

Ruolo del K-ras come fattore predittivo di Risposta ad anticorpi anti-EGFR nel carcinoma del colon retto

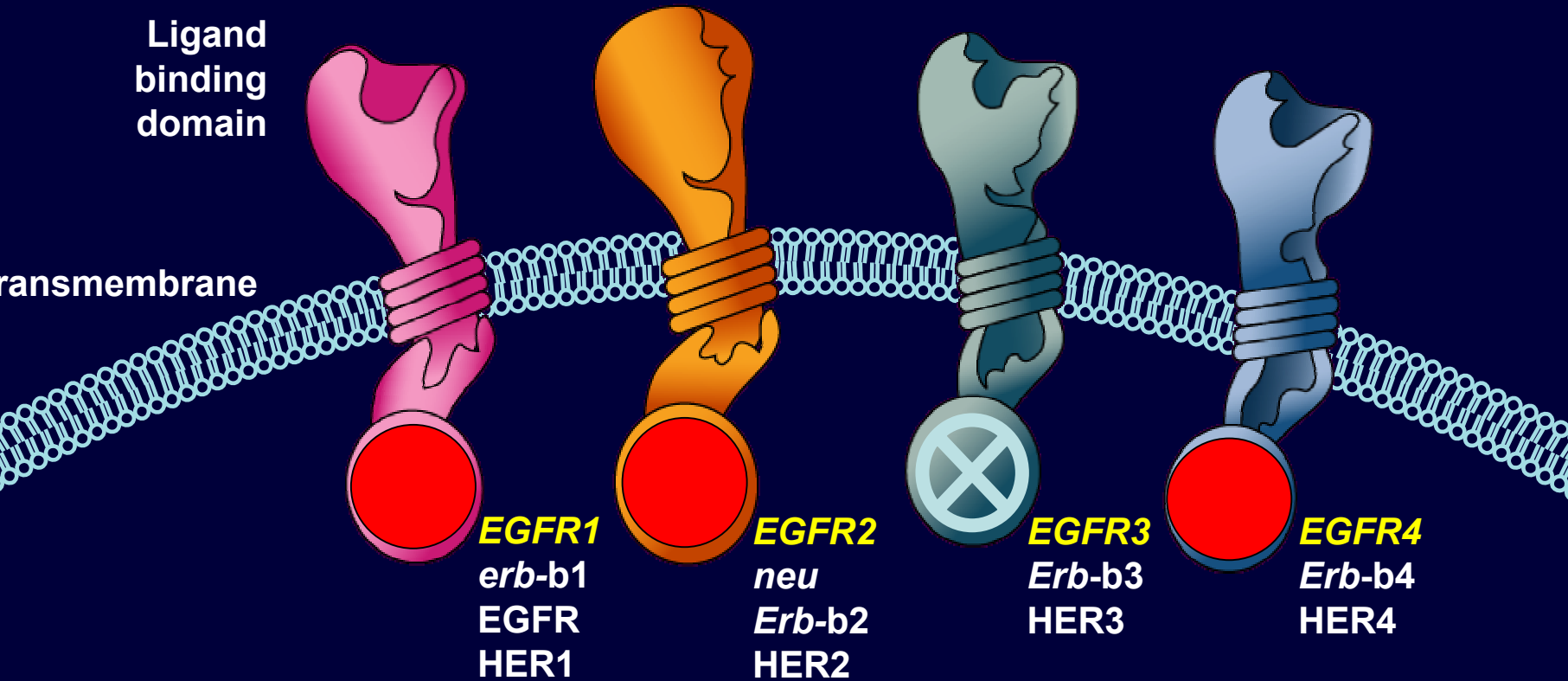
Antonio Marchetti

Pathology and Oncogenetic Unit,

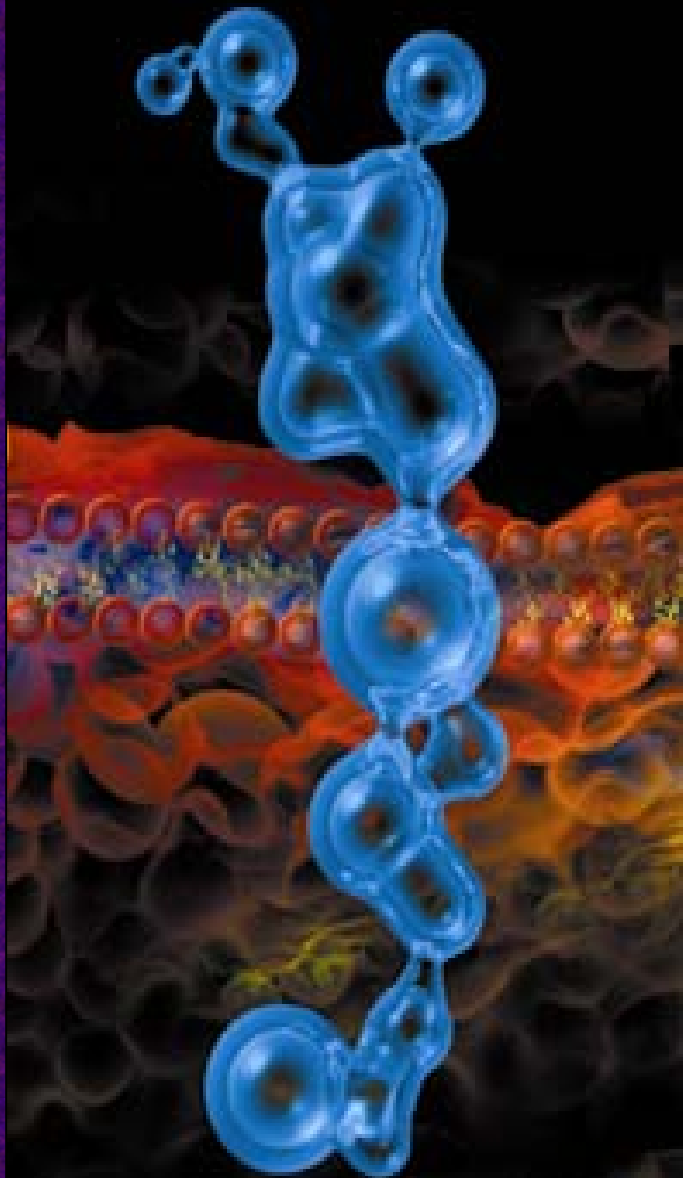
Center of Excellence on Aging

University-Foundation "G D'Annunzio", Chieti

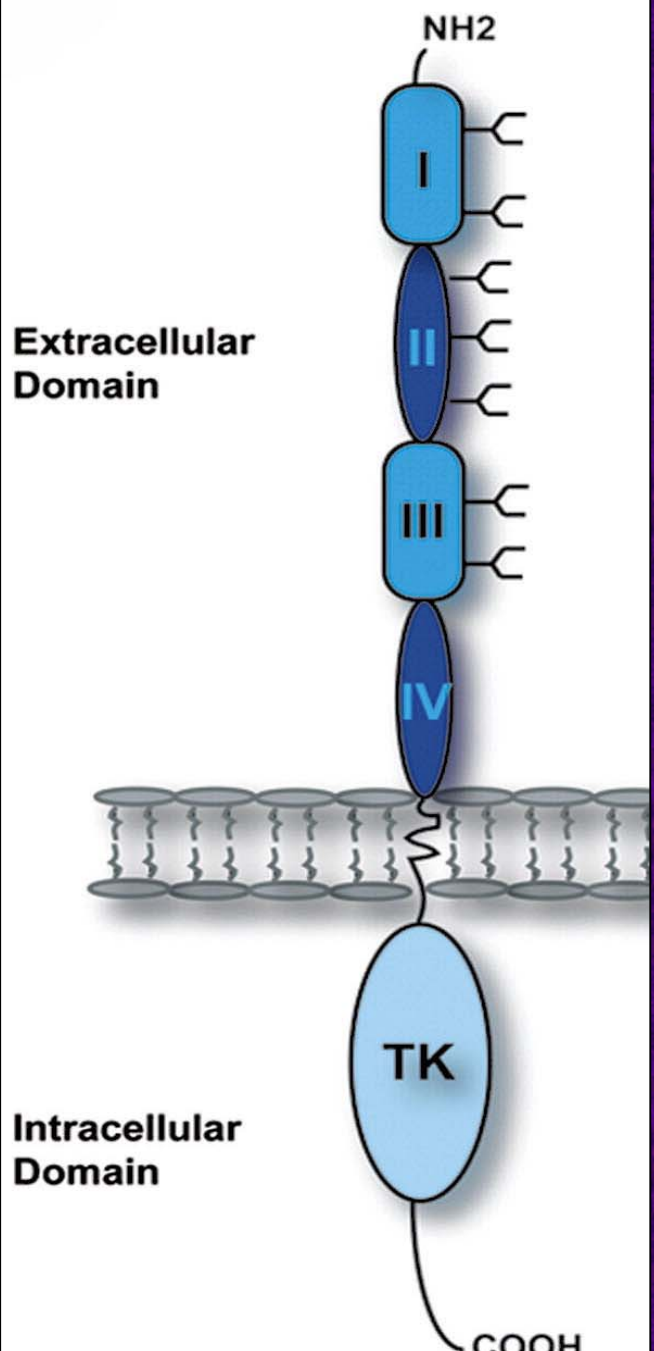
The EGFR/HER Family

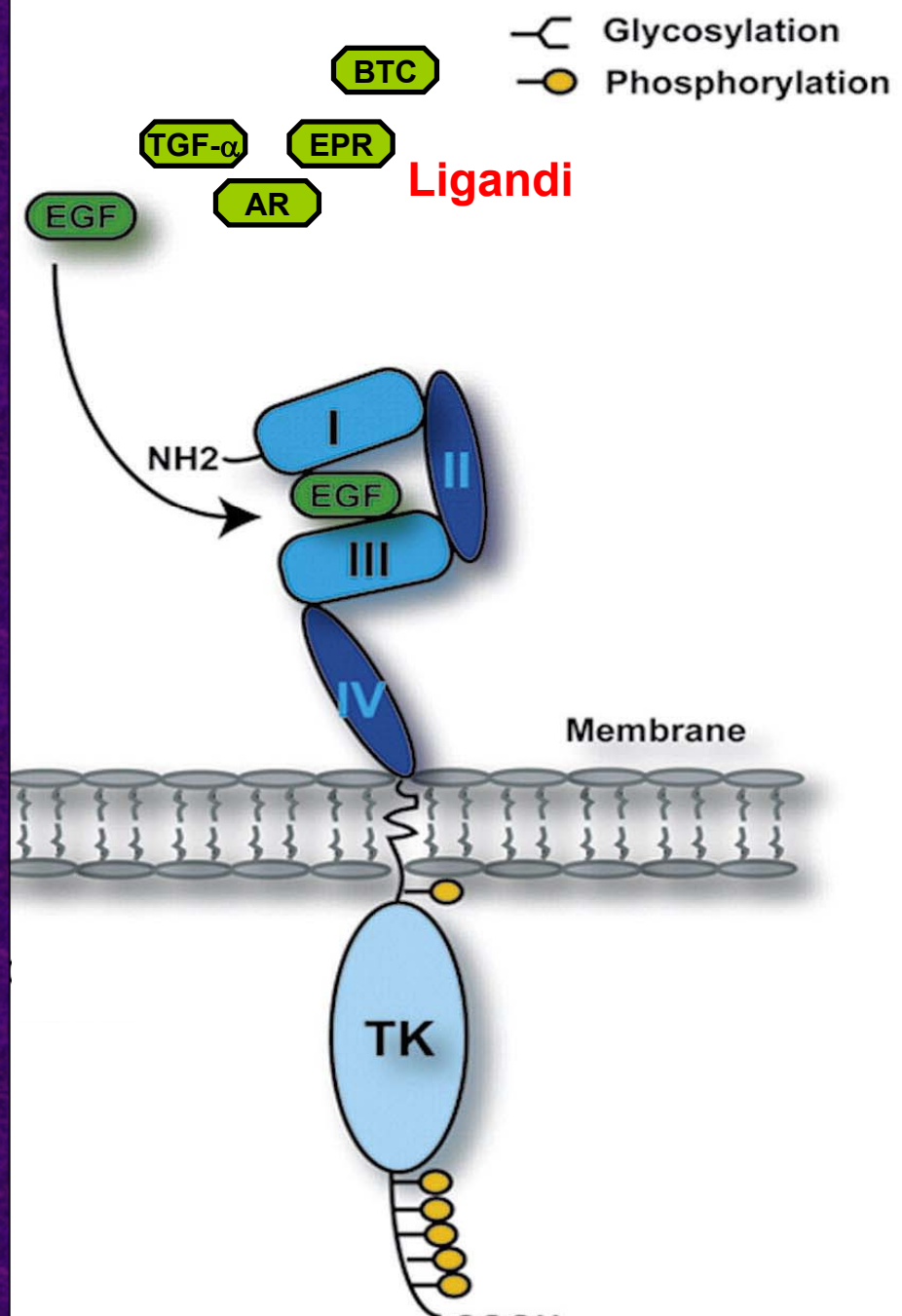
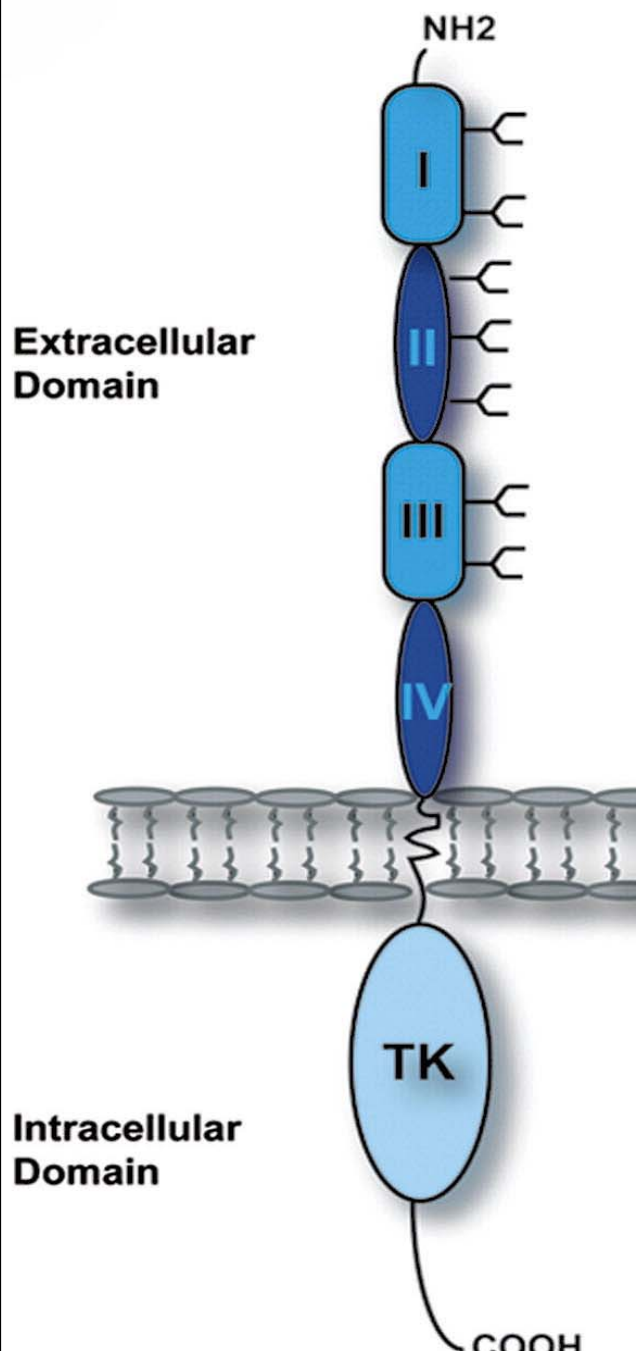


