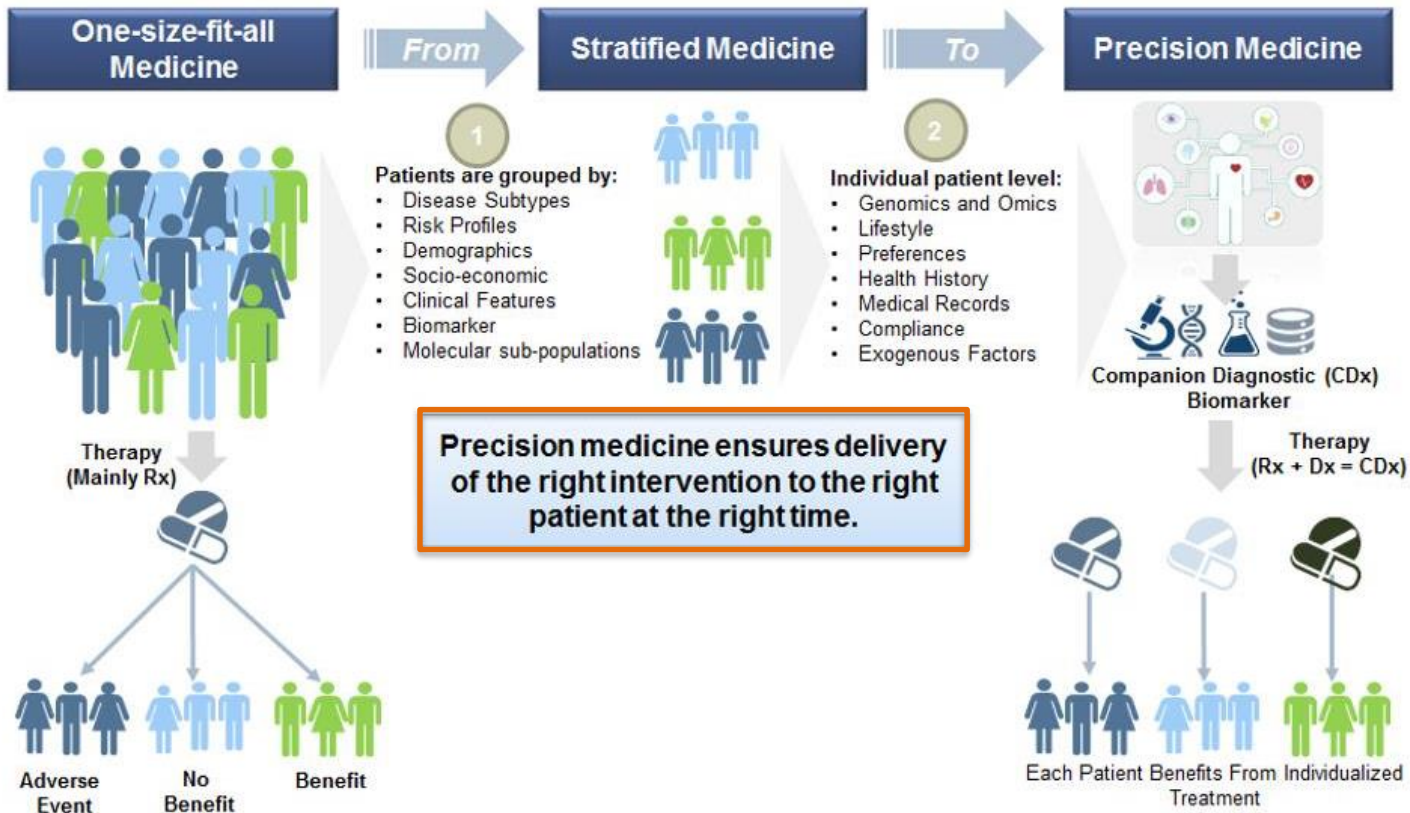


# Liquid biopsies as important tool for the implementation of precision oncology. The example of OncoBEAM EGFR mutation assessment in lung cancer.



