

# The Changing Landscape of Lung Cancer and its Treatment

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# Lung Cancer 2011

- **USA – 190,000 new cases**
  - 165,000 deaths
  - 165,300 cases of NSCLC
  - 115,000 cases of adenocarcinoma
  - 28,500 cases of lung cancer in never smokers
- **Global**
  - 550,000 deaths from lung cancer

# LUNG CANCER

## Histological Types

- ❖ **Non-small cell lung cancer (85%)**
  - ❖ **Adenocarcinoma**
  - ❖ **Squamous cell carcinoma**
  - ❖ **Large cell carcinoma**
- ❖ **Small cell lung cancer (15%)**

# LUNG CANCER

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  - ❖ Adenocarcinoma
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# Median survivals in SCLC

- Very-limited disease ~5 years
- Limited disease 18-24 months
- Extensive disease 10 months
  
- SCLC without treatment <3 months

# Major advances in treatment of SCLC in the last 2 decades

- Combined modality concomitant chemo-radiotherapy in limited disease
- Prophylactic cranial irradiation in limited and extensive disease
- Standard systemic therapy not changed in the last two decades: platinum plus etoposide
- No new systemic treatment approved after the registration of topotecan for second-line treatment (FDA 1998; EMEA 2006)

# Teniposide in the Treatment of Small-Cell Lung Cancer: The Influence of Prior Chemotherapy

By Giuseppe Giaccone, Michela Donadio, Gianmaria Bonardi, Franco Testore, and Alessandro Calciati

Fifty patients with small-cell lung cancer (SCLC) were treated with teniposide (VM26) at 120 to 140 mg/m<sup>2</sup> on days 1, 3, and 5, every 3 weeks. Twelve elderly patients were administered VM26 as first-line chemotherapy. Toxicity was manageable, myelosuppression being the major side effect. The response rate for 44 evaluable patients was 34% (36% for untreated patients); the median durations of response and survival were 230 and 208 days, respectively. Effectiveness of prior chemotherapy and time from last administration was found to influence patient response to VM26: 42% of responders to prior chemotherapy responded to VM26, while 0% of the nonresponders to prior che-

motherapy responded to the new agent. Moreover, among patients pretreated with chemotherapy, 12% of those recently treated (earlier chemotherapy ending  $\leq 2.6$  months before administration of VM26) responded to VM26, while 53% of patients treated  $> 2.6$  months earlier responded to VM26. Survival was influenced by common prognostic factors (performance status, weight loss, prior chemotherapy exposure). Selection of pretreated patients by type of exposure to prior chemotherapy may help in the testing of new drugs in this disease. *J Clin Oncol* 6:1264-1270. © 1988 by American Society of Clinical Oncology.

**Time from prior  
Chemotherapy**

**$\leq 2.6$  m**

**$> 2.6$  m**

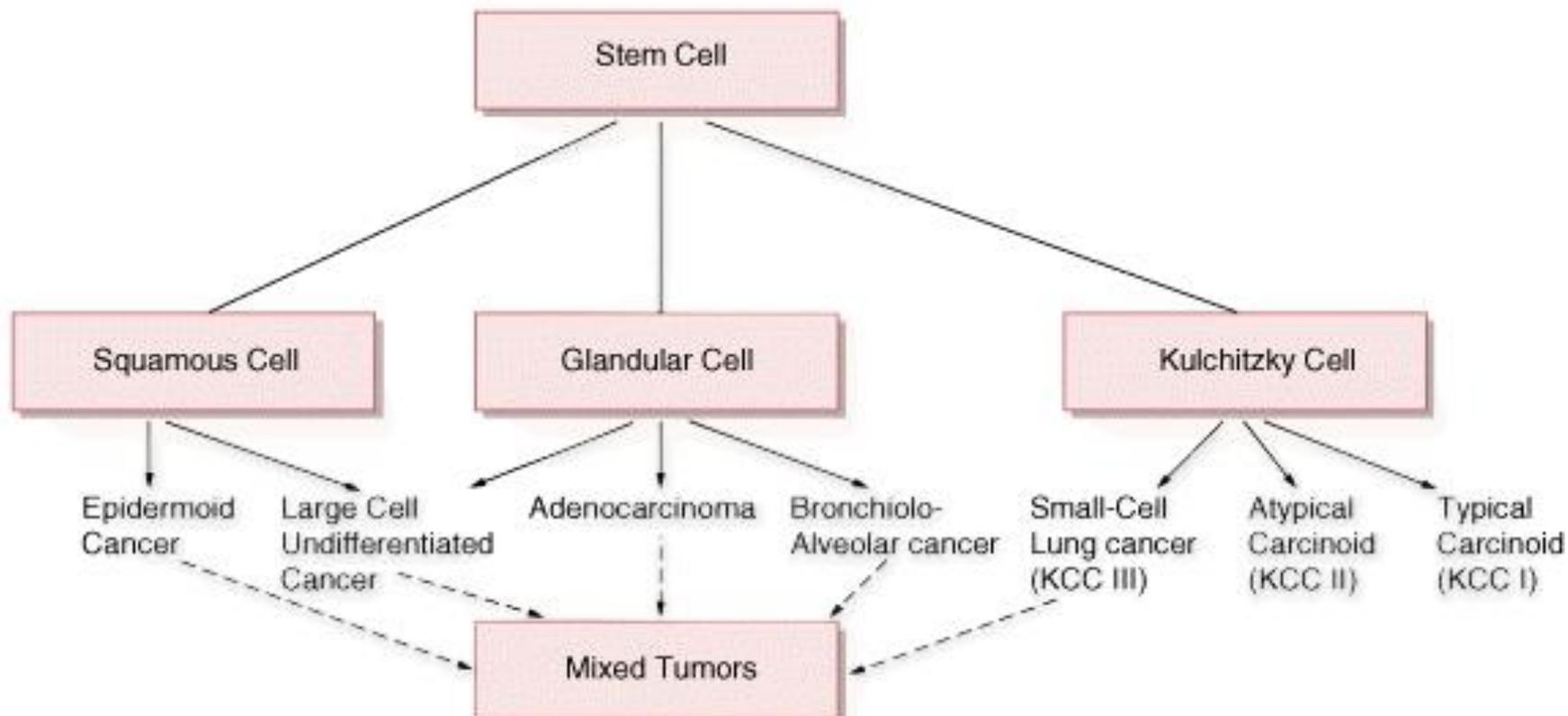
**Response to  
teniposide**

**12%**

**53%**

**(p=0.016)**

# Hypothesized model of origin of lung tumors

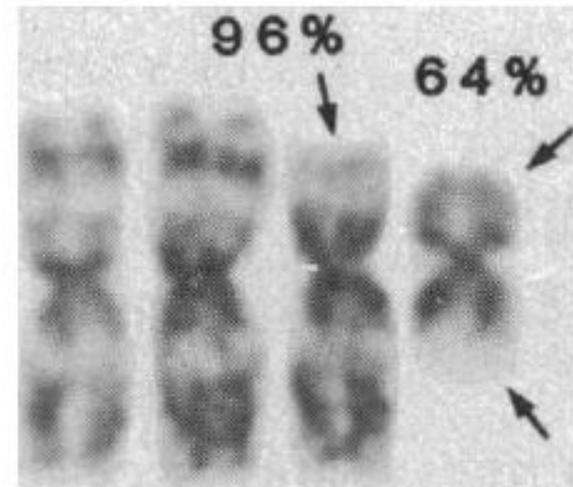
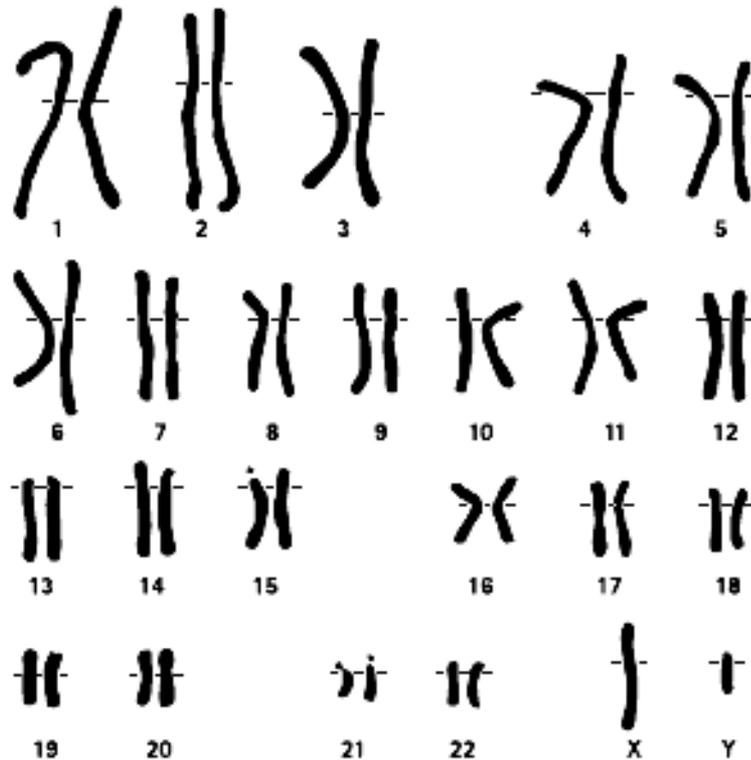


# Copy number change in SCLC

## □ Cytogenetic study

■ Establishment of cell line → triggered into metaphase

Chr 3 deletion in SCLC



**H69**

J WHANG-PENG

Science 1982;215:181-2

# SCLC, a forgotten disease ?

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- Difficulty in obtaining fresh material has hampered large genomic studies
- Novel technologies may allow use of FFPE material
- Better understanding the biology will help identify new targets and develop novel strategies

# Array-based comparative genomic hybridization of pulmonary neuroendocrine tumors

## Goals of the project

- To what extent are SCLCs and carcinoids genetically connected ?
- Can we identify potential treatment targets for SCLC?
- Are SCLC cell lines representative of SCLC tumors?

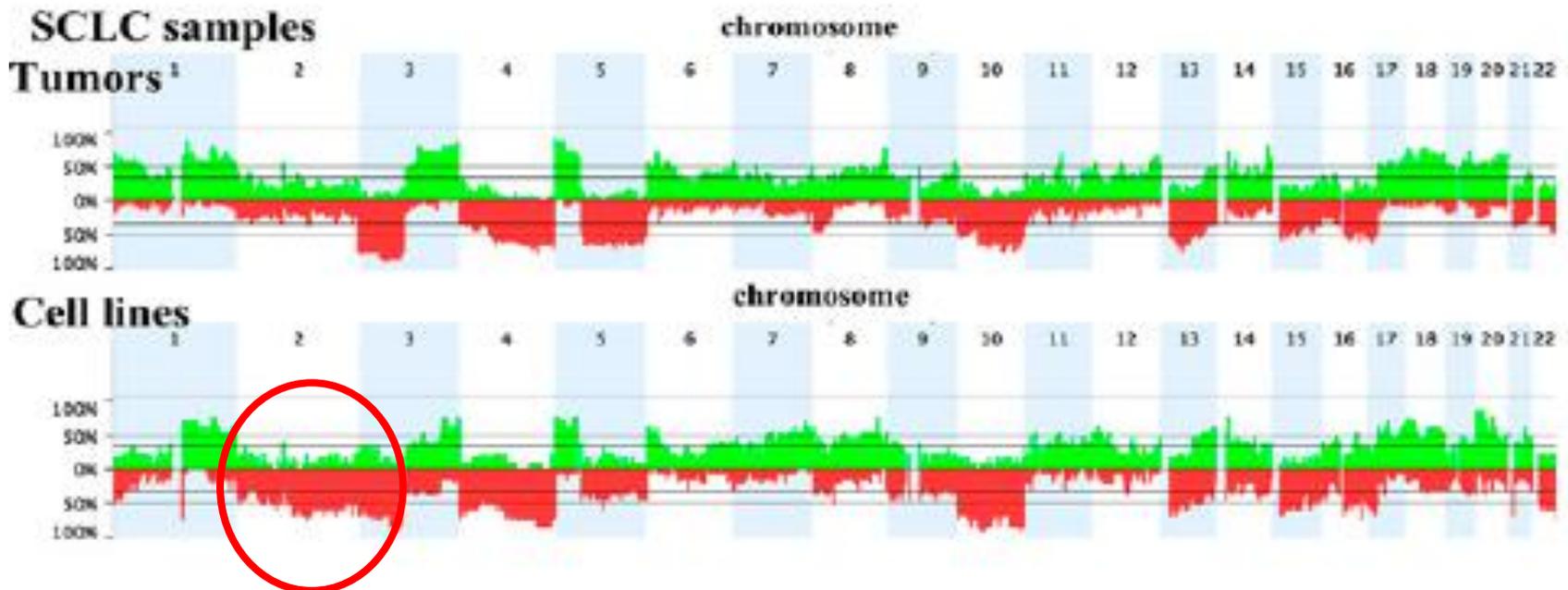
# Array CGH study of lung neuroendocrine tumors

	SCLC tumors	SCLC cell lines	Bronchial carcinoids	GI carcinoids
<b>Number</b>	33	13	19	9
<b>Cytogenetic bands with recurrent CN aberrancy</b>				
gains	122	98	86	92
losses	48	71	45	89
<b>Genes in cytogenetic bands with recurrent CN aberrancy</b>				
gains	8459	6851	536	3406
losses	5085	7232	1022	1178
<b>Cytogenetic bands with very high CN gain<sup>&amp;</sup></b>				
	4	11	0	0
<b>Genes in cytogenetic bands with very high CN gain<sup>&amp;</sup></b>				
	41	39	0	0

<sup>&</sup> 2-log ratio >3.0

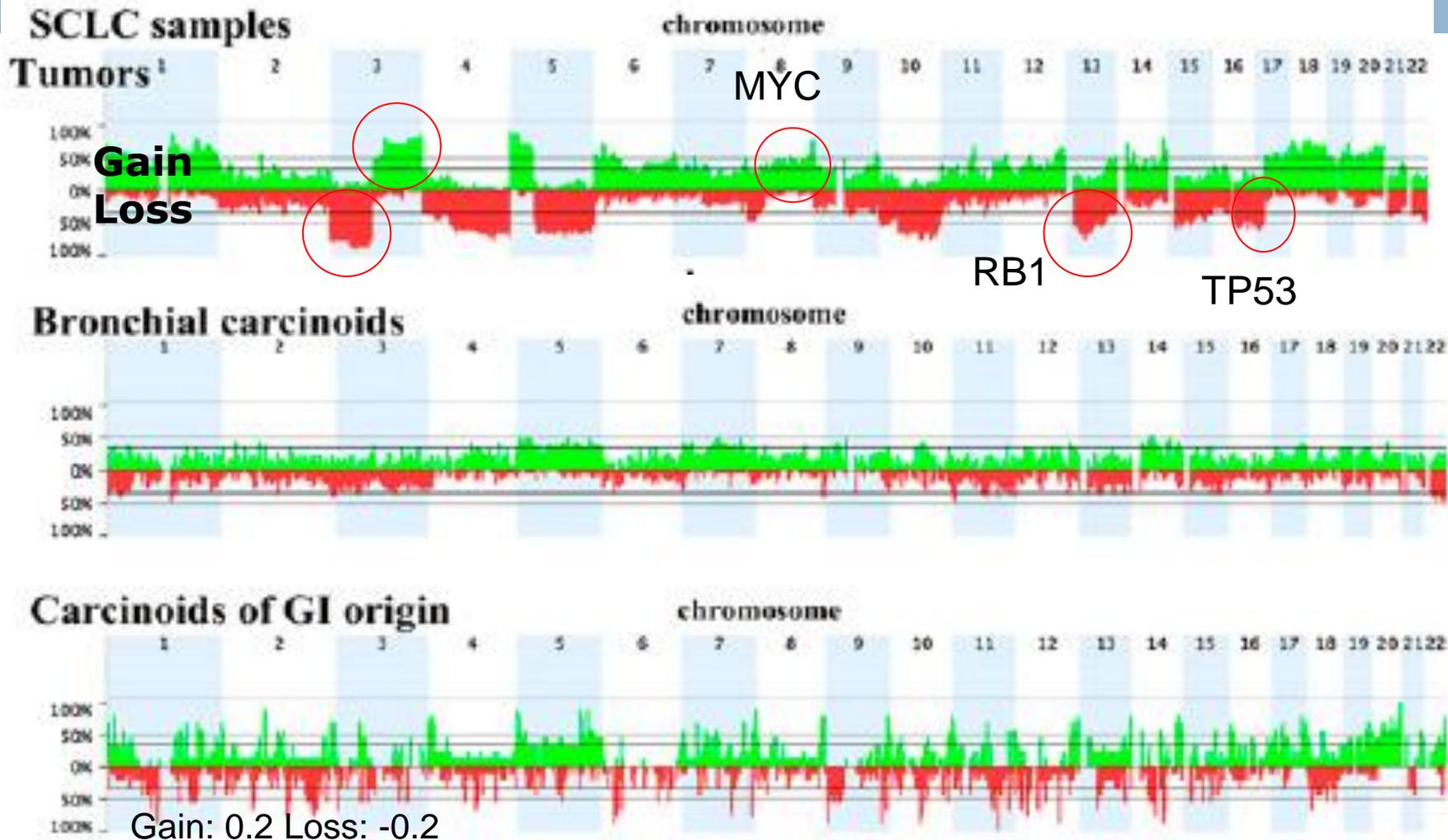
# SCLC cell lines may be representative of SCLC tumors genomically

