

# Thérapies ciblées : quel traitement pour quel patient ?

Prof Thierry Berghmans

Service des Soins Intensifs et Urgences  
Oncologiques & Clinique d'Oncologie Thoracique

Institut Jules Bordet, Centre des Tumeurs de  
l'Université Libre de Bruxelles

Bruxelles, Belgique

# Conflits d'intérêt

- Aucun conflit d'intérêt en relation avec le sujet traité

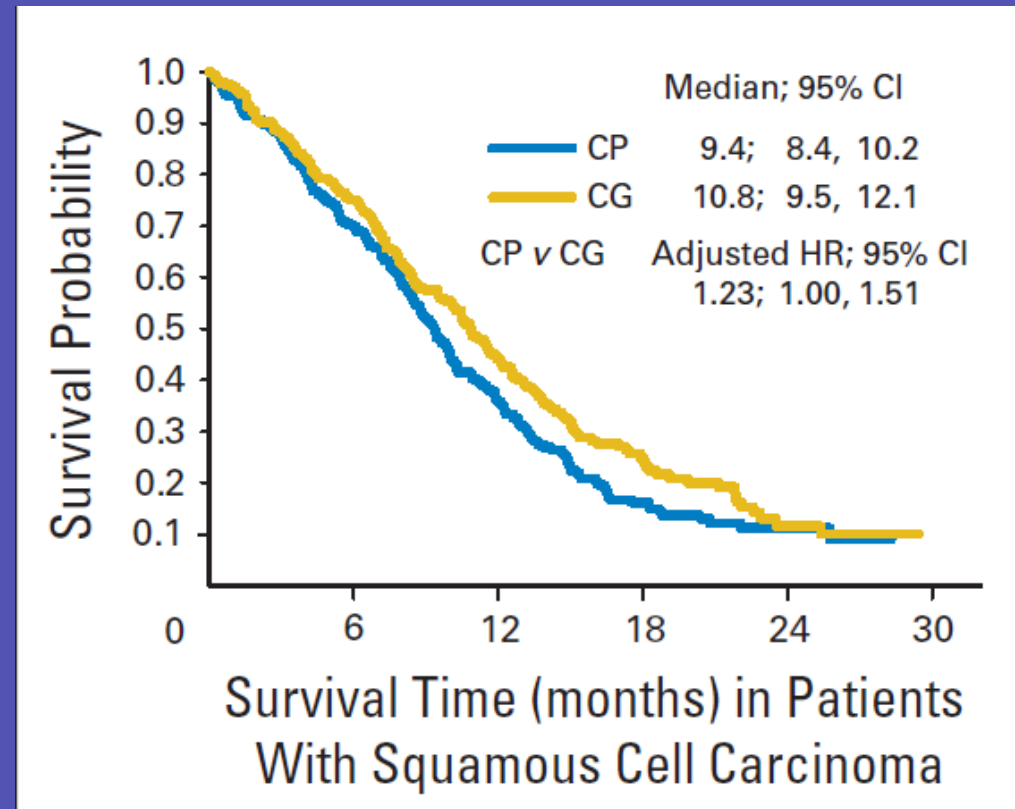
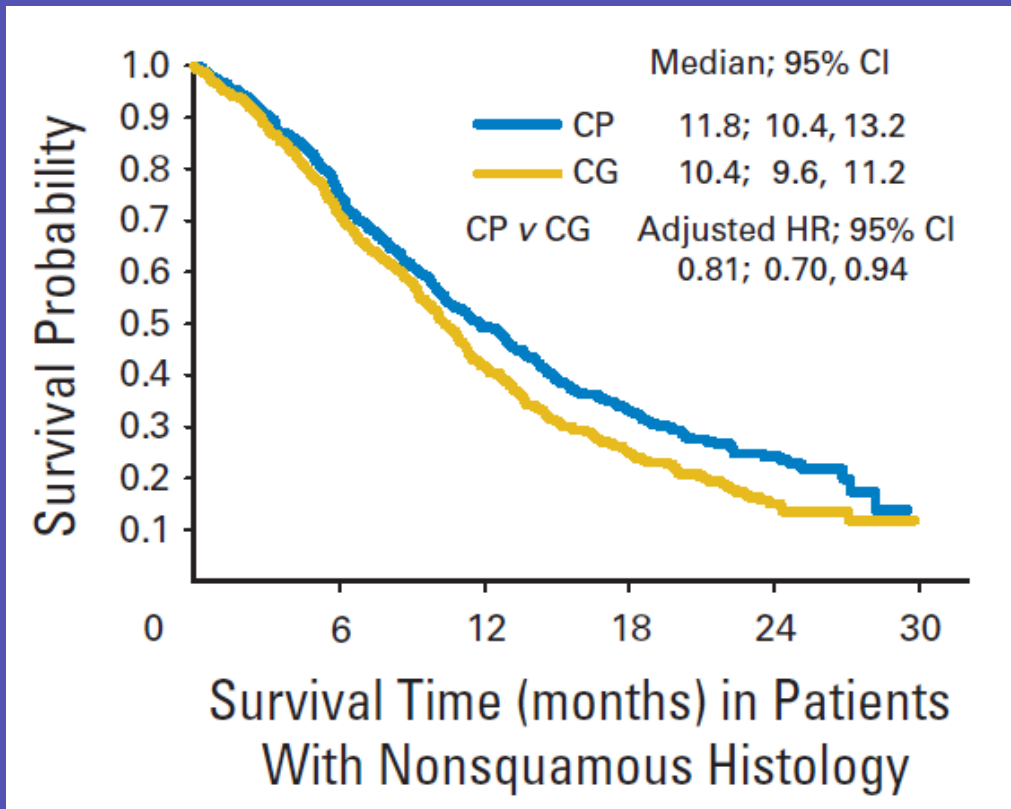
# Plan du cours

1. Qu'entend-on par thérapie ciblée?
2. Chez qui chercher une « cible »?
3. Place des traitements ciblés disponibles en routine:  
EGFR et ALK
4. Futur des thérapies ciblées
5. Quelques mots d'immunothérapie
6. Rôle des consultations multidisciplinaires

# 1. Qu'entend-on par thérapie ciblée?

# Qu'est-ce qu'un traitement "ciblé" ?

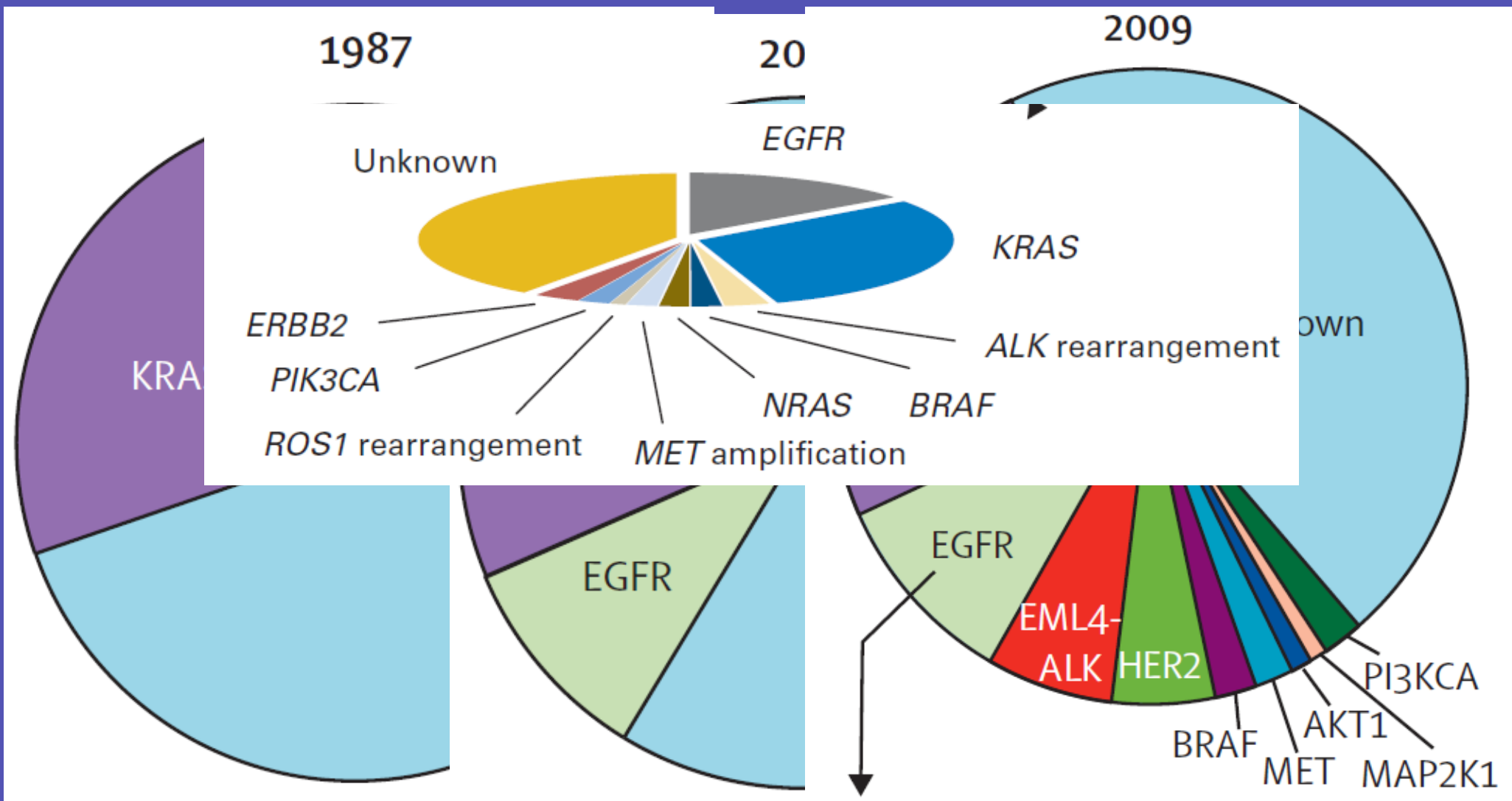
- **Groupe-Cible:** population avec caractéristiques prédisposant à sensibilité au traitement = Population enrichie



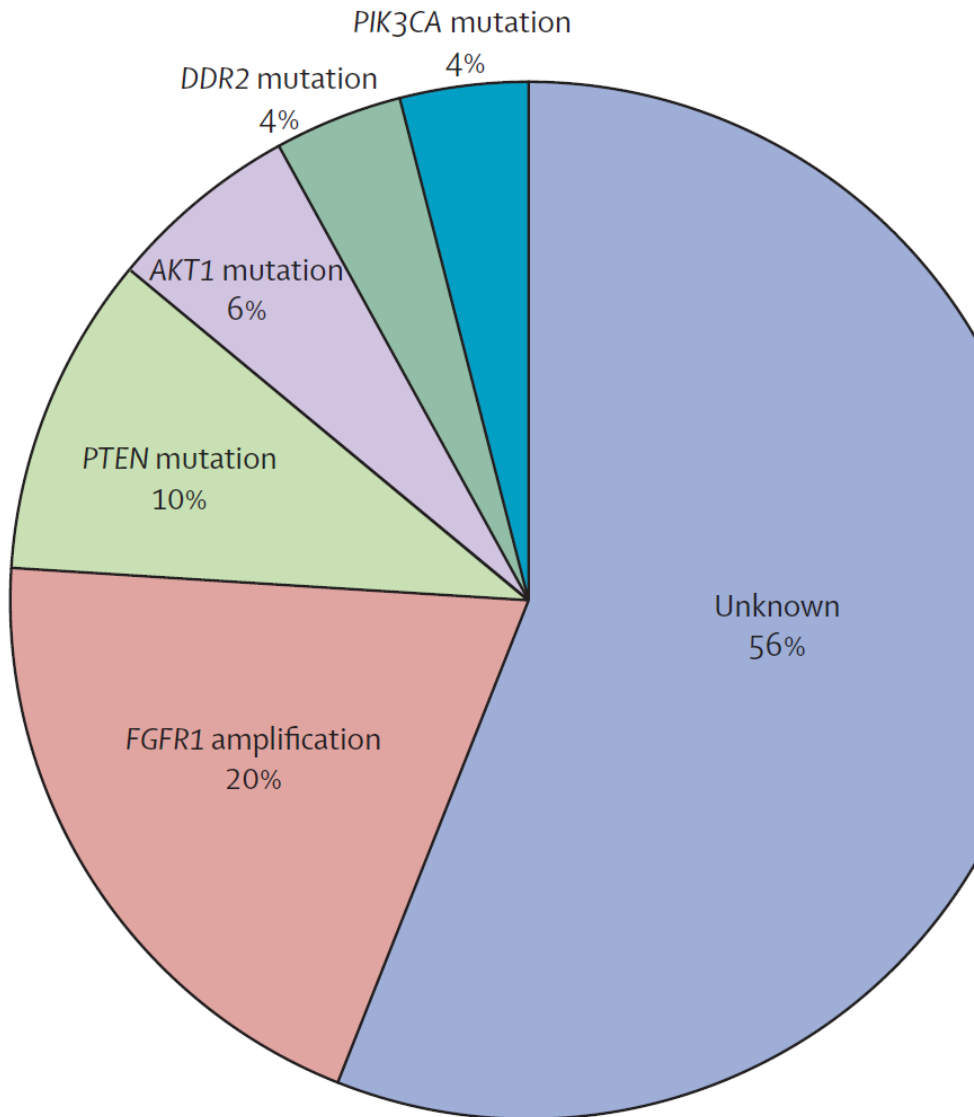
# Qu'est-ce qu'un traitement "ciblé" ?

- **Thérapies biologiques ciblées**: ciblent de manière « spécifique » des voies métaboliques vitales pour la cellule cancéreuse
- **Cibles**:
  - récepteurs de facteur de croissance
  - médiateurs intracellulaires appartenant à une cascade signalétique
  - facteurs de néo-angiogenèse
  - gènes...

# Evolution des connaissances et cibles moléculaires: adénocarcinomes



# Cibles moléculaires: épidermoïdes



**TABLE 1.** Key Candidate Genes in Squamous Cell Lung Cancer

Gene	Amplification		Mutation	
	Squamous	Adeno	Squamous	Adeno
FGFR1	20%	1%		
SOX	20%	n.r.		
PIK3CA	20%	5%	6%	1.5%
MDM2	10%	5%		
PDGFRA	8–10%	3–7%		
MET	6%	<5%		
P53			65%	40%
NRF2			10–15%	1–2%
PTEN			10.2%	1.7%
EPHA2			7%	0
LKB1			5%	13%
AKT			5%	0
EGFR vIII			5%	0
DDR2			3–4%	n.r.
PTEN loss				

IHC expression 70–75%  
 hypermethylation 30–35%  
 LOH 18–22%  
 not significantly different by histology



2. Chez qui chercher une « cible »?

# Molecular Testing Guideline for Selection of Lung Cancer Patients for EGFR and ALK Tyrosine Kinase Inhibitors

## *Guideline from the College of American Pathologists, International Association for the Study of Lung Cancer, and Association for Molecular Pathology*

Neal I. Lindeman,<sup>\*</sup> Philip T. Cagle,<sup>†</sup> Mary Beth Beasley,<sup>‡</sup> Dhananjay Arun Chitale,<sup>§</sup> Sanja Dacic,<sup>¶</sup> Giuseppe Giaccone,<sup>||</sup> Robert Brian Jenkins,<sup>\*\*</sup> David J. Kwiatkowski,<sup>††</sup> Juan-Sebastian Saldivar,<sup>‡‡</sup> Jeremy Squire,<sup>§§</sup> Erik Thunnissen,<sup>¶¶</sup> and Marc Ladanyi<sup>|||</sup>

### Question 1: Which Patients Should Be Tested for *EGFR* Mutations and *ALK* Rearrangements?

- 1.1a: Recommendation: *EGFR* molecular testing should be used to select patients for EGFR-targeted TKI therapy, and patients with lung adenocarcinoma should not be excluded from testing on the basis of clinical characteristics.
- 1.1b: Recommendation: *ALK* molecular testing should be used to select patients for ALK-targeted TKI therapy, and patients with lung adenocarcinoma should not be excluded from testing on the basis of clinical characteristics.

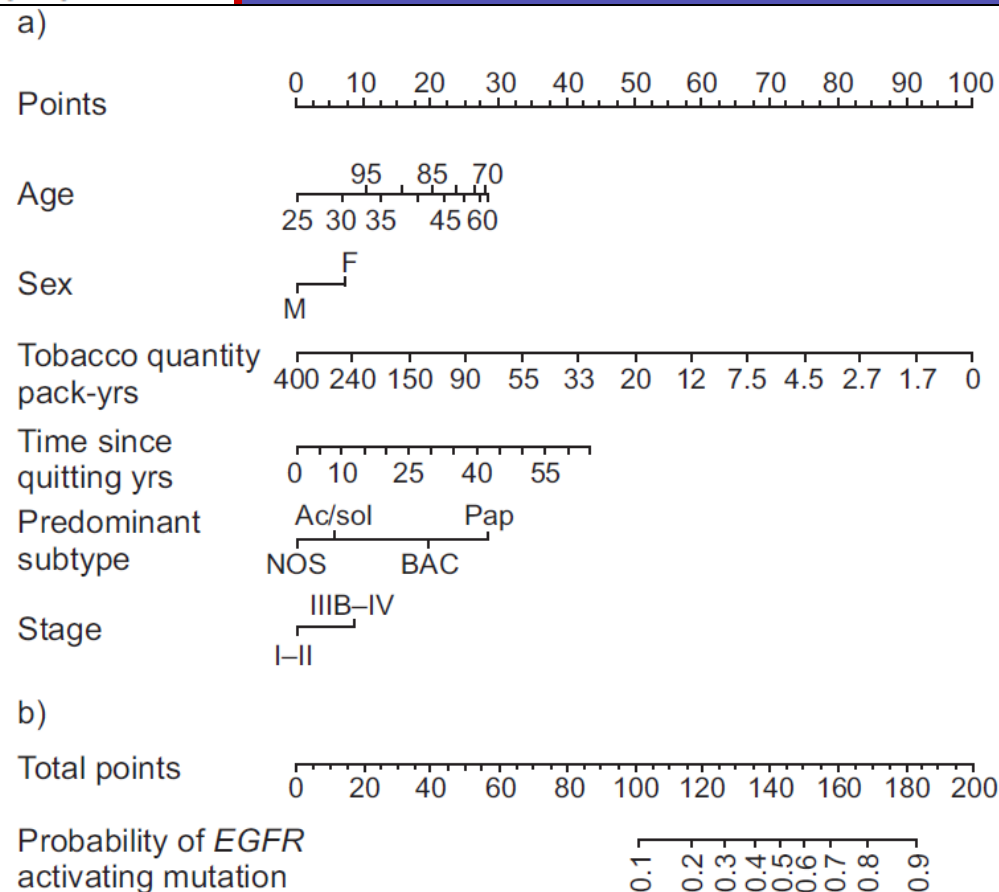
### Question 2: When Should a Patient Specimen Be Tested for *EGFR* Mutation or *ALK* Rearrangement?

- 2.1a: Recommendation: *EGFR* mutation testing should be ordered at the time of diagnosis for patients presenting with advanced-stage disease (stage IV according to the 7th edition TNM staging system) who are suitable for therapy or at time of recurrence or progression in patients who originally presented with lower-stage disease but were not previously tested.
- 2.1b: Suggestion: *ALK* rearrangement testing should be ordered at the time of diagnosis for patients presenting with advanced-stage disease (stage IV according to the 7th edition TNM staging system) who are suitable for therapy or at time of recurrence or progression in patients who originally presented with lower-stage disease but were not previously tested.

