

Cancer Bronchique Non à Petites Cellules

Les nouvelles stratégies thérapeutiques

2016-2020

Cours du GOLF
Lyon – 21 Septembre 2016

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**Assistance Publique
Hôpitaux de Marseille**



Disclosure slide

I provided consultations for Astra-Zeneca, Bristol-Myers Squibb, Boehringer–Ingelheim, Clovis Oncology, Eli Lilly Oncology, F. Hoffmann–La Roche Ltd, Novartis, Merck, MSD, Pierre Fabre and Pfizer.

2016-2020 ...



2016-2020 ...

- **Nouveauté(s) ?**
 - Connaissances scientifiques
 - Progrès technologiques
 - Classes thérapeutiques (Ph II ou III)
- **Aspects économiques**
 - Business
 - Contraintes économiques

Le cancer du poumon



- **45,222** nouveaux cas (**3^{ème}**)
 - 90% C. NAPC
 - 2/3 stades IV
- **30,555** décès (1^{er})
- **2 x C. colon** (17,833)
- **3 x C. sein** (11,913)

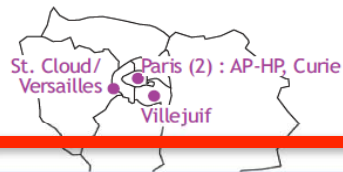
Les cancers en France en 2015, INCa 2016

Agenda

- **Le whole genome pour tous ?**
- L'immunothérapie, le graal ?
- Faire du neuf avec du vieux ?

LC genotyping program: France

- 28 platforms (2006)
- 10 routine biomarkers (+ 6 emerging bm)



Cancer du poumon

Mutations d'EGFR

23 336

8 924

Translocation d'ALK

18 861

709

19 347

3 330

1 105

1 005

23 336

18 861

5 026

6 750

861

TOTAL

89 254

Tableau 1. Nombre de recherches de marqueurs prédictifs de la réponse à une thérapie ciblée en 2013

Pathologie	Biomarqueur	Nombre de tests
Cancer du poumon	Mutations d'EGFR	23 336
	Translocation d'ALK	18 861
GIST	Mutations de PDGFRα	1 005
Cancer du poumon	Mutations d'EGFR	23 336
	Translocation d'ALK	18 861
Mélanome	Mutation de BRAF V600	5 026
Leucémies	Détection de BCR-ABL	6 750
	Mutations d'ABL	861
TOTAL		89 254

* i.e. Regional molecular genetics centers

Available at www.ecancer.fr

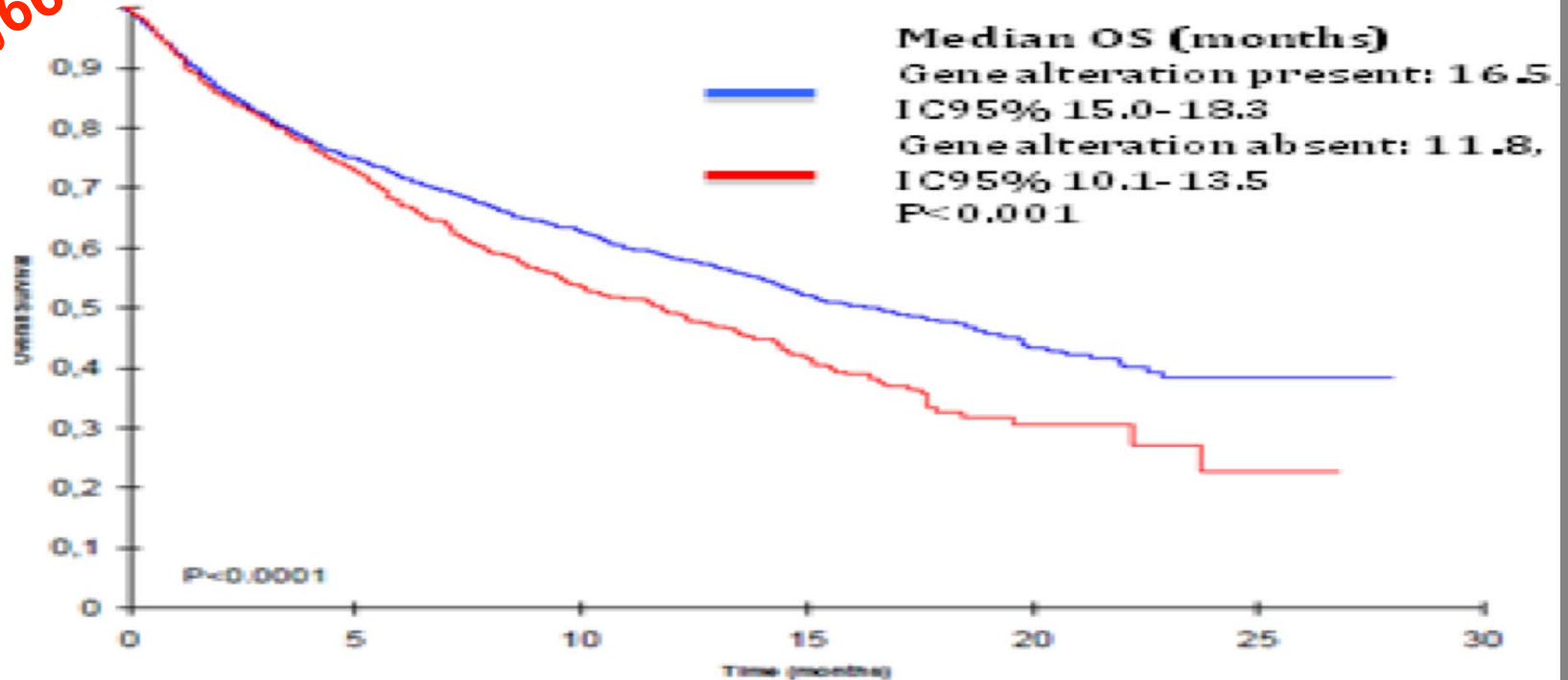
Back to the basics first?

<i>EGFR</i> mutation – no. (%)			
Positive	42 (9.9)	43 (10.1)	85 (10.0)
Negative	318 (74.8)	310 (72.9)	628 (73.9)
Unknown	65 (15.3)	72 (16.9)	137 (16.1)
<i>EMLA-ALK</i> translocation — no. (%)			
Positive	2 (0.5)	0	2 (0.2)
Negative	223 (52.5)	201 (47.3)	424 (49.9)
Unknown	200 (47.1)	224 (52.7)	424 (49.9)
<i>KRAS</i> mutation — no. (%)			
Positive	26 (6.1)	33 (7.8)	59 (6.9)
Negative	99 (23.3)	104 (24.5)	203 (23.9)
Unknown	300 (70.6)	288 (67.8)	588 (69.2)

Example from a large phase III trial in NSCLC

Impact of routine molecular profiling

17,664 pts



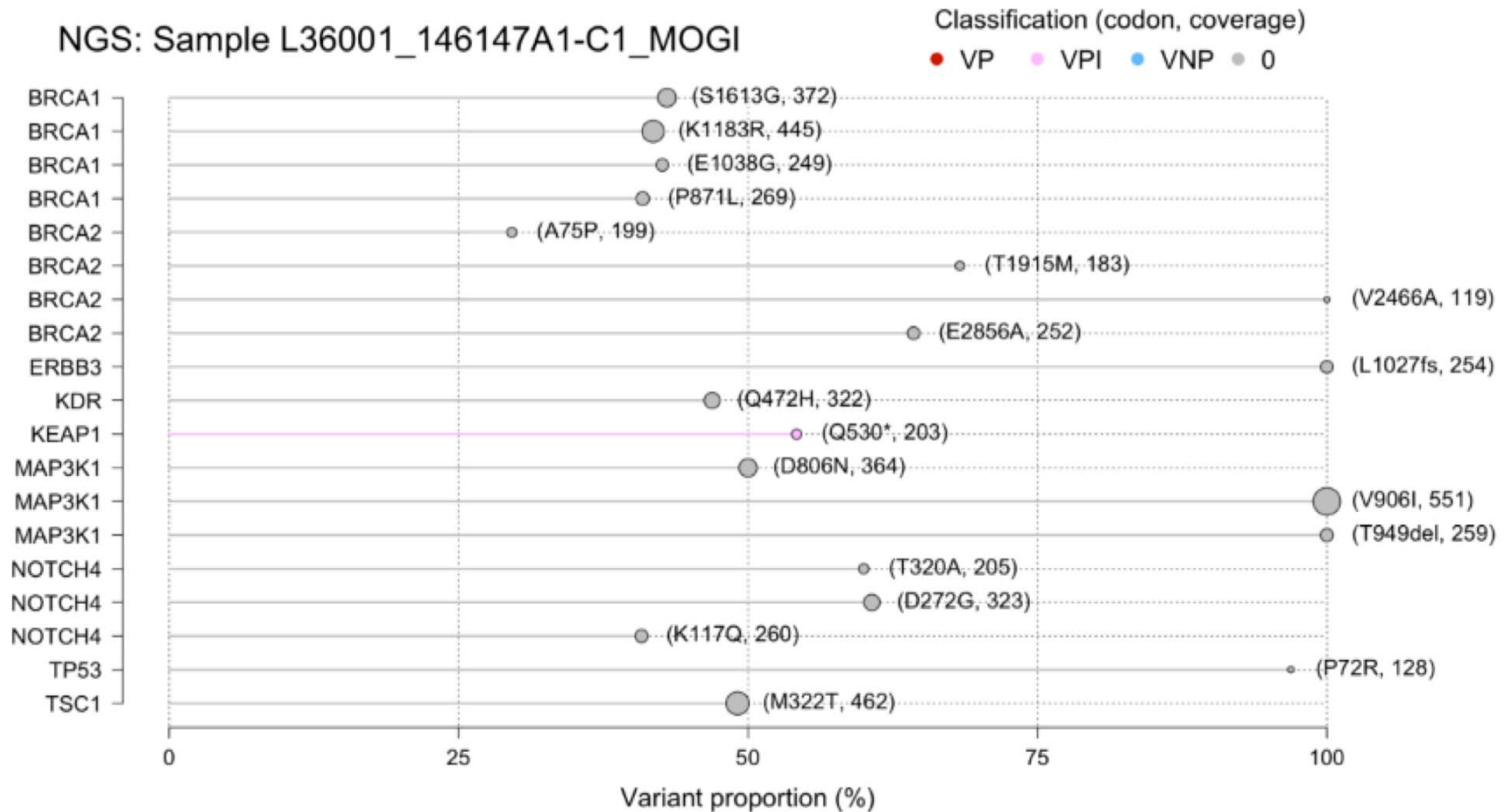
Number at risk

Driver + : 8488	2141	1428	584	165	8	0
Driver - : 1126	617	388	124	24	4	0



Barlesi et al, Biomarkers France, Lancet 2016

High throughput molecular genotyping



Images: NGS analyses from the SAFIR lung Unicancer IFCT trial

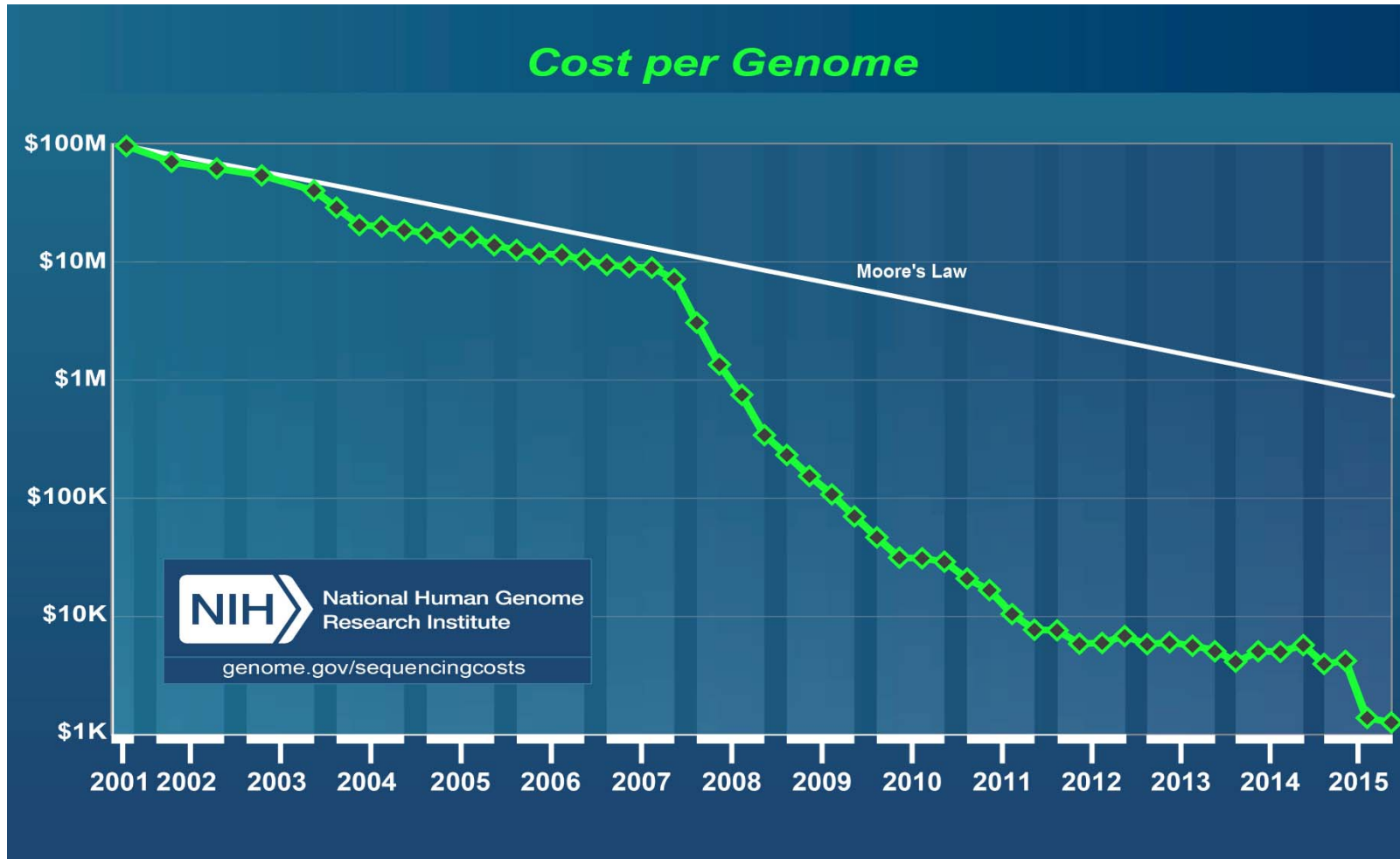
Larger genomic profiling (NGS)



Panel tumeurs solides			
gène	Exons / hotspots	Molécule associée	utilité clinique
AKT1	3	AKT inhibitors	essais cliniques
ALK	23+24+25	crizotinib, ALK inhibitors	AcSé, essais cliniques
BRAF	11+15	vemurafenib, dabrafenib	AMM
EGFR	18+19+20+21	anti EGFR	AMM
ERBB2 (HER2)	20	trastuzumab, neratinib	essais cliniques
ERBB4	E452K et R393W	Afatinib	essais cliniques
FGFR2	S252, N549, K659	FGFR inhibitors	essais cliniques
FGFR3	7+9+14 (R248 à S249 et G370 à Y	FGFR inhibitors	essais cliniques
HRAS	2+3+4	inhibiteurs de MEK	essais cliniques
KIT	8+9+11+13+17+18	imatinib	AMM
KRAS	2+3+4	panitumumab et cetuximab	AMM
MAP2K1 (MEK1)	2	inhibiteurs de MEK	essais cliniques
MET	2 + 14 à 20	crizotinib	AcSé
NRAS	2+3+4	panitumumab, MEK inhibitors, BRAF inhibitors	pré-AMM, essais cliniques
PDGFRA	12+14+18	imatinib	AMM
PIK3CA	9 + 20	PI3K inhibitors	essais cliniques