- SCLC cell lines have more losses in chr. 2
- Only 74 genes were in cytogenetic bands in which the difference of frequencies of CNA between SCLC tumors and cell lines was  $> 50\%$  and statistically significant



# Comparison of SCLC tumors, bronchial carcinoids, and carcinoids of GI origin

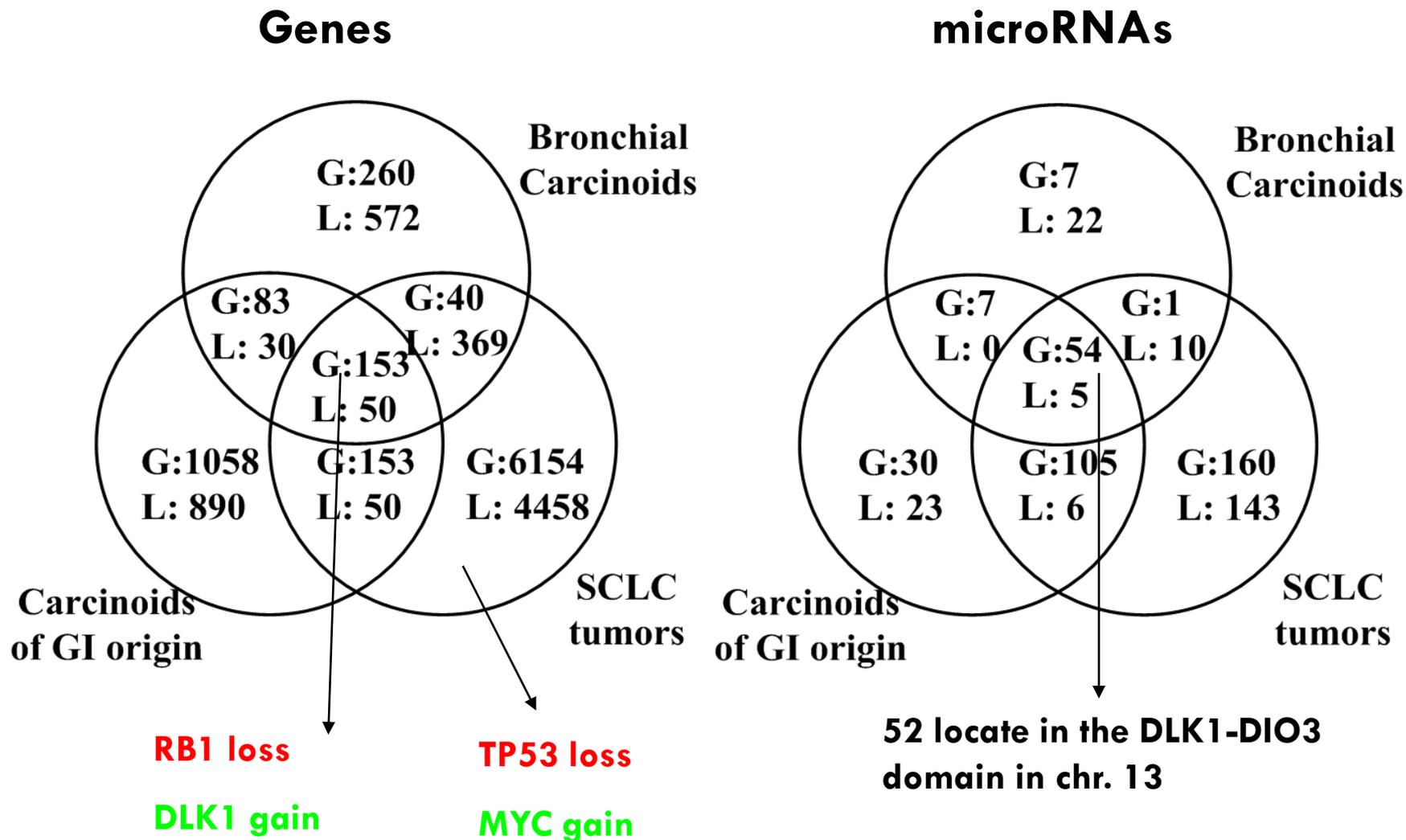
Frequency of CN alterations



Recurrent alteration: 35%

Voortman and Lee PNAS 2010

# Common copy number alterations of neuroendocrine tumors



# Potential drug targets for SCLC

## Genes with high frequencies of copy number alterations

Genes	Tumorscape		Our series	Genes	Tumorscape		Our series
	All cancers	SCLC	SCLC tumors		All cancers	SCLC	SCLC tumors
PIK3CA gain	21.6%	57.5%	75.8%	BCL2 gain	11.9%	47.5%	51.5%
AKT1 gain	16.3%	47.5%	63.6%	MCL1 gain	36.5%	57.5%	76.8%
PTEN loss	24.2%	62.5%	75.8%	PMAIP1 (Noxa) gain	11.7%	45%	66.8%
FRAP1 (mTOR) gain	13.4%	45.5%	54.5%	VHL loss	23.6%	80%	75.8%

# Cytogenetic bands with very high copy number gain<sup>&</sup> in SCLC tumors

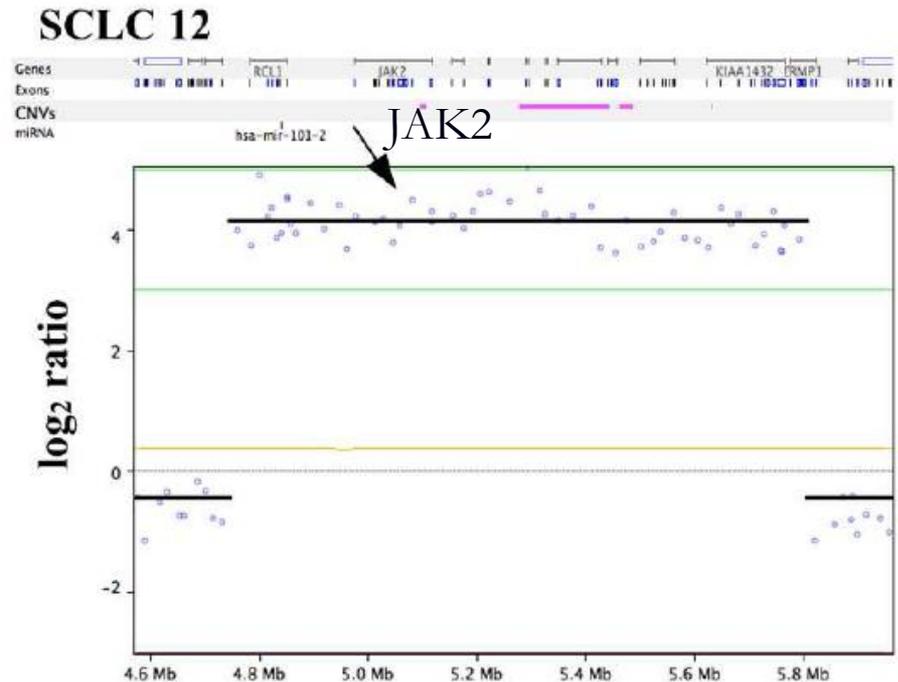
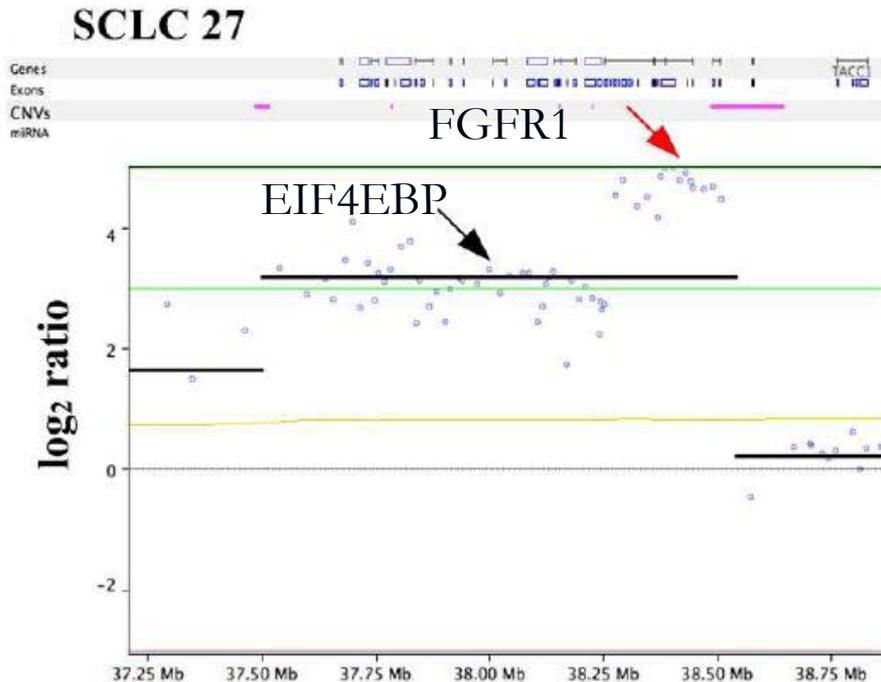
Cytogenetic band	Genes encoded	Candidate genes
8 p12 - p11.23	20	EIF4EBP1, FGFR1
9 p24.1	10	JAK2
9 p23 - p22.3	1	MPDZ
19 q13.12 - q13.13	10	*

<sup>&</sup>2-log ratio >3.0

\*chr. 19 encodes 10 transcription factors

# Potential drug targets

## High copy number gain



A  $\log_2$  ratio  $>4$  means that the copy number may be more than  $2^{(4+1)}=32$  copies

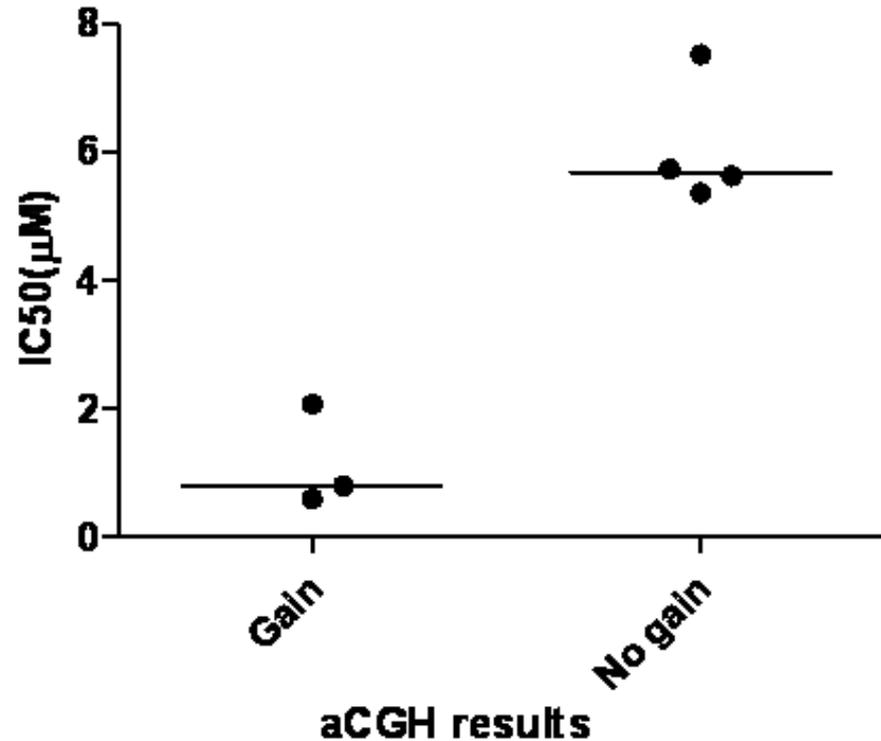
# Copy number gain of the JAK2 gene and the FGFR1 gene in SCLC tumors and cells

	Tumorscape	Our series	
		SCLC tumors	SCLC cell lines
JAK2	42.5%	27.8%	30.8%
FGFR1	27.5%	33.3%	30.8%

JAK2 and FGFR1 are potential targets in subgroups of SCLC

Tumorscape: aCGH data of 26 cancer types and >3000 specimens  
<http://www.broadinstitute.org/tumorscape/pages/portalHome.jsf>

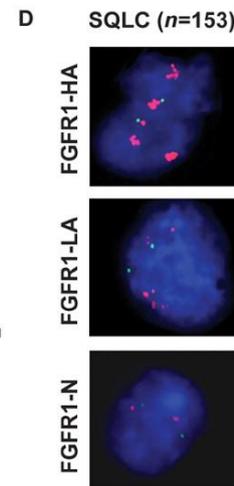
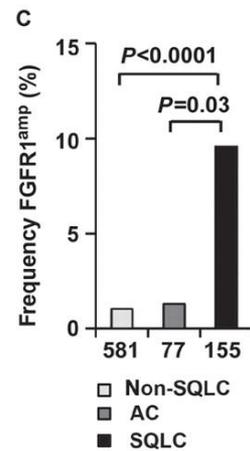
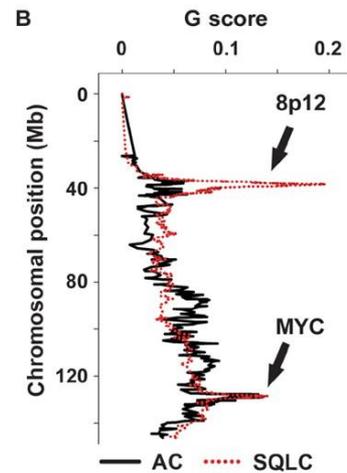
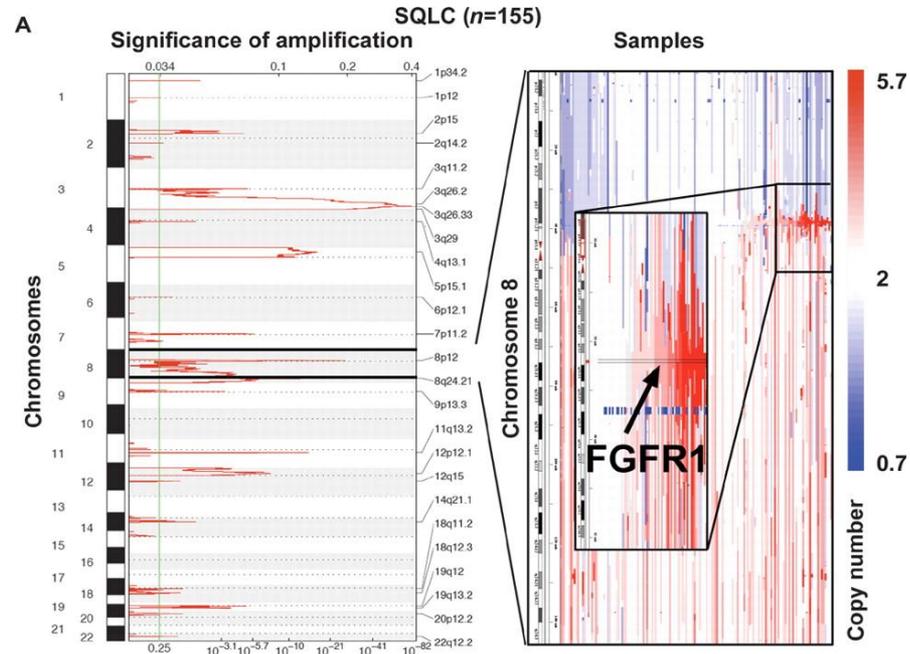
# Targeting FGFR1 in SCLC