# Facteurs prédictifs de la présence d'une mutation activatrice d'EGFR

Nomogram to predict the presence of *EGFR* activating mutation in lung adenocarcinoma

	Total (%)	Non-East Asian (%)
All subgroups	19	10
Smokers	11	4
Nonsmokers	54	35
Adenocarcinoma	42	16
Non-adenocarcinoma	3	1
Male	16	1
Female	46	20



Clin Cancer Res 2006: 4416S

Reference	N pts	EML4-ALK+ n (%)	ADC (n+/n)	SCC (n+/n)	Test	Clinical characteristics associated with <i>EML4-ALK</i>	
Soda (8), 2007	65	5 (7.7%)	4	1	PCR	-	
Koivunen (17), 2008	305	8 (2.6%)	8/208 (3.8%)	0/88	FISH	Non/light smoking (p 0.049)	
Inamura (15), 2008	221	5 (2.3%)	5/149 (3.4%)	0/72	PCR	-	
Perner (57), 2008	603	16 (2.7%)	-	-	FISH	-	
Wong (21), 2009	266	<p style="text-align: center;"><b>Facteurs prédictifs de ALK</b></p> <p style="text-align: center;"><b>Non fumeur/fumeur léger</b></p> <p style="text-align: center;"><b>Patients plus jeunes</b></p> <p style="text-align: center;"><b>Adénocarcinome</b></p>					oking (p 0.009), age (p
Shaw (18), 2009	141						0.005), male (p 0.036), t smoking (p< 0.001)
Inamura (16), 2009	363						0.006), light smoking (p status (p 0.039), histological tiation (p 0.008)
Takahashi (20), 2010	313						oking (p<0.001), histological differentiation (p<0.001)
Sun (19), 2010	52	3 (5.8%)	3/52 (5.8%)	-	PCR	-	
Martelli (29), 2009	120	9 (7.5%)	3/60 (5%)	4/48 (8.3%)	PCR	-	
Zhang (22), 2010	103	12 (11.7%)	10/62 (16.1%)	2/29 (6.9%)	PCR	Non smoking (p 0.03)	
Kwak (24), 2010	1500	82 (5.5%)	79	1	FISH	-	


# Autres exemples de facteurs prédictifs de la présence d'une cible moléculaire

Cible	Population
KRAS	Adénocarcinome, fumeur
HER-2	Adénocarcinome, non fumeur, sexe féminin
RET translocation	Adénocarcinome, non fumeur
ROS-1 translocation	Adénocarcinome, jeune, non fumeur
BRAF	Fumeur

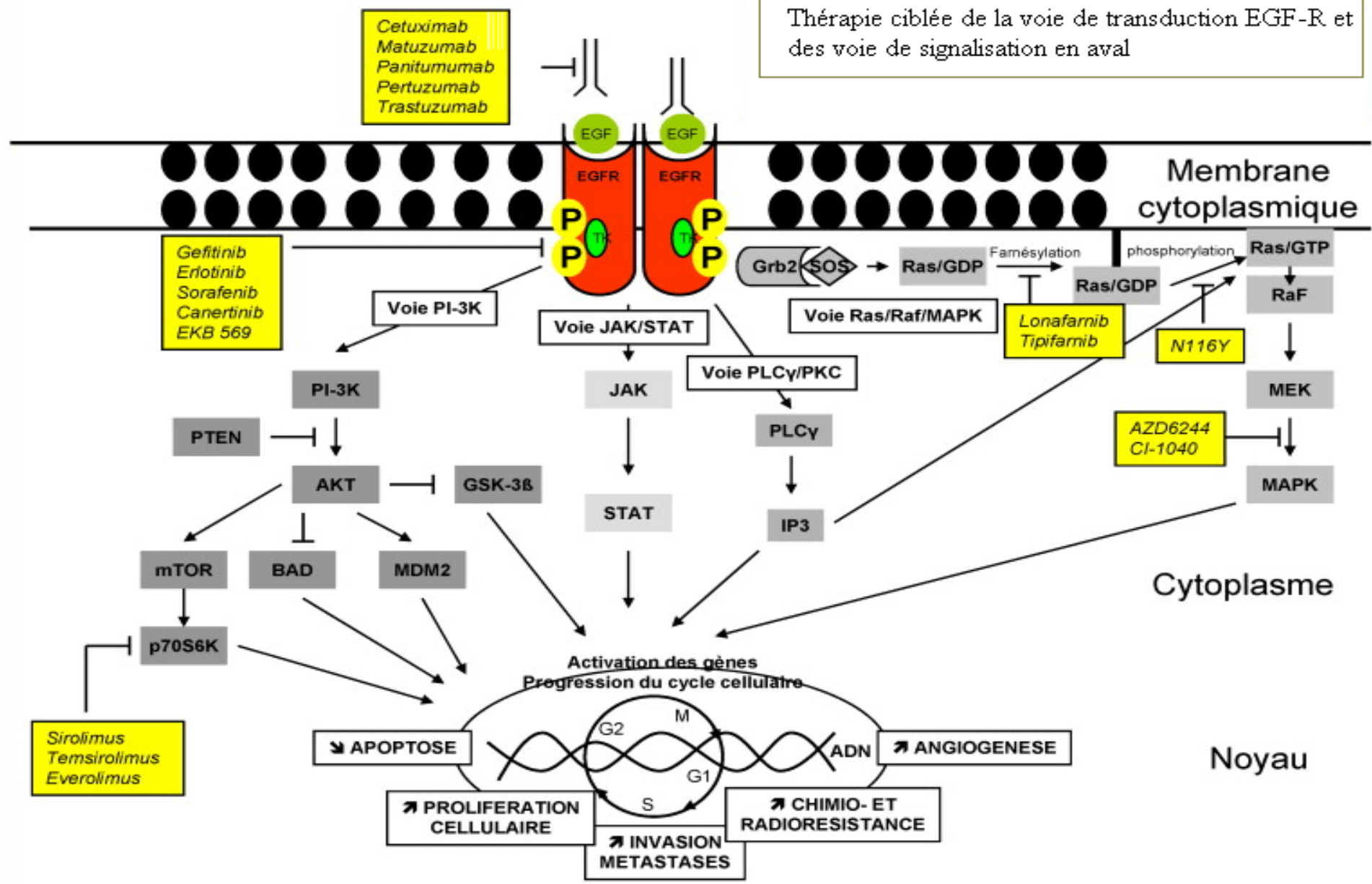
### 3. Place des traitements ciblés disponibles en routine: EGFR et ALK

EGFR

# Epidermal growth factor receptor

- Ubiquitaire mais présent en fortes quantités dans:
  - Tractus aérodigestif
  - Peau
- Nombreux ligands:
  - EGF
  - TGF $\alpha$ , amphiréguline,  $\alpha$ -celluline, épiréguline ...





# Molécules disponibles

- Au moins 4 molécules sur le marché
  - Gefitinib
  - Erlotinib
  - Afatinib
  - Icotinib (Chine)
- Nouveaux agents prometteurs de 3<sup>ème</sup> génération
  - AZD9291
  - CO-1686
  - HM61713

Référence	Chimiothérapie	N pts	% RO	p	SM	p	Phase
1 <sup>ère</sup> ligne en addition avec la chimiothérapie (population non sélectionnée)							
Giaccone (11)	Gemcitabine + CDDP + géfinitib 500 mg	365	50%	NS	9,9 m	NS	III
	Gemcitabine + CDDP + géfinitib 250 mg	365	51%		9,9 m		
	Gemcitabine + CDDP	363	47%		10,9 m		
Herbst (10)	Paclitaxel + CBDCA + géfinitib 500 mg	347	30%	NS	8,7 m	NS	III
	Paclitaxel + CBDCA + géfinitib 250 mg	345	30,4%		9,8 m		
	Paclitaxel + CBDCA	345	28,7%		9,9 m		
Herbst (12)	Paclitaxel + CBDCA	540	19,3%	NS	10,5 m	NS	III
	Paclitaxel + CBDCA + erlotinib	539	21,5%		10,6 m		
Gatzemeier (13)	Gemcitabine + CDDP + erlotinib	580	31,5%	NS	43 s	NS	III
	Gemcitabine + CDDP	579	29,9%		44,1 s		
En 1 <sup>ère</sup> ligne, en traitement séquentiel (population non sélectionnée)							
Takeda (14)	Doublet de platine puis Gefitinib	300	34%	NS	13,7 m	NS	III
	Doublet de platine	298	29%		12,9 m		
En 1 <sup>ère</sup> ligne, en comparaison à des soins de soutien chez des patients inéligibles pour une chimiothérapie conventionnelle							
Lee (18)	Erlotinib	350	4,3%		3,7m	0,46	III
	Soins de confort	320	2,2%		3,6m		
En maintenance, en l'absence de progression après 4 cycles à base de platine							
Cappuzzo (34)	Erlotinib	437	NA		12 m	S	III
	Placebo	447			11 m		
Gaafar (33)	Géfitinib	86	NA		10,9 m	NS	III
	Placebo	87			9,4 m		
Zhang (32)	Géfitinib	148	NA		18,7m	0,26	III
	Placebo	148			16,9m		
Pérol (31)	Gemcitabine	155	NA		?	NS	III
	Erlotinib	154			11,4 m		
	Observation	155			10,8m		
	Erlotinib + docétaxel ou pémétréxed	116			39%		

- Pas de place pour un ITK d'EGFR en 1<sup>ère</sup> ligne dans des populations non sélectionnées ou en l'absence de mutation
- Rôle en traitement de maintenance??

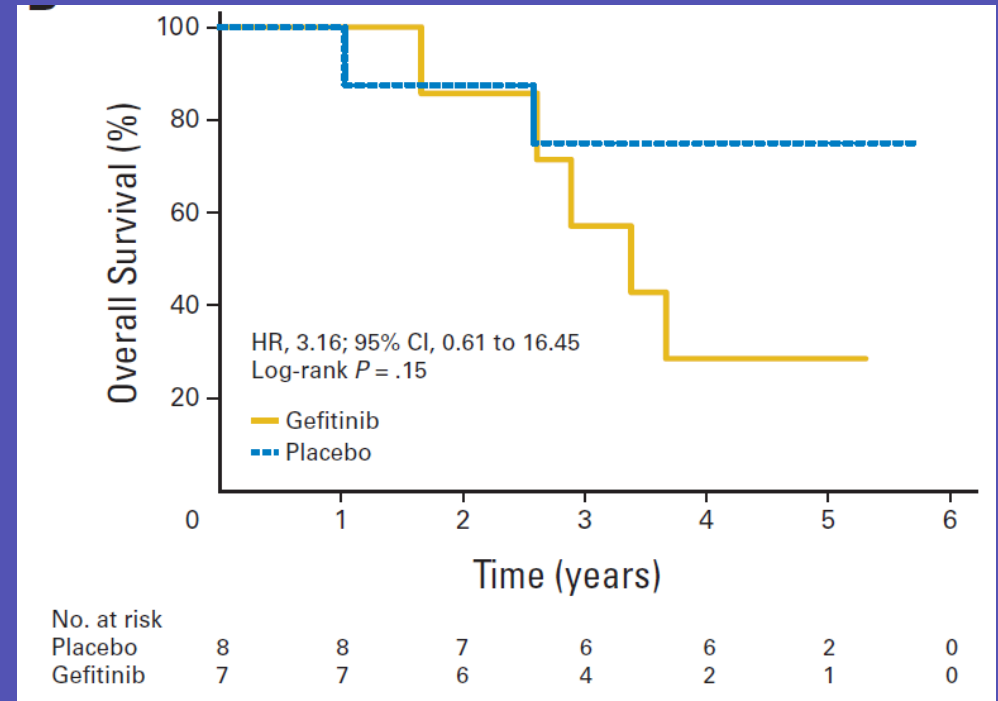
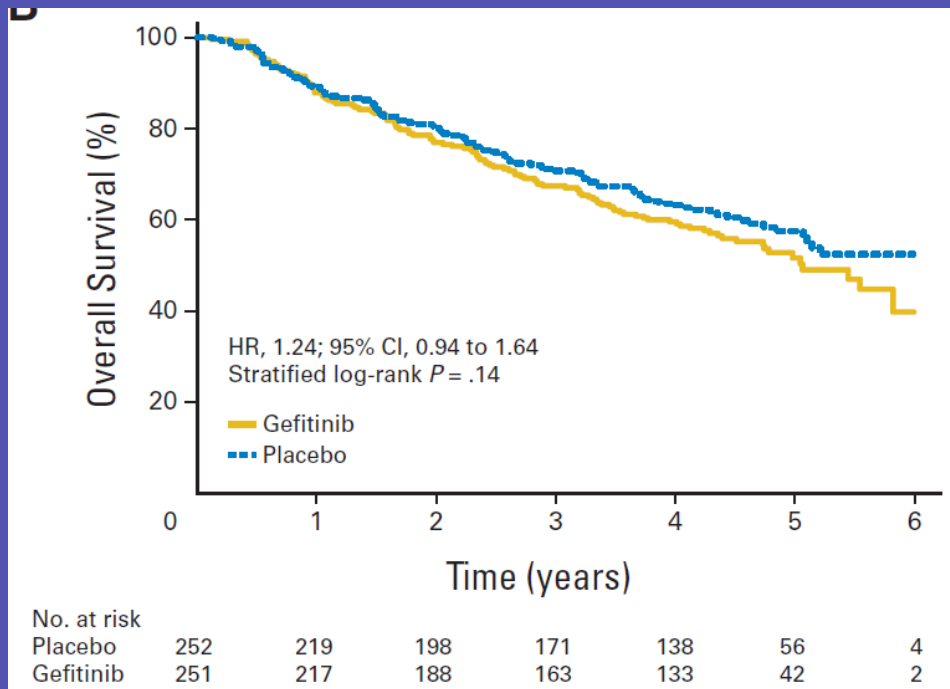
Référence	Chimiothérapie	N pts	% RO	p	SM	p	Phase
En rattrapage dans des populations non sélectionnées, versus soins de confort							
Shepherd (36)	Erlotinib	488	8,9%	S	6,7 m	S	III
	Placebo	243	1%		4,7 m		
Thatcher (37)	Géfitinib	1129	8%		5,6 m	0,09	III
	-	563	1,3%		5,1 m		
En rattrapage dans des populations non sélectionnées, versus chimiothérapie							
Maruyama (38)	Géfitinib	245	22,5%	0,009	11,5 m	0,33	III
	Docétaxel	244	12,8%		14 m		
Kim (39)	Géfitinib	723	9,1%	0,33	7,6m	NS	III
	Docétaxel	710	7,6%		8,0m		
Garassino (41)	Erlotinib	112	3%	0,001	5,4m	0,05	III
	Docétaxel	110	15,5%		8,2m		
Karampeazis (40)	Erlotinib	166	9%	0,47	10,1m	0,99	III
	Pémétréxed	166	11,4%		8,2m		
En rattrapage dans des populations « enrichies », versus chimiothérapie							
Sun (43)	Géfitinib	71	58,8%	<,001	22,2m	0,37	III
	Pémétréxed	70	22,4%		18,9m		
En rattrapage après échec d'un ITK d'EGFR							
Miller (42)	Afatinib	390	7%	0,007	10,8m	0,74	IIb/II I
	Placebo	195	0,5%		12,0m		

- Rôle limité pour un ITK d'EGFR en situation de rattrapage dans des populations non sélectionnées ou en l'absence de mutation

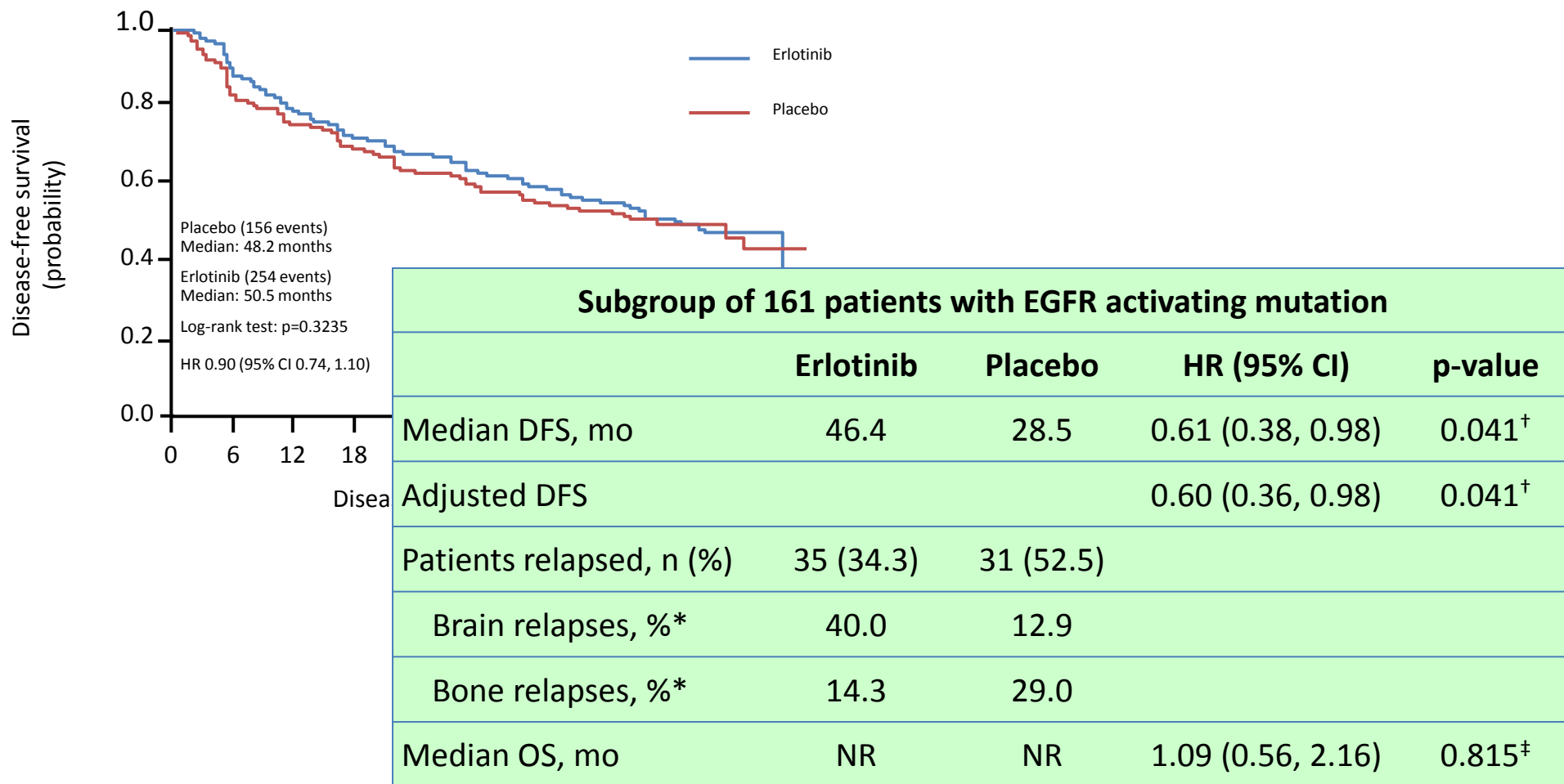
En cas de mutation activatrice d'EGFR

# Stades chirurgicaux

## Gefitinib Versus Placebo in Completely Resected Non-Small-Cell Lung Cancer: Results of the NCIC CTG BR19 Study



7501: A randomized, double-blind phase 3 trial of adjuvant erlotinib (E) versus placebo (P) following complete tumor resection with or without adjuvant chemotherapy in patients (pts) with stage IB-IIIa EGFR positive (IHC/FISH) non-small cell lung cancer (NSCLC): RADIANT results – Kelly K et al



# Stades locorégionaux

- Aucune donnée permettant de déterminer la place d'un ITK d'EGFR-TKI en cas de mutation activatrice chez les patients traités par CT-RT avec intention curative