Available www.ecancer.fr

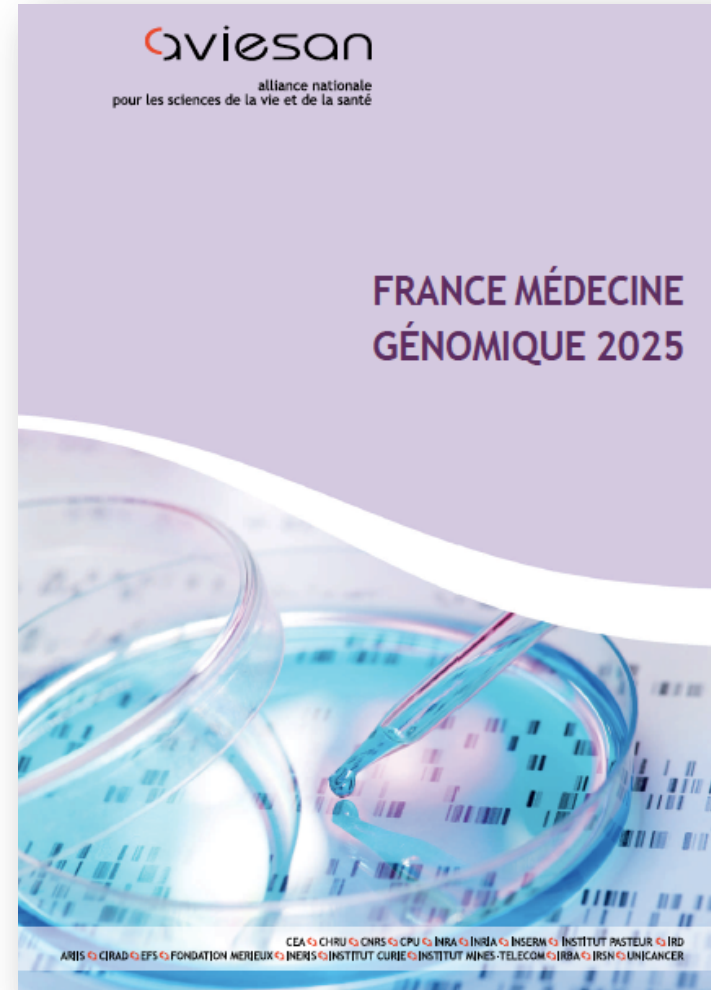
High throughput molecular genotyping



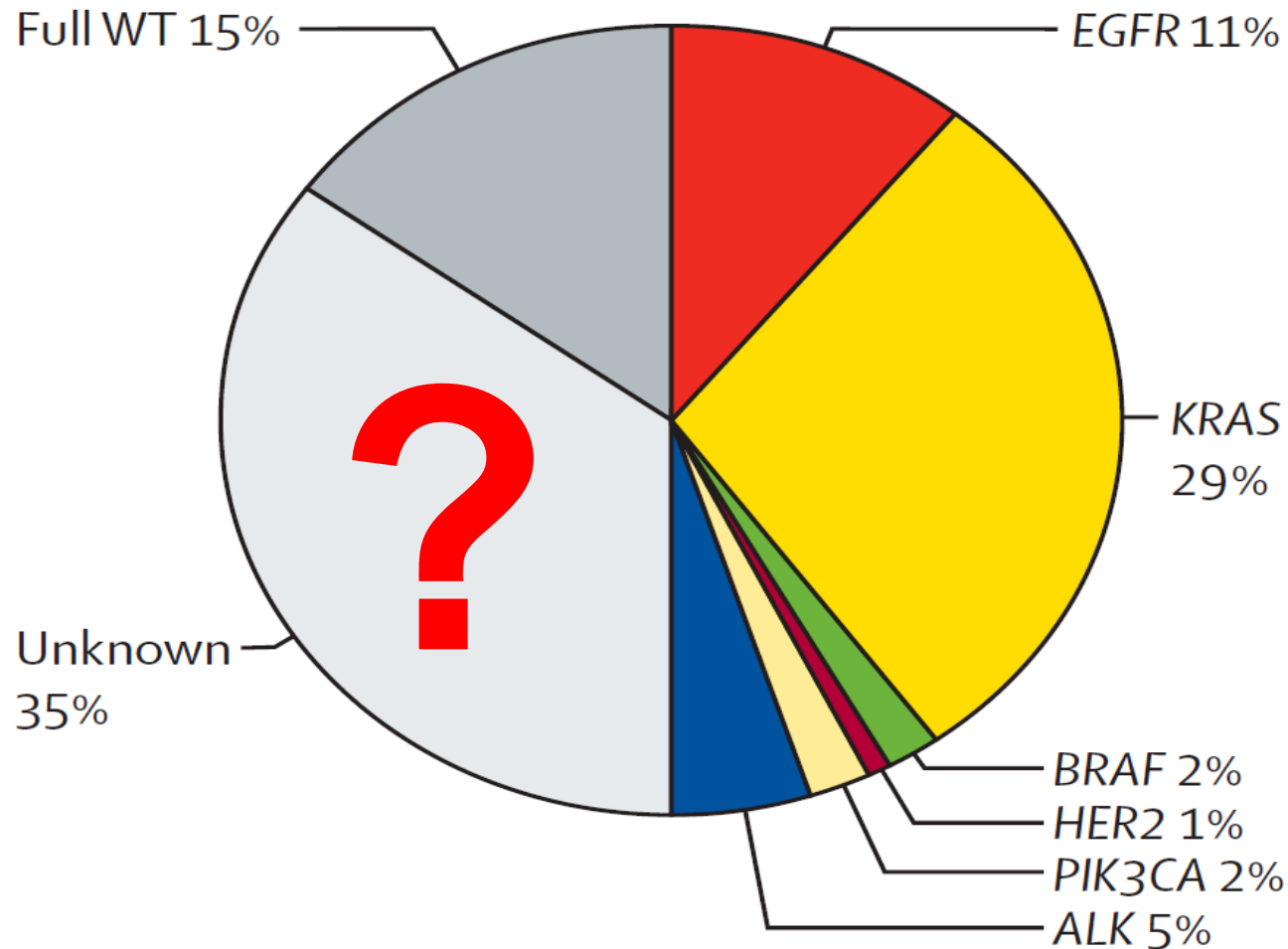
Images: NGS analyses from the NIH website

Enjeux et perspectives

- **Evolutions
des plateformes**

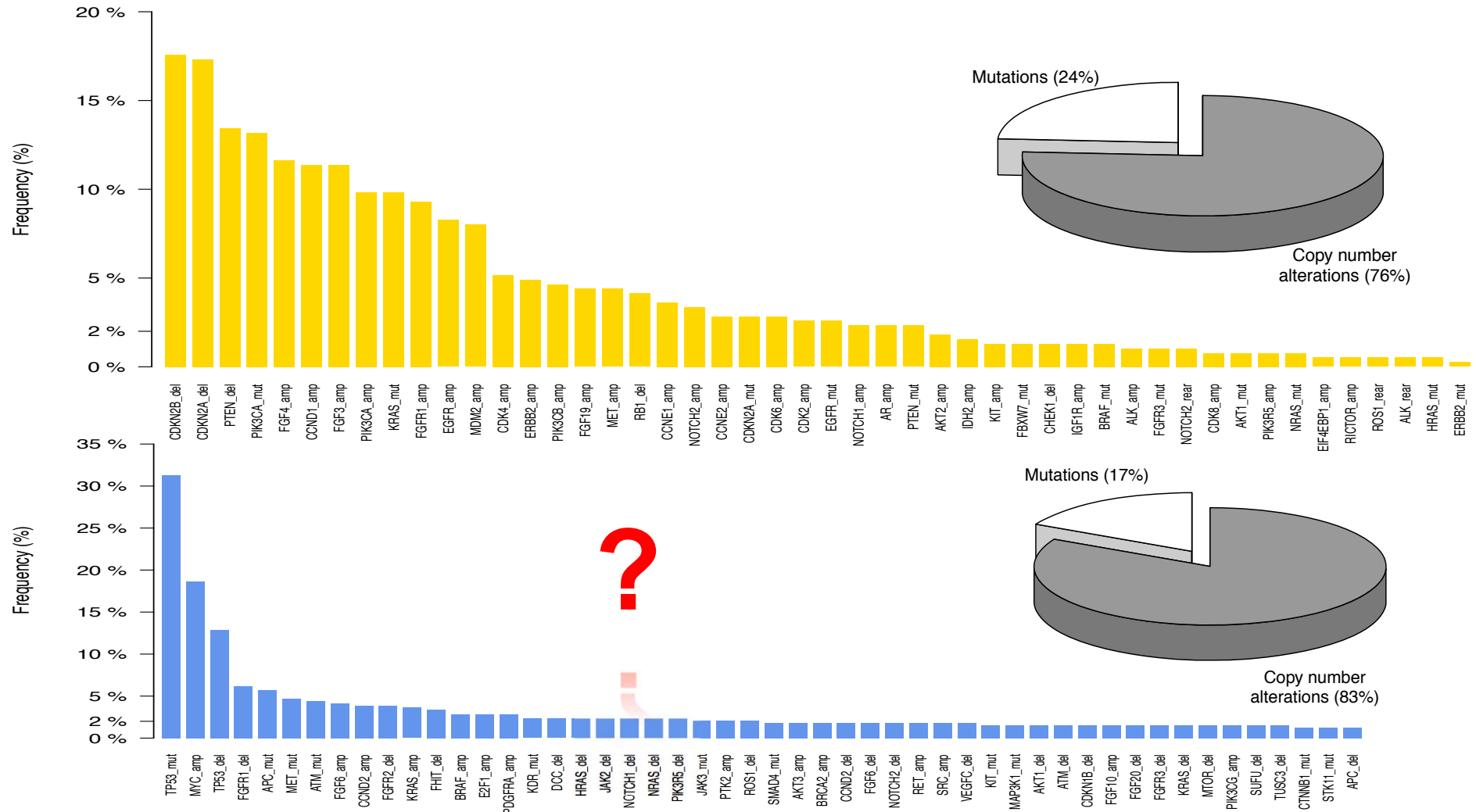


France (Biomarkers France)



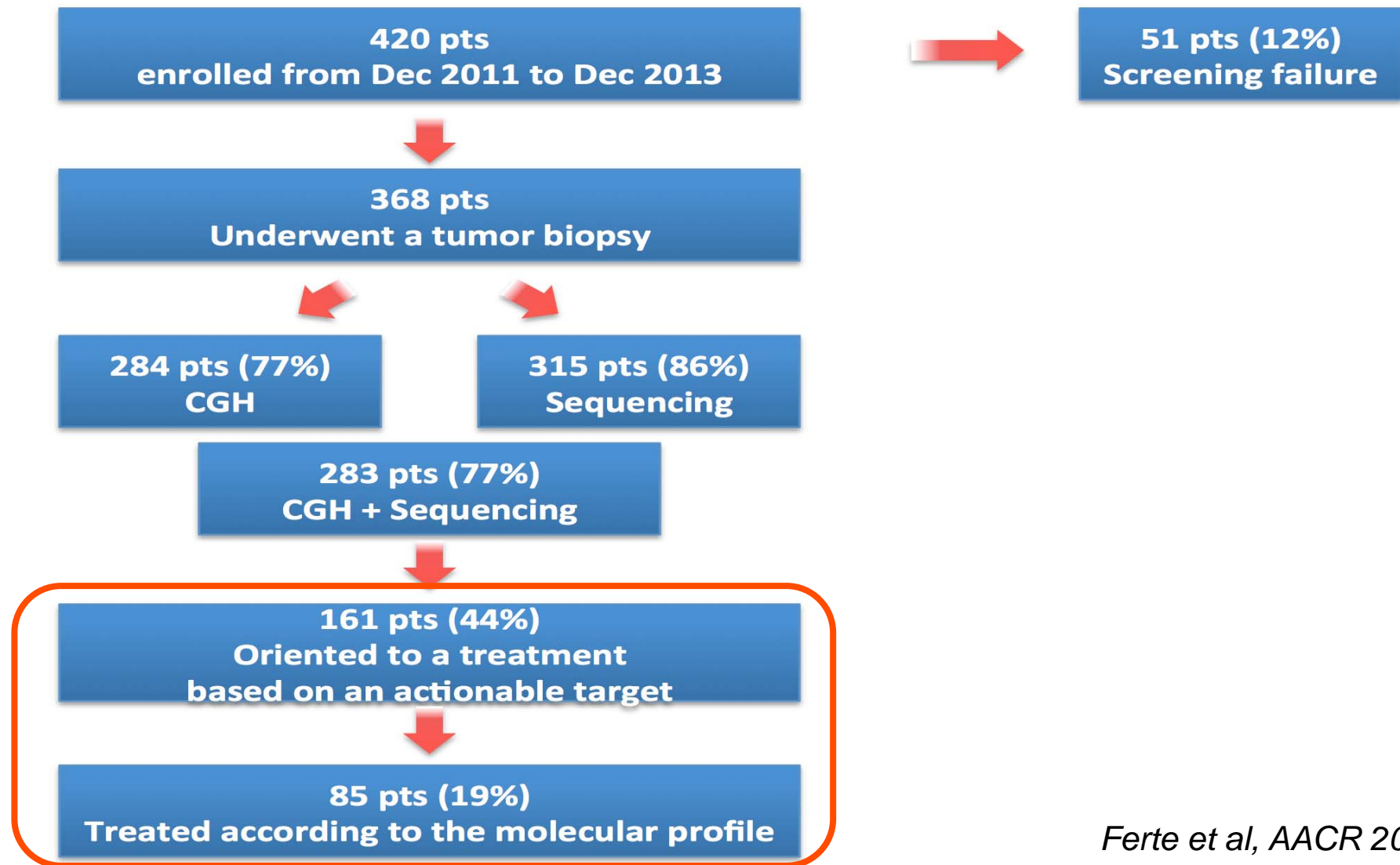
Barlesi et al, Biomarkers France, Lancet 2016 (in press)

How large should be the analysis?



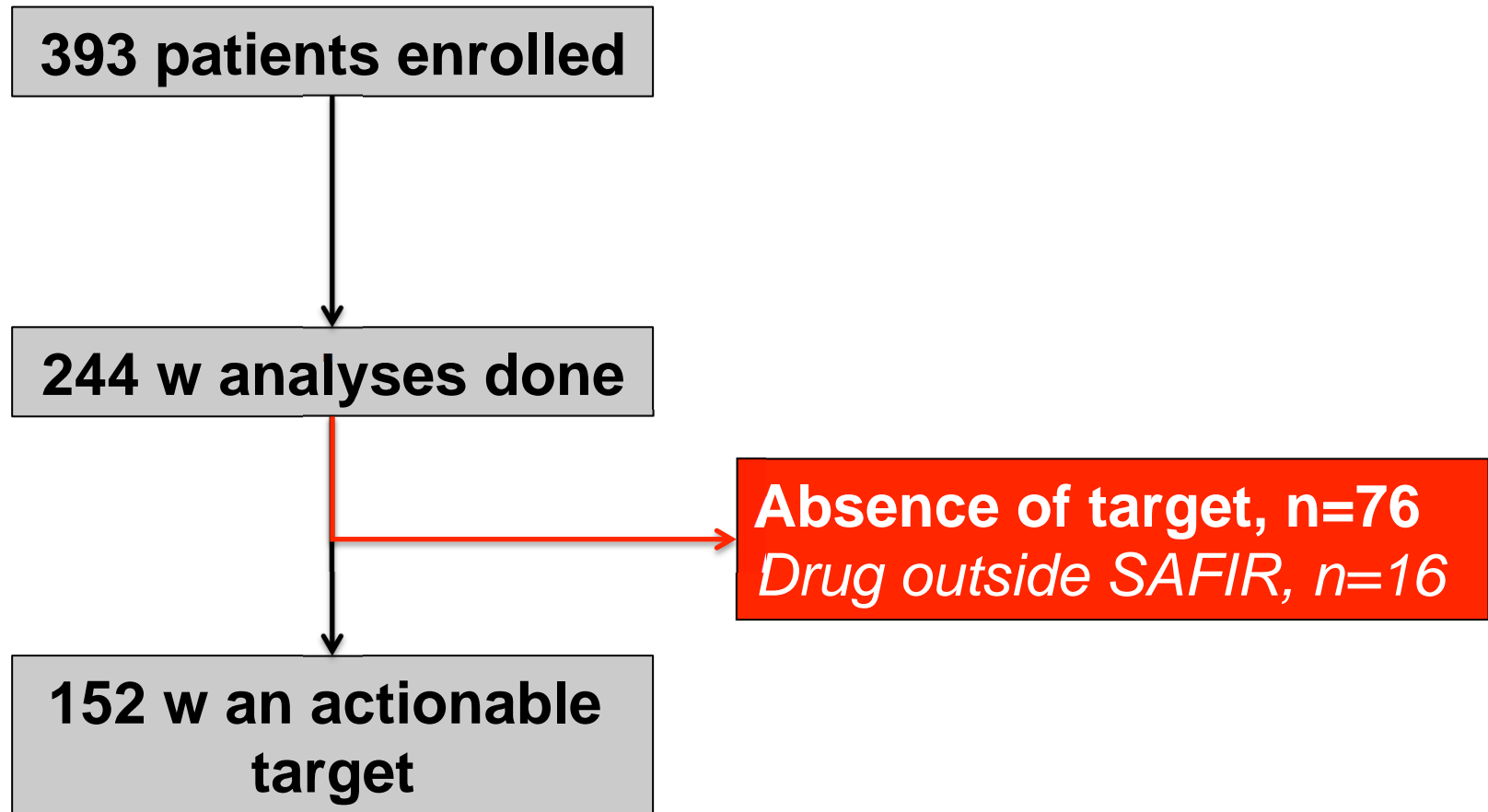
Ferte C et al, AACR 2014

Precision medicine for increased survival?



Ferte et al, AACR 2014

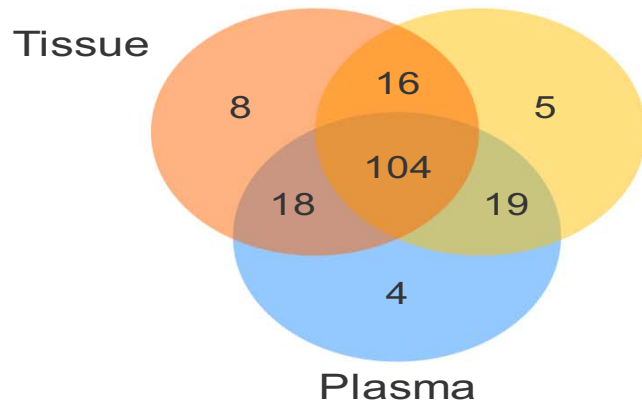
Trial status (April 2nd, 2016)



cfDNA for molecular genotyping

T790M		Tissue			Total
		Positive	Negative	Inadequate	
Plasma (BEAMing)	Positive	313	23	38	374
	Negative	74	17	17	108
Total		387	40	55	482

T790M-positive cases (n=181)



Tissue

Urine Total positive by tissue: 146 of 181

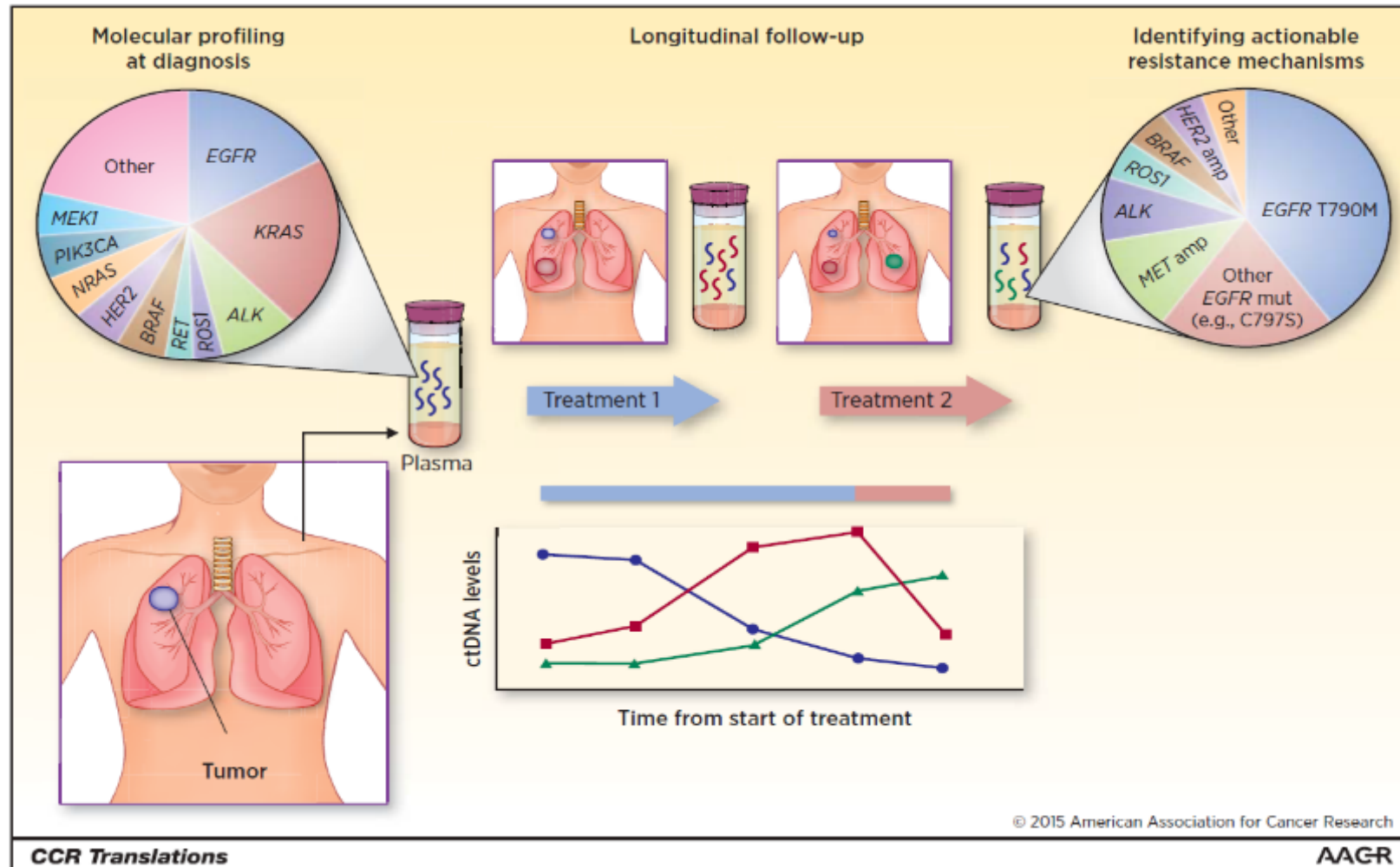
Total positive by plasma: 145 of 181

Total positive by urine: 144 of 181

104 (57%) were positive by all 3 sample types

Wakelee H et al, ASCO 2016 (abst 9001)

cfDNA as solution to monitor biomarkers?

