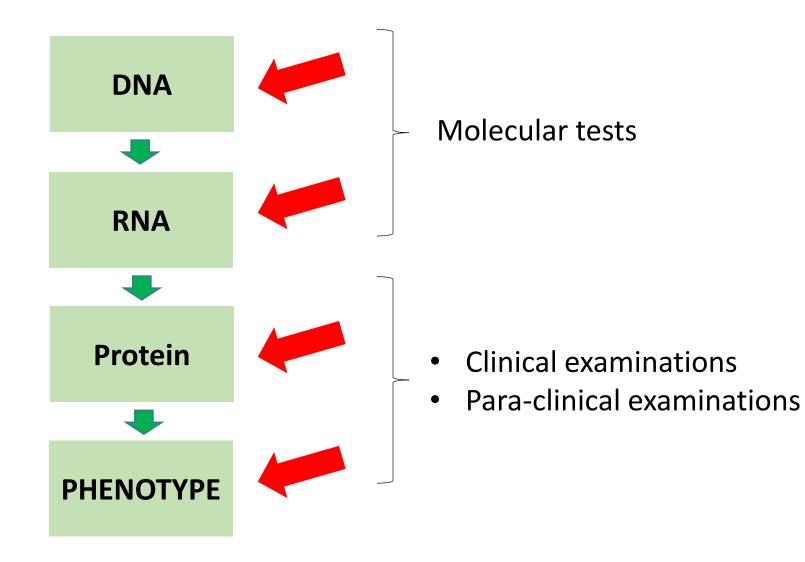
ỨNG DỤNG CHẨN ĐOÁN PHÂN TỬ TRONG NHÓM BỆNH UNG THƯ

Ts Bs Nguyễn Hữu Ngọc Tuấn

APPROACH



At the body level

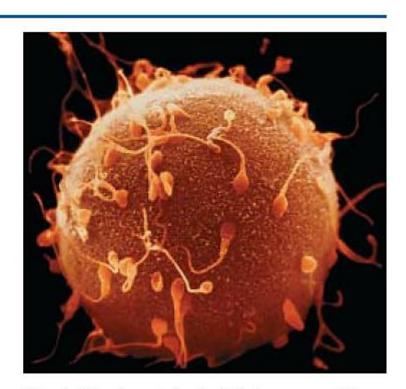




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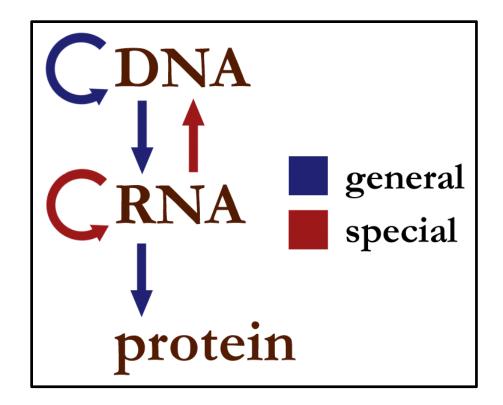
How does life begin?



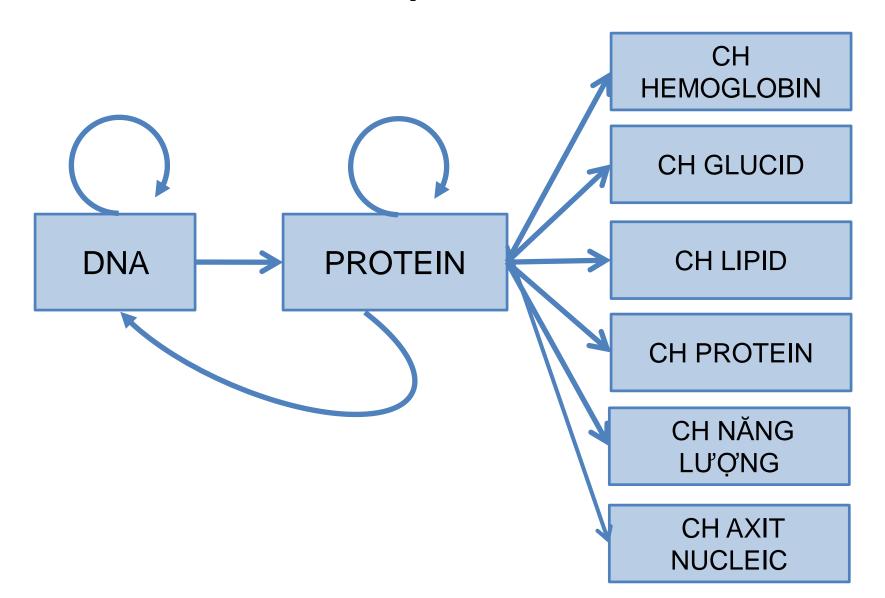


A single –200 micrometer (μ m) cell, the human egg, with sperm, which are also single cells. From the union of an egg and sperm will arise the 10 trillion cells of a human body.

