

# Long term therapeutic plan for NSCLC harboring EGFR mutation

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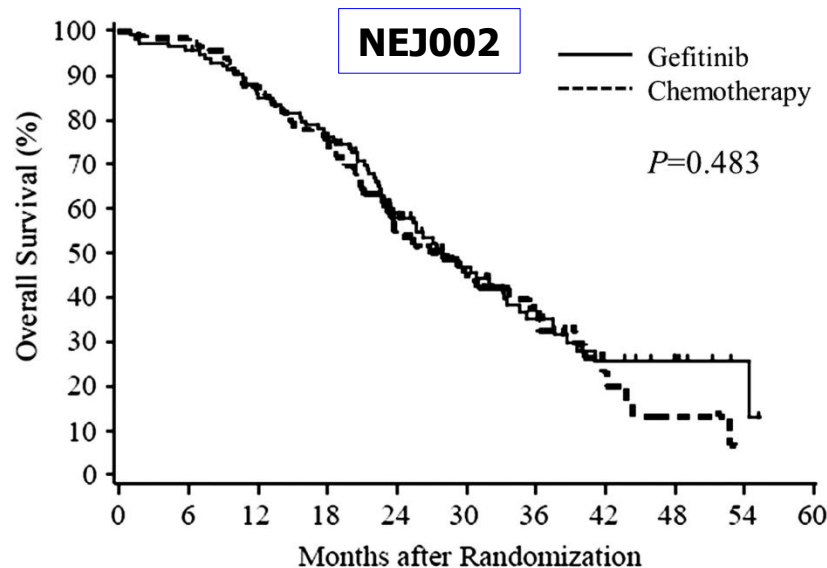


# EGFR-TKI therapy is effective as first-line treatment of advanced NSCLC with EGFR<sup>mut+</sup>

## A meta-analysis from six phase III RCTs

■ **Meta-analysis** of IPASS, First-SIGNAL, NEJ002, WJTOG3405, OPTIMAL, EURTAC

First-line therapy					
	N	EGFR-TKI	Chemotherapy	RR or HR	p-value
<b>RR</b>	1,021	66.6%	30.6%	5.68 [3.17-10.18]	<0.001
<b>PFS</b>	1,006	9.5mo	5.9mo	0.37 [0.27-0.52]	<0.001
<b>OS</b>	852	30.5mo	23.6mo	0.94 [0.77-1.15]	0.57



**Data from NEJ002 clinical trial**  
Stage IV, NSCLC EGFR<sup>mut+</sup>, Japan

<b>1-year survival</b>	≈ <b>86%</b>
<b>2-Year survival</b>	≈ <b>56%</b>
<b>3-Year survival</b>	≈ <b>35%</b>
<b>4-Year survival</b>	≈ <b>19%</b>
<b>5-Year survival</b>	≈ <b>10% (?)</b>

# Contents

- ① **First-line EGFR-TKI** in NSCLC with EGFR mutation
- ② **EGFR-TKI maintenance therapy**
- ③ **NCCN guideline 2013 updates**
- ④ **Clinical modes** of EGFR-TKI failure
- ⑤ **Continuation or Retreatment** of EGFR-TKI
- ⑥ Mechanisms of **EGFR-TKI resistance**
- ⑦ Reversible drug tolerance
- ⑧ Intratumoral heterogeneity
- ⑨ **Refractory brain metastases** to standard-dose EGFR-TKI
- ⑩ **Subsequent chemotherapy** after EGFR-TKI failure

# Efficacy of EGFR-TKI or chemotherapy as 1<sup>st</sup>-line therapy in NSCLC with EGFR<sup>mut+</sup>

Patient group	N	PFS (months)			OS (months)			
		Gefitinib	Chemotherapy	HR	Gefitinib	Chemotherapy	P	
Non-randomized pooled analysis								
<b>I-CAMP</b> <sup>1</sup>	Japanese	148	10.7	6.0	<b>0.35</b> [0.23-0.52]	27.7	25.7	<b>0.78</b>
Subset analysis of the phase 3 trials for patients selected by clinical background								
<b>IPASS</b> <sup>2,3</sup>	East Asian	261	9.5	6.3	<b>0.48</b> [0.36-0.64]	21.6	21.9	<b>0.99</b>
<b>First-SIGNAL</b> <sup>4</sup>	Korean	42	8.4	6.7	<b>0.61</b> [0.31-1.22]	30.6	26.5	
Phase 3 trials for patients selected by EGFR mutation status								
<b>NEJ002</b> <sup>5,9</sup>	Japanese	194	10.8	5.4	<b>0.30</b> [0.22-0.41]	27.7	26.6	<b>0.48</b>
<b>WJTOG3405</b> <sup>6,10</sup>	Japanese	172	9.2	6.3	<b>0.49</b> [0.34-0.71]	36.0	39.0	<b>NS</b>
			Erlotinib Chemotherapy			HR Erlotinib Chemotherapy P		
Phase 3 trials for patients selected by EGFR mutation status								
<b>OPTIMAL</b> <sup>7,11</sup>	Chinese	154	13.1	4.6	<b>0.16</b> [0.10-0.26]	HR 1.065		<b>0.68</b>
<b>EURTAC</b> <sup>8</sup>	European	173	9.7	5.2	<b>0.37</b> [0.25-0.54]	19.3	19.5	<b>0.87</b>

<sup>1</sup>Morita S, et al. Clin Cancer Res. 2009;15:4493-8

<sup>2</sup>Mok TS, et al. N Engl J Med 2009;361:947-57

<sup>3</sup>Fukuoka M, et al. J Clin Oncol. 2011;29:2866-74

<sup>4</sup>Lee JS, et al. J Thorac Oncol. 2009;4(suppl):abstr PRS.4.

<sup>5</sup>Maemondo M, et al. N Engl J Med. 2010;362:2380-8

<sup>6</sup>Mitsudomi T, et al. Lancet Oncol. 2010;11:121-8

<sup>7</sup>Zhou C, et al. Lancet Oncol. 2011;12:735-42

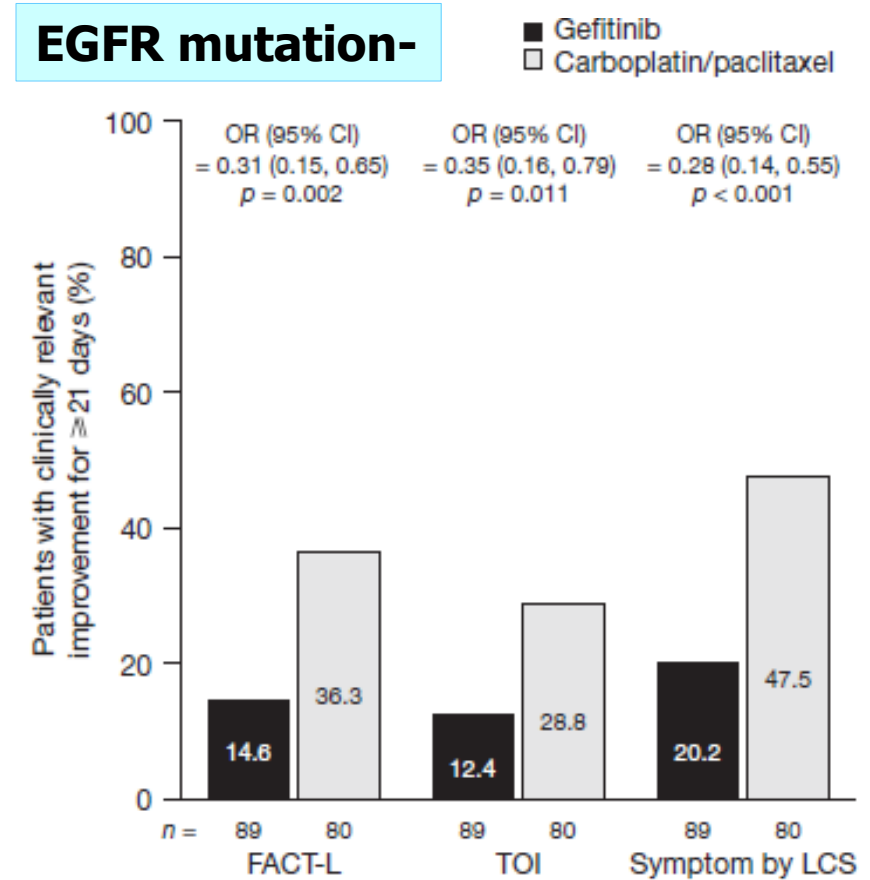
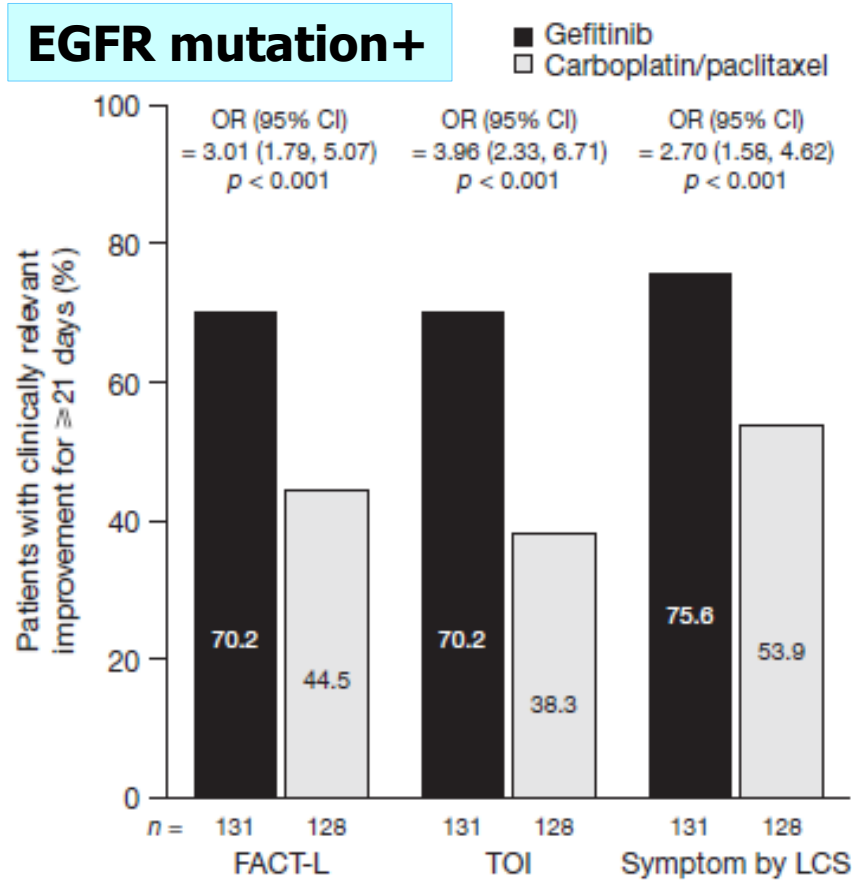
<sup>8</sup>Rosell R, et al. Lancet Oncol. 2012;13:239-46

