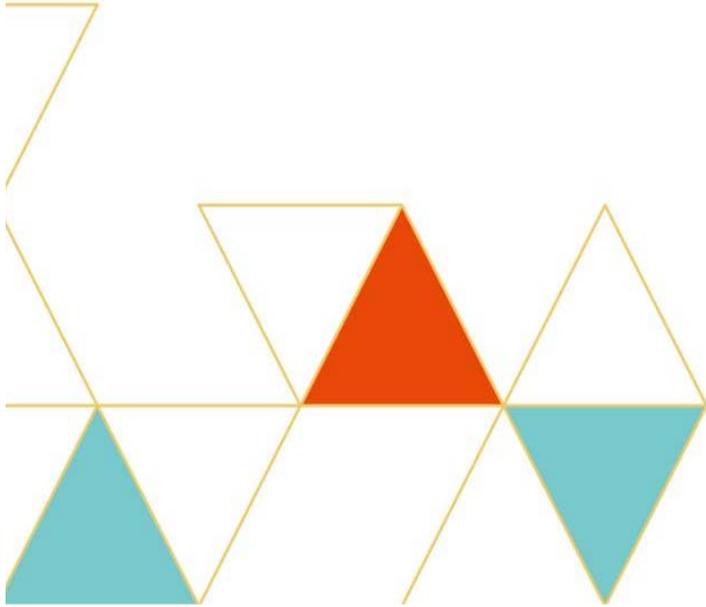


# FoundationOne<sup>®</sup> CDx Cancer Genomic Profile Product Overview



Kosuke Iijima

Department Manager of Foundation Medicine  
Business Department

Chugai Pharmaceutical Co., Ltd.