The central dogma of biology



Relationship of molecules

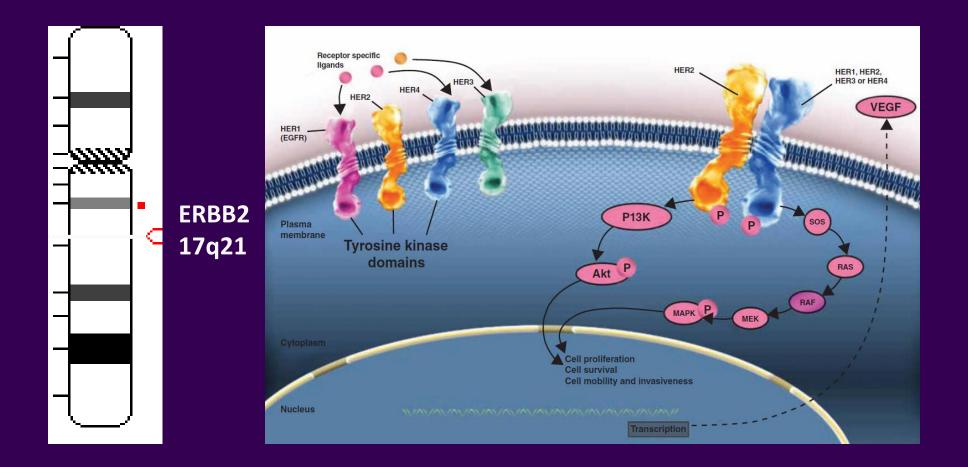


Cancer = Cellular Pathology

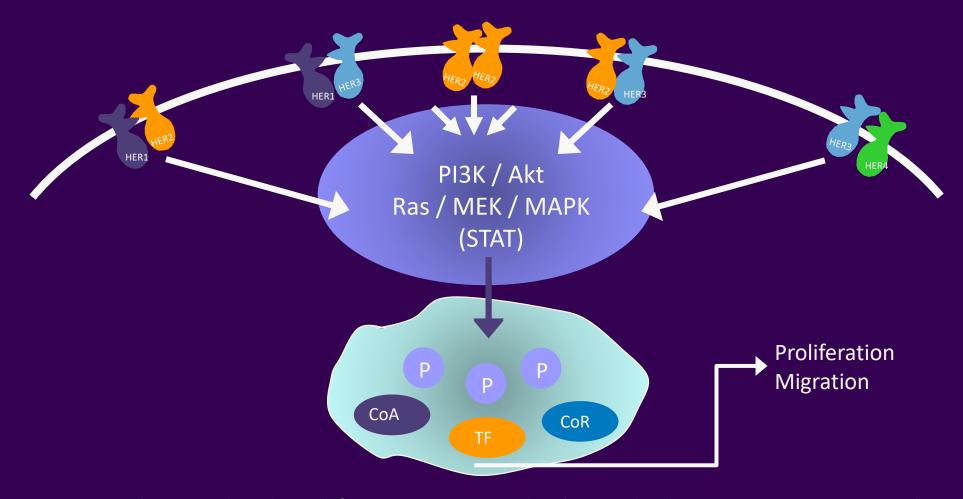
Cancer = Genomic Pathology

BREAST CANCER

HER2 is required for normal cell development

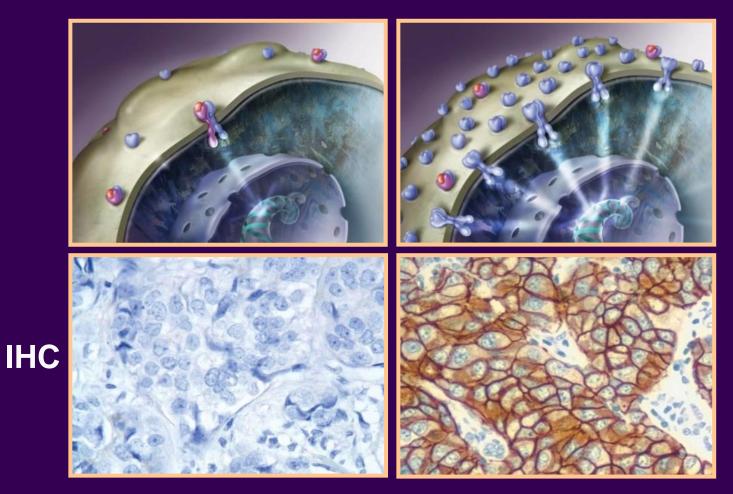


Overexpression of HER2 increases cellular proliferation and migration



HER2, human epidermal growth factor receptor 2; PI3K, phosphoinositide 3-kinase; MAPK, mitogen-activated protein kinase; STAT, signal transducer and activator of transcription; CoA, co-enzyme A; TF, tissue factor; CoR, co-repressor

Overexpression of HER2 in BC



Normal expression

Overexpression

Overexpression of HER2 is associated with reduced survival in patients with BC

• Herceptin improves the prognosis of patients with HER2-positive metastatic breast cancer



Overall survival possibility

 In a retrospective analysis of database records, women with HER2-positive disease who received Herceptin had a 44% reduction in risk of death compared to women with HER2-negative disease (multivariate analysis adjusted for patient and tumour characteristics: HR 0.56; 95% CI 0.45, 0.69; p<0.0001)