**PD173074, a pan-FGFR inhibitor, on 7 SCLC cell lines**

**J-H Lee unpublished data**

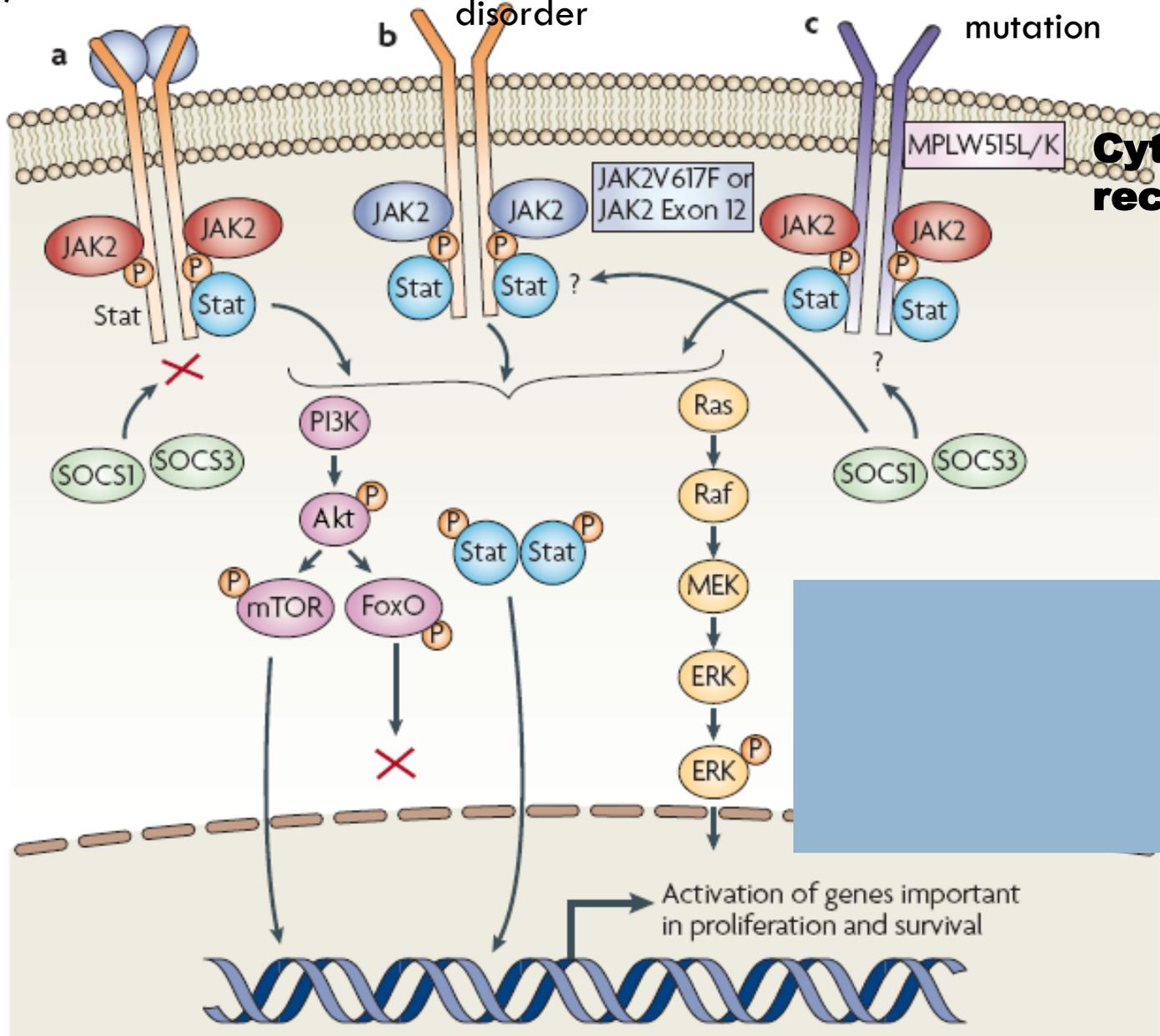
# FGFR1 is amplified in squamous cell lung cancer



IL-6, IFN

Myeloproliferative disorder

TPO receptor mutation

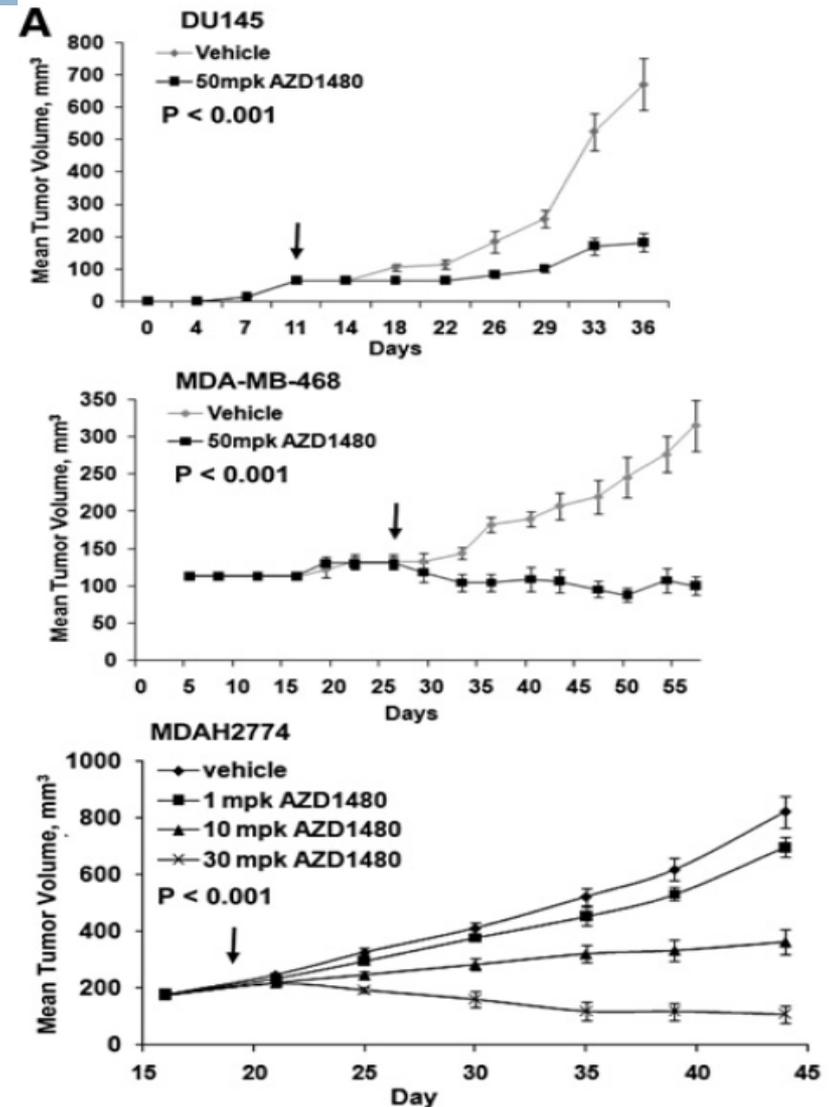


**Cytokine receptors**

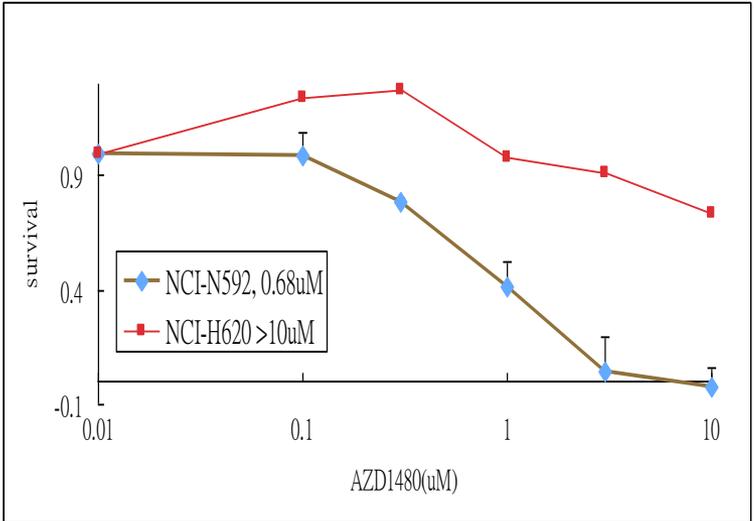
# Targeting JAK2 in SCLC

- AZD1480, a JAK1/JAK2 inhibitor
- AZD1480 slows cancer xenograft growth of many solid tumors
- AZD1480 shows little cytotoxicity in cell culture systems

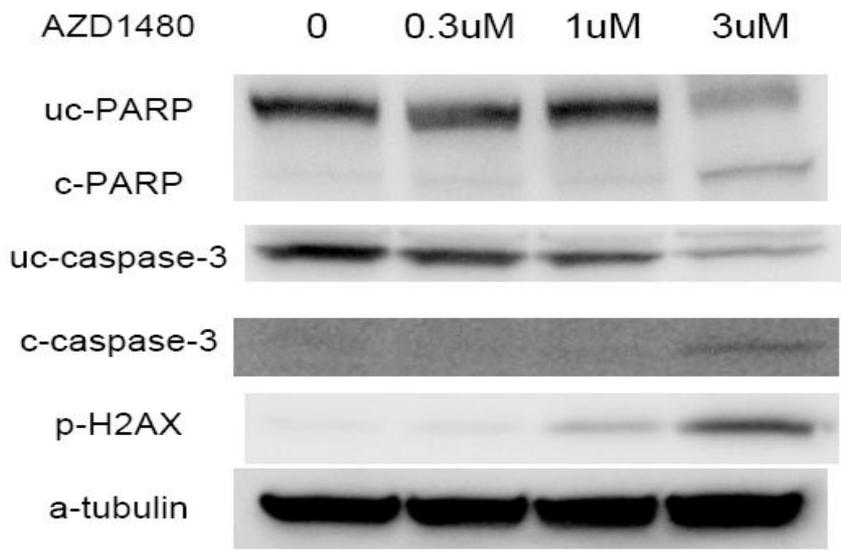
Cancer Cell 2009;16:487



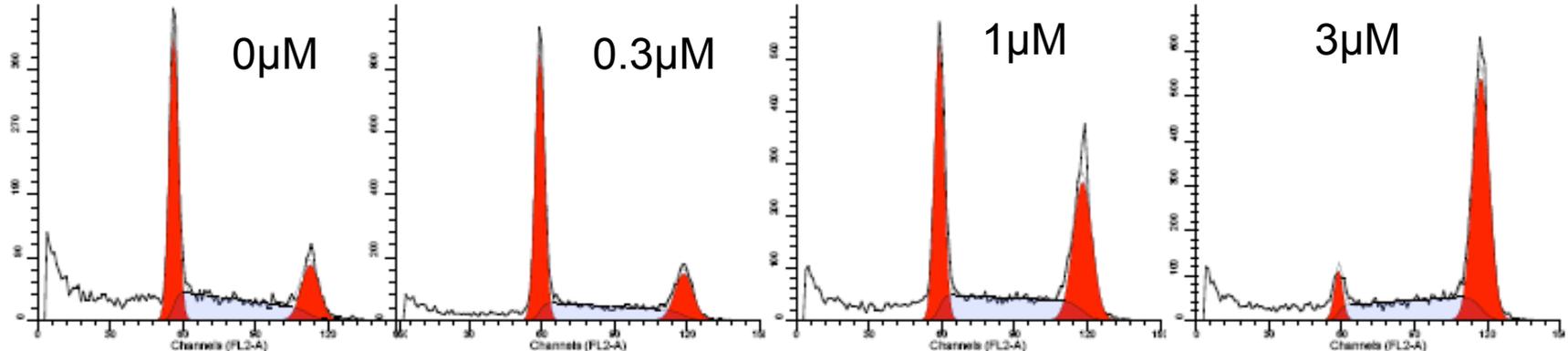
# AZD1480 kills a SCLC cell line carrying JAK2 gene amplification and induces G2/M arrest and apoptosis



NCI-N592: JAK2 amplification  
 NCI-H620: JAK2 deletion



NCI-N592 Treated with drug for 48 hours



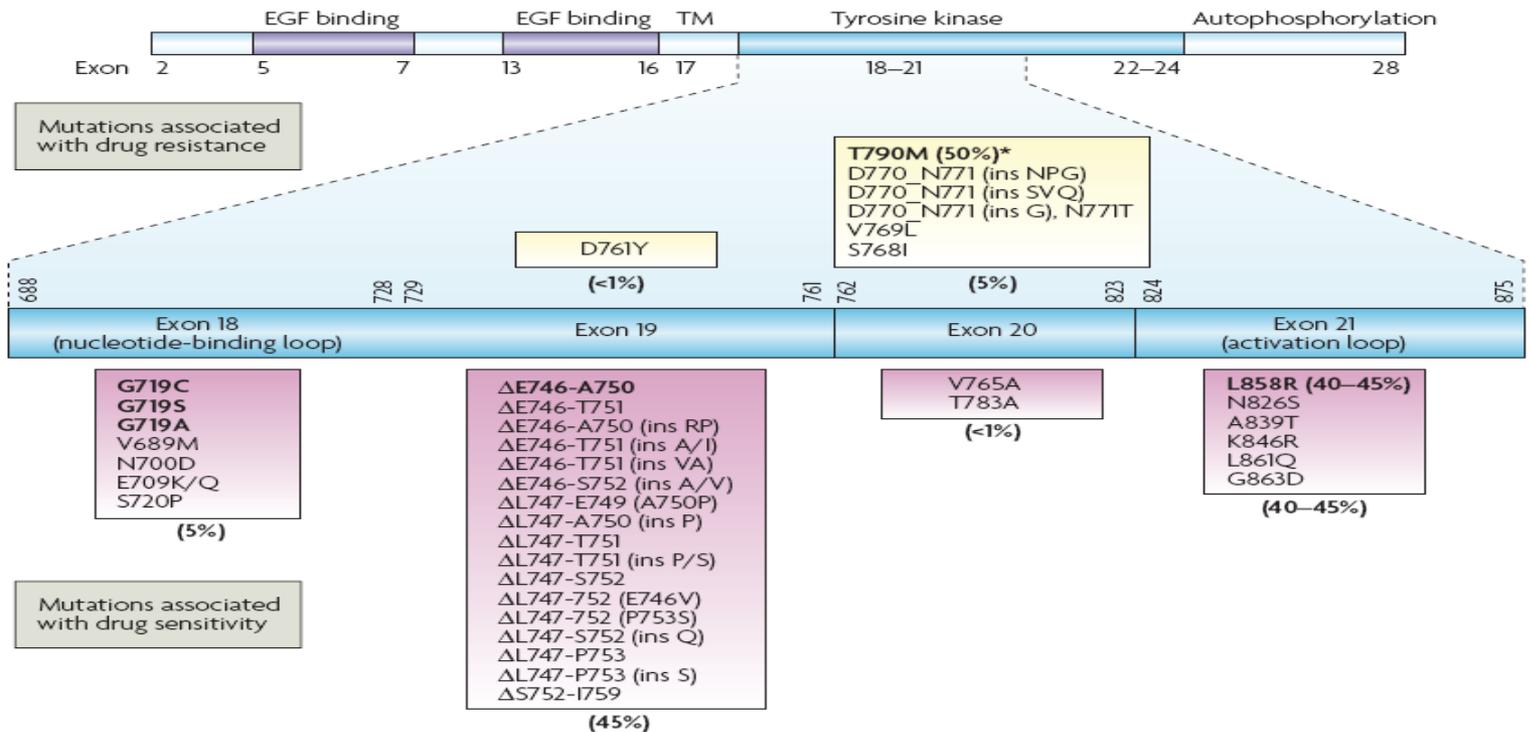
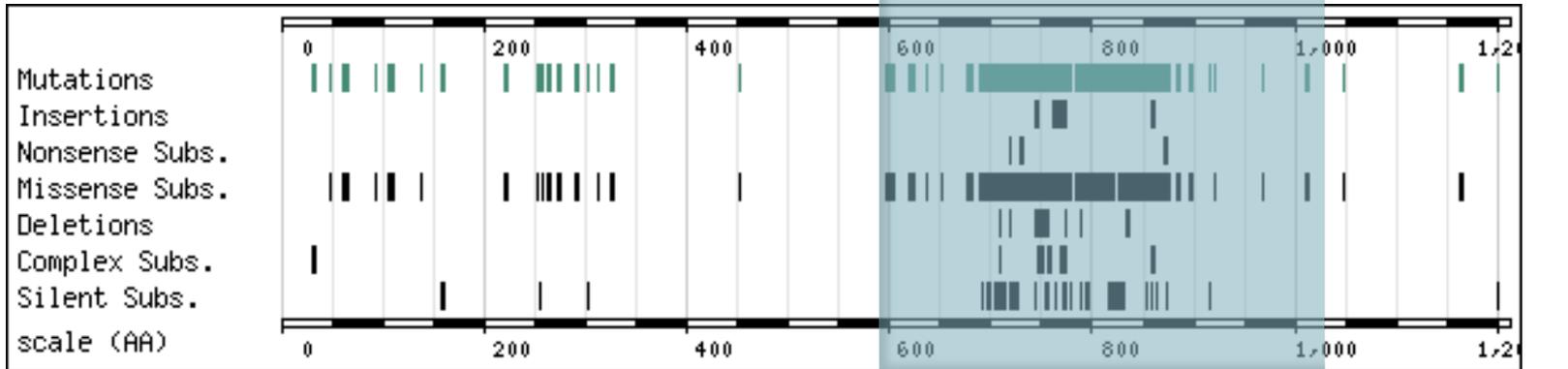
NCI-N592 Treated with drug for 24 hours

