### Molecular Detection of EGFR ex20ins in NSCLC: Why, When, and How?

Satellite Symposium Sponsored by Janssen

HKIAP Scientific Meeting Fall 2023



Approval code: EM-143197

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## **Dr Molly Ll**

**Clinical Assistant Professor** 

 Department of Clinical Oncology, The Chinese University of Hong Kong

Dr Molly Li received his MBBS from the University of Hong Kong with distinction in medicine and psychiatry.

Principal investigator or co-investigator of over 15 lung cancer clinical trials

Principal investigator of a number of translational studies:

- Resistance mechanism and tumour immune microenvironment of EGFRmutation positive lung cancer after osimertinib failure
- Intratumoral heterogeneity of ALK-mutated lung cancer)
- preclinical studies testing novel drug combinations in cell lines and patient derived xenograft models

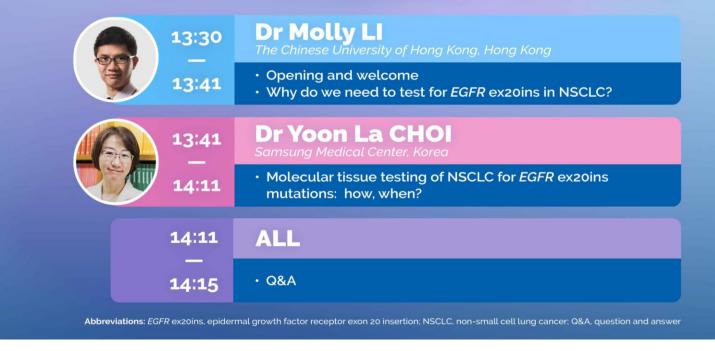
Selected as a participant of the ASCO Leadership Development Programme Asia Pacific 2023-2024.



THE INTERNATIONAL ACADEMY OF PATHOLOGY HONG KONG DIVISION FALL SCIENTIFIC MEETING 2023

MOLECULAR DETECTION OF EGFR ex20ins IN NSCLC: WHY, WHEN AND HOW?

#### 28<sup>th</sup> October (Saturday) | 13:30—14:15



- To highlight the clinical importance of molecular identification of EGFRex20ins in NSCLC
- To reflect on the recent evidence on the treatment of NSCLC with EGFRex20ins
- To demonstrate the limitation of molecular identification of EGFRex20ins by conventional methodologies (ie. PCR)
- To demonstrate the opportunities and challenges of identifying patients with EGFRex20ins by next generation sequencing (NGS)



香港中文大學 The Chinese University of Hong Kong



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### Calling for EGFR exon 20 insertion testing in NSCLC

Molly SC Li MBBS (HK), MRCP (UK), FHKCP, FHKAM Clinical Assistant Professor, Department of Clinical Oncology, CUHK

Email: <u>molly@clo.cuhk.edu.hk</u> Twitter/X: @mollylisc

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### Disclosure

Honorarium	AstraZeneca Novartis Amgen Pfizer Takeda ACE Oncology
Advisory Board	Amgen AnHeart AstraZeneca Pfizer Takeda
Research Funding	Gilead MSD Takeda

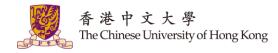






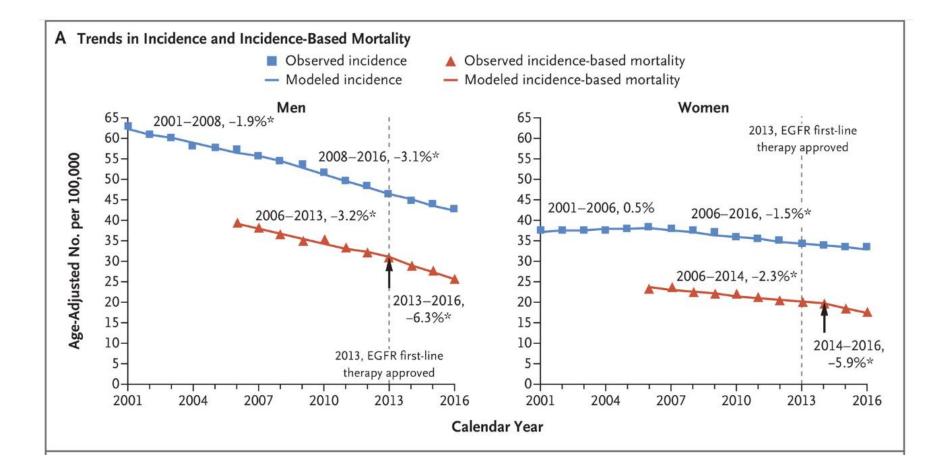
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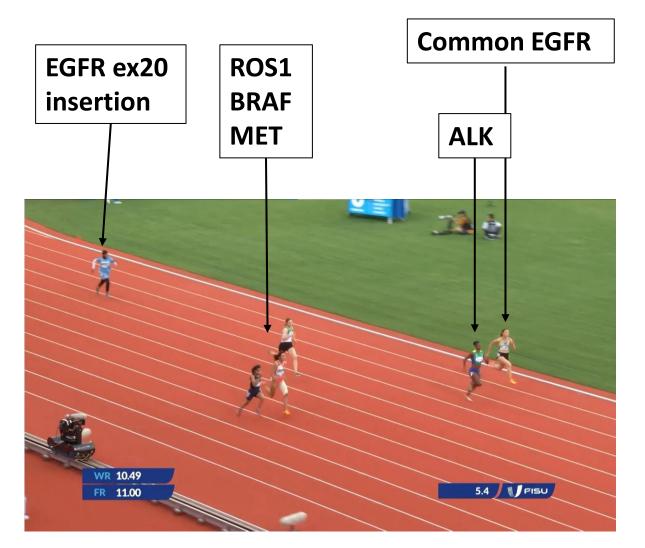
#### Personalized medicine is the cornerstone of Lung Cancer Treatment







Howlander et al NEJM 2020



#### Why?

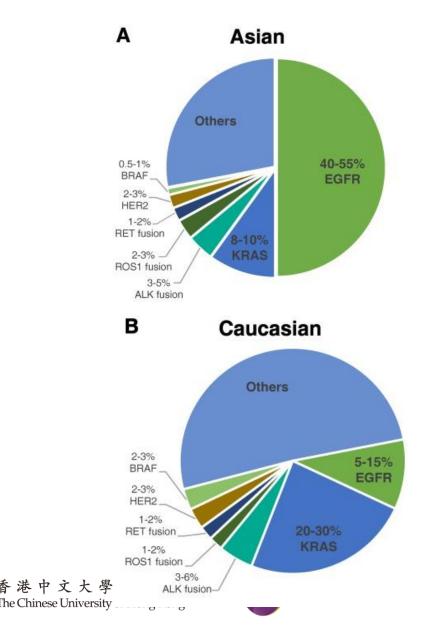
- It is a rare mutation
- EGFR ex20 ins are "TKI resistant"
- Approved EGFR PCR tests (more intended to capture common EGFR) can also detect EGFR exon 20 insertion

#### No harm in missing a case of EGFR exon 20ins?

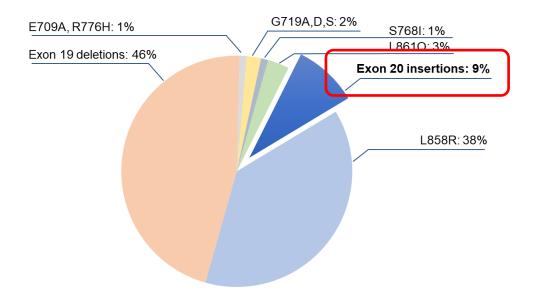




### **EGFR exon 20 insertion is not rare in Asia!**



#### Frequency of EGFR Mutations in NSCLC<sup>1</sup>

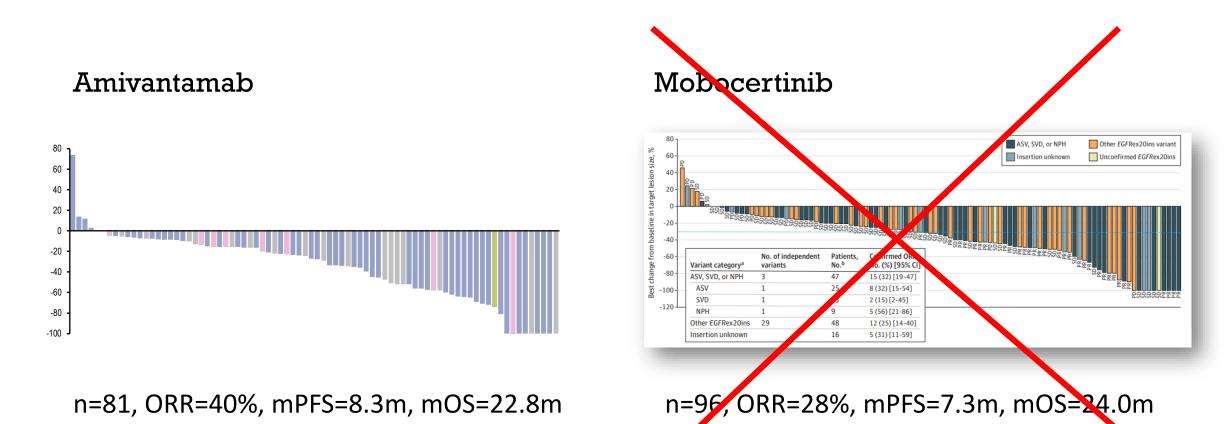


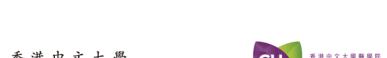
EGFR exon 20 ins is the  $3^{rd}$  most prevalent type of primary EGFR mutation in NSCLC