Months from diagnosis

36

48

24

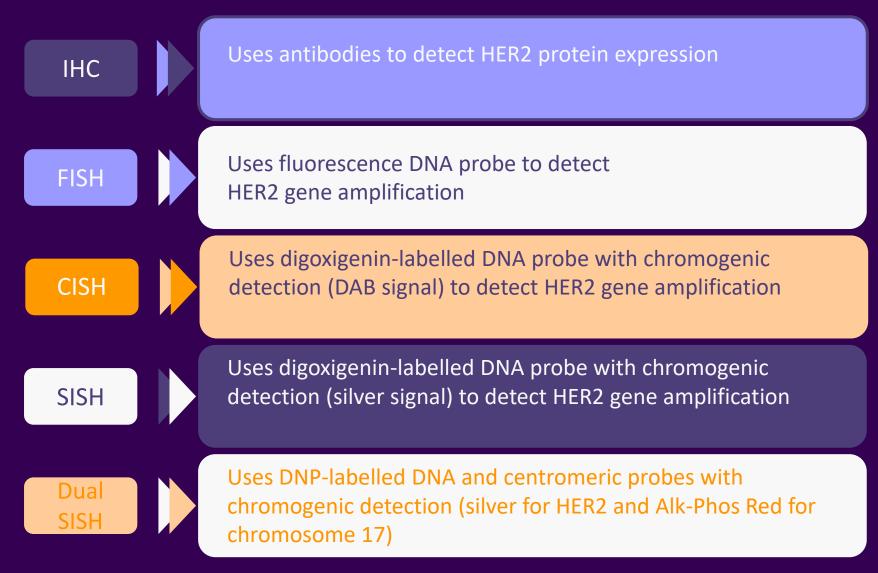
12

0.0 -

0

60

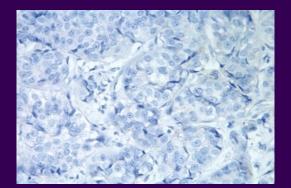
Summary of HER2-testing methods



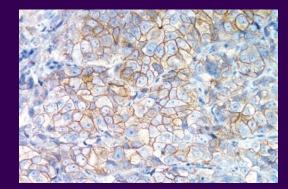
Advantages and disadvantages of HER2-testing methodologies (1)

Metho	d Advantages	Disadvantages	
IHC	Performed in majority of pathology laboratories	Susceptible to variations in testing protocol	
	Relatively easy, quick and cheap; can be automated	Score interpretation subjective and semi-quantitative	
	IHC-stained slides can be stored and re-assessed		
	Cell morphology can be seen in same section		
FISH	Less affected by pre-analytical factors and handling than IHC Score interpretation more quantitative than for IHC Identifies HER2-positive tumours (gene amplified) within IHC 2+ cases Automation available	Costly (more expensive than IHC) Signal decays over time Areas of invasive carcinoma may be difficult to identify Few pathologists and technologists are trained in the methodology and its interpretation	

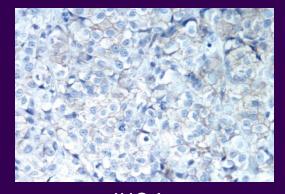
Interpretation of IHC results



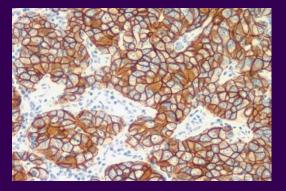
IHC 0 No staining or membrane staining in <10% of tumour cells



IHC 2+ Weak / moderate complete membrane staining in >10% of tumour cells



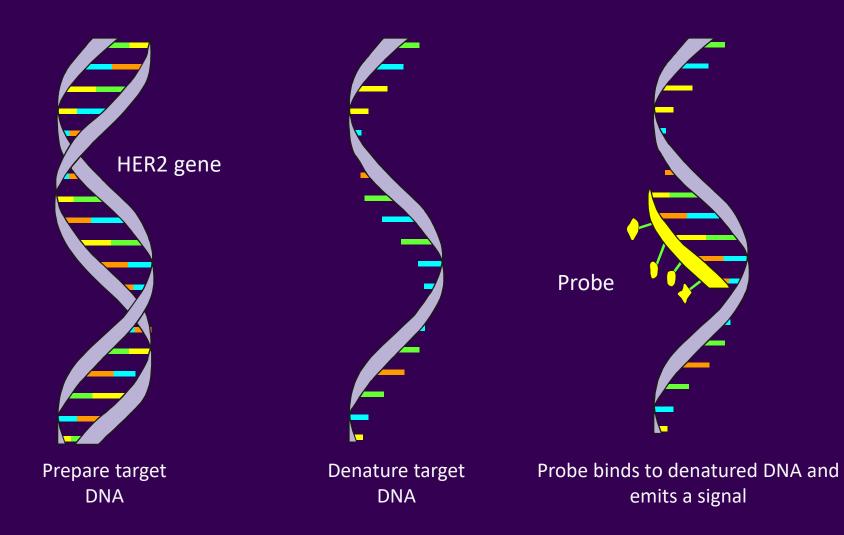
IHC 1+ Barely perceptible membrane staining in >10% of tumour cells; cells only stained in part of membrane



IHC 3+ Strong complete membrane staining in >10% of tumour cells

Images courtesy of Dako

Principles of ISH



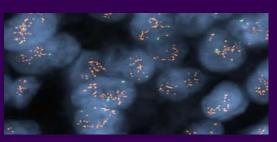
ISH: HER2 gene-amplification detection mechanisms

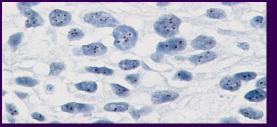
FISH positive^a

CISH positive^b

SISH positive^c

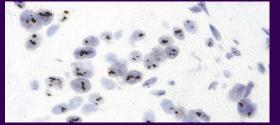
Dual SISH / Red Amplified HER2^c

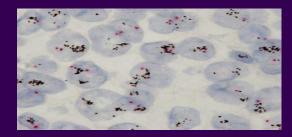




Fluorescence DNA probe

Digoxigenin-labelled DNA probe with chromogenic detection (DAB signal)





Dinitrophenol-labelled DNA probe with chromogenic detection (silver signal)

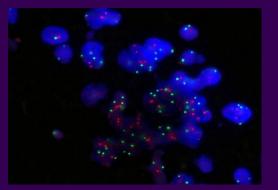
Dinitrophenol-labelled DNA and centromeric probes with chromogenic detection (silver and Alk-Phos Red)