 = Tyrosine Kinase Domain

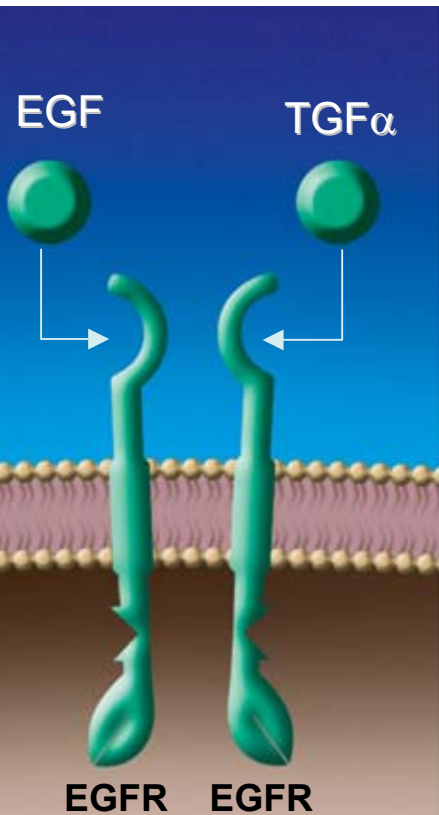


EGFR-1

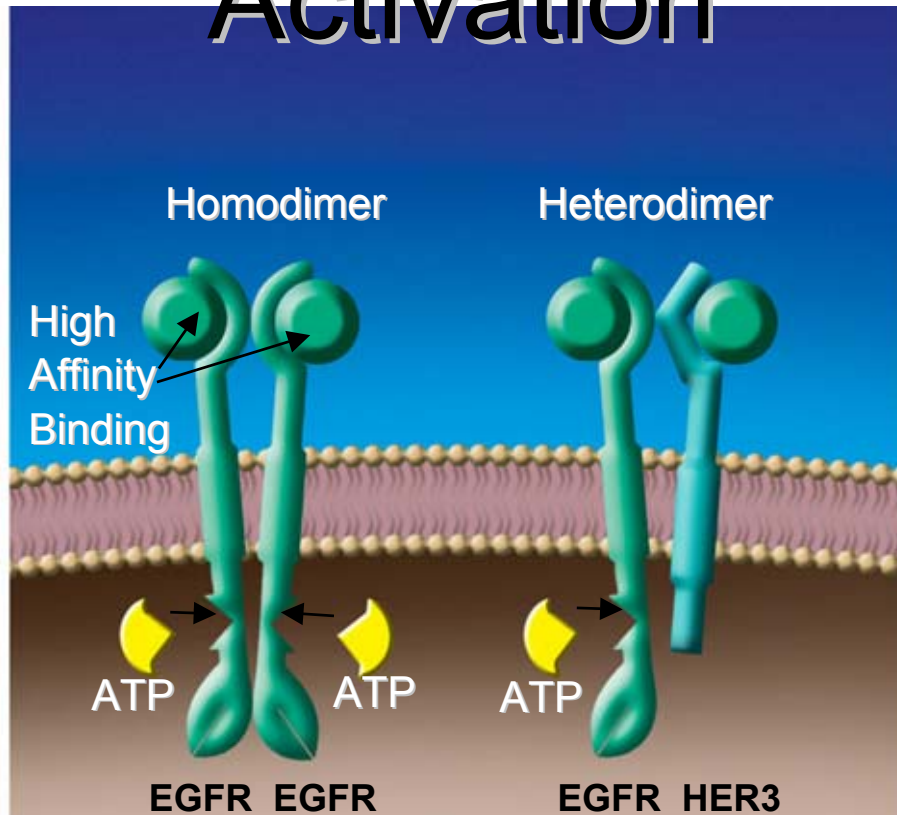




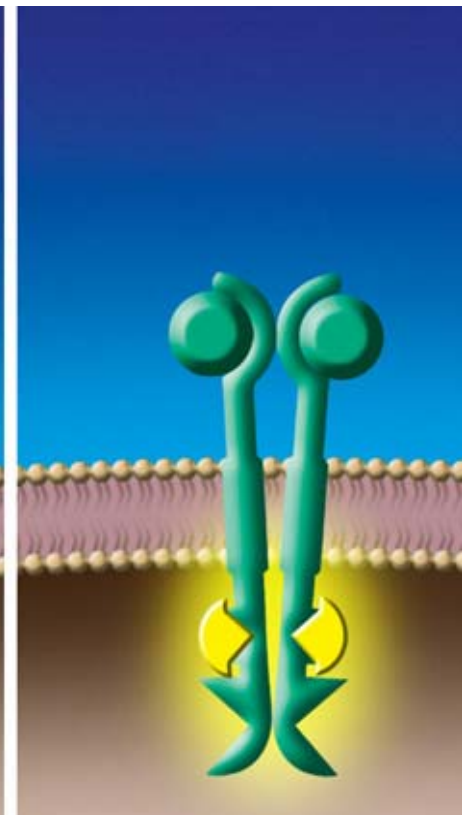
Ligand Binding and Dimerization Result in TK Activation



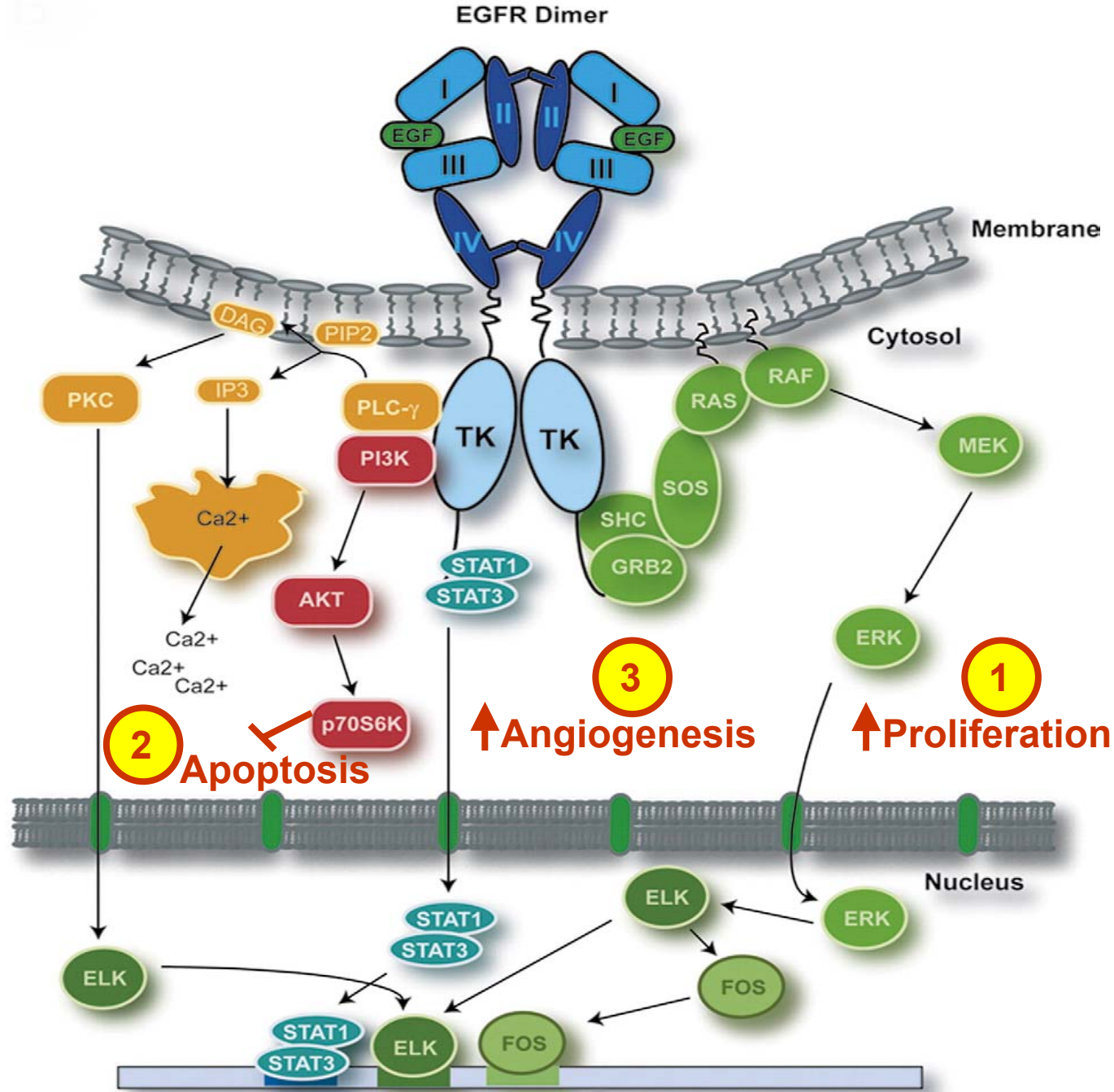
Ligand Binding



Dimerization



Phosphorylation and Activation



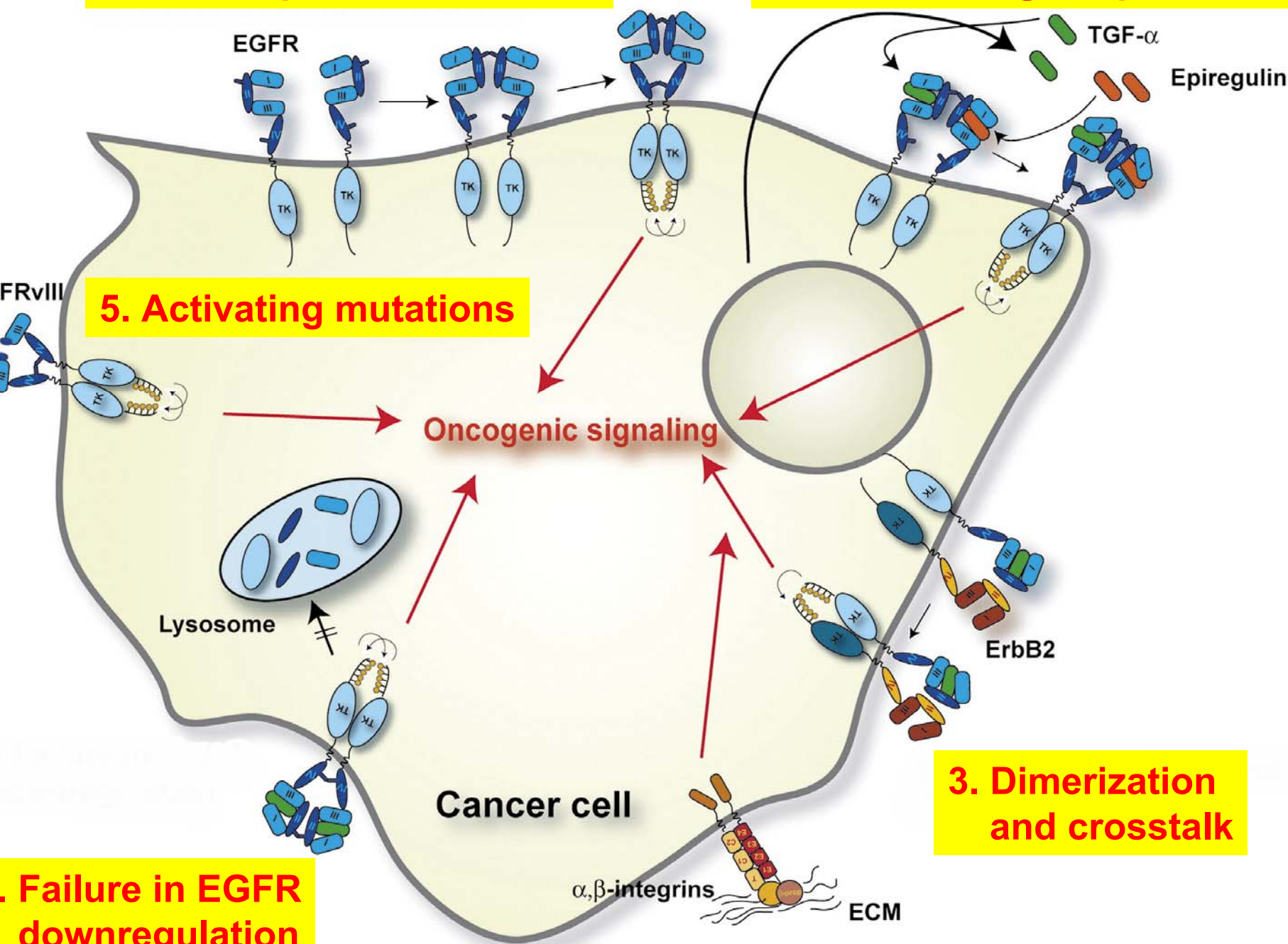
1. Overexpression of EGFR

2. Autocrine ligand production

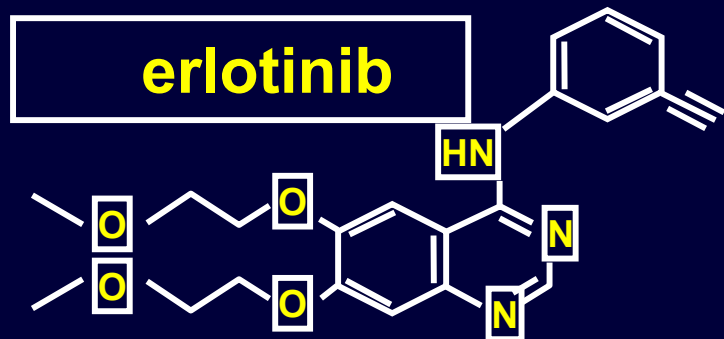
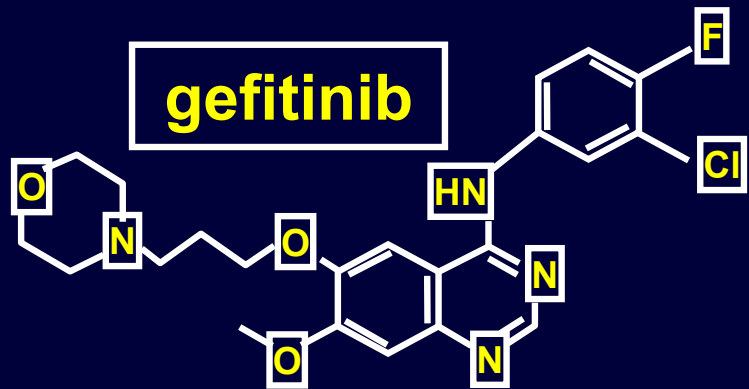
5. Activating mutations

3. Dimerization and crosstalk

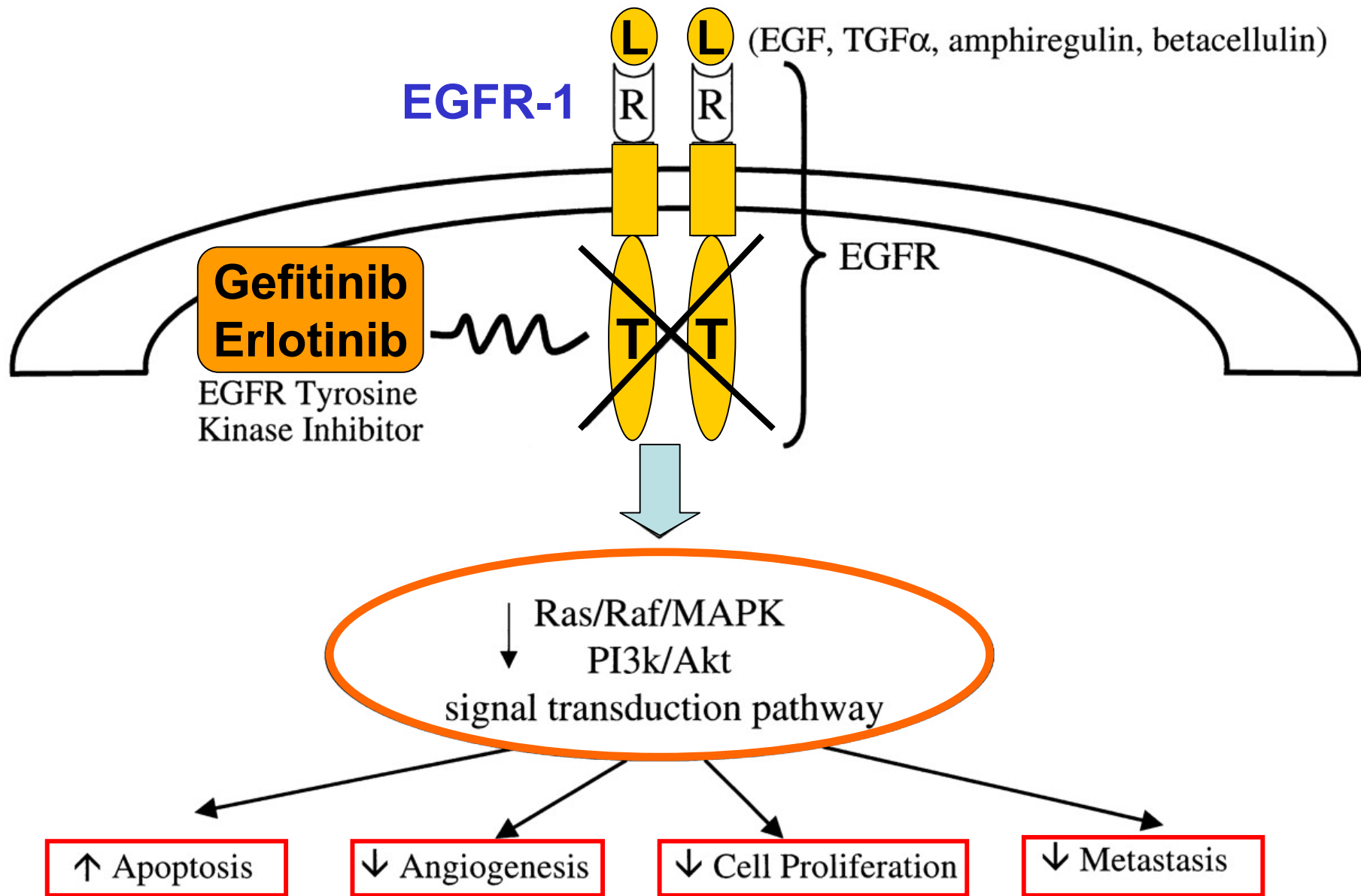
Failure in EGFR downregulation



EGFR Selective Small Molecule Tyrosine Kinase Inhibitors



- The tyrosine kinase activity requires ATP
- gefitinib and erlotinib compete for ATP binding
- Orally bioavailable small molecules



... questi inibitori del TKD, a differenza della chemioterapia classica, sono farmaci mirati ed hanno pertanto minimi effetti collaterali ...