Silvia Calabuig Fariñas  
FIHGU  
UV  
CIBERONC

## Precision Medicine

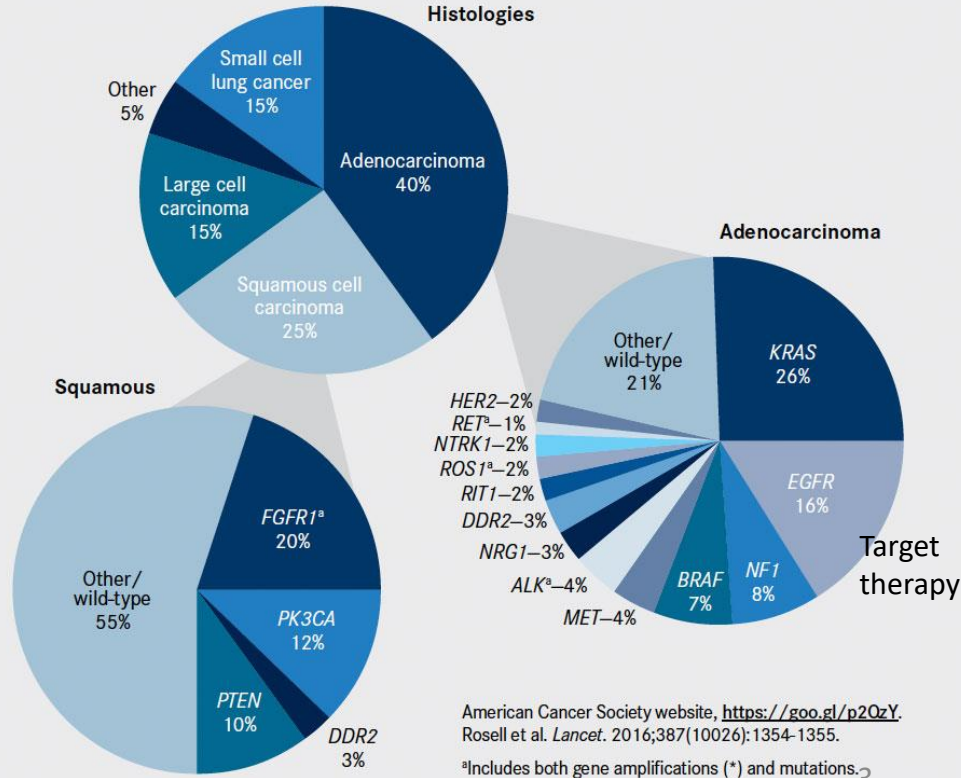


## Non-Small Cell Lung Cancer

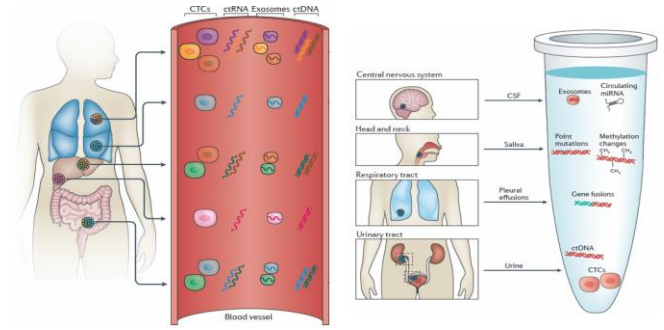
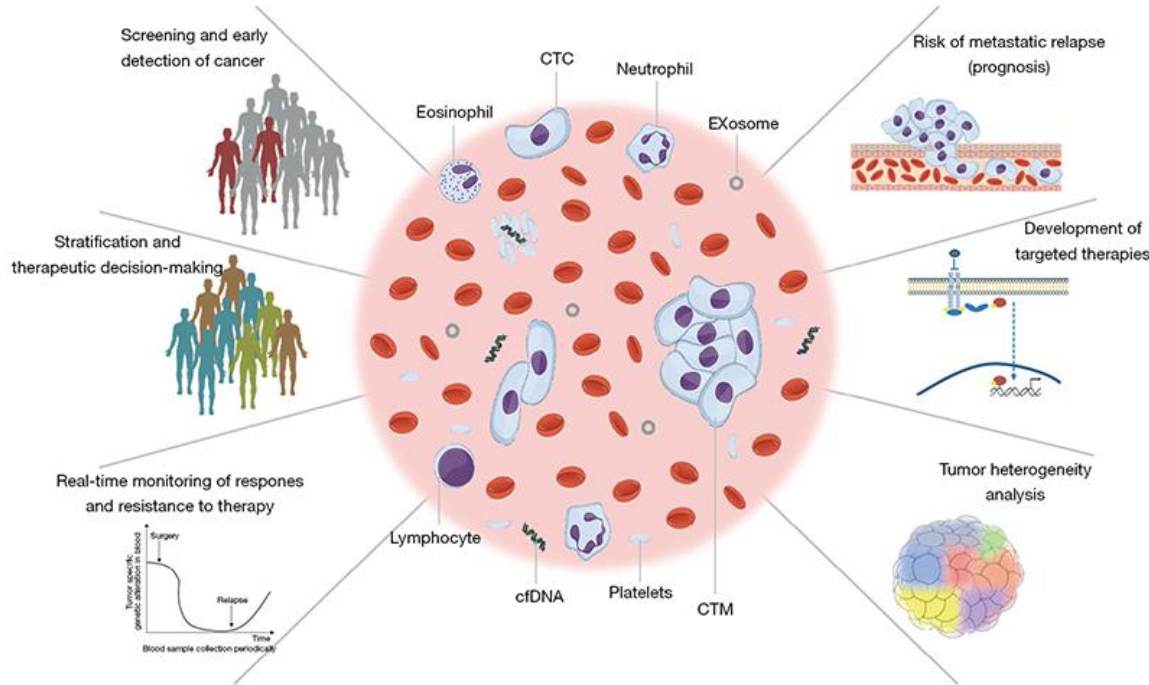


- **Leading cause of cancer deaths worldwide (1.6M deaths each year);**
- **In 15-20% of advanced NSCLC cases,** genetic tests are not performed due to: **unavailable** or **insufficient tissue**, poor PS, comorbidities.

### Lung Cancers and Their Molecular Drivers

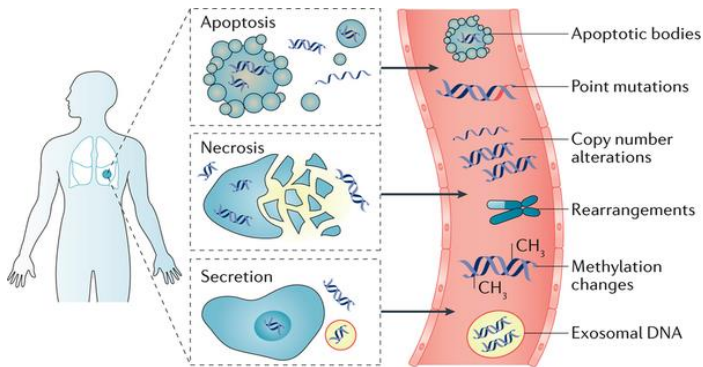


## Liquid biopsies as tool for the implementation of precision oncology.

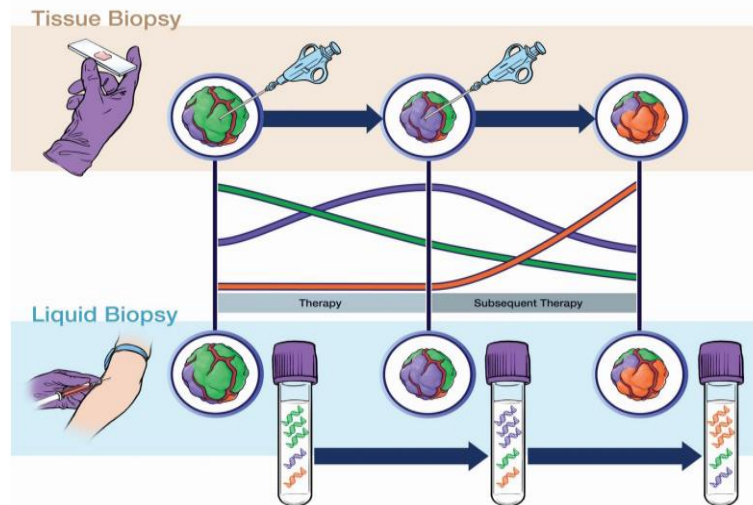
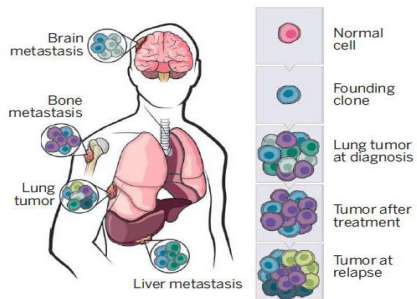


⇒ To analyze the clinical utility of plasma using cell-free circulating tumor DNA (ctDNA) for advanced-stage lung ADC patients, as a complement or alternative to tissue-based molecular profiling.