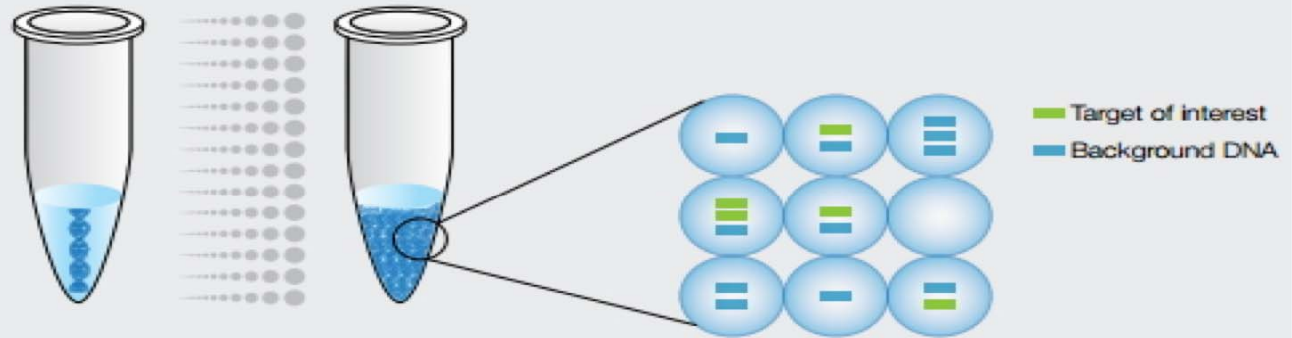


Tsuy DW et al, Clin Cancer Res (in press)

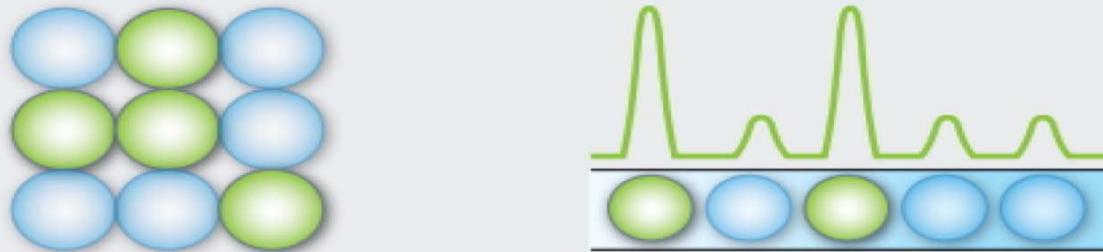
ddPCR development



Droplet Digital PCR



The sample is partitioned into 20,000 droplets, with target and background DNA randomly distributed among the droplets.

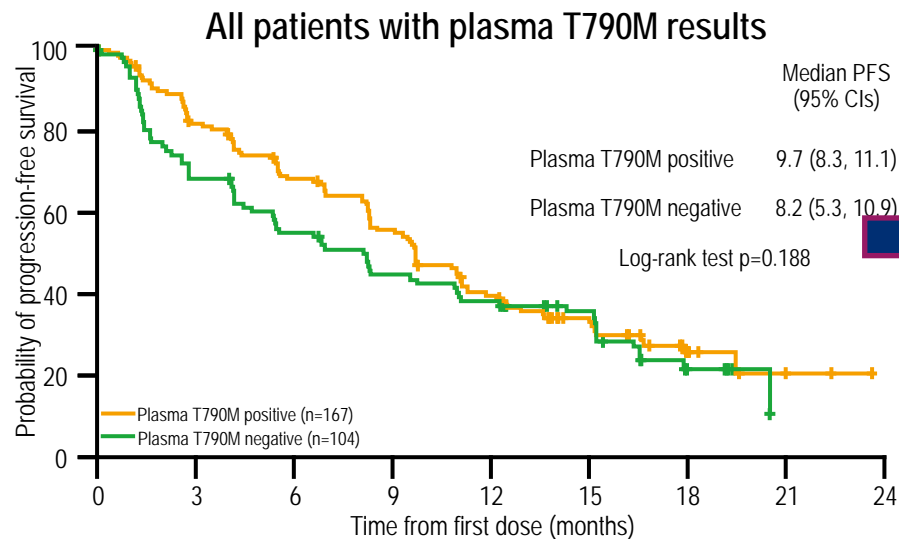
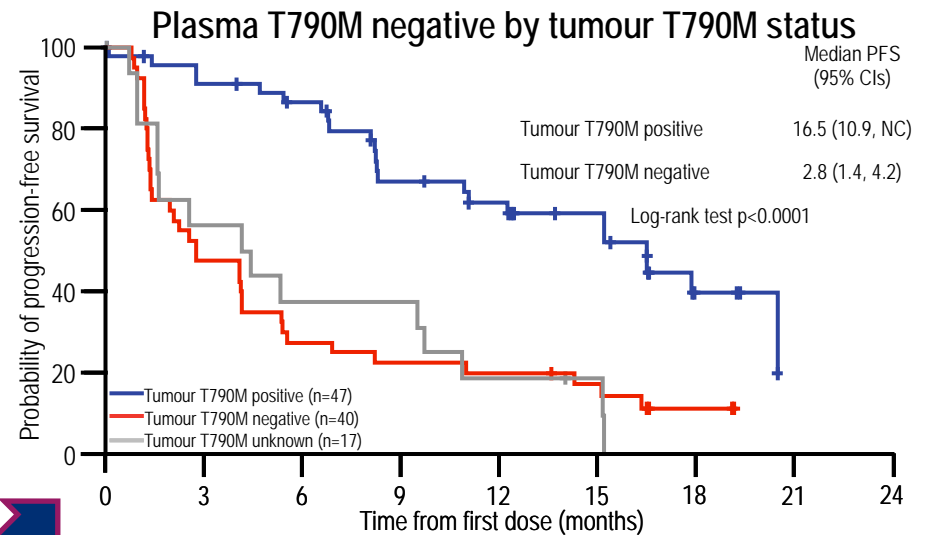


After PCR amplification, each droplet provides a fluorescent positive or negative signal indicating the target DNA was present or not present after partitioning. Each droplet provides an independent digital measurement.

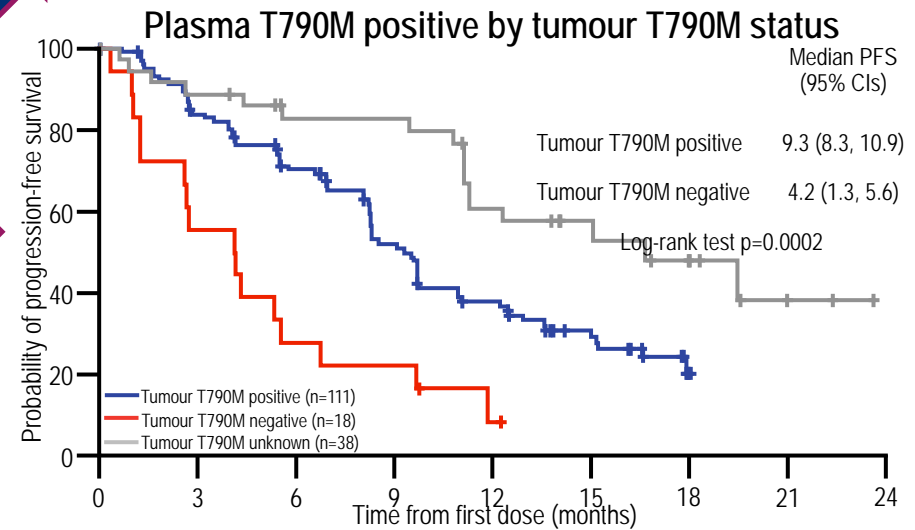
Downloaded from biodiscover.com

PFS by tumour and plasma T790M status

- In plasma T790M negative patients, tumour genotyping can distinguish those patients with better and worse outcomes
- Interestingly, a difference based on tumour genotype is also seen in plasma T790M positive cases

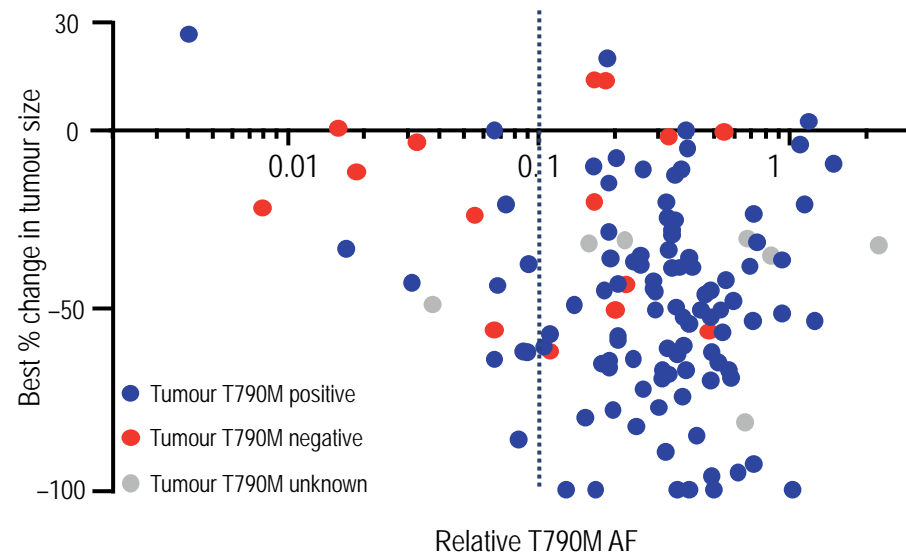
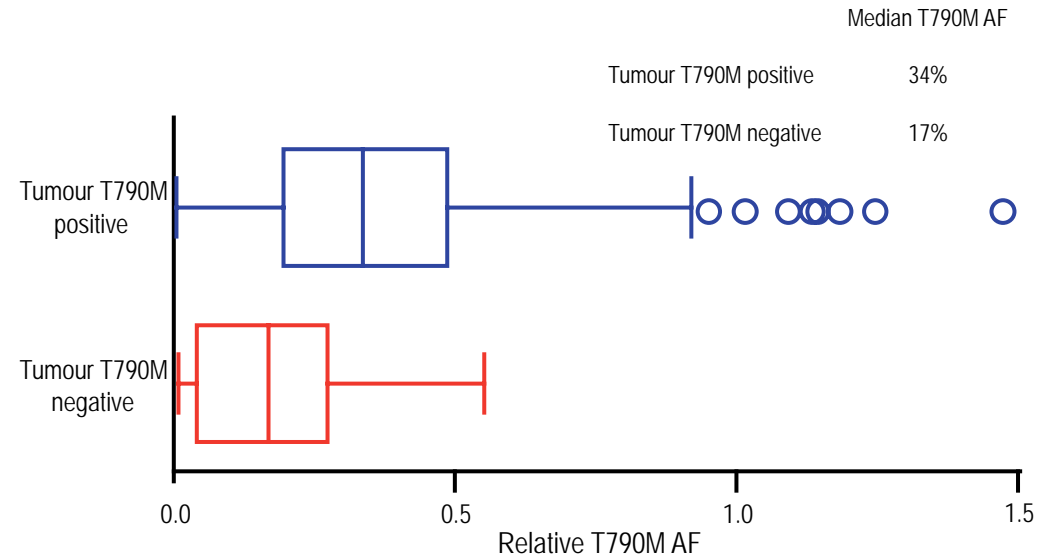


Data cutoff: 1 May 2015. Multiple doses included



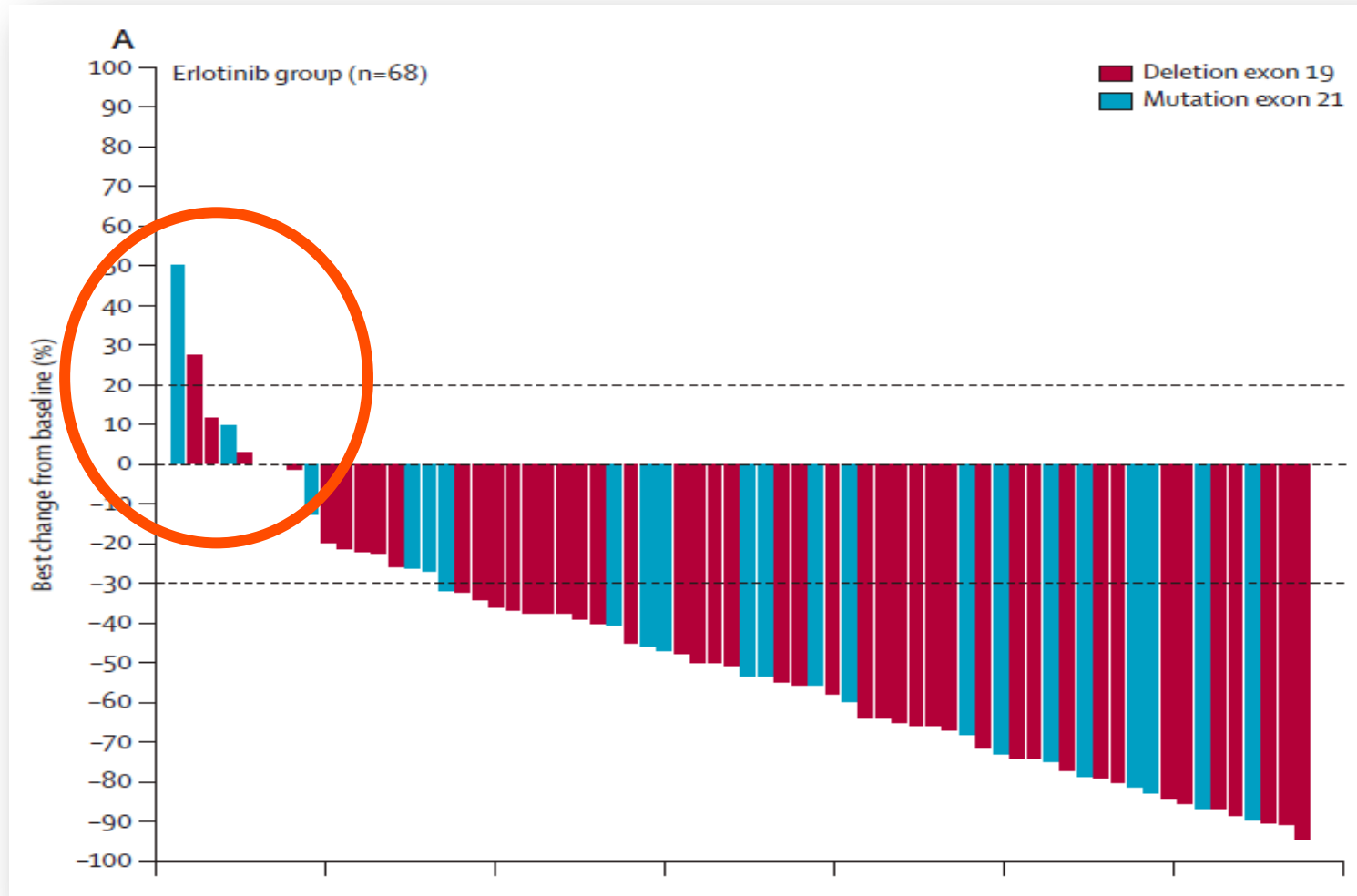
T790M heterogeneity in plasma “false positives”

- We hypothesised that cases T790M negative in tumour and T790M positive in plasma might have heterogeneous presence of T790M
- Relative T790M AF was calculated as a proportion of EGFR sensitising AF:
 - $\text{Relative T790M AF} = \text{T790M AF} / \text{sensitising AF}$
- Relative T790M AF was lower in cases with T790M negative in tumour, suggesting T790M may be present as a minor clone
- There was a trend toward lower response magnitude in the group with relative T790M AF <10% (p=0.08)



Data cutoff: 1 May 2015

Lung cancer EGFRm on 1G targeted therapy



Rosell R et al, Lancet Oncol 2012

Where comes the resistance from?