... si è cominciato ad utilizzare questi farmaci, a scopo compassionevole in varie forme neoplastiche in fase avanzata, incluso il ca. polmonare ...

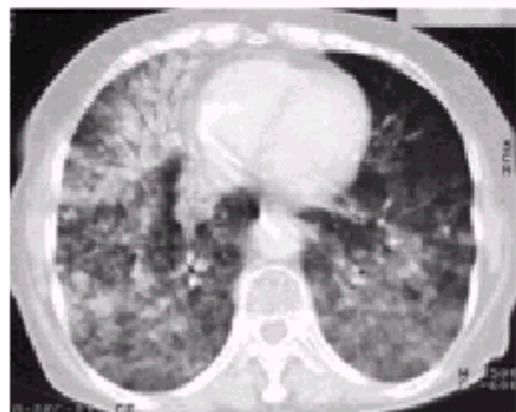
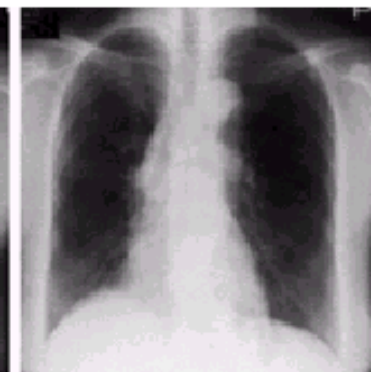
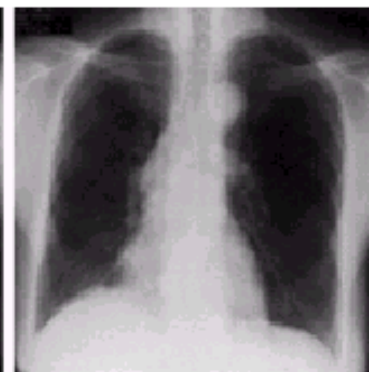
Before treatment

Day 3

Day 7

Day 14

Day 127



Before treatment

Day 28

Day 127

Phase III studies

ISEL (gefitinib)

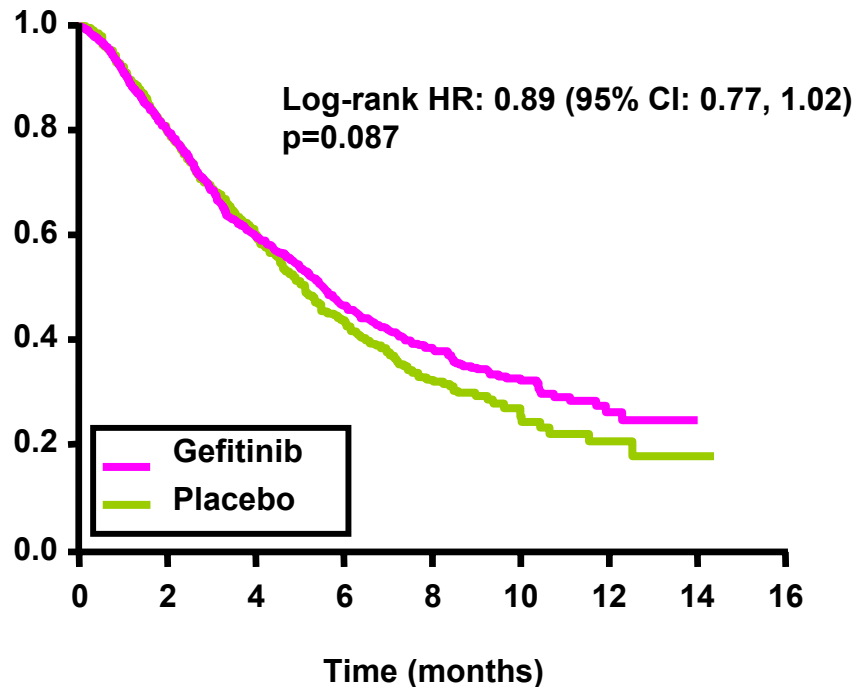
Previous treatment: platinum

Cases: 1129 gefitinib, 563 placebo

Response rate: Gefitinib: 8%, Placebo 1%.

Ov.survival: Gefitinib: 27 months, Placebo 21.

ISEL



BR-21 (erlotinib)

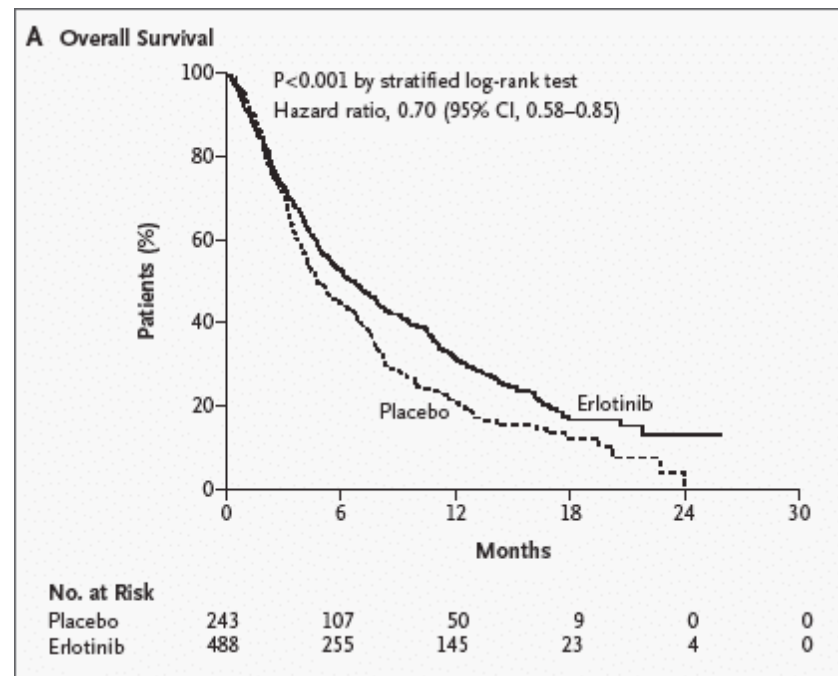
Previous treatment: platinum

Cases: 488 Erlotinib, 243 placebo

Response rate: Gefitinib: 9%, Placebo 1%.

Ov.survival: Erlotinib: 31 months, Placebo 22.

BR.21



Response in subgroup of patients



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ESTABLISHED IN 1812

MAY 20, 2004

VOL. 350 NO. 21

Activating Mutations in the Epidermal Growth Factor Receptor Underlying Responsiveness of Non–Small-Cell Lung Cancer to Gefitinib

Thomas J. Lynch, M.D., Daphne W. Bell, Ph.D., Raffaella Sordella, Ph.D., Sarada Gurubhagavatula, M.D.,
Ross A. Okimoto, B.S., Brian W. Brannigan, B.A., Patricia L. Harris, M.S., Sara M. Haserlat, B.A.,
Jeffrey G. Supko, Ph.D., Frank G. Haluska, M.D., Ph.D., David N. Louis, M.D., David C. Christiani, M.D.,
Jeff Settleman, Ph.D., and Daniel A. Haber, M.D., Ph.D.

SCIENCE VOL 304, 4 June 2004

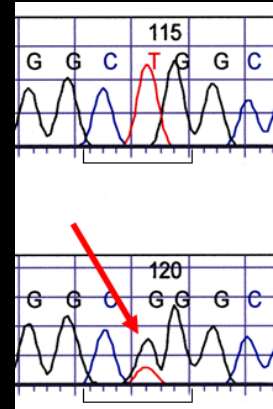
EGFR Mutations in Lung Cancer: Correlation with Clinical Response to Gefitinib Therapy

J. Guillermo Paez,^{1,2*} Pasi A. Jänne,^{1,2*} Jeffrey C. Lee,^{1,3*}
Sean Tracy,¹ Heidi Greulich,^{1,2} Stacey Gabriel,⁴ Paula Herman,¹
Frederic J. Kaye,⁵ Neal Lindeman,⁶ Titus J. Boggon,^{1,3}
Katsuhiko Naoki,¹ Hidefumi Sasaki,⁷ Yoshitaka Fujii,⁷
Michael J. Eck,^{1,3} William R. Sellers,^{1,2,4†}
Bruce E. Johnson,^{1,2†} Matthew Meyerson^{1,3,4†}

Pathology of EGFR Mutants



Mutations associated with drug sensitivity



L858R



- G719C
- G719S
- G719A
- V689M
- N700D
- E709K/Q
- S720P

(5%)

- ΔE746-A750
- ΔE746-T751
- ΔE746-A750 (ins RP)
- ΔE746-T751 (ins A/I)
- ΔE746-T751 (ins VA)
- ΔE746-S752 (ins A/V)
- ΔL747-E749 (A750P)
- ΔL747-A750 (ins P)
- ΔL747-T751
- ΔL747-T751 (ins P/S)
- ΔL747-S752
- ΔL747-752 (E746V)
- ΔL747-752 (P753S)
- ΔL747-S752 (ins Q)
- ΔL747-P753
- ΔL747-P753 (ins S)
- ΔS752-I759

48%

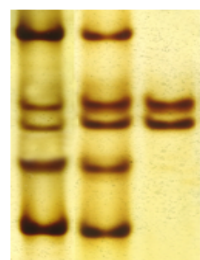
- V765A
- T783A

(<1%)

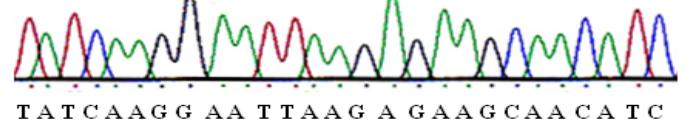
- L858R (40-45%)
- N826S
- A839T
- K846R
- L861Q
- G863D

43%

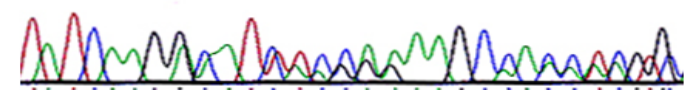
244T 89T 244N



244N



244T



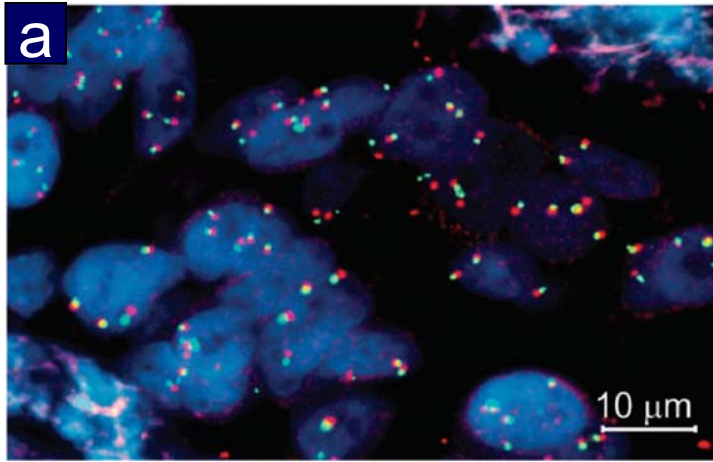
Ex 19 Deletion

Del-2239-2247 (TTAAGAGAA)

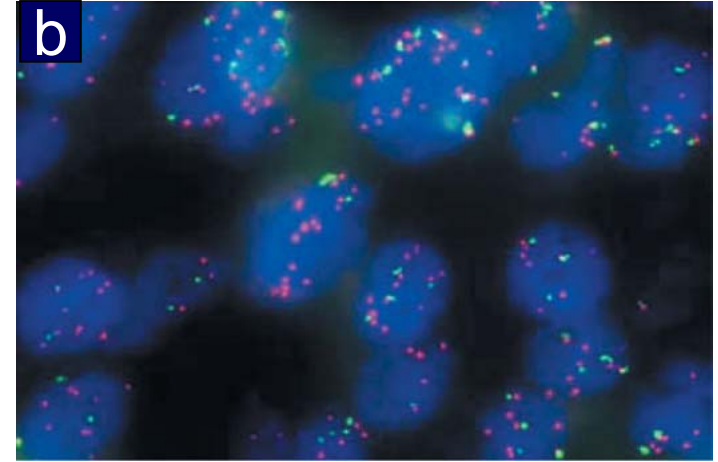
B

EGFR gene copy number

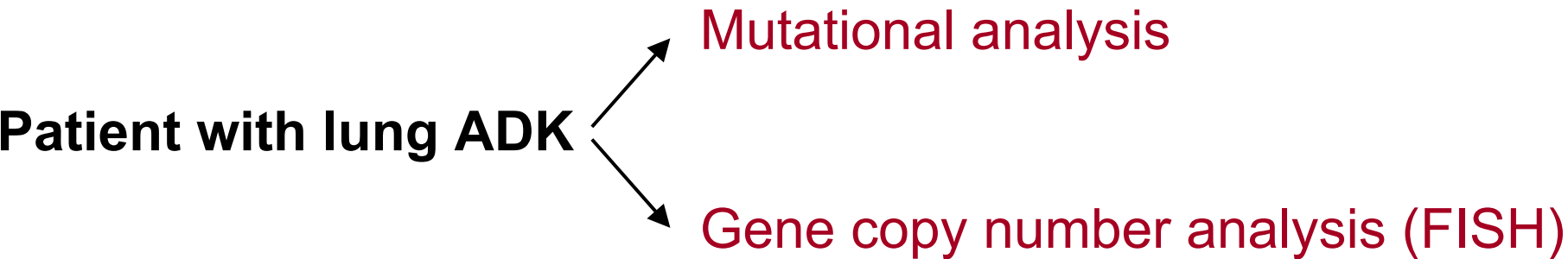
FISH



High polysomy



EGFR1 gene amplification



KRAS Mutations and Primary Resistance of Lung Adenocarcinomas to Gefitinib or Erlotinib

William Pao^{1,2*}, Theresa Y. Wang¹, Gregory J. Riely², Vincent A. Miller², Qiulu Pan³, Marc Ladanyi³, Lauren F. Zakowski³, Robert T. Heelan⁴, Mark G. Kris², Harold E. Varmus¹

¹ Program in Cancer Biology and Genetics, Memorial Sloan-Kettering Cancer Center, New York, New York, United States of America, ² Department of Medicine, Memorial Sloan-Kettering Cancer Center, New York, New York, United States of America, ³ Department of Pathology, Memorial Sloan-Kettering Cancer Center, New York, New York, United States of America, ⁴ Department of Radiology, Memorial Sloan-Kettering Cancer Center, New York, New York, United States of America

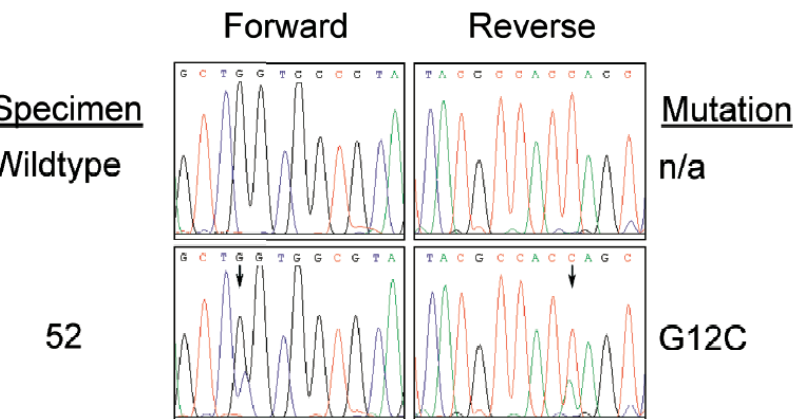


Table 1. EGFR and KRAS Mutation Status in Lung Adenocarcinomas Sensitive or Refractory to Gefitinib or Erlotinib