## Liquid biopsies as tool for the implementation of precision oncology

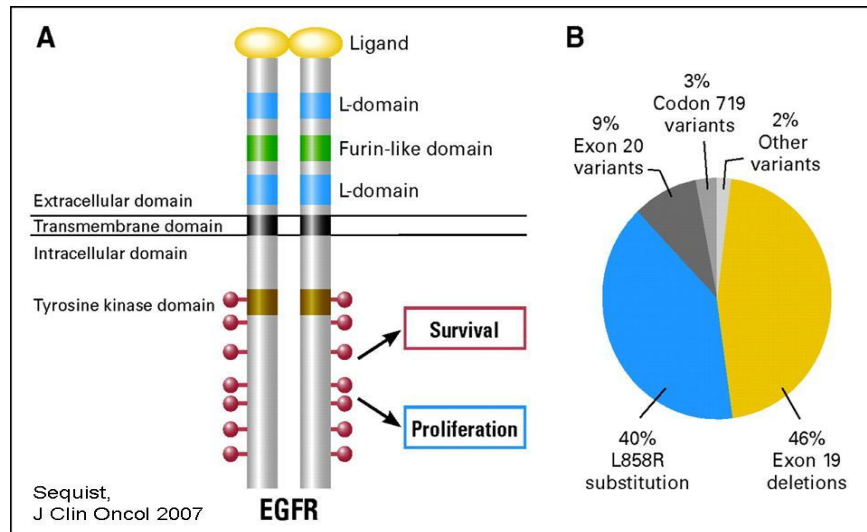


Nature Reviews | Cancer



## EGFR Sensitizing Mutations:

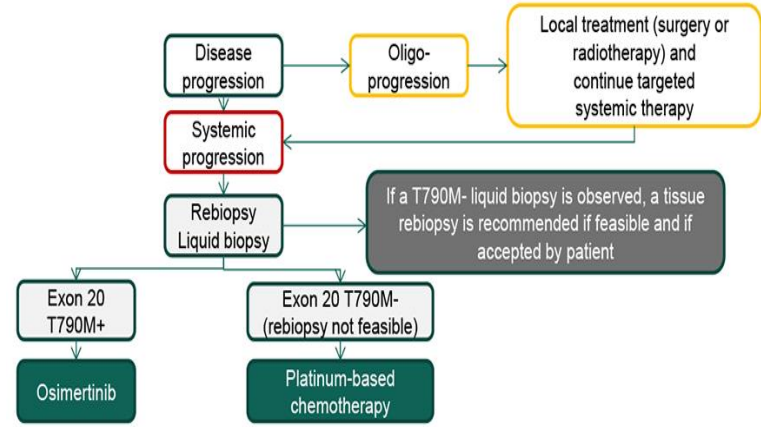
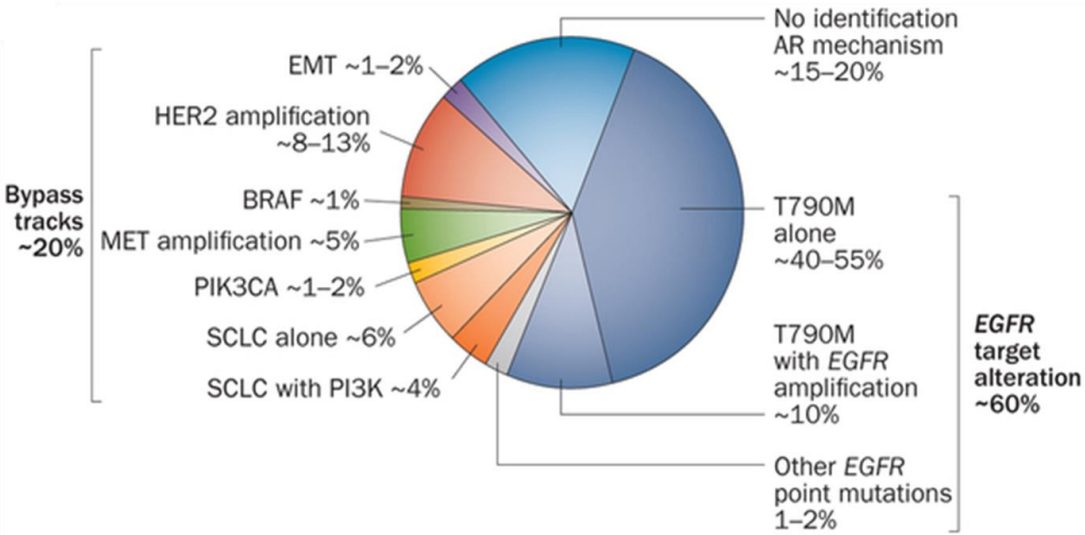
- Found in 10% to 30% of NSCLC pts[1]
- More common in never-smokers, adenocarcinomas, females, Asians[1,2]
- Predominantly located in *EGFR* exons 18-21[2]
  - ~ 85% of *EGFR* mutations are either deletions in exon 19 or a single-point mutation in exon 21 (L858R)[3]
- Specific *EGFR* mutation identified is important
  - There are sensitive mutations, primary resistance mutations (often exon 20), and acquired resistance mutations (T790M)[3]



<p><b>Erlotinib</b> <b>Gefitinib</b></p>	<p>1<sup>st</sup>-generation TKI EGFR inhibition</p>	<p>Activity range</p> <ul style="list-style-type: none"> <li>• Reversible binding to wild-type and mutant EGFR</li> <li>• Inactive on T790M mutant</li> </ul>
<p><b>Afatinib</b> <b>Dacomitinib</b></p>	<p>2<sup>nd</sup>-generation TKI ErbB family blockade</p>	<p>Activity range</p> <ul style="list-style-type: none"> <li>• Irreversible covalent binding to EGFR, ErbB2 and ErbB4 to inhibit all ErbB family signalling</li> <li>• Broader activity to overcome EGFR TKI-resistant mutations</li> </ul>

EGFR, epidermal growth factor receptor; TKI, tyrosine kinase inhibitor. Liao BC, et al. Curr Opin Oncol 2015;27:94–101.

# EGFR T790M the most common resistance mechanism to 1st and 2nd EGFR TKIs

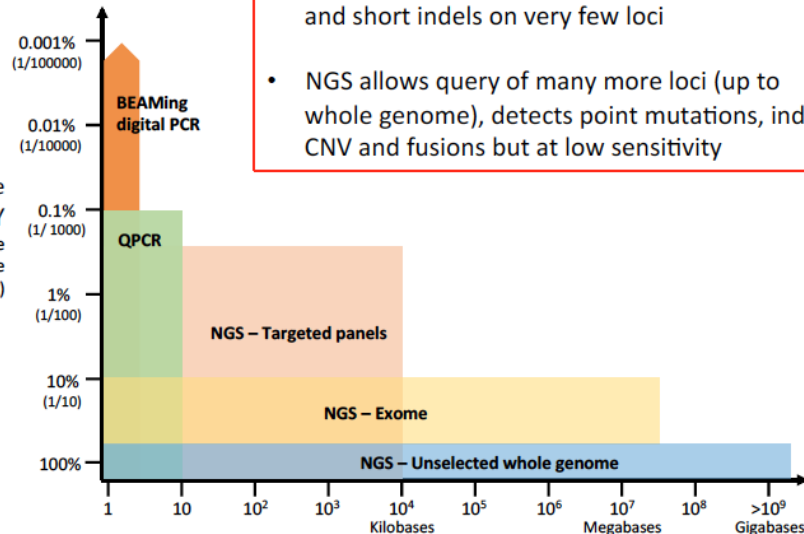


## Different methodologies

Technique	Sensitivity	Optimal Application
Sanger sequencing	> 10%	Tumor tissue
Pyrosequencing	10%	Tumor tissue
Next-generation sequencing	2%	Tumor tissue
Quantative PCR	1%	Tumor tissue
ARMS	0.10%	Tumor tissue
BEAMing, PAP, Digital PCR, TAM-Seq	0.01% or lower	ctDNA, rare variants in tumor tissue



Relative  
SENSITIVITY  
(Ability to see  
increasingly rare  
mutant alleles)



- Allele specific- and emulsion-PCR methods are highly sensitive but detect only point mutations and short indels on very few loci
- NGS allows query of many more loci (up to whole genome), detects point mutations, indels, CNV and fusions but at low sensitivity

	Type of mutation	EGFR exon	Mutations
OncoBEAM EGFR Kit v2 (RUO*) 36 EGFR mutations	Sensitising	18	G719A, G719S, G719C
		19	ΔK745, ΔE746, ΔL747
		21	L858R, L861Q
	Resistance	20	T790M, C797S



BREADTH  
(N° bases assayed per sample)



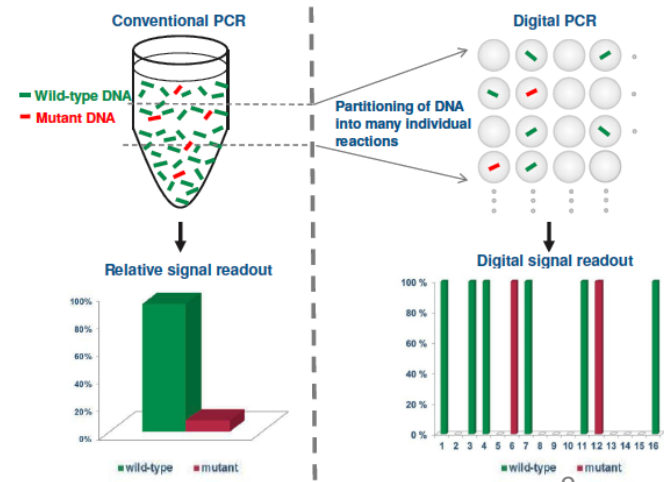
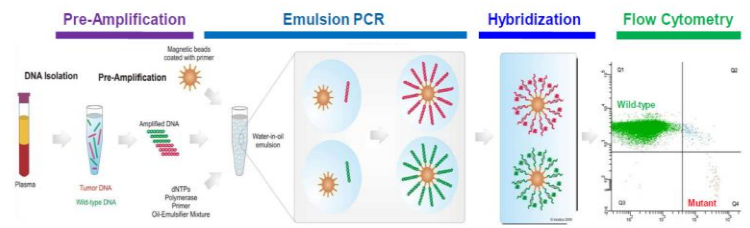
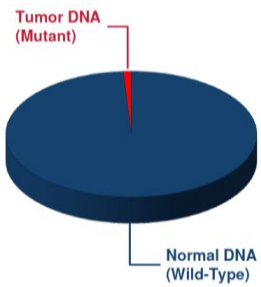
## Digital PCR