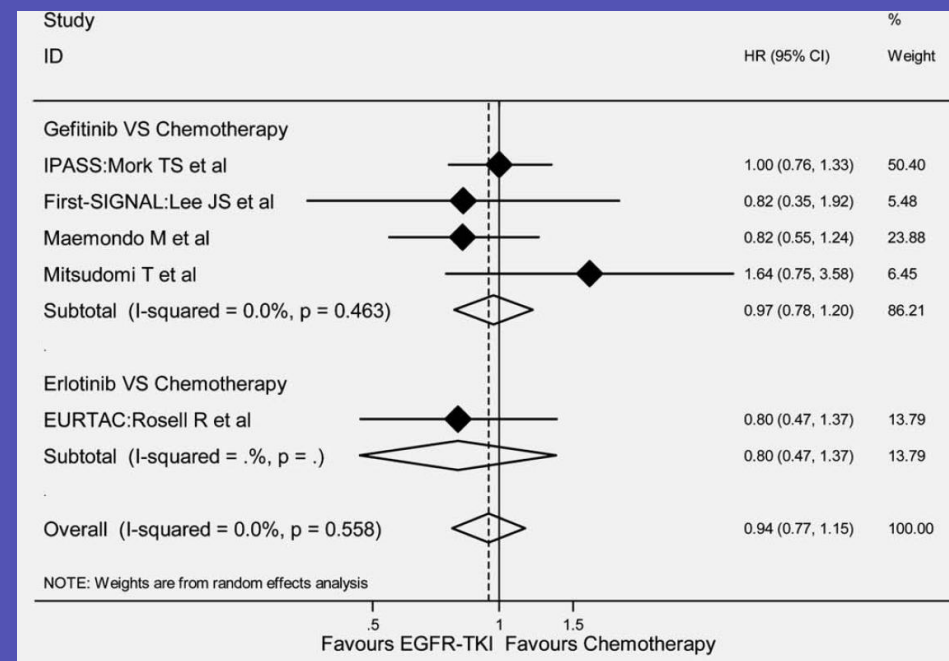
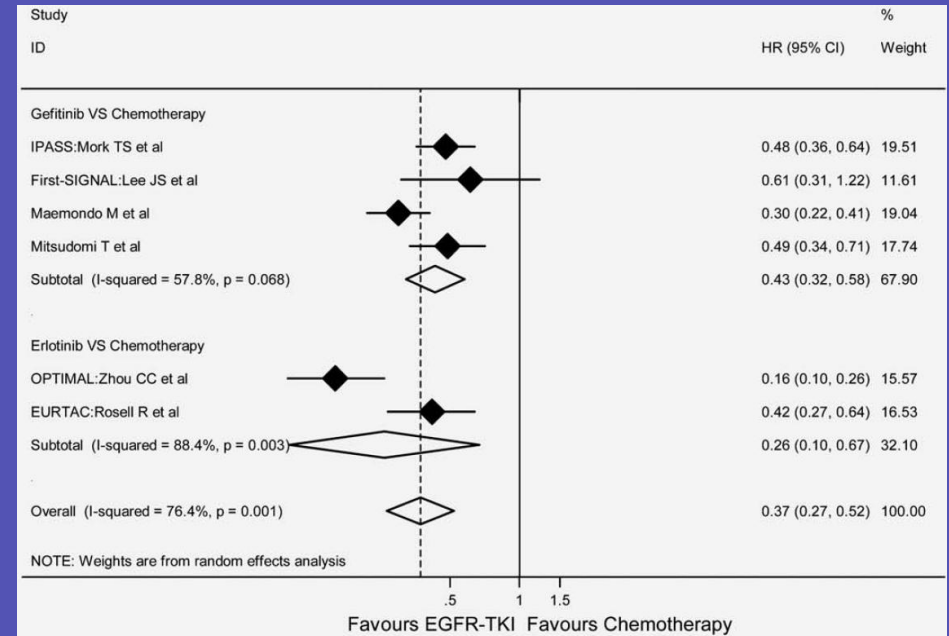
# Maladies avancées: études randomisées en 1ère ligne comparant un ITK d'EGFR à CT

- Amélioration du taux de réponse et de la SSP
- Pas d'effet sur la survie (cross-over)

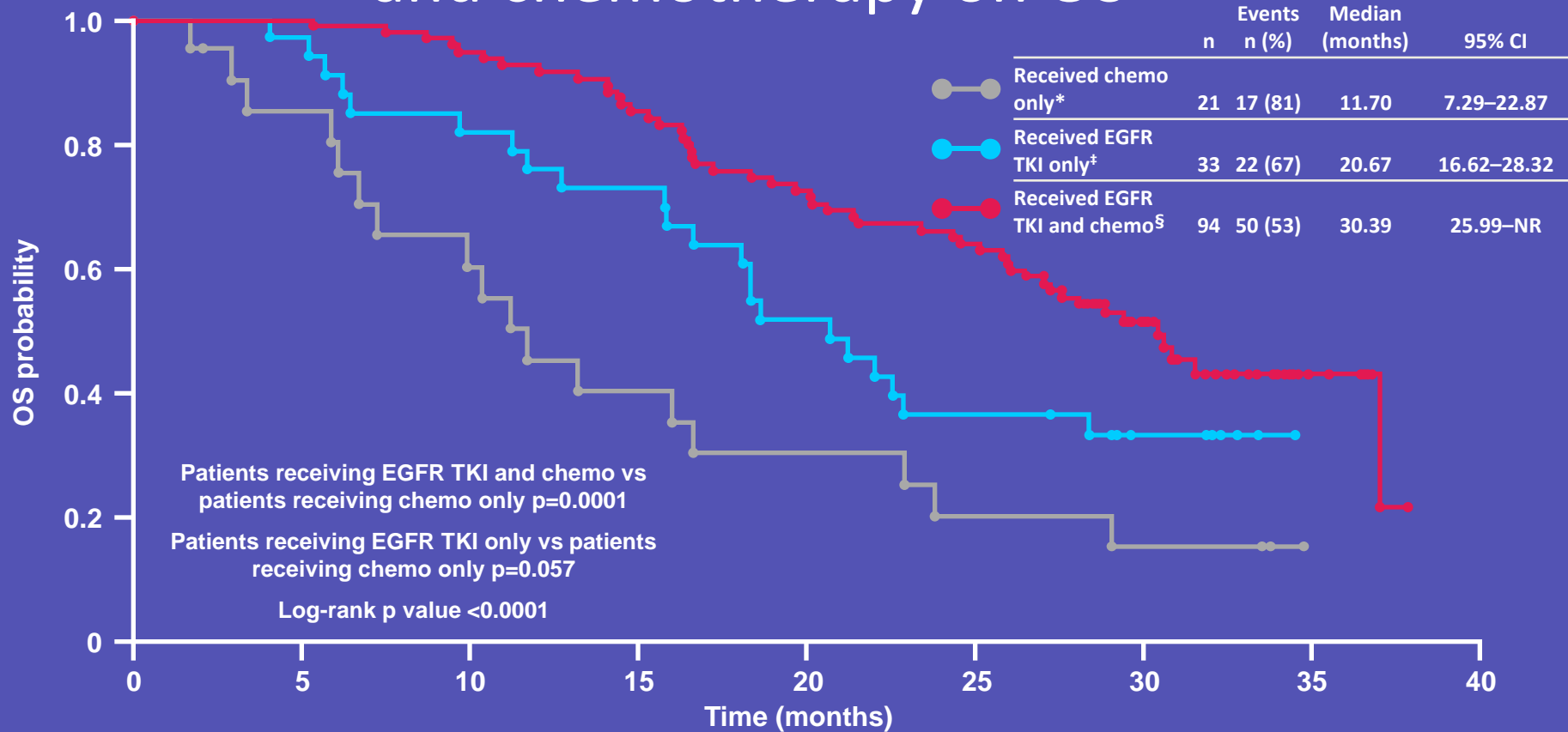
Maemondo, 2010 Inoue, 2013	Gefitinib	114	73,7%	S	27,7 m	0,48
	CBDCA + Pac	114	30,7%		26,6 m	
Mitsudomi, 2010	Gefitinib	86			Non atteint	0,21
	CDDP + Doc	86			30,9m	
Rosell, 2012	Erlotinib	86	58 %	S	19,3 m	NS
	Standard CT	87	15 %		19,5 m	
Zhou, 2011 (ASCO 2012)	Erlotinib	83	82 %	S	22,7m	0,69
	Gem + CBDCA	82	36 %		28,9m	
Sequist, 2013	Afatinib	230	56%	0,001	HR 1,12	0,60
	CDDP-PEM	115	23%			
Wu, 2014	Afatinib	242	67%	<0,001	22,1m	0,76
	CDDP-GEM	122	23%		22,2m	

# Cross-over

	ITK EGFR en rattrapage
Inoue	98.2%
Rosell	76%
Mitsudomi	59%
Lux Lung 3	75%
Lux Lung 6	56%



# Supplementary analysis: influence of EGFR TKI and chemotherapy on OS



## Patients at risk

	0	5	10	15	20	25	30	35	40
Patients receiving chemo only*	21	17	12	8	6	4	3	0	0
Patients receiving EGFR TKI only†	33	32	27	24	17	12	7	0	0
Patients receiving EGFR TKI and chemo‡	94	94	89	80	68	60	28	6	6

\*Chemo only, no EGFR TKI: patients from the GC arm who had no further treatment (n=16) or further chemotherapy (n=5)

†EGFR TKI only, no chemo: patients from the erlotinib arm who are still on treatment (n=7), had no further treatment (n=25) and who were re-challenged (n=1)

‡EGFR TKI and chemo: patients from the erlotinib arm who switched to chemo (n=43), patients from the GC arm who switched to erlotinib in any line (n=51)



- T790M et autres mutations associées avec résistance

→ ITK d'EGFR de 3<sup>ème</sup> génération (AZD9291)?



G719C  
G719S  
G719A  
V689M  
N700D  
E709K/C  
S720P

(5%)

ΔE746-A750

V765A

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

(40–45%)

- Mutations activatrices « classiques »
  - Del exon 19
  - Insertion exon 21 (L858R)

- Autres mutations activatrices

→ ITK d'EGFR

(45%)

Mutations asso  
with drug sensi

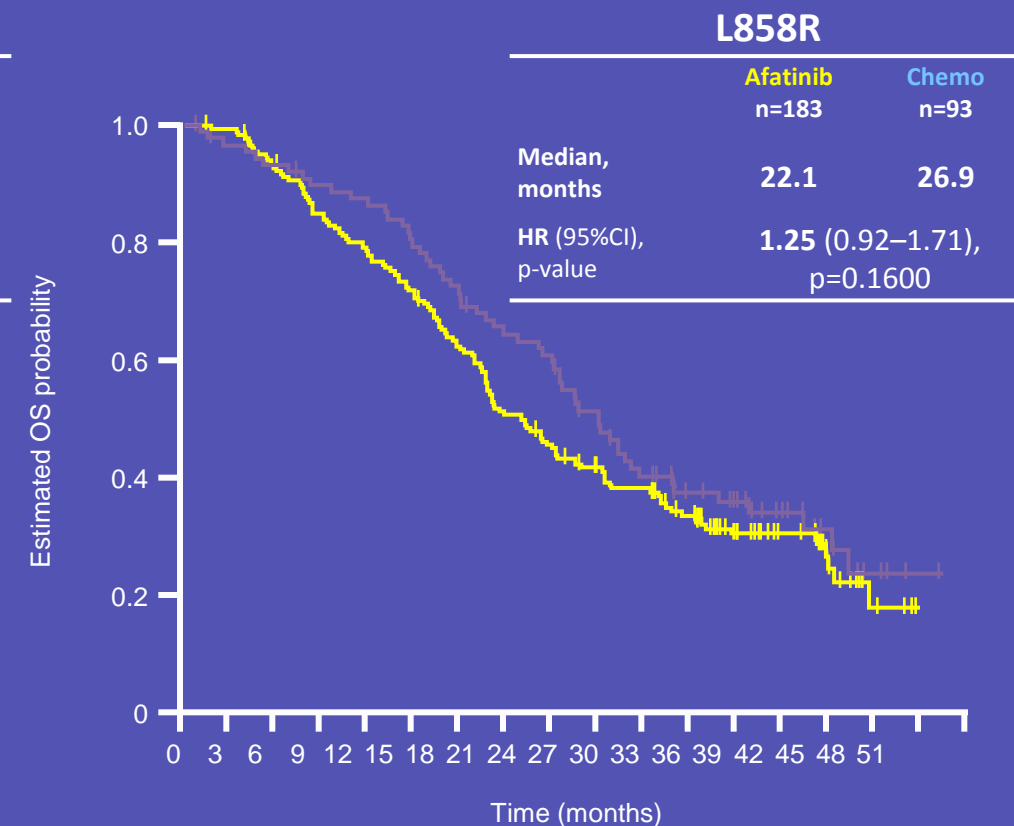
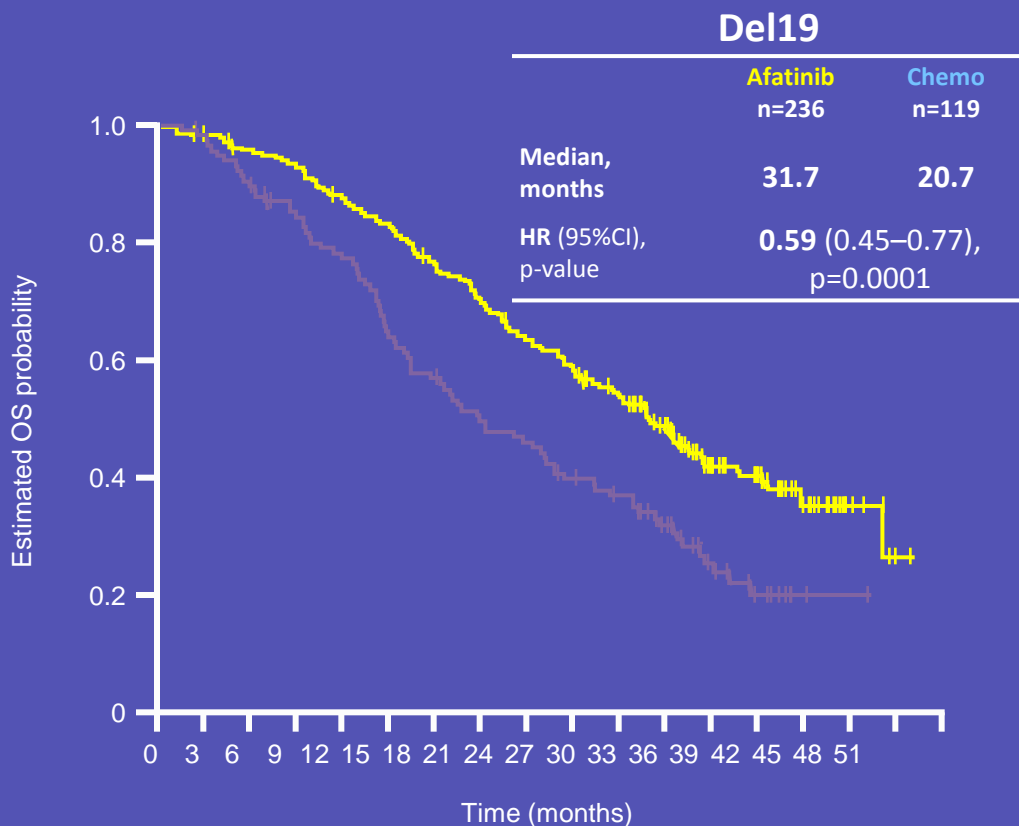
# Mutations activatrices classique: Sensibilité différente aux ITK d'EGFR?

- Del exon 19 semble plus sensible aux ITK de 1<sup>ère</sup>-2<sup>de</sup> génération mais résultats conflictuels

Effet de l'ITK d'EGFR sur la SSP en fonction du statut Del exon19/L858R		
	TKI	p
Inoue	Gefitinib	0,181
Mitsudomi	Gefitinib	0,68
Rosell	Erlotinib	0,07
Zhou	Erlotinib	0,054
Sequist	Afatinib	0,01

# Combined OS analysis: mutation categories

## Lux Lung 3 and 6



No of patients

	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45	48	51
Afatinib	236	230	223	217	202	192	173	160	145	131	117	90	50	38	22	6	1	0
Chemo	119	113	103	95	87	72	63	55	51	43	38	27	14	9	1	1	0	0

No of patients

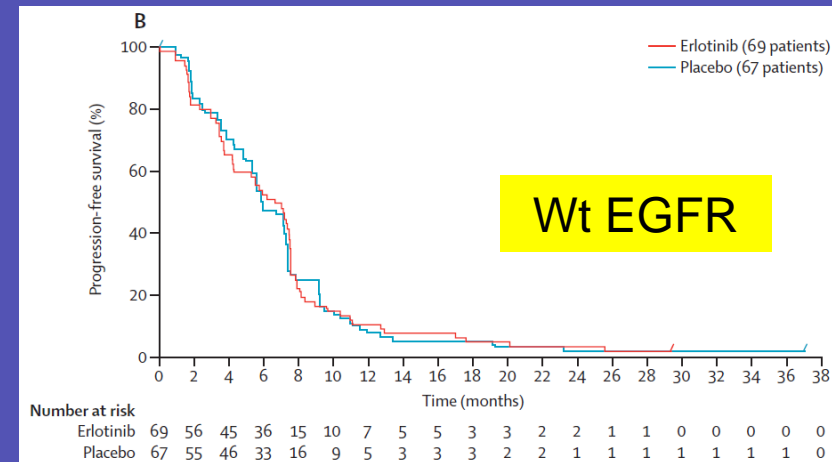
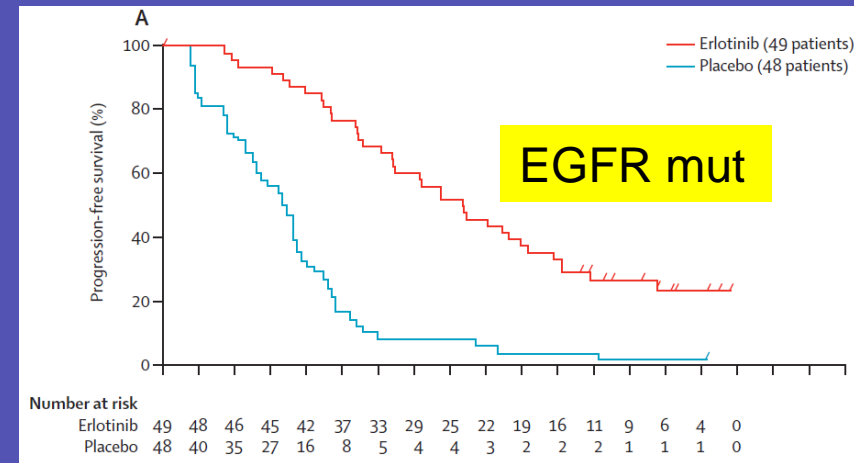
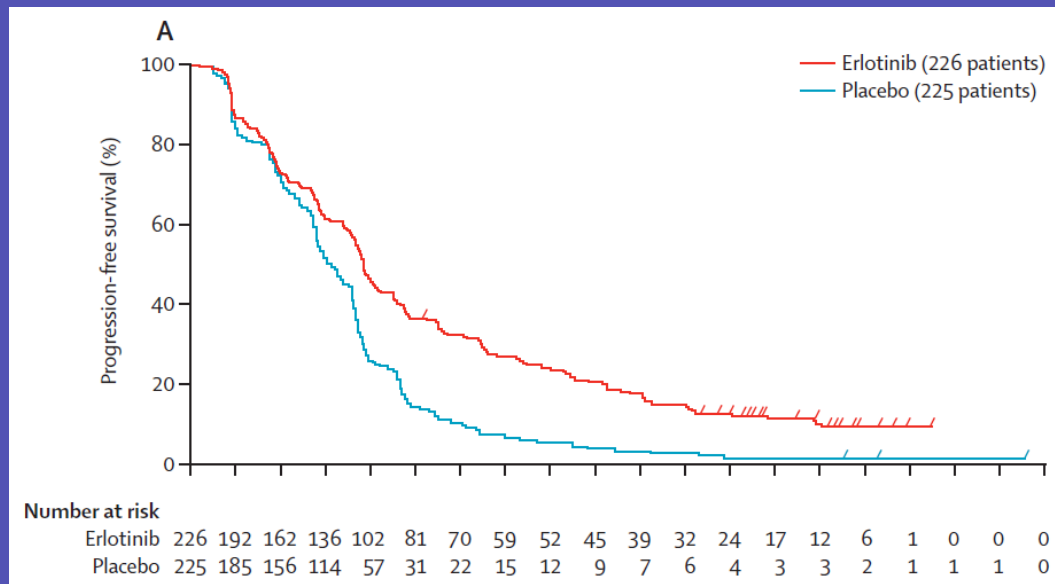
	0	3	6	9	12	15	18	21	24	27	30	33	36	39	42	45	48	51
Afatinib	183	181	167	154	141	128	111	91	80	70	64	51	27	20	11	3	0	0
Chemo	93	86	82	78	75	69	61	55	50	40	32	25	20	14	9	4	1	0