<sup>9</sup>Inoue A, et al. Ann Oncol. 2013;24:54-59

<sup>10</sup>Mitsudomi T, et al. J Clin Oncol 30, 2012 (suppl; abstr 7521)

<sup>11</sup>Zhou C, et al. J Clin Oncol 30, 2012 (suppl; abstr 7520)

# Health-related quality of life in patients on IPASS



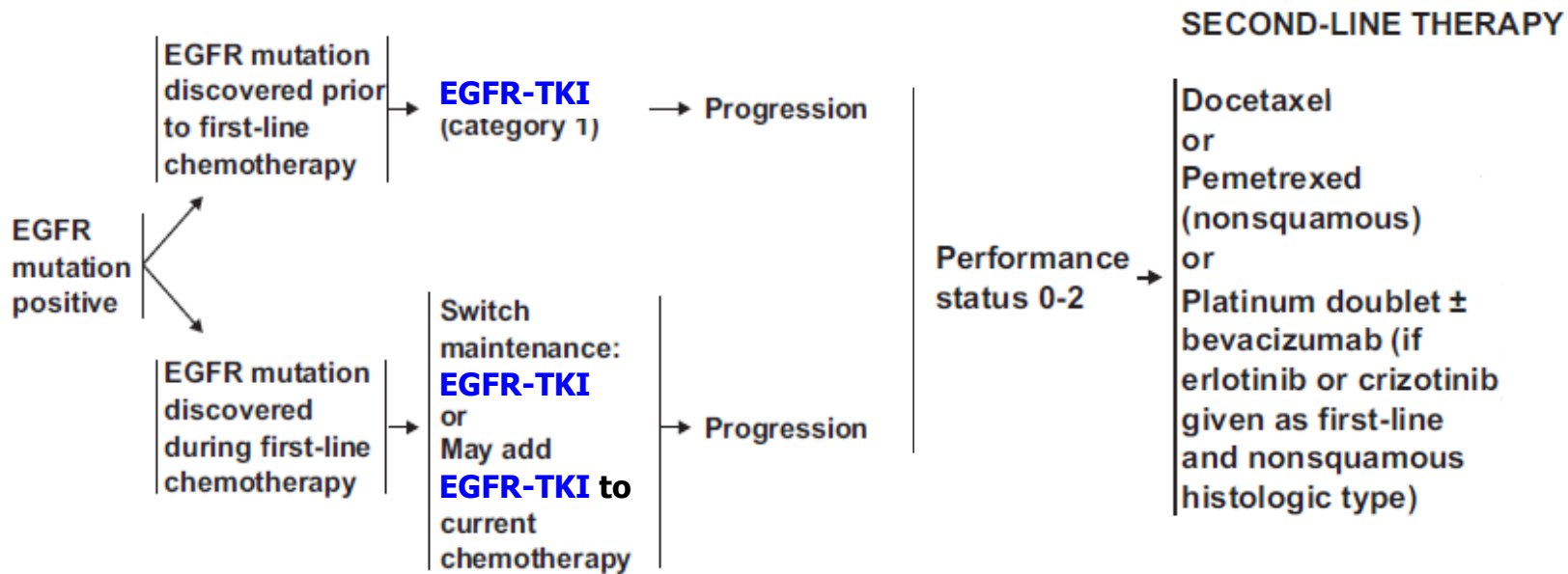
# Efficacy of switch maintenance to EGFR-TKI in NSCLC

Trial	Drug	Eligibility	N	Subgroup	PFS [Primary EP]	OS [Secondary EP]
<b>SATURN<sup>1</sup></b> □ Caucasian 84%	<b>Erlotinib</b>	① Stage IIIB-IV	889	Overall	HR <b>0.71</b> [0.62-0.82]	HR <b>0.81</b> [0.70-0.95]
		② Non-PD after 1 <sup>st</sup> -line CT, 4 cycles			2.87 vs. 2.59 mo	12.0 vs. 11.0 mo
		③ PS 0-1		EGFR <sup>mut+</sup>	HR <b>0.10</b> [0.04-0.25]	HR <b>0.83</b> [0.34-2.02]
				EGFR <sup>WT</sup>	HR <b>0.78</b> [0.63-0.96]	HR <b>0.77</b> [0.61-0.97]
<b>INFORM<sup>2</sup></b> □ Chinese 100%	<b>Gefitinib</b>	① Stage IIIB-IV	296	Overall	HR <b>0.42</b> [0.33-0.55]	HR <b>0.84</b> [0.62-1.14]
		② Non-PD after 1 <sup>st</sup> -line CT, 4 cycles			4.8 vs. 2.6 mo	18.7 vs. 16.9 mo
		③ PS 0-2		EGFR <sup>mut+</sup>	HR <b>0.17</b> [0.07-0.42]	
				EGFR <sup>WT</sup>	HR <b>0.86</b> [0.48-1.51]	

<sup>1</sup>Cappuzzo F, et al. Lancet Oncol. 2010;11:521-9

<sup>2</sup>Zhang L, et al. Lancet Oncol. 2012;13:466-475

# Treatment guideline for EGFR<sup>mut+</sup> NSCLC



# Clinical modes of EGFR-TKI failure and subsequent management in advanced NSCLC

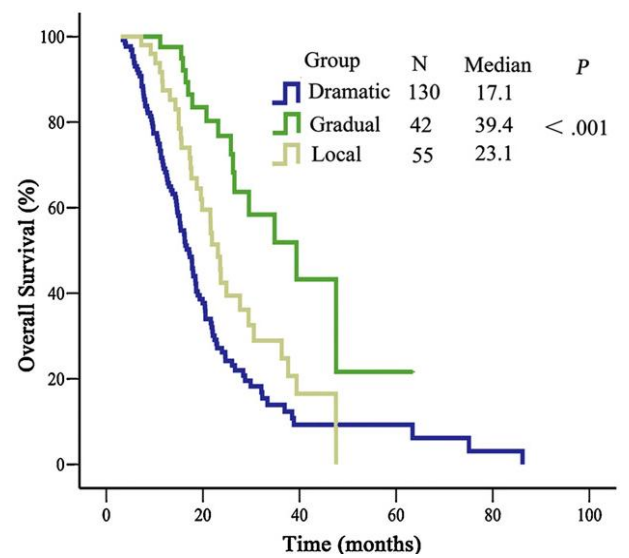
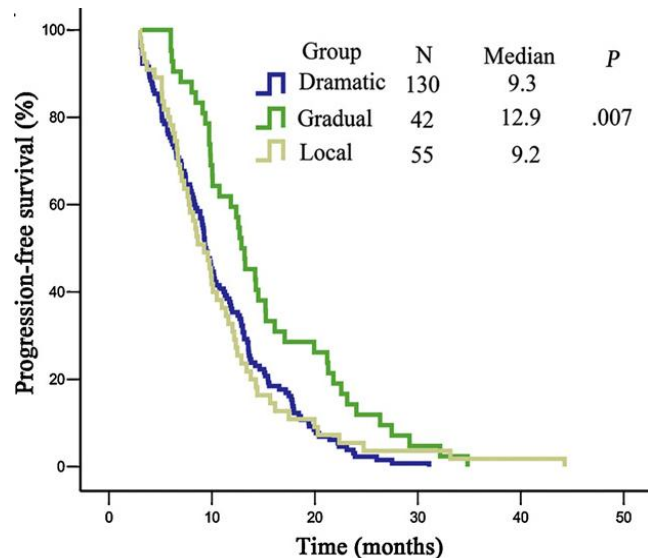
■ Stage III-IV NSCLC, First-line EGFR-TKI  
 ■ Duration of disease control  $\geq 3$  mo on EGFR-TKI

## EGFR-TKI failure

### Discriminating factors

**Disease control** on TKI  
 Tumor burden increment  
**[Doubling time]**  
**Symptom score**

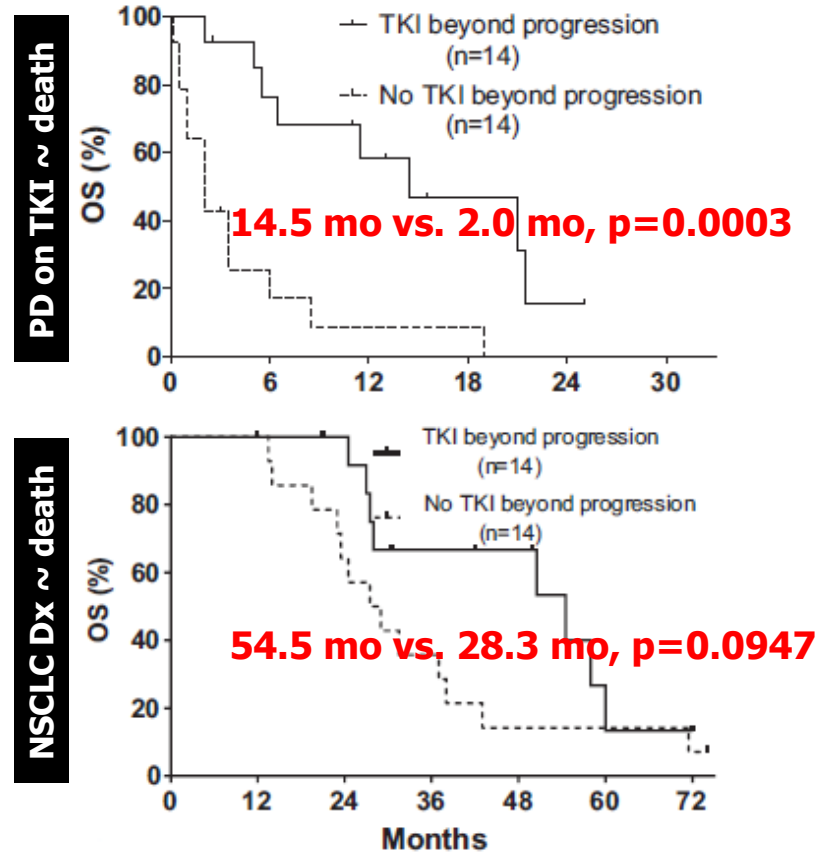
	Dramatic progression	Gradual progression	Local progression
Disease control on TKI	$\geq 3$ mo	$\geq 6$ mo	$\geq 3$ mo
Tumor burden increment [Doubling time]	Rapid	Minor	Solitary extra-cranial or limitation in intra-cranial lesion
Symptom score	$\geq 2$	$\leq 1$	$\leq 1$