> ^aImage courtesy of W Hanna; ^bimage from Invitrogen; ^cimage from Ventana

Polysomy

Gene amplification vs polysomy^a

XX	Normal 2 CEP17 2 HER2 genes
XX	Gene amplification 2 CEP17 >2 HER2 genes
XXXX	Polysomy >2 CEP17 >2 HER2 genes



The CEP17 probe identifies the centromere of chromosome 17

Polysomy means there are >2 CEP17 signals (green) and in consequence >2 HER2 gene signals (orange) detected per nucleus

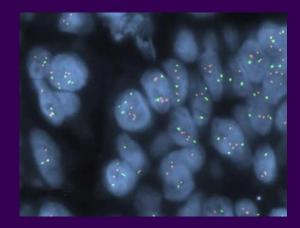
This can result in false-negative interpretation of ISH

FISH methodologies

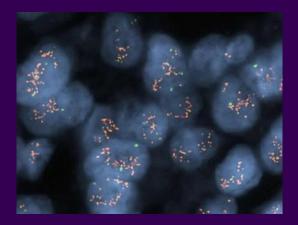
2 (1 for HER2 gene, 1 for CEP17) HER2:CEP17 ratio ≥2 recognised but can be scored as FISH negative te to scoring ratio), should be retested with IHC	1 for HER2 gene HER2 signals >4 Is not recognised Is scored as
recognised but can be scored as FISH negative	Is not recognised Is scored as
	Is scored as
araffinisation and Proteolytic digestion rehydration	•
	FISH probe mix
HER2 FISH pharmDx™ protocol	Wash
	· · · · · · · · · · · · · · · · · · ·

Hofmann et al 2008

FISH interpretation



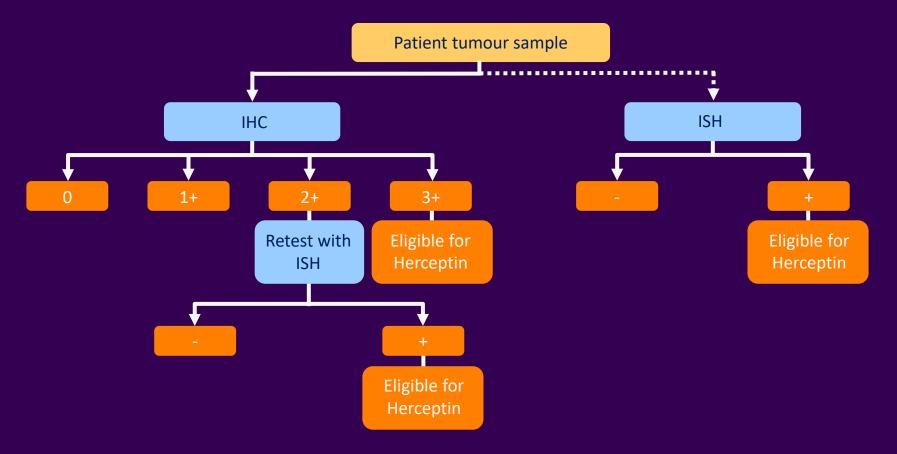
FISH negative (no amplification) Ratio of HER2 gene (orange) to CEP17 (green) signals is <2.0



FISH positive Ratio of orange to green signals is >2.0

Images courtesy of W Hanna using PathVysion

The HER2-testing algorithm



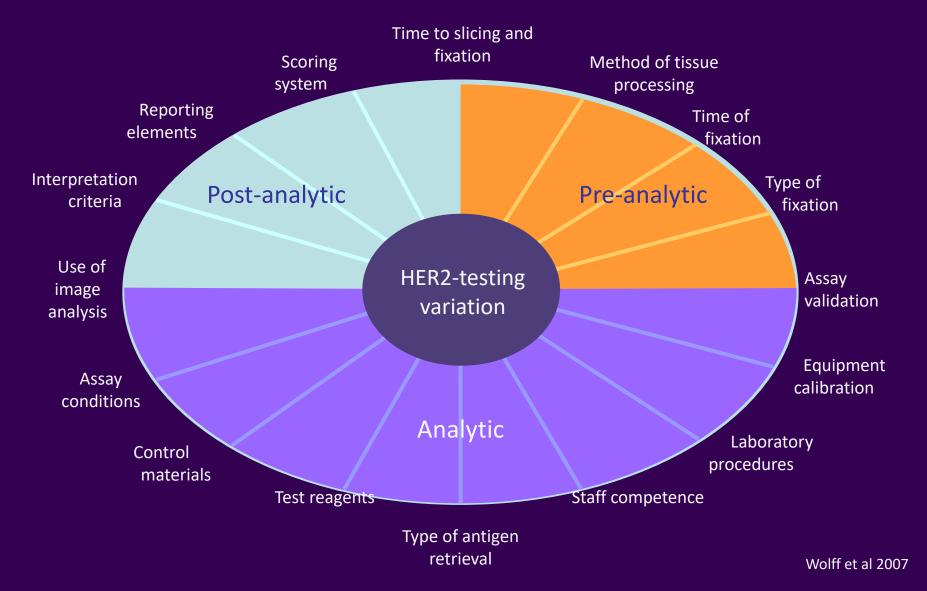
- If primary ISH testing is used, patients whose tumours overexpress HER2 (ie IHC 3+) may not be identified due to the HER2:FISH ratio being <2.0 (eg chromosome 17 polysomic cases, Hofmann et al 2007)
- ISH-detection mechanism can be fluorescent, chromogenic or silver