With the courtesy of Prof James Chih-Hsin Yang

# Y a-t-il un intérêt à combiner séquentiellement CT et ITK EGFR?

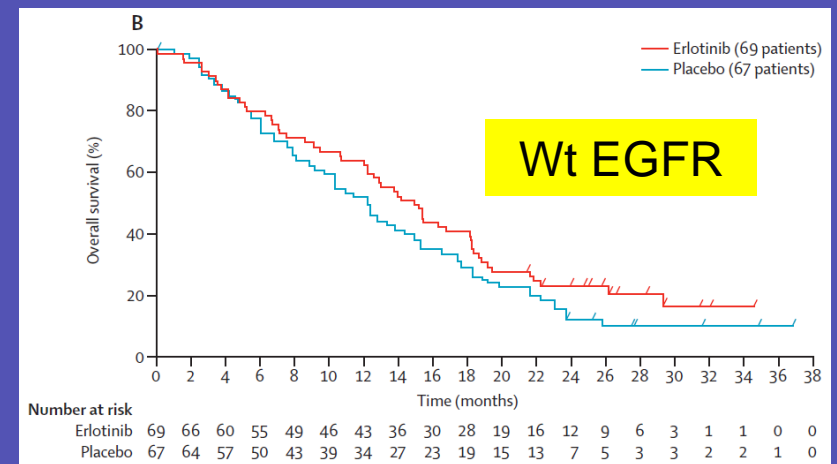
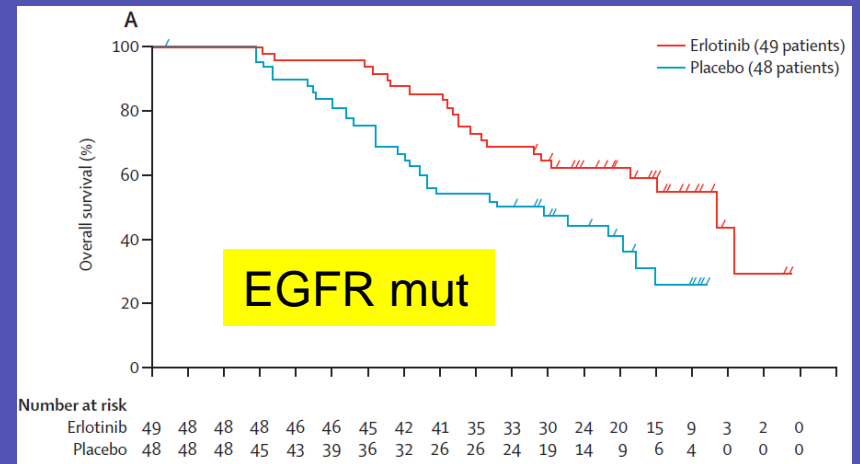
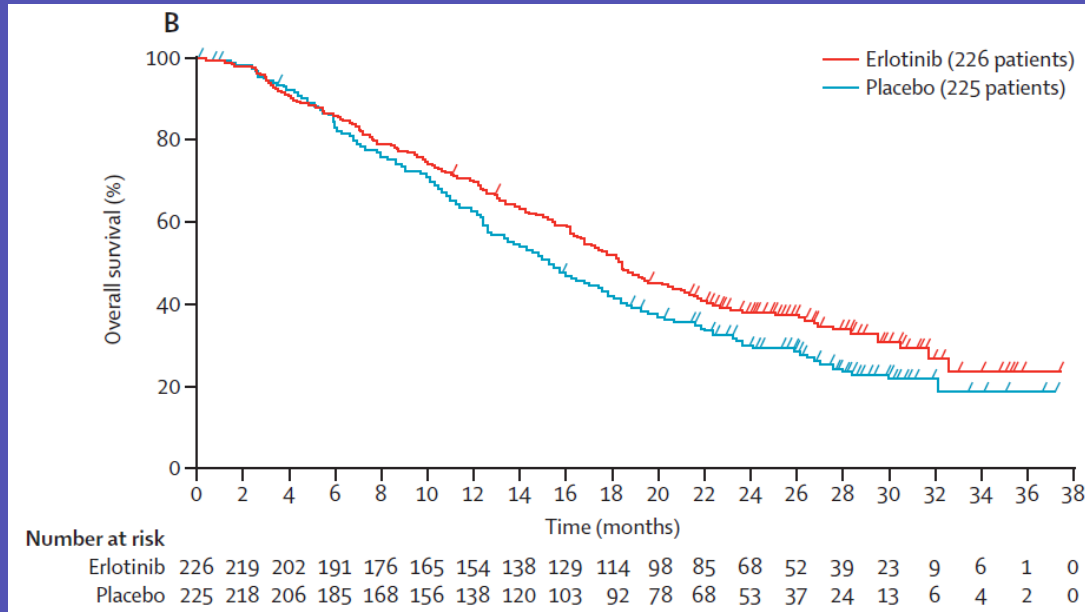
- Pas d'AMM!
- EGFR non muté: échec en concomitant en 1<sup>ère</sup> ligne
- CT et ITK intercalés: amélioration SSP

# Intercalated combination of chemotherapy and erlotinib for patients with advanced stage non-small-cell lung cancer (FASTACT-2): a randomised, double-blind trial

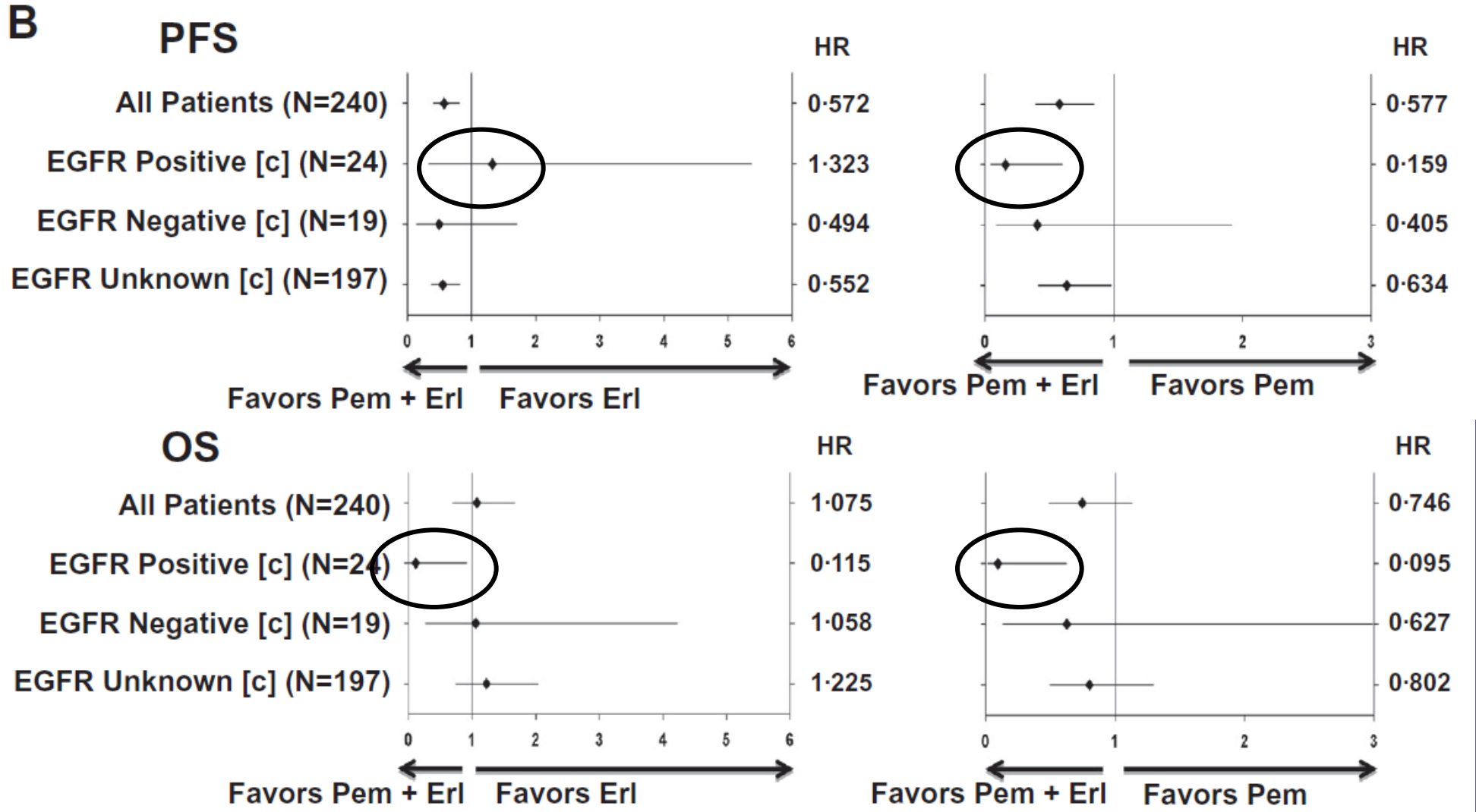




# Intercalated combination of chemotherapy and erlotinib for patients with advanced stage non-small-cell lung cancer (FASTACT-2): a randomised, double-blind trial



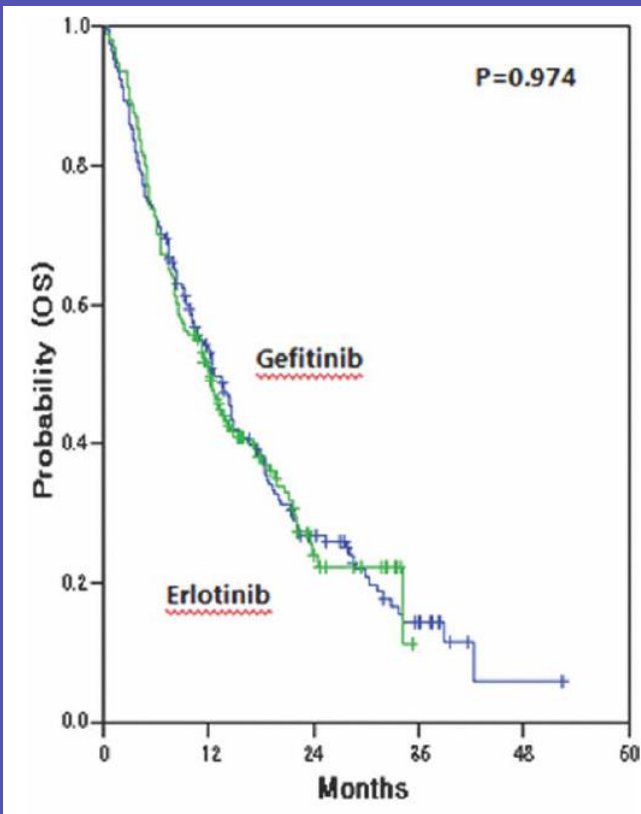
Three-arm randomised controlled phase 2 study comparing pemetrexed and erlotinib to either pemetrexed or erlotinib alone as second-line treatment for never-smokers with non-squamous non-small cell lung cancer ☆



# Quel ITK?

## Comparison of Gefitinib Versus Erlotinib in Patients With Nonsmall Cell Lung Cancer Who Failed Previous Chemotherapy

Seung Tae Kim, MD; Jeeyun Lee, MD, PhD; Jeong-hoon Kim, MD; Young-Woong Won, MD; Jong-mu Sun, MD; Jina Yun, MD; Yeon Hee Park, MD, PhD; Jin Seok Ahn, MD, PhD; Keunchil Park, MD, PhD; and Myung-Ju Ahn, MD, PhD



Cancer 2010

## Comparison of gefitinib and erlotinib in advanced NSCLC and the effect of EGFR mutations

Jenn-Yu Wu<sup>a</sup>, Shang-Gin Wu<sup>a</sup>, Chih-Hsin Yang<sup>b</sup>, Yih-Leong Chang<sup>c</sup>, Yeun-Chung Chang<sup>d</sup>, Ya-Chieh Hsu<sup>e</sup>, Jin-Yuan Shih<sup>e,\*</sup>, Pan-Chyr Yang<sup>e</sup>

	Réponse	p	Survie	p
Erlotinib	75,4%	0,07	15m	0,82
Géfitinib	61,9%		18,1m	

Lung Cancer 2011

**Erlotinib has better efficacy than gefitinib in adenocarcinoma patients without *EGFR*-activating mutations, but similar efficacy in patients with *EGFR*-activating mutations**

EXPERIMENTAL AND THERAPEUTIC MEDICINE 3: 207-213, 2012

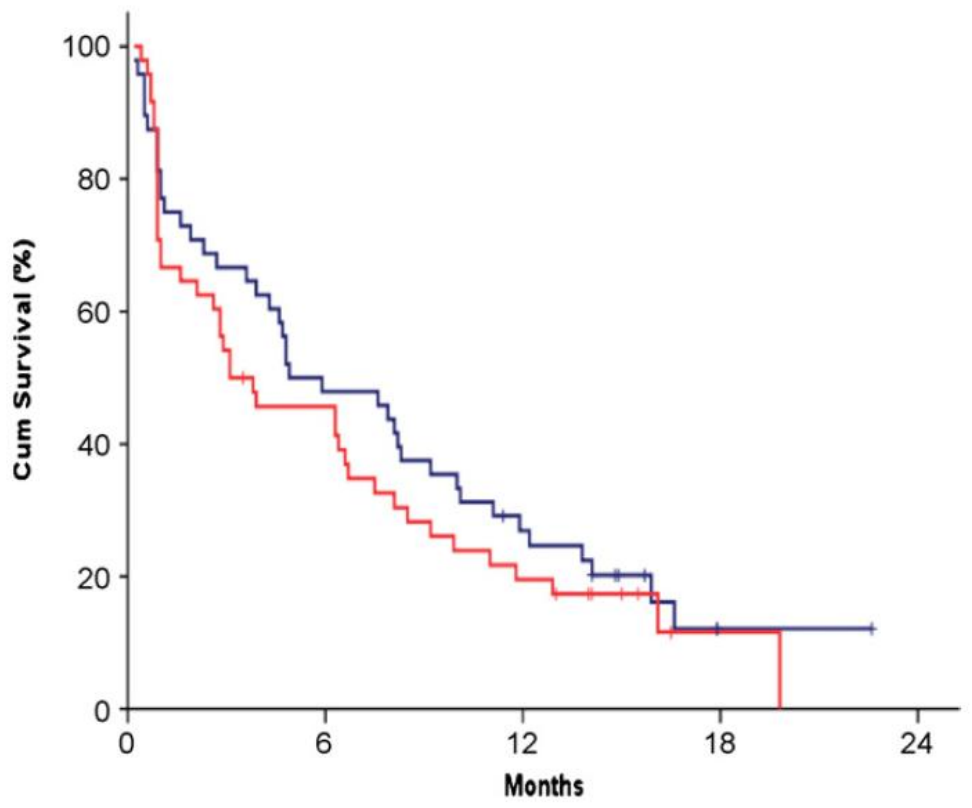
# Randomized phase II study of gefitinib versus erlotinib in patients with advanced non-small cell lung cancer who failed previous chemotherapy<sup>☆</sup>

Seung Tae Kim<sup>a,d,1</sup>, Ji Eun Uhm<sup>a,1</sup>, Jeeyun Lee<sup>a</sup>, Jong-mu Sun<sup>a</sup>, Insuk Sohn<sup>b</sup>, Seon Woo Kim<sup>b</sup>, Sin-Ho Jung<sup>c</sup>, Yeon Hee Park<sup>a</sup>, Jin Seok Ahn<sup>a</sup>, Keunchil Park<sup>a</sup>, Myung-Ju Ahn<sup>a,\*</sup>

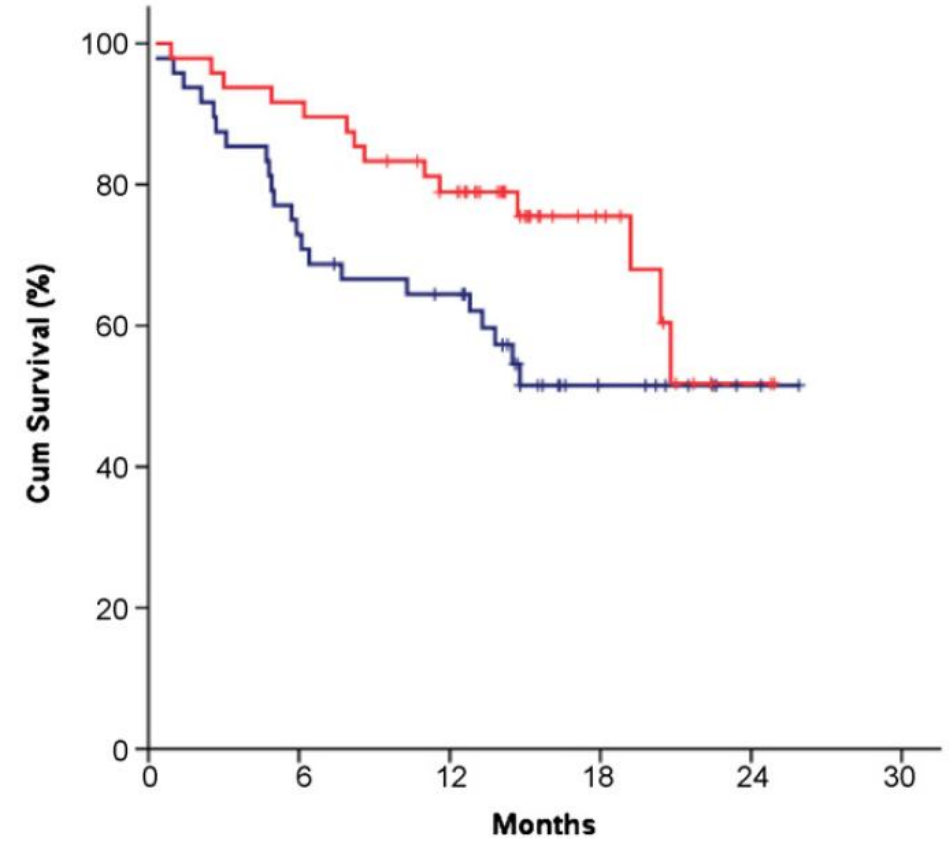
Best response rate and disease control rate of treatment groups.

Gefitinib (n = 48)		Erlotinib (n = 48)	
N	%	N	%
	2.1		
	45.8		
	25.0		
	25.0		
	2.1		
	47.9		
	72.9		

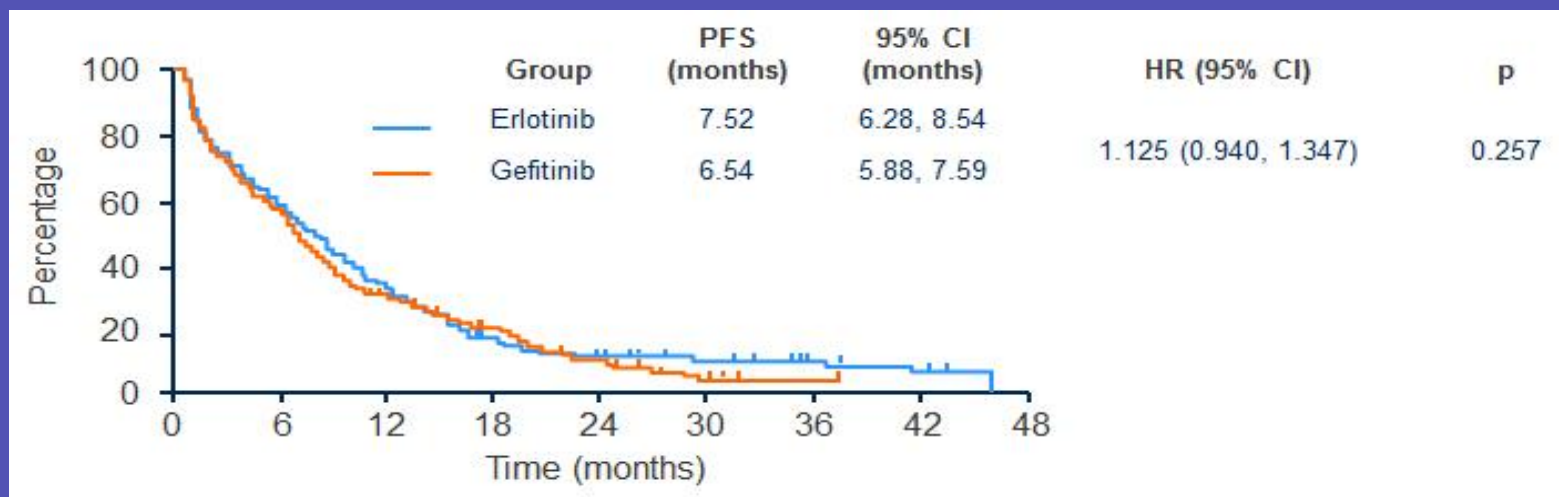
(A) Progression free survival according to treatment groups