*Beads*  
*Emulsion*  
*Amplification*  
*Magnetics*

- High Sensitivity
- Suitable for detection of specific point mutations, short indels
- No bioinformatic analysis
- One or few gene mutations could be analyzed
- Only enables monitoring of known mutations

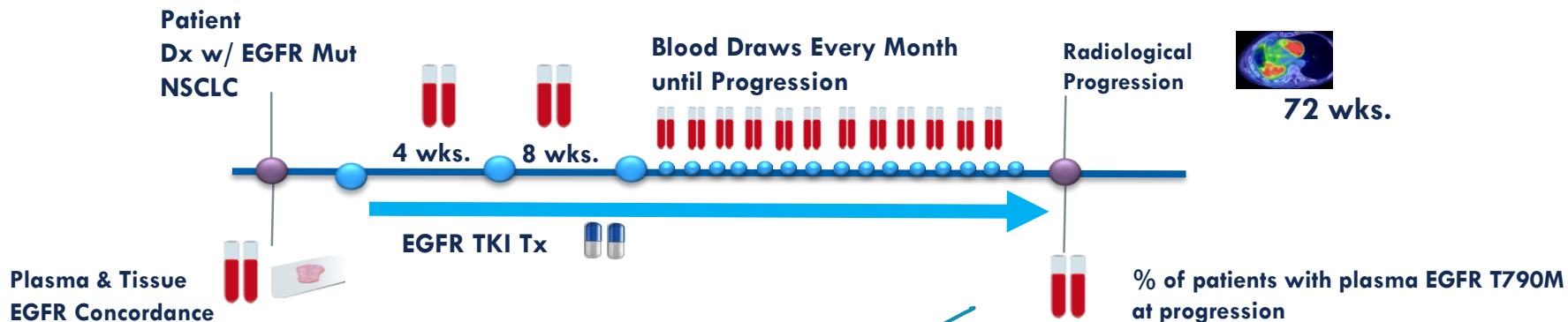
- We must be SCRUPULOUS in the analytical phase to avoid erroneous results
- Clinically irrelevant molecular changes due to the high sensitivity

Highly sensitive quantitative digital PCR technology that employs bead-based amplification in water-in-oil emulsions and allele-specific hybridization followed by flow cytometry for detecting small amounts of mutated DNA released by tumors into the blood circulation



## LungBEAM Trial – Study Design & Methods

- Therapy-naïve metastatic NSCLC patients with EGFR-positive sensitizing mutations (del19, L858R) tissue at baseline
- Standard EGFR TKI treatment and follow-up based on local guidelines
- CT every 8 weeks until 6 months and then every 12 weeks. No central review.
- Blood samples throughout standard first-line EGFR TKI treatment until PD or 72 wks.
- BEAMing test performed in a CLIA lab
- Clinicians blinded to blood results.



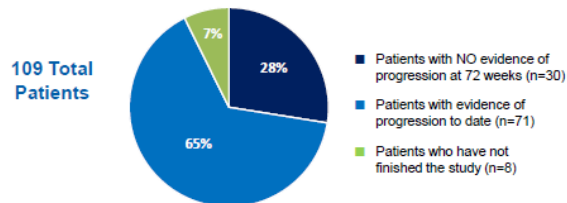
**Goals: Determine concordance at baseline and 'window' of T790M emergence with respect to time of disease progression.**

## LungBEAM Trial – Study

### Patient Characteristics

Patient cases recruited/analyzed for concordance:	109 (Nov 2015-May 2017)
Median Age:	65.5 (35-87)
Gender:	
Female	78 (71.5%)
Male	31 (28.5%)
Smoking status:	
Never Smoker	67 (61.5%)
Former Smoker	33 (30.2%)
Active Smoker	9 (8.3%)
M status:	
M1a	35 (32.1%)
M1b	74 (67.9%)
Metastases:	
brain only	10 (9.2%)
bone only	16 (14.7%)
Stage at diagnosis and tissue acquisition:	
M0	14 (12.8%)
M1	95 (87.2%)
Tissue EGFR mutation status:	
del19+	69 (63.3%)
L858R+	40 (36.7%)

### Positive Percent Agreement (PPA) at baseline of EGFR del19 and L858R mutations

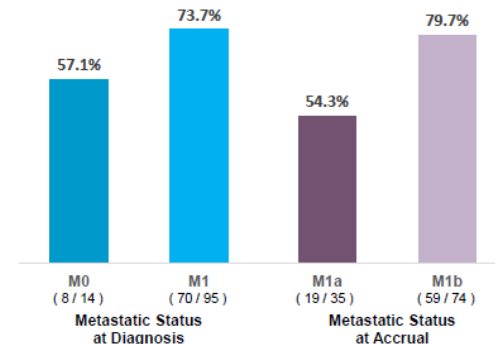


PPA at baseline of EGFR del19 and L858R mutational results between plasma and tissue was **71.6% (78/109)**

The **average number of days** between tissue biopsy and blood collection for:

Concordant cases	116 days
Discordant cases	306 days

Significant differences on PPA was observed according to the metastatic status of patients



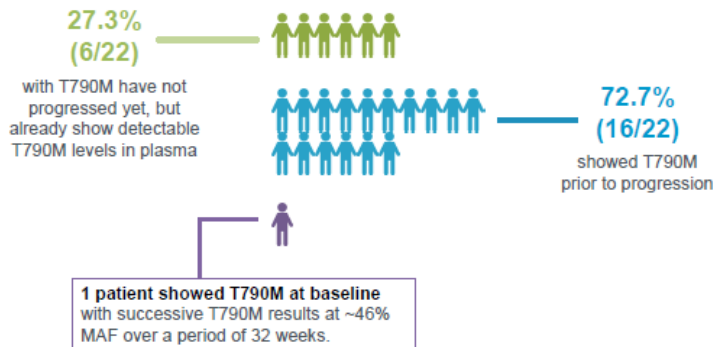
### Progression-free survival indicators

Clearance of EGFR mutations in plasma 8 weeks after initiation of EGFR TKI could be a favorable indicator for PFS:

	Number of Patients	MAF at baseline	Mean PFS
Detectable mutations at 8 weeks	13	13.5%	25.8 weeks
Clearance of mutations at 8 weeks	33	3.8%	40.3 weeks

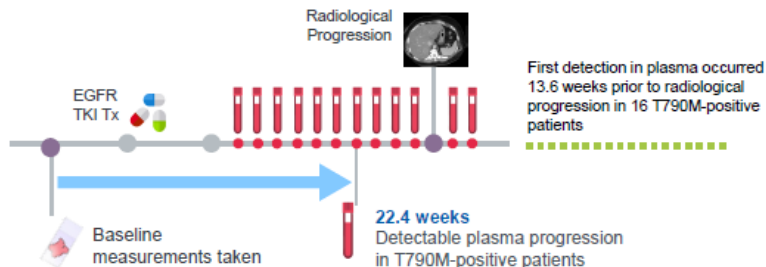
## Plasma T790M monitoring during first line EGFR TKI therapy