# EGFR-TKI treatment beyond progression in long-term Caucasian responders to erlotinib in advanced NSCLC

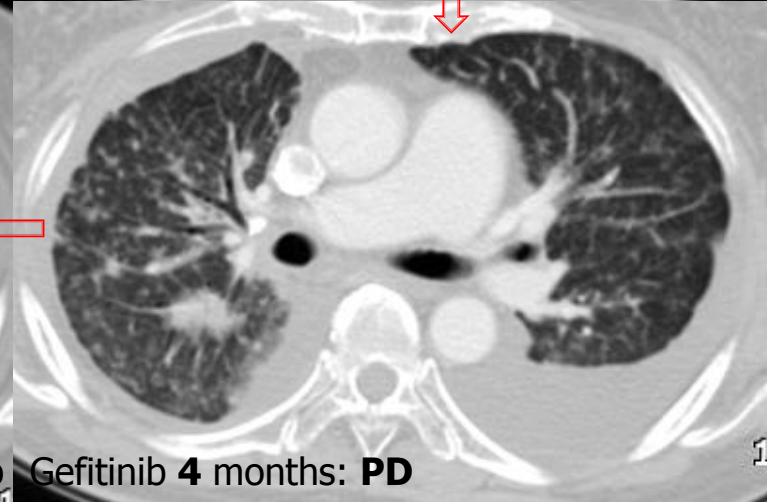
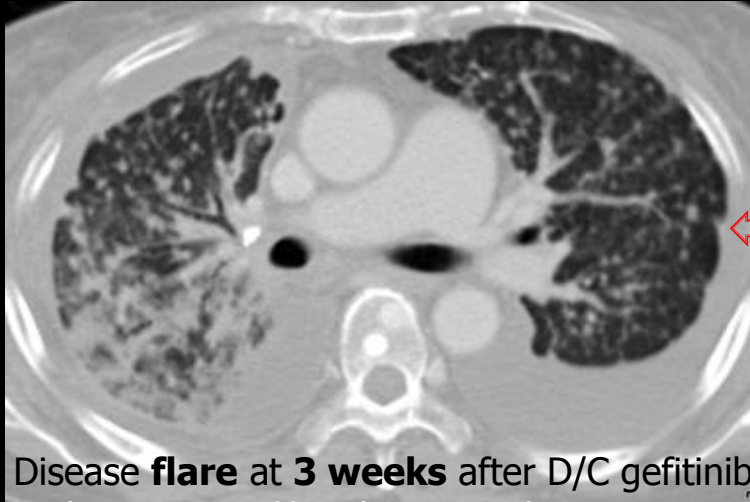
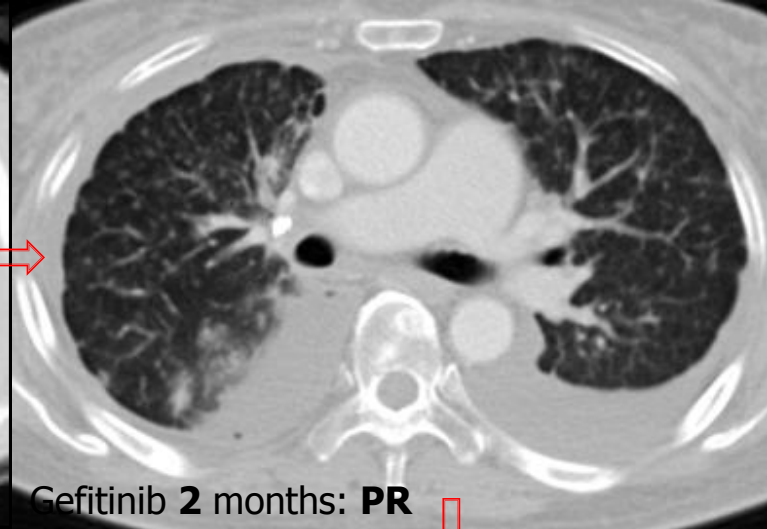
- A retrospective **case-control analysis** of **pairs matched** for sex, PS, smoking, histology
- Patients with stage IIIB-IV NSCLC, **at least SD on erlotinib  $\geq$  6 months**

	Continue TKI [n=14]	D/C TKI [N=14]
<b>EGFR mutation</b>		
(+) exon 19	4	1
(+) exon 21	0	1
(+) exon 18	1	0
Negative	6	3
Unknown	3	9
<b>Best response to erlotinib</b>		
CR	1	1
PR	10	9
SD	3	4
PFS	12.5 mo	12.0 mo
<b>Treatment after progression on TKI</b>		
Chemotherapy	11	9
Radiotherapy	4	3
BSC	4	5



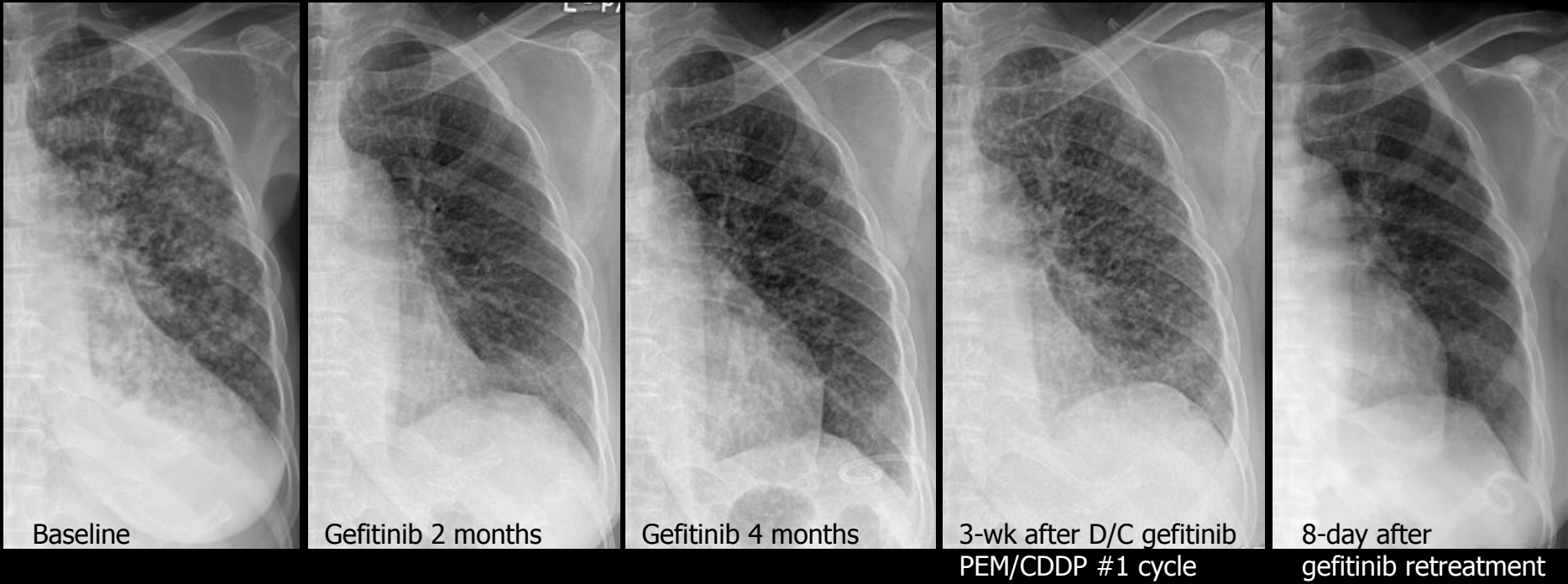
# A case of flare phenomenon after stopping EGFR-TKI

F/60, adenocarcinoma, EGFR exon19 del\_E746-A750



# A case of flare phenomenon after stopping EGFR-TKI

F/60, adenocarcinoma, EGFR exon19 del\_E746-A750

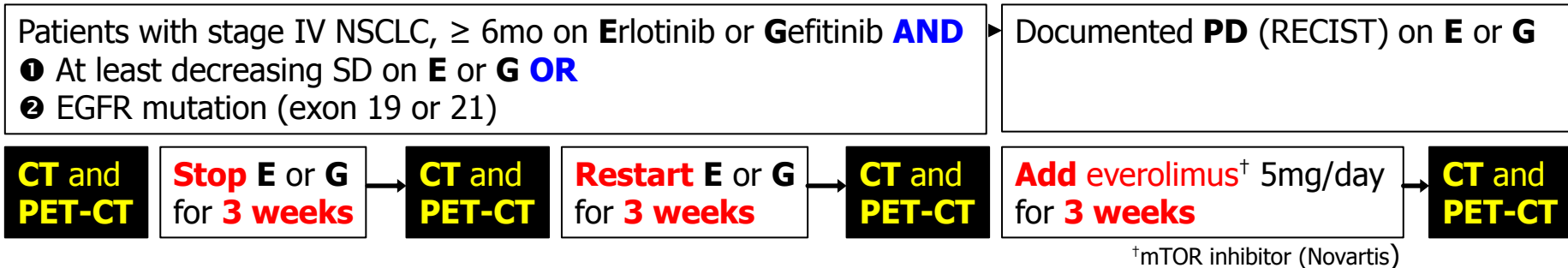


# Disease flare after TKI discontinuation in patients with EGFR-mutant lung cancer and acquired resistance to erlotinib or gefitinib

- A retrospective analysis (N=61)
- Patients with ❶ Lung cancer harboring a sensitizing mutation (G719X, exon 19 del, L858R)  
❷ Acquired EGFR-TKI resistance after initial benefit
- Definition of "**disease flare**": **hospitalization** or **death** attributable to disease progression after stopping TKI

- 14/61 (**23%**, 95% CI: 14-35) experienced a **disease flare**
- Median time to disease flare after TKI discontinuation was **8 days** (range 3-21)
- Factors associated with disease flare: ❶ **shorter time to progression on initial TKI** (p=0.002)  
❷ presence of **pleural involvement** (p=0.03)  
❸ presence of **CNS involvement** (p=0.01)
- No association with mutation type, presence of T790M, performance, sex, TKI, smoking status

# Prospective assessment of discontinuation and reinitiation of erlotinib(E) or gefitinib(G) in patients with acquired resistance to erlotinib or gefitinib followed by the addition of everolimus



**13** patients were enrolled, and **10 patients** completed the course

- Adenocarcinoma (11)/ NSCLC (2)
- Erlotinib (6)/ Gefitinib (7)
- Median time on erlotinib or gefitinib 16 months (8~79)
- EGFR L858R (2)/ exon 19 deletion (6)/ not available (5)

	After <b>stopping</b> E or G	After <b>restarting</b> E or G	After <b>adding</b> everolimus
Median $\Delta$ Tumor diameter	+9% (-13~+29%)	-1% (-14~+23%)	-8% (-34~+15%)
<b>Mean <math>\Delta</math> Tumor diameter</b>	<b>+9%</b>	<b>1%</b>	<b>-9%</b>
Median $\Delta$ Tumor volume	+50% (-4~+260%)	-1% (-27~15%)	-11% (-40~+26%)
<b>Mean <math>\Delta</math> Tumor volume</b>	<b>+61%</b>	<b>-4%</b>	<b>-10%</b>
Median $\Delta$ SUV <sub>max</sub>	+18% (-17~+87%)	-4% (-45~+62%)	-18% (-39~+82%)
<b>Mean <math>\Delta</math> SUV<sub>max</sub></b>	<b>+23%</b>	<b>-11%</b>	<b>-11%</b>
<b>Symptoms</b>	Generally aggravated	Generally stable or improved	