(B) Overall survival according to treatment group

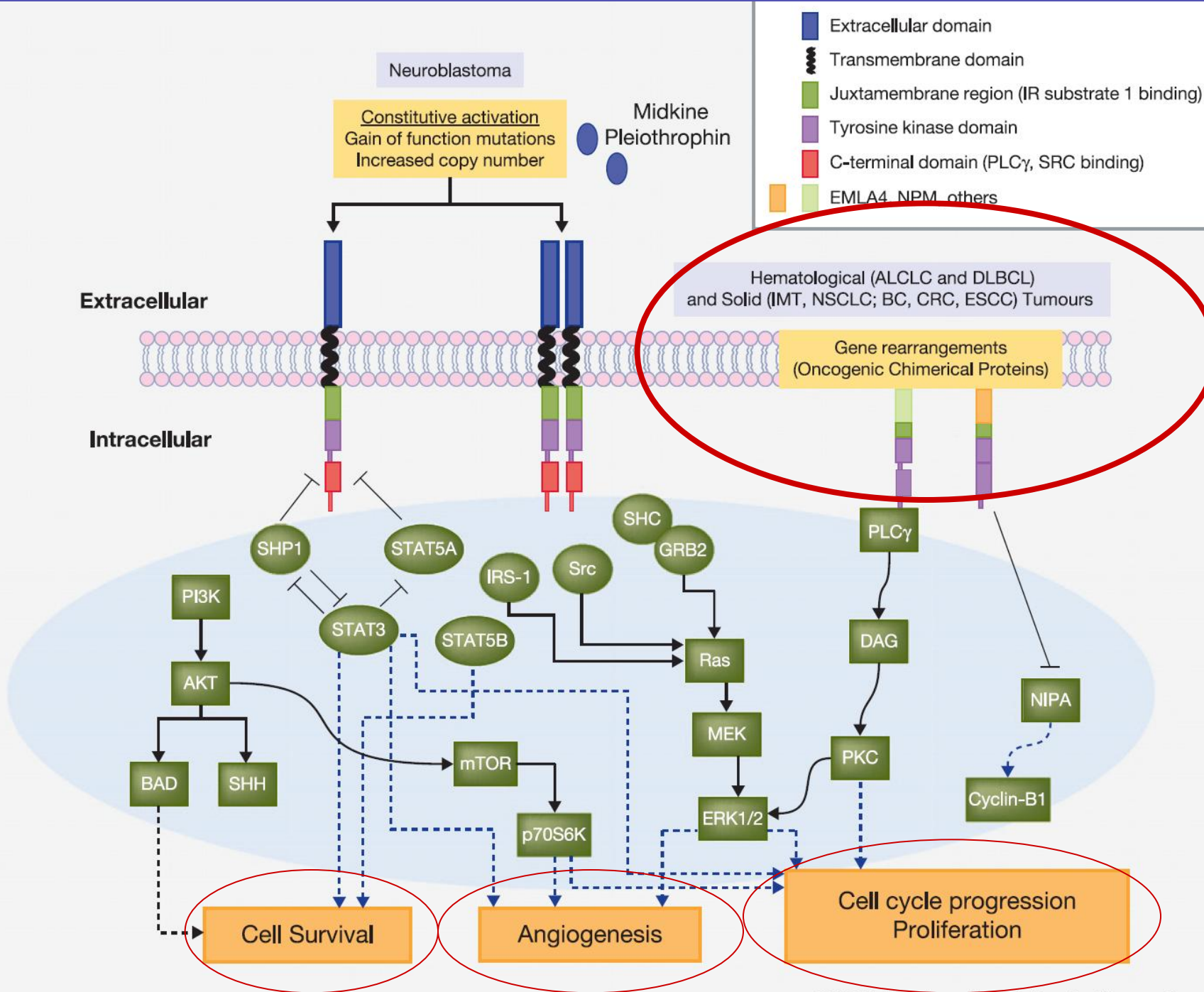


8041: Randomized phase III study comparing gefitinib (G) with erlotinib (E) in patients (pts) with previously treated advanced lung adenocarcinoma (LA):  
WJOG 5108L – Katakami N et al



- OS: 24,5 vs 22,8 m (HR 1,038, IC 95% 0,833-1,394)
- SSP et OS médianes en cas de mutation activatrice (erlotinib vs gefitinib)
  - SSP: 10,1 vs 8,9 m (p=0,532)
  - OS: 32,0 vs 26,6 m (p=0,111)

EML4-ALK



Prot aNle conserve domaine intracellulaire de ALK tandis que le partenaire intervient dans la dimérisation → dimérisation indépendante du ligand et activation constitutive de la kinase

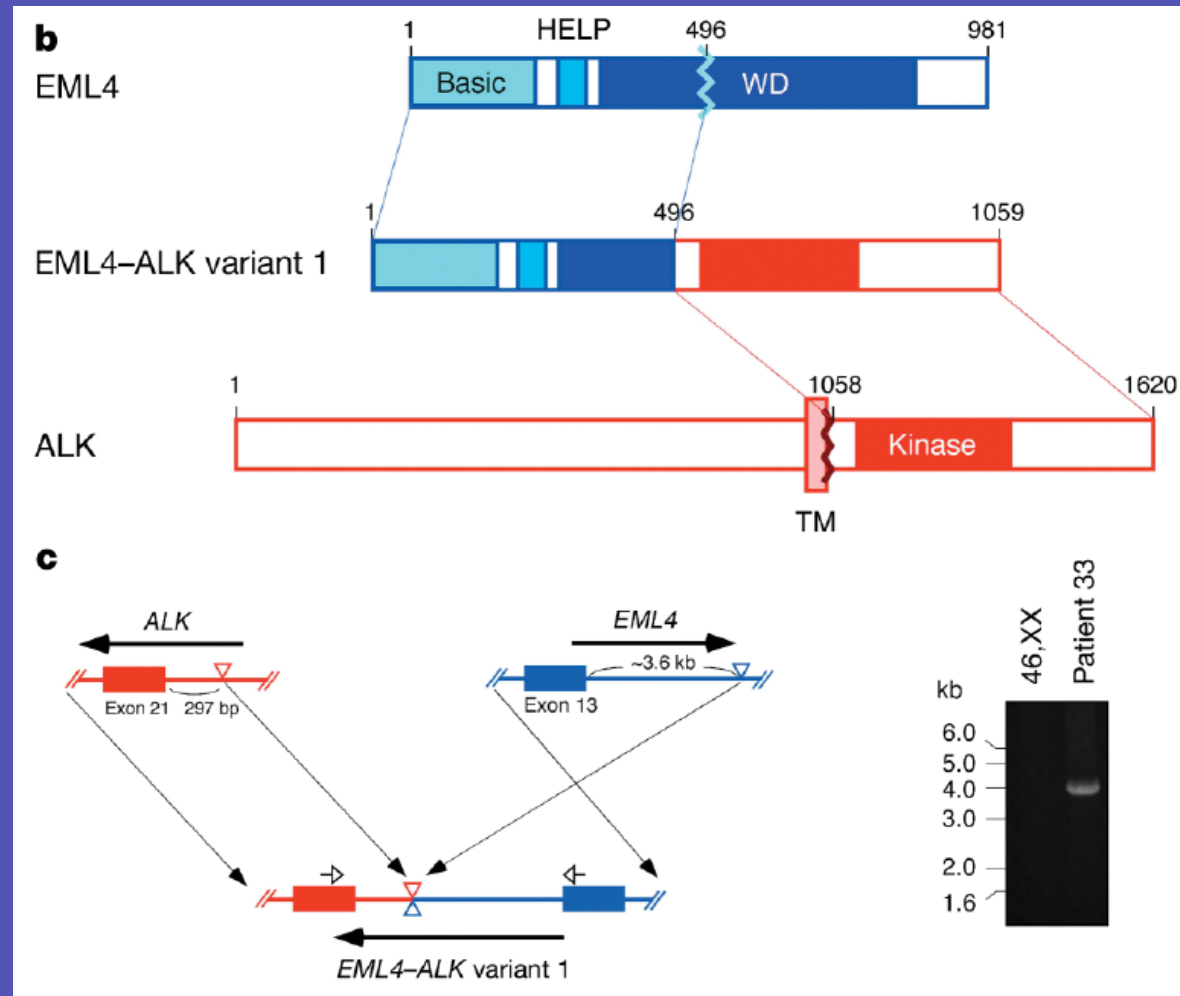
"ONCOGENE ADDICTION"

Grande et al; Mol Cancer Ther 2011

# 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>

echinoderm microtubule-associated protein-like 4

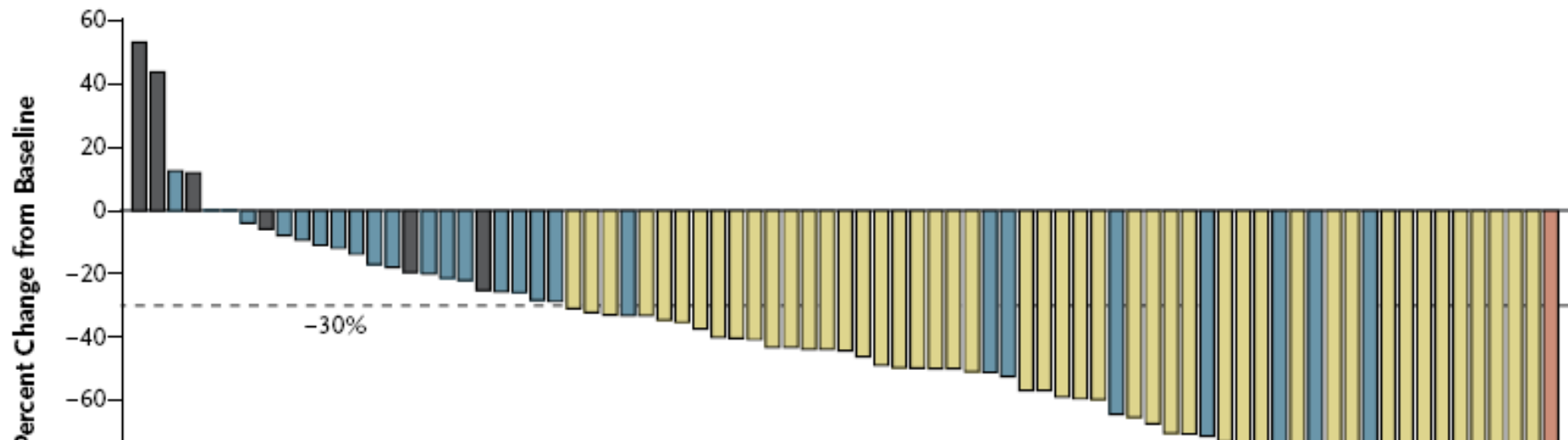




# Crizotinib

Kwak, NEJM 2010

A Percent Change in Tumor Burden



Référence	Ligne	N patients	Phase	RR
Camidge, 2012	Toutes	143 crizotinib naïfs 1ère ligne Rattrapage(2->3 lignes)	I	60,8% 63,6% 58,7-64,5%
Kim, 2012	Rattrapage	803 crizotinib naïfs 261 "population mature"	II	46% 60%

# Crizotinib versus Chemotherapy in Advanced ALK-Positive Lung Cancer

## Key patient inclusion criteria

- ALK+ by central FISH testing<sup>a</sup>
  - Stage IIIB/IV NSCLC
  - 1 prior chemotherapy (platinum-based)
  - ECOG PS 0-2
  - Measurable disease
- (n=318)



Crizotinib 250 mg BID, 21-day cycle (n=159)

## Stratification

- ECOG PS, brain metastases, prior EGFR TKI

Pemetrexed 500 mg/m<sup>2</sup> or docetaxel 75 mg/m<sup>2</sup> IV, day 1, 21-day cycle (n=159)

Crossover to crizotinib on profile 1005  
Secondary endpoints

- ORR, DCR, DR, OS, safety, pt reported outcomes

## Primary endpoint

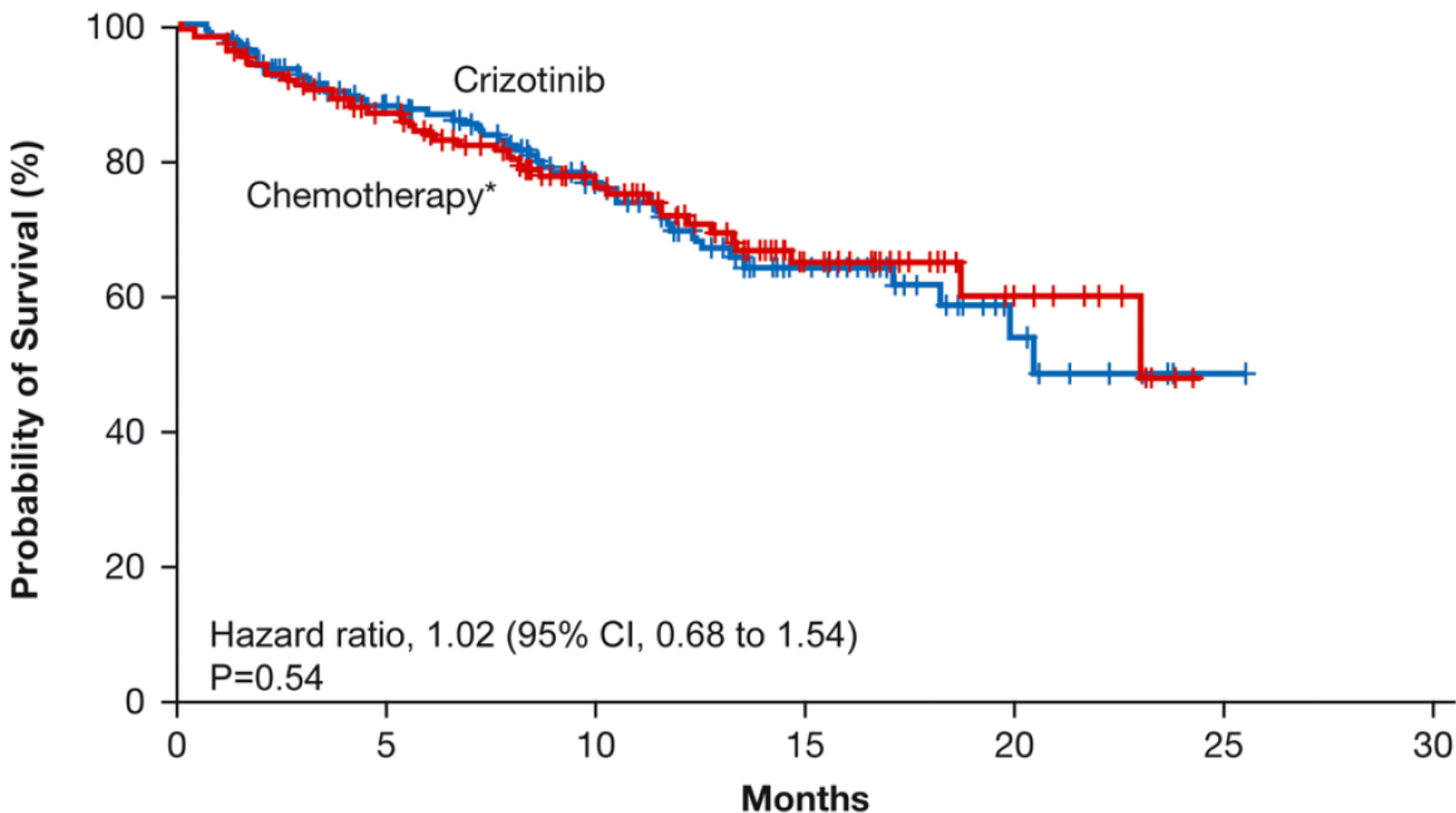
- PFS (RECIST 1.1 by independent radiology review)

<sup>a</sup>ALK status determined using standard ALK break-apart FISH assay

1.0

Crizotinib (n=173)      Chemotherapy (n=174)

Probability of survival



0.43

No. at risk

**No. at Risk**

	0	5	10	15	20	25	30
Crizotinib	173	129	83	37	11	1	0
Chemotherapy	174	129	84	34	10	0	0
<b>No. at risk</b>							
Crizotinib	172	93	38	11	2	0	
Pemetrexed	99	36	12	3	1	0	
Docetaxel	72	13	3	1	0		

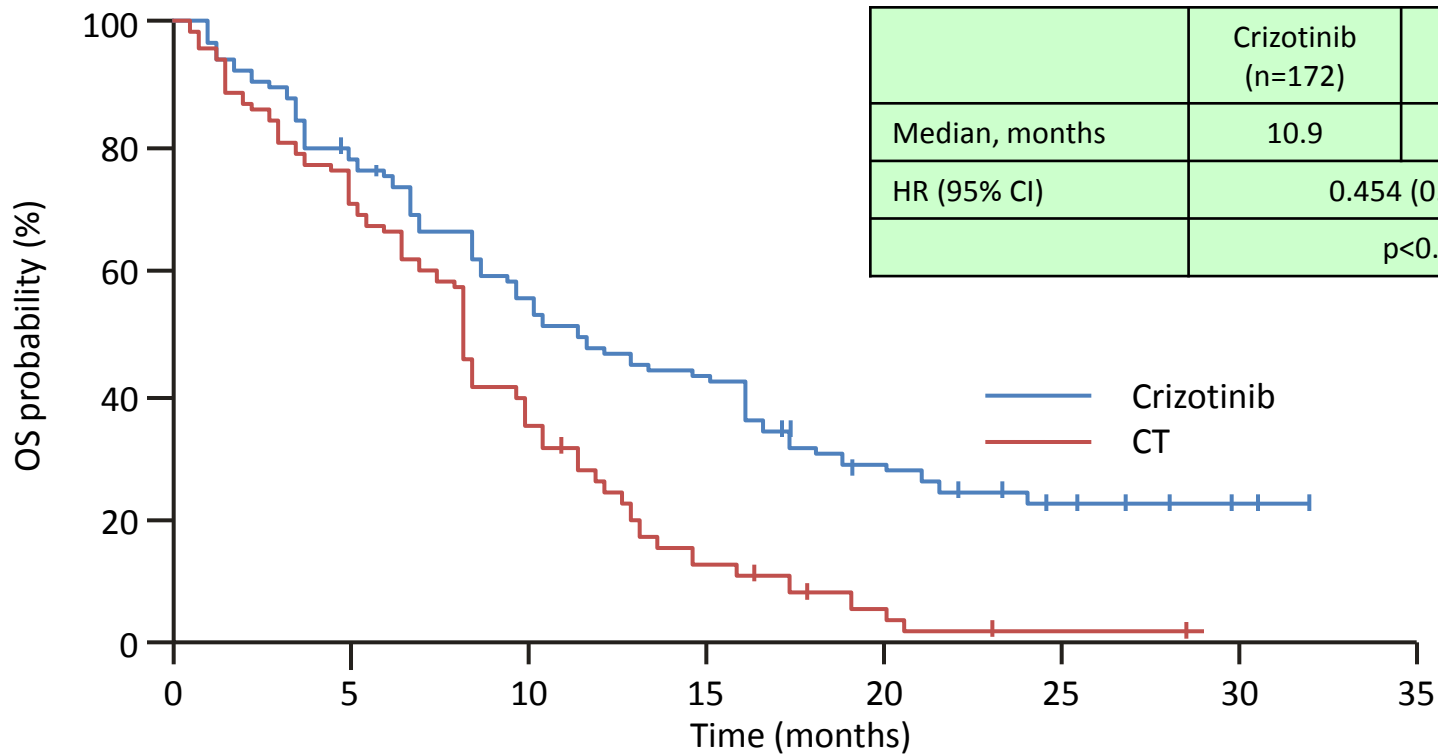
\*vs crizotinib

# 8002: First-line crizotinib versus pemetrexed-cisplatin or pemetrexed-carboplatin in patients (pts) with advanced *ALK*-positive non-squamous non-small cell lung cancer (NSCLC): results of a phase III study (PROFILE 1014)

- Key results**

- Addition of crizotinib significantly improved PFS but not OS compared with CT alone