2019/07/04

# Transformational Shift in Oncology

2009

2019



Cancer is an anatomical disease



Cancer is a genomic disease



Clinical trials are conducted solely by academia or a pharmaceutical company



Collaborative clinical trials, basket/umbrella trials



Few targeted therapy trials



> 600 therapies in development, thousands of clinical trials



Significant decrease in use of immunostimulants



Rapid adoption of immunotherapies

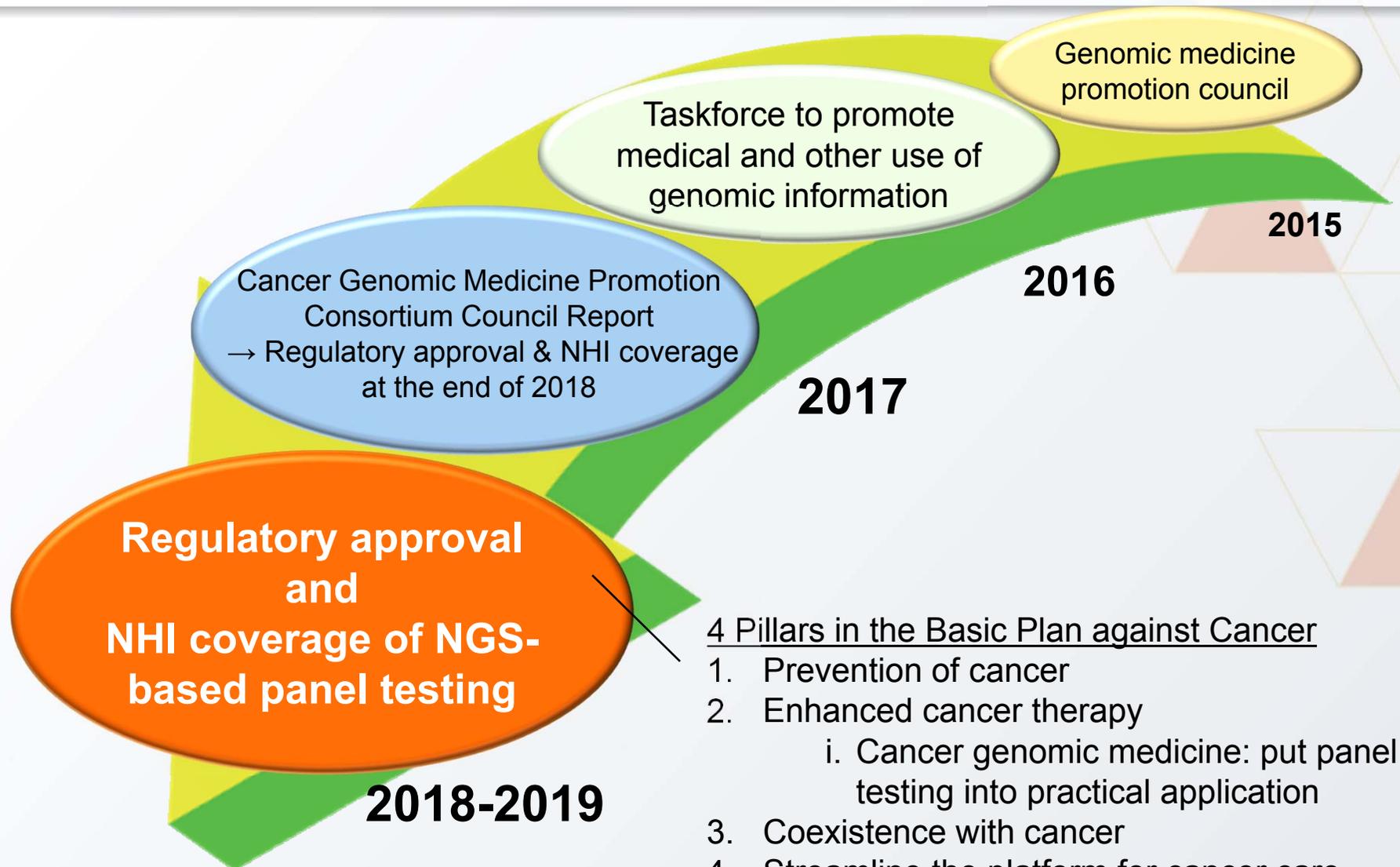


Disparate approaches to diagnostic assays



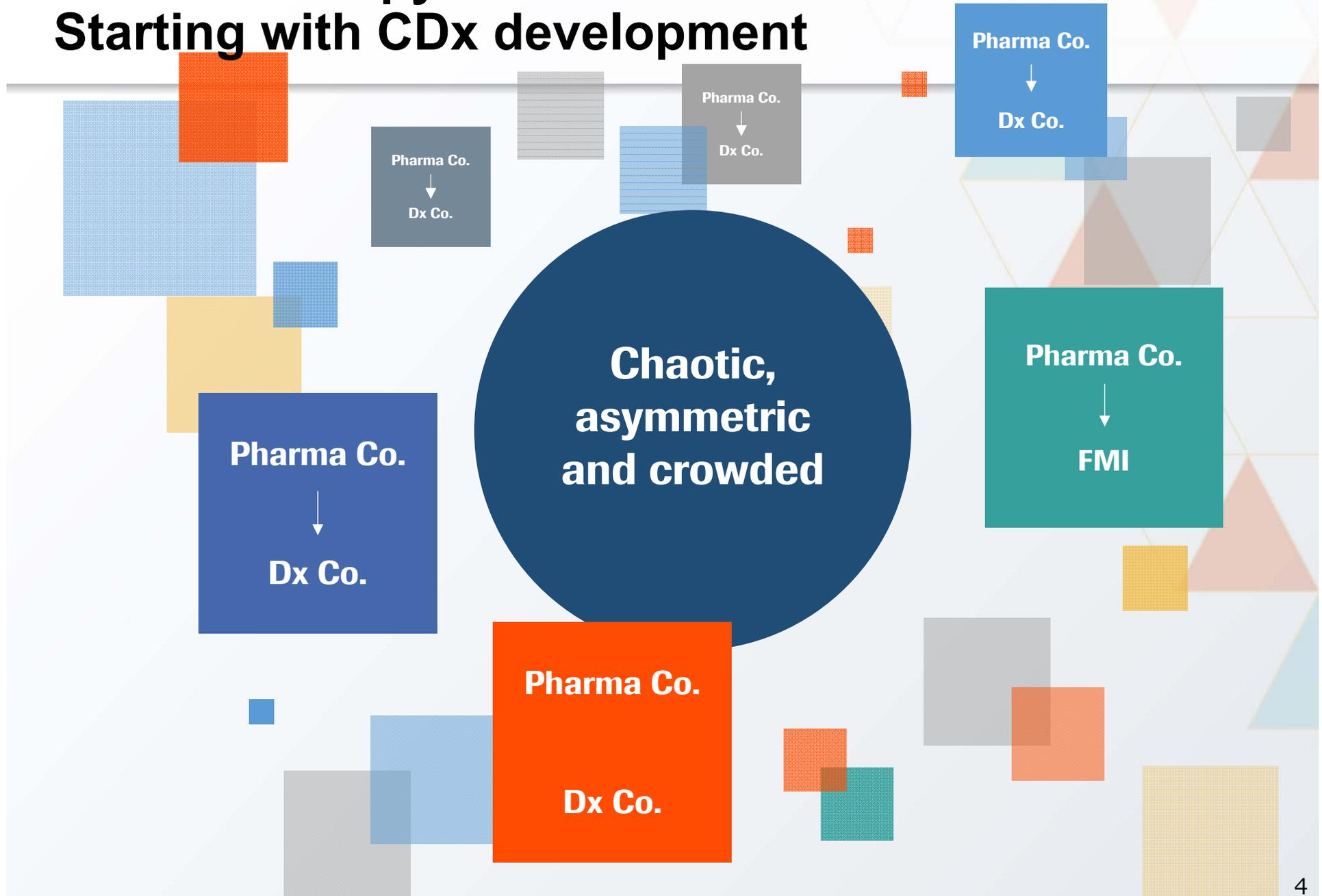
Emergence of comprehensive diagnostic assays

