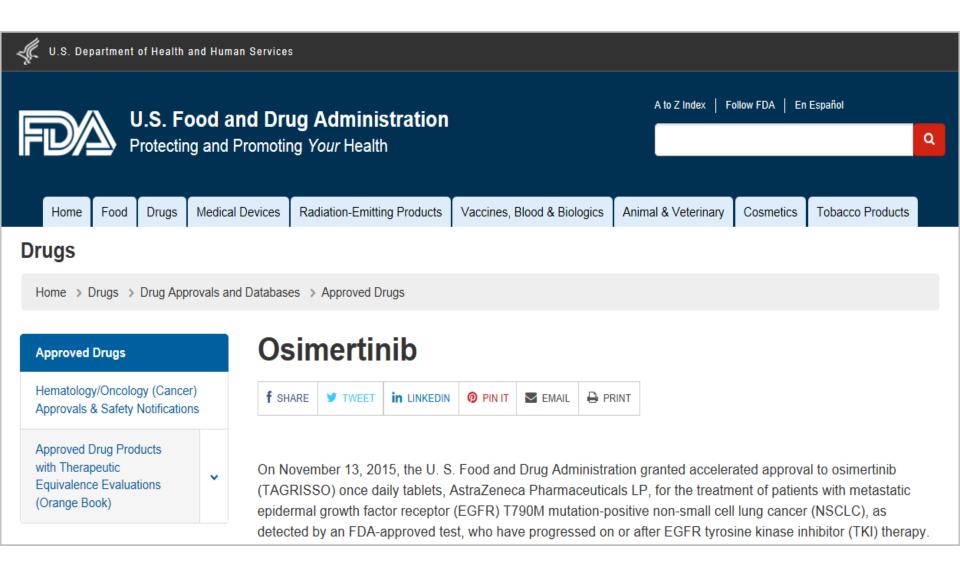




AURA (AZD9291): PFS according to status with respect to EGFR T790M

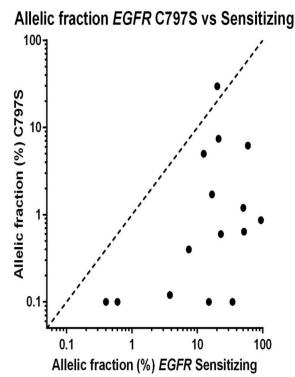


Osimertinib



C797S for acquired resistance to 3rd generation TKIs

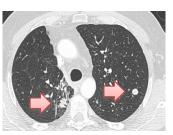
- 67 patients met the following two eligibility criteria for acquired resistance analysis:
 - T790M positive on plasma or tumor genotyping at enrollment
 - Detectable EGFR-TKI-sensitizing mutation in plasma at progression on AZD9291
- Of those, 15 (22%) had detectable C797S on ddPCR, all with detectable T790M
- C797S was more common with EGFR exon 19 del (13/43, 30%) vs those with L858R (2/24, 8%, p=0.06)



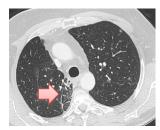
HER2 amplification at PD after AZD 9291

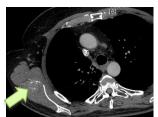
- 54-year-old man
- Former smoker (20 patientyears)
- Adenocarcinom a
- Metastatic to brain, bone
- EGFR Ex19del

PRE-GEFITINIB IMAGING UNAVAILABLE

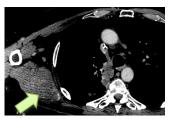








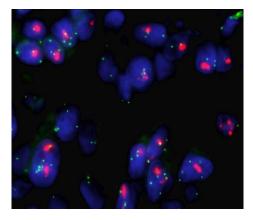




at PD after AZD 9291

HER2

HER2 amplification by CGH: log ratio x3.32



HER2/CEP17: 6.65
HER2 in red; centromere 17 in green

NGS: Ion Torrent Personal Genome Machine® CGH: Agilent technology HER2-FISH: Dako DNA probe

kit

Gefitinib

LUNG

EGFR Ex19del T790M negative



LUNG BIOPSY

EGFR Ex19del and T790M positive

No HER2 amplification (CGH) AZD9291 (80 mg)

_

SCAPULAR

EGFR Ex19del
T790M negative

SCAPULAR BIOPSY

EGFR Ex19del
T790M negative

LUNG

BIOPSY

EGFR Ex19del

T790M negative

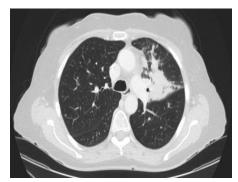
C797S negative

HER2 amplification

MET amplification at PD after AZD 9291

- 69-year-old female with EGFR-mutant NSCLC metastatic to liver, adrenal, bones who had progression after first-line chemotherapy and subsequent erlotinib
- Resistance biopsy was inadequate for genotyping, but plasma genotyping positive for L858R (26%) and T790M (4%)
- Initiated AZD9291 and responded on the first scan (-40%) but progressed after 24 weeks
- Resistance biopsy undergone for targeted NGS:
 - Positive for L858R, negative for T790M, positive for MET amplification
 - MET protein overexpression also seen on IHC

Pre-AZD9291 plasma genotype: L858R (26%) T790M (4%)





Progressio n tumor genotype: L858R T790M negative MET amplified

Baselin
Oxnard GR, WCLC 2015 e

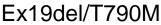
4 months

months

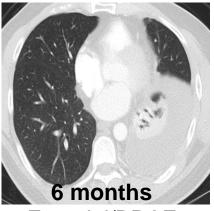
BRAF V600E at PD after AZD 9291

- 49-year-old male with metastatic NSCLC positive for EGFR exon 19 deletion
- Developed resistance to first-line erlotinib after 11 months, T790M positive biopsy
- Had a confirmed PR to AZD9291 but growth of lung mass, effusion after 5 months
- Targeted NGS of progression biopsy shows exon 19 deletion (8% of reads), no T790M, BRAF V600E (6% of reads)
 - A patient-derived xenograft is in development