1. Oncol Rep. 2017 Mar; 37(3): 1347–1358;

2. Arcila ME. Mol Cancer Ther. 2013;12(2):220-229

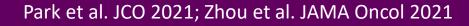
# Amivantamab and Mobocertinib are approved for patients with NSCLC harbouring EGFR ex20 ins



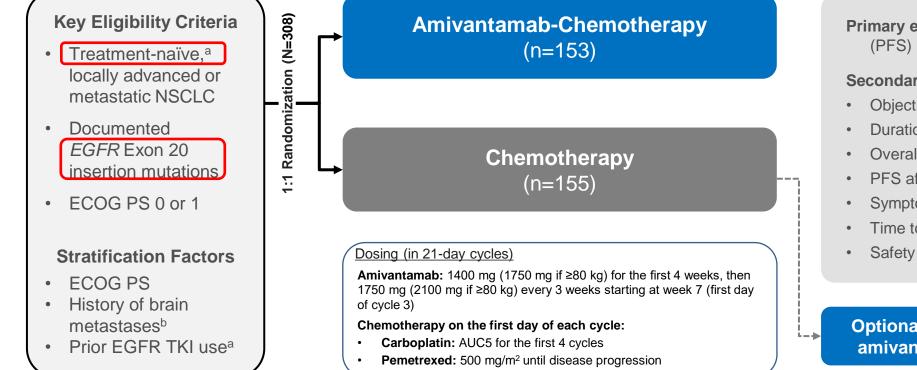


he Chinese University of Hong Kong

Faculty of Medicine



# PAPILLON: Phase 3 Study Design



Primary endpoint: Progression-free survival (PFS) by BICR according to RECIST v1.1<sup>c</sup>

#### Secondary endpoints:

- Objective response rate (ORR)<sup>c</sup>
- Duration of response (DoR)
- Overall survival (OS)<sup>c</sup>
- PFS after first subsequent therapy (PFS2)
- Symptomatic PFS<sup>d</sup>
- Time to subsequent therapy<sup>d</sup>

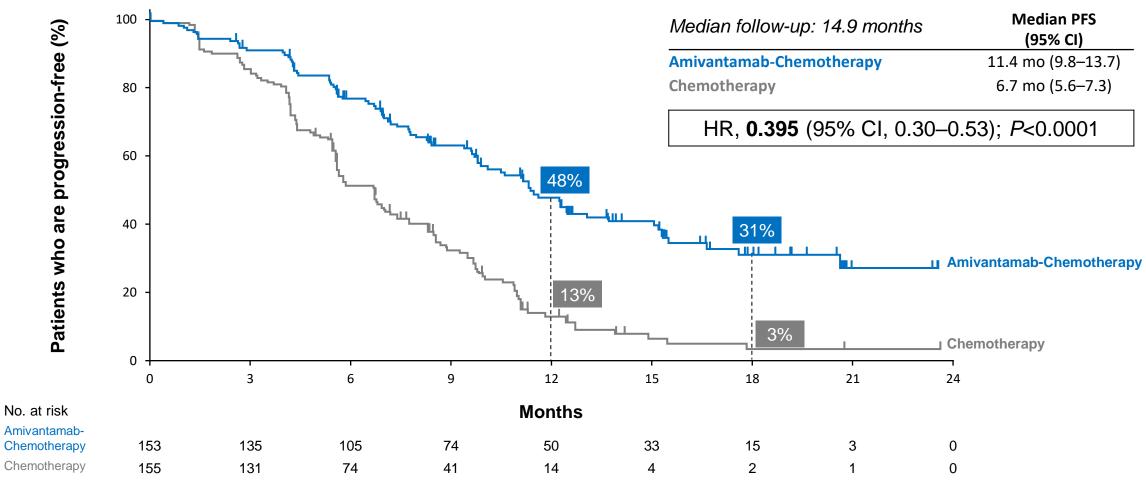
Optional crossover to 2<sup>nd</sup>-line amivantamab monotherapy<sup>e</sup>

#### Girard et al. ESMO 2023





### **Primary Endpoint: Progression-free Survival by BICR**



Girard et al. ESMO 2023





### Targeted therapy efficacy in different driver oncogenes

Gene	Targeted Therapy	Median PFS	ORR	Ref
EGFR	Gefitinib	10.9m	71%	Mok et al NEJM 2009
ALK	Crizotinib	10.9m	74%	Solomon et al NEJM 2014
BRAF V600E	Dabrafenib + Trametinib	10.2m	64%	Planchard et al. JTO 2022
MET ex14	Tepotinib	8.5m	46%	Paik et al. NEJM 2020
ERBB2 ex20	Trastuzumab Deruxtecan	8.2m	55%	Li et al NEJM 2022
EGFR ex20ins	Amivantamab + Chemotherapy	11.4m	73%	Girard et al. ESMO 2023





### The field of EGFR exon 20ins is advancing rapidly

#### Efficacy and Toxicity Cross-comparison in pretreated pts

			Efficacy		Toxicity		
Drug	Class	Ν	ORR (95% CI)	mPFS (95% Cl)	Rash all / gr 3	Diarrhoea	Other
Amivantamab	EGFR-MET bispecific mAb	81	40% (29-51%)	8.3 (6.5-10.0)	86% / 4%	12% / 4%	Infusion reactions, edema
Mobocertinib	EGFR TKI	114	28% (20-37%)	7.3 (5.5-9.2)	45% / 0%	91% / 21%	
Poziotinib	EGFR TKI	42	31% (19-46%)	5.5 (5.4-10.4)	90% / 34%	92% / 22%	
Sunvozertinib	EGFR TKI	104	61% (50-71%)	n.a.	80% / 1%	20% / 3%	
Zipalertinib	EGFR TKI	73	38% (27-49%)	10 (6-12)	80% / 1%	30% / 3%	
Furmonertinib	EGFR TKI	26	46% (27-67%)	n.a.	21% / 0%	86% / 0%	





### Any harm in missing EGFR ex20ins NSCLC?

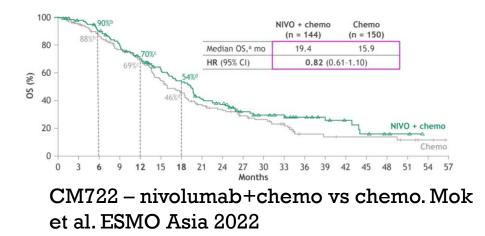
70yo/F, non-smoker p/w massive pleural effusion Cytology: TTF1+ adenocarcinoma

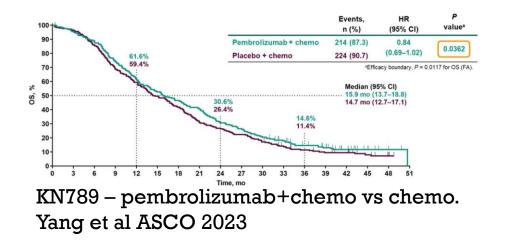
Multiplex PCR: EGFR/ALK/ROS1/BRAF/MET/RET/KRAS -ve PD-L1 TPS (22C3): 60%

1<sup>st</sup> line: Pembrolizumab monotherapy Further disease progression after 3 cycles

NGS testing: EGFR exon 20 H773\_V774dup

Switched to  $2^{nd}$  line chemotherapy  $\rightarrow$  responded









### Take Home Message

#### Call for EGFR exon 20 insertion testing in NSCLC

- Its not uncommon! (more common than ROS-1, RET, BRAF V600E...)
- There is approved targeted therapy
- Moving EGFR exon 20ins targeted therapy to the 1L setting
- May harm patients in missing EGFR exon 20 ins offering immunotherapy inappropriately







## **Professor Yoon-La (Yuna) CHOI**

Professor

- Department of Pathology & Translational Genomics, Samsung Medical Centre
- Department of Health Science and Technology, Samsung Advanced Institute for Health Science and Technology

Professor Choi obtained her M.D. and Ph.D in the Seoul National University College of Medicine. She has over 30 selected publication in high impact factor peer-reviewed journal. Her research interest include:

- molecular diagnostics development and validation,
- biomarker development and validation,
- companion diagnostics for target therapy and immunotherapy,
- cancer genomics and molecular pathology



2023. 10. 29 HKIAP\_EGFR exon20ins

삼성서올병원

### Molecular tissue testing for NSCLC for EGFRex20ins mutations: How, when? Perspective of Pathologist

#### Yoonla Choi

Department of Pathology & Translational Genomics Samsung Medical Center Sungkyunkwan University School of Medicine South Korea





### **DECLARATION OF INTERESTS**

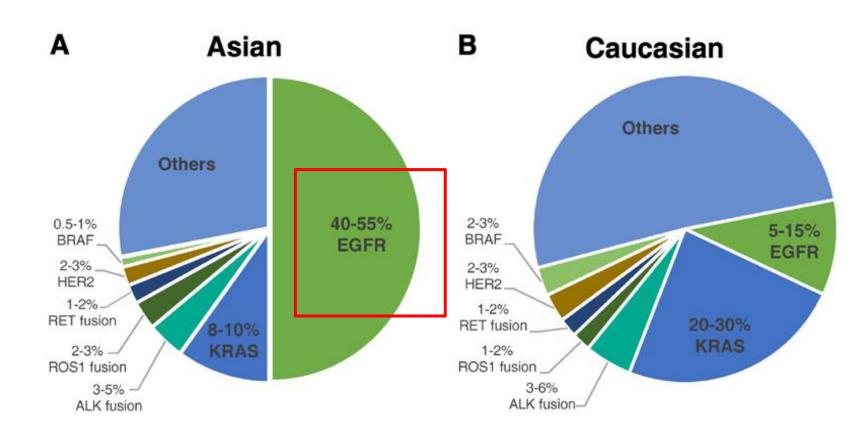
Lectures, Advisory boards, Consultant and Research support to institute

AstraZeneca, Roche, Novartis, Takeda, Janssen, MSD, Bayer, Merk, Pfizer, LG chemical, Yuhan, Panagene, Gencurix, Genecast

### **CONTENTS**

- 1. EGFR testing
- 2. EGFR Exon20 insertion
- 3. EGFR Exon20 insertion testing

# Frequency of oncogene-driven genes in NSCLC patients of Asian and Caucasian



- Asians display a 40-55% prevalence of EGFR mutation.
- This elevated frequency underscores the critical importance of EGFR testing in Asian NSCLC patients, emphasizing tailored treatment approaches for this population.
- Furthermore, when adjuvant TKI therapy is considered for NSCLC at an early stage, we need to test for EGFR mutations as well.