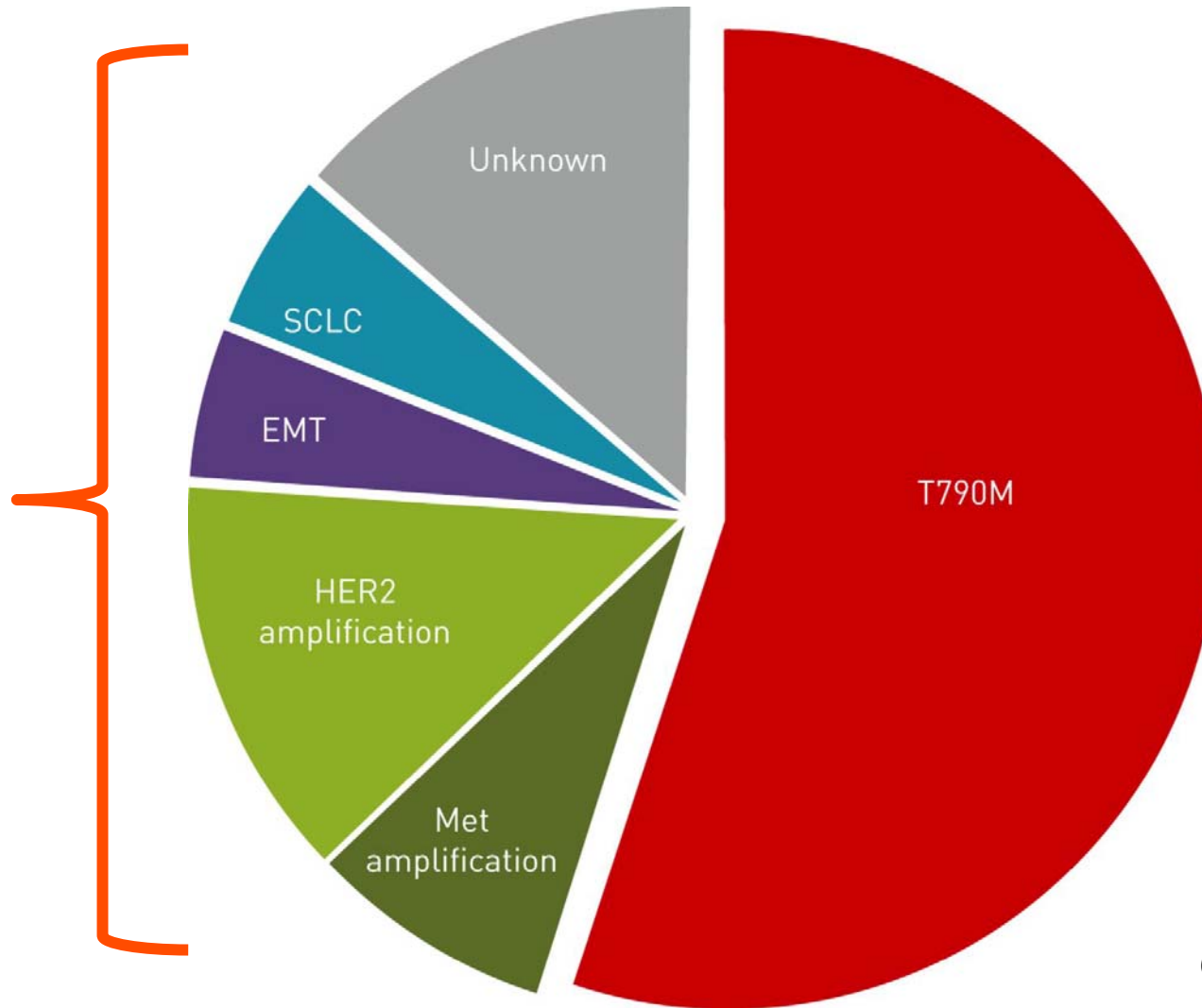
Tumor cells can follow distinct evolutionary paths to become resistant to epidermal growth factor receptor inhibition

Aaron N Hata^{1,2,14}, Matthew J Niederst^{1,2,14}, Hannah L Archibald¹, Maria Gomez-Caraballo¹, Faria M Siddiqui¹, Hillary E Mulvey¹, Yosef E Maruvka^{1,3}, Fei Ji⁴, Hyo-eun C Bhang⁵, Viveksagar Krishnamurthy Radhakrishna⁵, Giulia Siravegna^{6,7}, Haichuan Hu¹, Sana Raoof^{1,2}, Elizabeth Lockerman¹, Anuj Kalsy¹, Dana Lee¹, Celina L Keating⁵, David A Ruddy⁸, Leah J Damon¹, Adam S Crystal^{1,13}, Carlotta Costa^{1,2}, Zofia Piotrowska^{1,2}, Alberto Bardelli^{6,7}, Anthony J Iafrate⁹, Ruslan I Sadreyev^{4,9}, Frank Stegmeier⁵, Gad Getz^{1,3,9,10}, Lecia V Sequist^{1,2}, Anthony C Faber^{11,12} & Jeffrey A Engelman^{1,2}

Although mechanisms of acquired resistance of epidermal growth factor receptor (EGFR)-mutant non-small-cell lung cancers to EGFR inhibitors have been identified, little is known about how resistant clones evolve during drug therapy. Here we observe that acquired resistance caused by the *EGFR*^{T790M} gatekeeper mutation can occur either by selection of pre-existing *EGFR*^{T790M}-positive clones or via genetic evolution of initially *EGFR*^{T790M}-negative drug-tolerant cells. The path to resistance impacts the biology of the resistant clone, as those that evolved from drug-tolerant cells had a diminished apoptotic response to third-generation EGFR inhibitors that target *EGFR*^{T790M}; treatment with navitoclax, an inhibitor of the anti-apoptotic factors BCL-xL and BCL-2 restored sensitivity. We corroborated these findings using cultures derived directly from EGFR inhibitor-resistant patient tumors. These findings provide evidence that clinically relevant drug-resistant cancer cells can both pre-exist and evolve from drug-tolerant cells, and they point to therapeutic opportunities to prevent or overcome resistance in the clinic.

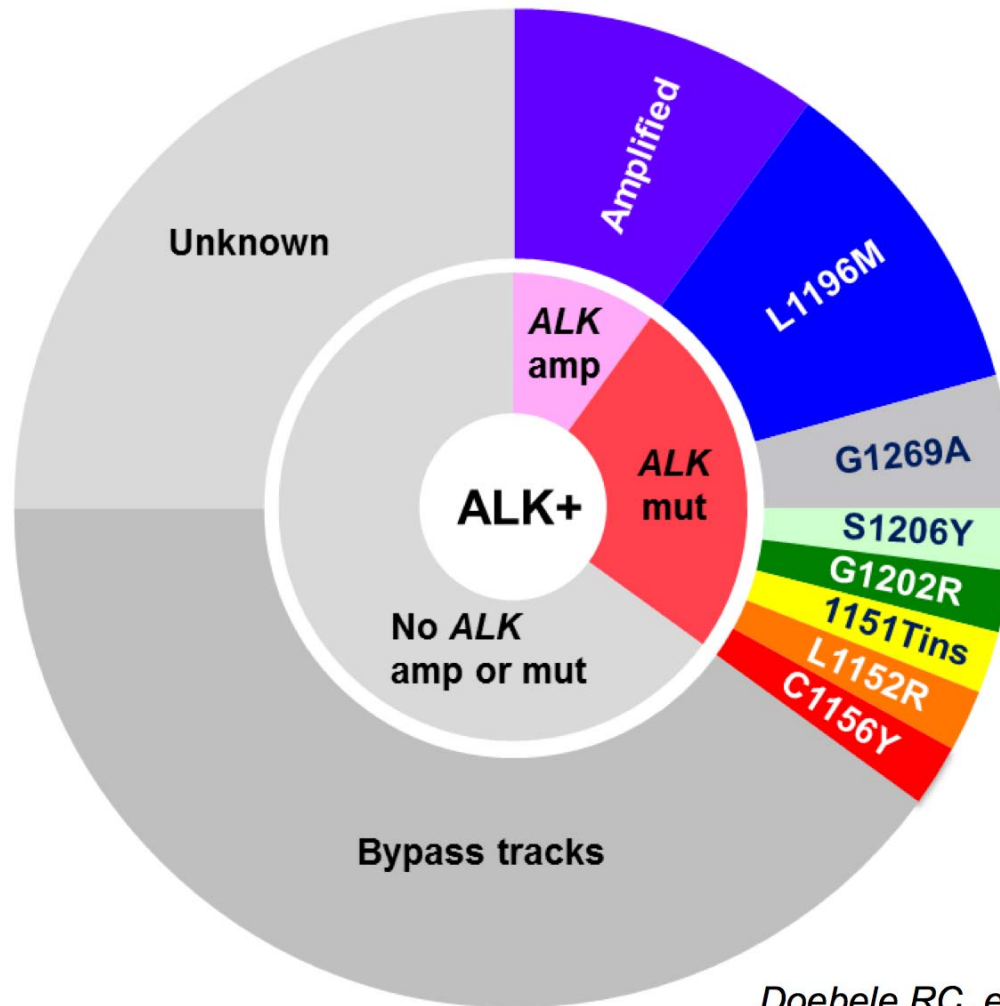
Hata A et al, Nature Med 2016

Résistance acquise aux EGFR TKI (1G)



Cortot A & Janne PA,
Eur Respir Rev 2014

Résistance acquise au crizotinib



*Doebele RC, et al. Clin Cancer Res 2012
Takeda M, et al. J Thorac Oncol 2013*

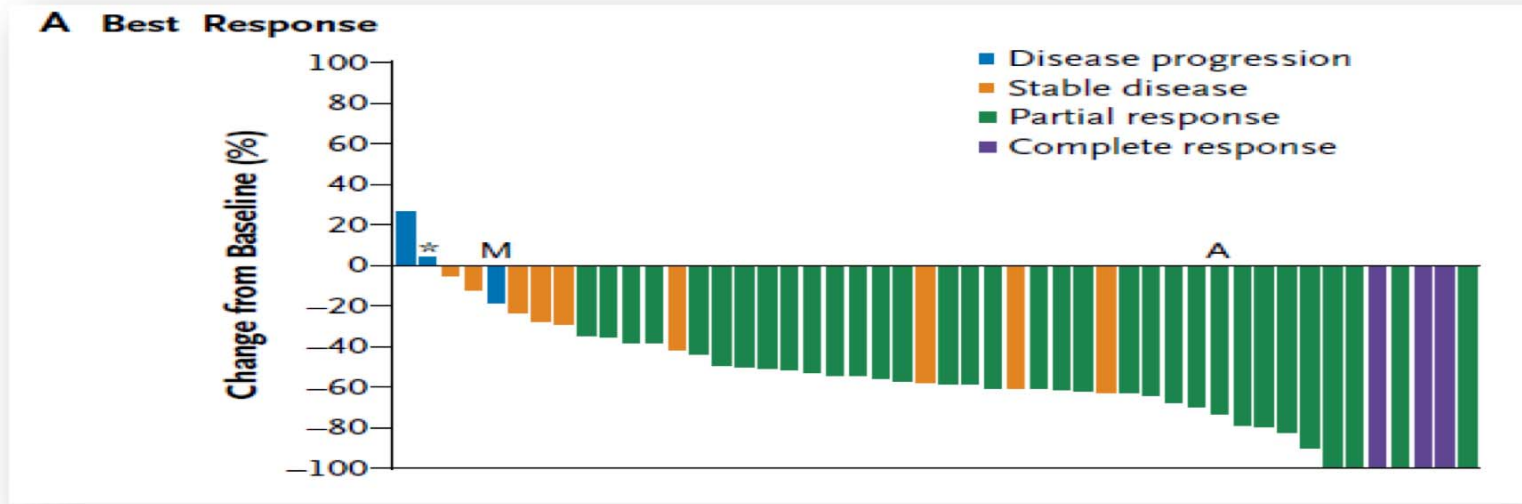
ALK mutations: frequencies

ALK inhibitor	Crizotinib	Ceretinib	Alectinib	PF-06463922
<i>Resistance mutation</i>				
G1123S		Resistant [113]	Sensitive [113]	
I151Tins	Resistant [79, 90]	Resistant [90]	Resistant [79]	Sensitive [87]
L1152P/R	Resistant [90]	Resistant [90]	Sensitive [114]	Sensitive [87]
C1156Y/T	Resistant [90]	Resistant [90]	Sensitive [87]	Sensitive [87]
I1171T/N	Resistant [90]	Sensitive [90, 115, 116]	Resistant [115, 116]	Sensitive [87]
F1174L/C	Resistant [90]	Resistant [90]	Sensitive [115]	Sensitive [87]
V1180L	Resistant [115]	Sensitive [115]	Resistant [115]	Sensitive [87]
L1196M	Resistant [78, 79, 83]	Sensitive [90]	Sensitive [87]	Sensitive [87]
G1202R	Resistant [79, 90]	Resistant [84, 90]	Resistant [79]	Sensitive [72, 87]
S1206Y	Resistant [79, 90]	Sensitive [90]	Sensitive [79]	Sensitive [87]
G1269A/S	Resistant [90]	Sensitive [90]	Sensitive [114]	Sensitive [87]

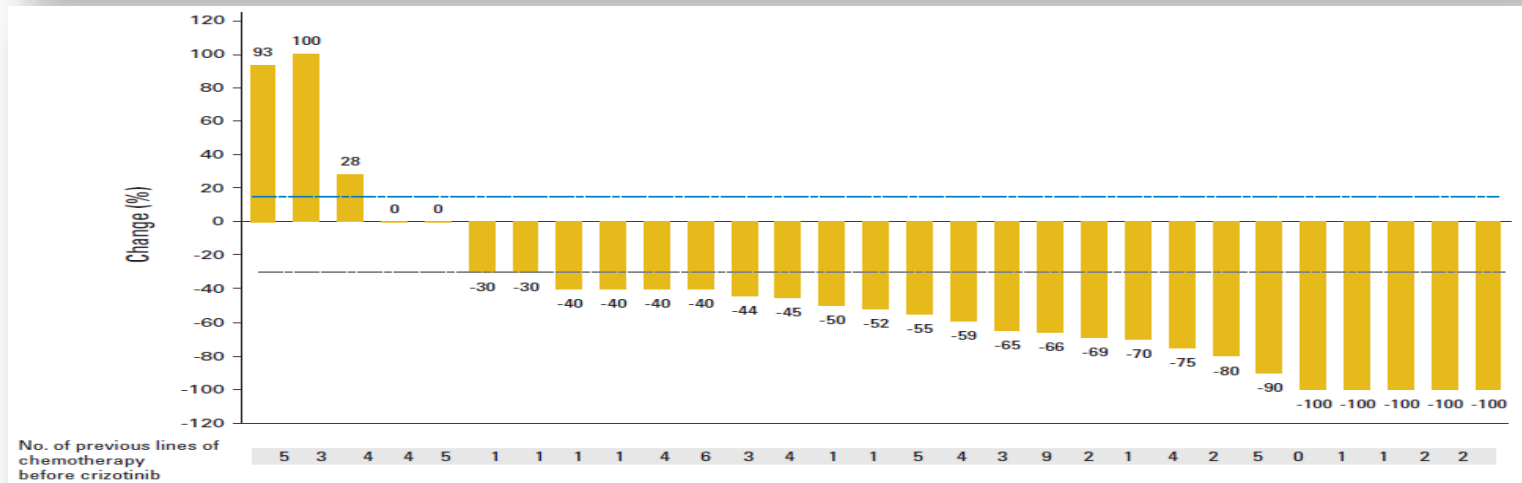
*Bayliss R et al,
Cell Mol Life Sci 2016*