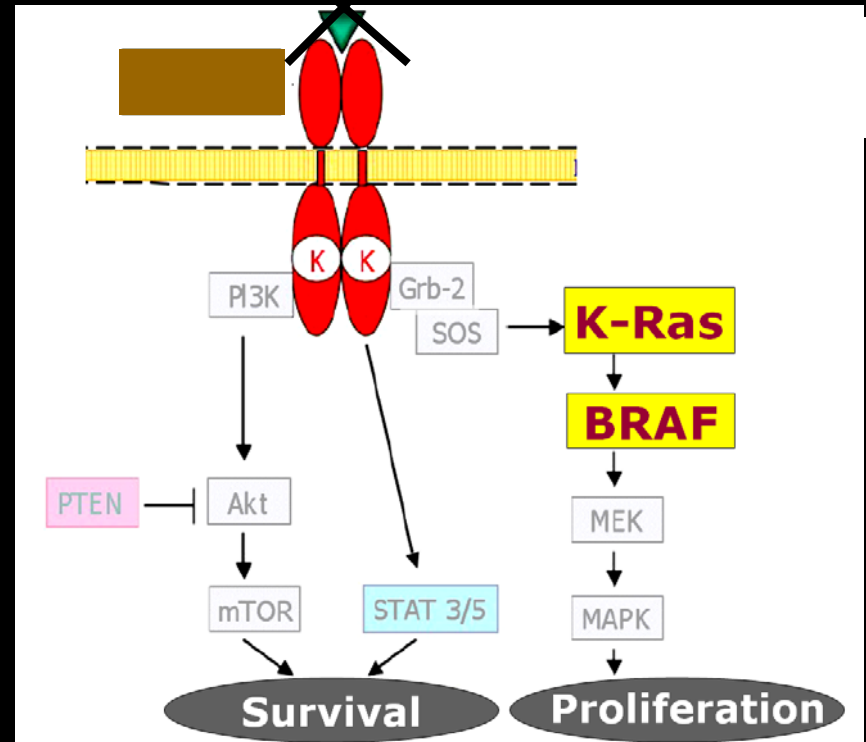
Drug	Gene Mutated	Proportion Sensitive	Proportion Refractory	p-Value
Gefitinib	EGFR	9/12	0/12	0.0034
	KRAS	0/12	5/12	0.0373
Erlotinib	EGFR	8/10	0/26	1.487×10^{-6}
	KRAS	0/9 ^a	4/26	0.5531 ^b
Gefitinib or Erlotinib	EGFR	17/22	0/38	6.801×10^{-11}
	KRAS	0/21 ^a	9/38	0.0201

Intrinsic resistance to TKI

K-ras mutations

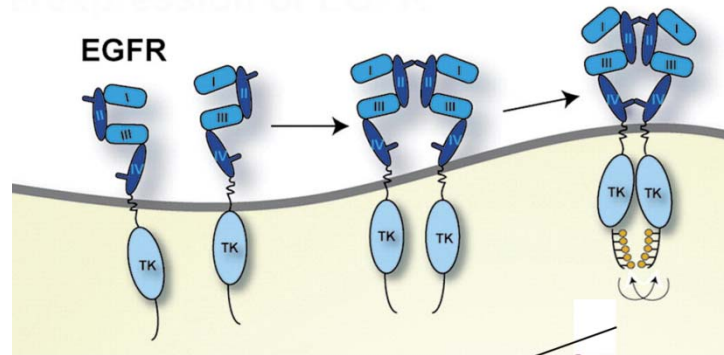


Resistance to TKI



Epidermal Growth Factor Receptor (EGFR) in Colorectal Cancer

Overexpression of EGFR

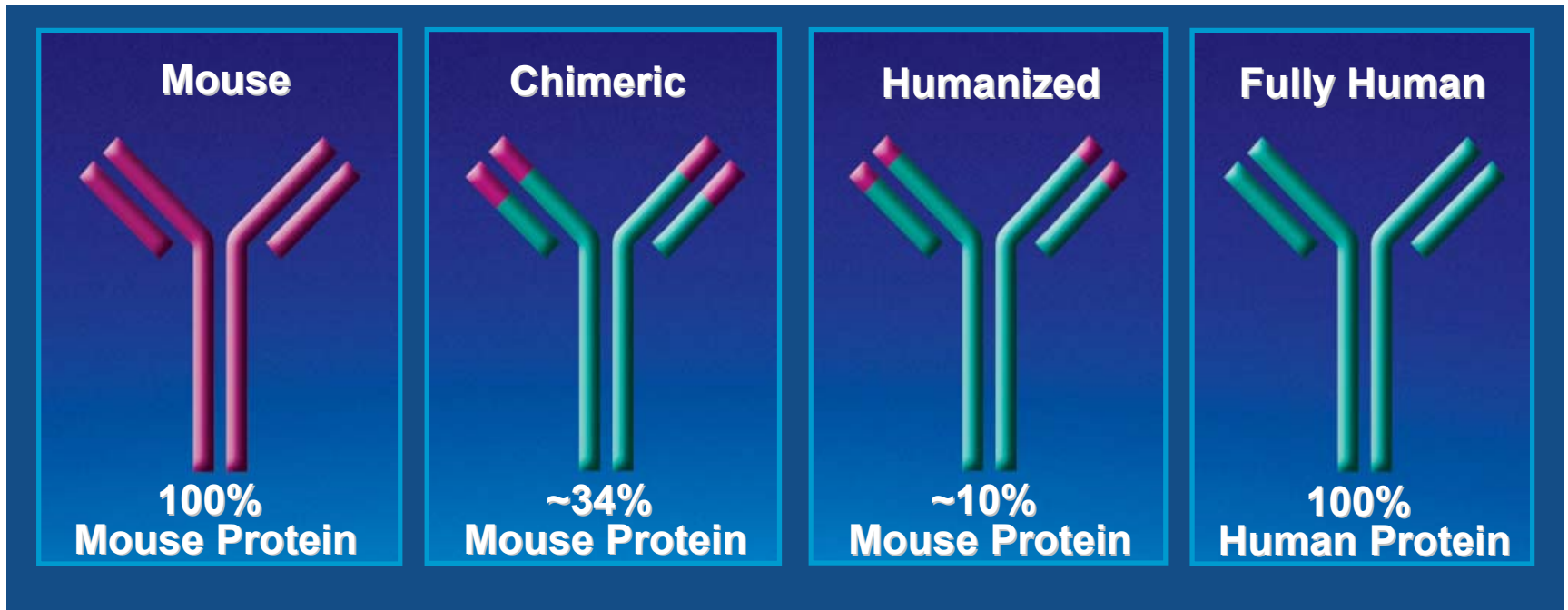


Oncogenic signalling

In vitro and in vivo data indicate that:

- preventing the binding of ligands to EGFR results in inhibition of tumor cell growth
- monoclonal antibodies can inhibit the binding of ligands

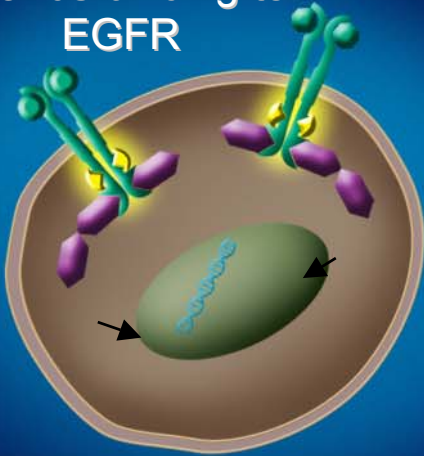
The Development of Human Monoclonal Antibodies



panitumumab

Panitumumab Inhibits Ligand Binding to EGFR and Dimerization

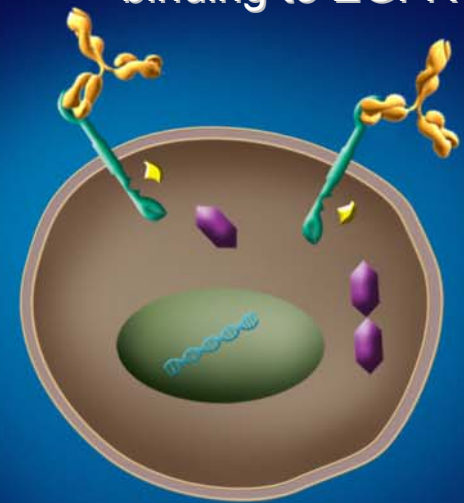
EGF, TGF α or other ligands binding to EGFR



Panitumumab



Inhibition of EGF and TGF α binding to EGFR



- A fully human* IgG2 monoclonal antibody to EGFR
- High affinity, $K_D = 5 \times 10^{-11}$ M
- Inhibits ligand-induced EGFR tyrosine phosphorylation

This may lead to:
↓ Cell proliferation
↓ Cell survival
↓ Angiogenesis
↓ Metastatic spread

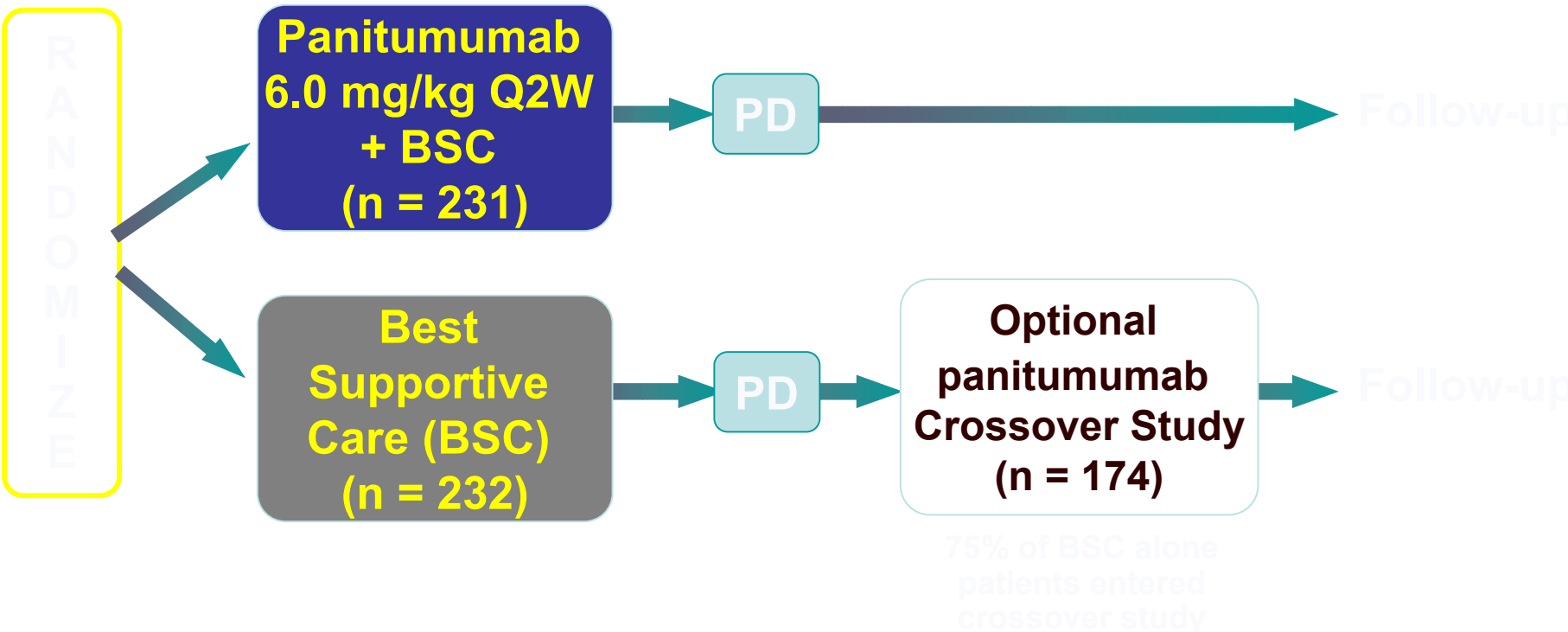
Panitumumab pharmacokinetics

- Panitumumab administered as a single agent or in combination with chemotherapy **exhibits nonlinear pharmacokinetics**
- Steady-state is obtained after **3 doses at 6 mg/kg** given **once every 2 weeks** without the need of a loading dose
- The **mean half-life** value during the dosing interval is **7.5 days** (range: 3.6 -10.9 days) for the **6 mg/kg dose**

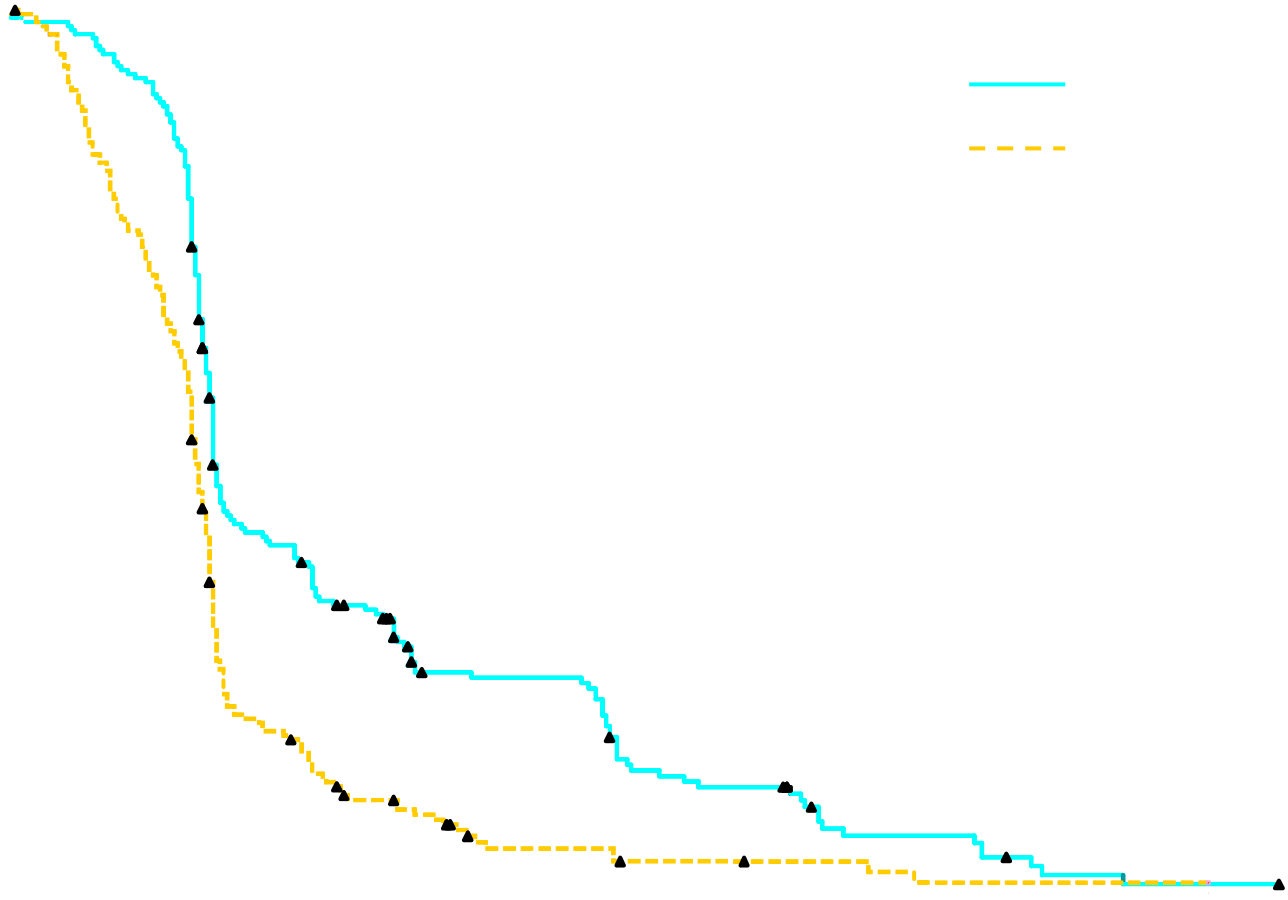
Based on preliminary data on outcomes in colorectal cancer, a large, randomized, phase 3 trial in EGFR-expressing tumors of patients with metastatic colorectal cancer resistant to 2 lines of chemotherapy was designed.