## PFS



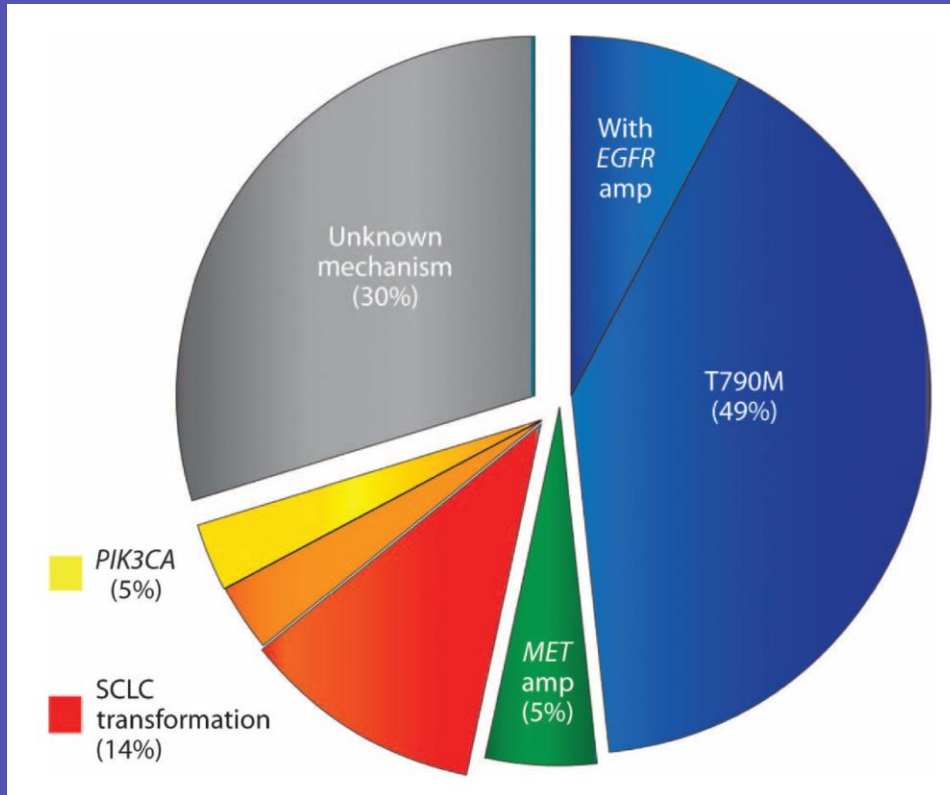
No. at risk  
Crizotinib  
CT

	0	5	10	15	20	25	30	35
Crizotinib	172	120	65	38	19	7	1	0
CT	171	105	36	12	2	1	0	0

# Nouveaux inhibiteurs de ALK

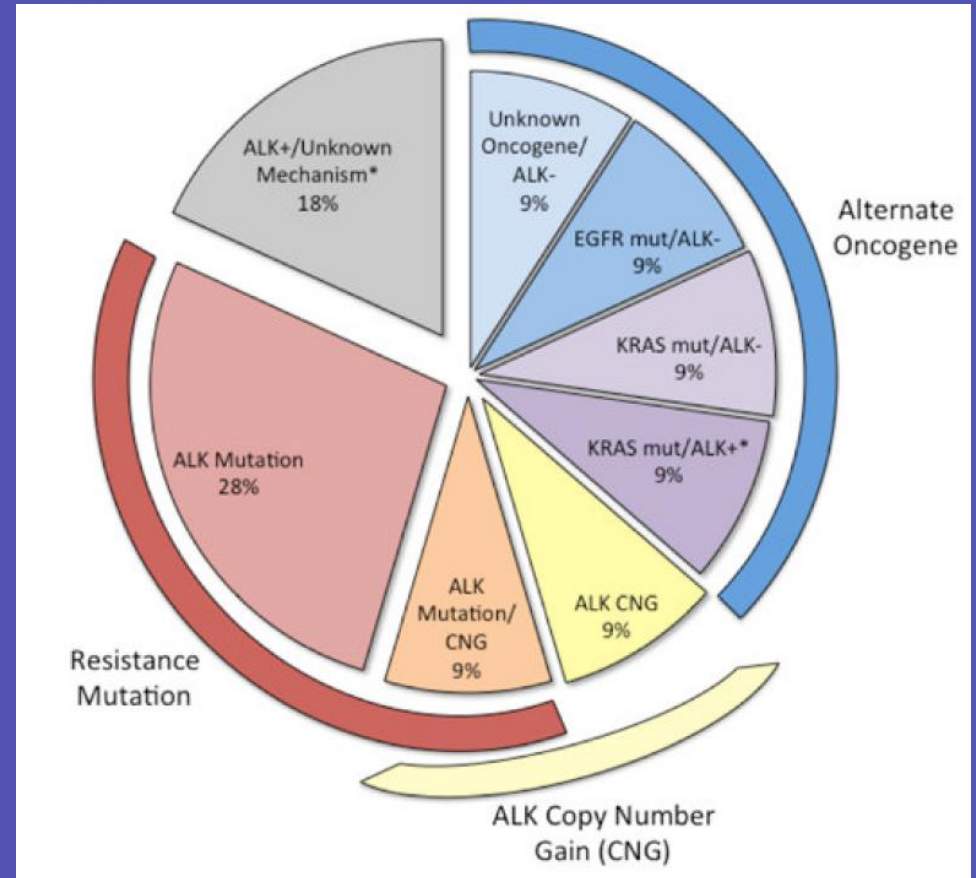
Référence	Molécule	Phase	RR
Kim, 2014	LDK378	I/II	58,5%
Kiura, 2012	CH5424802	I	85%
Nishio, 2012	CH5424802	II	85%
Gettinger, 2012	AP26113	I/II	73%
Felip, 2012	AUY922	II	32%

# Problèmes des résistances



EGFR

Sequist et al, Sci Transl Med 2011



ALK

Doebele et al, Clin Cancer Res 2012

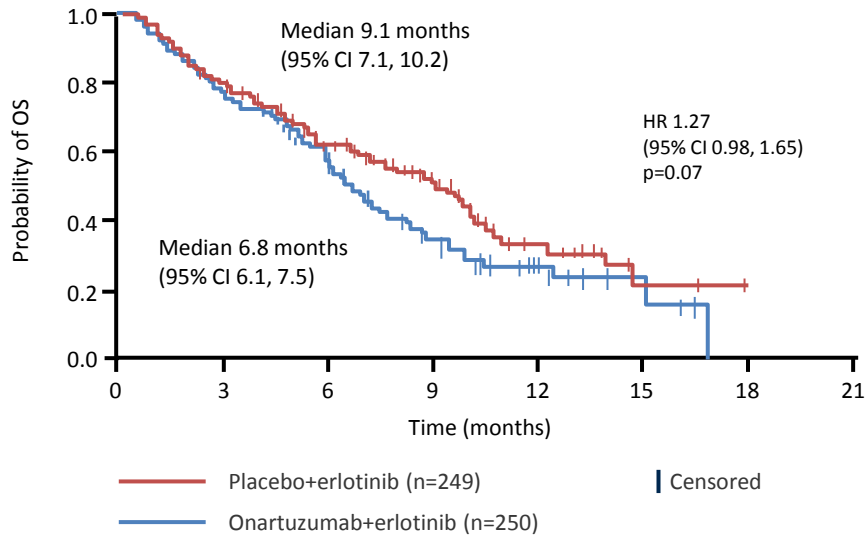
# Quelle option à la progression ?

Biopsier site de progression pour objectiver type de résistance et adapter traitement

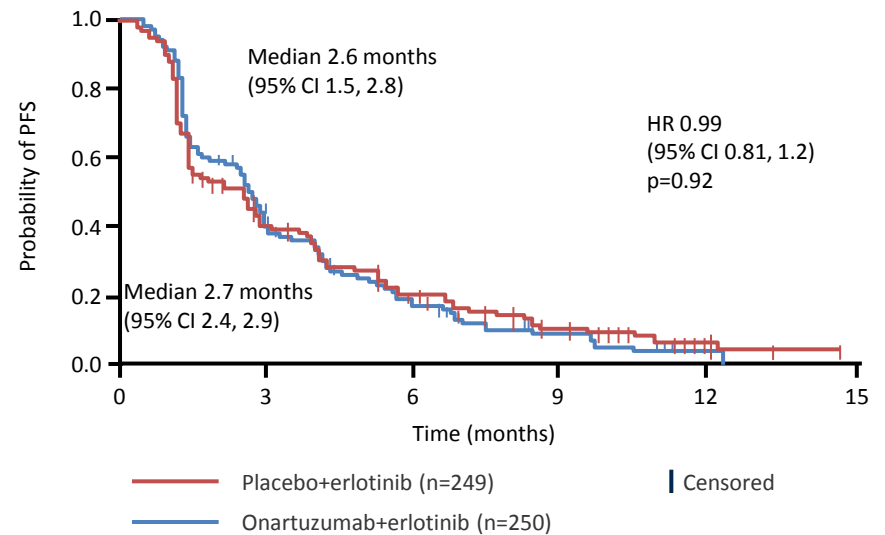
- EGFR
- ITK 3<sup>ème</sup> génération actif sur T790M
  - AZD9291
  - CO-1686
- Inhibiteur cMET
- ...
- ALK
- Inhibiteur 2<sup>de</sup> génération
  - Céritinib
  - CH5424802
  - AP26113
  - AUY922
- ...

# 8000: Onartuzumab plus erlotinib versus erlotinib in previously treated stage IIIb or IV NSCLC: Results from the pivotal phase III randomized, multicenter, placebo-controlled METLung (OAM4971g) global trial

## OS



## PFS

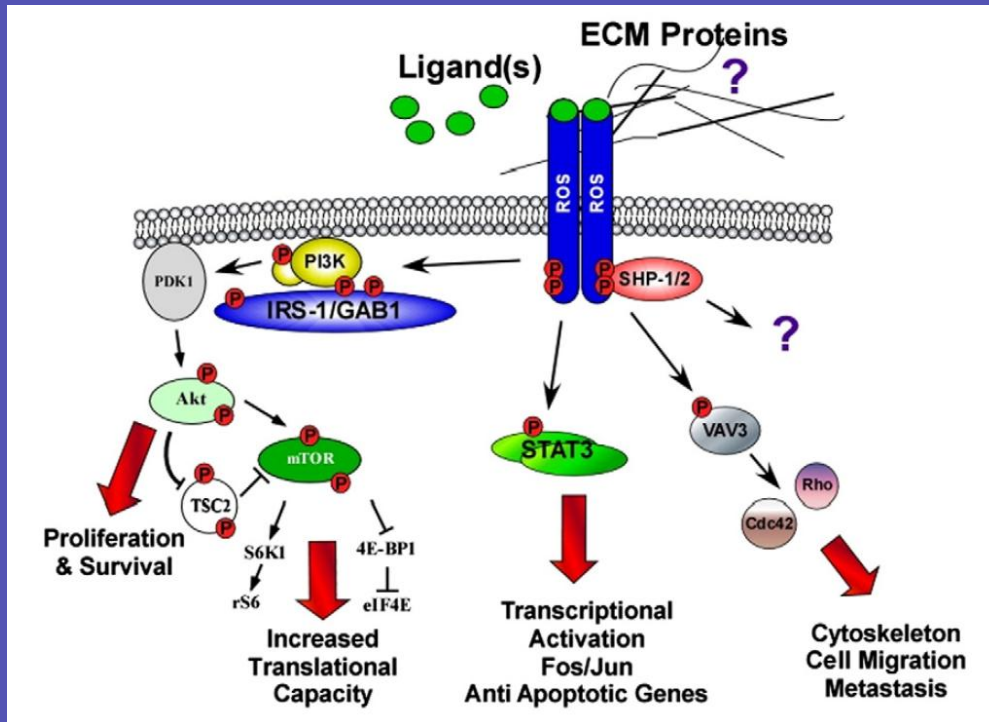


ORR 8.8% placebo+erlotinib vs. 6.4% onartuzumab+erlotinib



# 4. Futur des thérapies ciblées

# ROS1



- Phase I
- 14 patients avec translocation ROS-1 translocation traités par crizotinib
- 9 (64%) réponse objective

# KRAS

## Selumetinib plus docetaxel for *KRAS*-mutant advanced non-small-cell lung cancer: a randomised, multicentre, placebo-controlled, phase 2 study

### Facteur prédictif de résistance aux ITK

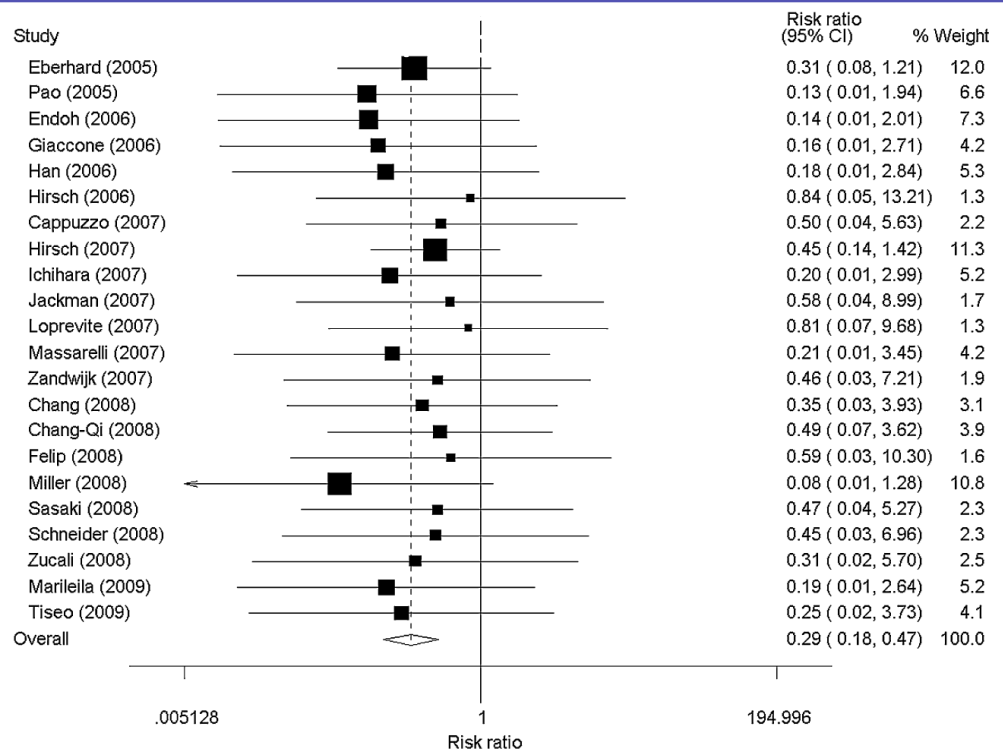
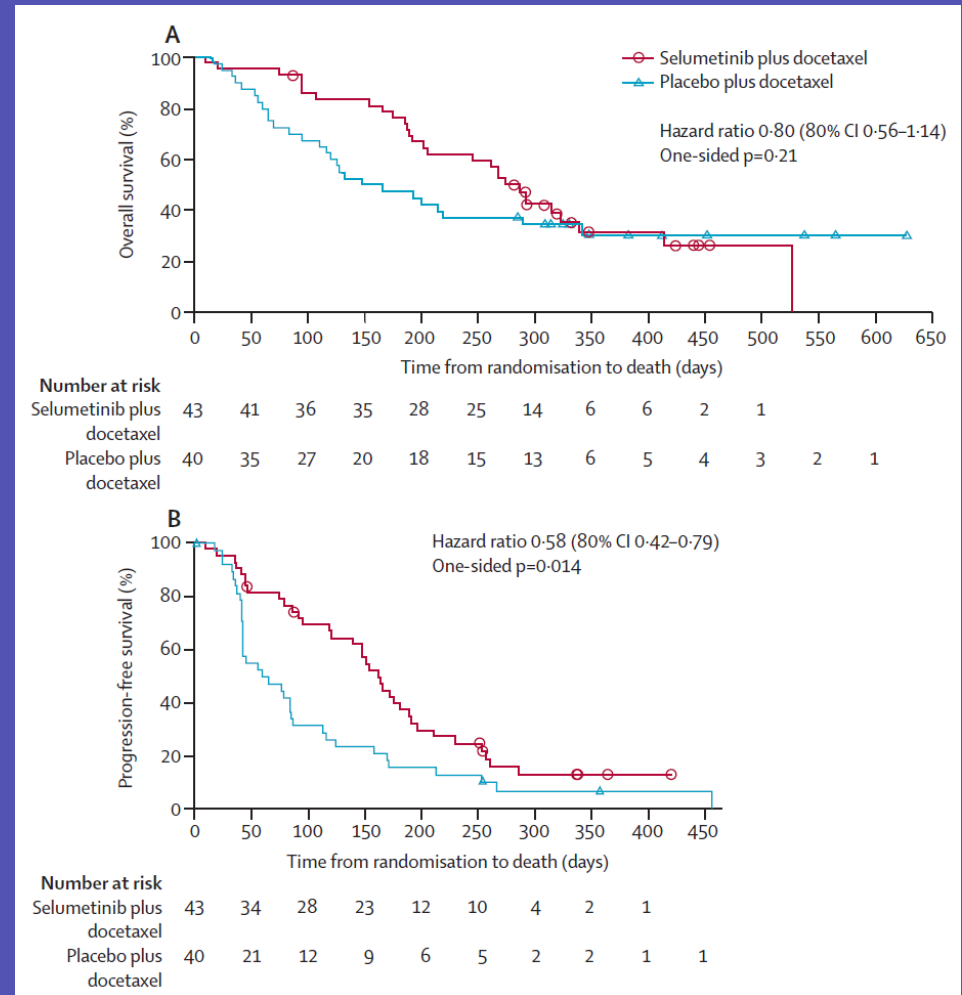
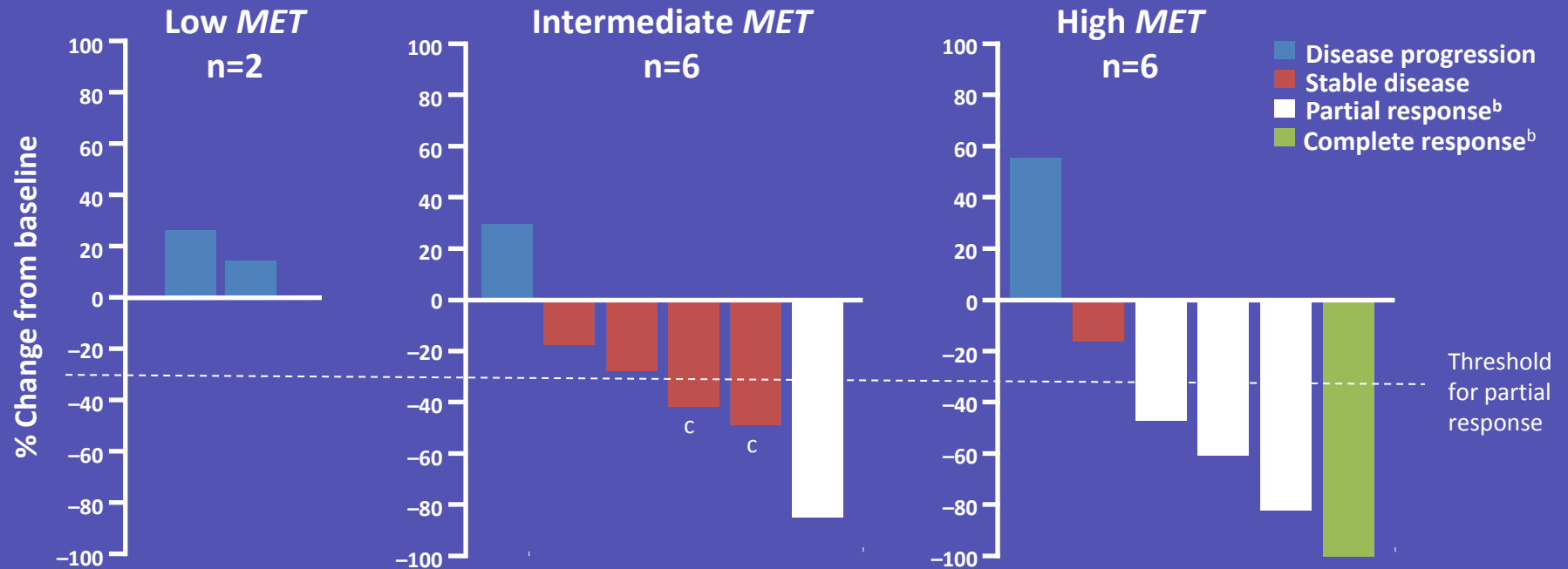


Fig. 1. The forest plot of relative ratios for overall response rate between *KRAS* mutant and wild-type patients in all studies.