Lung cancer ROS1 on targeted therapy

US Cohort

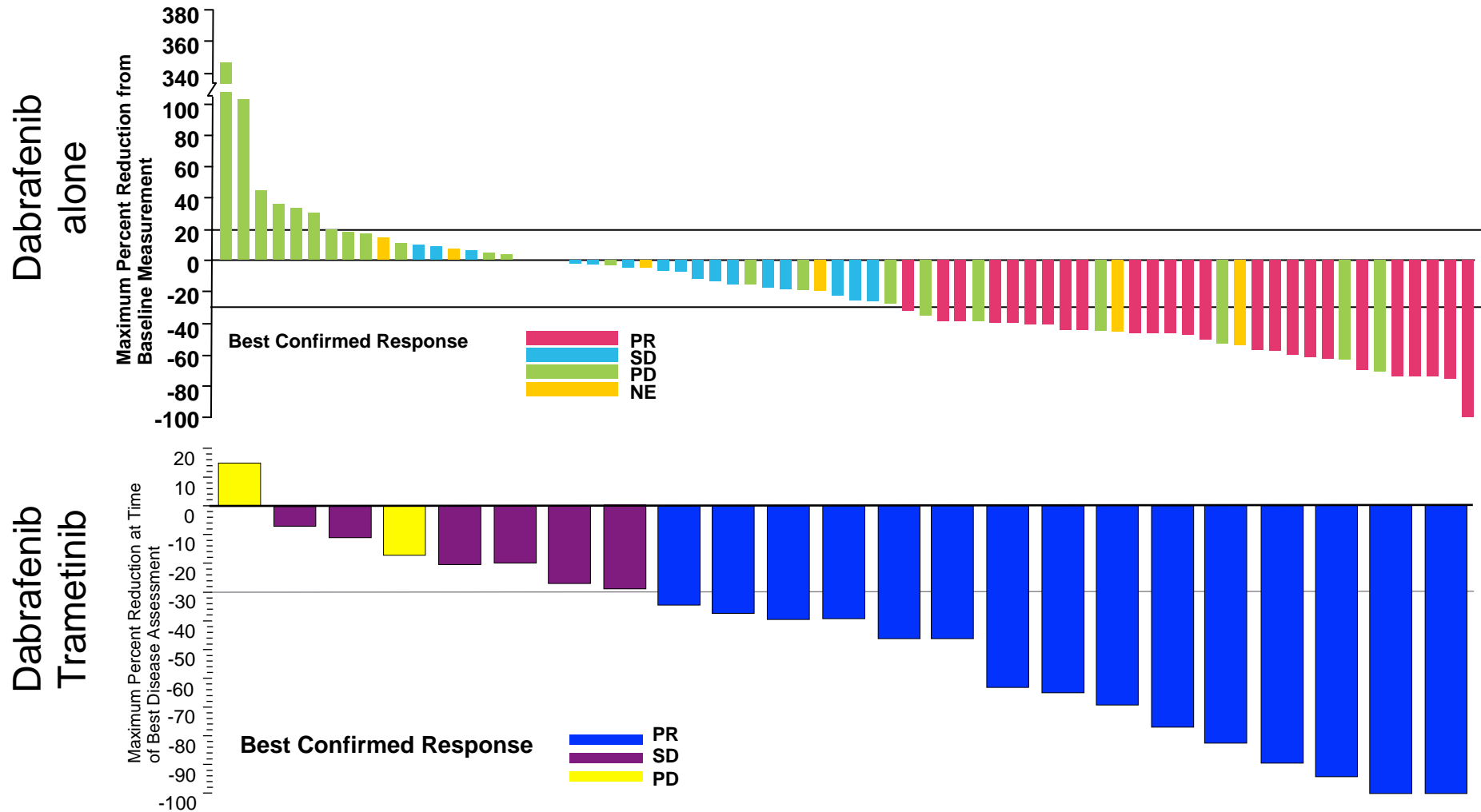


EU Cohort



Shaw A et al, NEJM 2014; Mazieres et al, J Clin Oncol 2015

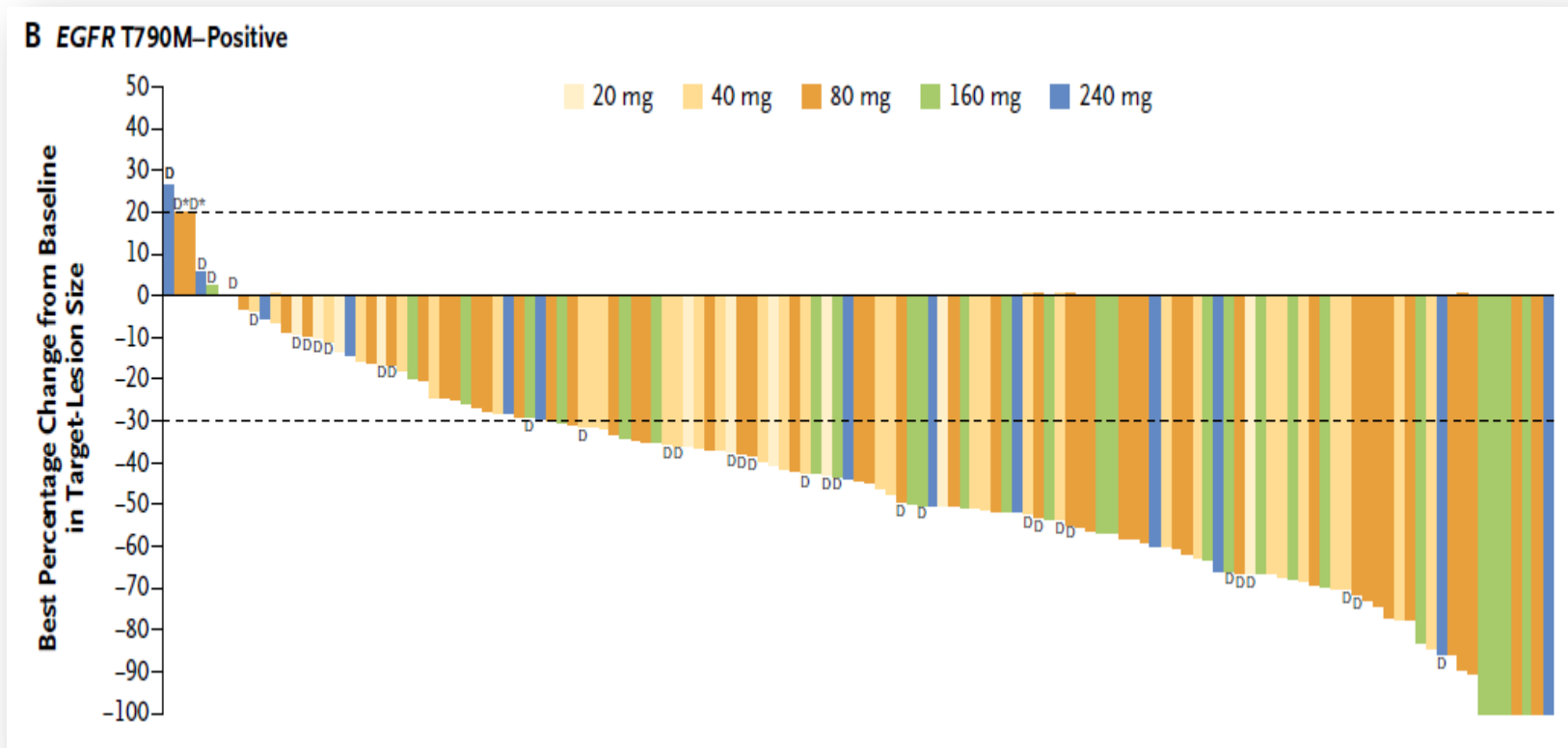
Lung cancer BRAFm on targeted therapy



Planchard et al, ASCO 2013; Planchard et al, ASCO 2016 & Lancet Oncol 2016 (in press)

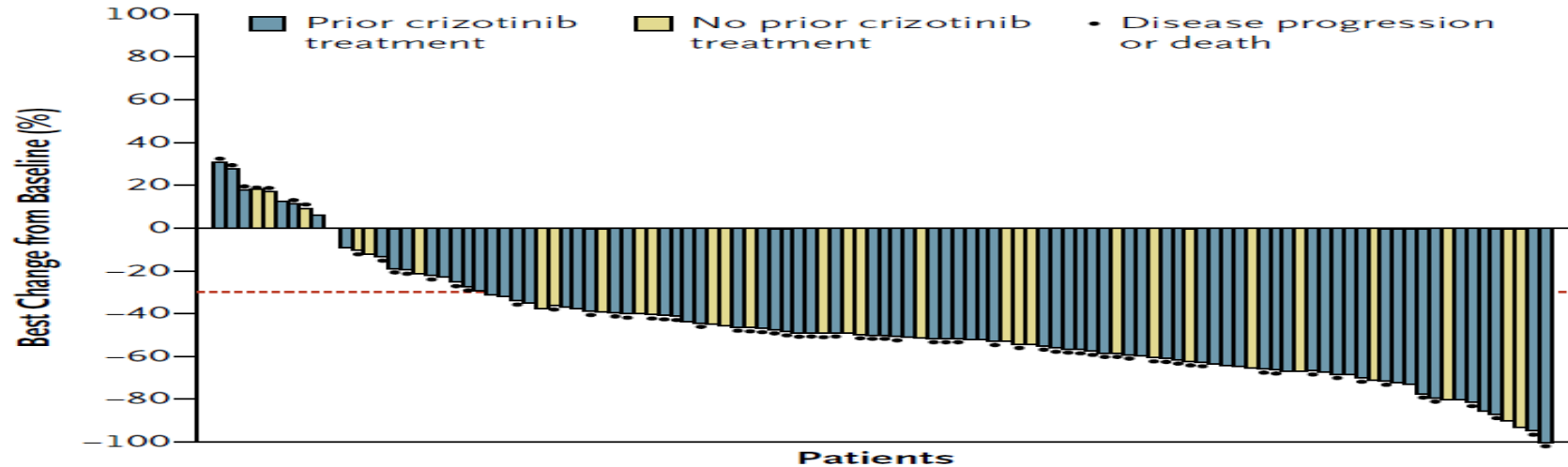
Lung cancer EGFRm on targeted therapy

AZD9291
Osimertinib



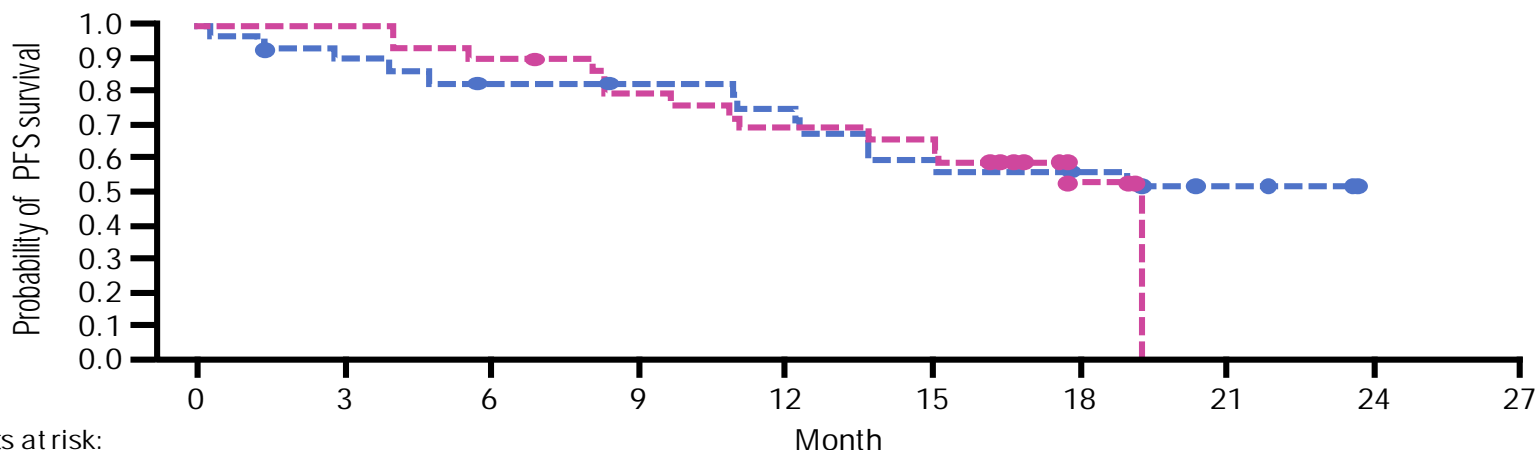
Janne P, NEJM 2015

Lung cancer ALKrearr on targeted therapy



Seto et al, Lancet Oncol 2013; Shaw A et al, NEJM 2014

3G EGFR-TKI in 1st line (AURA program)



Number of patients at risk:

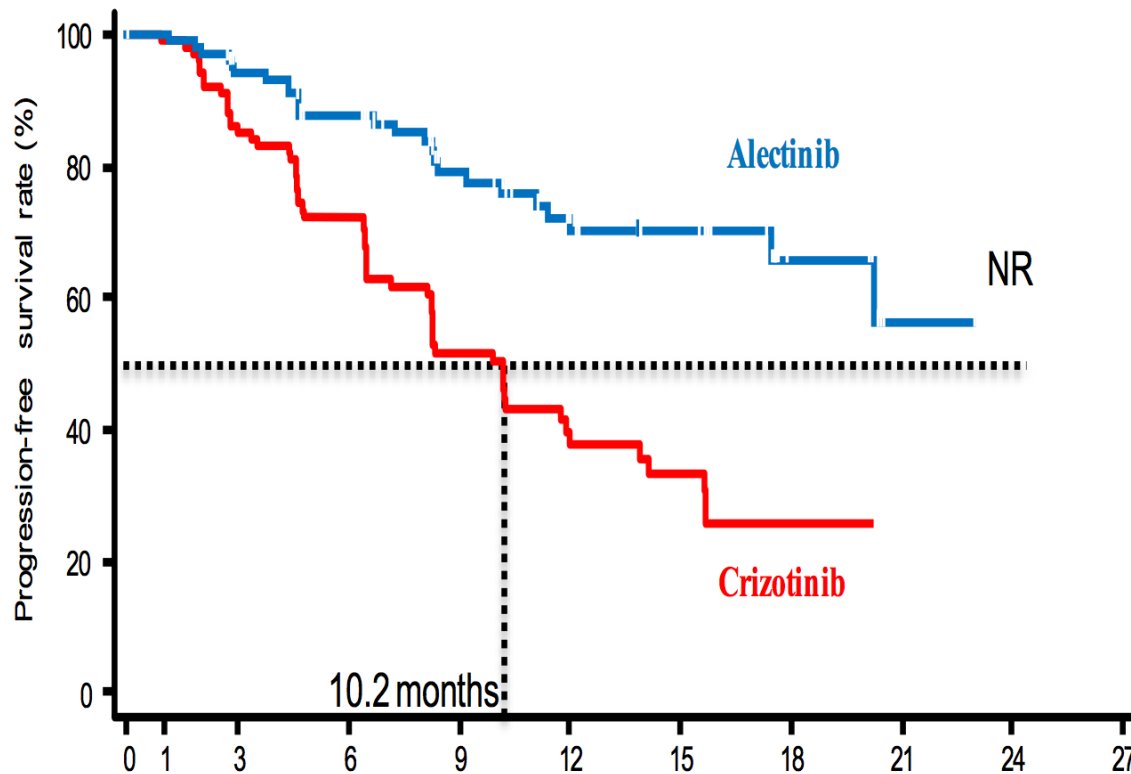
--- 1st line 80 mg
 --- 1st line 160 mg

Month	0	3	6	9	12	15	18	21	24	27
1 st line 80 mg	30	26	23	22	20	16	14	7	0	0
1 st line 160 mg	30	29	27	23	20	19	7	0	0	0

	80 mg n=30	160 mg n=30	Total N=60
Median PFS,* months (95% CI)	NC (12.3, NC)	19.3 (11.1, 19.3)	19.3 (13.7, NC)
Remaining alive and progression-free, [†] % (95% CI)			
12 months	75 (55, 88)	69 (49, 83)	72 (59, 82)
18 months	57 (36, 73)	53 (32, 70)	55 (41, 67)

Ramalingam S, et al. ELCC 2016; Abstract LBA1_PR

3G ALKi Alectinib in 1st line



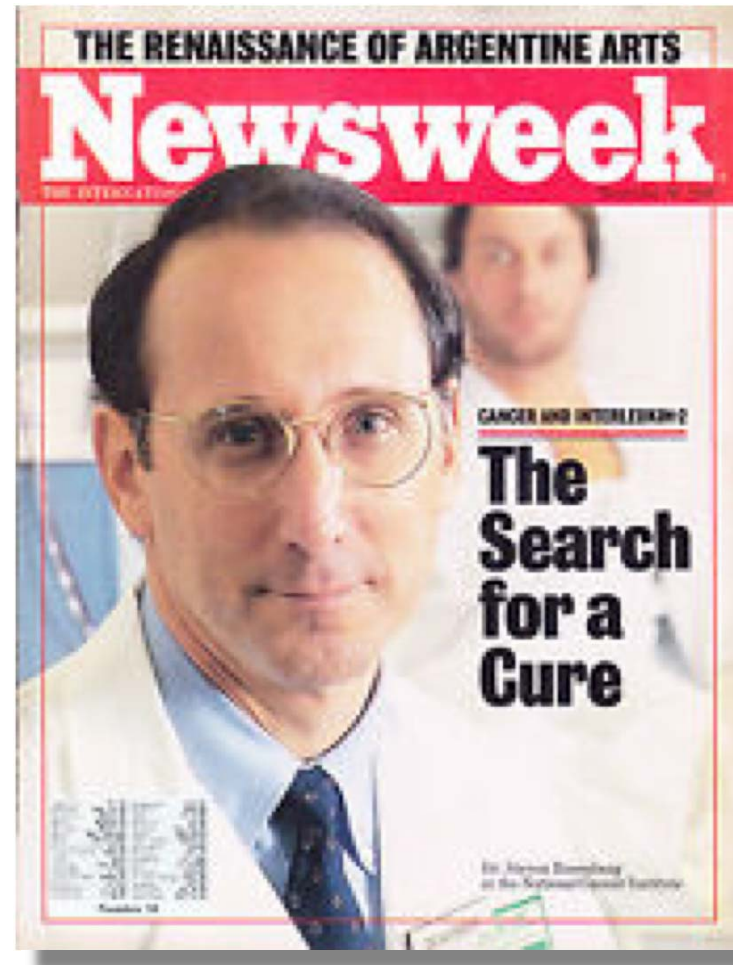
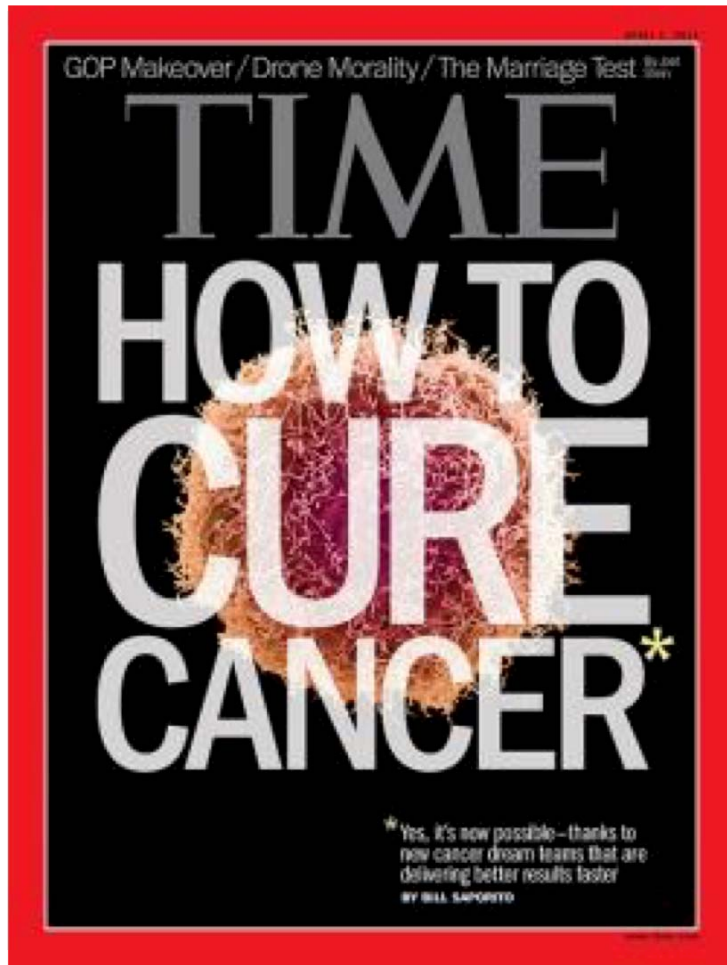
	Alectinib (N=103)	Crizotinib (N=104)
Events, n (%)	25 (24.3%)	58 (55.8%)
Median, mo [95% CI]	NR [20.3 - NR]	10.2 [8.2 - 12.0]
P-value	<0.0001	
HR [95% CI]	0.34 [0.17 - 0.71]	