# Retreatment of gefitinib in patients with NSCLC who previously controlled to gefitinib

- A single arm, open-label, prospective phase II trial
- Advanced or metastatic NSCLC **previously controlled (CR/PR or SD  $\geq$ 3mo) by gefitinib**
- **At least one prior cytotoxic chemotherapy after a previous EGFR-TKI failure**
- Retreatment with gefitinib 250mg/day until PD or intolerant AE

## □ Baseline characteristics (n=23)

Clinical factor	Value	
<b>Sex</b>	Female	20 (86.9%)
	Male	3 (13.0%)
<b>Histology</b>	ADC	22 (95.7%)
	SQC	1(4.3%)
<b>Smoking</b>	Ex-smoker	2 (8.7%)
	Never	21 (91.3%)
<b>Response to initial gefitinib</b>	PR	10 (43.5%)
	SD	13 (56.5%)
<b>Gefitinib holiday</b>	$\leq$ 6mo	9 (39.1%)
	6-12mo	8 (34.8%)
	$\geq$ 12mo	6 (26.1%)

Initial diagnosis ~ gefitinib retreatment, **median 25.5 mo**

## □ Outcomes of gefitinib retreatment

Index	Outcome
<b>ORR</b>	21.7% (5/23)
<b>DCR</b>	65.2% (15/23)
<b>TTP</b>	103 day (95% CI, 70-134)
<b>OS</b>	343 day (95% CI, 235-409)

### Favorable factor for better TTP

DC on retreatment  
Good PS

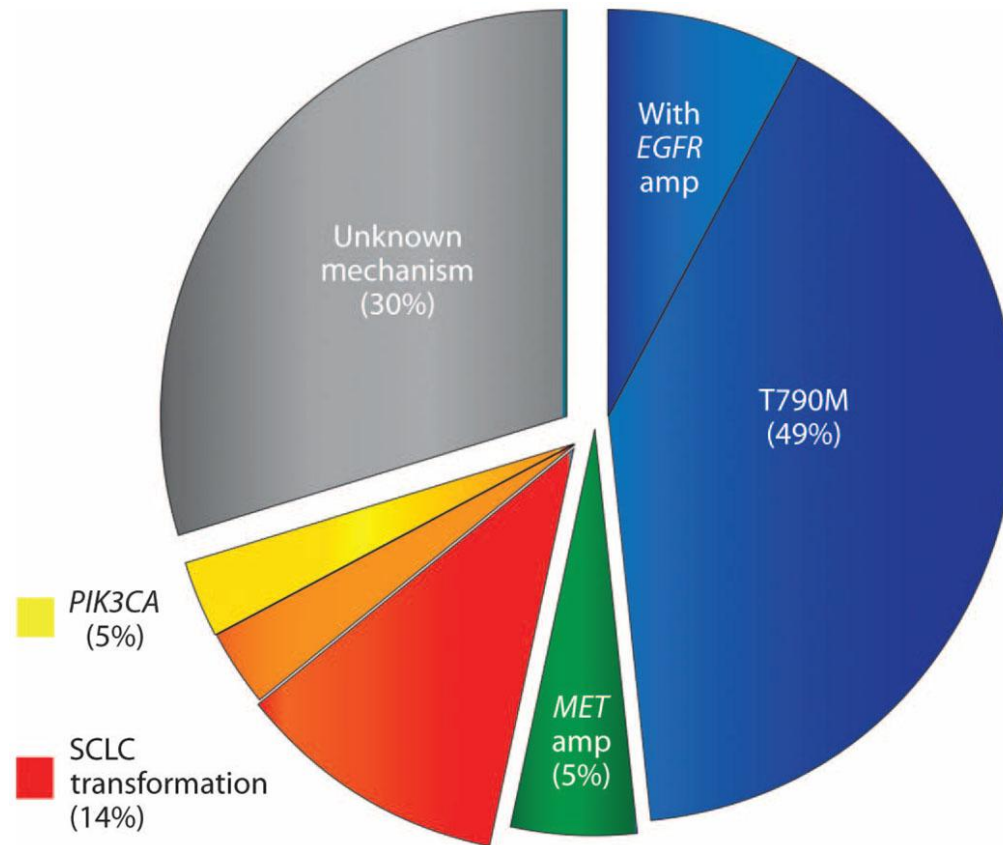
### PR on initial gefitinib

Skin rash

### Favorable factor for better OS

DC on retreatment  
Good PS

# Frequency of observed EGFR-TKI resistance mechanisms



**Biopsy on progression** to determine mechanism of acquired resistance, because proportion of patients will **transform to SCLC** at progression



PIK3CA (5%)  
SCLC transformation (14%)  
MET amp (5%)

# Molecular basis of drug resistance

- ① Drug efflux
- ② Acquisition of drug binding **target-deficient** mutants
- ③ Engagement of **alternative survival pathways**
- ④ Cancer **stem cells**
- ⑤ **Epigenetic mechanisms**

“One gene-one outcome” models may be an oversimplification of acquired resistance. Instead, clinical drug resistance may be due to changes in expression of **a large number of genes** that have a **cumulative impact on chemosensitivity**

Redmond KM, et al. Front Biosci. 2008;13:5138-54  
Trumpp A, et al. Nat Clin Pract Oncol. 2008;5:337-47  
Glasspool RM, et al. Br J Cancer. 2006;94:1087-92

# Retreatment with erlotinib: Regain of TKI sensitivity following a drug holiday for patients with NSCLC who initially responded to EGFR-TKI treatment

- A retrospective analysis
- Stage IV NSCLC with an **initial response** to EGFR-TKI (n=14: erlotinib 10, erlotinib+sorafenib 3, gefitinib 1)
- **Platinum based chemotherapy** after initial EGFR-TKI failure
- Retreatment with erlotinib at renewed progression

## □ Baseline characteristics (n=14)

Clinical factor		Value
<b>Sex</b>	Female	9 (64%)
	Male	5 (36%)
<b>Initial EGFR</b>	Exon 19 del	11 (79%)
	L858R	1 (7%)
	Unknown	2 (14%)
<b>Rebiopsy EGFR</b>	Exon 19 del	6 (43%)
	Exon 19 del + <b>T790M</b>	<b>5 (36%)</b>
	L858R	1 (7%)
	Unknown	2 (14%)
<b>Response duration to initial EGFR-TKI</b>		
		12.5 mo (95% CI,7-34)
<b>TKI holiday</b>		9.5 mo (range, 3-36)

## □ Outcomes of erlotinib retreatment<sup>†</sup>

Index	Outcome
<b>ORR</b>	36% (5/14)
<b>DCR</b>	86% (12/14)
<b>PFS</b>	6.5 mo (range, 1-16+)

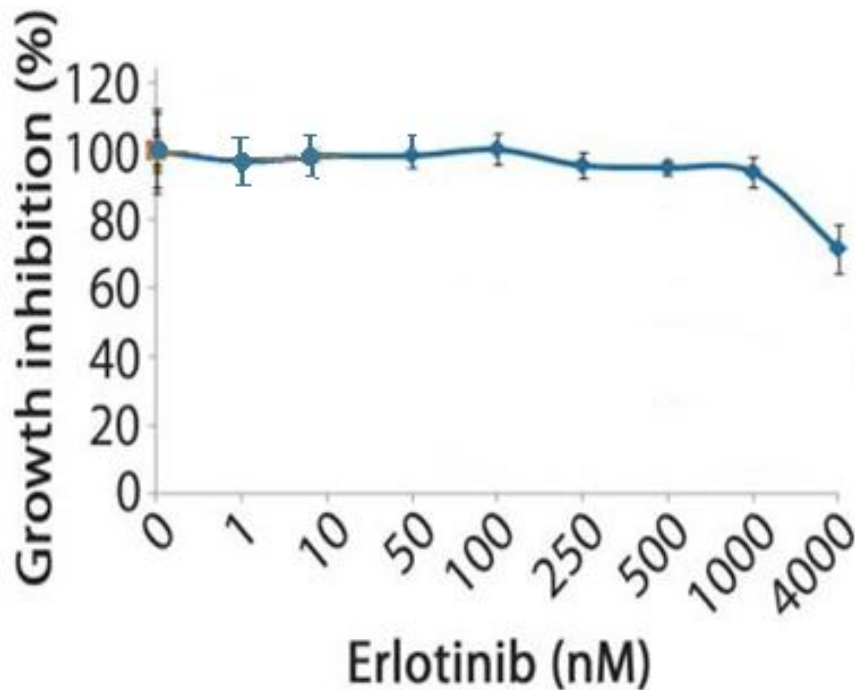
<sup>†</sup>3 patients received erlotinib + cetuximab  
2/3 showed PD, 1/3 PR

## □ Response in patients with **EGFR<sup>ex19 del</sup>+T790M**

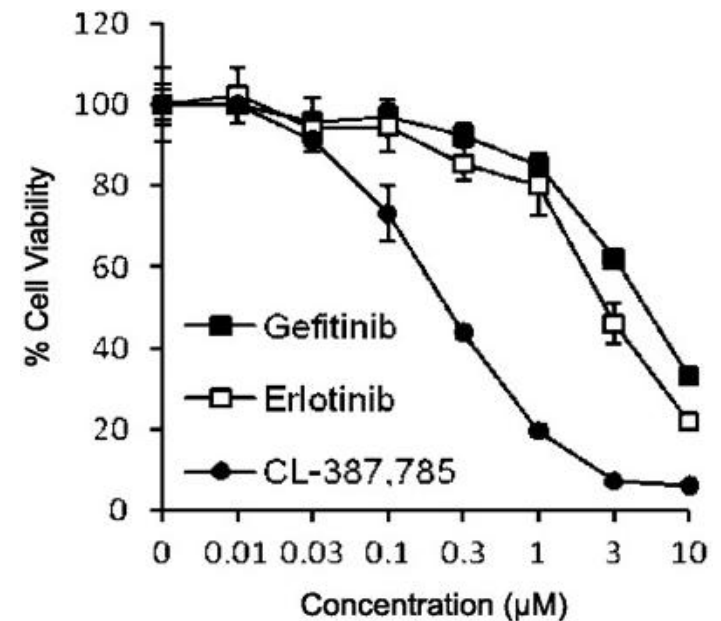
<b>PR</b>	<b>40% (2/5)</b>
SD	20% (1/5)
PD	40% (2/5)

# Cancer cells harboring dual EGFR mutation (sensitizing+T790M) have resistance to EGFR-TKI

PC-9 (EGFR exon19 del/T790M)



H1975 (EGFR L858R/T790M)