Concordance between IHC and FISH is 75-100%

Study	No. of cases	Overall concordance, ^a %
Di Palma et al 2007	161	93
Ricardo et al 2007	161	83, 82 ^b
van de Vijver et al 2007	209	81
Vocaturo et al 2006	111	76
Sapino et al 2003	106	85, 80 ^b
Dandachi et al 2002	171	92
Tanner et al 2000	157	98

^aFor IHC results, 3+ scores are considered positive; ^bstudy used different antibodies for IHC; therefore, concordance data are presented per antibody Only studies with >100 cases are shown

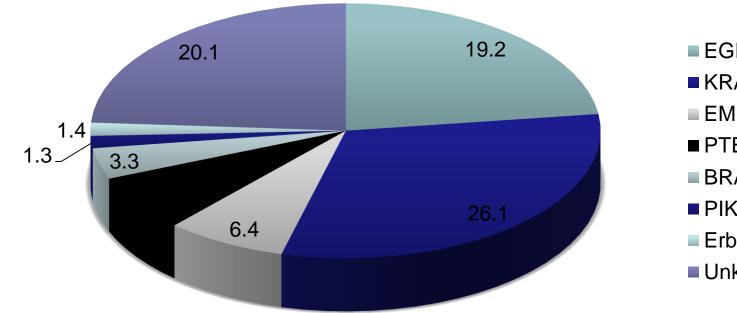
Sources of variation in HER2 testing

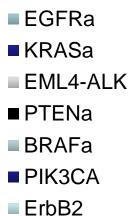


NON SMALL CELL LUNG CANCER

Lung cancer genetics – increasing complexity

Incidence of individual mutations for western NSCLC (adenocarcinoma)





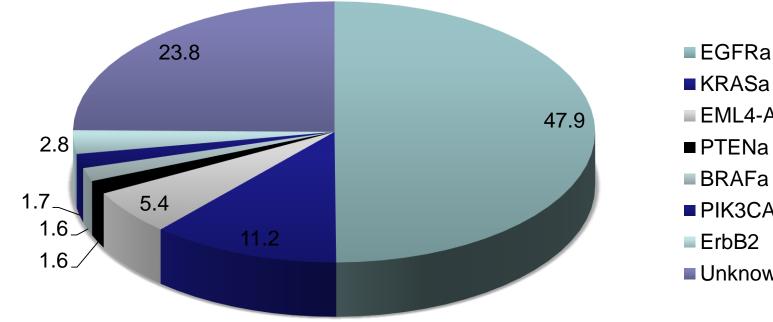
Unknown

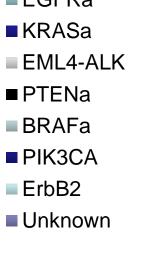


After Dearden et al., Ann Oncol 2013.

Lung cancer genetics – increasing complexity

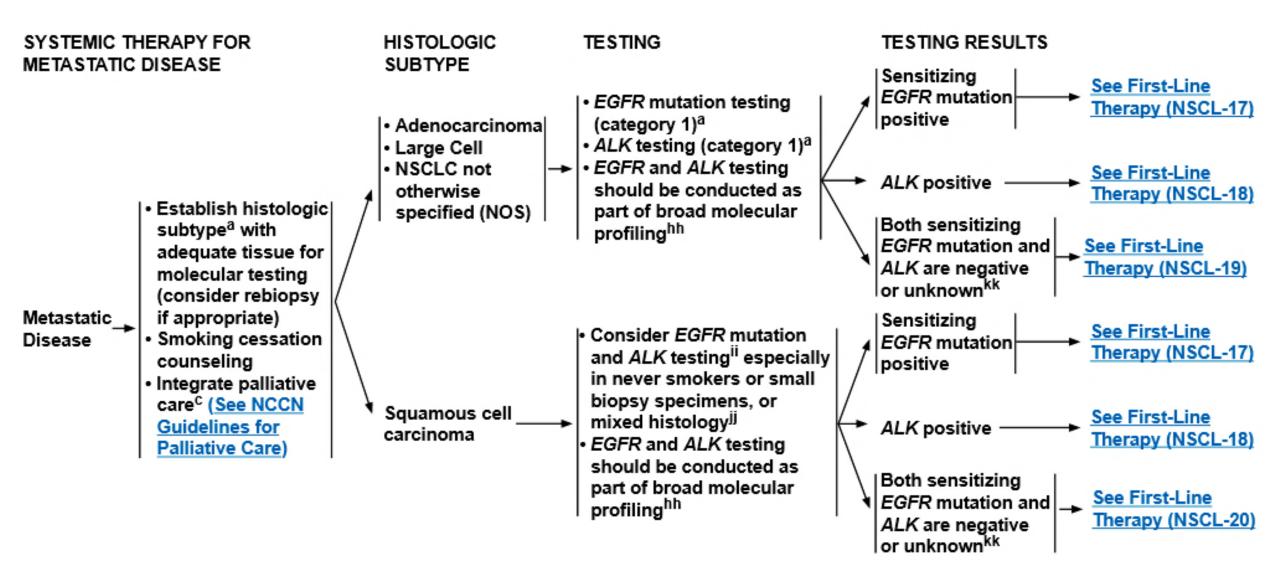
Incidence of individual mutations for asian NSCLC (adenocarcinoma)