Nokihara H et al, Abst #9008 ASCO 2016

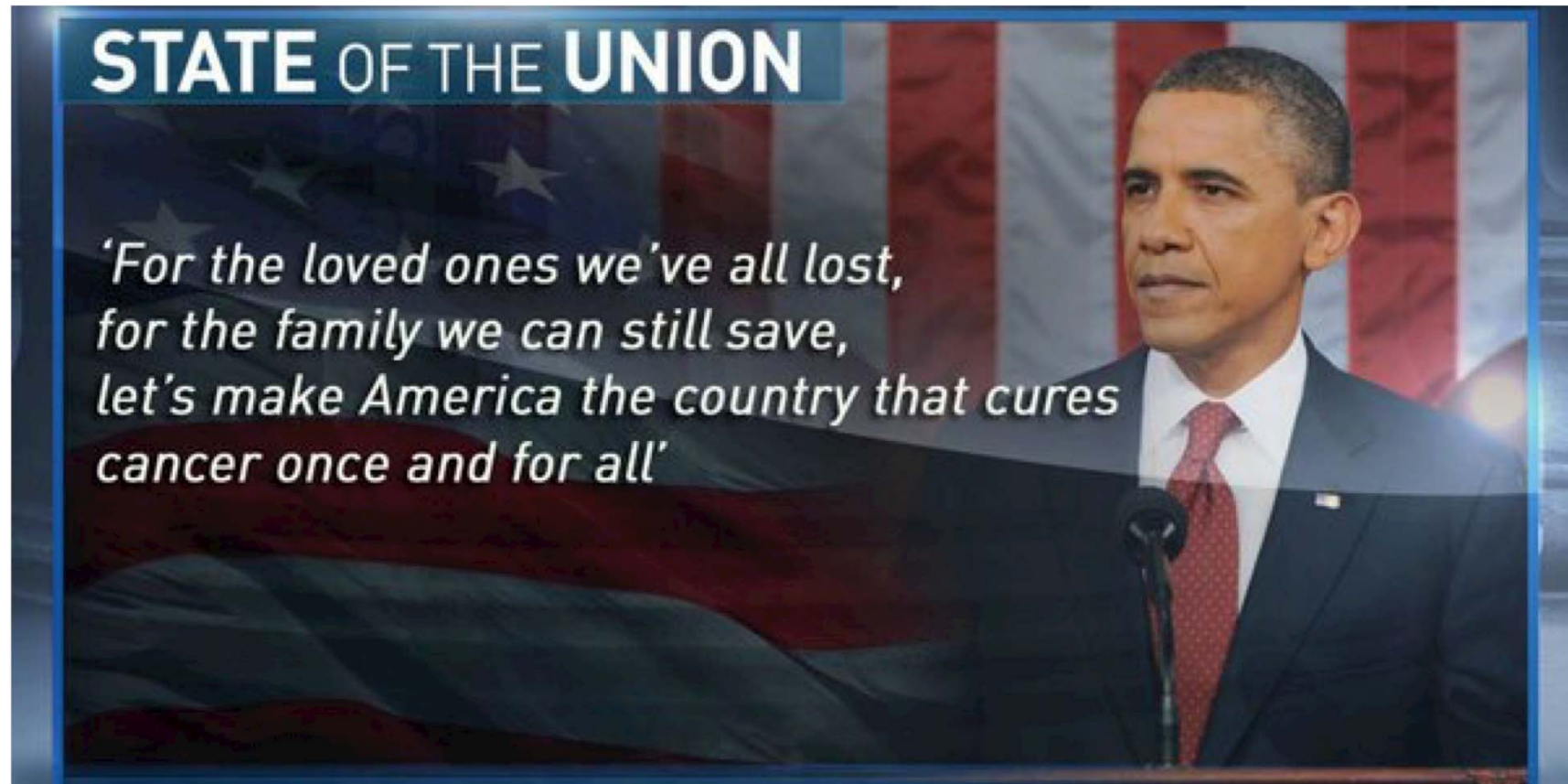
En résumé ... 2020

- **Poussée technologique**
- **Réorganisation (régionale)**
- **Peu de nouvelles cibles actionnables**
 - **KRAS: untargetable target?**
 - **Process décision (plus) complexe**
- **Passage en 1^{ère} ligne TKI 3G**

Les attentes ne sont pas là ...



US Cancer Moonshot

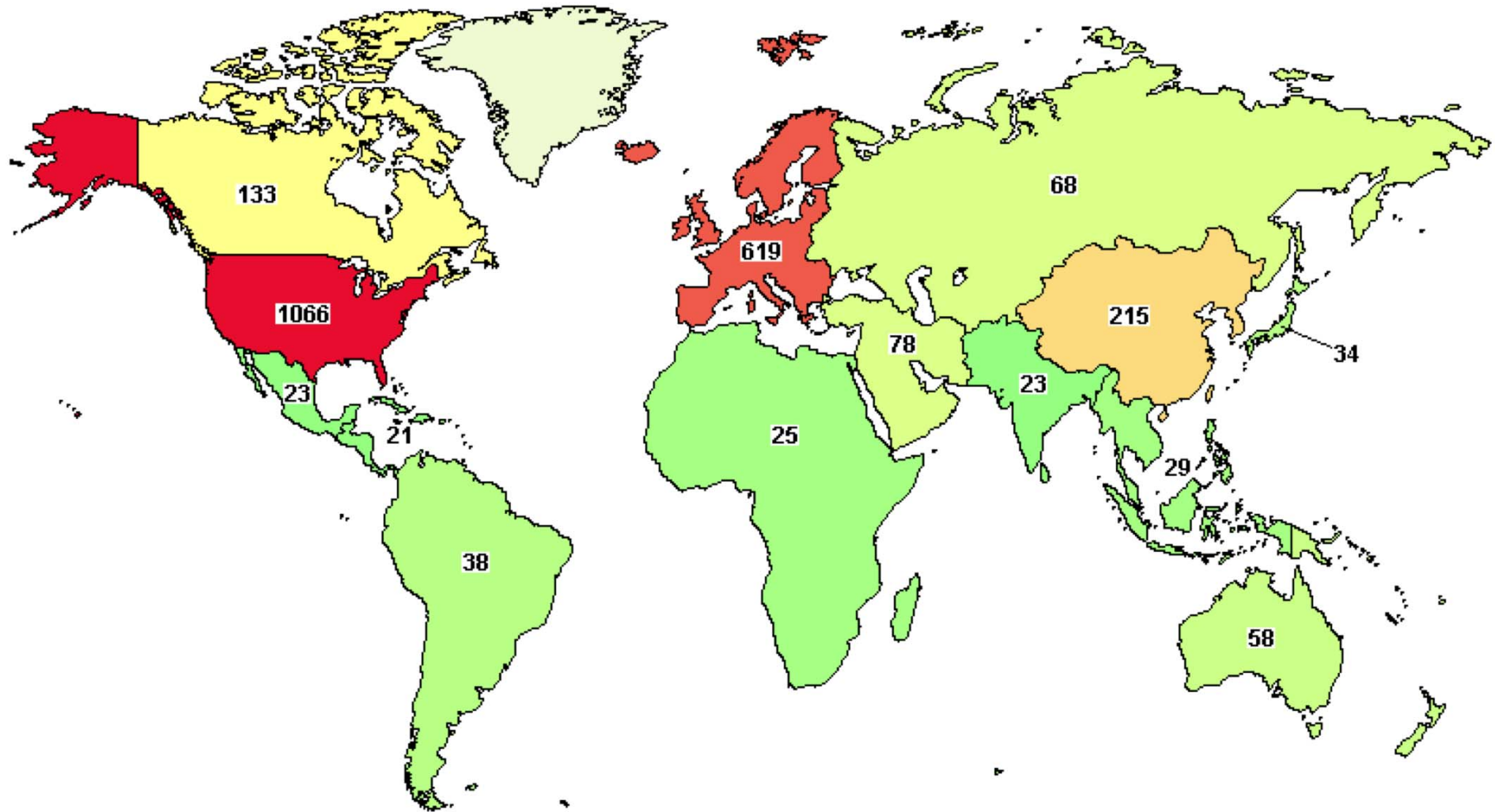


Obama, January 2016

Agenda

- Le whole genome pour tous ?
- **L'immunothérapie, le graal ?**
- Faire du neuf avec du vieux ?

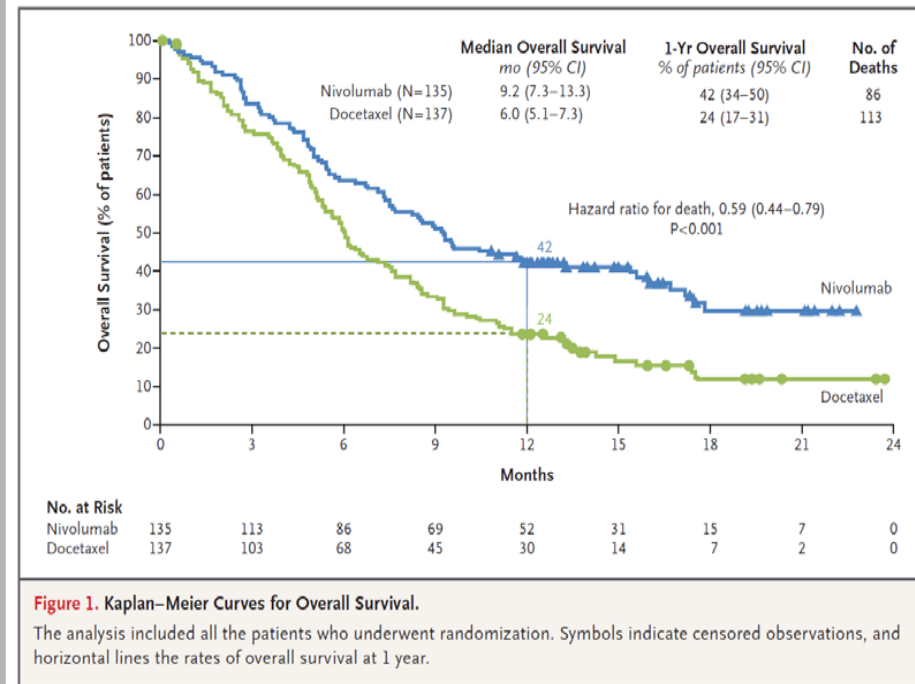
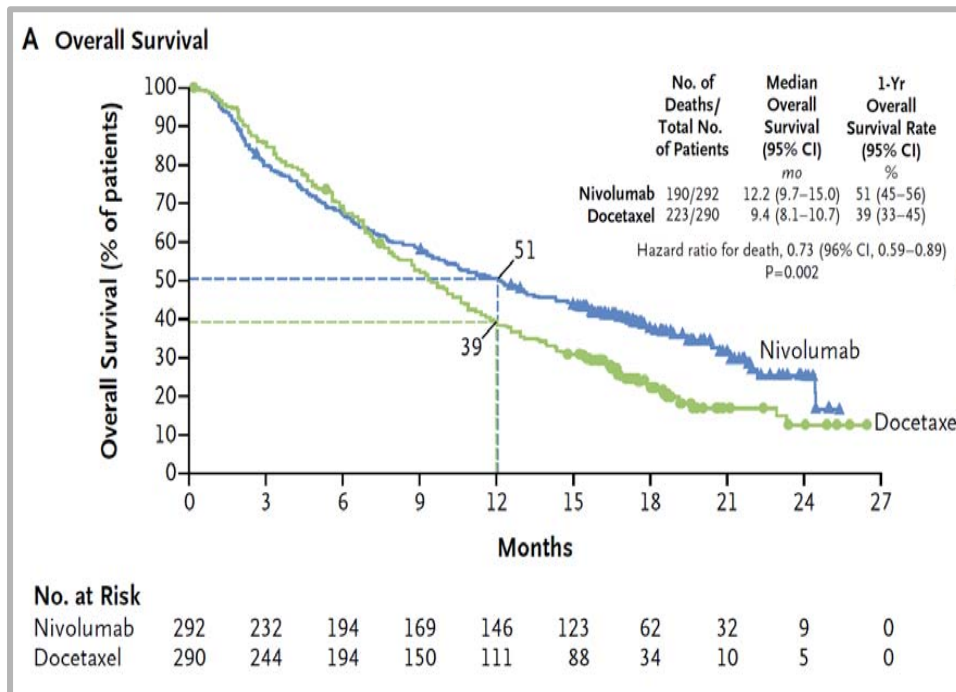
Consequences



PD1 inhibitor 2L: Nivolumab (PhIII)

Nsq-NSCLC

Sq-NSCLC

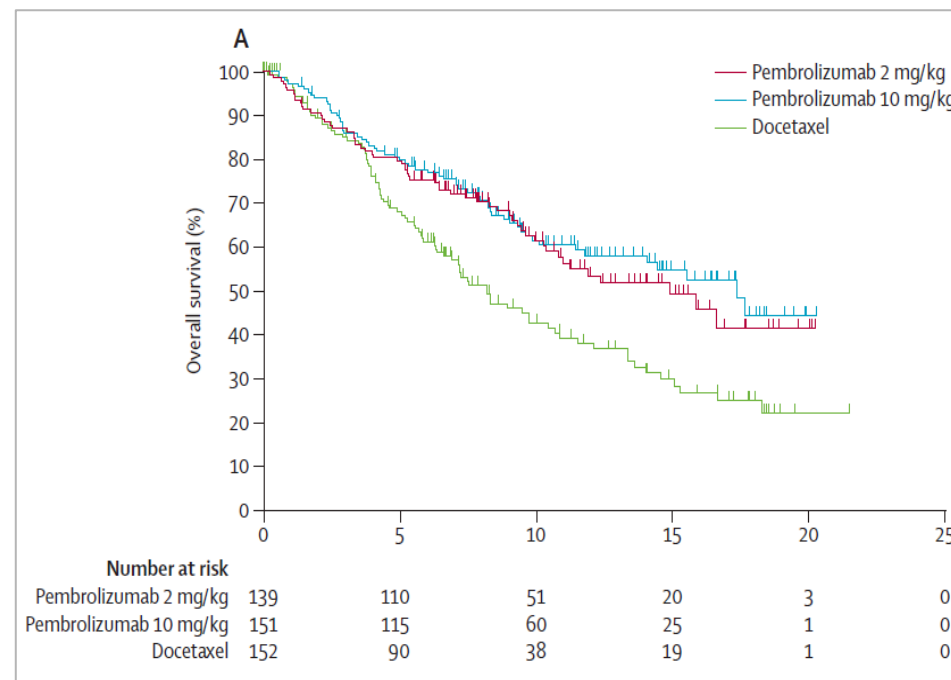
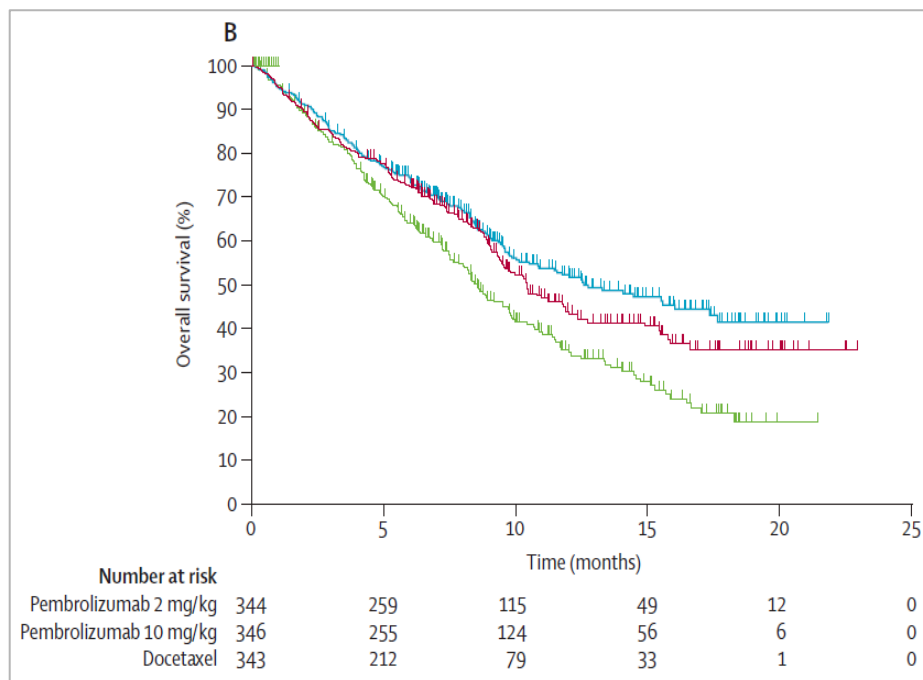


Brahmer J et al, NEJM 2015; Borghaei H et al, NEJM 2015

PD1 inhibitor 2L: Pembrolizumab (PhIII)

AII NSCLC

NSCLC w PD-L1+ >50%



HR for OS (doc vs 2): 0,71 (0,58-0,88)
 HR for OS (doc vs 10): 0,61 (0,49-0,75)

HR for OS (doc vs 2): 0,54 (0,38-0,77)
 HR for OS (doc vs 10): 0,50 (0,36-0,70)

Herbst R et al., Lancet 2016

PD-L1 inhibitor 2L: Atezolizumab (PhIII)

Wednesday, Aug 31, 2016

Phase III Study Showed Genentech's Cancer Immunotherapy
TECENTRIQ™ (Atezolizumab) Helped People with a Specific
Type of Lung Cancer Live Significantly Longer Compared to
Chemotherapy

- **TECENTRIQ showed significant improvement in overall survival for people regardless of their PD-L1 status**

To be presented at ESMO (Presidential session, Sunday 9th)

Intégration immunothérapie

Drug	FDA Approval	EMA Approval	Price (France)
Nivolumab (Sq) 2L	Mar 2015	Jul 2015	
Nivolumab (Nsq) 2L	Oct 2015	Apr 2016	
Pembrolizumab (All PD-L1+) 2L	Oct 2015	Submitted	
Atezolizumab (All) 2L	2017?	2018?	