Study Design

Van Cutsem E et al JCO 2007; 25(13):1658-1664

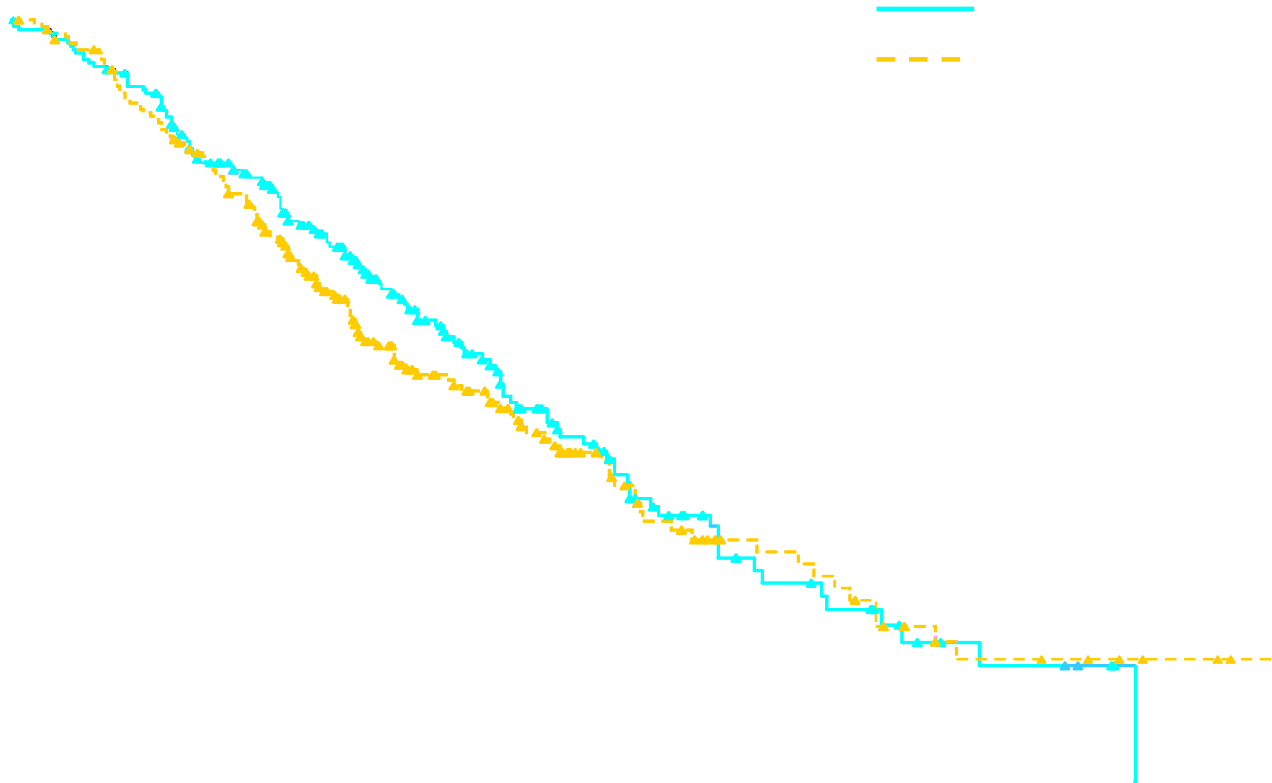


Progression-Free Survival



Overall Survival

(All Randomized Analysis Set)



Skin Toxicities Are the Most Common Adverse Events With Panitumumab



- **Dermatologic toxicities**

- **12%**

grade 3

89% of patients

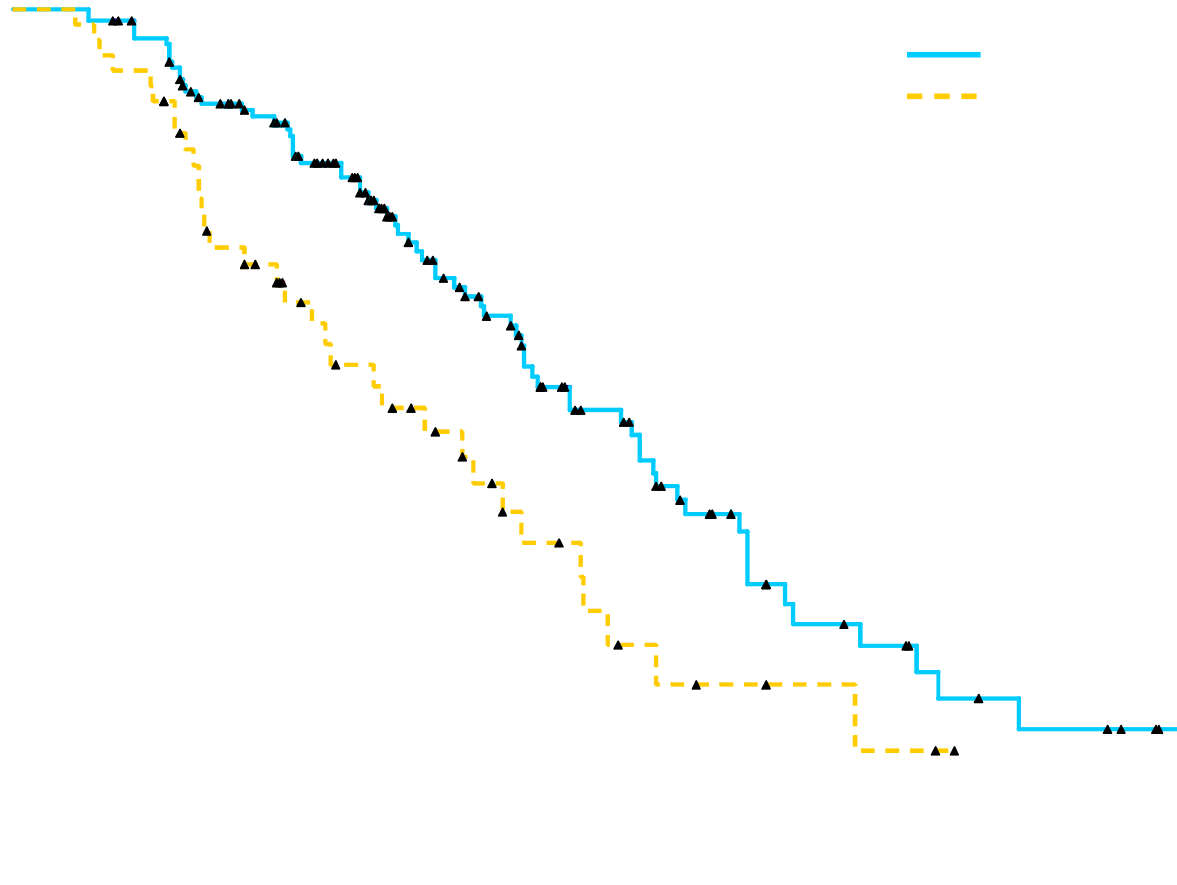
Skin Toxicity was the Most Commonly Reported Adverse Event

Panitumumab + BSC
(n = 229)

BSC alone
(n = 234)

Adverse Event	All grades	Grade 3-4	All grades	Grade 3-4
All Skin/Integument toxicity	90%	16%	9%	0%
Skin	90%	14%	6%	0%
Erythema	65%	5%	1%	0%
Acneiform dermatitis	57%	7%	1%	0%
Pruritus	57%	2%	2%	0%
Skin exfoliation	25%	2%	0%	0%
Rash	22%	1%	1%	0%
Skin fissures	20%	1%	<1%	0%
Dry skin	10%	0%	0%	0%
Acne	13%	1%	0%	0%
Nail				
Paronychia	25%	2%	0%	0%
Other nail disorder	9%	0%	0%	0%
Hair	9%	0%	1%	0%
Growth of eyelashes	6%	0%	0%	0%
Eye	15%	<1%	2%	0%

Skin Toxicity in the Panitumumab Patients



US FDA Label

‘Panitumumab has been approved for treatment of EGFR-expressing metastatic colorectal carcinoma in patients with disease progression or following chemotherapy containing fluoropyrimidine, oxaliplatin, and irinotecan’

In search for a predictive marker

- **CLINICAL MARKER:** Skin toxicity

- **BIOLOGICAL MARKERS:**

- EGFR mutations
- Immunohistochemical expression
- EGFR copy number
- K-ras mutations

EGFR mutations in CRC

N Engl J Med 2004;351:2883.

EGFR mutations: 1/293 (0,34%)

Clin Cancer Res 2005;11:1368–1371.

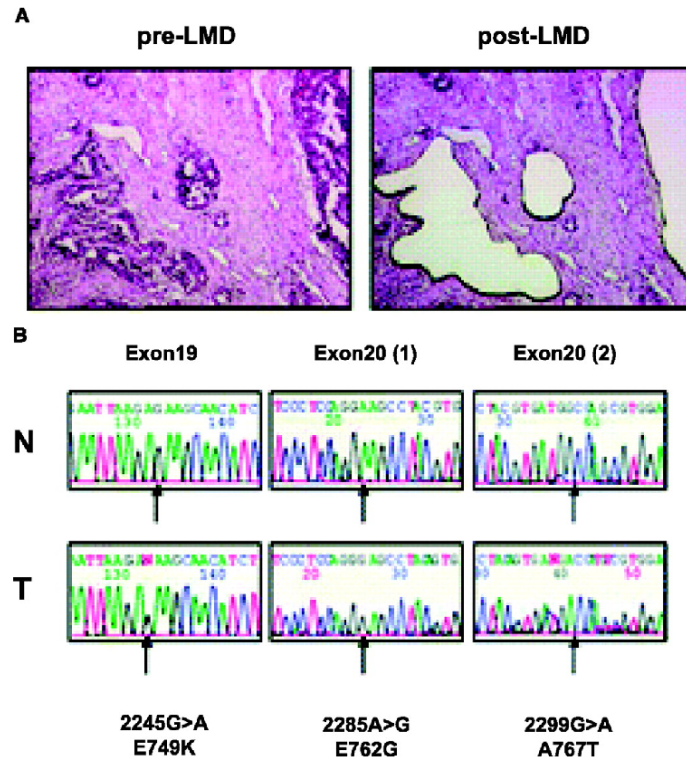
EGFR mutations: 4/33 (12%)

Somatic Mutations of Epidermal Growth Factor Receptor in Colorectal Carcinoma

Hisashi Nagahara,^{1,2} Koshi Mimori,¹ Mitsuhiro Ohta,¹ Tohru Utsunomiya,¹ Hiroshi Inoue,¹ Graham F. Barnard,³ Masaichi Ohira,² Kosei Hirakawa,² and Masaki Mori¹

1Department of Surgery, Medical Institute of Bioregulation, Kyushu University, Beppu, Japan; 2Department of Surgical Oncology, Osaka City University Graduate School of Medicine, Osaka, Japan; and 3Department of Medicine, University of Massachusetts, Worcester, Massachusetts

**Laser
microdissection**



EGFR mutations

4/33 (12%)

G-A transitions

Assessing *EGFR* Mutations

Adenocarcinoma

GGACTCTGGATCCCAGAAGGTGAGAAAGTTAAAATTCCCGTCGCTATCAAGGAATTAAGAGAAGCAACATCTCCGAAAGCCAACAAGGAAATCCTCGA

C

A

C

T

A

Normal Tissue (normal lymph node)

GGACTCTGGATCCCAGAAGGTGAGAAAGTTAAAATTCCCGTCGCTATCAAGGAATTAAGAGAAGCAACATCTCCGAAAGCCAACAAGGAAATCCTCGA

C

A

A

C

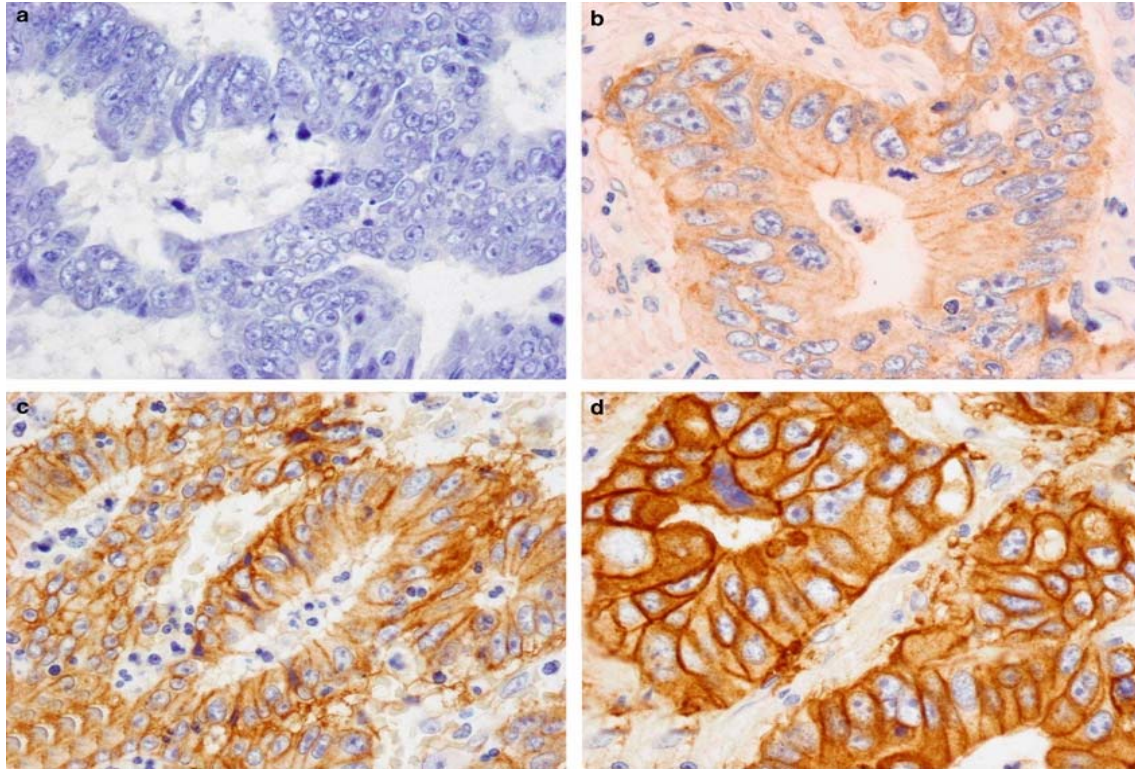
T

T

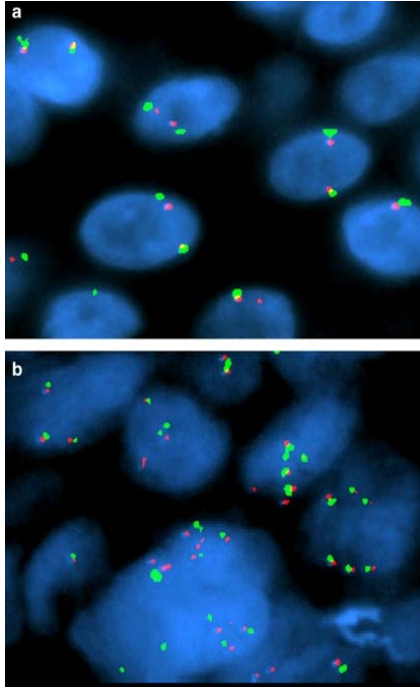
**Artifactual mutations
in normal tissues**

Antonio Marchetti, M.D.
Lara Felicioni, M.D.

EGFR IHC expression



EGFR copy number



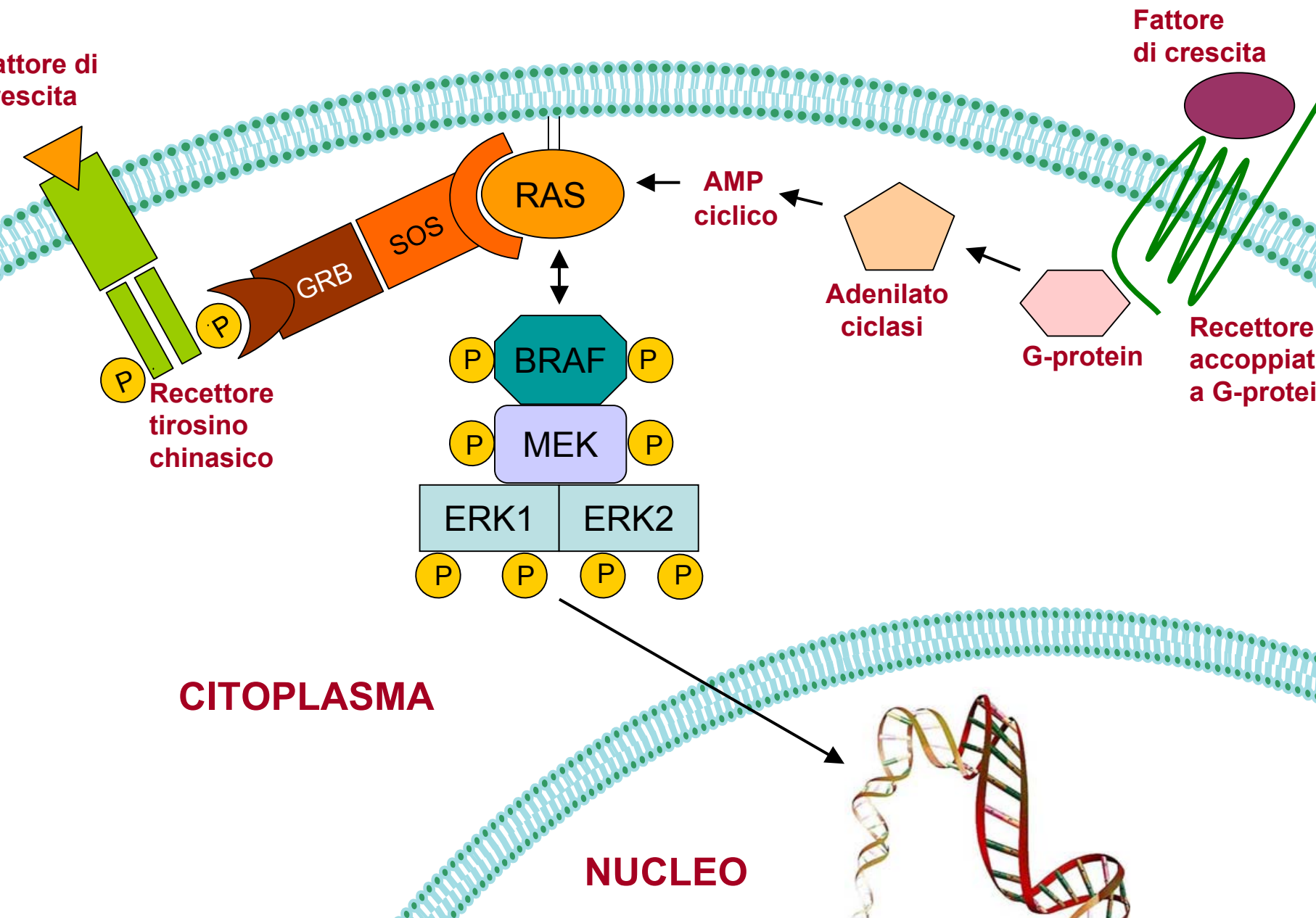
Moroni M et al. *Lancet Oncol* 6:279–286, 2005.
Italiano A et al. *Ann Surg Oncol* 15:649–654, 2008.