# Governmental Activities for Genomic Medicine

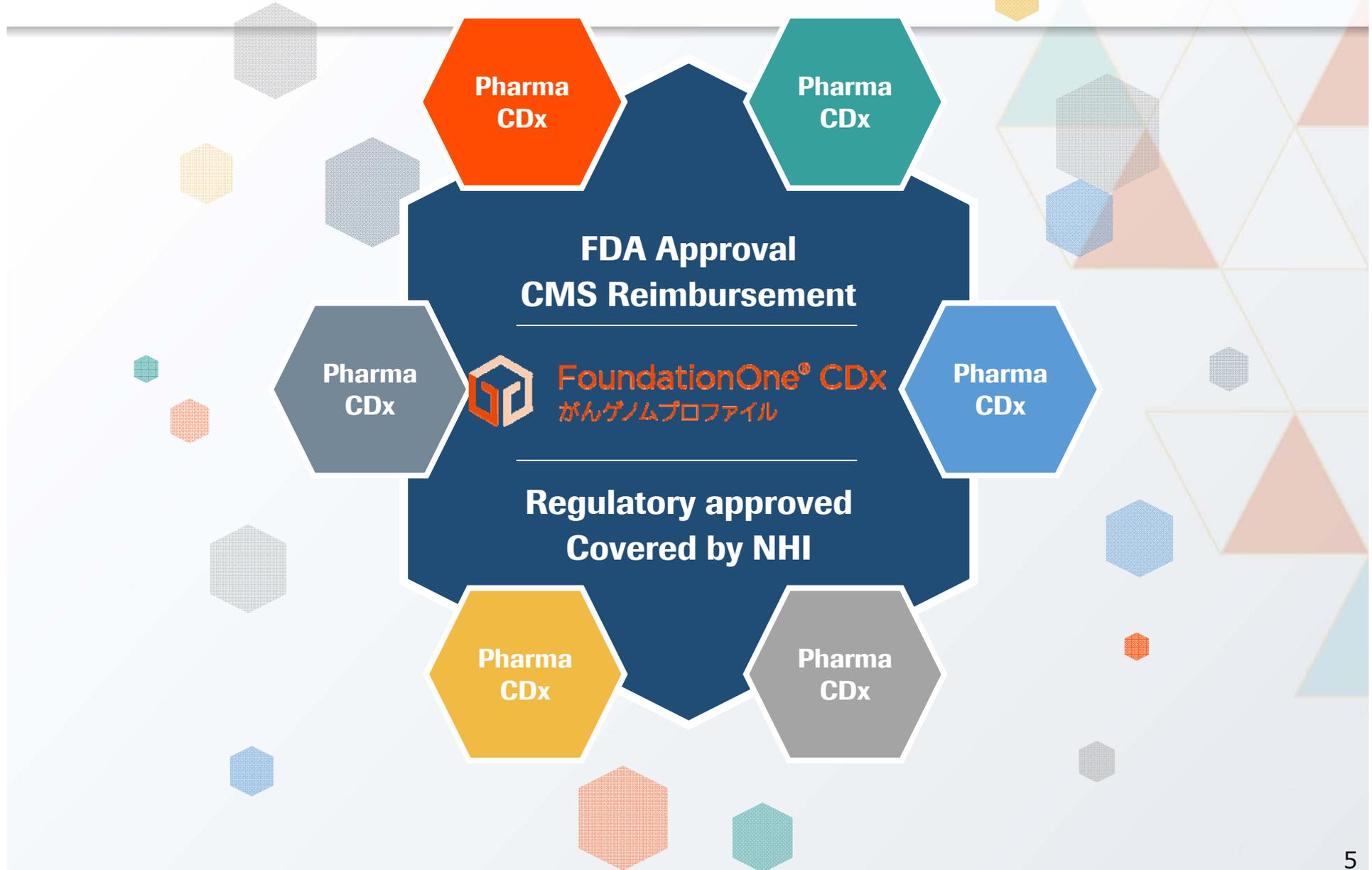


NGS: Next-generation sequencer  
NHI: National health insurance

# Chaotic therapy... Starting with CDx development



# Deliver Novel Platform Solution





FoundationOne® CDx

# Intended Usage or Indications

- Obtain comprehensive genome profiling with tumor tissues in patients with solid tumors
- Detect gene alterations to support the assessment of drug indications listed in the table below

**CGP**

**CDx**

Alterations	Cancer type	Relevant drugs
<i>EGFR</i> exon 19 deletions and <i>EGFR</i> exon 21 L858R alterations	Non-small cell lung cancer (NSCLC)	afatinib dimaleate, erlotinib hydrochloride, gefitinib, osimertinib mesylate
<i>EGFR</i> exon 20 T790M alterations		osimertinib mesylate
<i>ALK</i> fusion genes		alectinib hydrochloride, crizotinib, ceritinib
<i>BRAF</i> V600E and V600K alterations	Malignant melanoma	dabrafenib mesylate, trametinib dimethyl sulfoxide, vemurafenib
<i>ERBB2</i> copy number alterations (HER2 gene amplification positive)	Breast cancer	trastuzumab (genetical recombination)
<i>KRAS/NRAS</i> wild-type	Colorectal cancer	cetuximab (genetical recombination), panitumumab (genetical recombination)
<i>NTRK1/2/3</i> fusion gene	Solid tumors	entrectinib

# FoundationOne CDx Cancer Genomic Profiling



PATIENT  
Chugal Unique ID

TUMOR TYPE  
Lung adenocarcinoma

REPORT DATE  
01 Mar 2019

ORF#  
XXXXXXXX

**PATIENT**

DISEASE: Lung adenocarcinoma  
NAME: Not Given  
DATE OF BIRTH: Not Given  
SEX: Not Given  
MEDICAL RECORD #: Not Given

**PHYSICIAN**

ORDERING PHYSICIAN: Not Given  
MEDICAL FACILITY: Not Given  
ADDITIONAL REPORT#: Not Given  
MEDICAL FACILITY ID: Not Given  
PATHOLOGIST: Not Given

**SPECIMEN**

SPECIMEN SITE: Not Given  
SPECIMEN ID: Not Given  
SPECIMEN TYPE: Not Given  
DATE OF COLLECTION: Not Given  
SPECIMEN RECEIVED: Not Given

**Companion Diagnostic (CDx) Associated Findings**

GENOMIC FINDINGS DETECTED	APPROVED THERAPEUTIC OPTIONS IN JAPAN
<b>EGFR</b> L858R	Afatinib maleate Erlotinib hydrochloride Gefitinib Osimertinib mesilate

TUMOR TYPE  
Lung adenocarcinoma

REPORT DATE  
01 Mar 2019

ORF#  
XXXXXXXX

**Findings**

status - MS-Stable  
Tumor Mutational Burden - TMB-Intermediate (11 Muts/Mb)

**Findings**

If genes analyzed, please refer to the Appendix on L858R

Genes with no reportable alterations: KRAS, ALK, ERBB2, ROS1

Clinical Benefit: 18 Clinical Trials  
Lack of Response

**OTHER ALTERATIONS & BIOMARKERS IDENTIFIED**

Results reported in this section are not prescriptive or conclusive for labeled use of any specific therapeutic product. See professional services section for additional information.