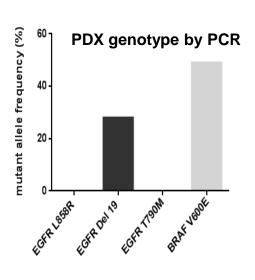




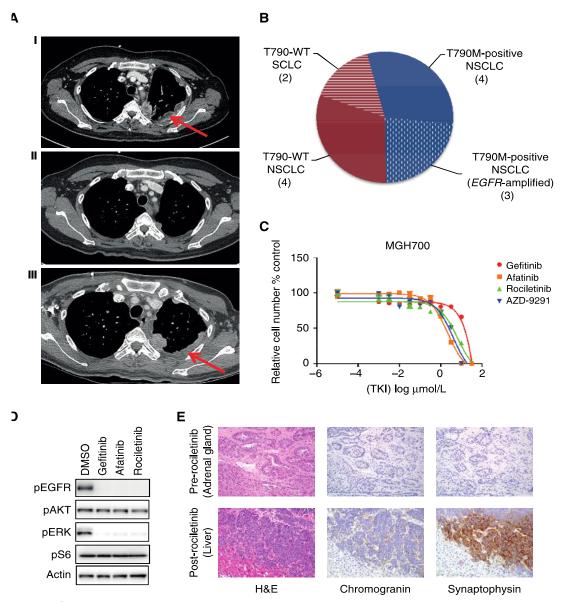




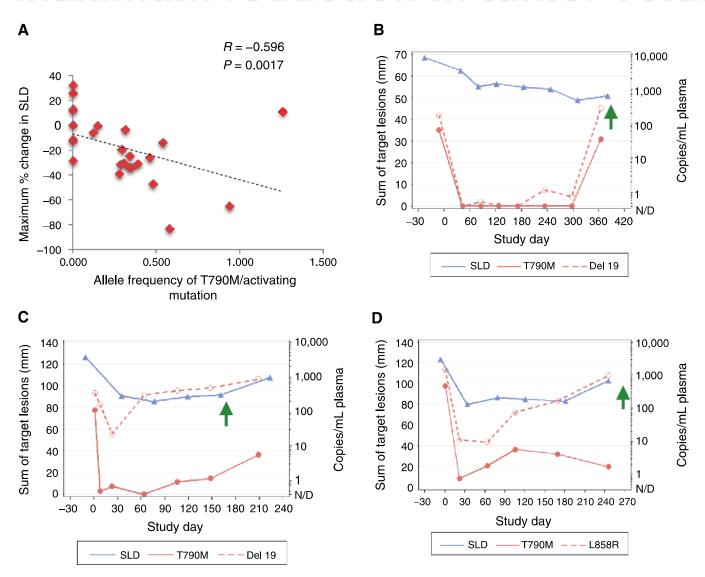
Ex19del/BRAF V600E



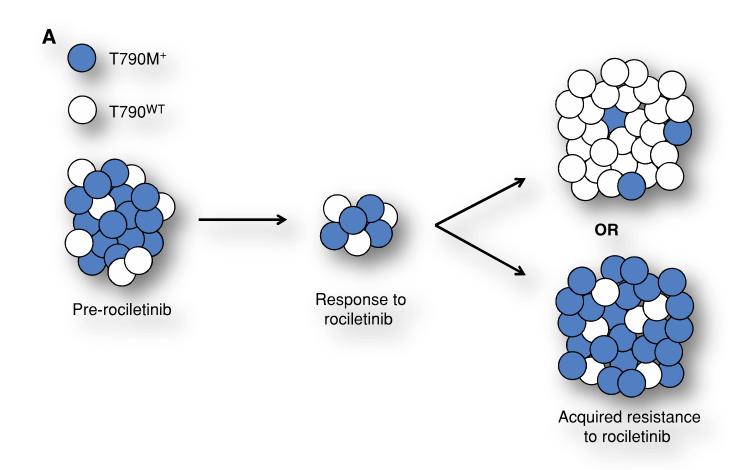
Rociletinib resistance: 12 case series



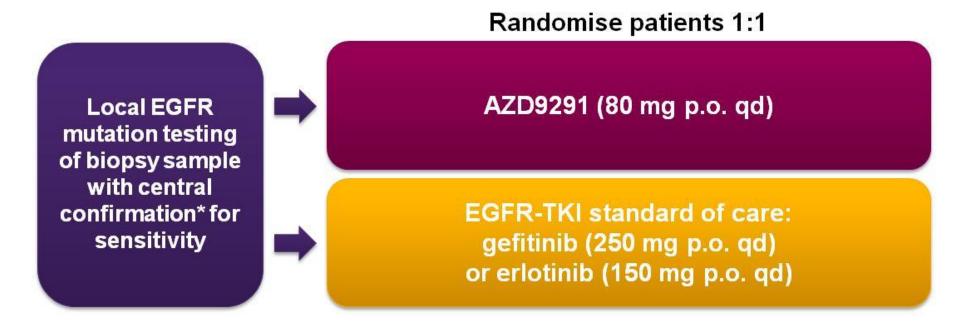
Correlation with allelic fraction of T790M and maximum reduction in tumor volume



Rociletinib resistance: intratumoral heterogeneity



FLAURA: Study design



Algorithm at PD during EGFR TKIs