### Molecular biomarker analysis : Testing methodologies

- NGS
  - Broad-based genomic testing approaches that efficiently utilize limited biopsy tissue while maximizing diagnostic genomic information are most commonly NGS-based.

#### PCR (RT-PCR, real-time PCR, digital PCR)

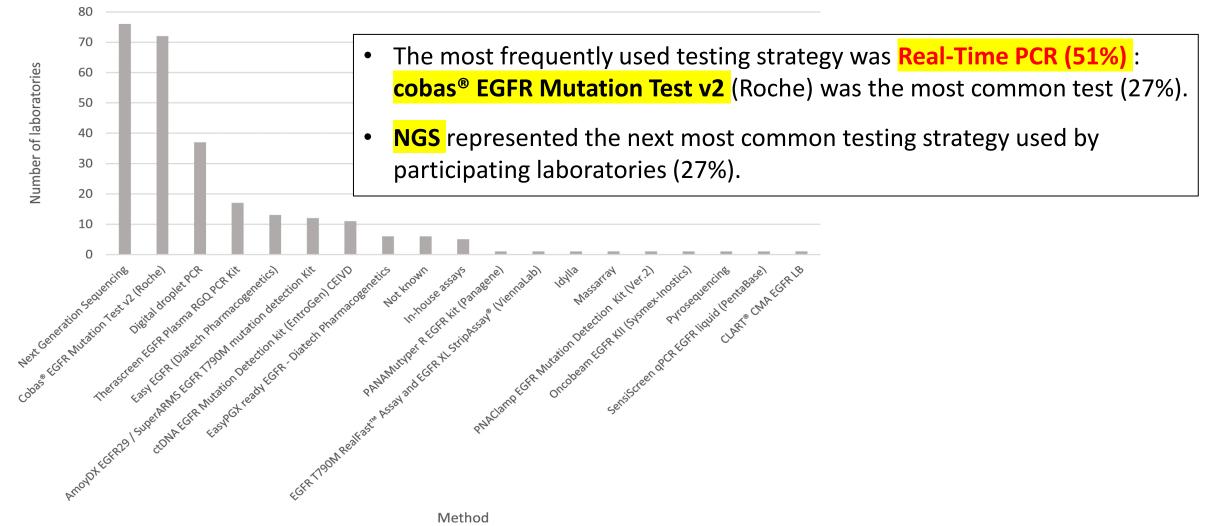
• Only those specific alterations that are targeted by the assay are assessed and the potential for mutations outside of the testing scope of the specific assay must be considered.

#### Sanger sequencing

- Requires the greatest degree of tumor enrichment
- Not appropriate for assays in which identification of subclonal events (eg, resistance mutations)

#### EGFR testing methods among 264 laboratories from 45 countries

Summary of testing methods used by participating laboratories



Usage of epidermal growth factor mutation testing and impact on treatment patterns in non-small cell lung cancer: An international observational study

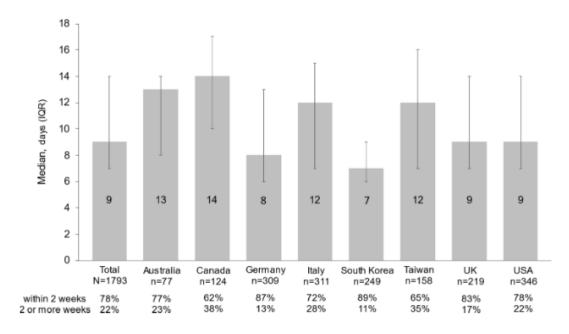
Janakiraman Subramanian<sup>a,\*,1</sup>, Natasha B. Leighl<sup>b,2</sup>, Yoon-La Choi<sup>c,3</sup>, Teh-Ying Chou<sup>d,4</sup>, Jeffrey Gregg<sup>e,5</sup>, Rina Hui<sup>f,6</sup>, Antonio Marchetti<sup>g,7</sup>, Mark Silvey<sup>h,8</sup>, Rebecca Makin<sup>h,9</sup>, Liane Gillespie–Akar<sup>h,10</sup>, Aliki Taylor<sup>i,11</sup>, Doreen A Kahangire<sup>i,12</sup>, Tom Bailey<sup>h,13</sup>, Maiyan Chau<sup>j,14</sup>, Neal Navani<sup>k,15</sup>

- A cross-sectional medical chart review was completed May–August 2020 in <u>Australia</u>, <u>Canada, Germany</u>, Italy, South Korea, Taiwan, UK, and USA.
- **1,793** patients with advanced (stage IIIb/IIIc/IV) NSCLC
- 78% of EGFRm test results were received ≤ 2 weeks (7-14d) from order to result
- Median time from Dx to EGFRm test result : 18 d(10–22 d)
- **37 %** of patients received a systemic treatment prior to EGFRm result;

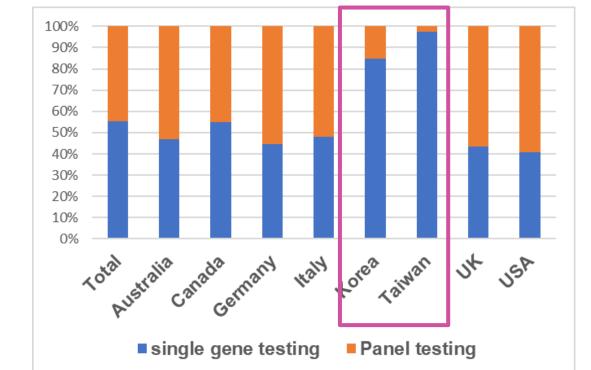
#### Real world EGFR testing patterns

Time from EGFR mutation test request to receipt of test result in patients with NSCLC

### Time from EGFRm test request to receipt of test result in patients with NSCLC



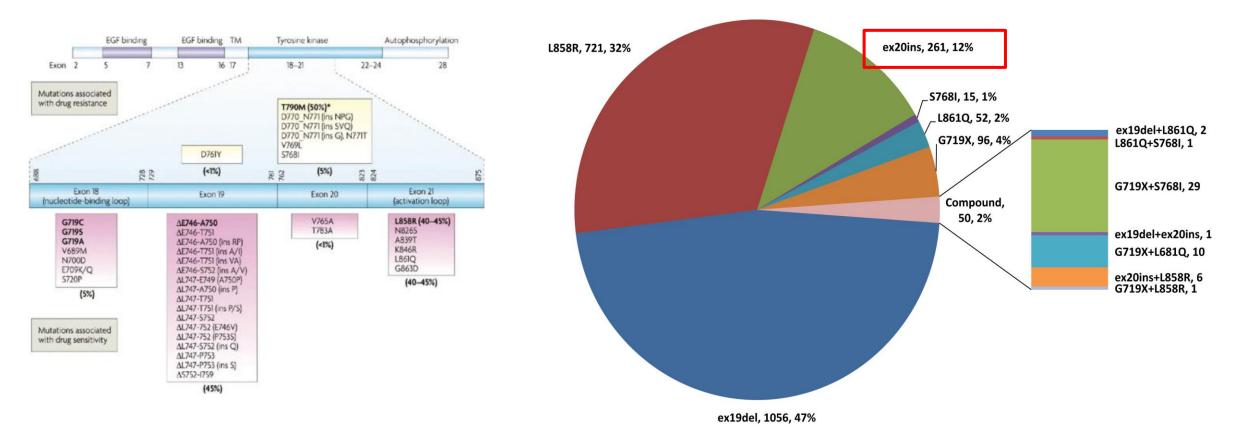
It took 7 to 14 days to get the results after requesting the test.



While panel and single gene tests are equally prevalent in other countries, Korea and Taiwan are more likely to perform single gene tests.