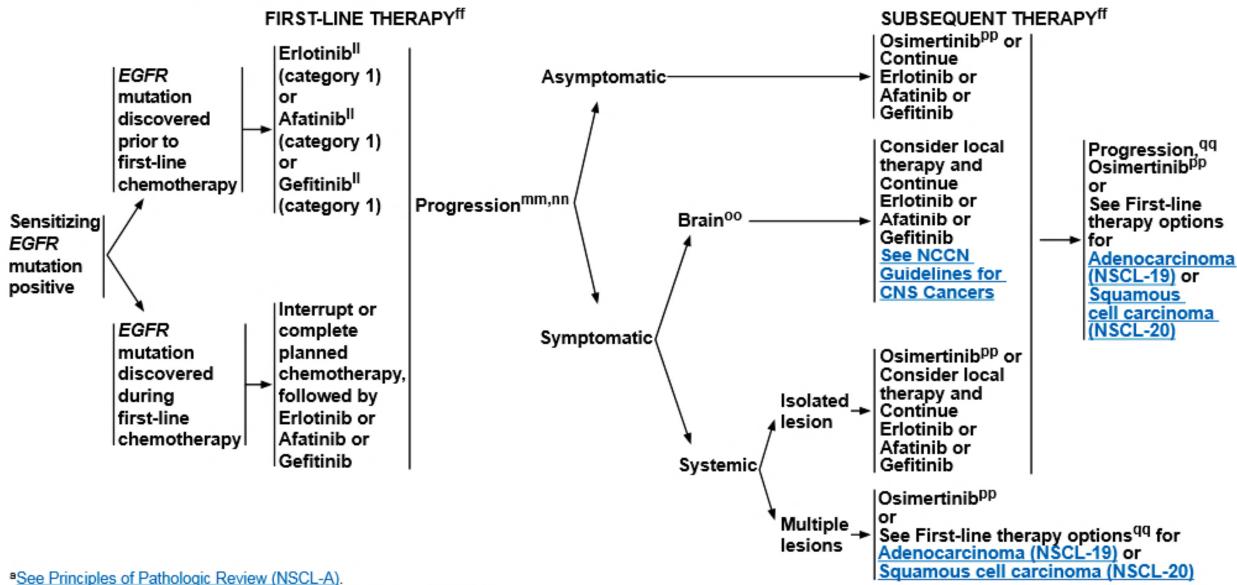




After Dearden et al., Ann Oncol 2013.