## Amplification cMET

8001: Efficacy and safety of crizotinib in patients with advanced c-MET-amplified non-small cell lung cancer (NSCLC)



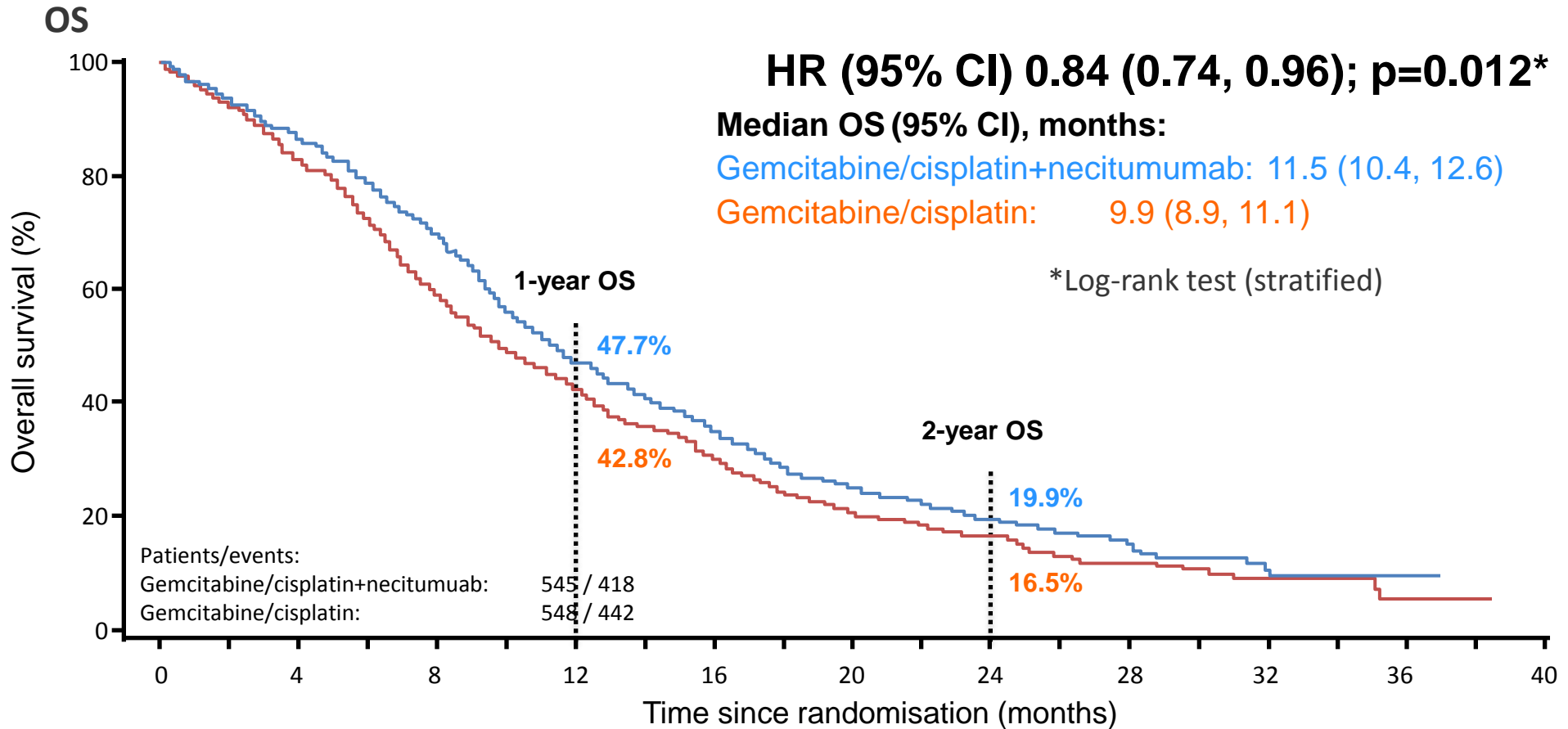
## Autres exemples de molécules intéressantes

- HER2: trastuzumab + CT, afatinib
- PI3K: inhibiteurs mTOR
- RET: cabozantinib, vandetanib, sunitinib, sorafenib
- BRAF (V600E): vemurafenib
- FGFR1: brivanib
- DDR2: activité *in vitro* imatinib, nilotinib, dasatinib

# Quelle place pour les anticorps monoclonaux?

- Anti EGFR
  - Cible connue mais pas de sélection adéquate du groupe-cible
  - Pas d'AMM en Europe
- AntiVEGFR
  - Cible mal connue
  - Bévacizumab améliore un peu survie mais toxicité ++
  - Groupe-cible à définir

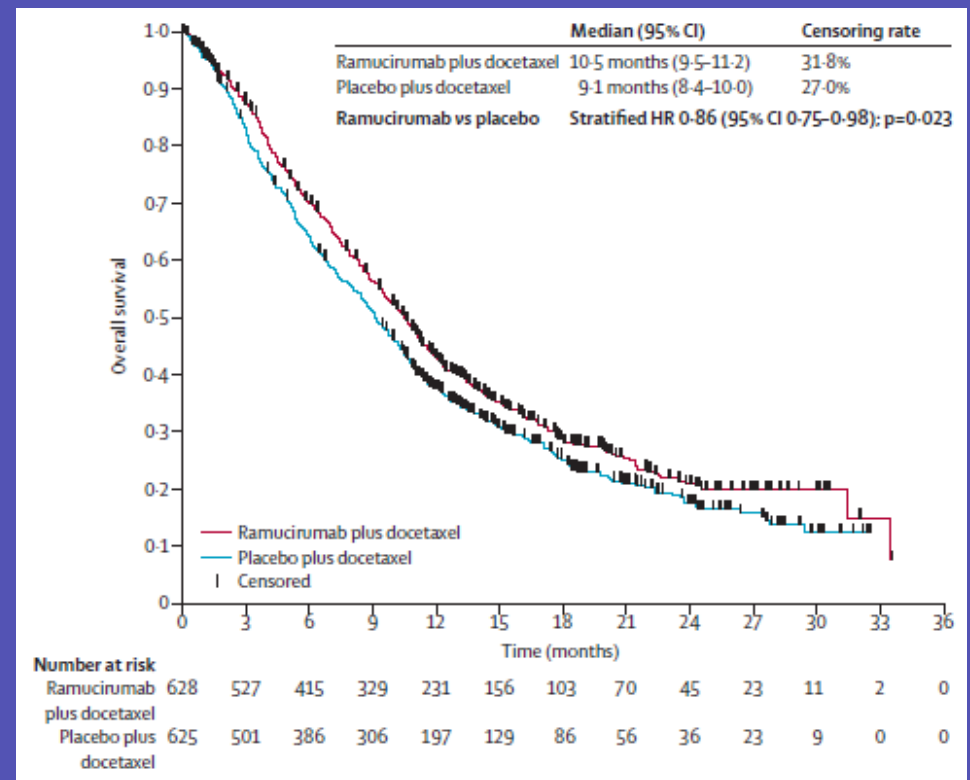
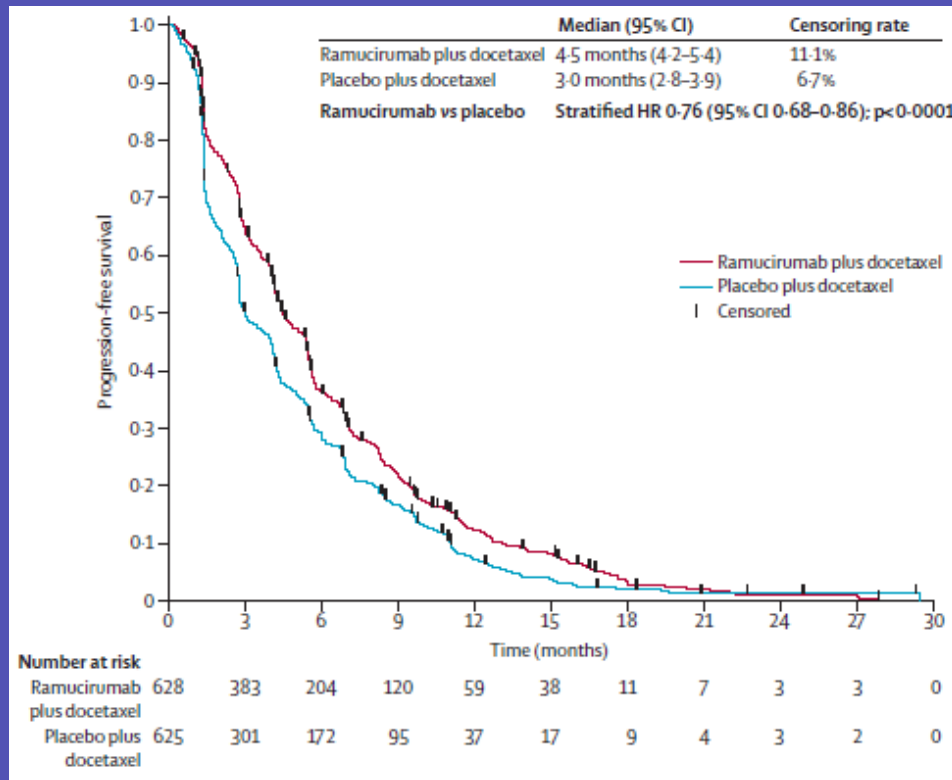
8008: A randomized, multicenter, open-label, phase III study of gemcitabine-cisplatin (GC) chemotherapy plus necitumumab (IMC-11F8/LY3012211) versus GC alone in the first-line treatment of patients (pts) with stage IV squamous non-small cell lung cancer (sq-NSCLC)



Follow-up time (median): Gemcitabine/cisplatin+necitumumab: 25.2 months; gemcitabine/cisplatin: 24.8 months

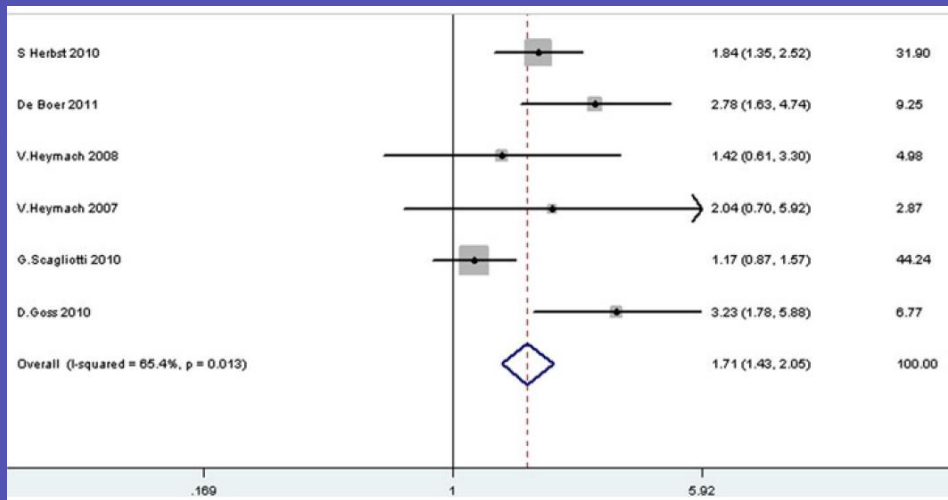
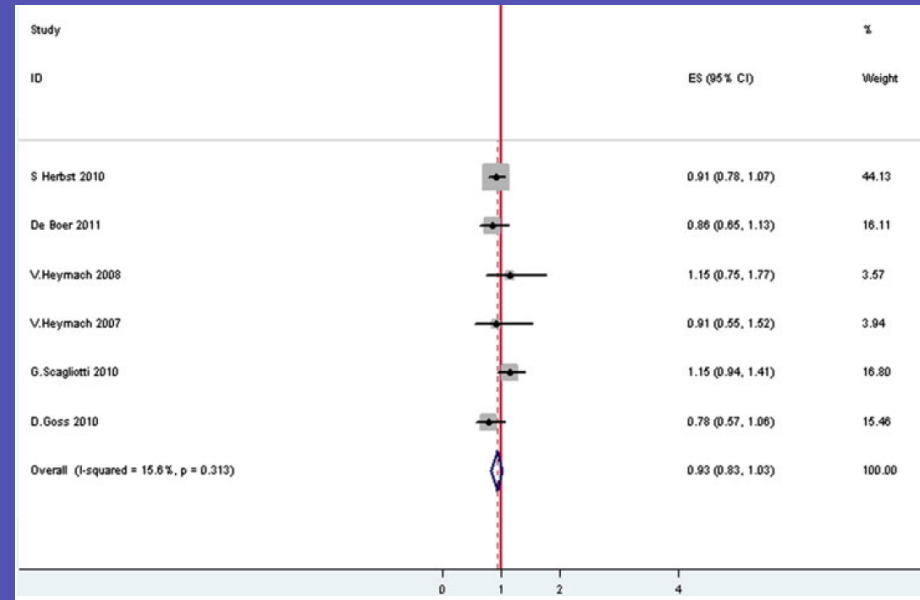
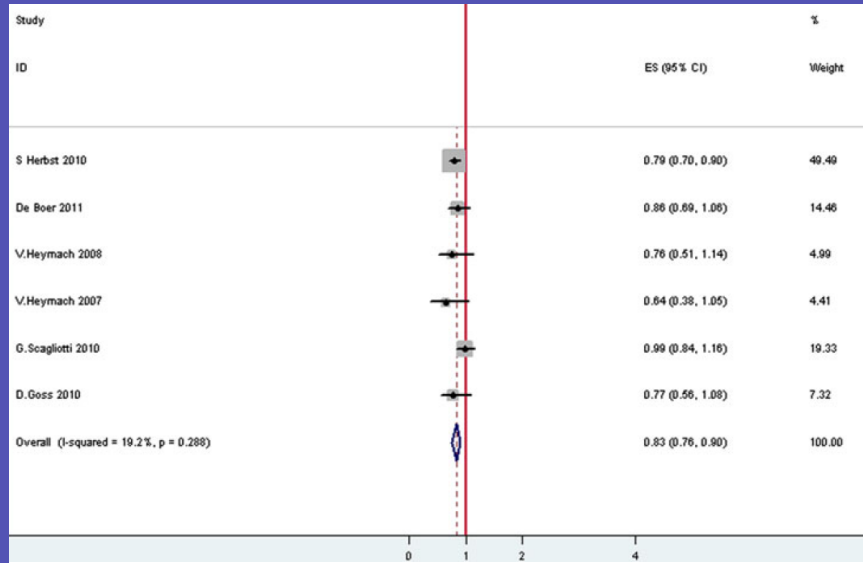
# Ramucirumab plus docetaxel versus placebo plus docetaxel for second-line treatment of stage IV non-small-cell lung cancer after disease progression on platinum-based therapy (REVEL): a multicentre, double-blind, randomised phase 3 trial

Edward B Garon, Tudor-Eliade Ciuleanu, Oscar Arrieta, Kumar Prabhaskar, Konstantinos N Syrigos, Tuncay Goksel, Keunchil Park, Vera Gorbunova, Ruben Dario Kowalyszyn, Joanna Pikiel, Grzegorz Czyzewicz, Sergey V Orlov, Conrad R Lewanski, Michael Thomas, Paolo Bidoli, Shaker Dakhil, Steven Gans, Joo-Hang Kim, Alexandru Grigorescu, Nina Karaseva, Martin Reck, Federico Cappuzzo, Ekaterine Alexandris, Andreas Sashegyi, Sergey Yurasov, Maurice Pérol





# Chemotherapy plus multitargeted antiangiogenic tyrosine kinase inhibitors or chemotherapy alone in advanced NSCLC: a meta-analysis of randomized controlled trials



Toxicities	Chemotherapy plus multitargeted antiangiogenic TKI group (CMATKI)		Chemotherapy group (C)		CMATKI vs. C	
	n1	a1 (%)	n0	a0 (%)	OR (95% CI)	p
Rash	1,555	668 (43.0)	1,518	328 (21.6)	2.78 (2.37-3.26)	0
Diarrhoea	1,681	632 (37.6)	1,641	397 (24.2)	1.92 (1.65-2.24)	0.003
Hypertension	1,681	236 (14.0)	1,641	88 (5)	2.90 (2.19-3.84)	0.291
Nausea	1,555	350 (32.5)	1,518	236 (15.5)	0.71 (0.60-0.83)	0.201
Vomiting	1,469	199 (13.5)	1,477	254 (17.2)	0.75 (0.61-0.92)	0.206
Hemorrhage	1,681	222 (13.2)	1,641	179 (10.9)	1.27 (0.98-1.56)	0.025
Fatigue	1,681	551 (51.3)	1,641	552 (33.6)	0.95 (0.82-1.11)	0.321
Cough	1,091	247 (22.6)	1,056	206 (19.5)	1.08 (0.87-1.34)	0.772
Constipation	1,469	223 (15.2)	1,477	235 (15.9)	0.95 (0.78-1.17)	0.245
Anorexia	1,595	373 (23.4)	1,600	344 (21.5)	1.12 (0.95-1.33)	0.029
Alopecia	1,295	321 (24.8)	1,245	328 (26.3)	0.91 (0.75-1.10)	0.413

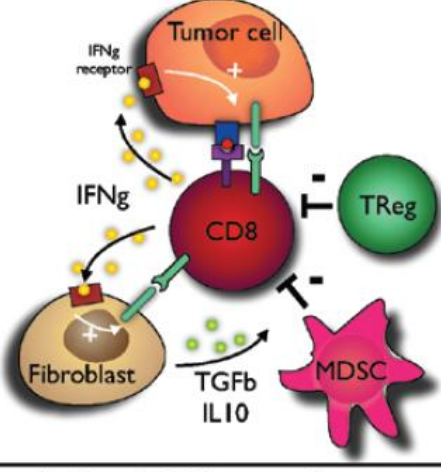
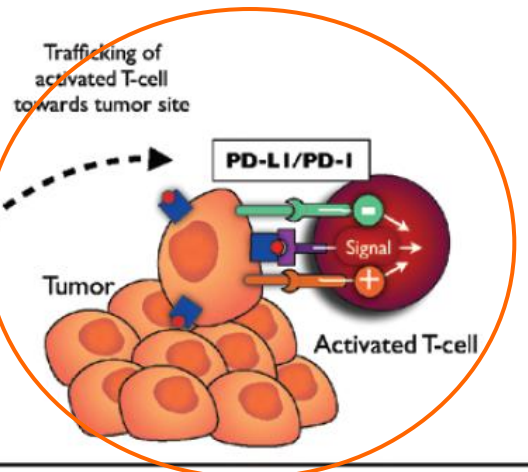
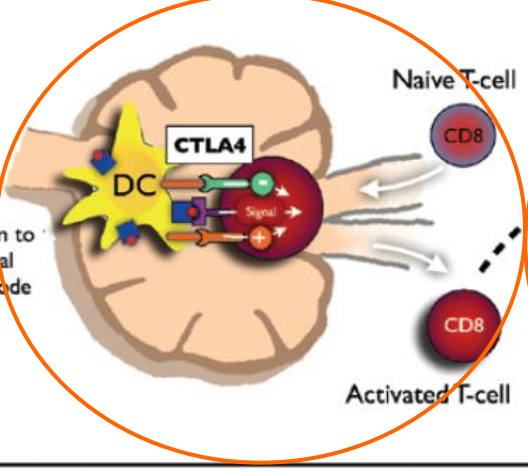
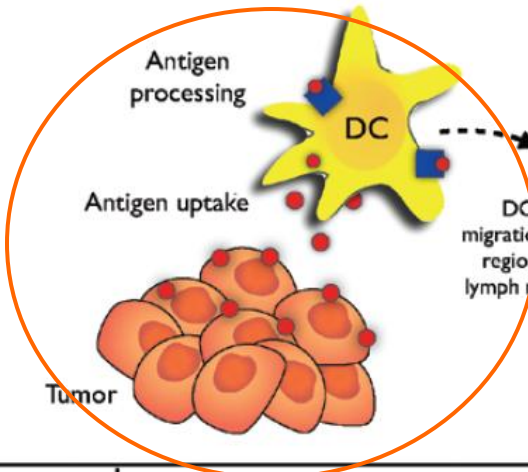
## 5. Quelques mots d'immunothérapie