Of the 23\* T790M-positive patients:



\*21 of the 23 patients also had an accompanying del19 or L858R mutation

## Timing of T790M Detection in Plasma



A total of 23 patients showed T790M-positive (only 16 had radiological progression):

Average time to T790M-positive plasma:	22.4 weeks
Average T790M mutant allelic fraction (MAF):	0.11 %

So far, 71 patients have experienced Radiological Progression. The average time was:

non-T790M patients (n=55):	26.7 weeks
T790M-positive patients (n=16):	35.8 weeks

## CONCLUSIONS

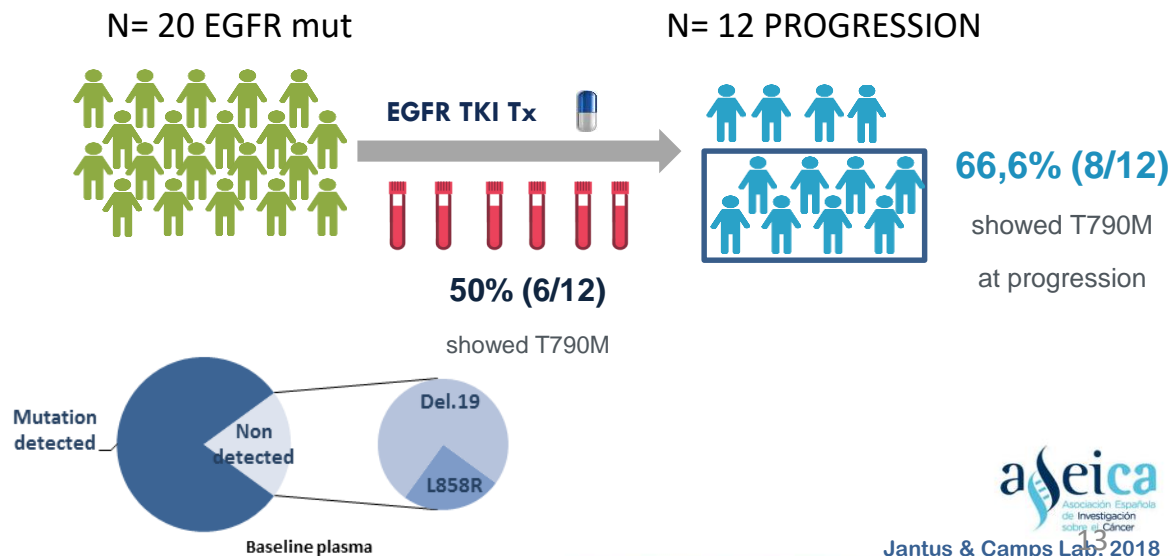
- Overall, these initial results show high concordance of plasma and tissue del19 and L858R EGFR mutation status at baseline (~72%).
- Early detection of T790M ctDNA using a highly sensitive method may assist in anticipating T790M resistance to first-line anti-EGFR therapy.
- Clearance of EGFR mutations from plasma 8 weeks after initiation of anti-EGFR therapy could be indicator of a more favorable PFS.

Characteristics	N=20
<b>Age at diagnosis (median, range)</b>	70 [47-85]
<b>Gender</b>	
Male	5
Female	15
<b>Smoking status</b>	
Never smoker	12
Former smoker	4
Active smoker	4
<b>EGFR mutation status</b>	
Del.19	8
L858R	12
<b>Progression</b>	
No	8
Yes	12
<b>Dead</b>	
No	15
Yes	5

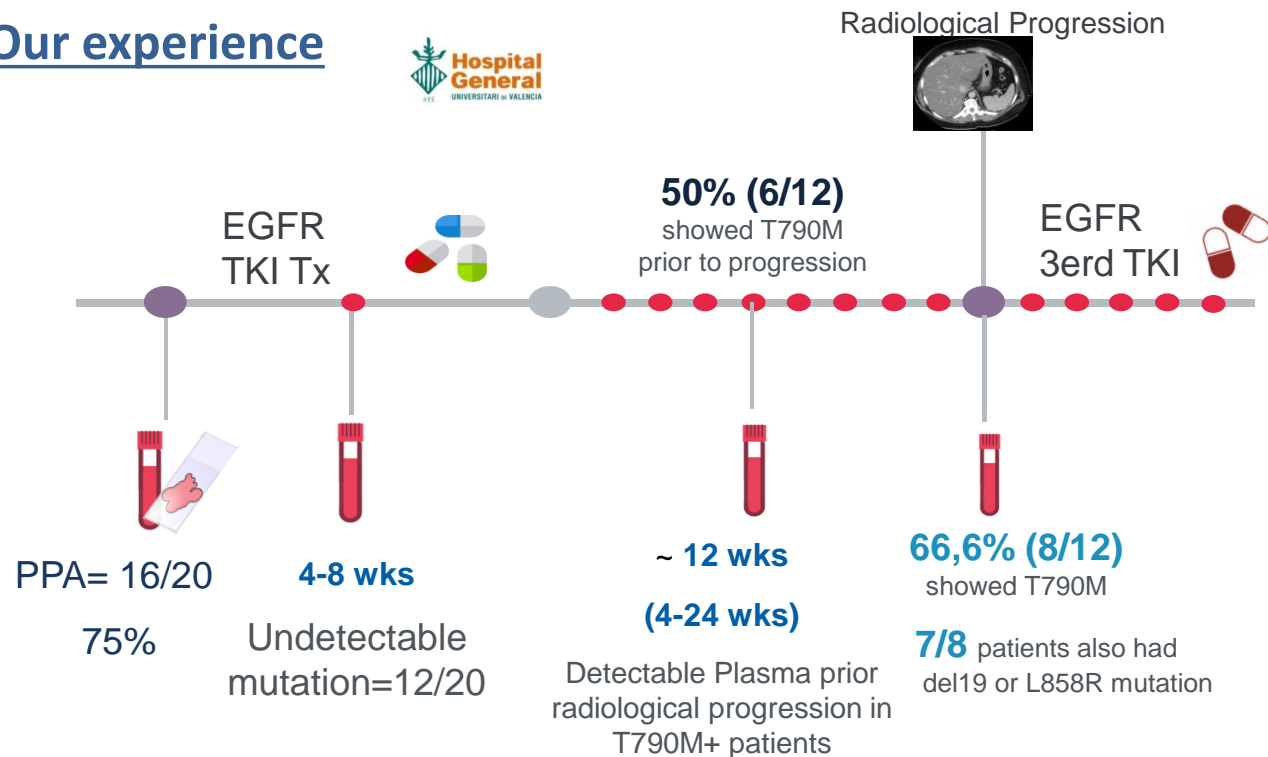
## Our experience



- Patients recruited between 2014-2018.
- The Positive Percent Agreement (PPA) at baseline of EGFR del19 and L858R mutational results between plasma and tissue was **16/20 = 75%**