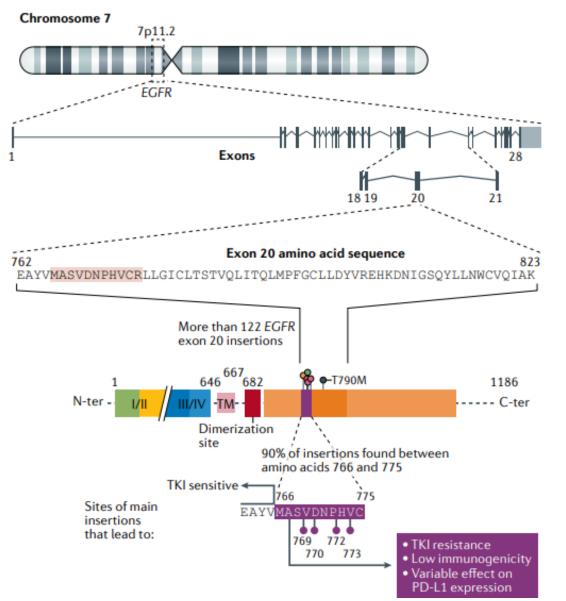
#### Single gene test vs. Panel test

### EGFRex20ins are the third most common type of EGFR mutation



- Detected following *EGFR* exon 19 deletions (47%) and EGFR L858R (32%).
- CGP performed on 14,483 NSCLC cases in the course of clinical care identified 2251 cases with *EGFR* mutations.
- 263 of these cases were *EGFR*ex20ins, representing **12%** of all *EGFR*-mutant NSCLC and **1.8%** of all NSCLC cases tested.

#### EGFR gene and protein domain organization and summary of exon 20 ins



- In EGFR exon 20, in addition to p.T790M mutation, in-frame insertions and 3–21-bp duplications (corresponding to 1–7 amino acid) can occur within amino acids 762 and 774.
- At least 122 different EGFR exon 20 insertions have been identified, with ~90% occurring in the region comprising amino acids 766–775.
- Different insertion sites are associated with differing disease phenotypes and patient characteristics, sensitivity to current therapies and, potentially, tumour immunogenicity.

#### EGFR mutation testing in NSCLC 2023

NCCN NCCN NCCN Network®

Comprehensive Cancer Network® NCCN Guidelines Version 4.2023 Non-Small Cell Lung Cancer

#### TESTING RESULTS<sup>II,mm</sup>

EGFR exon 19 deletion or exon 21 L858R mutation positive	NSCL-20
EGFR S768I, L861Q, and/or G719X mutation positive	NSCL-23
EGFR exon 20 insertion mutation positive	NSCL-24
KRAS G12C mutation positive	NSCL-25
ALK rearrangement positive	NSCL-26
ROS1 rearrangement positive	NSCL-29
BRAF V600E mutation positive	<u>NSCL-31</u>
NTRK1/2/3 gene fusion positive	NSCL-32
METex14 skipping mutation positive	NSCL-33
RET rearrangement positive	NSCL-34
ERBB2 (HER2) mutation positive	NSCL-35
PD-L1 ≥1% and negative for actionable molecular biomarkers above	NSCL-36
PD-L1 <1% and negative for actionable molecular biomarkers above	NSCL-37

 A recent guideline lists several specific mutation variants of EGFR based on the therapeutic value of each variant.

#### EGFR exon20 insertion : Patient characteristics

		,	AII		R Exon sertion	EC	nmon SFR ation	
		Ν	%	Ν	%	N	%	<i>p</i> -value*
Cohort Size	N	2209	100.00%	53	2.40%	1978	89.5%	
Age at advanced NSCLC diagnosis	Mean (SD) Median (Q1-Q3) Min-Max	62.00 (55	(11.15) 5.00-71.00) - 94.00	58.00 (	3 (13.15) (50.0-70.0) 0 - 82.00	62.00 (5	(10.93) 55.0-70.0) - 94.00	0.023
Gender	Female Male	1321 888	59.80% 40.20%	28 25	52.83% 47.17%	1198 780	60.57% 39.43%	0.320
Smoking Status	Never smoker Former Current Unknown	1451 505 246 7	65.69% 22.86% 11.14% 0.32%	30 13 10 0	56.60% 24.53% 18.87% 0.00%	1314 450 208 6	66.43% 22.75% 10.52% 0.30%	0.221
Performance Score ECOG at diagnosis	0 1 2 3 4 Unknown	521 1157 143 31 5 352	23.59% 52.38% 6.47% 1.40% 0.23% 15.93%	10 29 5 3 0 6	18.87% 54.72% 9.43% 5.66% 0.00% 11.32%	471 1037 123 26 3 318	23.81% 52.43% 6.22% 1.31% 0.15% 16.08%	0.134
TNM Classification	IIIB/C IV	126 2083	5.70% 94.30%	4 49	7.55% 92.45%	108 1870	5.46% 94.54%	0.533
Histology	NSQ SQ Other specified NOS	2139 33 13 24	96.83% 1.49% 0.59% 1.09%	50 0 <3 <3	94.34% 0.00% <5.66% <5.66%	1923 26 9 20	97.22% 1.31% 0.46% 1.01%	0.040

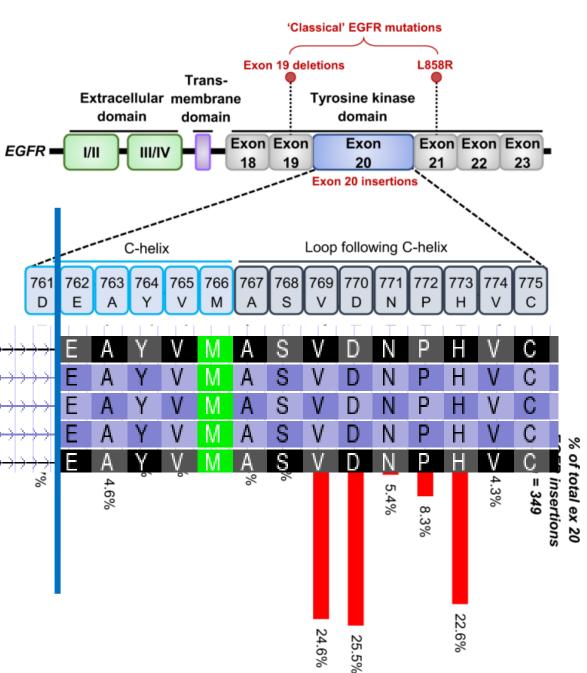
Table 1: Characteristics of Exon 20 insertion patients vs common EGFR mutations

NSCLC, Non-small cell lung cancer; SD, standard deviation; NSQ, Non-squamous cell carcinoma; SQ, squamous cell carcinoma; NOS, Not otherwise specified. \*Student's t-test was used to compare age at advanced NSCLC diagnosis between groups. For categorical variables, Chi-square test or Fisher exact test when any of expected cell count < 5 was used.

#### EGFR exon 20

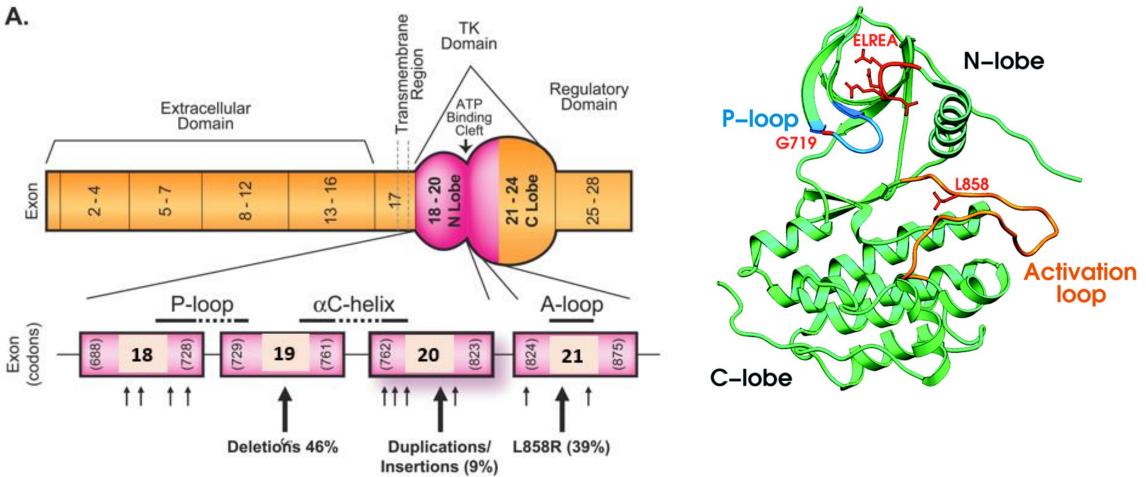
Exon20 starts 762E

- Heterogenous at the molecular level
- Inframe insertions or duplications of 3~21 bp (761~775 aa of the EGFR)