**Tumor antigen uptake and DCs antigen processing**

**T-cell priming in regional lymph node**

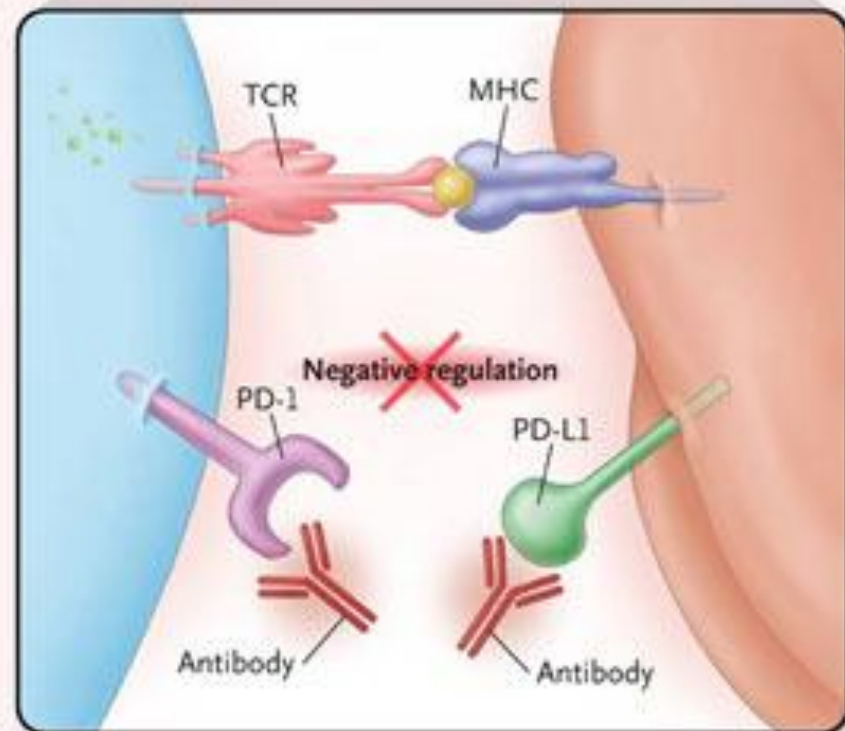
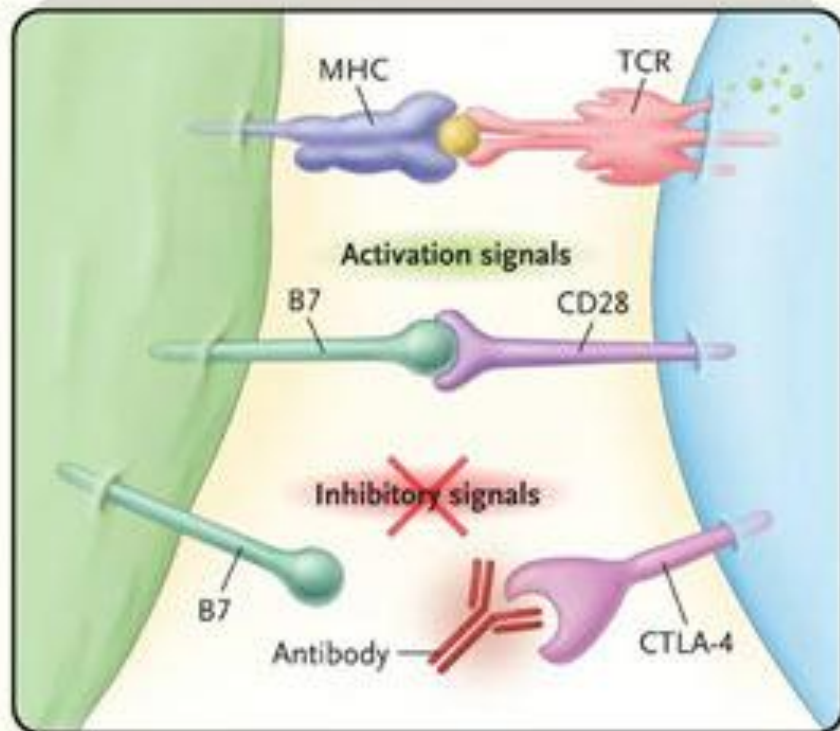
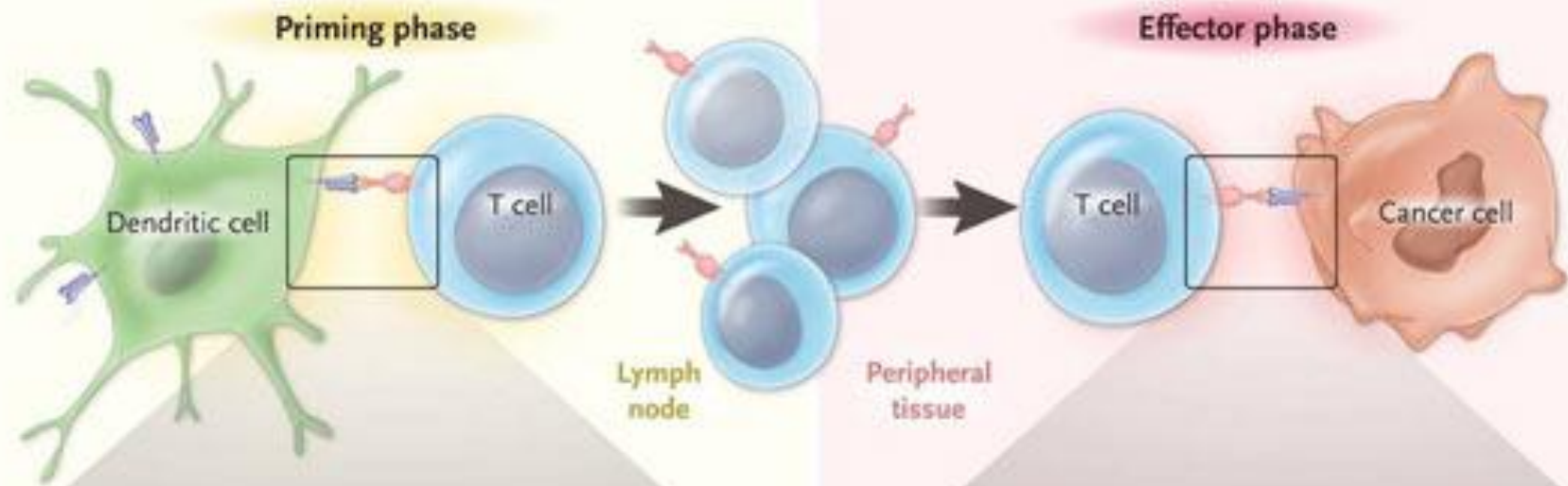
**T-cell activation in the tumor**

**T-cell modulation in the tumor environment**



immune system modulation in NSCLC	> <u>down-regulation of MHC-I</u>	> <u>up-regulation of PD-L1</u> through activation of PI3K/Akt ? MAPK ? Alk ?	> <u>up-regulation of TRegs</u> > <u>up-regulation of MDSCs</u>	> <u>IL-10 and TGFb</u> increased concentration by tumor environment > <u>up-regulation of PD-L1</u> by IFNg secreted by activated T-cell
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immuno modulation by NSCLC drugs	<p><b>immunogenic cell death</b> irradiation</p> <p><b>vaccination strategies</b> MAGE-A3, MUC-1, rHU-EGF</p>	<p><b>up-regulation of MHC-I</b> paclitaxel, gemcitabine, erlotinib</p> <p><b>DC maturation</b> paclitaxel, docetaxel, bevacizumab</p> <p><b>anti-CTLA4</b> Ipilimumab, Tremelimumab</p>	<p><b>anti-PD-I</b> MDX-1106, CT-011, MK-3475</p> <p><b>anti-PD-L1</b> MPDL-3280A, MDX-1105</p> <p><b>up-regulation of PD-L1</b> paclitaxel, etoposide</p>	<p><b>TReg inhibition</b> cisplatin, paclitaxel, bevacizumab</p> <p><b>MDSC inhibition</b> cisplatin, docetaxel, gemcitabine</p> <p><b>down-regulation of PD-L1</b> by PI3Ki ? MEKi ? Crizotinib ?</p>
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## 8007: Safety and clinical activity of MK-3475 as initial therapy in patients with advanced non-small cell lung cancer (NSCLC)

	<b>RECIST v1.1</b>	<b>Immune-related response criteria</b>
ORR %(IC 95%)	26 (14-42)	47 (32-62)
SSP médiane (IC 95% CI), sem	27 (13,6-45,0)	37 (27-NR)

- Effets secondaires >5% patients: fatigue (22%), prurit (13%), hypothyroïdisme (9%), dermatite acnéiforme (7%), diarrhée (7%), dyspnée (7%), rash (7%)

8022: Safety and response with nivolumab (anti-PD-1; BMS-936558, ONO-4538) plus erlotinib in patients (pts) with epidermal growth factor receptor mutant (EGFR MT) advanced NSCLC

	<b>Nivolumab+erlotinib (n=21)</b>
<b>ORR</b>	19%
<b>SSP</b>	
SP (IC 95%) à 24 sem	51% (28-70)
SSP Médiane (range), sem	29,4 (4,6-81,7+)
<b>OS</b>	
Survie à 1 an (IC 95%)	73% (46-88)
Survie médiane (range), sem	Non atteinte (10,7+-86,9+)

## 6. Rôle des consultations multidisciplinaires

## O27.07 MOLECULAR MULTIDISCIPLINARY TUMOR BOARD (MMTB) FOR LUNG CANCER PATIENTS: 2-YEAR EXPERIENCE REPORT

- Cliniciens, unité phase I, pathologistes et biologistes.
- 245 pts
- Oncogenic drivers
  - ALK 10%
  - EGFR mut 24%
  - KRAS mut 30%
  - PI3KCA mut 0,4%
  - BRAF mut 3%
  - HER2 mut 1%
  - FGFR1 amplification 3%
  - Autres mutations 14%
- Recommendations
  - ✧ Inclusion étude 75 pts (31%)
  - ✧ Prescription selon AMM 49 pts (20%)
  - ✧ Prescription hors AMM 18 pts (7%)
  - ✧ Expanded access program 18 pts (7%).
- Ont reçu le traitement ciblé proposé
  - ✧ D'emblée 63 (42%)
  - ✧ A la progression 16 (11%)

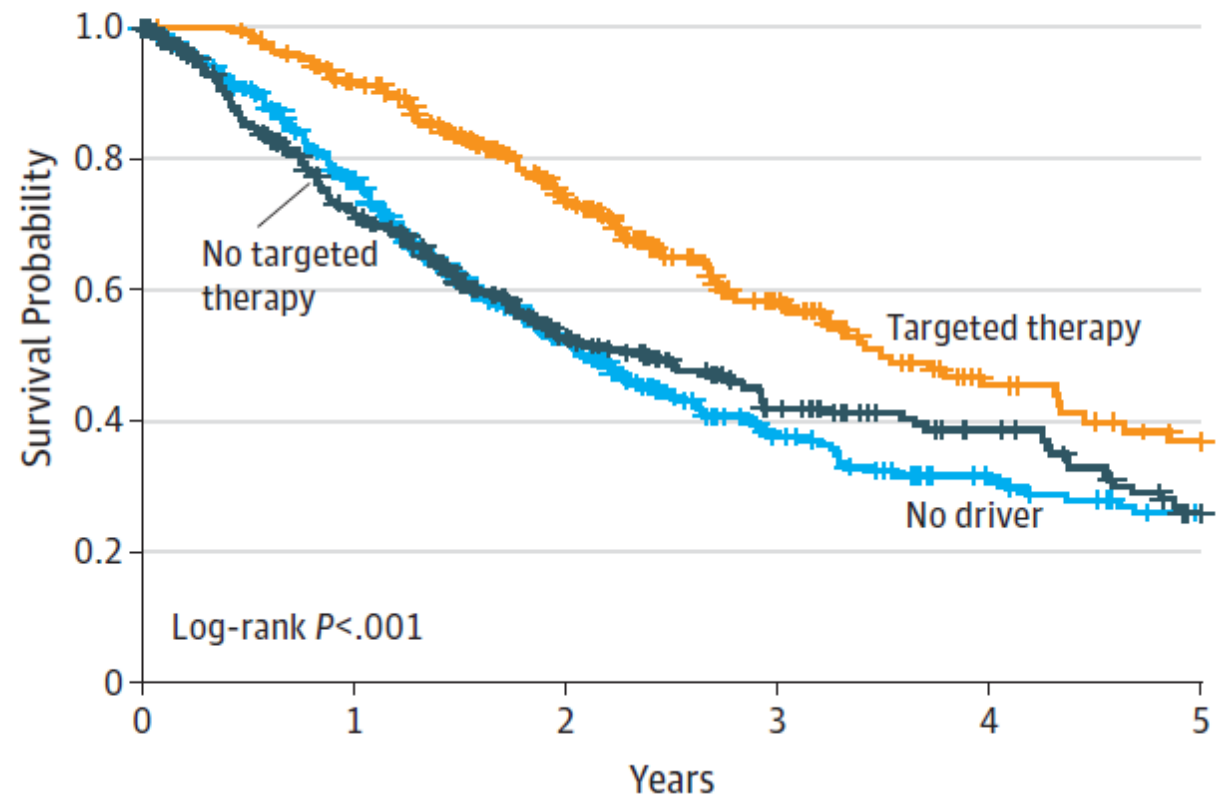


# Quelle importance?

Table 2. Oncogenic Drivers Identified and Targeted Treatments Received

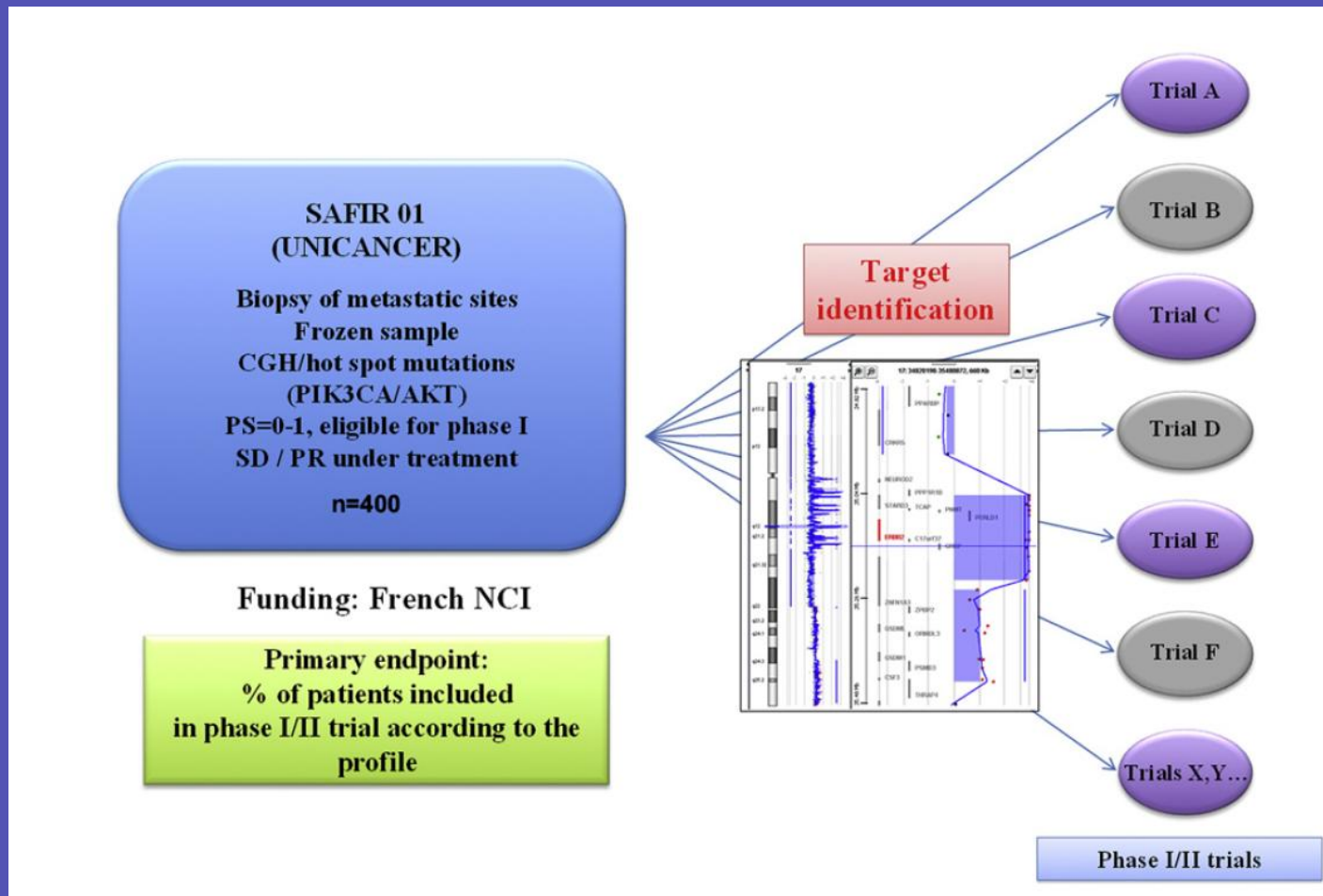
Gene With Mutational or Structural Change	Genotyping, N
	Any (n = 1007) <sup>a,b</sup>
Any gene(s)	623 (62) [59-65]
Singletons <sup>d</sup>	
<i>KRAS</i>	245 (24) [22-27]
<i>EGFR</i> (sensitizing) <sup>e</sup>	175 (17) [15-20]
exon19 del	103 (10) [9-12]
<i>L858R</i>	64 (6) [5-8]
<i>G719X</i>	5 (0.5) [0.2-1]
<i>L861Q</i>	5 (0.5) [0.2-1]
<i>ALK</i> (rearrangement)	80 (8) [6-10]
<i>EGFR</i> (other) <sup>f</sup>	35 (4) [3-5]
<i>ERBB2</i> (formerly <i>HER2</i> )	23 (2) [2-4]
<i>BRAF</i>	18 (2) [1-3]
<i>V600E</i>	14 (1) [0.8-2]
Non- <i>V600E</i>	4 (0.4) [0.1-1]
<i>PIK3CA</i>	7 (0.7) [0.3-2]
<i>MET</i> (amplification)	6 (0.6) [0.2-1]
<i>NRAS</i>	5 (0.5) [0.2-1]
<i>MEK1</i>	2 (0.2) [0.03-1]
<i>AKT1</i>	0 [0-1]
Doubletons	
>1 gene	27 (3) [2-4]

**A** Patients with an oncogenic driver mutation who did and did not receive targeted therapy, and patients without an oncogenic driver



# The challenge to bring personalized cancer medicine from clinical trials into routine clinical practice: The case of the Institut Gustave Roussy

Monica Arnedos<sup>a,b</sup>, Fabrice André<sup>a,b</sup>, Françoise Farace<sup>b,c</sup>, Ludovic Lacroix<sup>b,c</sup>, Benjamin Besse<sup>a,b</sup>, Caroline Robert<sup>a,b</sup>, Jean Charles Soria<sup>a,b</sup>, Alexander M.M. Eggermont<sup>d,\*</sup>



# Designs and challenges for personalized medicine studies in oncology: focus on the SHIVA trial

Christophe Le Tourneau • Maud Kamal • Olivier Trédan • Jean-Pierre Delord • Mario Campone • Anthony Goncalves • Nicolas Isambert • Thierry Conroy • David Gentien • Anne Vincent-Salomon • Anne-Lise Pouliquen • Nicolas Servant • Marc-Henri Stern • Anne-Gaëlle Le Corroller • Sébastien Armanet • Thomas Rio Frio • Xavier Paoletti

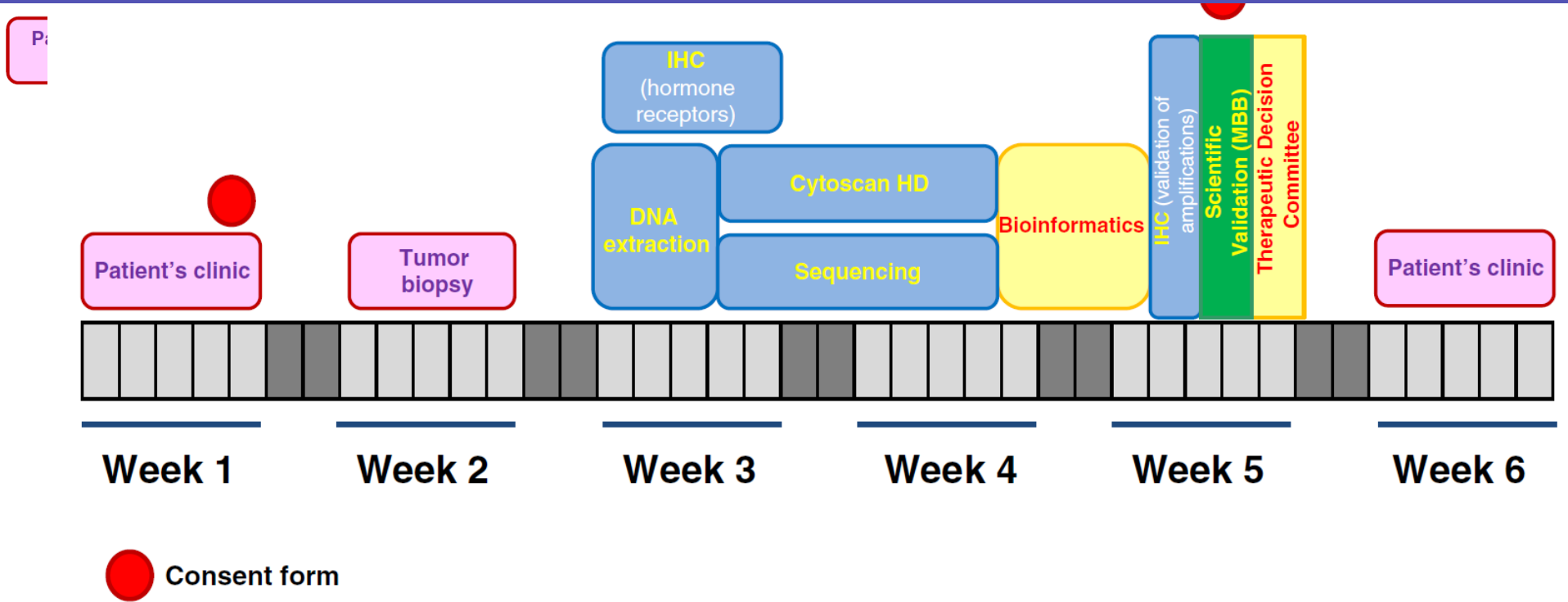


Fig. 3 Timelines for the establishment of the molecular profile in the SHIVA trial



# Conclusions

- Les thérapies ciblées sont actives si la cible est connue!
  - Les avancées en biologie moléculaire amènent à une complexification de la prise en charge des CBNPC qui d'une maladie fréquente évolue vers de nombreuses maladies rares
- ⇒ prise en charge multidisciplinaire (concertations multidisciplinaires moléculaires)