## Our experience



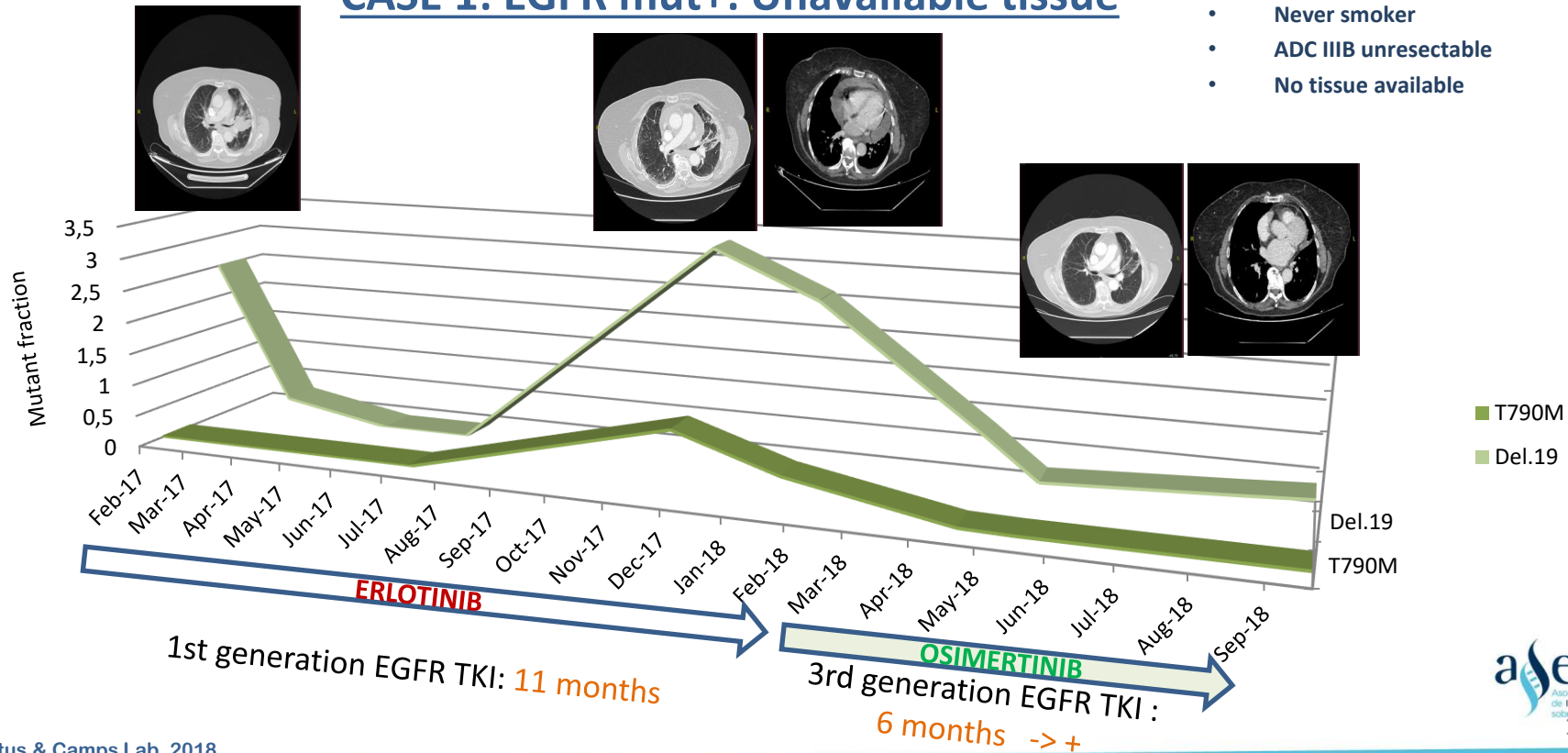
- High **concordance** of plasma and tissue at baseline.

- **Clearance** of mutations from plasma after initiation of targeted therapy could be **indicator of a more favorable PFS.**

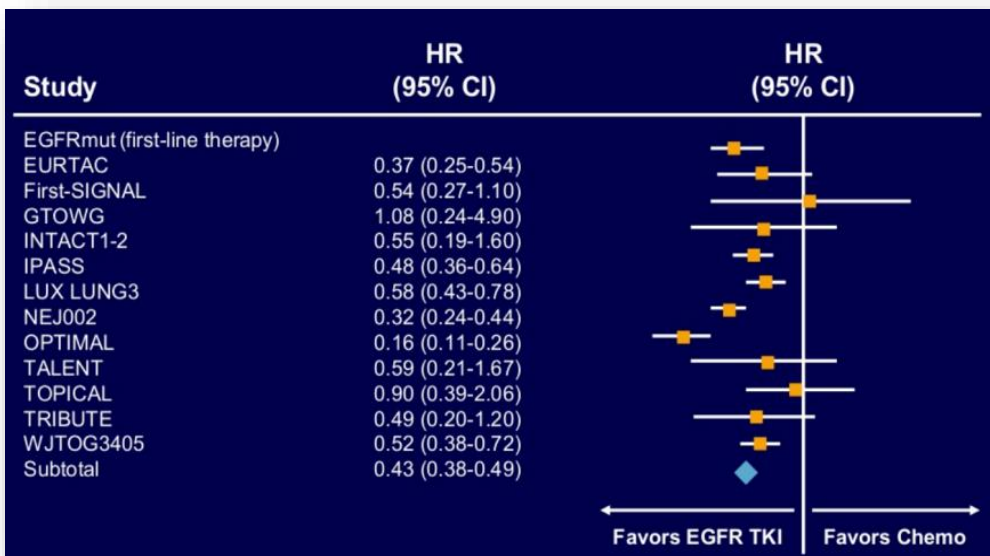
- **Early detection** of resistance mutations -> highly sensitive method may assist in anticipating resistance to anti-targeted therapy.

## CASE 1: EGFR mut+: Unavailable tissue

- 85 years old woman
- Never smoker
- ADC IIIB unresectable
- No tissue available



## First-line Treatment With EGFR TKIs vs. Chemotherapy in EGFR-Mutated NSCLC



"If repeat biopsy is not feasible, plasma biopsy should be considered"

"Testing should be conducted as part of broad molecular profiling"

NCCN 2017 NSCLC Practice Guidelines

"Key new recommendations include the inclusion of additional genes (*ERBB2*, *MET*, *BRAF*, *KRAS*, and *RET*)...and the use of cell-free DNA to "rule in" targetable mutations when tissue is limited or hard to obtain.

AMP/CAP/IASLC 2018 Molecular Testing Guidelines for Lung Cancer

"Even for patients who are able to undergo a traditional tissue biopsy, a liquid biopsy may be safer, quicker, and more convenient—and perhaps even more informative."

2017 ASCO Clinical Cancer Advances

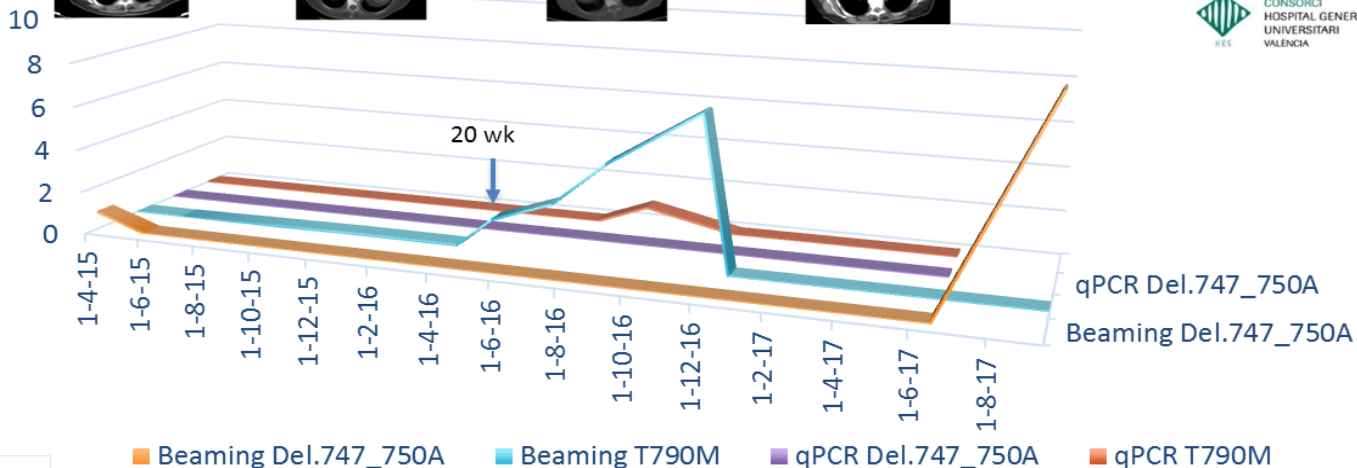
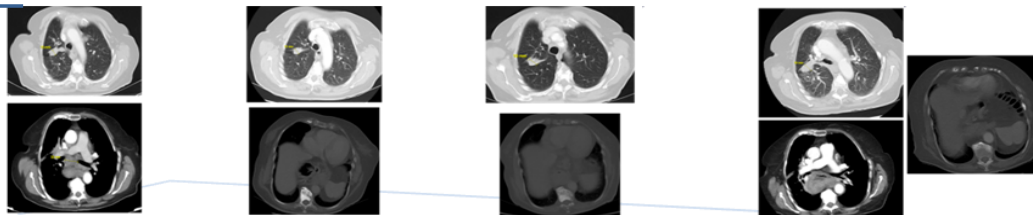