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# EGFR mutations

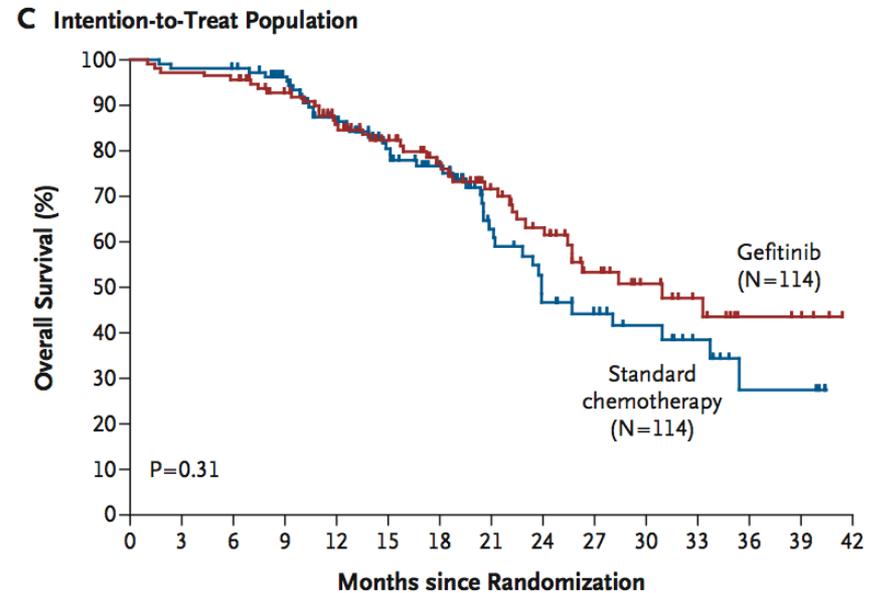
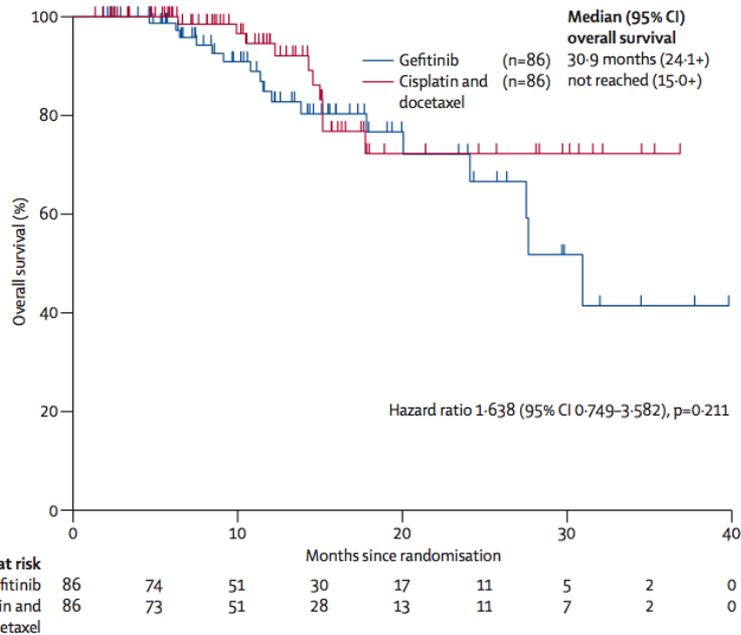
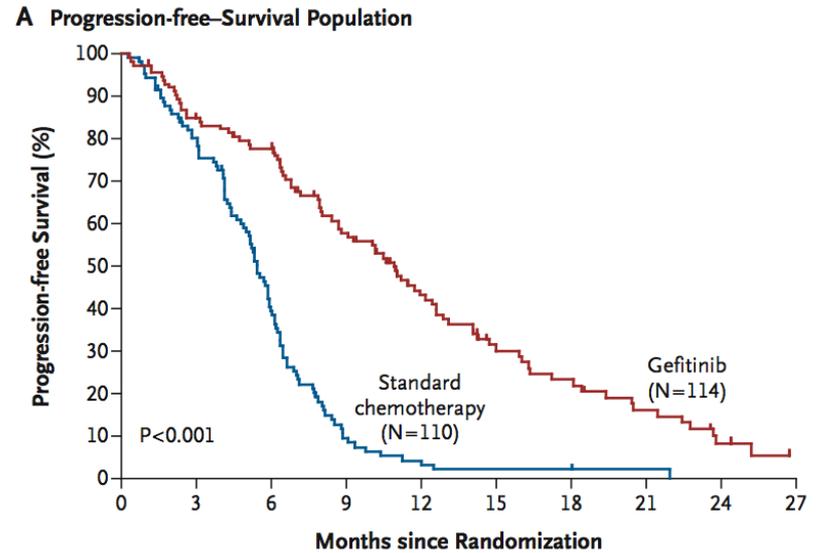
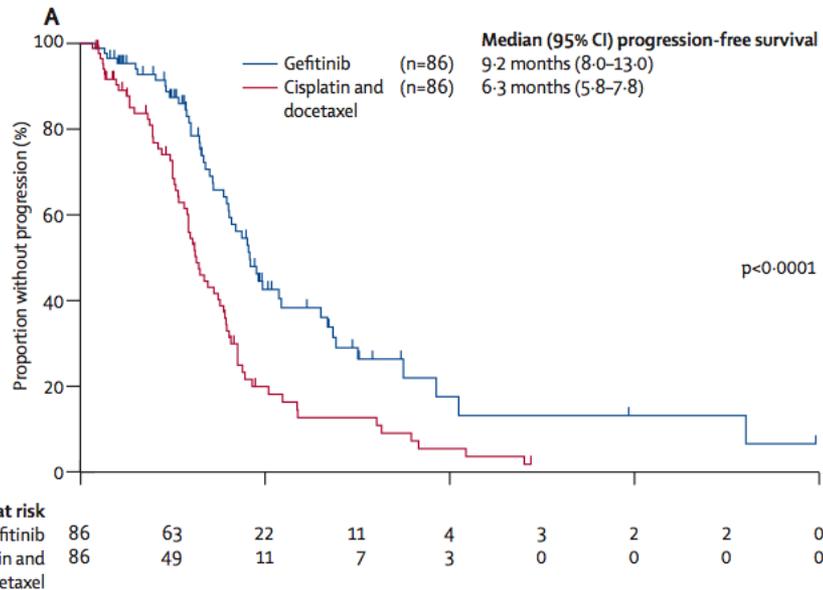


## Prospective clinical trials comparing EGFR-TKIs to chemotherapy in first-line in advanced NSCLC with EGFR mutations

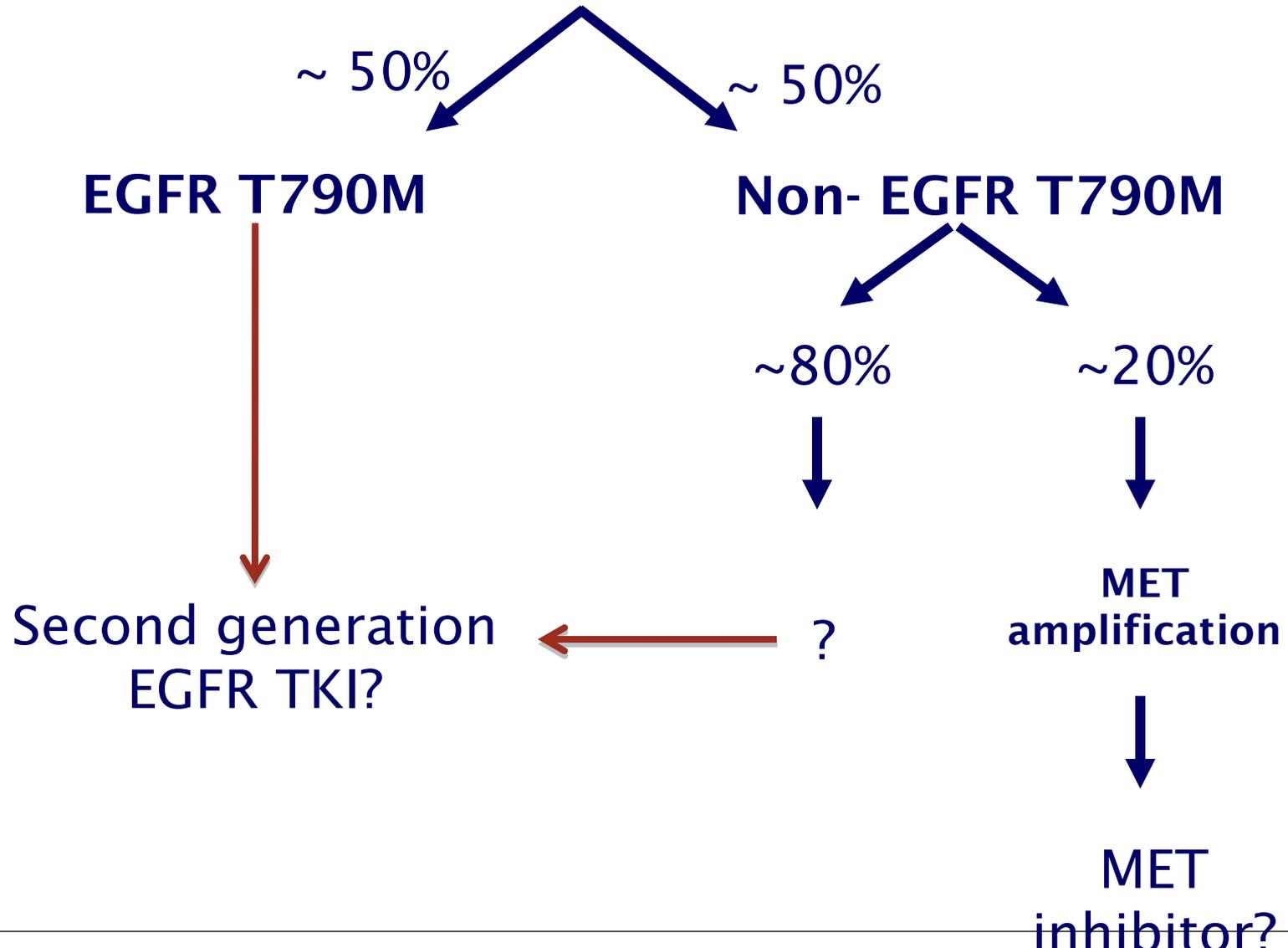
Author	Treatments	# patients	Response rate	PFS (m)	OS (m)
Mitsudomi (Lancet Oncol 2010)	Gefitinib	86	62.1	9.2	30.9
	DDP/DXL	86	23.2 (p<.001)	6.3 (p<.001)	NR (p=NS)
Maemondo (NEJM 2010)	Gefitinib	114	73.7	10.8	30.5
	CRB/PXL	110	30.7 (p<.001)	5.4 (p<.001)	23.6 (p=NS)
Zhou (ESMO 2010)	Erlotinib	82	83	13.1	NA
	CRB/GEM	72	36 (p<.0001)	4.6 (p<.0001)	NA
EURTAC	Erlotinib CRB/GEM DDP/DXL				

NR=not reached; NA=not available; NS=not significant

# EGFR mutant selected



# Mechanisms of Acquired Resistance to EGFR TKIs



# PF-00299804

## *in vitro* Activity against EGFR, HER2 and KRASmut NSCLC Cell Lines

Cell Line	EGFR mutation	ERBB2 mutation	K-ras mutation	Gefitinib IC <sub>50</sub>	PF00299804 IC <sub>50</sub>
A549	WT	WT	G12S	> 10 μM	> 10 μM
H441	WT	WT	G12V	> 10 μM	4 μM
Calu-3	WT	WT HER2+++	WT	1.4 μM	0.063 μM
H1819	WT	WT HER2+++	WT	0.42 μM	0.029 μM
H1781	WT	Ins G776V,C	WT	> 10 μM	0.275 μM
HCC 827	Del E746_A750	WT	WT	0.008 μM	0.002 μM
HCC 4006	Del L747_E749	WT	WT	0.050 μM	0.004 μM
PC-9	Del E746_A750	WT	WT	0.023 μM	0.002 μM
H3255	L858R	WT	WT	0.075 μM	0.007 μM
H3255 GR	L858R/T790M	WT	WT	> 10 μM	0.119 μM
H1975	L858R/T790M	WT	WT	> 10 μM	0.44 μM

- MTS 72-Hour Proliferation Assay; 6-12 wells per assay; all experiments repeated at least 3X



# Case Presentation

- 52 y/o AA male
  - ▣ Never smoker
  - ▣ Diagnosed with NSCLC adenocarcinoma
  - ▣ Stage IV, multiple lesions in LLL and RUL
- No Past Medical History
- No Family History
  
- Treatment:
  - ▣ carboplatin/paclitaxel/bevacizumab
    - 8 cycles, initial Partial Response (PR) then Progressive Disease (PD)
  - ▣ erlotinib 150 mg PO daily, (6 weeks) PD
  - ▣ Referred for evaluation at NCI

# Treatment Course

- Screened for PF-00299804 trial:
  - KRAS wild type (wt)
  - [EGFR wt]
- Enrolled on trial PF-00299804 for patients who have failed EGFR TKIs
- Started on PF-00299804
  - 45 mg PO daily

## Pretreatment and after 4 cycles of PF-00299804

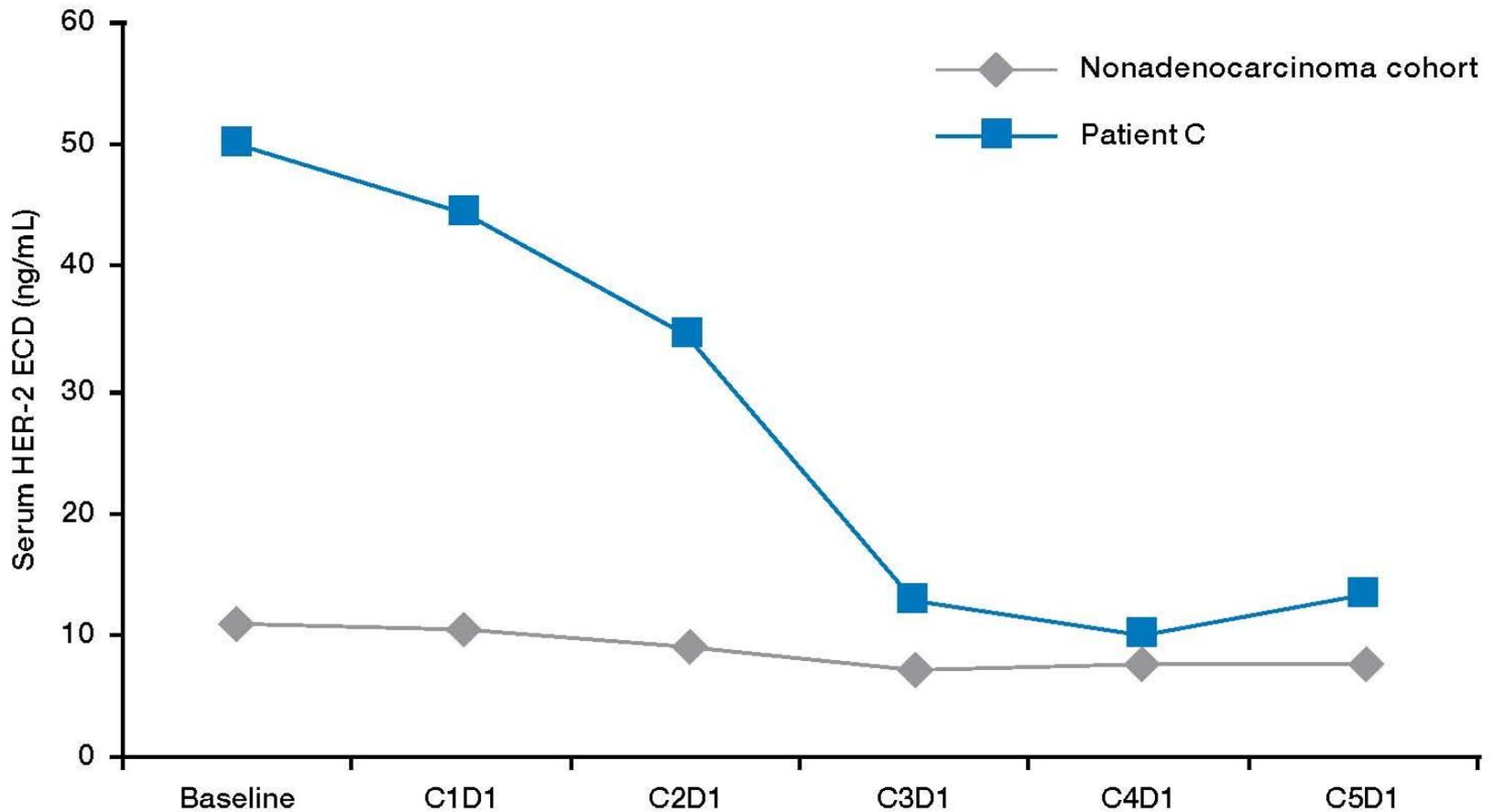


October 21, 2008



January 16, 2009

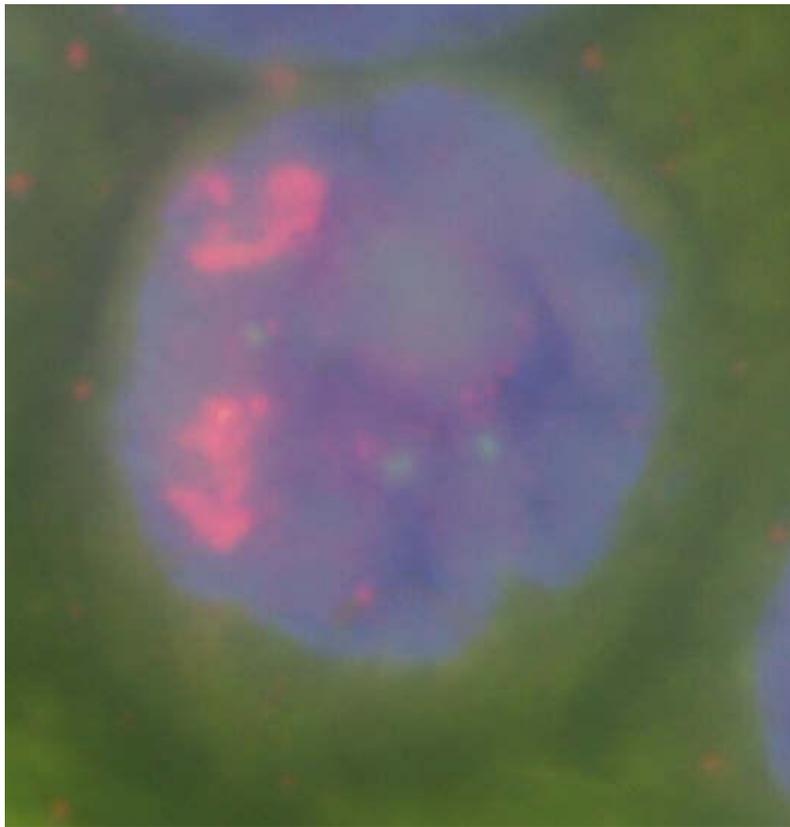
# Serum HER-2 levels at baseline and through 4 cycles of PF-00299804 therapy



C = cycle; D = day; ECD = extracellular domain; EGFR = epidermal growth factor receptor; WT = wild type.