#### EGFR exon 20

C-helix : **761**-766 Exon20 : **762**-823



#### Heterogeneity of EGFR exon20 insertion mutations in NSCLC

1	774	p.V774_C775insHV	p.V774_C775insPHV	p.V774_C775insPR	
/	773 H	p.H773_V774dup p.H773_V774insAH p.H773_V774insG	p.H773_V774insGH p.H773_V774insGNPH p.H773_V774insH	p.H773_V774insNPH p.H773_V774insNPHT p.H773_V774insPH	p.H773_V774insQ p.H773_V774insTQPP p.H773dup
/	772 P	p.P772_H773dup p.P772_H773ins? p.P772_H773insDHP p.P772_H773insDNP	p.P772_H773insNP p.P772_H773insPNP p.P772_H773insPR p.P772_H773insQCP	p.P772_H773insQV p.P772_H773insTHP p.P772_H773insV	p.P772_H773insVDN p.P772_H773insX p.P772_H773insYNP
	L <sup>Z</sup> N	p.N771_H773dup p.N771_P772ins? p.N771_P772insG	p.N771_P772insH p.N771_P772insHH p.N771_P772insN	p.N771_P772insRH p.N771_P772insTQPNP	p.N771_P772insV p.N771dup
	0 2 2	p.D770_N771insAPW p.D770_N771insAWT p.D770_N771insD p.D770_N771insG p.D770_N771insGD	p.D770_N771insGF p.D770_N771insGL p.D770_N771insGT p.D770_N771insH p.D770_N771insMATP	p.D770_N771insPAW p.D770_N771insSN p.D770_N771insSVD p.D770_N771insSVE p.D770_N771insSVP	p.D770_N771insSVQ p.D770_N771insVDSVDNP p.D770_N771insY p.D770_P772dup p.D770Efs*61
	769 V	p.V769_D770insANV p.V769_D770insASV p.V769_D770insCV p.V769_D770insERG	p.V769_D770insGGTR p.V769_D770insGRV p.V769_D770insGSV	p.V769_D770insGV p.V769_D770insGVG p.V769_D770insGVV	p.V769_D770insMASVD p.V769_N771dup p.V769dup
	768 S	p.S768_D770dup	p.S768_V769insAWT	p.S768_V769insVAS	
	767 A	p.A767_S768insIA	p.A767_V769dup	p.A767_S768insTLA	p.A767_S768insYVM
	765	p.V765_M766insMAS			
	764 Y	p.Y764_V765insHH			
	763 A	p.A763_Y764insFQEA			
Ì	761 D	p.D761_E762insEAFQ			
ļ	756 N	p.N756dup			

- Over 100 different *EGFR* exon 20 insertion mutations have been reported in NSCLC
- The majority are composed of 1 to 4 AA insertions located in the loop following the C-helix

Exon

Exon

Exon 21

Exon 20

Exon 19

11

EGFR

### EGFR exon20 insertion

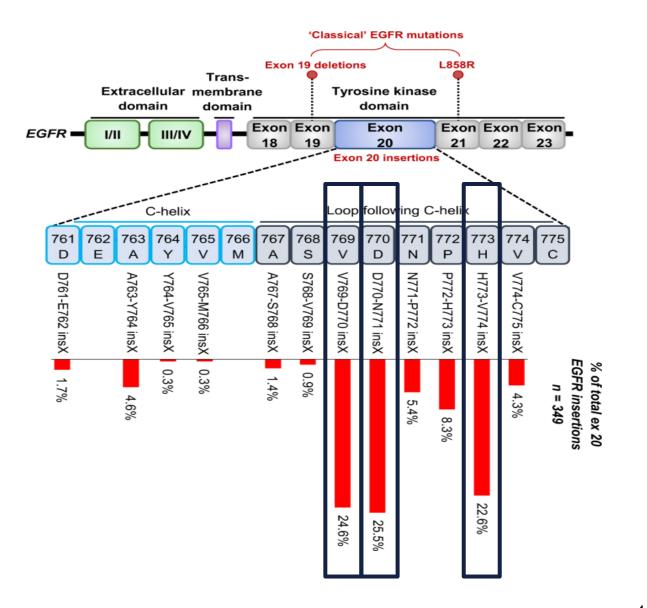
#### **cobas**<sup>®</sup> EGFR Mutation Test v2

p.V769\_D770insASV

p.D770\_N771insSVD

p.D770\_N771insG

p.H773\_V774insH



#### EGFR exon20 insertion CDx

Only three diagnostic products are CDx for Exon20 insertion mutations

Diagnostic name	Method	Drug	Biomarker
Oncomine Dx Target Test	Amplicon NGS	Rybrevant(amivantamb)	Exon 20 insertion mutations
Guardant360 CDx	Hybridization NGS	Rybrevant(amivantamb)	Exon 20 insertion mutations
FoundationLiquid CDx	Hybridization NGS	Exkivity (mobocertinib)	Exon 20 insertion mutations

- Cobas EGFR Mutation Test v2
- FoundationOne CDx
- ONCO/Reveal Dx Lung & Colon Cancer Assay
- therascreen EGFR RGQ PCR Kit

Exon 19 deletion or exon 21 L858R substitution mutation

T790M mutation

### EGFR exon20 insertion Testing: RWD

Exon 18 G719X [c.2156G>C p.G719A or c.2155G>A p.G719S or c.2155G>T p.G719C, associated with sensitivity to anti-EGFR therapy]138Exon 19 (in-frame deletion, associated with sensitivity to anti-EGFR therapy]141Exon 20 [c.2303G>T p.S768I, associated with low sensitivity to anti-EGFR therapy]134Exon 20 [c.2369C>T p.T790M, associated with resistance resistance to anti-EGFR therapy]141Exon 20 [c.2369C>T p.T790M, associated with resistance resistance to anti-EGFR therapy]141Exon 20 [c.2369C>T p.T790M, associated with resistance resistance to anti-EGFR therapy]141Exon 20 [c.2369C>T p.T790M, associated with resistance resistance to anti-EGFR therapy]141Exon 21 [c.2573T>G p.L858R, associated with sensitivity anti-EGFR therapy]142Exon 21 [c.2582T>A p.L861Q, associated with sensitivity142	6. Which <i>EGFR</i> mutations does your laboratory report?	Freq*
associated with sensitivity to anti-EGFR therapy]138Exon 19 (in-frame deletion, associated with sensitivity to anti-EGFR therapy]141Exon 20 [c.2303G>T p.S768I, associated with low sensitivity to anti-EGFR therapy]134Exon 20 [c.2369C>T p.T790M, associated with resistance resistance to anti-EGFR therapy]141Exon 20 [c.2369C>T p.T790M, associated with resistance resistance to anti-EGFR therapy]141Exon 20 (insertion/duplication, typically associated with resistance to anti-EGFR therapy)141Exon 21 [c.2573T>G p.L858R, associated with sensitivity anti-EGFR therapy]142	Exon 18 G719X [c.2156G>C p.G719A or	
Exon 19 (in-frame deletion, associated with sensitivity to anti-EGFR therapy] 141   Exon 20 [c.2303G>T p.S768I, associated with low sensitivity to anti-EGFR therapy] 134   Exon 20 [c.2369C>T p.T790M, associated with resistance resistance to anti-EGFR therapy] 141   Exon 20 [c.2369C>T p.T790M, associated with resistance resistance to anti-EGFR therapy] 141   Exon 20 [c.2369C>T p.T790M, associated with resistance resistance to anti-EGFR therapy] 141   Exon 20 (insertion/duplication, typically associated with resistance to anti-EGFR therapy) 131   Exon 21 [c.2573T>G p.L858R, associated with sensitivity anti-EGFR therapy] 142	c.2155G>A p.G719S or c.2155G>T p.G719C,	
anti-EGFR therapy]141Exon 20 [c.2303G>T p.S768I, associated with low sensitivity to anti-EGFR therapy]134Exon 20 [c.2369C>T p.T790M, associated with resistance resistance to anti-EGFR therapy]141Exon 20 (insertion/duplication, typically associated with resistance to anti-EGFR therapy)141Exon 20 (insertion/duplication, typically associated with resistance to anti-EGFR therapy)131Exon 21 [c.2573T>G p.L858R, associated with sensitivity anti-EGFR therapy]142	associated with sensitivity to anti-EGFR therapy]	138
Exon 20 [c.2303G>T p.S768I, associated with low sensitivity to anti-EGFR therapy] 134   Exon 20 [c.2369C>T p.T790M, associated with resistance resistance to anti-EGFR therapy] 141   Exon 20 (insertion/duplication, typically associated with resistance to anti-EGFR therapy) 131   Exon 21 [c.2573T>G p.L858R, associated with sensitivity anti-EGFR therapy] 142	Exon 19 (in-frame deletion, associated with sensitivity to	
sensitivity to anti-EGFR therapy]134Exon 20 [c.2369C>T p.T790M, associated with resistance resistance to anti-EGFR therapy]141Exon 20 (insertion/duplication, typically associated with resistance to anti-EGFR therapy)131Exon 21 [c.2573T>G p.L858R, associated with sensitivity anti-EGFR therapy]142		141
Exon 20 [c.2369C>T p.T790M, associated with resistance resistance to anti-EGFR therapy] 141   Exon 20 (insertion/duplication, typically associated with resistance to anti-EGFR therapy) 131   Exon 21 [c.2573T>G p.L858R, associated with sensitivity anti-EGFR therapy] 142	Exon 20 [c.2303G>T p.S768I, associated with low	
resistance to anti-EGFR therapy] 141   Exon 20 (insertion/duplication, typically associated with resistance to anti-EGFR therapy) 131   Exon 21 [c.2573T>G p.L858R, associated with sensitivity anti-EGFR therapy] 142	sensitivity to anti-EGFR therapy]	134
Exon 20 (insertion/duplication, typically associated with resistance to anti-EGFR therapy) 131   Exon 21 [c.2573T>G p.L858R, associated with sensitivity anti-EGFR therapy] 142	Exon 20 [c.2369C>T p.T790M, associated with resistance	
resistance to anti-EGFR therapy)131Exon 21 [c.2573T>G p.L858R, associated with sensitivity anti-EGFR therapy]142	resistance to anti-EGFR therapy]	141
Exon 21 [c.2573T>G p.L858R, associated with sensitivity anti-EGFR therapy] 142	Exon 20 (insertion/duplication, typically associated with	
anti-EGFR therapy] 142	resistance to anti-EGFR therapy)	131
	Exon 21 [c.2573T>G p.L858R, associated with sensitivity	
Exon 21 [c.2582T>A p.L861Q, associated with sensitivity	anti-EGFR therapy]	142
	Exon 21 [c.2582T>A p.L861Q, associated with sensitivity	
sensitivity to anti-EGFR therapy] 137	sensitivity to anti-EGFR therapy]	137
Any mutation in exons 18-21 21	Any mutation in exons 18-21	21
Other mutations, specify: 5	Other mutations, specify:	5