Microsatellite Status: MS-Stable<sup>§</sup>  
Tumor Mutational Burden: 11 Muts/Mb<sup>§</sup>  
CDKN2A/B loss<sup>§</sup>  
EGFR amplification<sup>§</sup>

PTCH1 T416S  
RBM10 Q494\*  
TP53 R267P

<sup>§</sup> Refer to appendix for limitation statements related to detection of any copy number alterations, gene rearrangements, MSI or TMB result in this section. Please refer to appendix for Explanation of Clinical Significance Classification and for variants of unknown significance (VUS).

Background information on the patient, the medical facility etc.

Summary of detected alterations

- Approval status of corresponding targeted therapies
- Ongoing clinical trials targeting detected alterations

BIOMARKER FINDINGS	THERAPIES WITH CLINICAL BENEFIT (IN PATIENT'S TUMOR TYPE)	THERAPIES WITH CLINICAL BENEFIT (IN OTHER TUMOR TYPE)
<b>Tumor Mutational Burden -</b> TMB-Intermediate (11 Muts/Mb)	Atezolizumab Durvalumab Nivolumab Pembrolizumab	Avelumab Cemiplimab-rwc
9 Trials see p. 14		
<b>Microsatellite status -</b> MS-Stable	No therapies or clinical trials. see Biomarker Findings section	
GENOMIC FINDINGS	THERAPIES WITH CLINICAL BENEFIT (IN PATIENT'S TUMOR TYPE)	THERAPIES WITH CLINICAL BENEFIT (IN OTHER TUMOR TYPE)
<b>EGFR -</b> amplification, L858R	Afatinib Dacomitinib Erlotinib Gefitinib Osimertinib	Cetuximab Lapatinib Panitumumab
4 Trials see p. 16		
<b>PTCH1 -</b> T416S	none	Sonidegib Vismodegib
5 Trials see p. 17		

Summary of references on detected alterations and potential therapies

# From Organs to Gene Alterations/Biomarkers

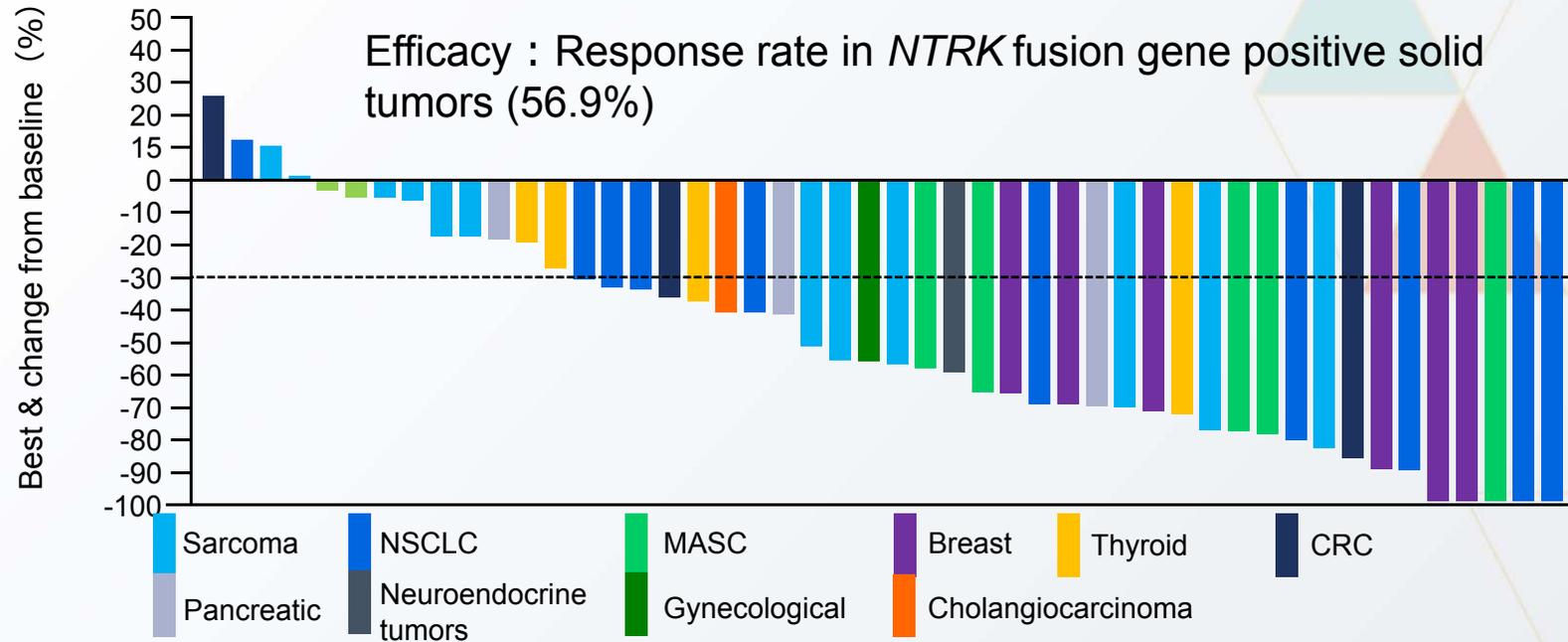
*Tumor agnostic approval + New histology independent medicines*