# Stain Tumor for Her2/Neu

## HER2/Neu FISH



## PATHOLOGY

- IHC 2+
  
- Her2/Neu Amplified
  - ▣ Cells counted 20
  - ▣ HER2 = 10.3
  - ▣ CEP17 = 2.4
  - ▣ HER2/CEP17 Ratio: 4.3

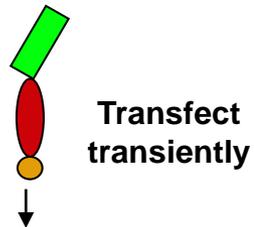
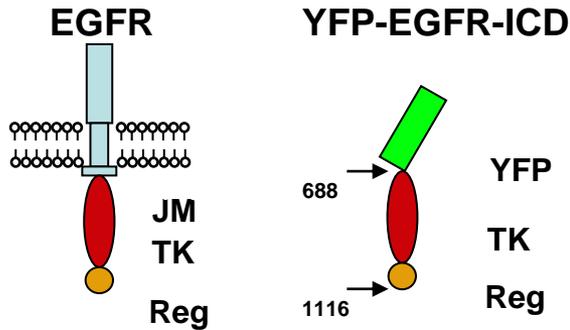
# EGFR mutations in African-Americans

<i>Patient number</i>	<i>Gender</i>	<i>Histology</i>	<i>Age at diagnosis</i>	<i>Smoking history (PY)</i>	<i>Stage</i>	<i>EGFR mutation</i>	<i>EGFR (IHC)</i>	<i>KRAS mutation</i>	<i>Response to EGFR-TKI</i>
<b>Non-smokers</b>									
NS-001	Male	Ad	52	<1	IIIB	<b><u>N771GY</u></b>	WP	WT	PD
NS-002	Female	Ad	64	Never	IIIA	<b><u>L858R</u></b>	SP	WT	NA
NS-003	Male	Ad	50	Never	IV	<b><u>Del19</u></b>	NA	WT	PR
NS-004	Female	Ad	45	Never	IV	<b><u>WT</u></b>	NA	WT	PD
NS-005	Male	Ad	48	Never	IIIB	<b><u>767A-769V dup</u></b>	NA	WT	NA
NS-006	Male	Ad	49	Never	IV	<b><u>WT</u></b>	SP	WT	PD
NS-007	Female	Ad	71	Never	IV	<b><u>WT</u></b>	NA	WT	PR
NS-008	Female	Ad	58	Never	IV	<b><u>Del19</u></b>	NA	WT	PR
<b>Smokers</b>									
S-001	Female	NOS	54	30	IV	<b><u>WT</u></b>	WP	13 (GGC > GAC)	NA
S-002	Female	Ad	58	35	IV	<b><u>WT</u></b>	SP	WT	NA
S-003	Female	SCC	51	30	IB	<b><u>WT</u></b>	NA	WT	NA
S-004	Male	NOS	68	50	IV	<b><u>WT</u></b>	NA	WT	PD
S-005	Male	NOS	49	15	IV	<b><u>WT</u></b>	NA	WT	PD
S-006	Female	Ad	58	35	IV	<b><u>WT</u></b>	NA	WT	NA
S-007	Male	Ad	73	50	IIIB	<b><u>WT</u></b>	NA	NA	NA
S-008	Female	Ad	65	50	IIIB	<b><u>WT</u></b>	NA	12 (GGT > TGT)	NA

# Mutated YFP-tagged EGFR intracellular domain activate the downstream signal

Harada T et al Oncogene 2010

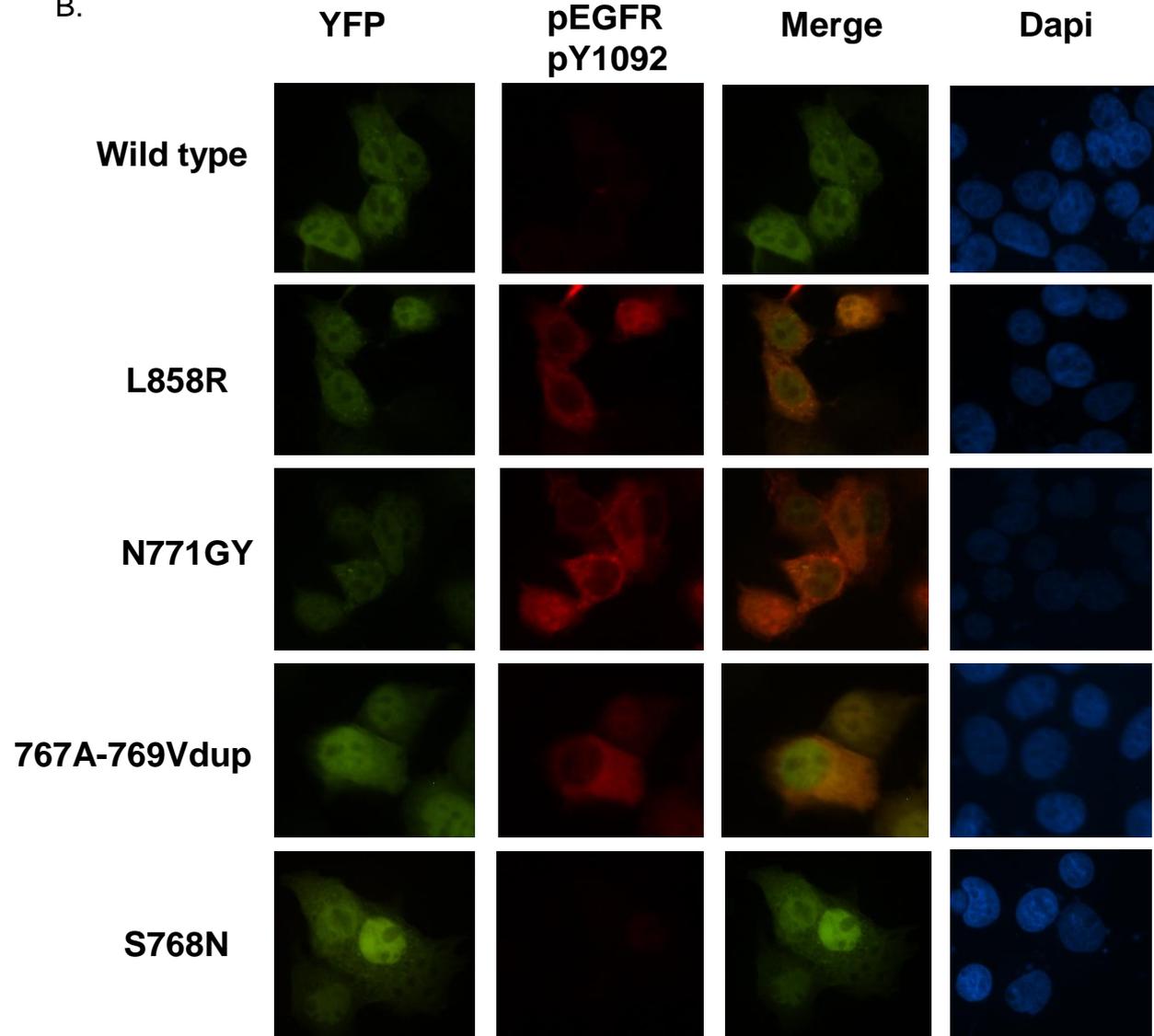
A.



Immunofluorescence

Observation under fluorescence microscope

B.



# erlotinib

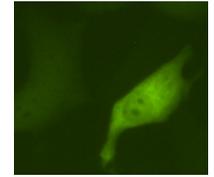
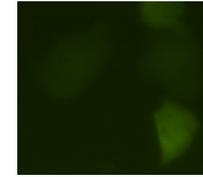
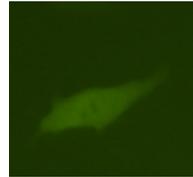
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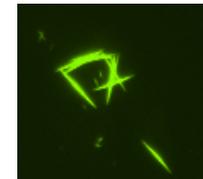
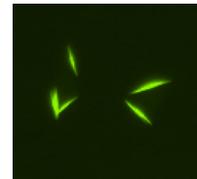
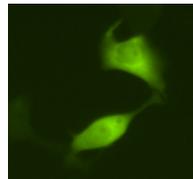
300

3000 nM

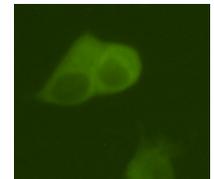
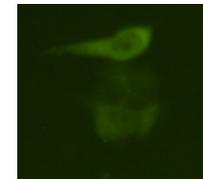
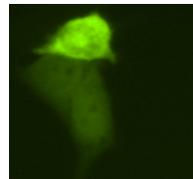
Wild type



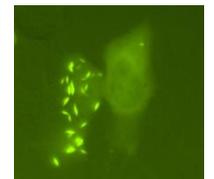
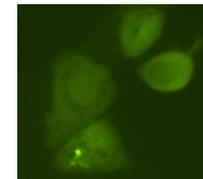
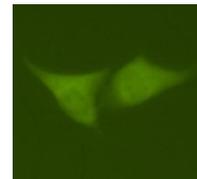
L858R



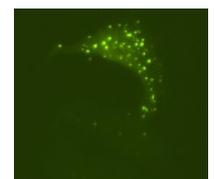
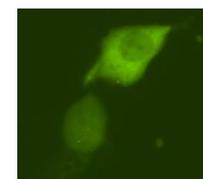
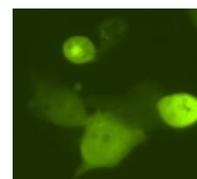
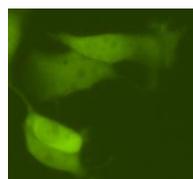
L858R/T790M



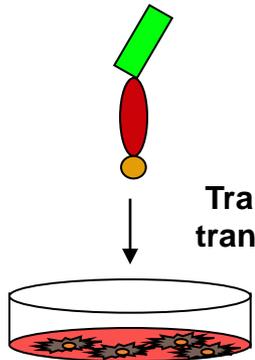
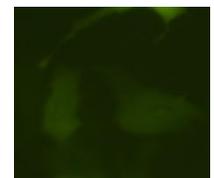
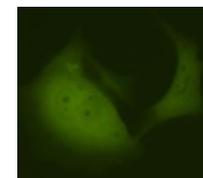
N771>GY