En 1ère ligne dès 2017 ?



Bristol-Myers Squibb

Bristol-Myers Squibb Announces Top-Line Results from CheckMate -026, a Phase 3 Study of Opdivo (nivolumab) in Treatment-Naïve Patients with Advanced Non-Small Cell Lung Cancer

08/05/2016

Opdivo did not meet trial primary endpoint of progression-free survival in patients expressing PD-L1 = 5%

PRINCETON, N.J.--(BUSINESS WIRE)-- [Bristol-Myers Squibb Company](#) (NYSE:BMJ) announced today that CheckMate -026, a trial investigating the use of *Opdivo* (nivolumab) as monotherapy, did not meet its primary endpoint of progression-free survival in patients with previously untreated advanced non-small cell lung cancer (NSCLC) whose tumors expressed PD-L1 at = 5%. The company will complete a full evaluation of the CheckMate -026 data and work with investigators on the future presentation of the results.

To be presented at ESMO (Presidential session)

En 1ère ligne dès 2017 !



Merck's KEYTRUDA[®] (pembrolizumab) Demonstrates Superior Progression-Free and Overall Survival Compared to Chemotherapy as First-Line Treatment in Patients with Advanced Non-Small Cell Lung Cancer

KEYNOTE-024 Studied Patients Whose Tumors Expressed High Levels of PD-L1

June 16, 2016 06:45 AM Eastern Daylight Time

KENILWORTH, N.J.--(BUSINESS WIRE)--Merck (NYSE:MRK), known as MSD outside the United States and Canada, today announced that the KEYNOTE-024 trial investigating the use of KEYTRUDA[®] (pembrolizumab), in patients with previously untreated advanced non-small cell lung cancer (NSCLC) whose tumors expressed high levels of PD-L1 (tumor proportion score of 50 percent or more), met its primary endpoint. In this trial, KEYTRUDA was superior compared to chemotherapy for both the primary endpoint of progression-free survival (PFS), and the secondary endpoint of overall survival (OS). Based on these results, an independent Data Monitoring Committee (DMC) has recommended that the trial be stopped, and that patients receiving chemotherapy in KEYNOTE-024 be offered the opportunity to receive KEYTRUDA.

To be presented at ESMO (Presidential session)

En 1ère ligne dès 2017 !

Essai combinaison

Cx +/- ICI

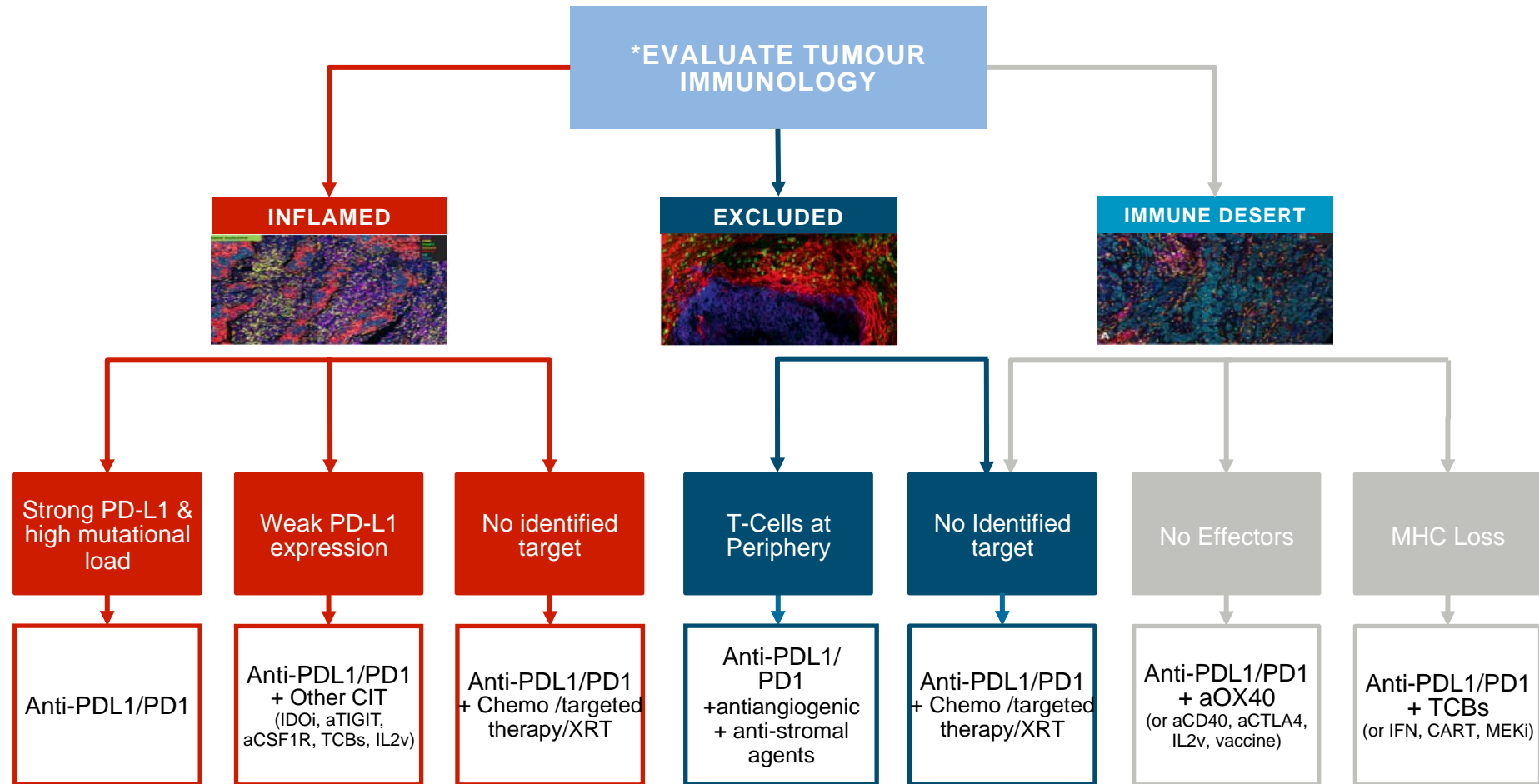
To be possibly presented at ESMO (Presidential session)

PD-L1 expression as a predictive factor?

Drug	Biomarker body	Rx line	Definition of 'Positive' # %	N Positive %	Positive Predictive outcome	ORR % IHC pos cases	ORR % IHC neg cases
Highly +	Nivolumab	1st	≥5% in >100 cells	59%	Yes	31%*	10%
	Nivolumab	≥2 nd	≥5%	49%	No	15%	14%
	Nivolumab	1st	≥1% ≥5% in >100 cells	56% #20	No	13% 19%	17% 14%
Weakly +	Nivolumab	≥2 nd	≥5%	33%##	Yes	24%	14%
	Nivolumab	≥2 nd	≥5% Also studied TIICs	67%	Yes	<i>No data For lung</i>	<i>No data For lung</i>
	Pembrolizumab	any	'Strong' ≥50% 'Weak' 1-49%	25% #40 40%	Yes Yes	37% 17%	9%
Negative	Pembrolizumab	1st	≥50% ≥1%	? #40	Yes	47% 26%	? 20%
	MPDL3280A	≥2 nd	≥10% TIICs*** ≥5% TIICs ≥1% TIICs	13% 28% #40 66%	Yes	83% 46% 31%	18% 18% 20%
	MEDI-4736	≥2 nd	<i>Data not available</i>	41%	yes	25%	3%

Kerr K et al, J Thorac Oncol (in press)

Personnalized IO? > 2020

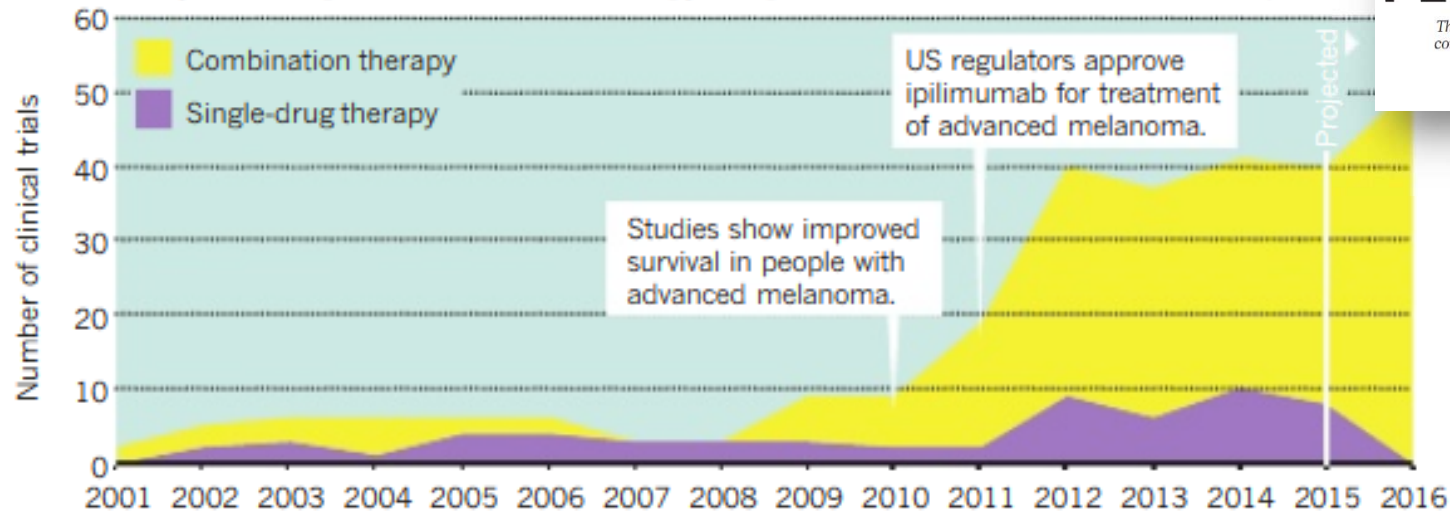


Modified from Kim and Chen, Ann Oncol 2016

Enjeux et Perspectives

COMBINATORIAL EXPLOSION

Ipilimumab, the first approved checkpoint inhibitor, has been tested in dozens of clinical trials since 2001. And like many other drugs in its class, it is increasingly being tested in combination with other therapies.



THE PERFECT BLEND

The next frontier in cancer immunotherapy lies in combining it with other treatments. Scientists are trying to get the mix just right.

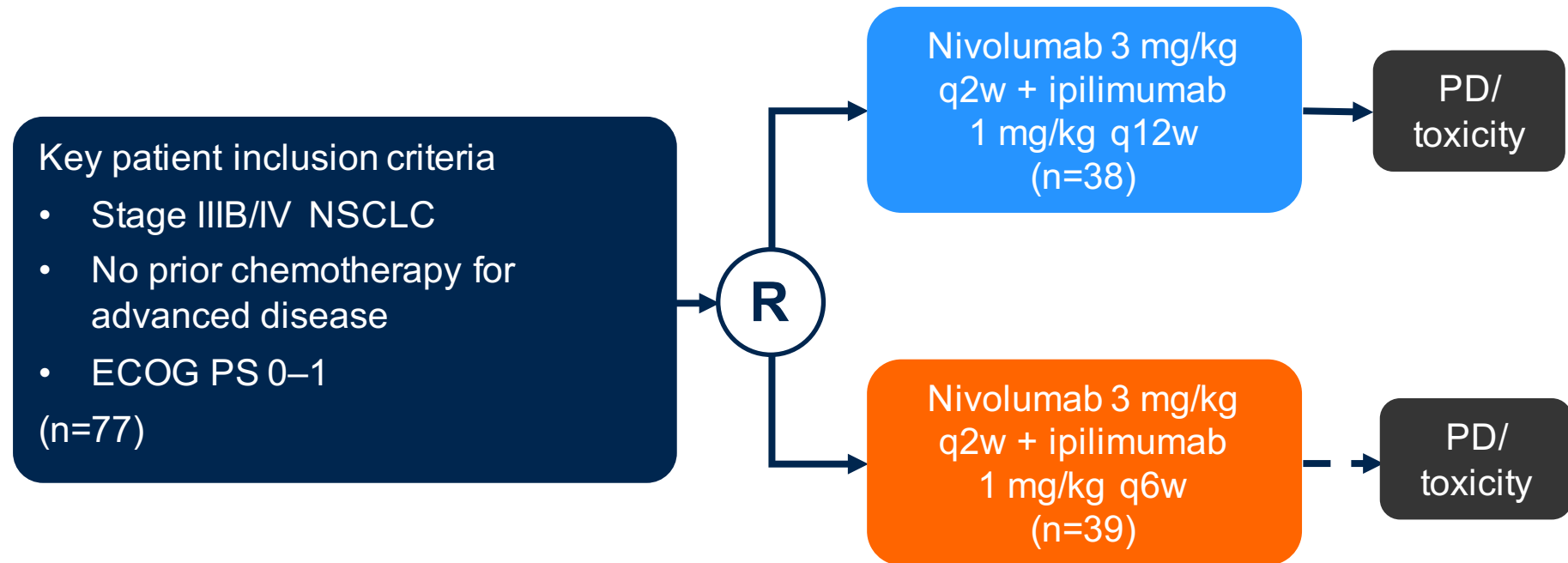
BY HEIDI LEFORD



« There will be not enough money on this earth to test all the possible combinations »

Ira Mellman, Vice-President, Cancer Immunology, Genentech Inc.
AACR, New Orleans April 2016

PD-1 inhibitor 1L: Nivolumab



Primary endpoint

- Safety/tolerability

Secondary endpoints

- ORR, PFS, OS, efficacy by PD-L1 expression

Hellman et al., ASCO 2016 (abst 3001)

PD-1 inhibitor 1L: Nivolumab/Ipilimumab

	Nivo 3 q2w + Ipi 1 q12w (n=38)	Nivo 3 q2w + Ipi 1 q6w (n=39)	Nivo 3 q2w (n=52)
Confirmed ORR, %	47	39	23
Median DOR, months (95%CI)	NR (11.3, NR)	NR (8.4, NR)	NR (5.7, NR)
Median follow-up, months (95%CI)	12.9 (0.9, 18.0)	11.8 (1.1, 18.2)	14.3 (0.2, 30.1)
mPFS, months (95%CI)	8.1 (5.6, 13.6)	3.9 (2.6, 13.2)	3.6 (2.3, 6.6)
1-year OS rate, % (95%CI)	NC	69 (52, 81)	73 (59, 83)

Efficacy is enhanced with increasing PD-L1 expression:

≥1% tumour PD-L1 expression: **57% ORR**; 83–90% 1-year OS rate

≥50% tumour PD-L1 expression: **92% (12/13) ORR**

Hellman et al., ASCO 2016 (abst 3001)

En résumé ... 2020

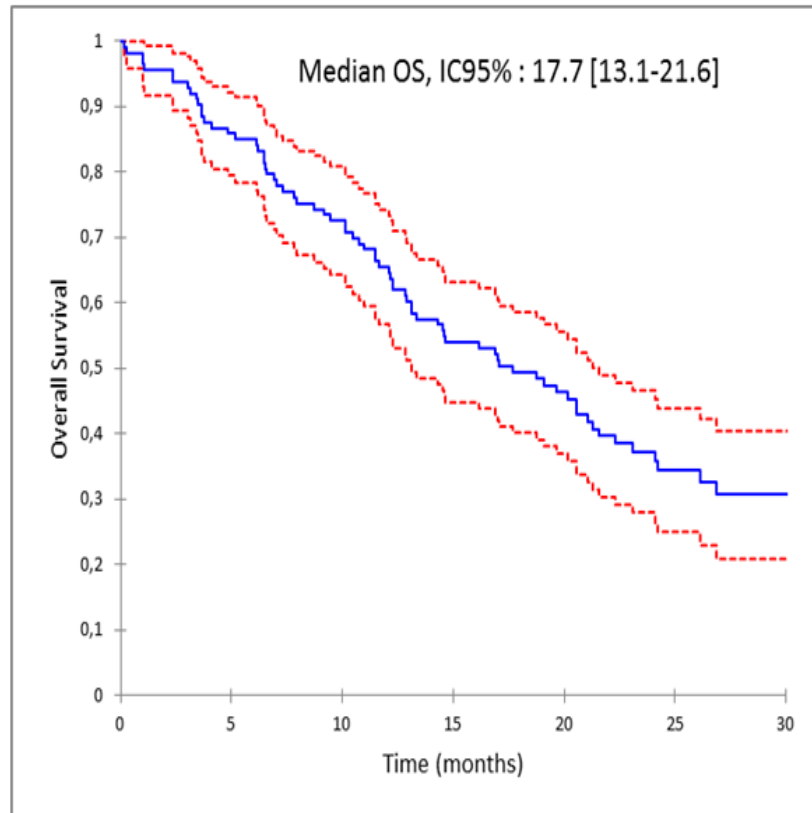
- **ICI 1^{ère} ligne 2017-2018 (20% ?)**
- **ICI 2^{ème} ligne 2016 (sans sélection)**
- **Combinaisons**
 - ICIs (2018-2020)
 - ICI + CT (2018-2020)

Agenda

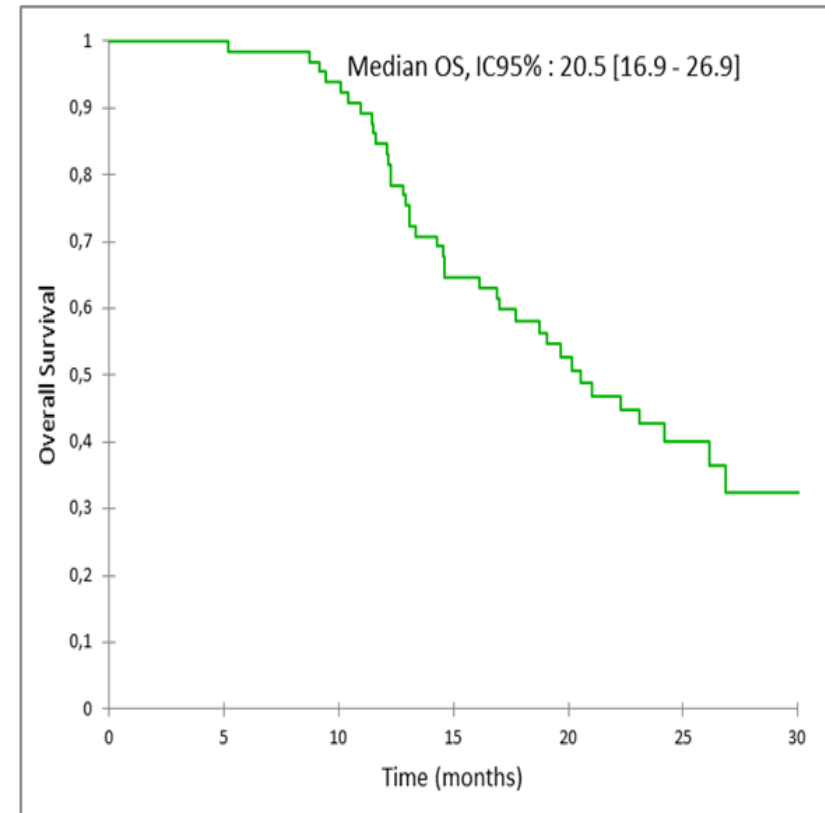
- Le whole genome pour tous ?
- L'immunothérapie, le graal ?
- **Faire du neuf avec du vieux ?**