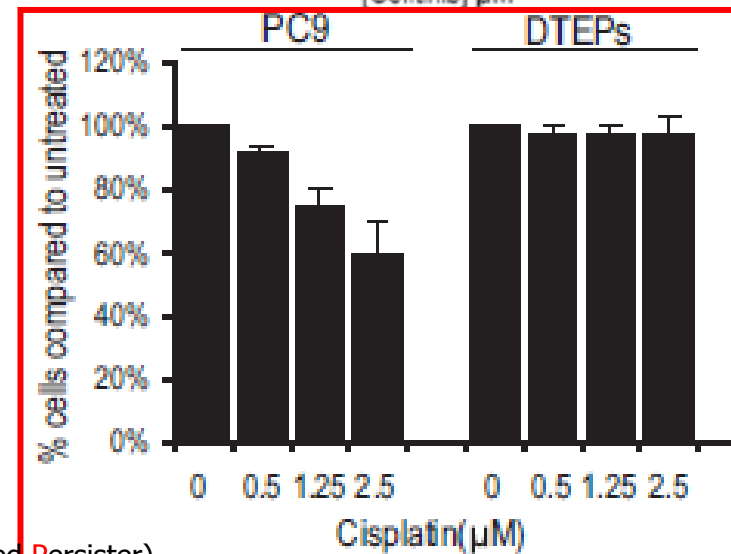
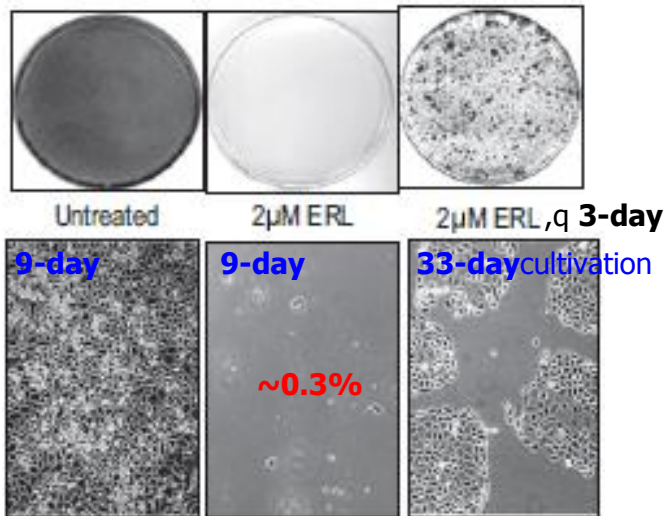
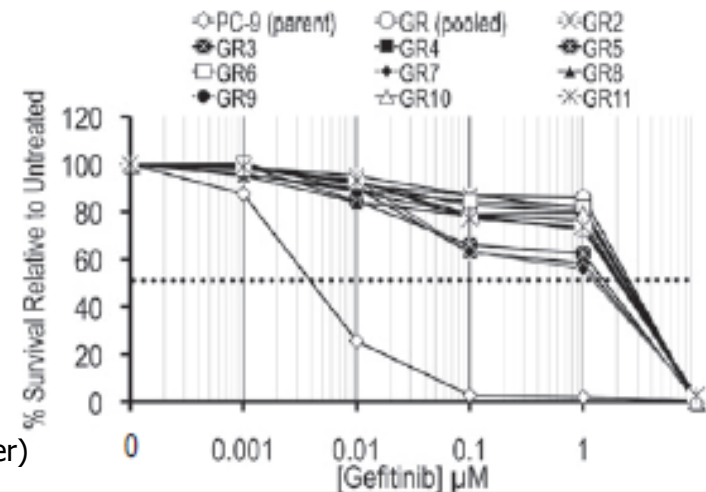
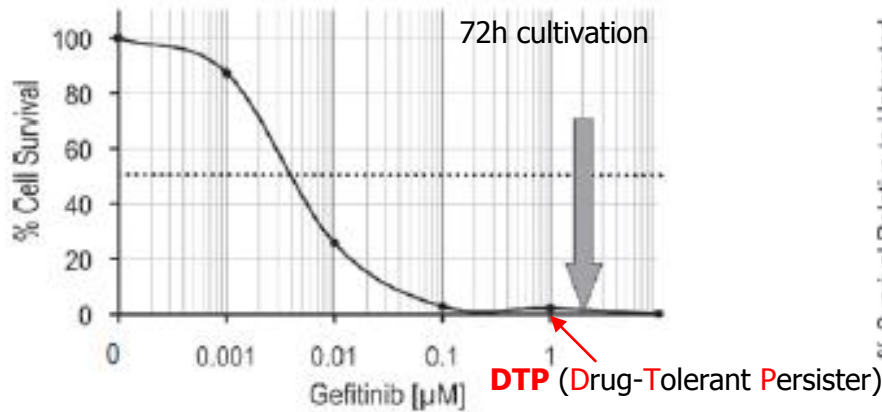


Chmielecki J, et al. *Sci Transl Med*. 2011;3:90ra59

Yamada T, et al. *Clin Cancer Res*. 2010;16:174-83

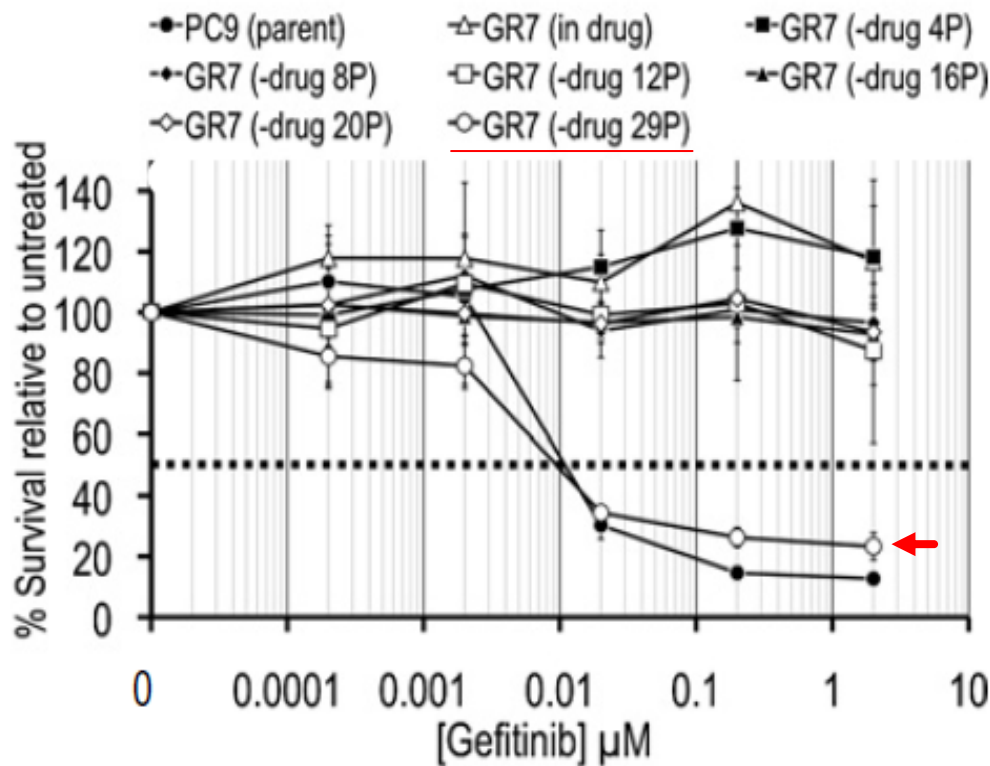
# A chromatin-mediated reversible drug-tolerant state in cancer cell subpopulations

## PC-9 (harboring EGFR exon19 deletion)



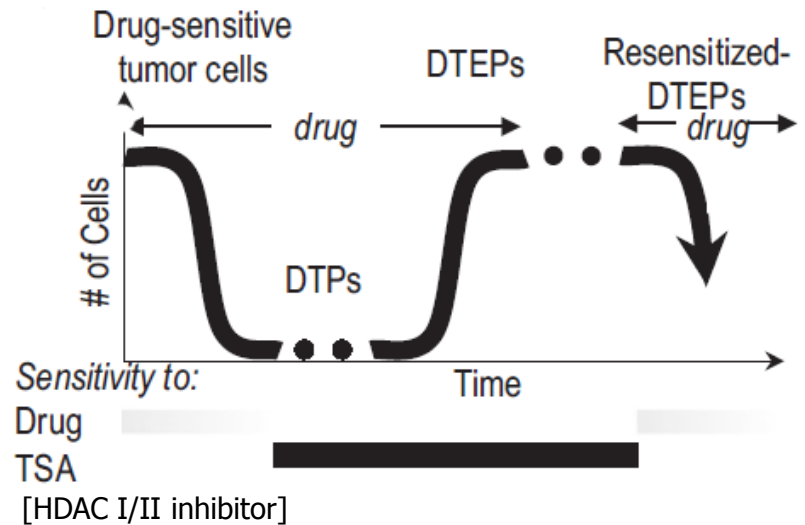
# A chromatin-mediated reversible drug-tolerant state in cancer cell subpopulations