767-769dup



S768N



Transfect transiently

drug treatment  
20h

Observation under  
fluorescent microscope

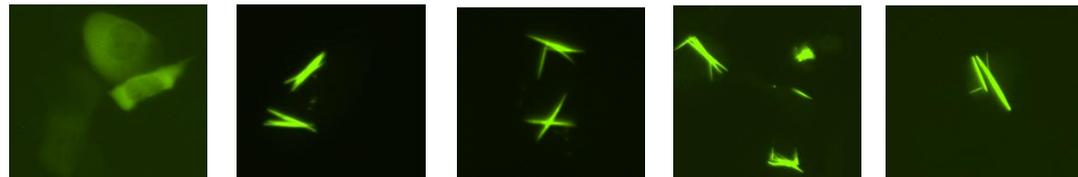
PF00299804

0 1 10 50 100 nM

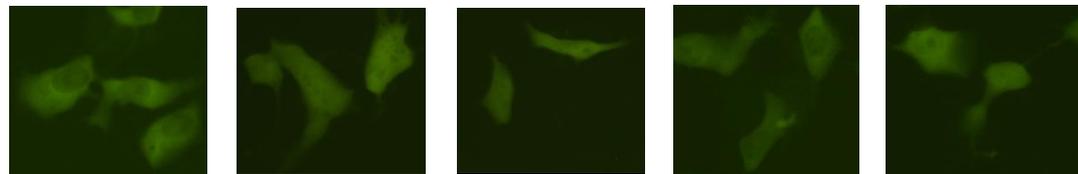
Wild type



L858R



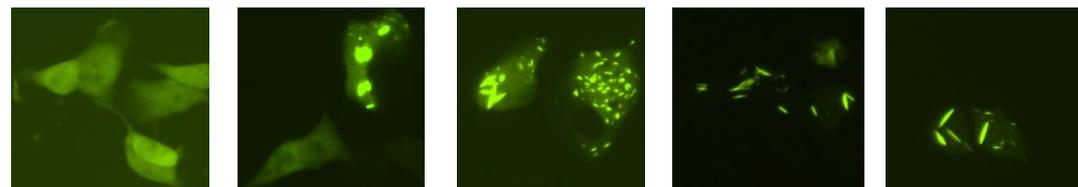
L858R/T790M



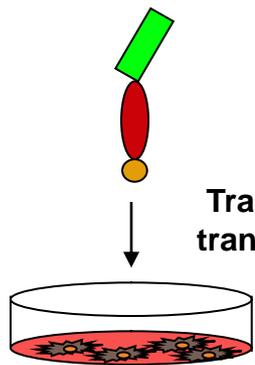
N771>GY



767-769dup



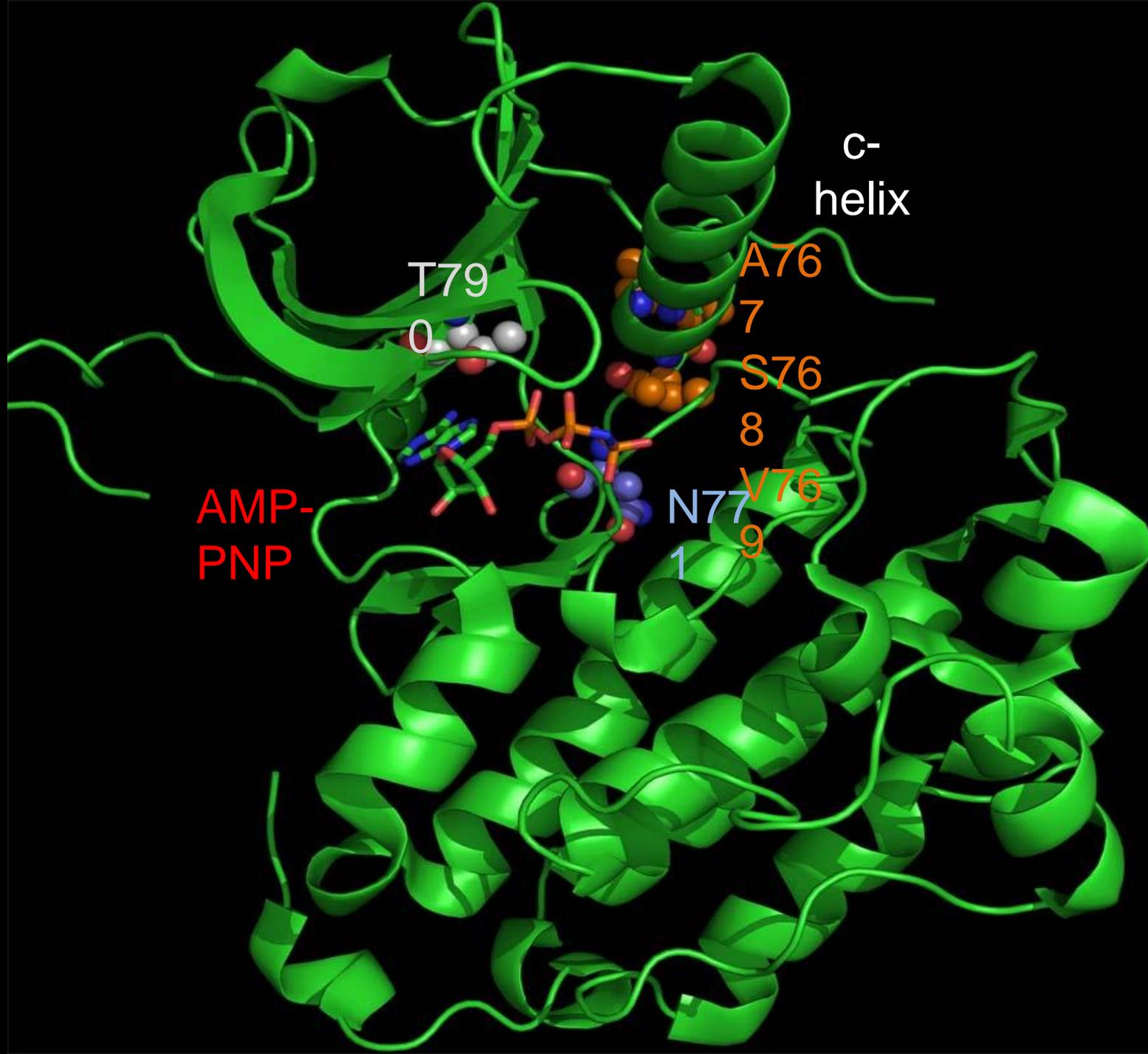
S768N



Transfect transiently

drug treatment  
20h

Observation under  
fluorescent microscope



c-  
helix

T79  
0

A76  
7

S76  
8

N77  
1

V76  
9

AMP-  
PNP

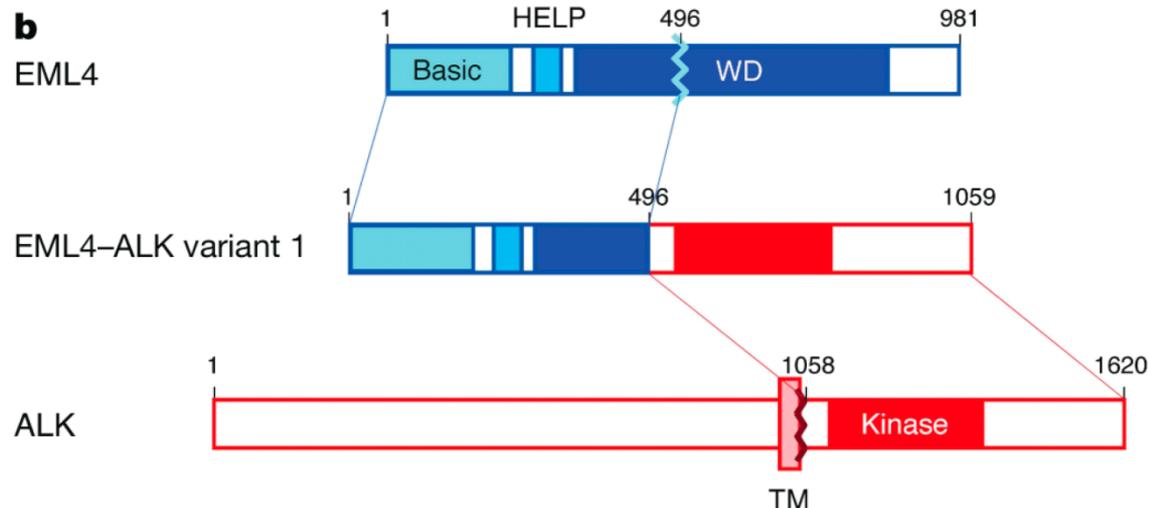
# *EML4-ALK* Translocations in NSCLC

Vol 448 | 2 August 2007 | doi:10.1038/nature05945

nature

## Identification of the transforming *EML4-ALK* fusion gene in non-small-cell lung cancer

Manabu Soda<sup>1,2</sup>, Young Lim Choi<sup>1</sup>, Munehiro Enomoto<sup>1,2</sup>, Shuji Takada<sup>1</sup>, Yoshihiro Yamashita<sup>1</sup>, Shunpei Ishikawa<sup>5</sup>, Shin-ichiro Fujiwara<sup>1</sup>, Hideki Watanabe<sup>1</sup>, Kentaro Kurashina<sup>1</sup>, Hisashi Hatanaka<sup>1</sup>, Masashi Bando<sup>2</sup>, Shoji Ohno<sup>2</sup>, Yuichi Ishikawa<sup>6</sup>, Hiroyuki Aburatani<sup>5,7</sup>, Toshiro Niki<sup>3</sup>, Yasunori Sohara<sup>4</sup>, Yukihiko Sugiyama<sup>2</sup> & Hiroyuki Mano<sup>1,7</sup>



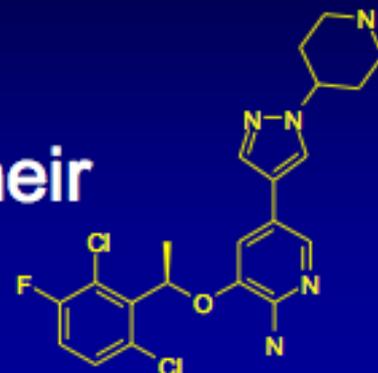
***EML4-ALK*  
frequency:**

**~4% (64/1709)**

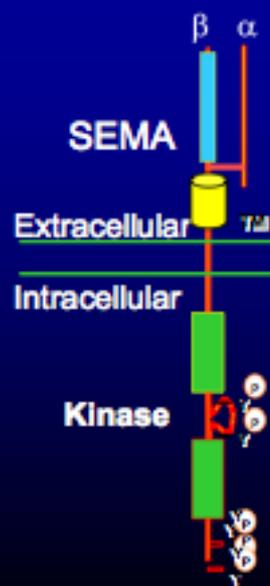
**Primarily lung  
adenocarcinoma**

# PF-02341066

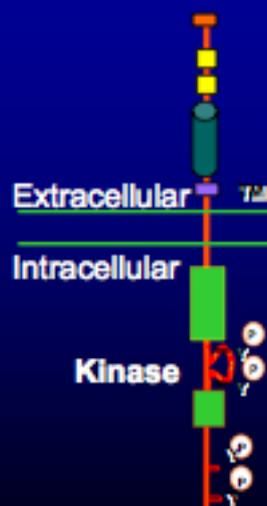
Potent & selective ATP competitive oral inhibitor of MET and ALK kinases and their oncogenic variants



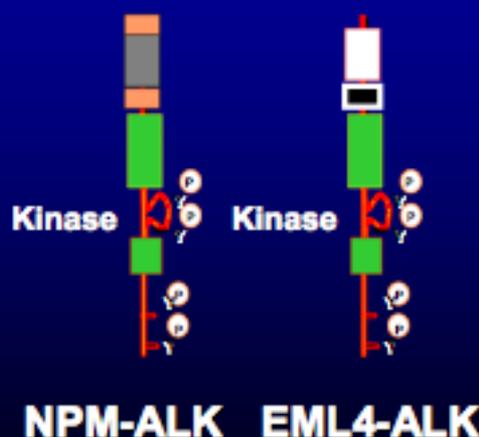
## MET/HGFR



## ALK



## Cytoplasmic Fusion Variants of ALK

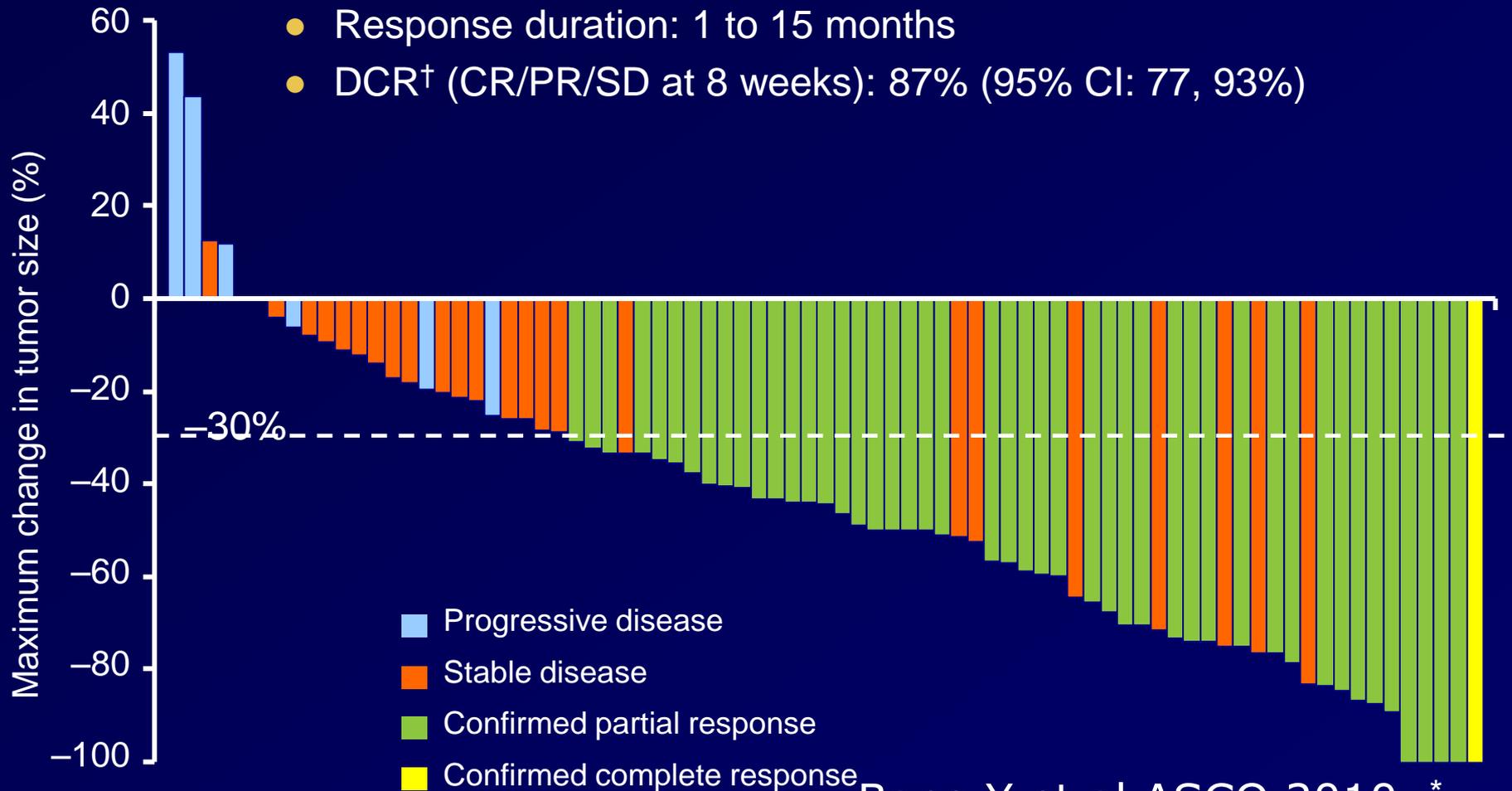


Kinase	IC50 (nM) Mean	Selectivity Ratio <sup>a</sup>
c-Met	8	—
ALK	20	2X
RON	248	31X
Axl	308	39X
Tie-2	448	56X
Trk A	580	73X
Trk B	399	50X

PF2341066 was >100X selective for Met/ALK across a panel of 150 additional kinases.

# Tumor Responses to Crizotinib for Patients with *ALK*-positive NSCLC

- Objective response rate : 57% (95% CI: 46, 68%)
- Response duration: 1 to 15 months
- DCR<sup>†</sup> (CR/PR/SD at 8 weeks): 87% (95% CI: 77, 93%)

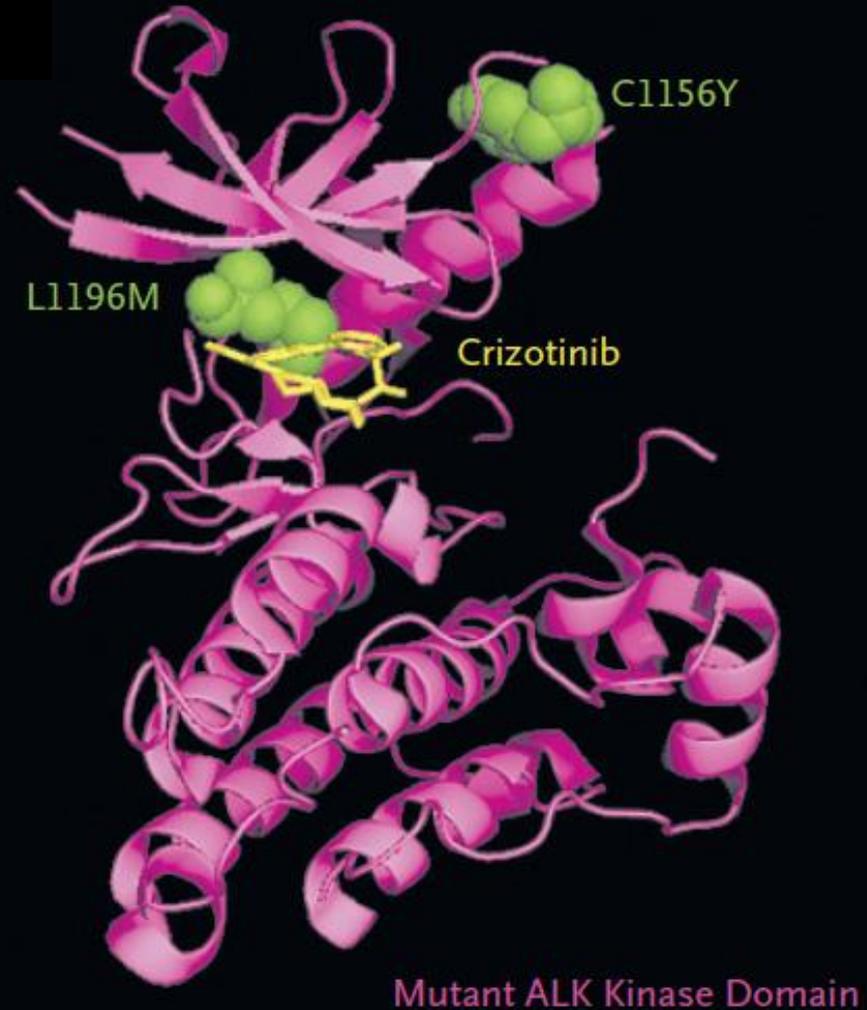
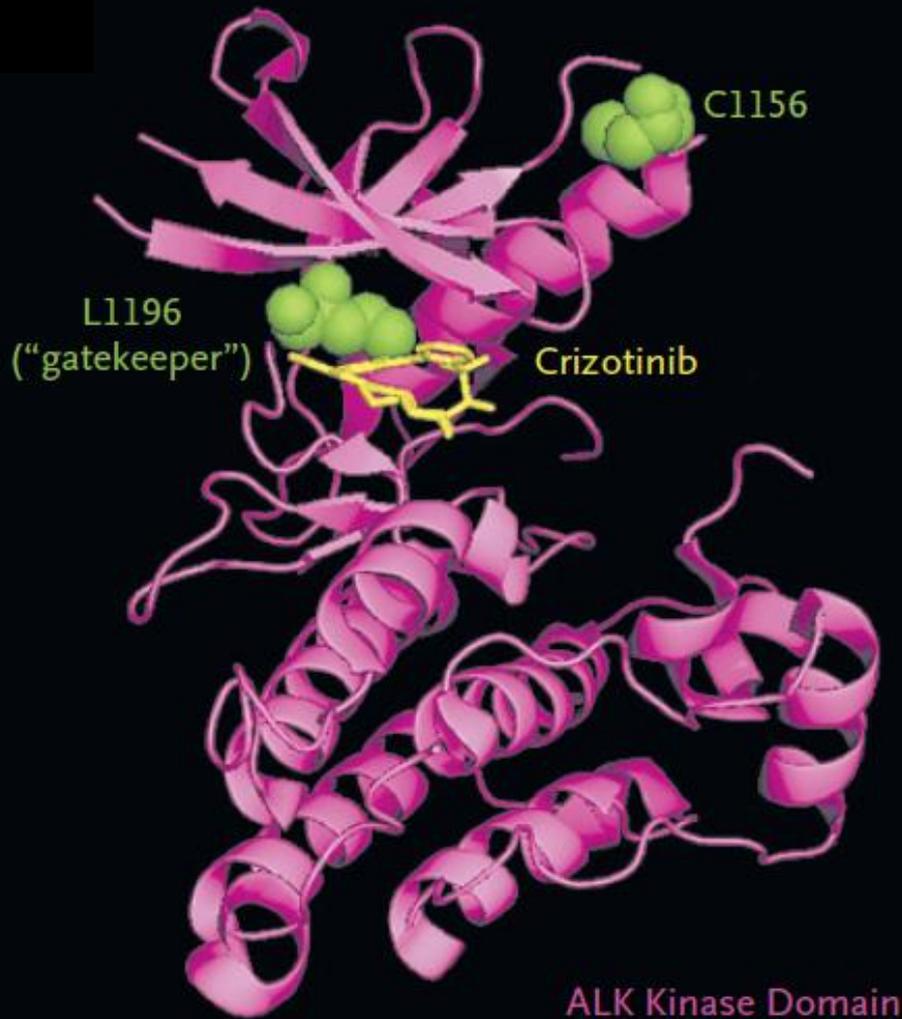