Angiogenics (BUCIL)

OS of all eligible patients (n=113)

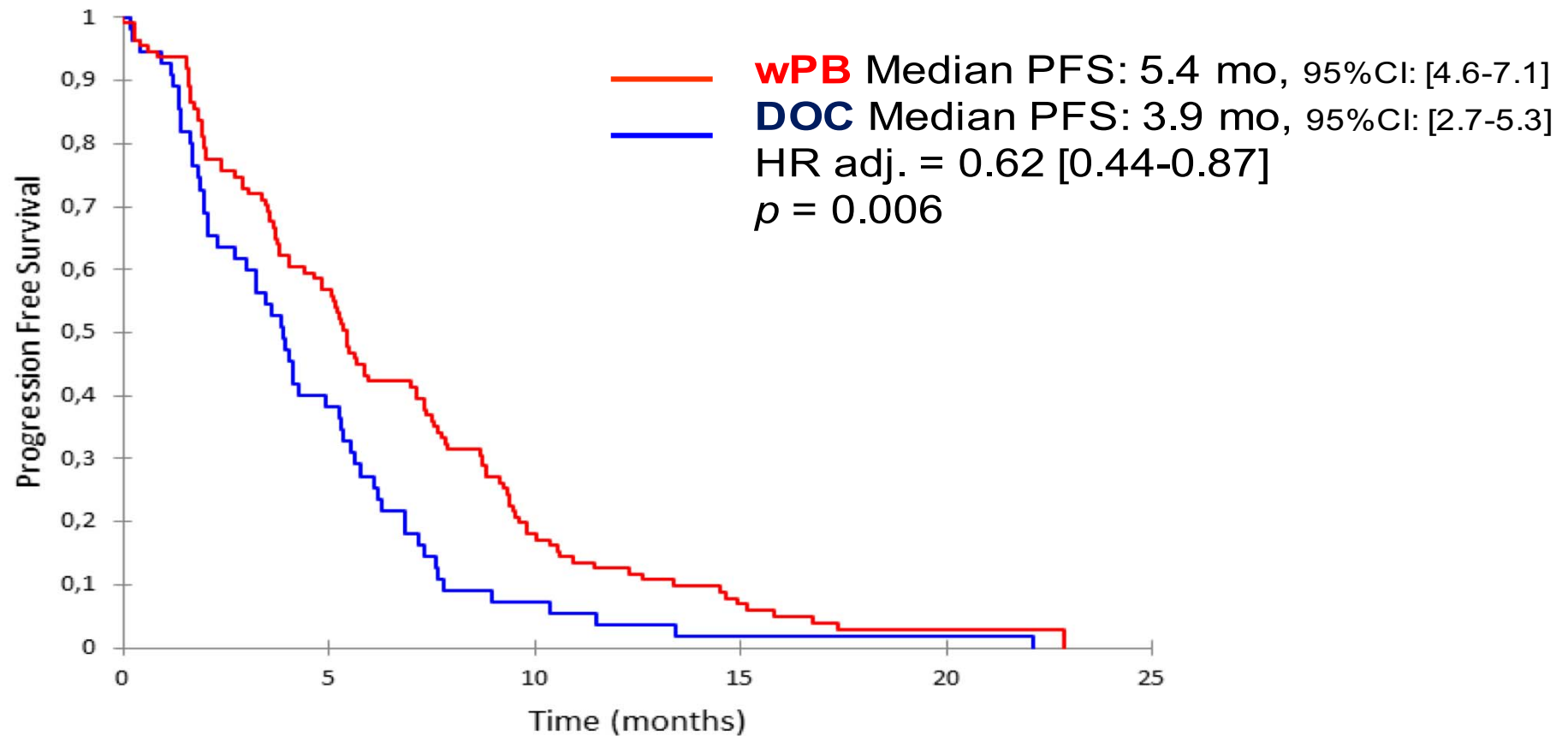


OS of patients receiving sequence 2 (n=65)



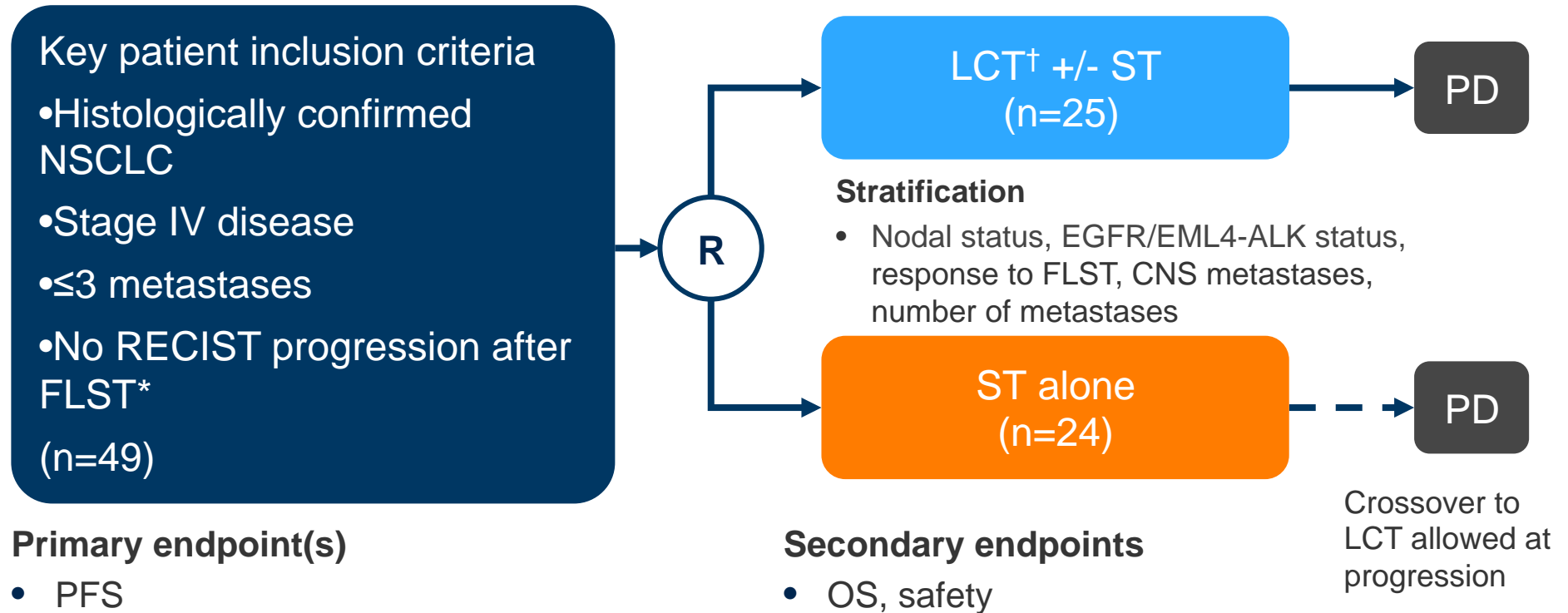
Barlesi F et al, Abst #9077 ASCO 2016

Angiogenics (ULTIMATE)



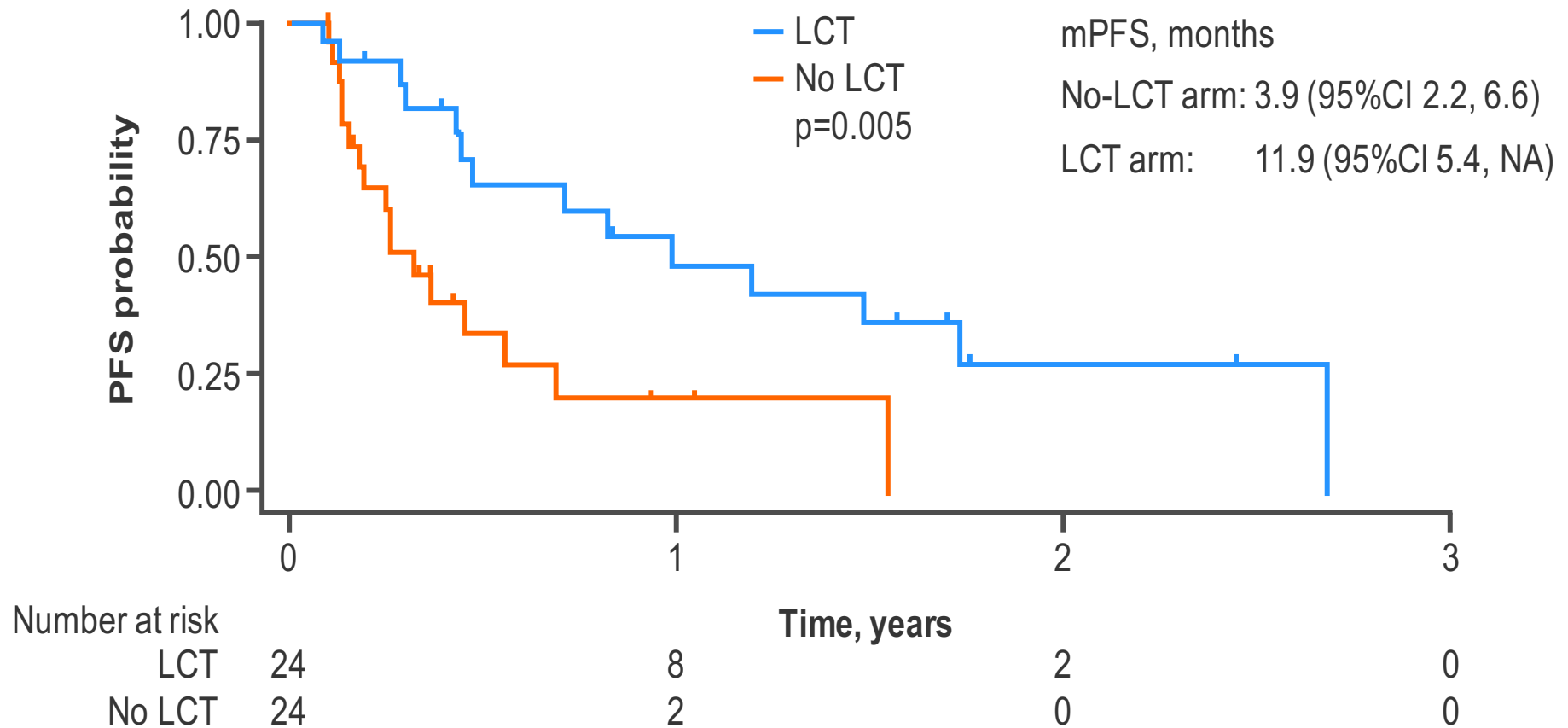
Cortot A et al, Abst #9077 ASCO 2016

Enjeux et Perspectives



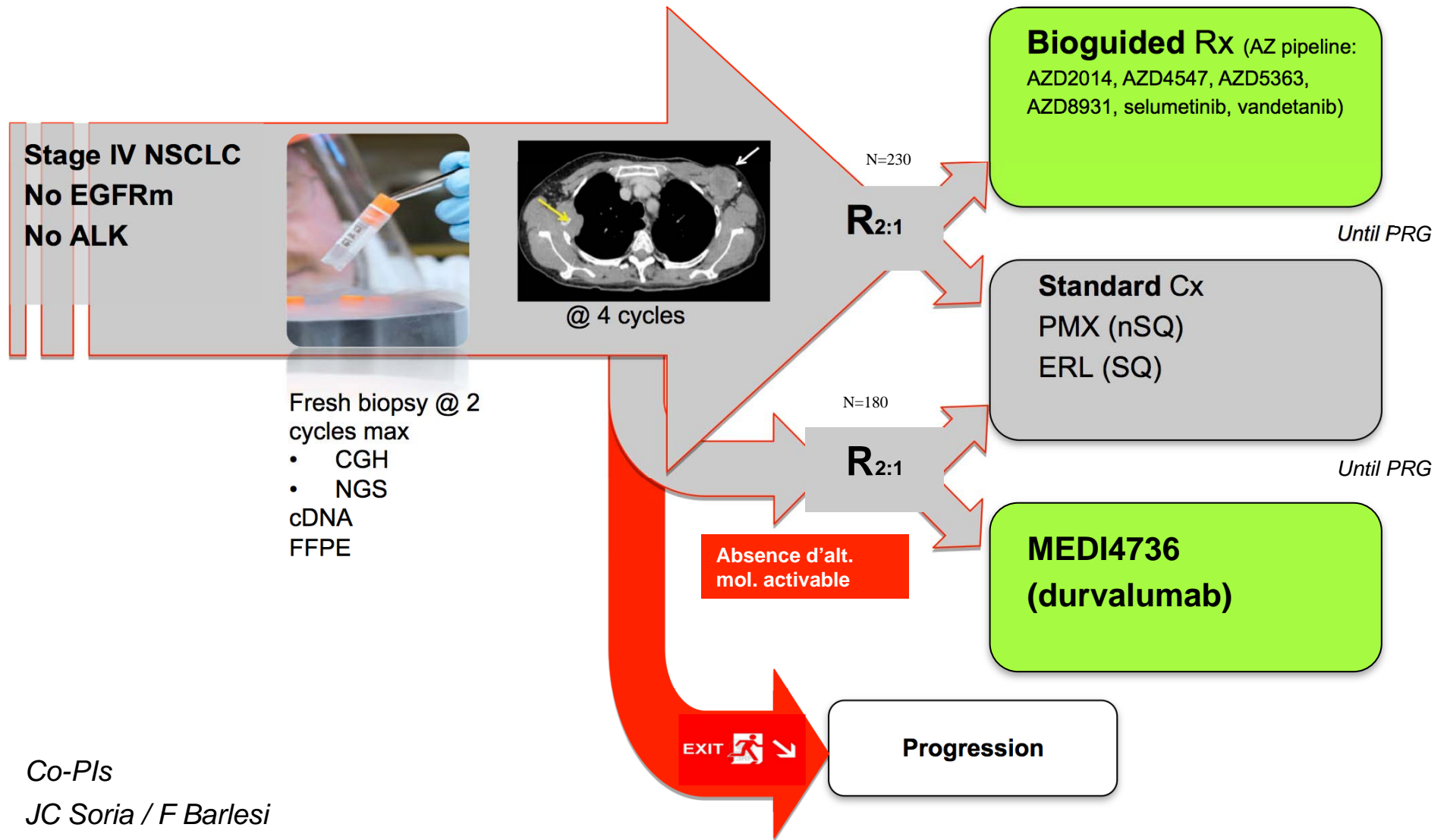
Gomez et al, ASCO 2016 (abst 9004)

Enjeux et Perspectives



Gomez et al, ASCO 2016 (abst 9004)

IFCT-UNICANCER SAFIR 02 lung trial



Co-PIs

JC Soria / F Barlesi

Conclusions

- **2017-2018:**
 - « **NGS** » **accessible facilement**
 - **TKI 3G 1L**
 - **ROS1, BRAF, MET activables**
 - **Complexification décision (RCP bio mol)**
 - **Peu d'autres cibles (essais précoces)**
 - **ICI (mono PD-L1 high+ et combo?) 1L**

Conclusions

- **2019-2020:**
 - Profils prédictifs (Cx, ICI, etc) ?
 - Nouveaux réarrangements
 - Nouveaux TKI 3G ou 4G (paninhib)
 - Nouveaux inhibiteurs (CDK, JAK) ?
 - IO: ICI mono vs combo (ICI, Cx, AA)
 - Impact contraintes économiques ??

Conclusions



Ministère des Affaires sociales et de la Santé

Rechercher

Affaires sociales | Prévention en santé | Santé et environnement | Soins et maladies | Système de santé et médico-social

Accueil > Études et statistiques > Publications > Recueils, ouvrages et rapports > Recueils annuels > Panoramas de la DREES > Les dépenses de santé en 2015 - Résultats des comptes de la santé

Les dépenses de santé en 2015 - Résultats des comptes de la santé

publié le : 05.09.16

Comptes et analyses économiques | 2016

A+ A- 



En 2015, la consommation de soins et de biens médicaux (CSBM) est de 194,6 milliards d'euros. Elle progresse de 1,8 %, soit légèrement moins rapidement que le PIB en valeur (+1,9 %), contrairement à la période 2012-2014 où sa croissance était supérieure à celle du PIB. La France consacre, au total, 11 % de son PIB à la santé, tout comme la Suède, l'Allemagne et les Pays-Bas.

La Sécurité sociale finance plus des trois quarts de la CSBM et les organismes complémentaires 13,3 %. La part restant à la charge des ménages recule pour la quatrième année consécutive et atteint 8,4 % en 2015. Les ménages consacrent ainsi un peu moins de 250 euros par habitant à leur consommation de santé, soit moins que la plupart de leurs voisins européens.

Les dépenses de santé en 2015 – édition 2016 présentent également un éclairage sur les dépenses de prévention sanitaire, qui représenteraient plus de 4,8 % de la CSBM, en 2014.

Sommaire

Avant-propos

Ouvrage

Données

Contributeurs

Dans la même rubrique

La complémentaire santé : acteurs, bénéficiaires, garanties - édition 2016

La protection sociale en France et en Europe en 2014 - Résultats des comptes de la protection sociale - Édition 2016

Les établissements de santé - Édition 2016

Les retraités et les retraites - édition 2016

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