The screenshot shows the FDA website's 'Drugs' section. The main headline, highlighted with a red box, reads: 'FDA grants accelerated approval to pembrolizumab for first tissue/site agnostic indication'. Below the headline are social media sharing buttons for Facebook, Twitter, LinkedIn, Pinterest, Email, and Print. A link to a podcast is provided: 'Listen to the FDA D.I.S.C.O. podcast about this approval'. The article text states: 'On May 23, 2017, the U.S. Food and Drug Administration granted accelerated approval to pembrolizumab (KEYTRUDA, Merck & Co.) for adult and pediatric patients with unresectable or metastatic, microsatellite instability-high (MSI-H) or mismatch repair deficient (dMMR) solid tumors that have progressed following prior treatment and who have no satisfactory alternative treatment options or with MSI-H or dMMR colorectal cancer that has progressed following treatment with a fluoropyrimidine, oxaliplatin, and irinotecan.' A concluding sentence reads: 'This is the FDA's first tissue/site-agnostic approval.'

<https://www.fda.gov/drugs/informationondrugs/approveddrugs/ucm560040.htm>

Annals of Oncology, Volume 29, Issue suppl\_8

# Rozlytrek: The Second Drug in Japan Approved Across All Solid Tumors



## Study design

- Phase II, global, multicenter, open-label basket study (STARTRK-2)
  - Target: People aged 18 older with ***NTRK* fusion-positive\*** metastatic or relapse solid tumors. **N=51 including 1 Japanese.**
  - Administer oral entrectinib 600 mg/day
  - Primary endpoint: objective response rate (Time Frame: approx. 24 months)
- \* *NTRK* fusion-positive status was determined with a nucleic acid-based diagnostic test. **FoundationOne CDx Cancer Genomic Profile**, confirmed to be equivalent to this diagnostic test, is marketed as a companion diagnostic.

Source: Roche's Virtual Pipeline Event from ESMO 2018

## Outcomes

- **ORR was 56.9%** (95% CI: 42.3%–70.7%) by INV assessment per RECIST v1.1

## Clinically Significant Adverse Reactions

- 1 Cardiac disorders (4.8%) Cardiac disorders such as cardiac failure, ventricular extrasystoles, and myocarditis may occur. 2 Prolonged QT interval (frequency unknown) 3 Cognitive disorder, ataxia (28.6%) Cognitive disorder, confusional state, mental status changes, memory impairment, hallucinations, ataxia, dysarthria, etc., may occur. 4 Interstitial lung disease (1.6%)

Source: package insert

# CGP: Comprehensive Genomic Profiling

## Which position of usage benefits the patient?

- Finding matched therapy and selecting optimal treatment
- Understanding pathogenesis from clonal evolution
- Understanding pathogenesis including passenger alterations, not only searching for driver alterations
- Understanding pathogenesis from multiple genetic alterations
- Comprehensive treatment plan from biomarkers (TMB, MSI, LOH), etc.