5. If your laboratory uses an FDA-cleared or approved companion diagnostic for <i>EGFR</i> , specify the manufacturer:	Freq (63)
Roche cobas EGFR Mutation Test	-
Roche cobas EGFR Mutation Test v2	50
Qiagen therascreen EGFR RGQ PCR Kit	13

9. If your laboratory-developed test uses primers or probes by a commercial vendor, specify:	Freq (148)
AmoyDx	2
Applied Biosystems	2
Biocartis Idylla	50
DxS EGFR mutation	-
EntroGen	4
Illumina	5
Integrated DNA Technologies	5
Qiagen PCR	-
Qiagen Pyro	-
Qiagen RGQ PCR	6
Roche	14
Sequenom Oncocarta	-
Thermo Scientific	4
TrimGen	1
Other, specify:	8
Not applicable (N/A)	47

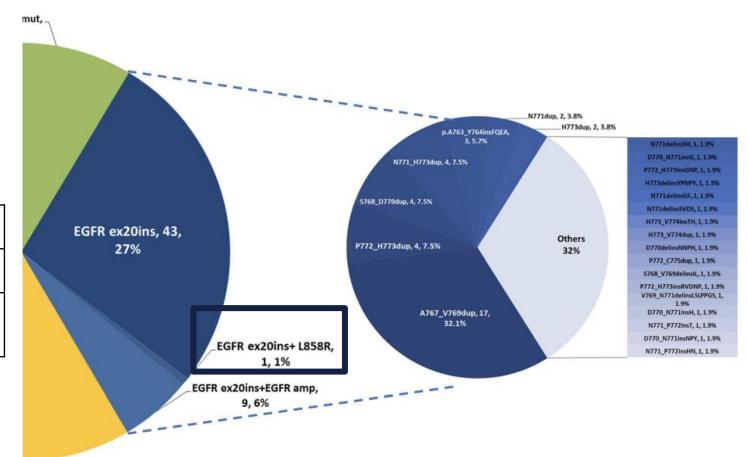
### **cobas® EGFR Mutation Test v2 – Potential for false positive results**

Updated 11/08/2022

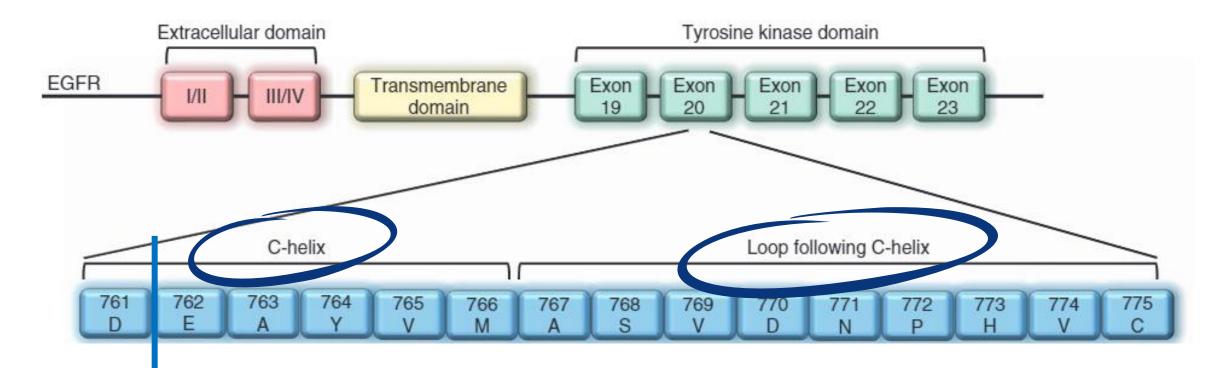
#### Device Indication for a Specific Group of Oncology Therapeutic Products

Diagnostic Name (Manufacturer)	Indication(s) - Sample Type	PMA (Approval Date)	Device Indication for a Specific Gr	roup of Oncology Therapeutic Products and Trade Name (Generic) – <u>NDA/BLA</u>
cobas EGFR Mutation Test v2 [Roche Molecular Systems, Inc.] Non-Small Cell Lung Cancer (NSCLC) - Tissue or Plasma P120019/S031 (10/27/2020) Non-small Cell lung cancer (lissue): "Identifying patients with NSCLC whose tumors have EGFR exon 19 deletions or exon 21 (L8588 mutations and are suitable for treatment with a tyrosime kinase inhibitor approved by FDA for this indication: Tarceva (erlotinib) - NDA 2021743 Tagrisso (osimertinib) - NDA 208065 Iressa (geftinib) - NDA 208065 Iressa (geftinib) - NDA 201292 Vizimpro (darcomitinib) - NDA 20129 Vizimpro (darcomitinib) - NDA 20129 Vizimpro (darcomitinib) - NDA 20129 Vizimpro (darcomitinib) - NDA 20129 Vizimpro (darcomiti			"Identifying patients with NSCLC who mutations and are suitable for treatm List of tyrosine kinase inhibitors appro Tarceva (erlotinib) - <u>NDA 021743</u> Tagrisso (osimertinib) - <u>NDA 208065</u> Iressa (gefitinib) - <u>NDA 206995</u> Gilotrif (afatinib) - <u>NDA 201292</u> Vizimpro (dacomitinib) - <u>NDA 211288</u>	nent with a tyro <mark>sine kinase inhibitor approved by FDA for that indication"</mark>
		tial for False "Mutation Detected" s customers using: Catalog Number		

SMC (2022)	Cobas	ddEGFR/Sanger	
E20ins only	50	33	
E20ins + L858R	26	0	
E20ins + E19del	36	U	

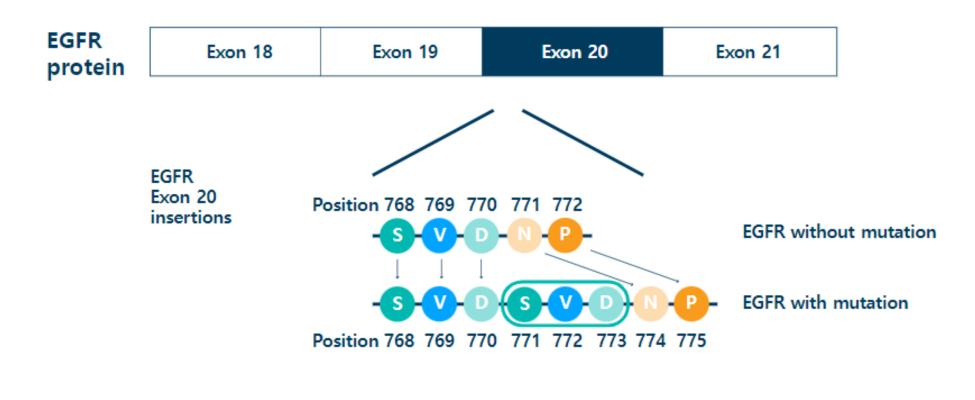


### HETEROGENEITY OF EGFR EXON 20 INSERTION MUTATIONS IN NSCLC



Over 100 different *EGFR* exon 20 insertion muts have been reported in NSCLC and the majority are composed of 1 to 4 AA insertions located in the loop following the C-helix

### EGFR EXON20 INSERTION NOMENCLATURE

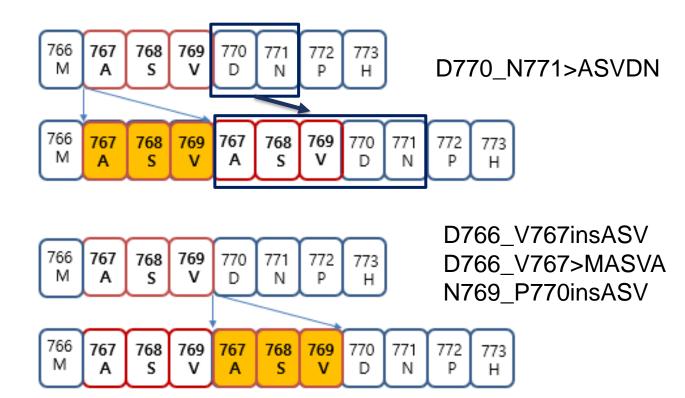


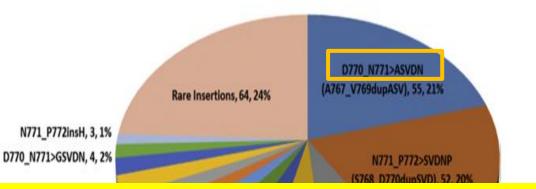
When SVD is newly inserted, how should it be named?