Indicated number passage in drug free media



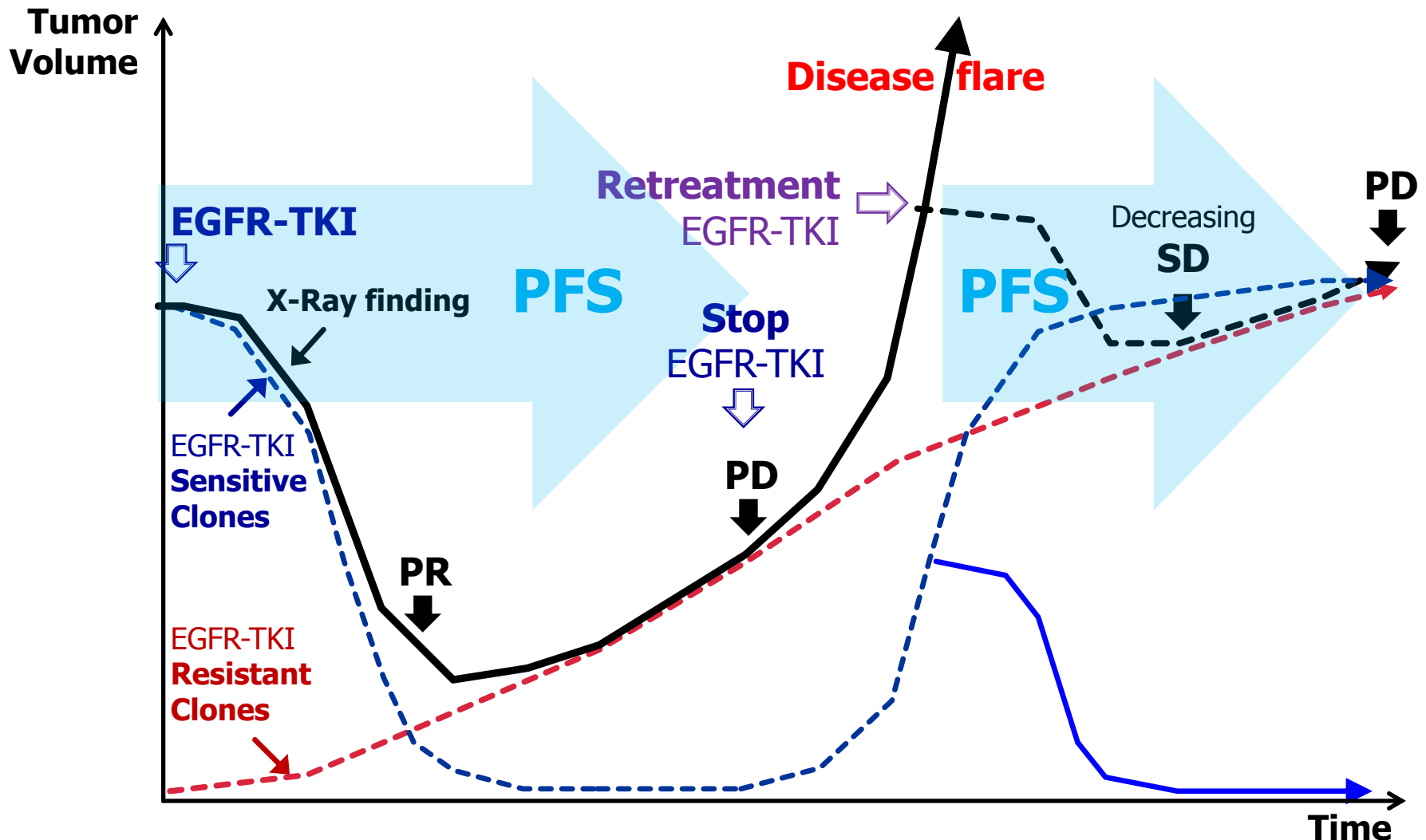
Involved mechanisms establishing DTEPs

- ① Stem cell marker+ (CD133)
- ② Histone demethylation, deacetylation
- ③ IGF-1R signaling
- ④ EGFR T790M in several weeks later(?)



Dynamic phenotypic heterogeneity of a nongenetic nature under a lethal drug exposure

# Real practice for EGFR-mutant NSCLC



# EGFR-TKI in the treatment of EGFR mutant NSCLC metastatic to the brain

- Incidence of brain metastases (BM) in patients with NSCLC: approximately **25~30%**<sup>1</sup>
- Different incidence of BM: EGFR **mutant: 64%** (9/14) vs. **wild type: 31%** (30/96)<sup>2</sup>
- Local therapy: Surgery/SRS/WBRT
- Systemic therapy: **EGFR-TKI** > chemotherapy in NSCLC **with EGFR-mutation**

	<b>N</b>	<b>Drug</b>	<b>ORR of Brain metastasis</b>
Porta R, et al. <sup>3</sup>	17	Erlotinib	<b>82.4%</b> (14/17)
Park SJ, et al. <sup>4</sup>	28	Erlotinib or Gefitinib	<b>82.1%</b> (23/28)
Li Z, et al. <sup>2</sup>	9	Gefitinib	<b>88.9%</b> (8/9)

## ■ **Dose escalation** for **refractory** brain metastases to standard-dose EGFR-TKI

- ① Switching Erlotinib > Gefitinib in ratio of CSF/Serum EFGR-TKI<sup>5</sup>
- ② Gefitinib 500mg ▶ 750mg ▶ 1,000mg ▶ 1,250mg<sup>6</sup>
- ③ Erlotinib Alternating 300mg/d ◀▶ 150mg/d<sup>7</sup>
- ④ Erlotinib Weekly 1,000~1,500mg pulse<sup>8</sup>

<sup>1</sup>Langer CJ, et al. J Clin Oncol. 2005;23:6207-19

<sup>2</sup>Li Z, et al. J Clin Oncol. 2011;29:e18065

<sup>3</sup>Porta R, et al. Eur Respir J. 2011;37:624-31

<sup>4</sup>Park SJ, et al. Lung Cancer. 2012;77:556-60

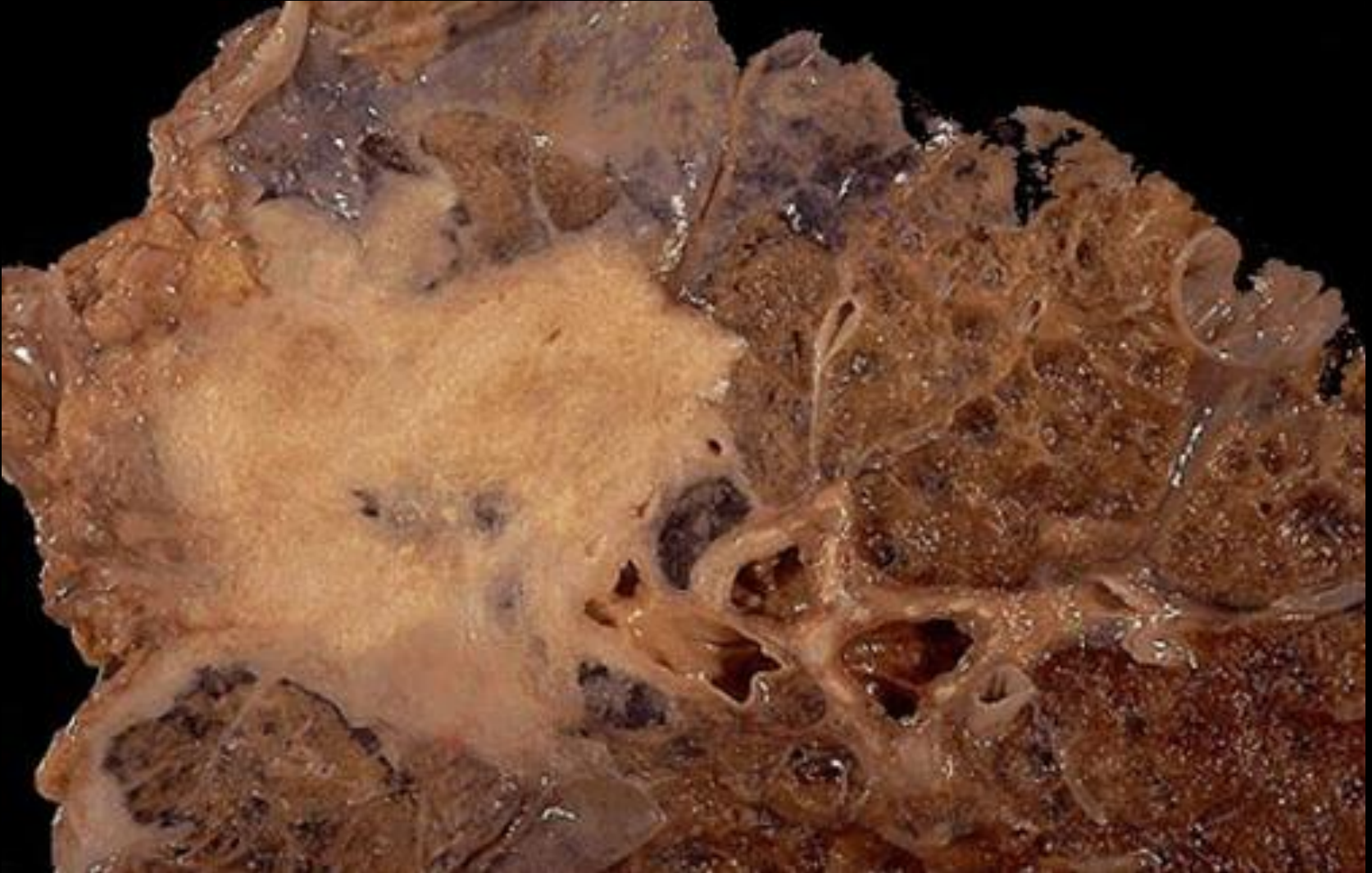
<sup>5</sup>Katayama T, et al. J Thorac Oncol. 2009;4:1415-9

<sup>6</sup>Jackman DM, et al. J Clin Oncol. 2006;24:4517-20

<sup>7</sup>Hata A, et al. J Thorac Oncol. 2011;6:653-4

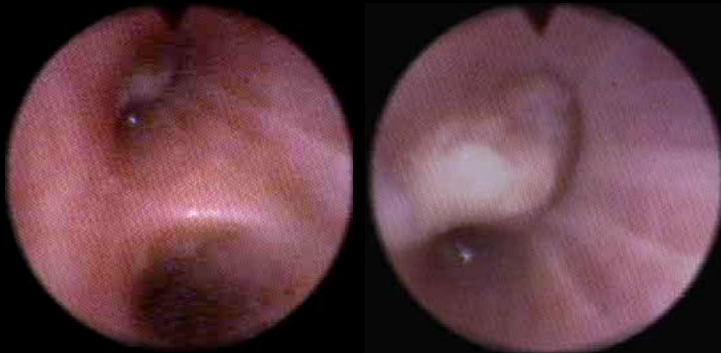
<sup>8</sup>Clarke JL, et al. J Neurooncol. 2010;99:283-6

# Intratumoral heterogeneity in NSCLC, probably be!



# A case of intratumoral heterogeneity in NSCLC

Lung percutaneous needle biopsy:  
Adenocarcinoma, well differentiated  
EGFR: positive, TTF-1: positive, ALK: negative



Bronchoscopic biopsy: Squamous cell carcinoma  
P63: positive, TTF-1: negative

**PNA Clamping for EGFR**  
**Deletion mutation in exon 19**  
No mutation in exon 18, 20, 21



F/55, never smoker  
cough and pain



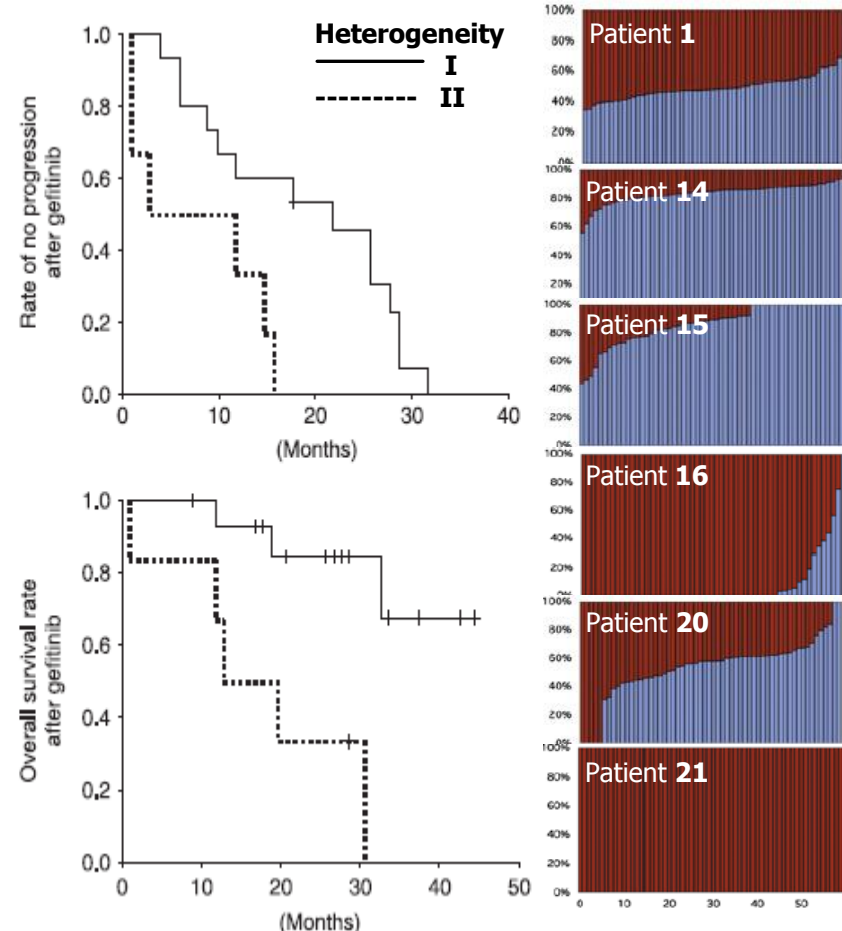
After gefitinib 250mg/d  
for 2 months

**Squamous cell carcinoma, molecular testing recommendation:**  
EGFR mutation and ALK testing are not routinely recommended  
**except in never smokers and small biopsy specimens**