## CASE 2: T790M mut.

Mutant Fraction



%mutant allele



Methods	Sample, n	T790M, %	PPA, %	NPA, %	OPA, %	Therapy Response Rate, %	
						Tissue T790M	Plasma T790M
cobas EGFR Mutation Test	72	41	73	67	70	61	59
cobas EGFR Mutation Test	110	21	64	98	86	52	44
BEAMing Digital PCR	77	66	73	NA	NA	-	-
BEAMing Digital PCR	216	60	70	69	70	62	63
Droplet Digital PCR	60	58	77	63	72	NA	NA
NGS	63	41	93	94	78	NA	NA



NPA: negative percentage agreement; OPA: overall percentage agreement; PPA: positive percentage agreement.

Passaro A et al. *Pharmacol Res.* 2017;117:406-415.

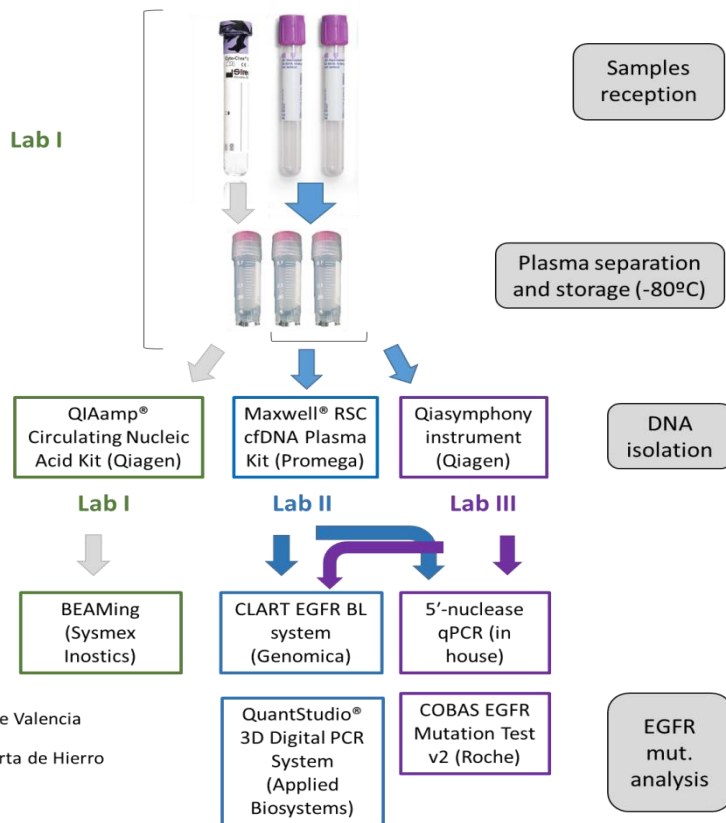
Using the Cobas tissue test as a reference, sensitivity for the detection of T790M was increased for ddPCR and NGS compared with AS-PCR. In AURA I, ctDNA (BEAMing) **“rescue” ~30% of tissue T790M-negative tumors**

**Is test approved by FDA the best?**

## RING Study



Grupo Español de Cáncer de Pulmón  
Spanish Lung Cancer Group



- Lab Oncología Molecular, Fundación Htal Gral Univ de Valencia
- Lab de Biopsia Líquida, Fundación del Htal. Univ. Puerta de Hierro
- Pangaea Oncology

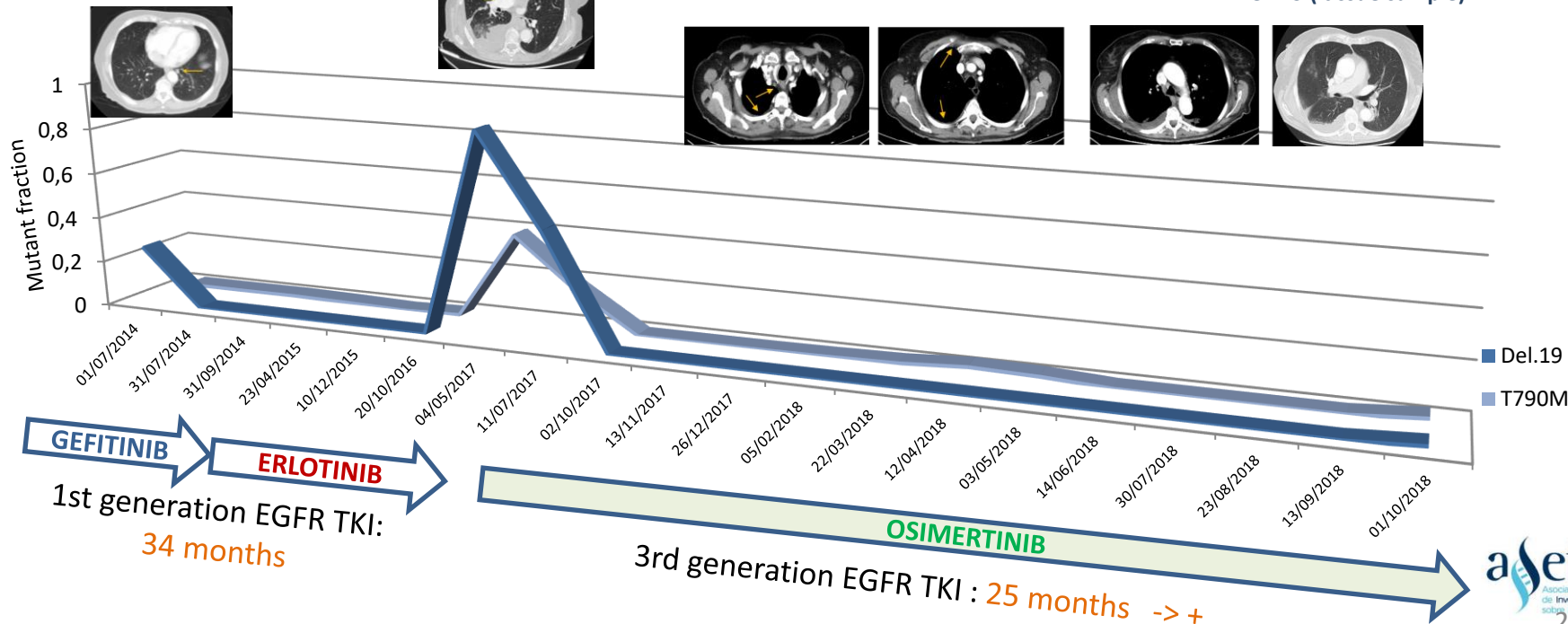
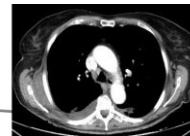
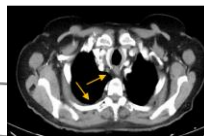
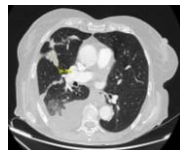
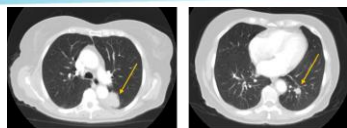
“T790M mutation testing in blood by different methodologies”