D770\_S771 ins SVD S767\_S768 ins SVD S768\_D770 dup

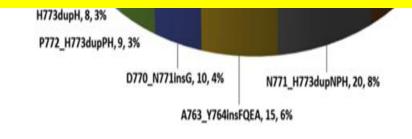
### HGVS nomenclature : 3' RULE

For all descriptions, the **most C-terminal position** possible of the reference sequence is arbitrarily assigned to have been changed





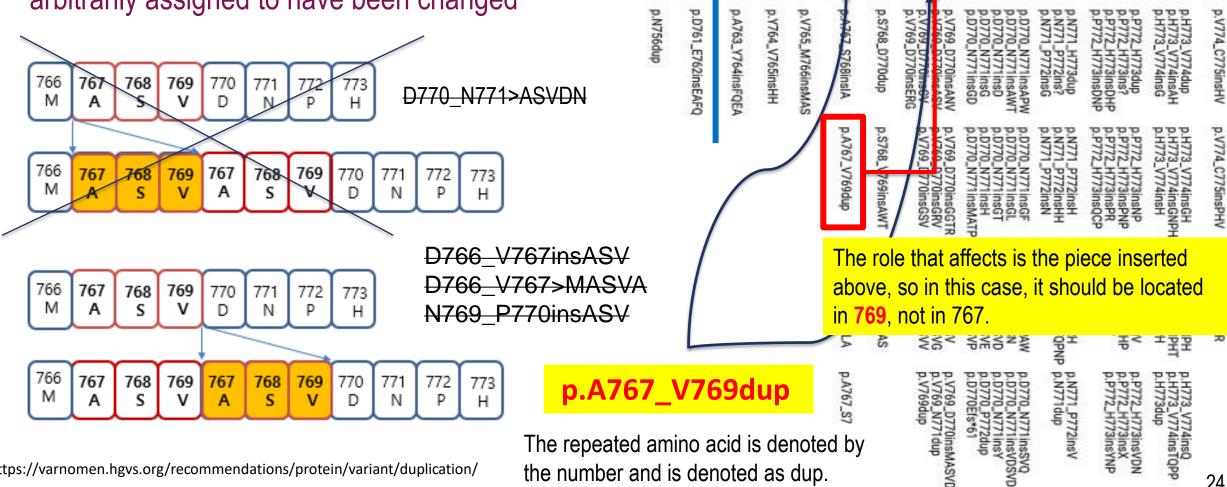
A protein's structure, however, is affected by those at the back. Therefore, in the case of duplication, it should be marked as inserted in the back, i.e., 3', not the front.



J Thorac Oncol. 2018 Oct;13(10):1560-1568

### **3' RULE**

For all descriptions, the most C-terminal **position** possible of the reference sequence is arbitrarily assigned to have been changed



https://varnomen.hgvs.org/recommendations/protein/variant/duplication/

the number and is denoted as dup.

Exon

22

Exon

23

Exon

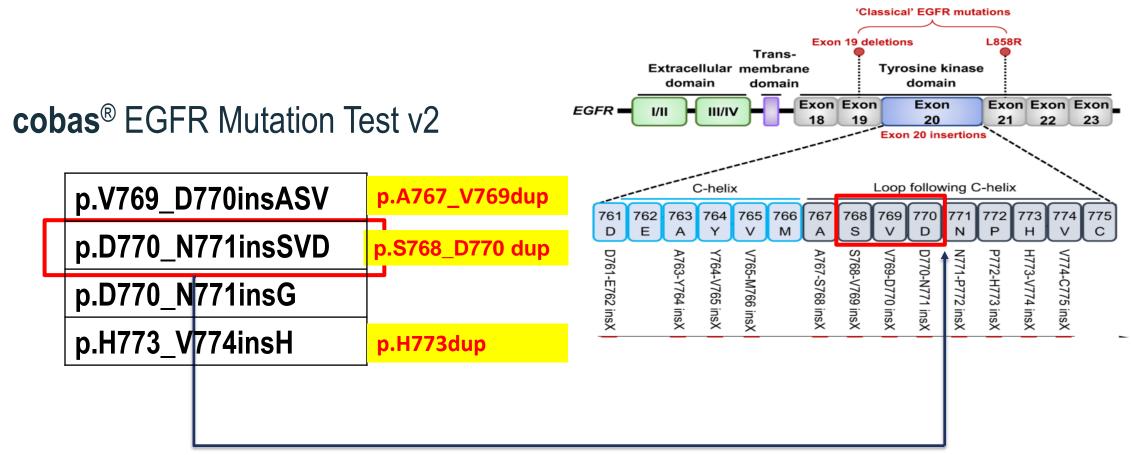
21

Exon 20

Exon Exon

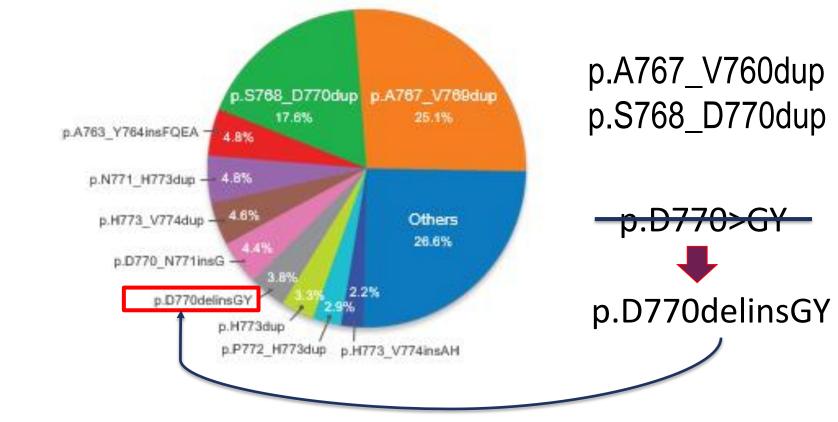
19

### EGFR EXON20INS

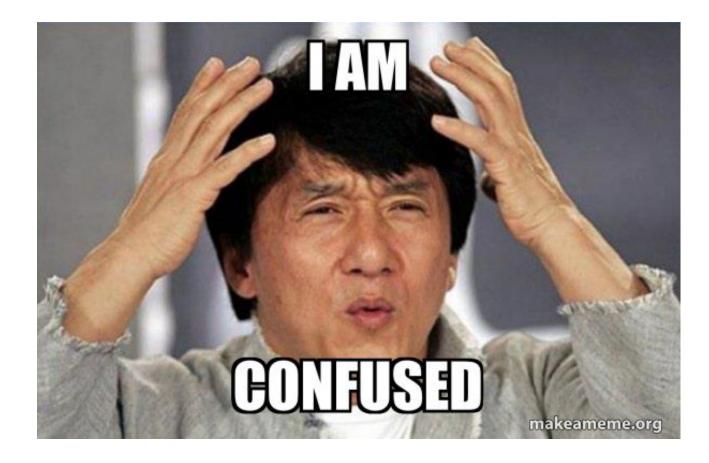


### EGFR exon20 insertion mutations are highly diverse

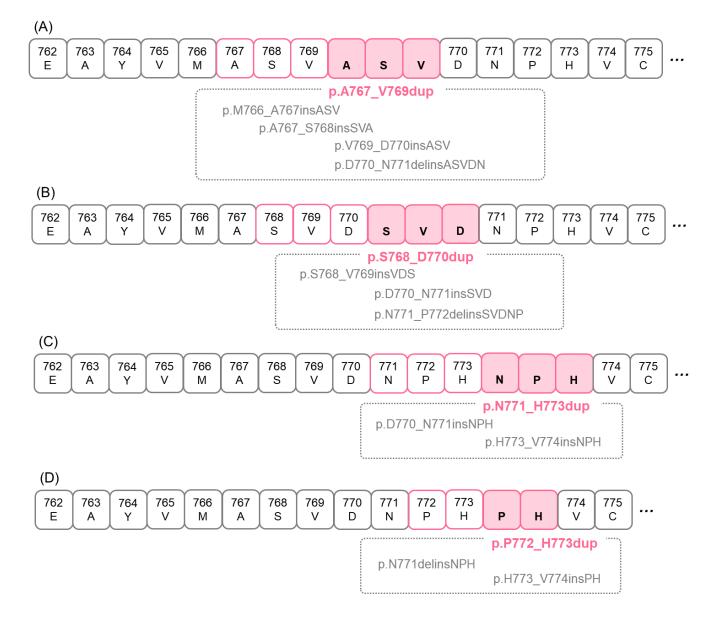
### **Frequencies of different EGFR ex20ins**



Out of 547 Chinese patients with NSCLC, a total of 85 unique EGFR exon20ins variants were identified

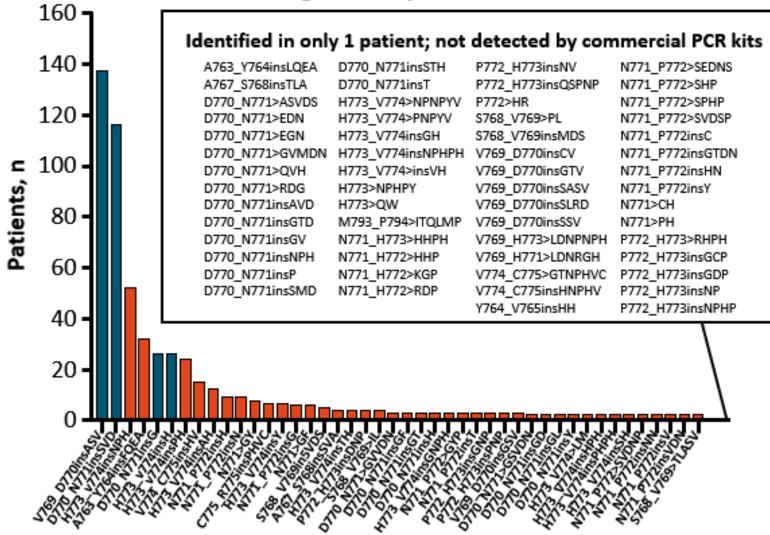


### HGVS nomenclature : 3' RULE



- Gray texts in a dotted box are mutations confusingly used in the literature, which should be reported according to the HGVS nomenclature (pink text) to avoid confusion.
- Most annotations that require correction according to the HGVS nomenclature were "duplications (dup)", which were incorrectly expressed as "insertion (ins)" or "deletion-insertion (delins)".