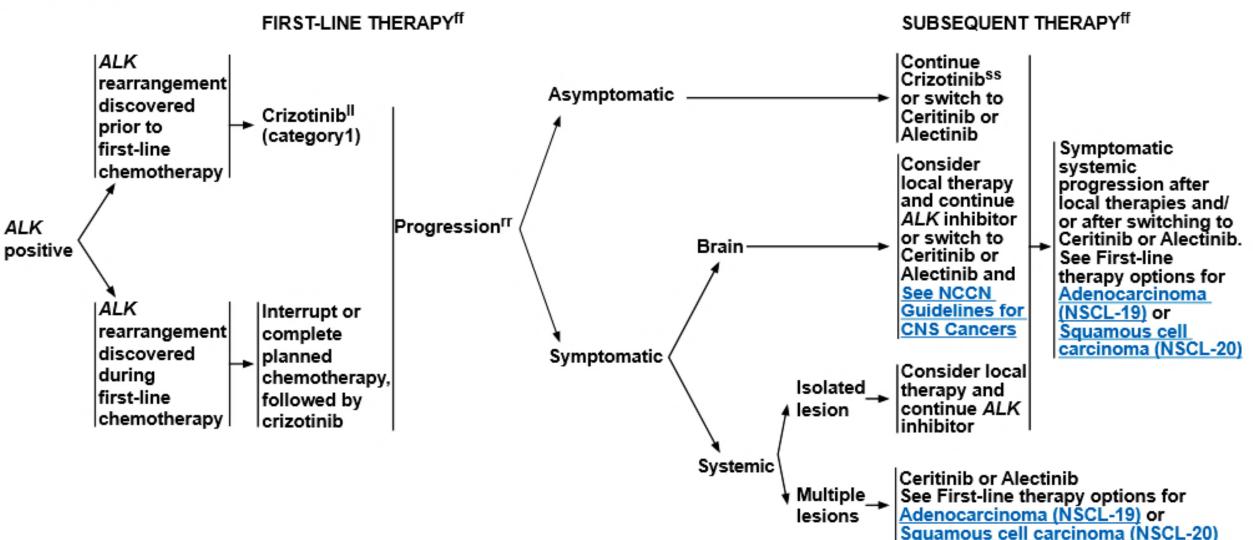


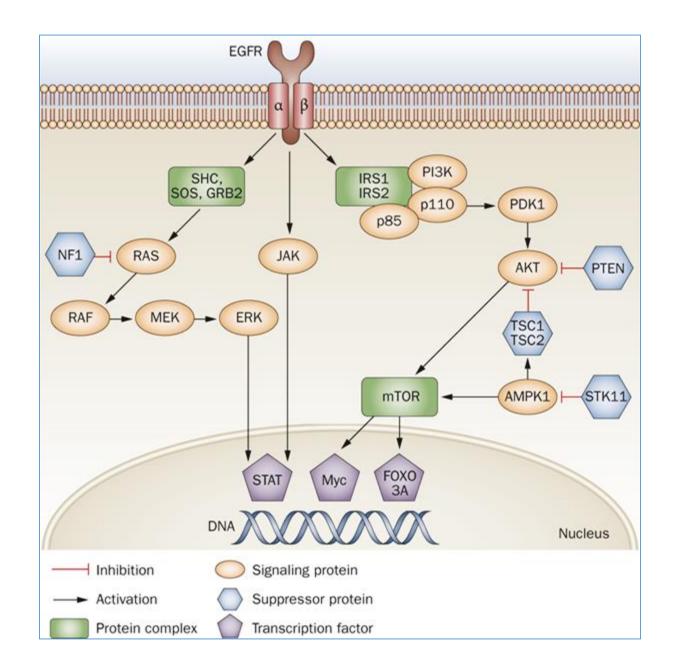
SENSITIZING EGFR MUTATION POSITIVE^a

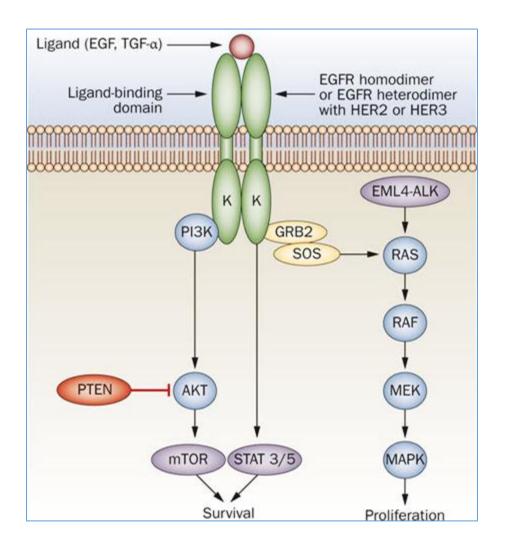


0.10.01

ALK POSITIVE^a



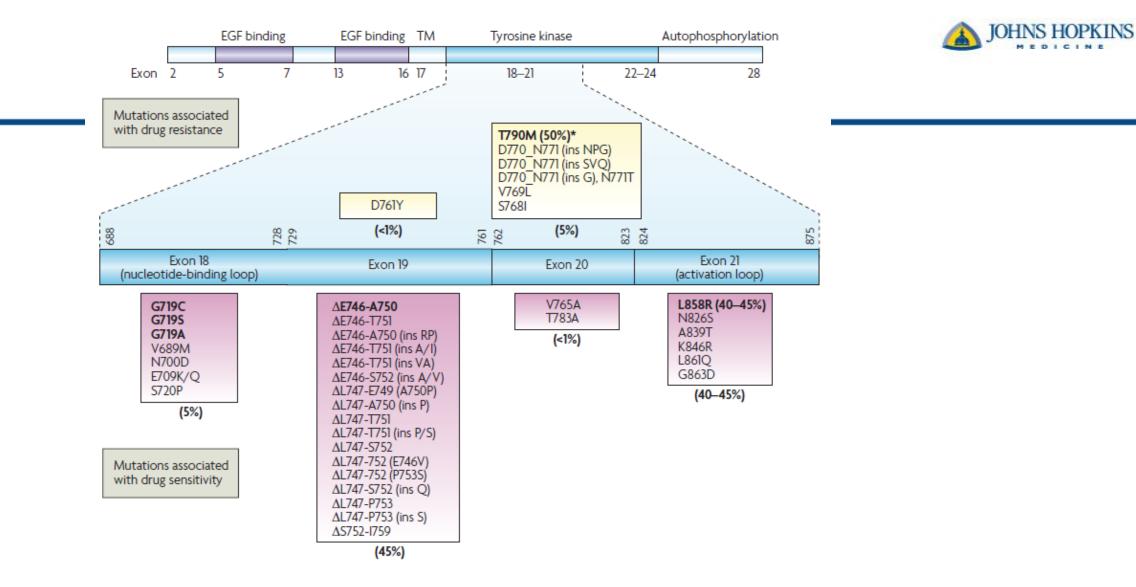




		Clinical dose of gefitinib 250 mg day ⁻¹ G	(G) or erlotinib (150 mg day ⁻¹ L ^E
In vitro equivalent (µM) 0	0.0001 0.001 0.001	0.1 1	10
Sensitivity	Hypersensitive	Sensitive	Insensitive
NSCLC cell lines	NCI-H3255 (1 nM) PC9 (4 nM)	NCI H2170 (200 nM) NCI H2073 (250 nM)	NCI-H1975 (12 μM) NCI-H1650 (9 μM) NCI-H460 (20 μM)
Genetic signatures	EGFR: L858R EGFR: Δ(E746-A750)	EGFR: wild type	EGFR: T790M PTEN loss KRAS

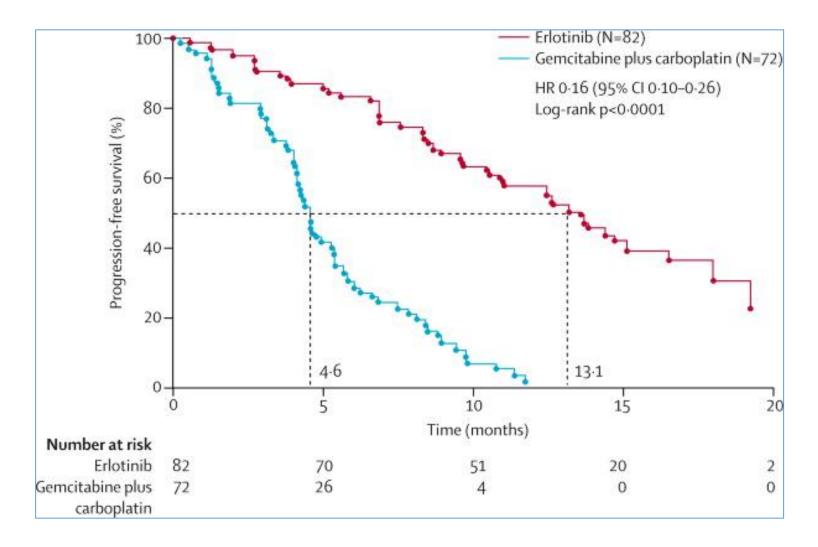
Sharma et al. Nature Reviews Cancer 7, 169–181 (March 2007) | doi:10.1038/nrc2088