A study to compare 3 purification methods and 8 techniques to determine p.T790M (including NGS)

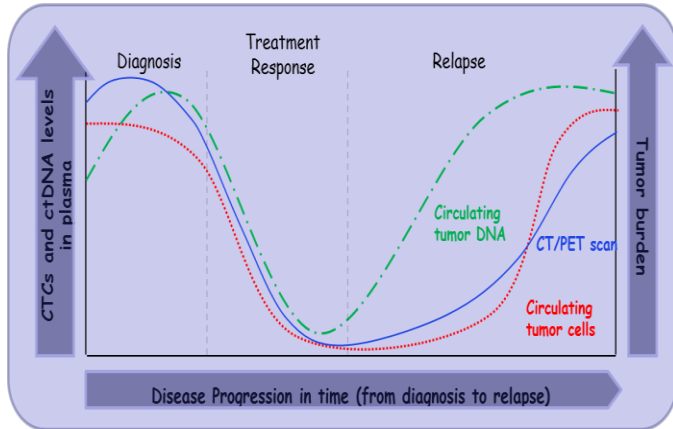
Protocol code: GECP 17/03; ESR-17-12988  
PI: Mariano Provencio

## CASE 3: Monitoring of EGFR mut+

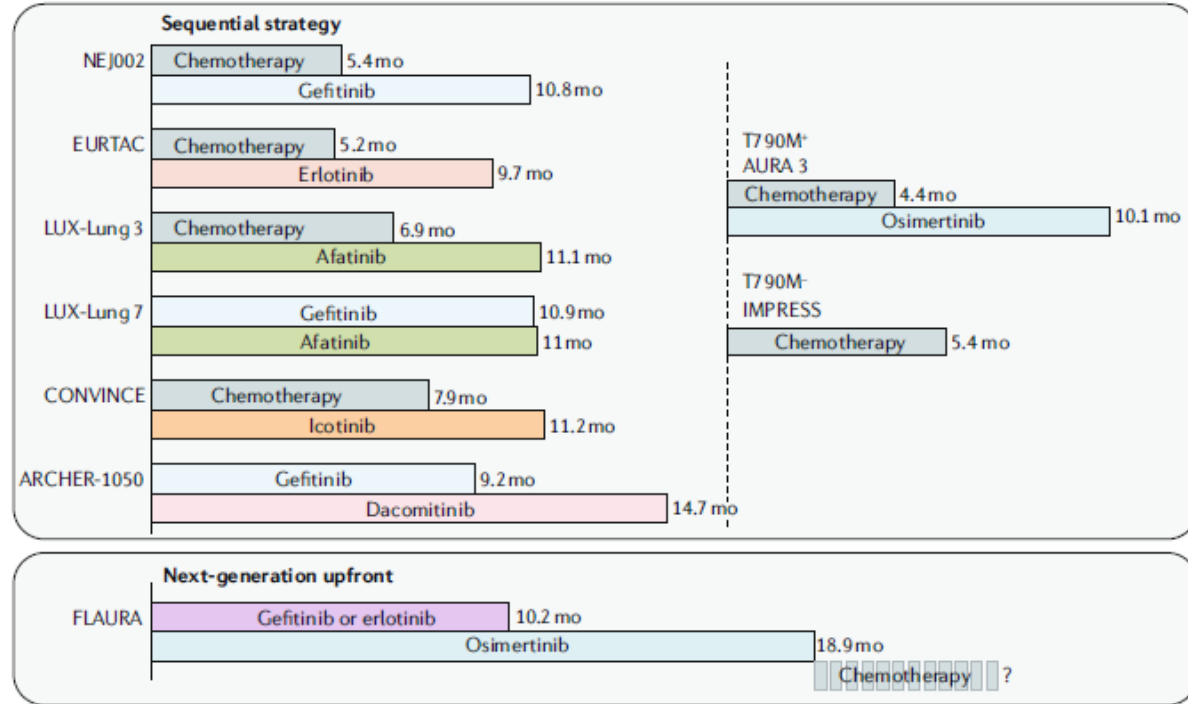
- 76 years old woman
- Never smoker
- Poor differentiated lung carcinoma E-IV
- Del. 19 ( tissue sample)



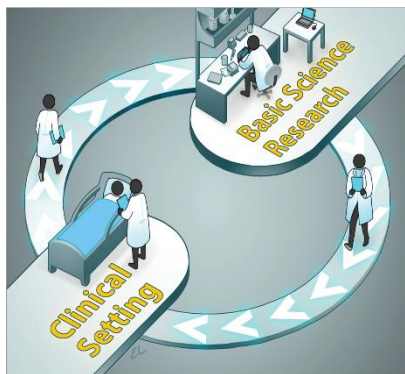
## Optimal sequencing treatment



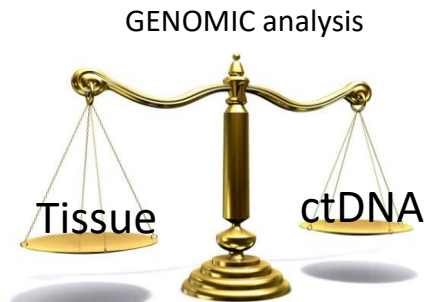
### a EGFR



## Take home message



From bench to bedside



Complementary biopsies

Next Generation Sequencing



Digital PCR



RTqPCR



BEAMing

Different methodologies



Multidisciplinary team



Quality controls



## Translational Res.



### MOLECULAR ONCOLOGY LAB.

Carlos Camps, Eloisa Jantus, Silvia Calabuig, Rafael Sirera, Sandra Gallach, Eva Escorihuela, Marais Mosqueda, Alejandro Herreros, Elena Duréndez, Héctor Amado, Clara (Ning Dong), Bruno (Feiyu Zhang), Susana Torres, Andrea Moreno.

### CLINICAL INVEST.

Vicente Castellano, Paula Matoses, Ana Saval, Lola Serra

## Clinical Res.



### MEDICAL ONCOLOGY

Carlos Camps, Alfonso Berrocal, Ana Blasco, M<sup>a</sup> José Safont, Cristina Caballero, M<sup>a</sup> José Godes, Vega Irazo, Francisco Aparisi, Miriam Lobo, Mireia Gil, Alberto Cunquero.

THORACIC SURGERY: Ricardo Guijarro, Eva del Olmo

PATHOLOGY: Miguel Martorell, Atilio Navarro, Lara Navarro



**21:00 YOUNG ONCOLOGISTS NETWORKING EVENING**  
(Only for assistants under 45 years old)

Venue: ADEIT | Fundación Universidad-Empresa  
de la Universidad de Valencia  
Plaza Virgen de la Paz, 3, 46001 Valencia

**Liquid biopsies as important tool  
for the implementation of precision oncology.  
The example of OncoBEAM EGFR mutation  
assessment in lung cancer.**

Silvia Calabuig Fariñas.

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