# Intratumor heterogeneity of EGFR mutations in lung cancer and its correlation to the response to gefitinib

- 21 patients **with EGFR<sup>mut+</sup>** NSCLC treated with gefitinib for **postoperative recurrence**
- Multipoint microsampling (laser capture microdissection: 50-60 area, 30-50 cells/area) and SNaP shot assay

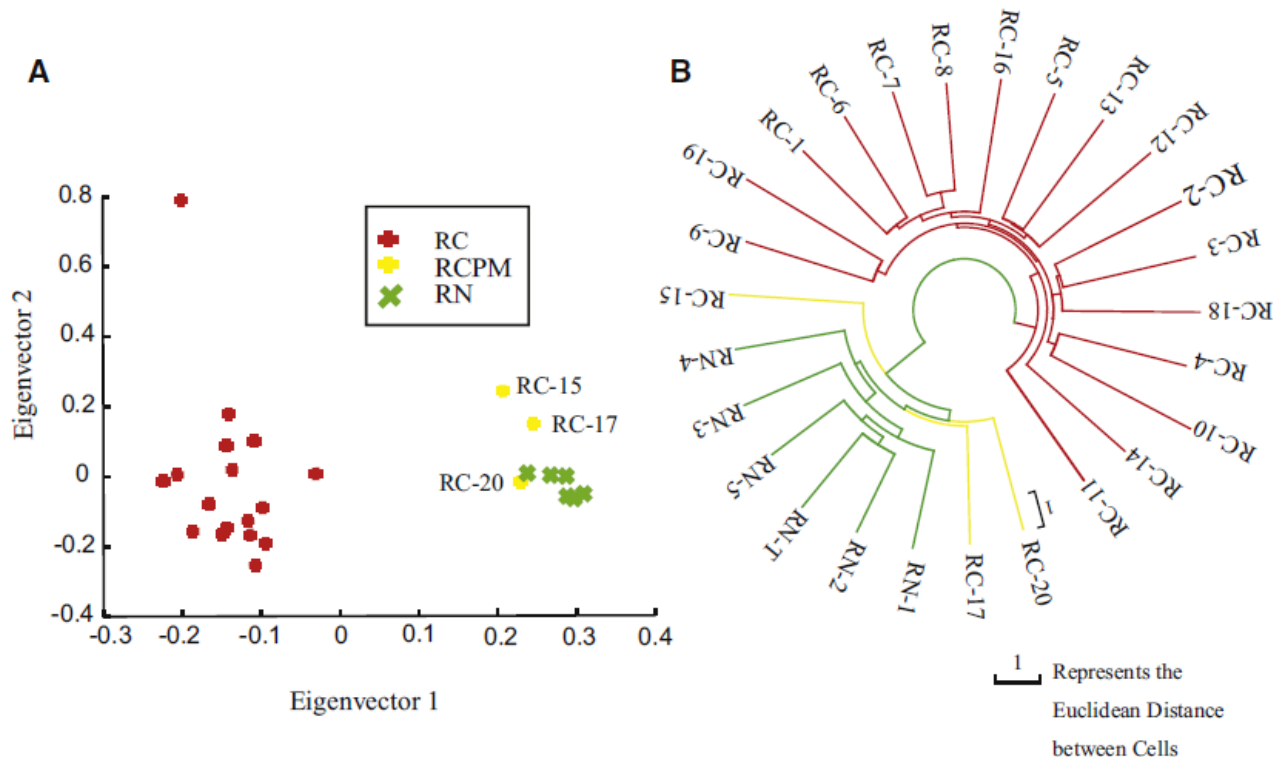
No.	Histo	EGFR mutation	Wild type area(%)	Mutated allele(%)	Heterogeneity	Gefitinib response
1	Ad	L858R	0.00	47.9	I	PR
2	Ad	L858R	0.00	48.0	I	PR
3	Ad	L858R	0.00	50.4	I	CR
4	Ad	L858R	0.00	53.2	I	CR
5	Ad	L858R	0.00	53.9	I	CR
6	Ad	delE746-A750	0.00	58.6	I	CR
7	Ad	delE746-A750	0.00	58.7	I	PR
8	Ad	delE746-A750	0.00	61.5	I	PR
9	Ad	L858R	0.00	74.2	I	PR
10	Ad	delE746-A750	0.00	75.1	I	PR
11	Ad	delE746-A750	0.00	81.2	I	PR
12	Ad	delE746-A750	0.00	82.8	I	PR
13	Ad	L861Q, G719S	0.00	82.9	I	PR
14	Ad	delL747-T751	0.00	84.5	I	PR
15	Ad	delE750-K758	0.00	93.9	I	PR
16	Ad	L858R	74.07	24.1	II	SD
17	Ad	delE746-A750	6.67	48.3	II	PR
18	Ad	delE746-A750	11.54	49.8	II	CR
19	Ad	delE746-T751 insIP	8.77	54.3	II	PR
20	Ad	delE746-A750	8.62	58.1	II	PR
21	AdSq	delA859-L883 insV	100.00	0.0	II	PD





# Single-cell exome sequencing reveals single-nucleotide mutation characteristics of a kidney tumor

- A 59-year-old Chinese male with stage IV **clear cell renal cell carcinoma (ccRCC)**
- Isolation of single cells (20 cancer cells: RC1~20 and 5 normal cells: RN1~5) under inverted microscope
- Principle component analysis (PCA) mutation profiling on the 260 somatic mutation sites

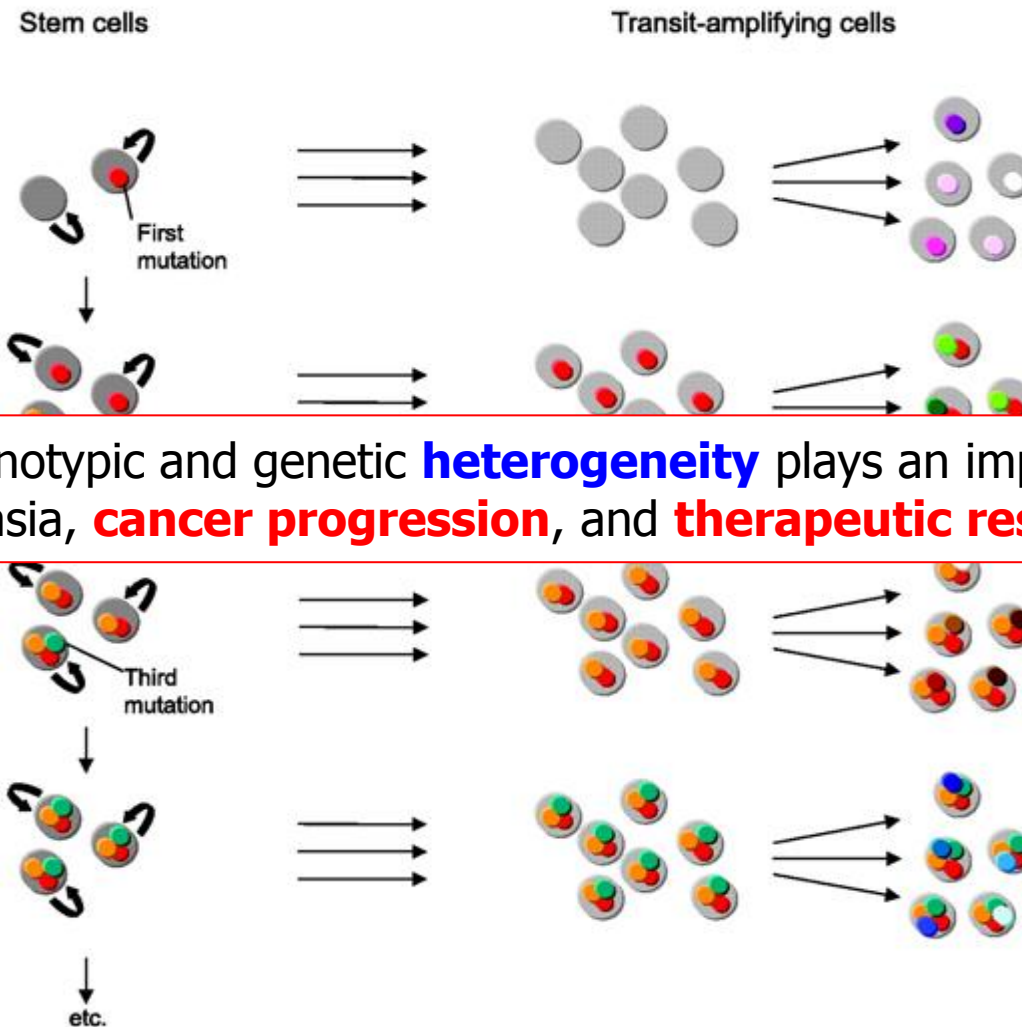


(A) PCA of cancer cell (RC), normal control cell (RN), and normal cells picked as cancer cells (RCPM)

(B) Neighbor joining phylogenetic tree constructed using sites of somatic mutation data by Euclidean distance  
RN-T (normal tissue DNA) as control