\*Partial response patients with 100% change have non-target disease present

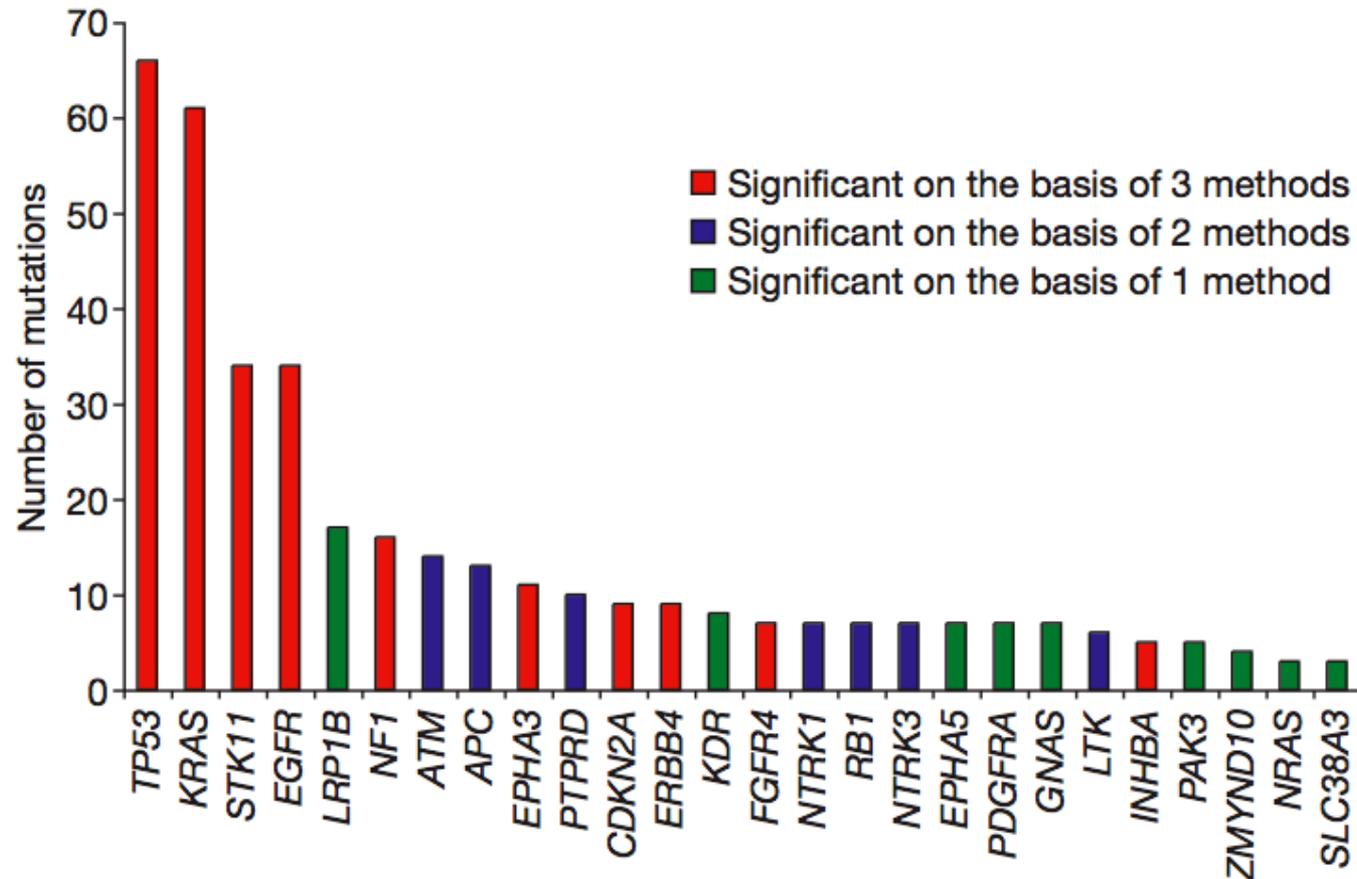
# Main patient characteristics

- NSCLC screened = 1500
- NSCLC with Alk translocation = 82 + 2
- Male 52%
- White 56%, Asian 35%
- $\geq$  prior lines 41%
- Adenocarcinoma 96%
- Never-smoking 76%

# Predicted Crystal Structure of the Kinase Domain of ALK and Resistant Mutants

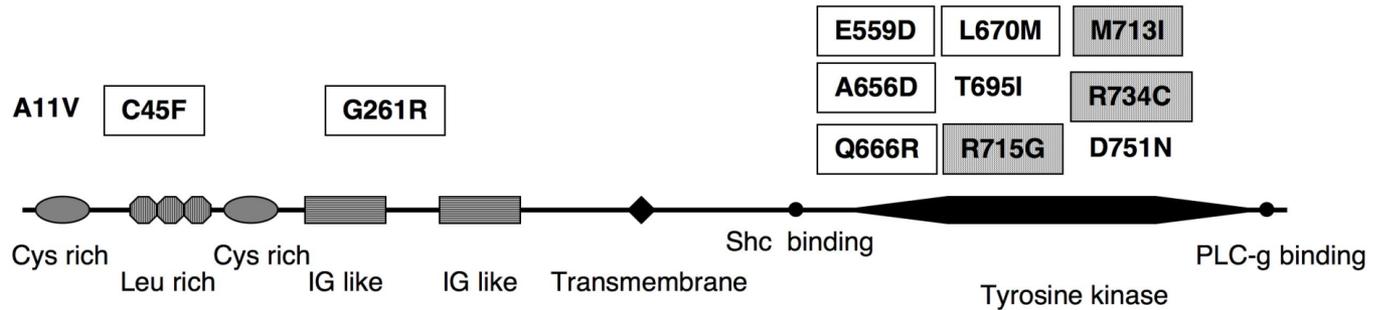


# Significantly mutated genes in adenocarcinoma of the lung



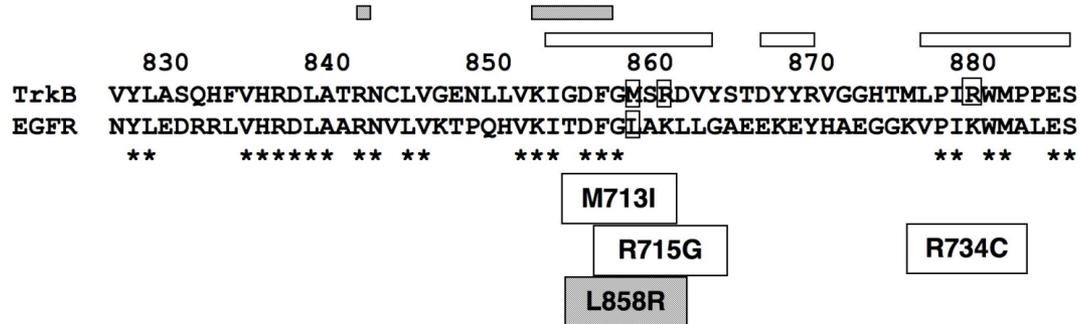
# TRKB domain and reported mutations

A



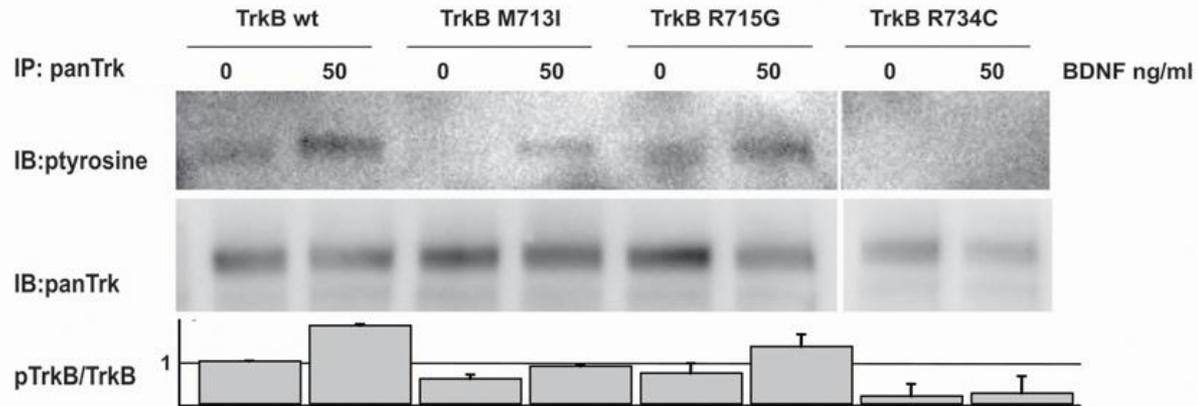
B

ATP\_binding  
Act\_loop

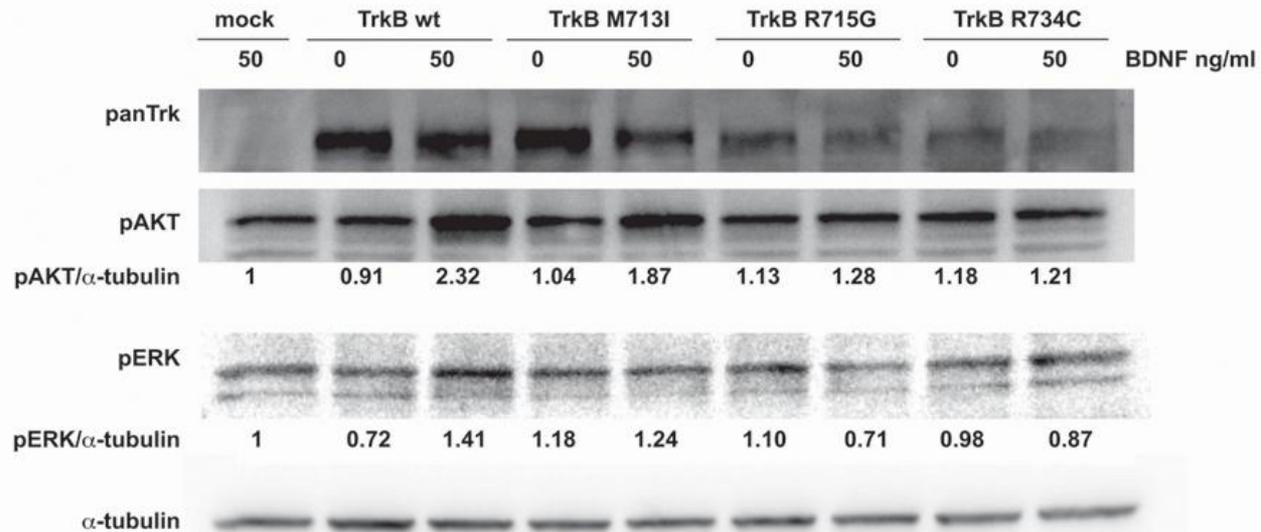


# TRKB in NIH3T3 transfected cells

**A**



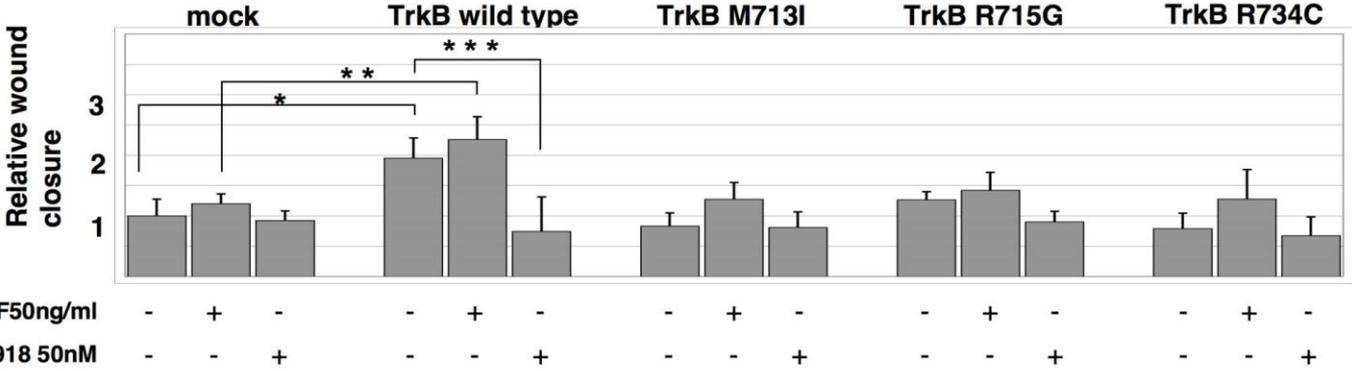
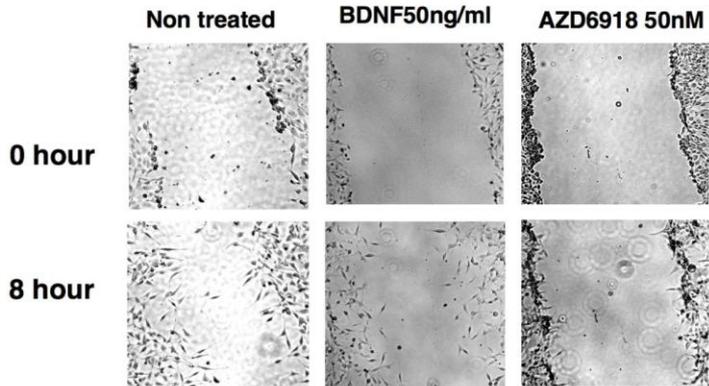
**B**



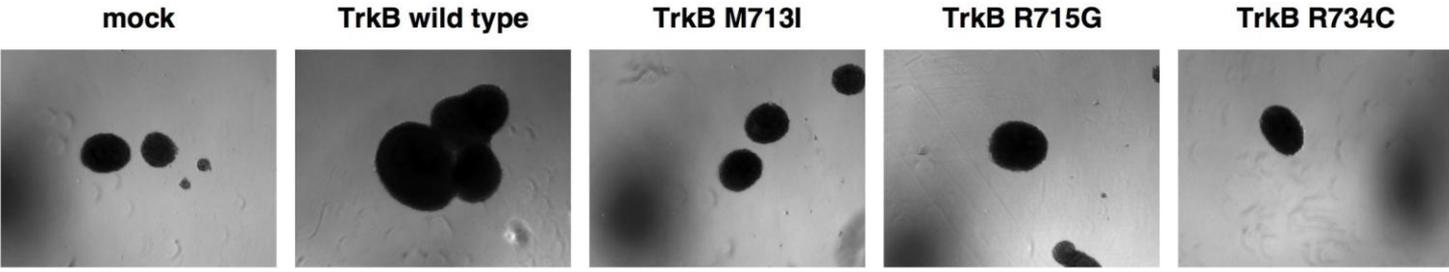
# Cell migration and anchorage independent growth