The Role of K-Ras in Patient Selection for Therapeutic EGFR Inhibitors

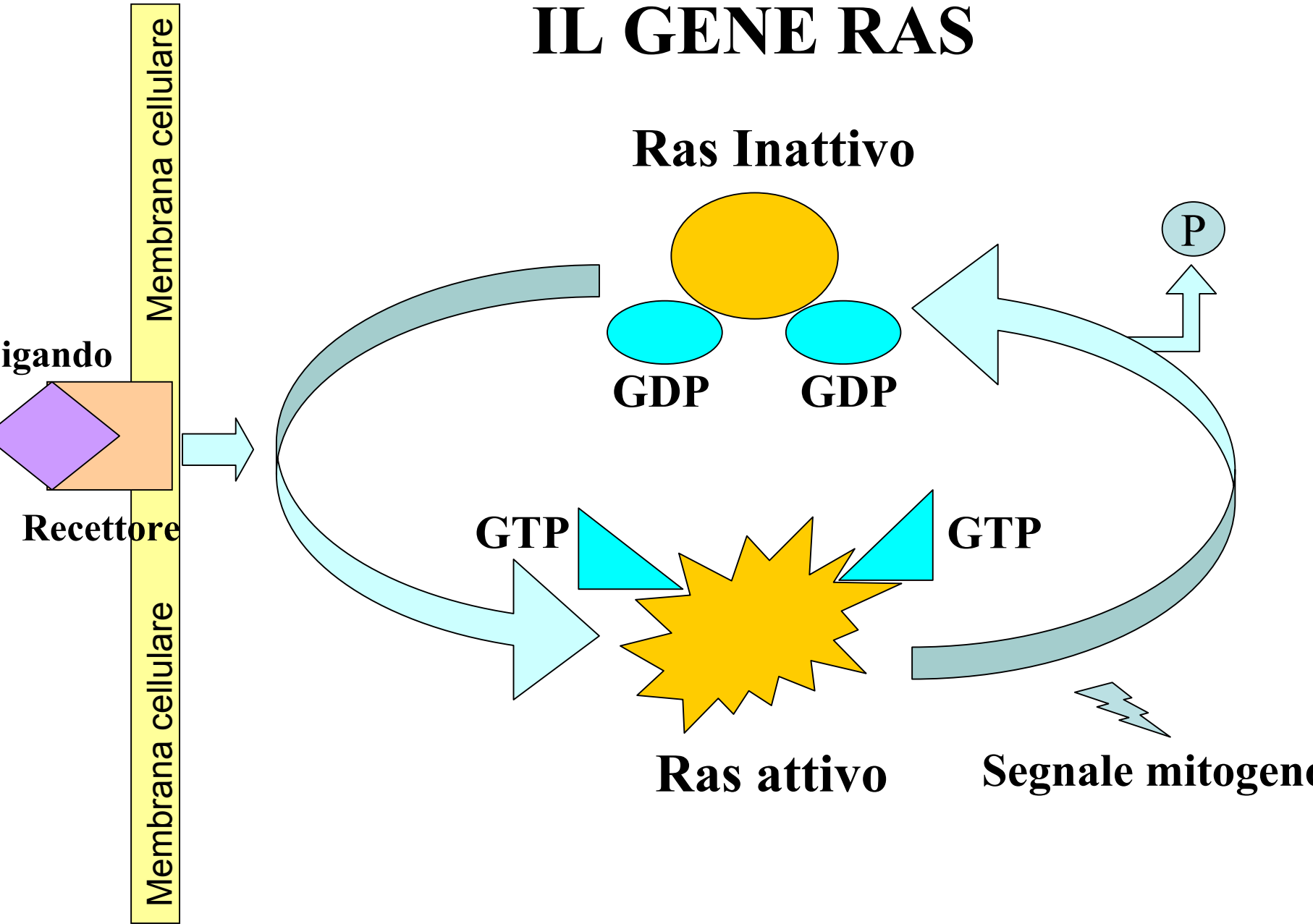
Geni della famiglia *RAS*

H-RAS *K-RAS* *N-RAS*

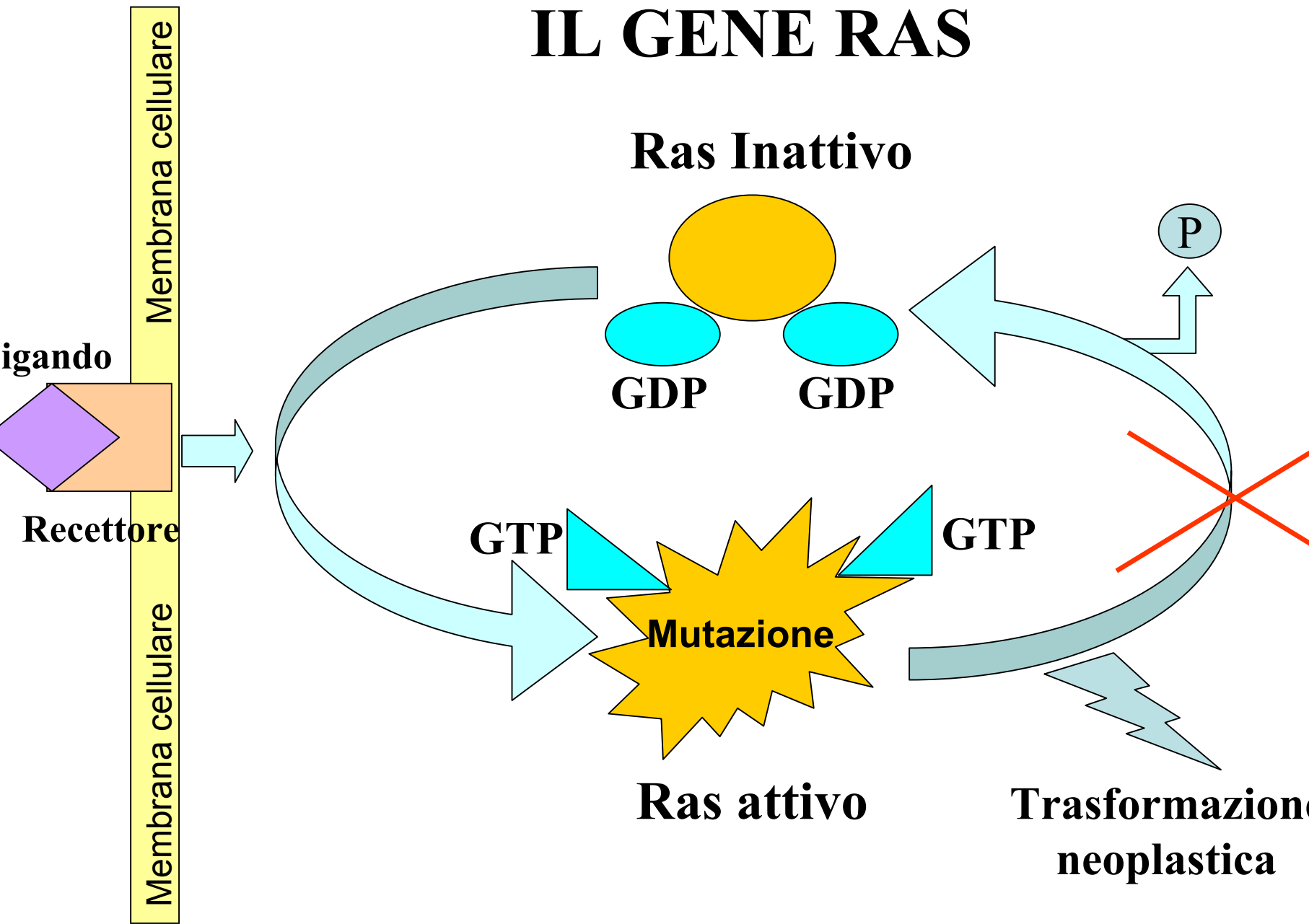
- **Codificano proteine di 21 kd ad attività GTPasica**



IL GENE RAS



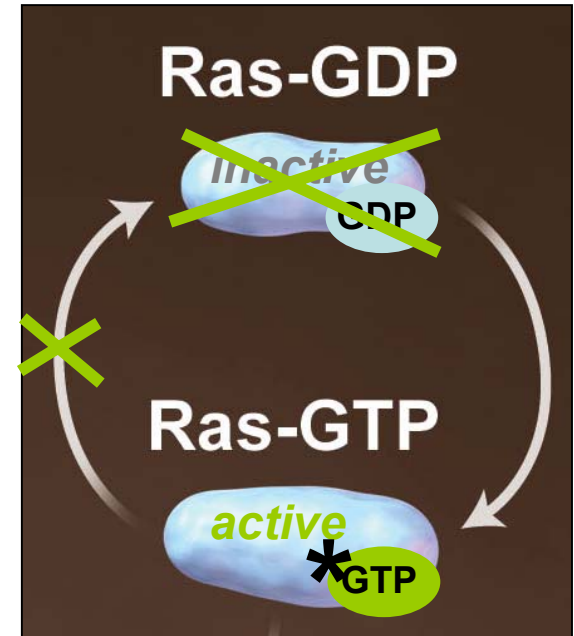
IL GENE RAS



Mutated Ras May Lead to Uncontrolled Cell Signaling and Cancer

– K-ras mutations

- Mutations at codon 12, 13 (exon 2) and 61 (exon 3) near the GTP binding site.
- Mutated proteins lose their GTPase activity.
- This type of mutation is often called “dominant activating”
- There is no way for the cell to “turn off” a protein that has a dominant activating mutation



ABNORMAL

- Growth
- Proliferation
- Differentiation

1. Esteller M, et al. *J Clin Oncol*. 2001; 19:299-304.

2. Schubert S, et al. *Nature Rev Cancer*. 2007; 7:295-308.

Mutated *KRAS* is Prevalent in Many Different Tumor Types

Cancer Type	Reported Incidence of Mutated <i>KRAS</i>
Pancreatic	72 – 90%
Colon	32 – 51% (40%)
Lung	15 – 33%
Ovarian	5 – 50%
Gall bladder	14 – 38%
Multiple myeloma	16 – 33%

K-Tas mutazioni in CRC revealed by direct sequencing

Studio	n° casi studiati	n° mutazioni	%	Codone 12 (%)	Codone 13 (%)
Urosevic et al. [1993]	37	17	46		
Breivik et al. [1994]	251	99	39		
Andreyev et al. [1993]	679	225	33		
Rajagopalan et al. [2002]	330	169	51		
Brink et al. [2003]	737	271	37	72	22
Moroni et al. [2005]	31	10	32	60	40
Ogino et al. [2005]	30	10	33	60	40
Lièvre et al. [2006]	30	13	43		
Benvenuti et al. [2007]	48	16	33	63	37
Wojcik et al. [2008]	163	57	35	66	22
Benvenuti et al. [2008]	175	70	44		
Freeman et al. [2008]	62	24	39		
Totale	2573	981	38	64	32

Single-Arm Studies Support the Hypothesis for *KRAS* as a Biomarker for EGFR Inhibitors

Reference	Treatment (panitumumab or cetuximab)	No of patients (WT:MT)	Objective Response N (%)	
			MT	WT
Relièvre, et al. (AACR Proceedings, 2007)	cmab ± CT	76 (49:27)	0 (0)	24 (49)
S. Benvenuti, et al. (Cancer Res, 2007)	pmab or cmab or cmab + CT	48 (32:16)	1 (6)	10 (31)
W. De Roock, et al. (ASCO Proceedings, 2007)	cmab or cmab + irinotecan	113 (67:46)	0 (0)	27 (40)
D. Finocchiaro, et al. (ASCO Proceedings, 2007)	cmab ± CT	81 (49:32)	2 (6)	13 (26)
F. Di Fiore, et al. (Br J Cancer, 2007)	cmab + CT	59 (43:16)	0 (0)	12 (28)
S. Khambata-Ford, et al. (J Clin Oncol, 2007)	cmab	80 (50:30)	0 (0)	5 (10)
	TOTAL	457 (290:167)	3 (1.7)	91 (20)

KRAS Analysis of a Phase 3, Randomized Trial Comparing Panitumumab vs Best Supportive Care (BSC) in Colorectal Cancer

Amado R et al. *JCO*:26 (10) April 2008.

Hypothesis:

R
A
N
D
O
M
I
Z
E

Panitumumab
6.0 mg/kg Q2W
+ BSC

PD

Follow-up

BSC

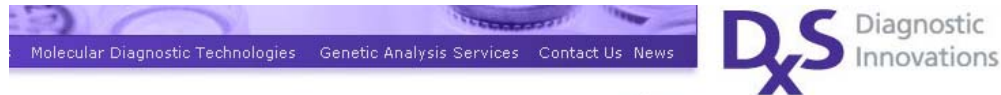
PD

Follow-up

Optional
Panitumumab
Crossover Study

Method Used to Detect *KRAS* Mutational Status

- DNA was isolated from fixed tumor samples
- Mutant *KRAS* was detected using a *KRAS* mutation kit (DxS Ltd, Manchester, UK) that used allele-specific, real-time PCR



TheraScreen: K-RAS MUTATION TEST KIT

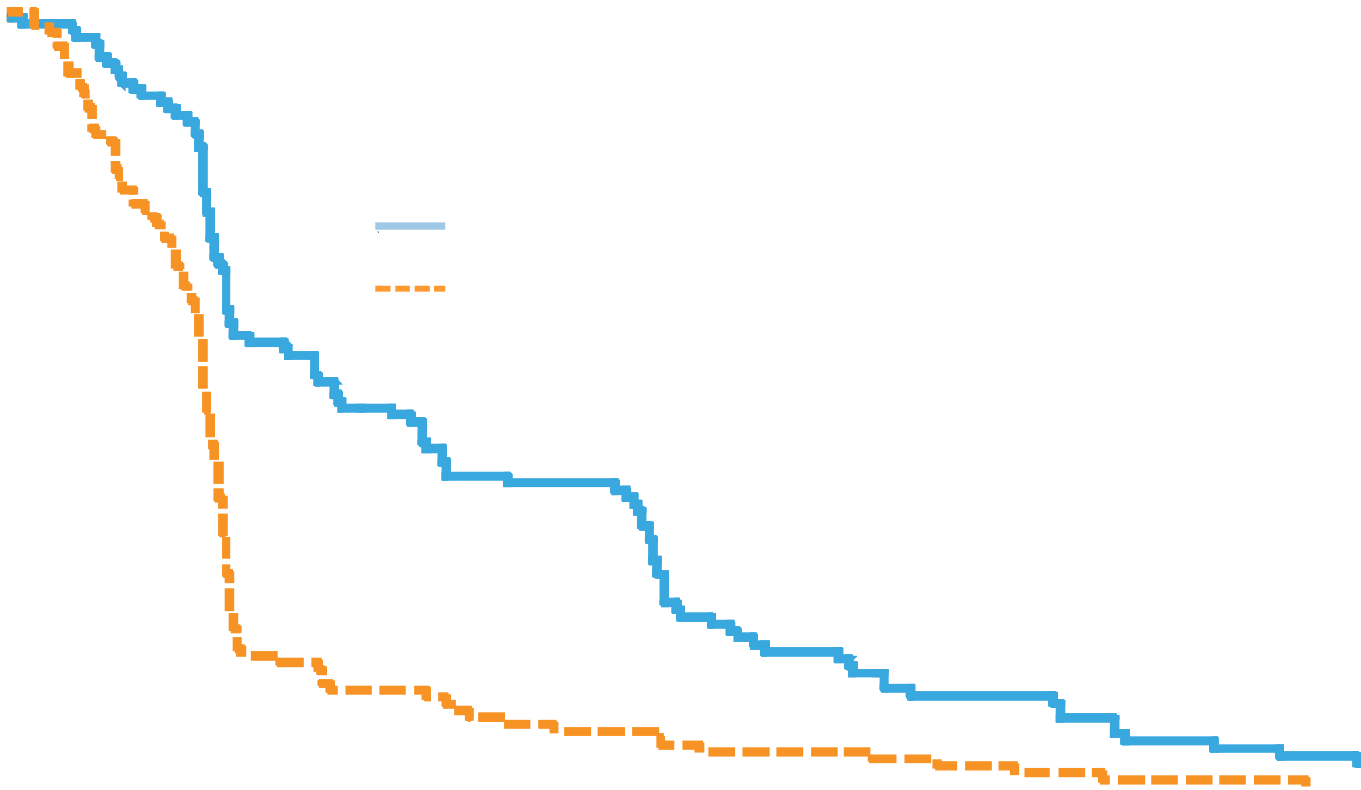
TheraScreen: K-RAS Mutation kit is the first molecular companion diagnostic to support targeted therapies in colorectal cancer. The kit detects seven mutations in codons 12 and 13 of the K-RAS oncogene.