EGFR mutations

Nat Rev Cancer (2007)7:169



OPTIMAL trial

cobas[®] EGFR Mutation Test v2 (IVD)

• The very first FDA-approved liquid biopsy assay, expanding targeted therapies to patients unable to contribute a tissue biopsy

INTENDED USE

The **cobas® EGFR Mutation Test v2** is a real-time PCR test for *the in vitro* qualitative detection and identification of mutations in exons 18, 19, 20, and 21 of the epidermal growth factor receptor (EGFR) gene in DNA derived from formalin-fixed paraffinembedded (FFPET) **tumor tissue and/or plasma** from non-small cell lung cancer (NSCLC) patients.

The **cobas**[®] EGFR Mutation Test v2 for use with plasma is a real-time PCR test for the **in vitro** qualitative and semi-quantitative measurement of mutations in exons 18, 19, 20, and 21 of the EGFR gene in human plasma. The EGFR test is further indicated for **serial measurement of EGFR mutation status** as an aid in the management of NSCLC cancer patients.

FFPET specimens are processed using the cobas[®] DNA Sample Preparation Kit and plasma specimens are processed using the **cobas[®] cfDNA Sample Preparation Kit**. The **cobas[®]** EGFR Mutation Test v2 and cobas z 480 analyzer are used for automated amplification and detection.



Semi Quantitative Index

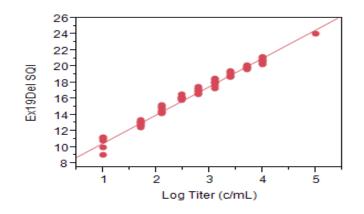
• New reporting tool for management of NSCLC patients

What is a Semi Quantitative Index (SQI)?

The SQI is a semi-quantitative measure of the amount of mutant cfDNA in a sample that can be used to measure the presence of EGFR mutations over time



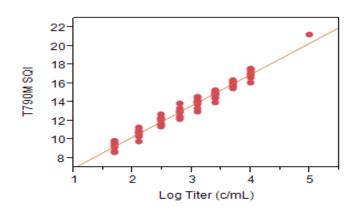
Linearity of mutant DNA in K2 EDTA Plasma: Ex19 Del cell line DNA



SI = 7.042 + 3.507 * Log Copies per mL R² = 0.981



Linearity of mutant DNA in K2 EDTA Plasma: T790M cell line DNA



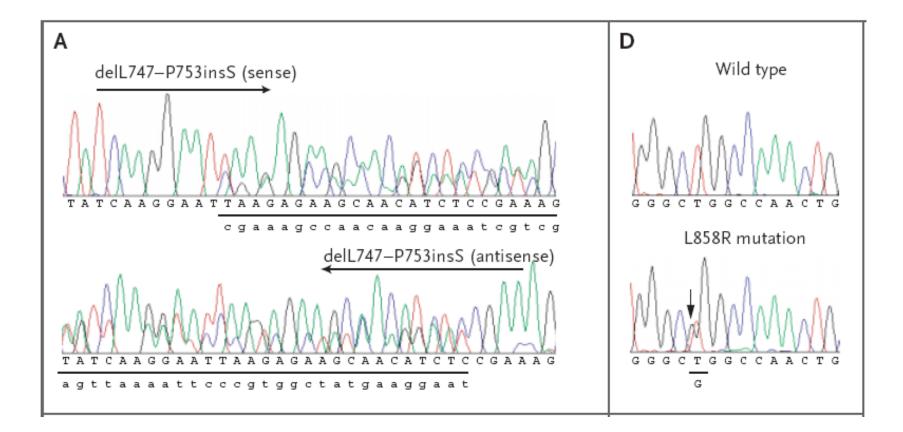
SQ = 3.593 + 3.352 *Log Copies per mL R² = 0.973

SQI TREND

Designed to reflect a change in the amount of mutant cfDNA over time per corresponding target mutation within a patient

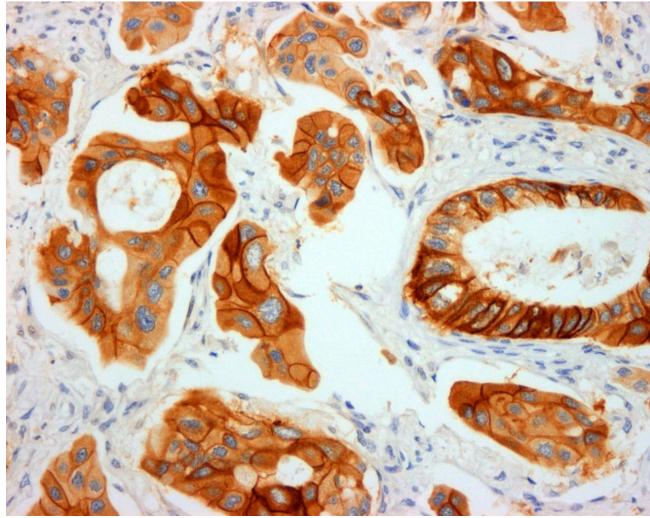
Direct sequencing of EGFR mutations





NEJM (2004)350:2129

Adenocarcinoma with positive staining for EGFR exon 21 L858R mutation-specific antibody (x200)



Cooper W A et al. *J Clin Pathol* Published Online First: 11 June 2013 doi:10.1136/jclinpath-2013-201607

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Warwick

Medical School