#### FoundationInsights<sup>®</sup>: Expected PCR Detection Results



#### Expected to be detected by

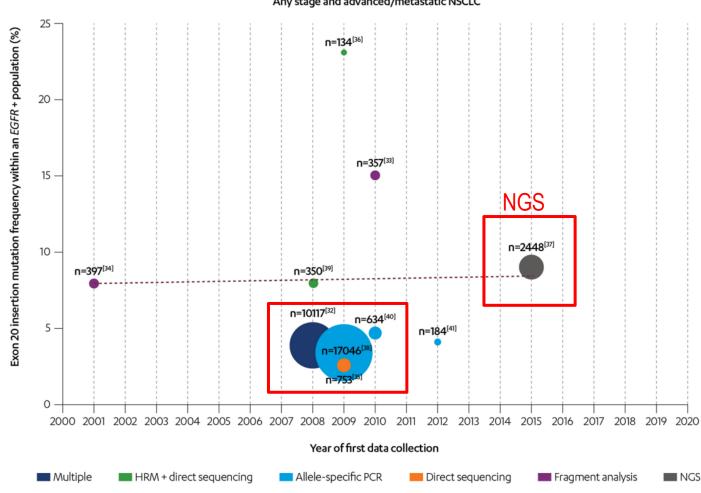
Cobas®, therascreen

Not detected by above kit

Method	Kit	
qRT-PCR	Cobas® EGFR Mutation Test v2	
	therascreen EGFR RGQ PCR	

### EGFR exon20 insertion : Frequency

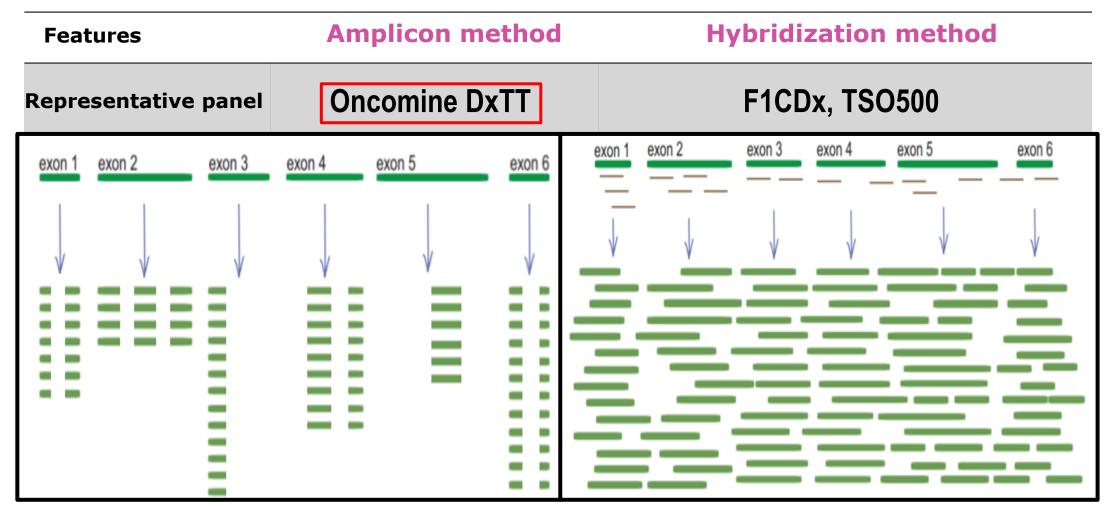
Frequency of exon 20 insertion mutations in any stage of EGFR-mutated NSCLC



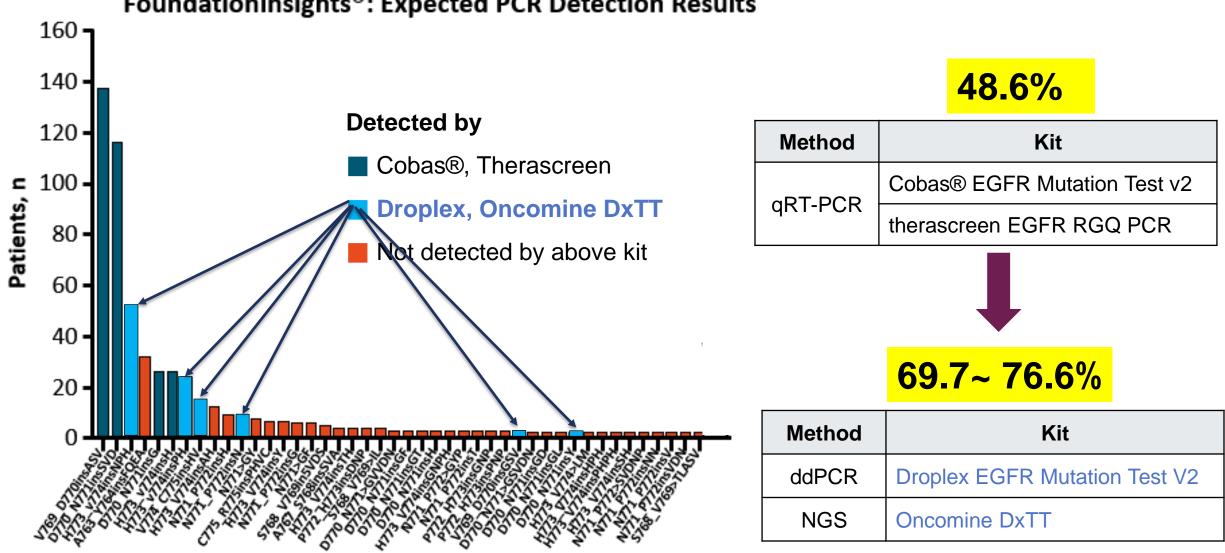
#### Any stage and advanced/metastatic NSCLC

- 1.6% in the overall NSCLC population •
- 9.0% in an EGFR-positive population ٠
- 2-fold increase in frequency when • NGS is applied.

### **Amplicon vs. Capture Sequencing**



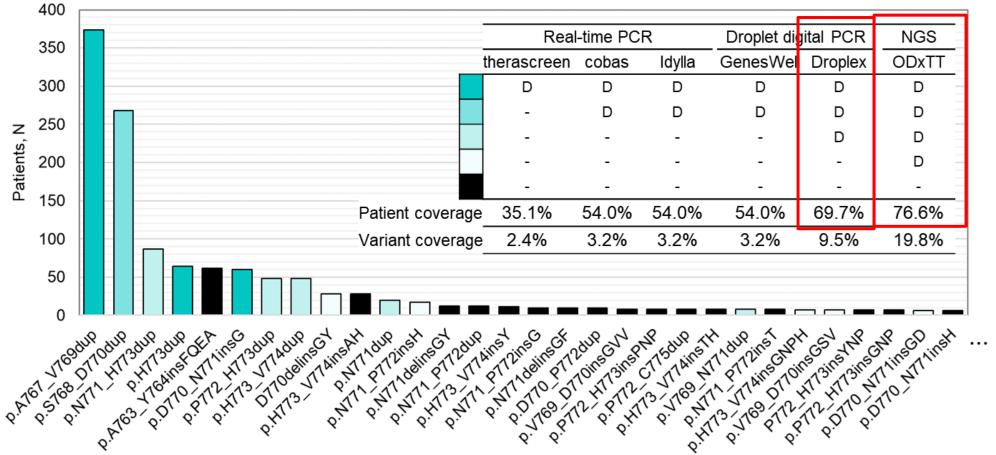
Yatabe Y, Sunami K, Goto K et al. Multiplex gene-panel testing for lung cancer patients. Pathol Int 2020;70;921-931.



#### FoundationInsights<sup>®</sup>: Expected PCR Detection Results

### Detection coverages of EGFR exon20 insertion mutations

Total 1,418 EGFR exon20 insertion patients



- Oncomine Dx Target Test : 76.6%
- Droplex EGFR Mutation Test v2 : 69.7%

Data from FoundationInsights database (n=625), Geneseeq Technology Inc. (n=517), Phase I/II trial of mobocertinib (n=95), Phase II study of poziotinib (n=47), phase I study of sunvozertinib (n=51), SMC (n=83)

### **Summary**

- Testing for *EGFR* is **widely available** throughout the world.
- More than half are PCR-based single gene tests, and NGS tests are growing.
- A PCR-based single gene test may miss 30 to 50% of EGFR exon20 insertions compared with other activating mutations.
- It is possible to have different levels of coverage of mutation variants on the same platform (including NGS).
- In some cases, the numbering and variable notation for EGFR exon20 insertion mutations do not comply with the guidelines, which can be confusing.

# KSP 2024

The 76<sup>th</sup> Annual Fall Meeting of the Korean Society of Pathologists

| The 1<sup>st</sup> International Conference of KSP |

#### SAVE THE DATE

Oct. 30 ~ Nov. 1, 2024 Lotte Hotel Seoul, Korea

www.pathology.or.kr/english



## Thank you