RESULTS

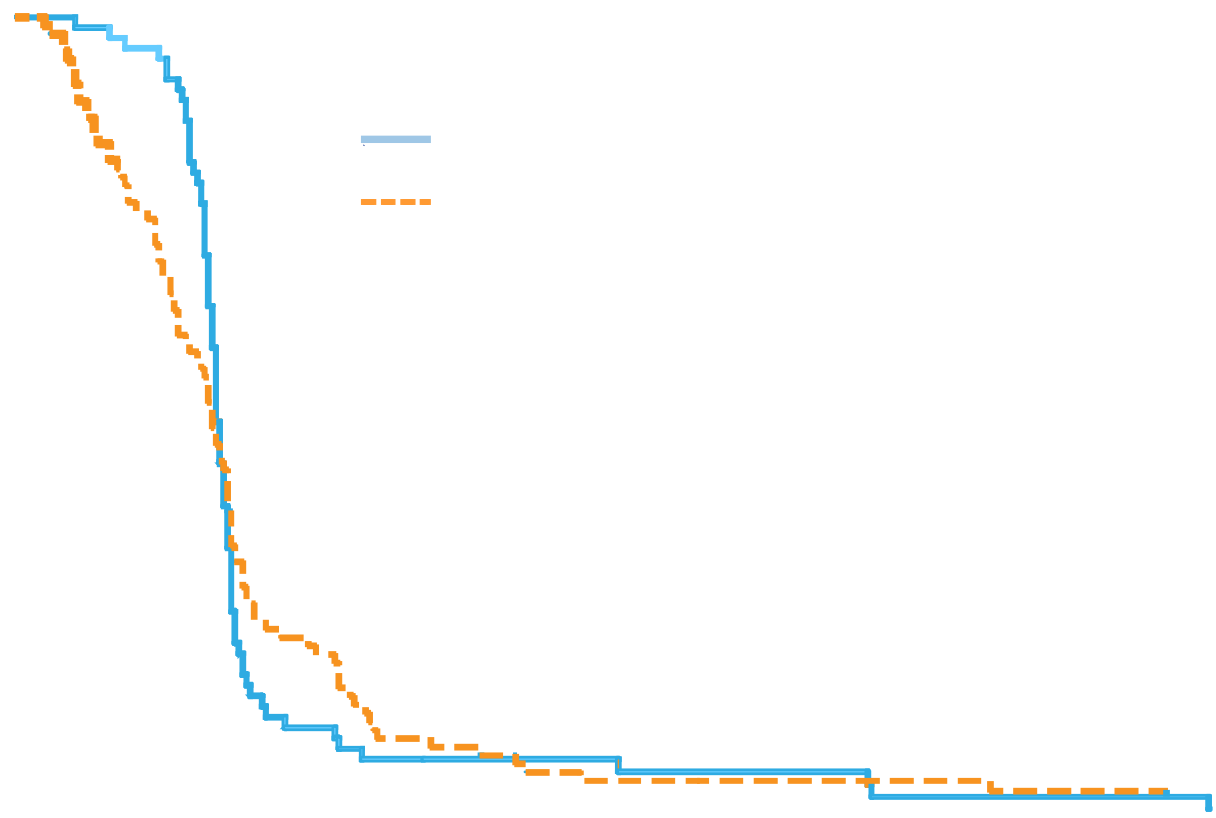
Prevalence of Mutant *KRAS*

	Total
Patients included in <i>KRAS</i> analysis, n (%)	427 (92)
Wild-type <i>KRAS</i> , n (%)	243 (57)
Mutant <i>KRAS</i> , n (%)	184 (43)

Treatment



Mutant KRAS Subgroup PFS by Treatment



Objective Tumor Response (Central Radiology)

Response	<i>KRAS</i>					
	All Evaluable n (%)		Mutant n (%)		Wild-type n (%)	
	Pmab (N = 208)	BSC (N = 219)	Pmab (N = 84)	BSC (N = 100)	Pmab (N = 124)	BSC (N = 119)
CR	0 (0)	0 (0)	0 (0)	0 (0)	0 (0)	0 (0)
PR	21 (10)	0 (0)	0 (0)	0 (0)	21 (17)	0 (0)
SD	52 (25)	22 (10)	10 (12)	8 (8)	42 (34)	14 (12)
PD	104 (50)	149 (68)	59 (70)	60 (60)	45 (36)	89 (75)
CR, PR, SD	73 (35)	22 (10)	10 (12)	8 (8)	63 (51)	14 (12)

PR partial response;

Objective Tumor Response (Central Radiology)

Response	<i>KRAS</i>					
	All Evaluable n (%)		Mutant n (%)		Wild-type n (%)	
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CR, PR, SD	73 (35)	22 (10)	10 (12)	8 (8)	63 (51)	14 (12)

CR, complete response; PR partial response;

SD, stable disease;

Panitumumab (Vectibix)

Anticorpo monoclonale anti-EGFr

- **Parere positivo del Committee for Medicinal Products for Human Use (CHMP): 20 settembre 2007**
- **Decisione della Commissione Europea : 5 dicembre 2007**
- **Vectibix è indicato** *« come monoterapia per il trattamento di pazienti con carcinoma coloretale metastatico esprimenti il recettore per il fattore di crescita epidermico (EGFR) dopo fallimento di regimi chemioterapici contenenti fluoropirimidine,*

Meta-Analysis

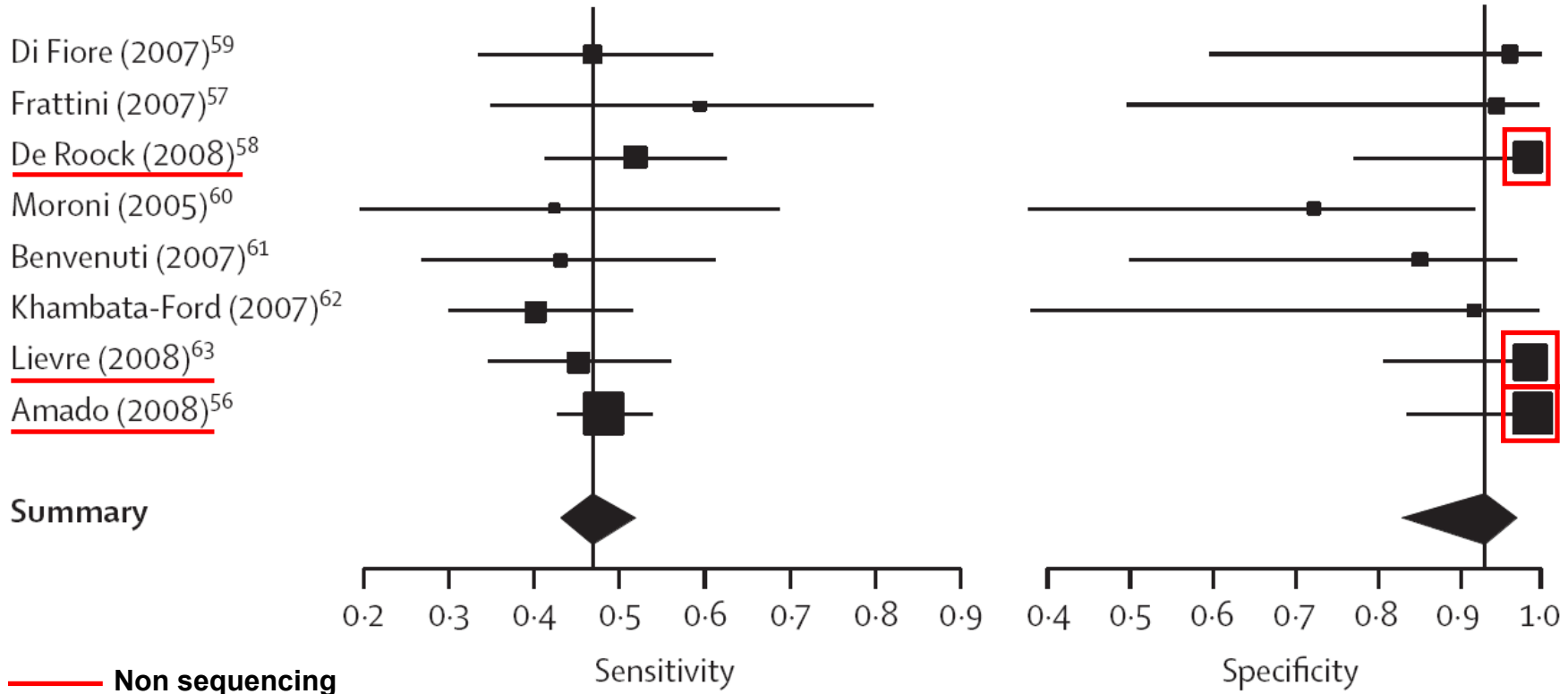
Linardeou H. *Lancet Oncol*, October 2008; 9: 962–72

	Patients included in analysis, n	k-RAS-mutation positive, n (%)	Study design	Mutation analysis	Previous treatment*	Study treatment
Tomado (2008) ^{56†‡}	376	147 (39)	Prospective	<u>DxS</u>	I,O,F	P§±BSC
Attini (2007) ^{57¶}	23	9 (39)	Retrospective	<u>Ex2 bi-ds</u>	≥1 chemotherapy	C+I-based; or C+CAF
de Roock (2008) ⁵⁸	108	42 (39)	Retrospective	<u>AD+sequencing</u>	I	C+I; or C alone**
Fiore (2007) ⁵⁹	59	22 (37)	Retrospective	<u>Ex2 bi-ds</u>	≥1 chemotherapy	C+I; or C+O
Coroni (2005) ^{60¶}	20	7 (35)	Retrospective††	<u>Ex2 bi-ds</u>	≥1 chemotherapy	C+I based; or P alone C alone
Menvenuti (2007) ^{61¶}	37	13 (37)	Retrospective	<u>Ex2 bi-ds</u>	I	C+I based; or P alone C alone
Shambata-Ford (2007) ⁶²	80	30 (38)	Prospective	NR	≥1 chemotherapy	C‡‡
Levre (2008) ^{63§§¶¶}	114	36 (32)	Retrospective	<u>AD+sequencing</u>	≥1 chemotherapy	C+I; or C+FOLFIRI

Comparison of all studies and subgroups for metastatic colorectal cancer

	Sensitivity (95% CI)	p value	Specificity (95% CI)	p value	+LR	-LR	Predictive odds ratio
Overall (n=8)	0.47 (0.43-0.52)	..	0.93 (0.83-0.97)	..	6.82	0.57	12.01
MoAb							
Panitumumab (n=4)	0.48 (0.41-0.55)	0.25	0.84 (0.54-0.96)	0.86	2.95	0.62	4.74
Cetuximab (n=4)	0.41 (0.30-0.54)	..	0.86 (0.48-0.98)	..	2.94	0.68	4.30
MoAb combination							
Panitumumab (n=4)	0.48 (0.42-0.54)	0.72	0.84 (0.58-0.95)	0.21	2.95	0.62	4.74
Cetuximab±chemotherapy (n=10)	0.46 (0.40-0.53)	..	0.93 (0.83-0.97)	..	6.75	0.57	11.74
chemotherapy							
MoAb (n=8)	0.46 (0.41-0.52)	0.54	0.87 (0.69-0.95)	0.14	3.59	0.62	5.82
MoAb+chemotherapy (n=6)	0.49 (0.41-0.57)	..	0.96 (0.85-0.99)	..	11.05	0.53	20.67
Detection method*							
Sequencing (n=4)	0.47 (0.35-0.60)	0.82	0.87 (0.62-0.96)	0.02	3.65	0.61	6.01
Other (n=3)	0.48 (0.42-0.54)	..	0.99 (0.89-1.00)	..	33.36	0.52	63.58

Forest plots representing all studies for *k*-RAS-mutation sensitivity and specificity in colorectal carcinoma



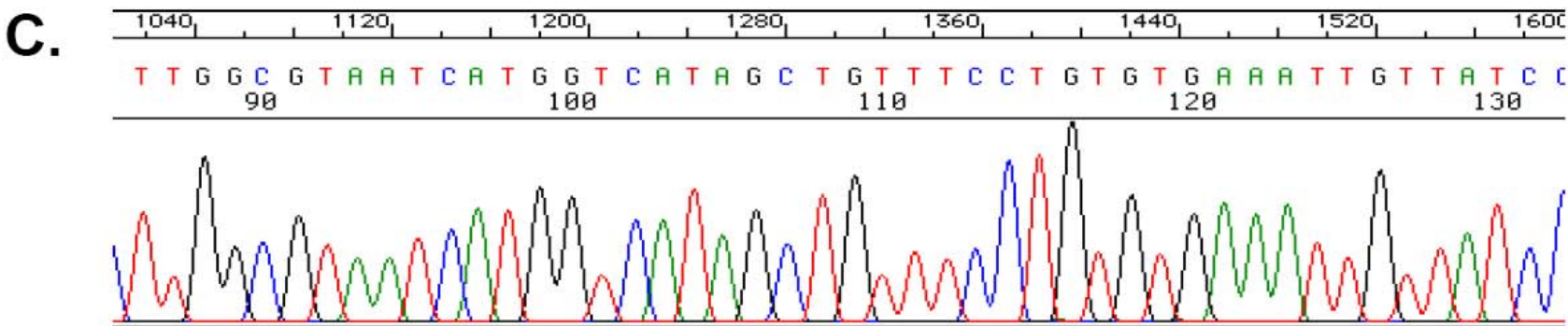
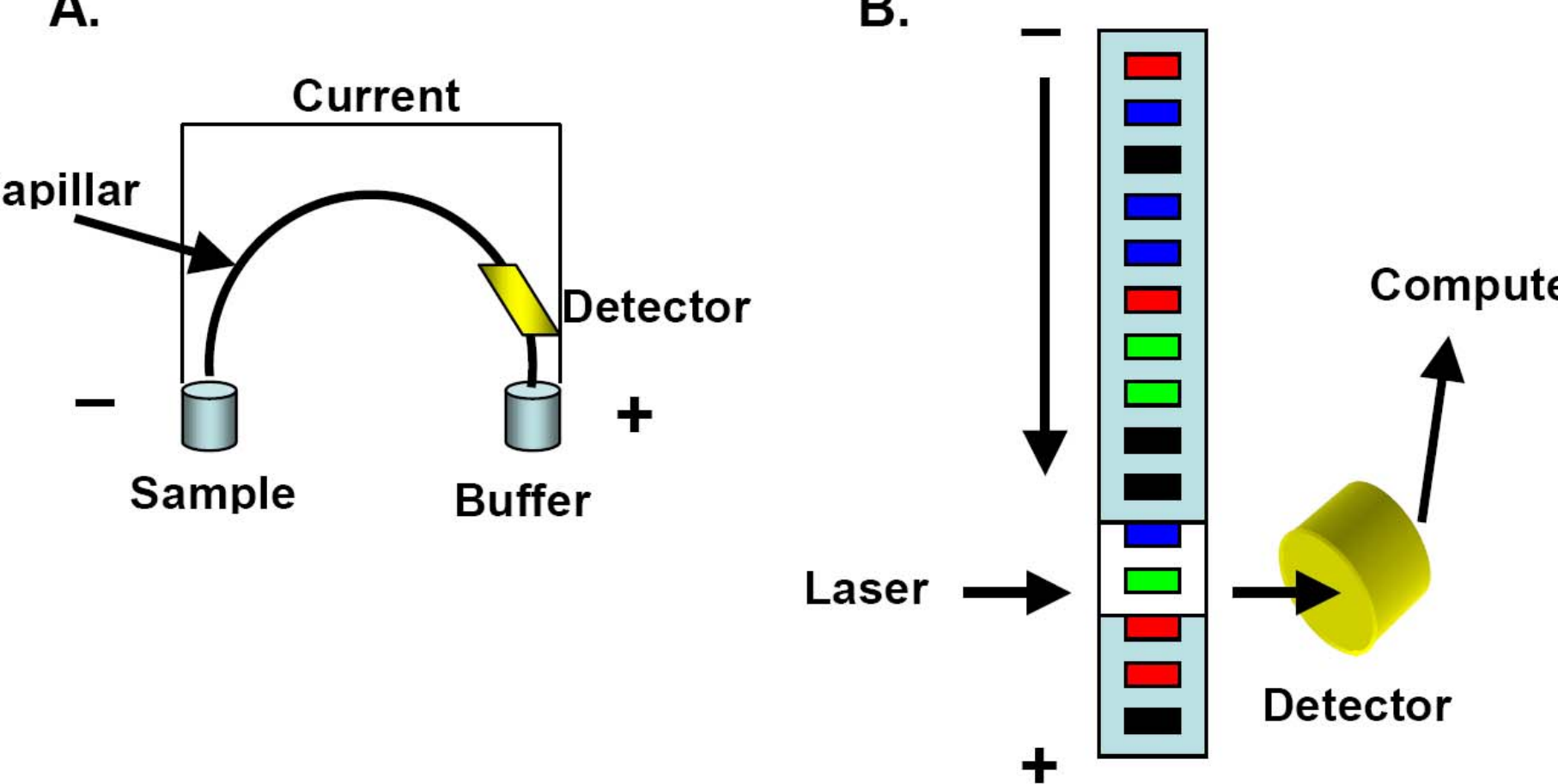
The main problem is the low sensitivity of the K-ras marker

Resistenza farmacologica su base mutazionale.