# Emergence and maintenance of tumor diversity

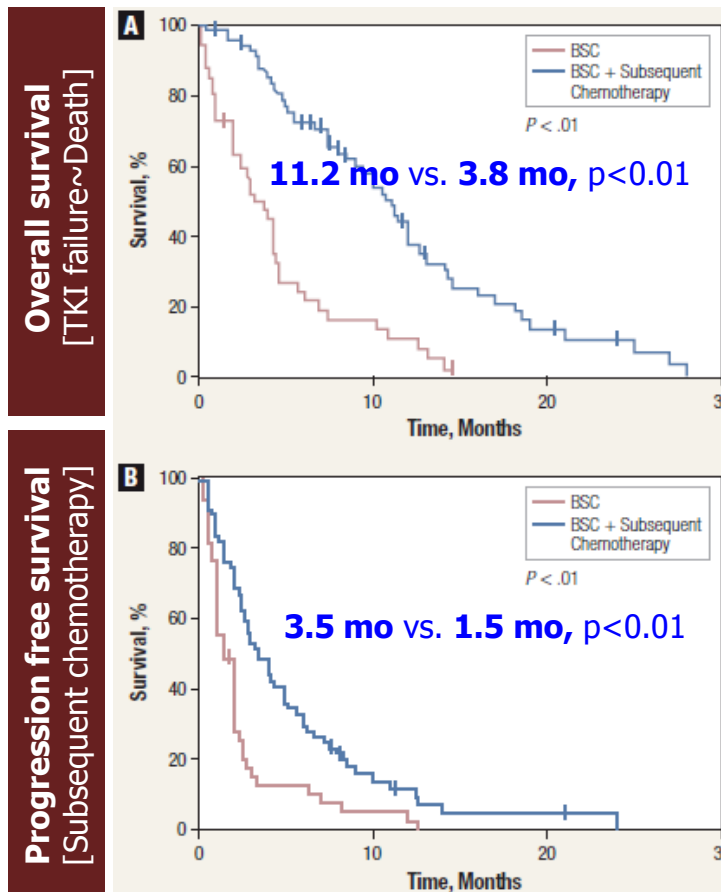
## Intratumor heterogeneity



This phenotypic and genetic **heterogeneity** plays an important role in neoplasia, **cancer progression**, and **therapeutic resistance**!

# Subsequent chemotherapy improves survival outcome in advanced NSCLC with acquired TKI resistance

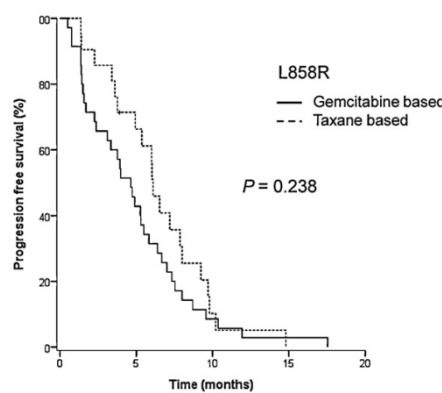
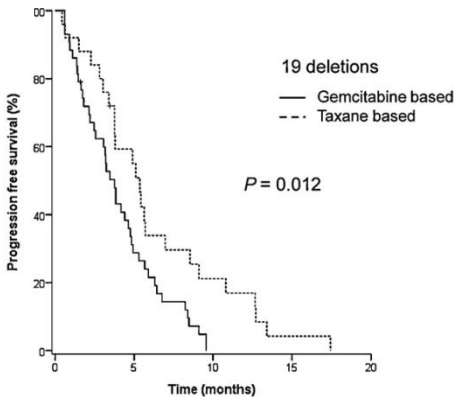
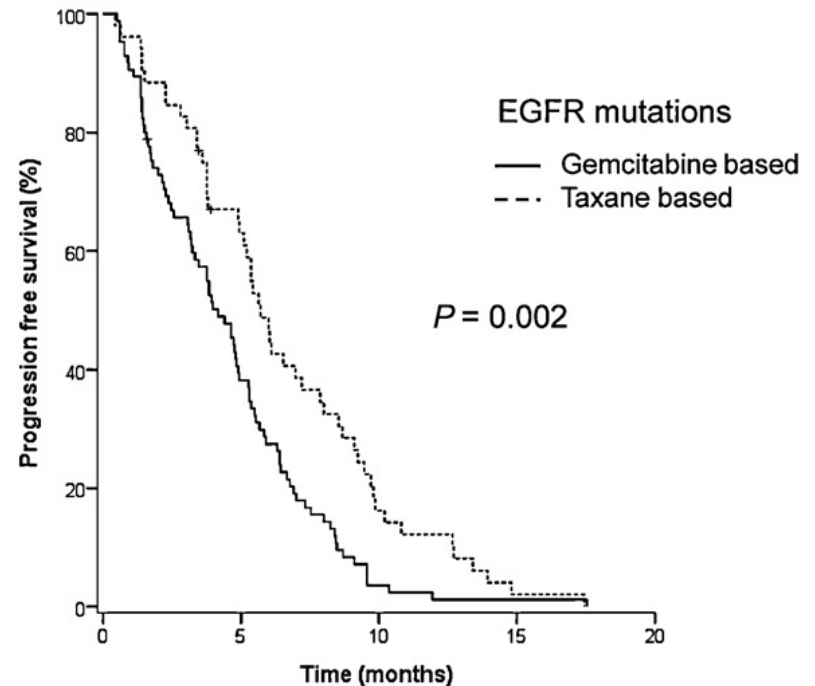
- A retrospective analysis in Taiwan
- 114 patients with stage IIIB-IV NSCLC, disease control on EGFR-TKI



# EGFR mutations as a predictive marker of cytotoxic chemotherapy

- A retrospective analysis, **1<sup>st</sup>-line chemotherapy**, Year 2005-2010, in Korea
- Advanced/metastatic NSCLC (n=217: ADC 174, SQC 4, Others 39)
- **Gemcitabine-based vs. Taxane-based regimen: ORR and PFS** according to **EGFR mutation status**

		GEM	TAX	P-value
<b>Wild type EGFR</b>	N	46	34	
	<b>ORR</b>	34.8%	35.3%	0.962
	<b>DCR</b>	78.3%	73.5%	0.623
	<b>PFS</b>	4.4mo	4.4mo	0.362
<b>Mutant EGFR</b>	N	85	52	
	<b>ORR</b>	30.6%	38.5%	0.344
	<b>DCR</b>	<b>71.8%</b>	<b>88.5%</b>	<b>0.022</b>
	<b>PFS</b>	<b>4.1mo</b>	<b>5.7mo</b>	<b>0.002</b>



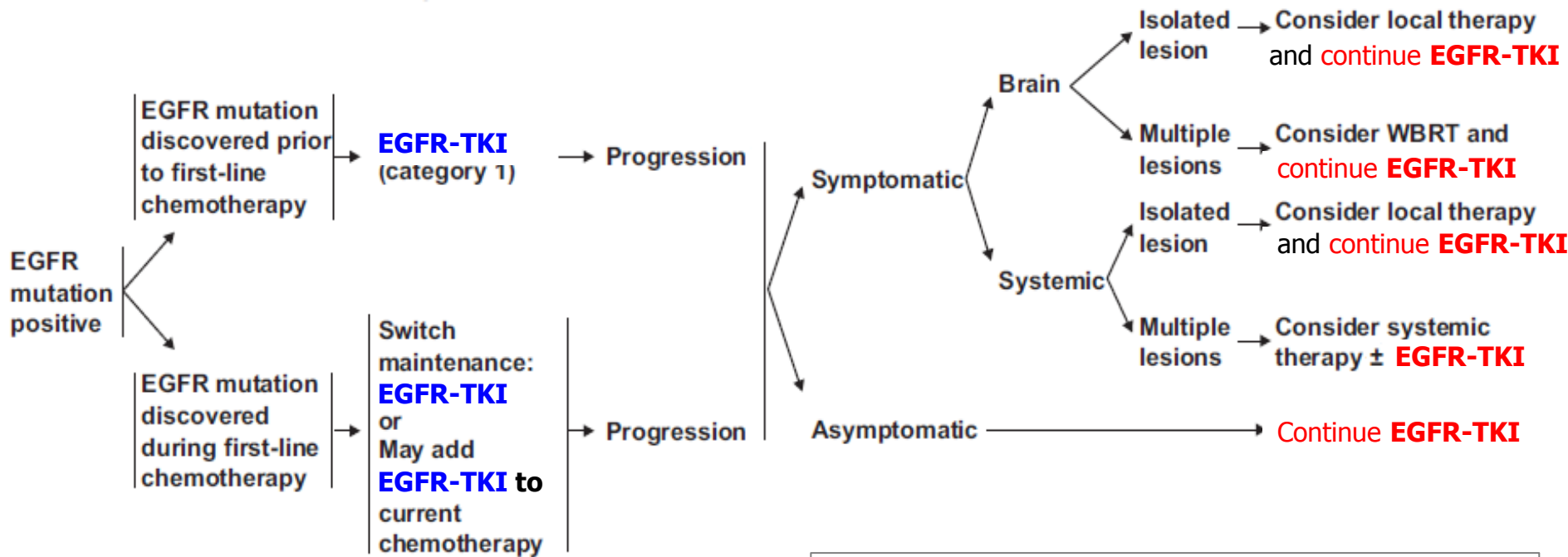
# Retreatment of patients with same chemotherapy: implications for clinical mechanism of drug resistance

- Advanced cancer patients **retreated with same chemotherapy** previously received
- Retreatment** for the progressive disease **after complete remission**
- Medline review of 15 reports
- Median off-treatment 48 weeks, **Response rate 18-100% (median 51%)**

Cancer type	Drug	Number	Holiday	Response	Duration
AML	Cytosine, ADR	24	31 wk	37% CR	13 wk
AML	Cytosine, daunorubicin	136	46 wk	51% CR	20 wk
Harry cell leukemia	Chlorodeoxyadenosine	4	76 wk	50% CR, 25% PR	72 wk
CML	Chrorodeoxyadenosine	6	58 wk	16% CR, 32% PR	-
Multiple myeloma	Melphalan, procarbazine, PD, VCR	20	>48 wk	80% PR	-
HD					84 wk
Breast cancer					12 wk
SCLC					-
SCLC	Cytosine, ADR, etoposide	37	34 wk	16% CR, 45% PR	26 wk
SCLC	Cytosine, ADR, etoposide or VCR	12	30 wk	16% CR, 33% PR	19 wk
SCLC	Carboplatin, epirubicin, etoposide	6	58 wk	33% CR, 67% PR	-
Ovarian cancer	Platinum, paclitaxel	10	36 wk	30% CR+PR	10 wk
Ovarian cancer	Cytosine, ADR, cisplatin, methylmelamine	11	88 wk	36% CR, 36% PR	38 wk
Colorectal cancer	5-FU, leucovorin	49	50 wk	18% CR+PR	21 wk
Urothelial cancer	MTX, VBL, ADR, cisplatin	8	52 wk	71% PR	-

**Transient resistance to chemotherapy!**

# Summary



**Flare phenomenon or after EGFR-TKI holiday, consider retreatment with EGFR-TKI!**

**Chemotherapy! Singlet vs. platinum doublet? Taxane seems to be more efficacious than non-taxane: if so, taxane vs. pemetrexed?**



**Thank you for your attention...**