**A**

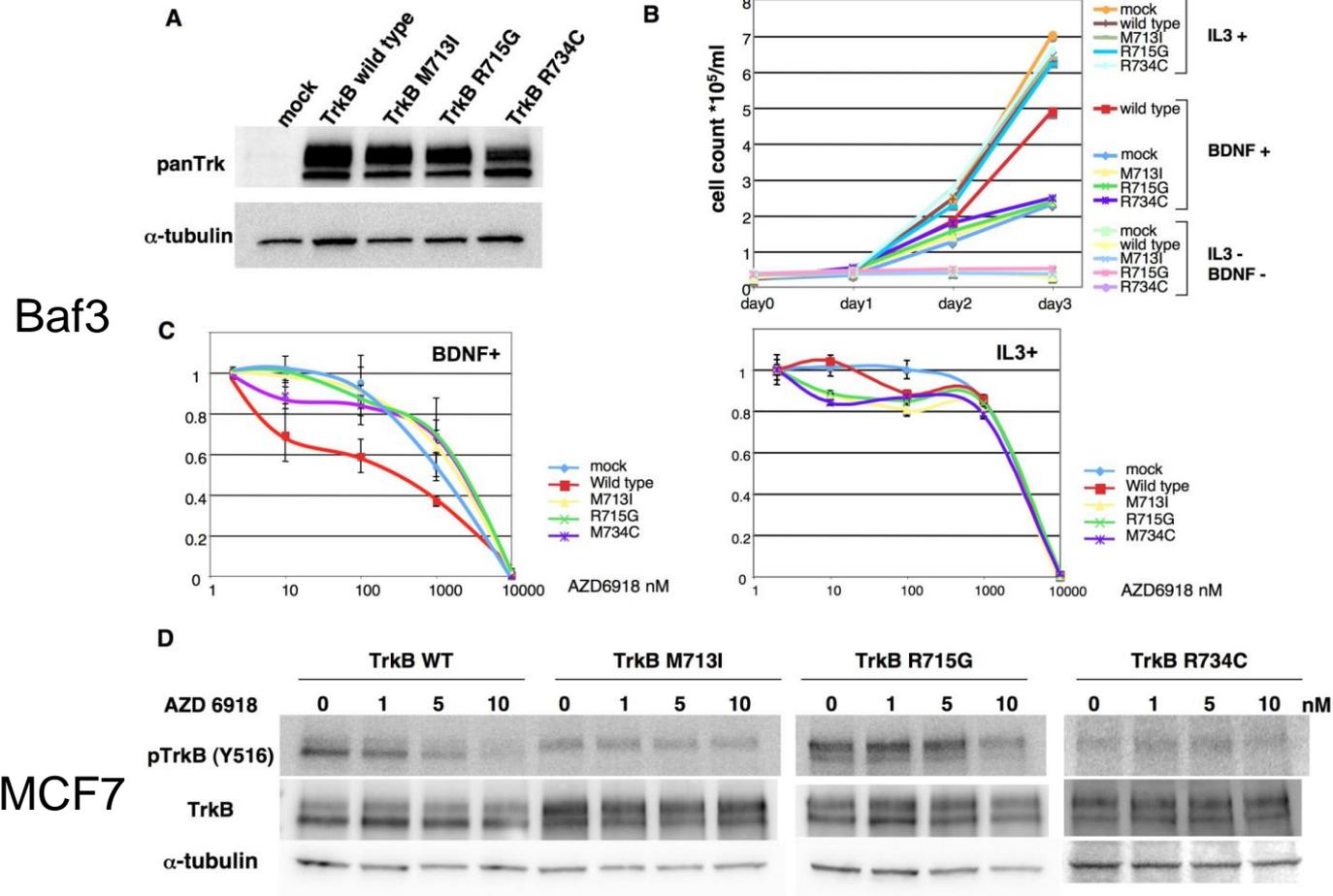
NIH3T3/TrkB wild type



**B**



# Functional characterization of TrkB



# Summary of TRKB mutations in lung cancer

	Histology	Pt.No.	mutations	country	
<b>This study</b>	ADC	17	ND	Asia	
	LCNEC	28	ND	(Japan)	
	SCLC	10	ND		
	ADC	12	ND	Europe	
	SCC	6	ND	(Netherland)	
	LCC	5	ND		
	<b>Cell lines</b>				
	ADC	17	ND		
	SCC	1	ND		
	LCC	3	ND		
SCLC	8	ND			
TC	1	ND			
<b>Marchetti et al.</b>	ADC	228	ND	Europe	
	SCC	184	ND	(Italy)	
	LCC	31	ND		
	LCNEC	29	4(13.8%)		
	SCLC	39	ND		
	TC	17	ND		
	AC	10	ND		
<b>Ding et al.</b>	ADC	188	6(3.2%)	USA	
<b>total</b>	NSCLC(without cell lines)	728	10(1.39%)		
	SCLC(without cell lines)	49	ND		

ADC=adenocarcinoma; LCC=large cell carcinoma; LCNEC=large cell neuroendocrine carcinoma; SCC=squamous cell carcinoma; SCLC=small cell carcinoma; TC=typical carcinoid; AC=atypical carcinoid; ND=not detected; NSCLC=non-small cell lung cancer

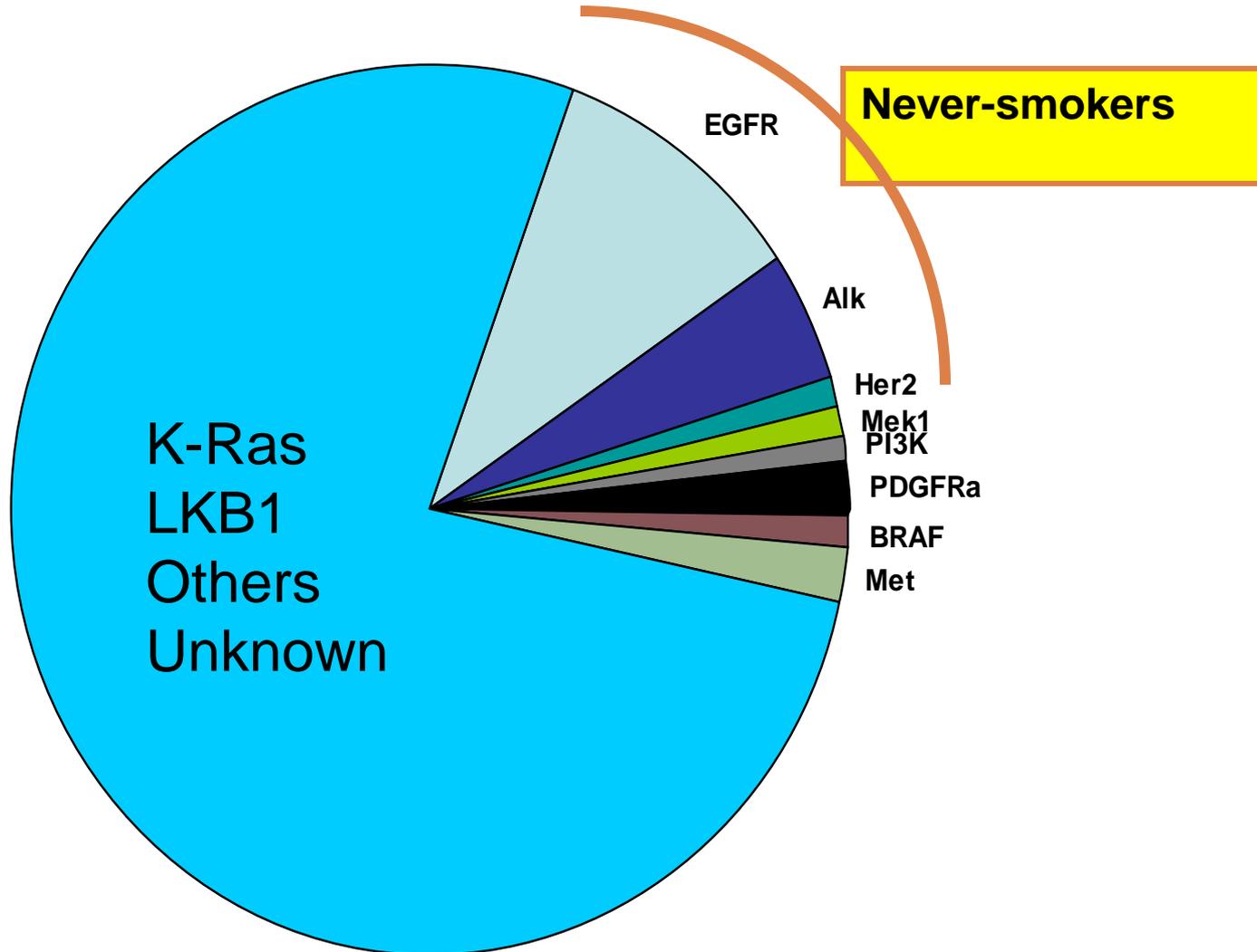
# “Oncogene addiction” may explain clinical responses to many kinase-targeted therapeutics

<u>Tumor type</u>	<u>Mutated/amplified kinase</u>	<u>Drug</u>
NSCLC	EGFR	gefitinib/erlotinib
NSCLC	Alk	PF-02341066
CML	BCR-ABL	imatinib/dasatanib
GIST	c-Kit/PDGFR	imatinib/sunitinib
CMML	PDGFR-b	imatinib
Breast cancer	Her2	trastuzumab/lapatanib
Gastric cancer	c-MET	PHA-665752*
Gastric cancer	FGFR2	AZD2171*
Melanoma	B-Raf	PLX4032
Melanoma	Her4	lapatinib*
Basal cell carcinoma	PTCH1/SMO	GDC-0449

\* Pre-clinical data only

“Addiction” to mutationally activated kinases can be faithfully modeled in cell culture.

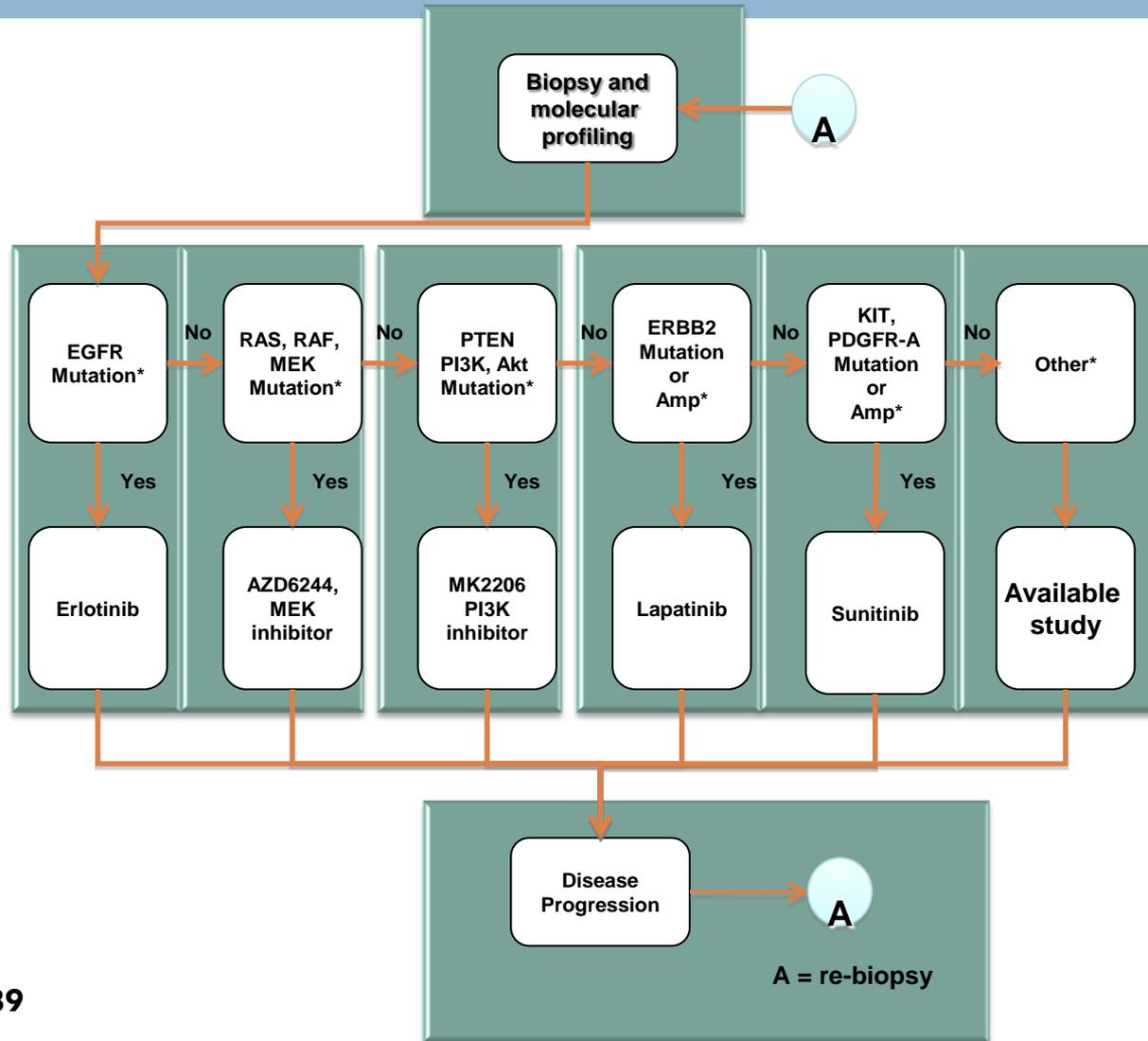
# “Drugging” the genome in NSCLC



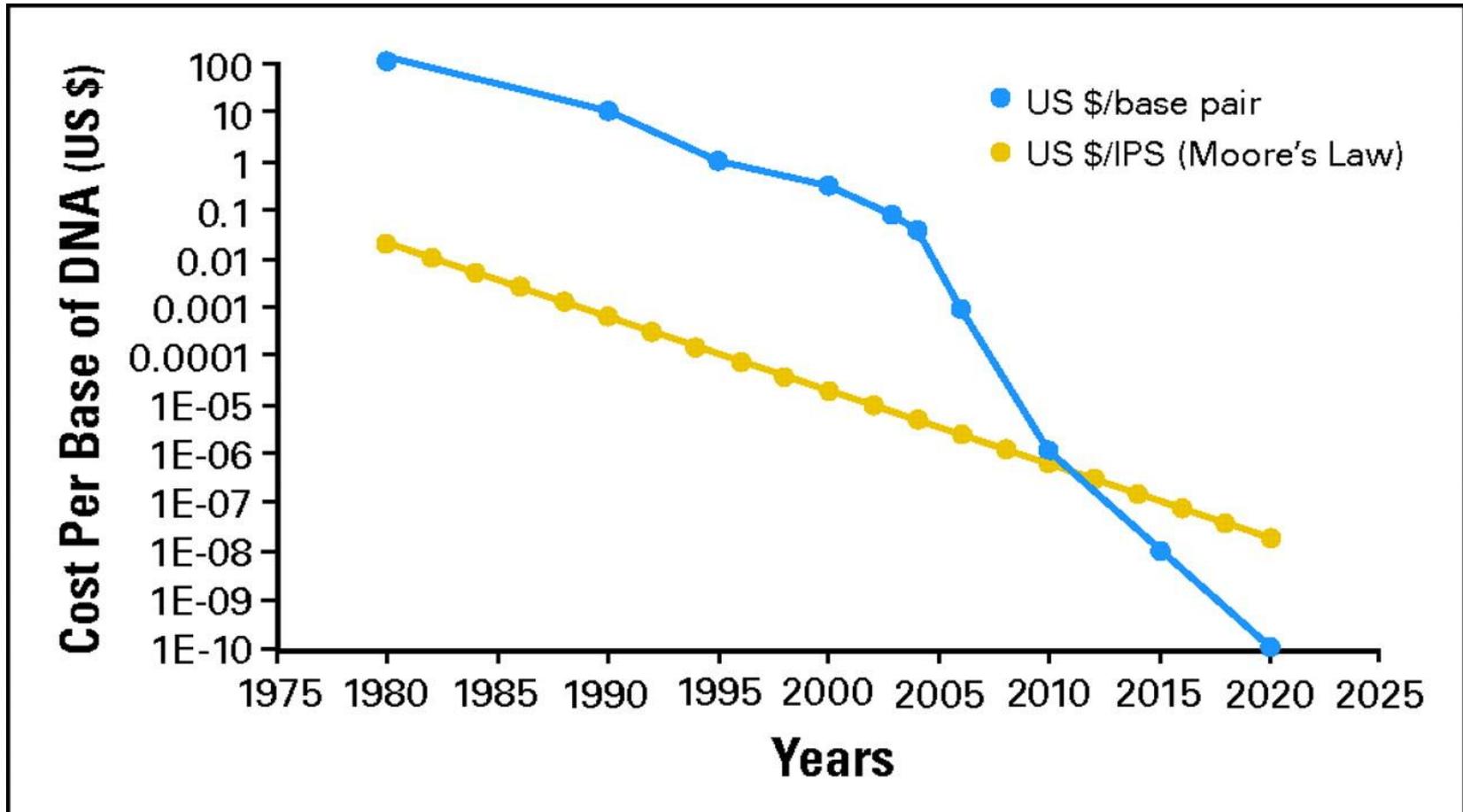
# Molecularly Targeted Treatment of Advanced Thoracic Malignancies

**Patient selection:**  
Molecular +  
NSCLC, SCLC and  
thymic malignancies

**Treatment:**  
Targeted therapies



Advances in massively parallel technologies have dramatically reduced the cost of sequencing.



MacConaill L E , Garraway L A JCO 2010;28:5219-5228





# Acknowledgments

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# The Changing Landscape of Lung Cancer and its Treatment