Extremely sensitive techniques are needed

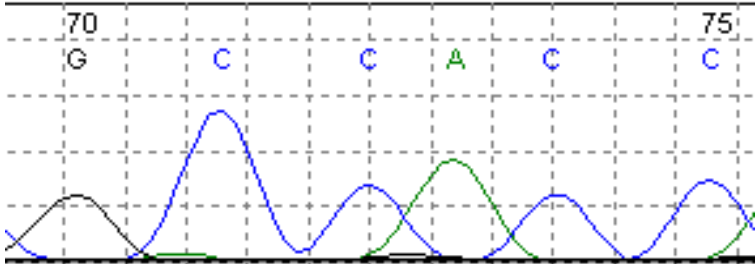
Direct sequencing of the PCR products



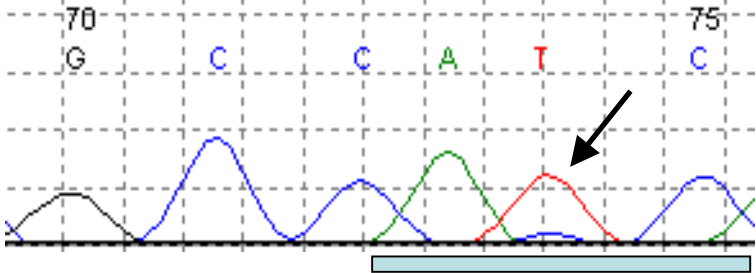


Codone 12 (GGT)

Tessuto normale



Tumore



Codone 12 (GAT)

The direct sequencing is not the the most sensitive method available

Detection limit of sequencing = 1:5 (20%)

Neoplastic cells in a tumor samples 70%

The mutation is usually heterozigous 35%

Tumor policlonal for mutation (50% of the neoplastic cells) 17%



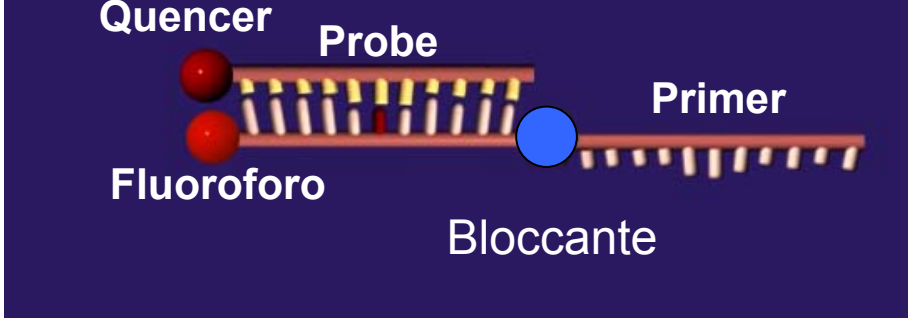
TheraScreen: K-RAS MUTATION TEST KIT

TheraScreen: K-RAS Mutation kit is the first molecular companion diagnostic to support targeted therapies in colorectal cancer. The kit detects seven mutations in codons 12 and 13 of the K-RAS oncogene.

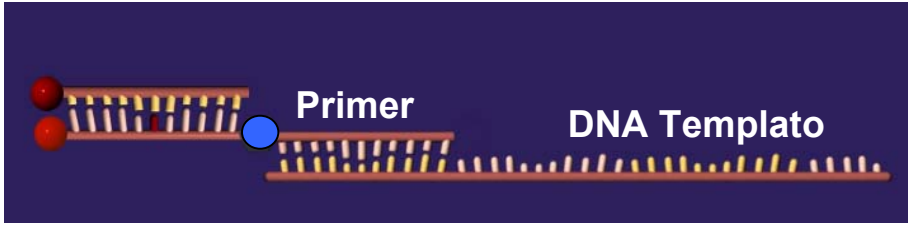
Technological principles

The Kit combines ARMS (allele specific PCR) with the scorpions real time PCR technology

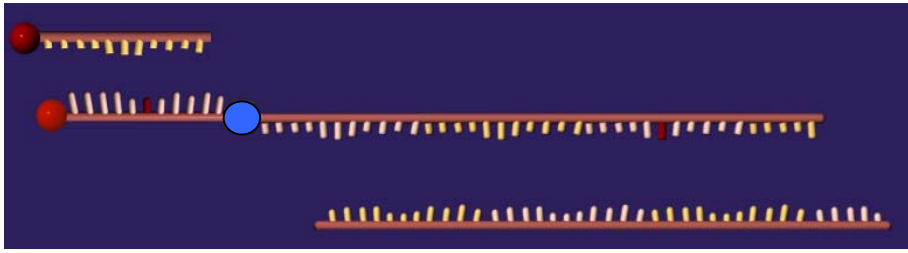
A Sonda scorpion



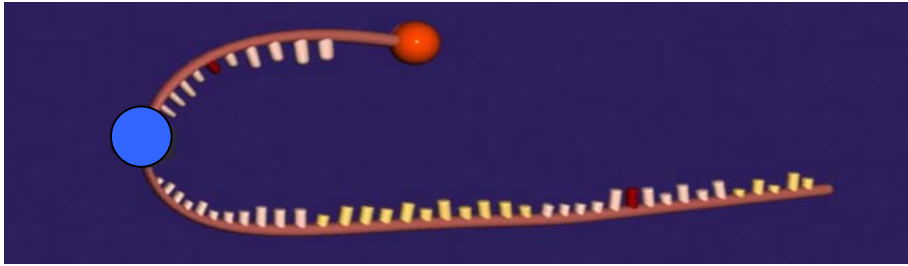
B Ibridazione al template



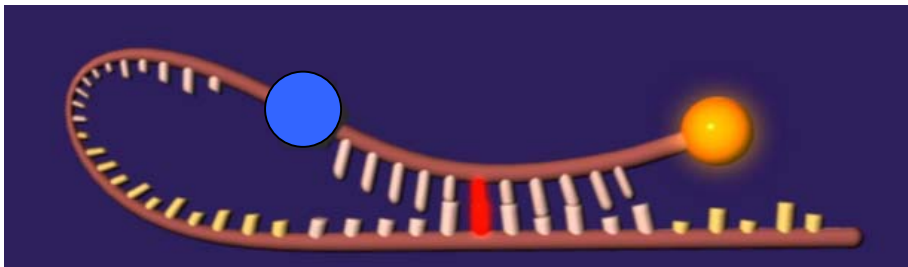
C Denaturazione



D Formazione dell'ansa



E Ibridazione ed emissione di fluorescenza



SENSITIVITY

The **TERASCREEN** assays can detect 1% of mutant in a background of wild type genomic DNA.

The test identifies 7 somatic mutations in codons 12 and 13

Codon 12	Gly 12 Asp
	Gly 12 Ala
	Gly 12 Val
	Gly 12 Ser
	Gly 12 Arg
	Gly 12 Cys
Codon 13	Gly 13 Asp

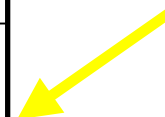
Amado R et al. *JCO*:26 (10) April 2008

39% of K-ras mutations

15% in codon 13

K- ras mutations in CRC revealed by direct sequencing

Studio	n° casi studiati	n° mutazioni	%	Codone 12 (%)	Codone 13 (%)
Urosevic et al. [1993]	37	17	46		
Breivik et al. [1994]	251	99	39		
Andreyev et al. [1993]	679	225	33		
Rajagopalan et al. [2002]	330	169	51		
Brink et al. [2003]	737	271	37	72	22
Moroni et al. [2005]	31	10	32	60	40
Ogino et al. [2005]	30	10	33	60	40
Lièvre et al. [2006]	30	13	43		
Benvenuti et al. [2007]	48	16	33	63	37
Wojcik et al. [2008]	163	57	35	66	22
Benvenuti et al. [2008]	175	70	44		
Freeman et al. [2008]	62	24	39		
Totale	2573	981	38	64	32



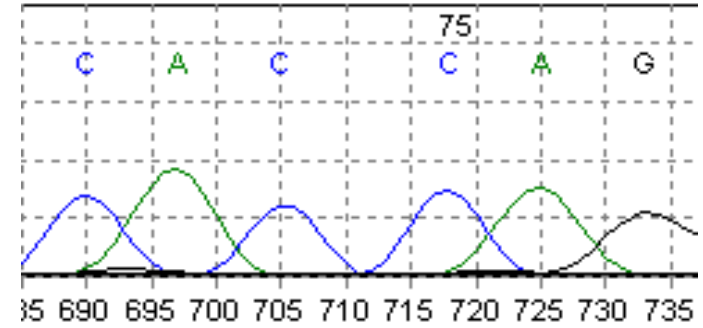
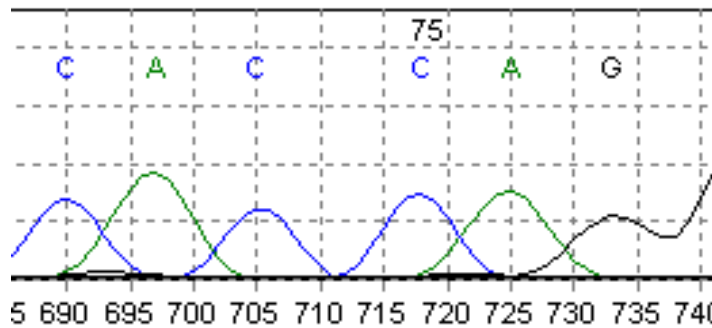
Sensitive detection of K-ras mutations

Enriched sequencing

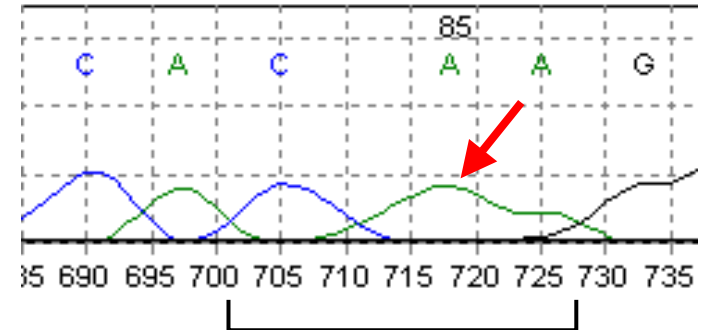
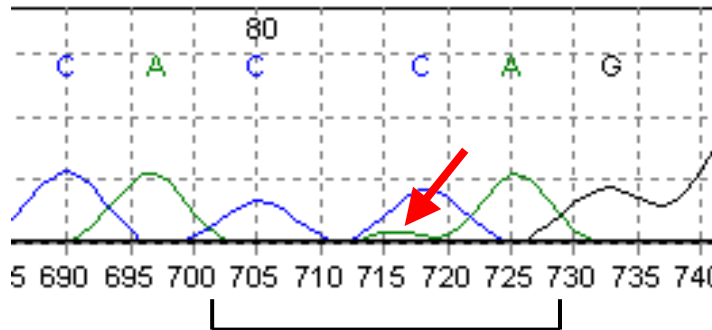
Normal Sequencing

Enriched sequencing

Normal



Tumor
379T



Codon 12

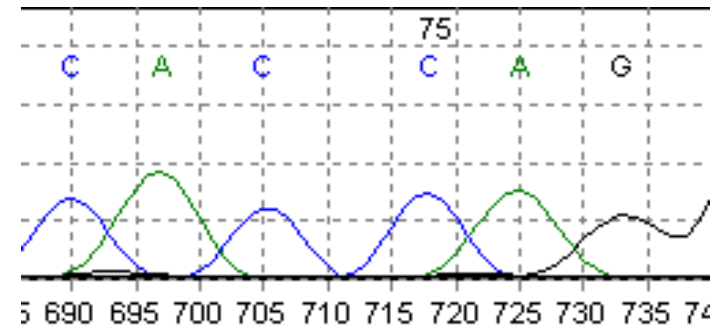
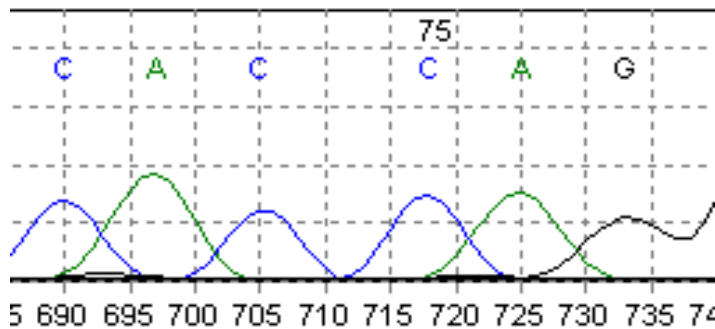
Codon 12

Sensitive detection of K-ras mutations

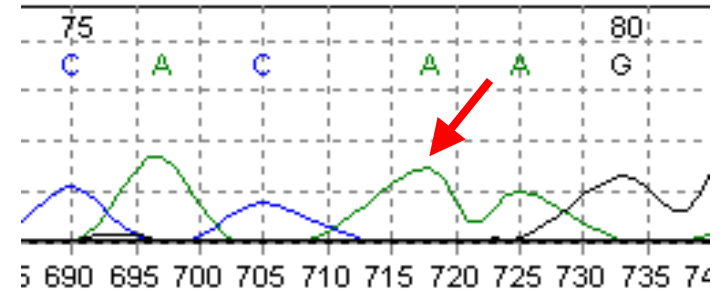
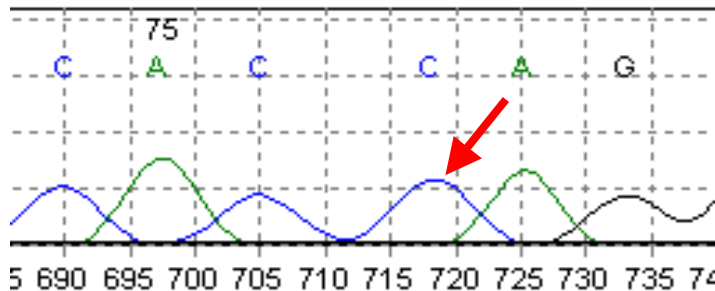
Normal Sequencing

Enriched sequencing

Normal



Tumor
382T



Detection limit of Enriched sequencing = 1:1000 (0,001%)

In a series of 90 colorectal carcinomas

METHOD	(%) of K-ras mutations
Direct sequencing:	39 %
Enriched sequencing	54 %

In a series of 126 NSCLC treated with Erlotinib :

METHOD	(%) of K-ras mutations
Direct sequencing:	17 %
Enriched sequencing	30 %



Overall survival
(multivariate
analysis)

	HR (95% CI)	<i>P</i>
KRAS mut (Direct sequencing)	2.45 (0.92-6.319)	0.044
KRAS mut (Enriched)	3.19 (1.43-7.16)	0.005

Ferdinand I ([June 2, 1423](#) – [January 25, 1494](#)),

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Ferdinand I of Naples

From Wikipedia, the free encyclopedia

Ferdinand I of Naples should not be confused with Ferdinand I of the Two Sicilies, a latter king of Naples.

Ferdinand I ([June 2, 1423](#) – [January 25, 1494](#)), also called **Don Ferrante**, was the [King of Naples](#) from [1458](#) to [1494](#). He was the natural son of [Alfonso V of Aragon](#).

Contents [hide]

- 1 Biography
- 2 Ferdinand's reputation
- 3 Marriages and children
- 4 Notes
- 5 External links



Biography [edit]

In order to arrange a good future for Ferdinand, King Alfonso had him married in 1444 to a feudal heiress, [Isabella of Taranto](#), who besides being the elder daughter of Tristan di Chiaramonte (Tristan de Clermont-Lodeve), Count of [Copertino](#), and Catherine of [Baux Orsini](#), was the niece and heiress presumptive of childless prince [Giovanni Antonio del Balzo Orsini of Taranto](#). She was a granddaughter of Queen [Mary of Enguien](#) (mother of Giovanni and Catherine), who had been Queen Consort of Naples (Queen of Jerusalem and Sicily) in 1406-14.



Ferdinand I of Naples.

He was autopsied by an Italian team. They concluded that King Ferrante I had died of a large pelvic tumor, either prostate cancer or colorectal cancer.

Two years later, in 1996, genetic research of a small part of the tumor revealed a KRAS mutation, which is rare in prostate cancer but is frequently encountered in colorectal tumors. Concluded is that King Ferrante I of Naples died from a colorectal tumor and is therefore the earliest documented person in history who died from this cancer.”

K-ras mutations in the tumour of Ferrante I of Aragon, King of Naples

A. Marchetti, S Pellegrini, G. Bevilacqua, G. Fornaciari.

Saturday 4 May 1996, vol. 347 No. 9010

K-ras mutation: G→A