### ASCO2019 CSS: Targeting Breast Cancer: Breaking the Code

*TP53*: Poor prognostic factor but promotes early progression of CDK4/6i

*FGFR1*: Promotes early progression of CDK4/6i

*RB1*: Possible CDK4/6i resistant mutation

*PTEN loss*: Lowers IC50 of CDK4/6i to promote tolerization

PD expected soon after administration of palbociclib

### Earlier treatment plan from CGP

Tolerization by enhancement during administration of palbociclib

### Monitoring of ctDNA CGP

TMB: Tumor Mutational Burden, MSI: MicroSatellite Instability,  
LOH: Loss of Heterozygosity

# Positioning of NGS Panel Test Directly Relates to Use and Quality of Clinico-Genomic DB

 **flatiron**  
Real-world, longitudinal patient-level clinical data from Electronic Health Records (EHRs) from cancer clinics



 **FOUNDATION MEDICINE®**  
Deep NGS profiling across hundreds of cancer-related genes for each patient's tumor



>42,000 patients (as of 2019.2)

## ■ Filing with FDA

- U.S. FDA APPROVES IBRANCE® (PALBOCICLIB) FOR THE TREATMENT OF MEN WITH HR+, HER2- METASTATIC BREAST CANCER

- ✓ Approval of expanded indication based predominately on real-world data

“.... The approval is based on data from electronic health records and postmarketing reports of the real-world use of IBRANCE in male patients sourced from three databases: IQVIA Insurance database, Flatiron Health Breast Cancer database and the Pfizer global safety database. ....”

Access: 28 June, 2019.

Note) IBRANCE for male with HR+, HER2- breast cancer is NOT approved in Japan

[https://www.pfizer.com/news/press-release/press-release-detail/u\\_s\\_fda\\_approves\\_ibrance\\_palbociclib\\_for\\_the\\_treatment\\_of\\_men\\_with\\_hr\\_her2\\_metastatic\\_breast\\_cancer](https://www.pfizer.com/news/press-release/press-release-detail/u_s_fda_approves_ibrance_palbociclib_for_the_treatment_of_men_with_hr_her2_metastatic_breast_cancer) 11

# Cancer Precision Medicine

**Transform genomic medicine  
into true precision medicine**

## **Personalized Health Care**

Fit one target to  
One Organ  
e.g. Targeted therapies

## **Genome Medicine**

Fit one target to  
Multiple Organs  
e.g. Comprehensive  
Universal diagnostics

## **Precision Medicine**

Focus on  
Individual patients  
e.g. Insight and Decision  
for patients based on  
Mixed genome and  
clinical info

# Cancer Genomic Medicine

## — Clinical Implementation and Challenges —

Manabu Muto, M.D., Ph.D., Professor

Therapeutic Oncology Course, Kyoto University Graduate School of Medicine  
Department of Clinical Oncology, Kyoto University Hospital



# COI Disclosure

Presenter: Manabu Muto

The presenter has the following conflict of interests relating to this presentation.

<u>COI description</u>	<u>Company name, etc.</u>
Directorial/advisory roles:	None
Presentation fees:	None
Manuscript fees:	None
Joint research fees:	Mitsui Knowledge Industry, Sysmex Corporation, Riken Genesis

# Agenda

1. Cancer and Genetics
2. Cancer Genomic Medicine in Healthcare Settings
3. Cancer Genomic Medicine in Current Clinical Practice
4. Future Challenges in Cancer Genomic Medicine

# **Cancer Genomic Medicine**

Several years from now,  
cancer genomic medicine will likely  
become standard practice.

# ***Precision Medicine***

An emerging approach for disease treatment and prevention that takes into account individual variability in genes, environment, and lifestyle for each person.

Source: NIH website (as of June 25, 2019)

<https://ghr.nlm.nih.gov/primer/precisionmedicine/definition>

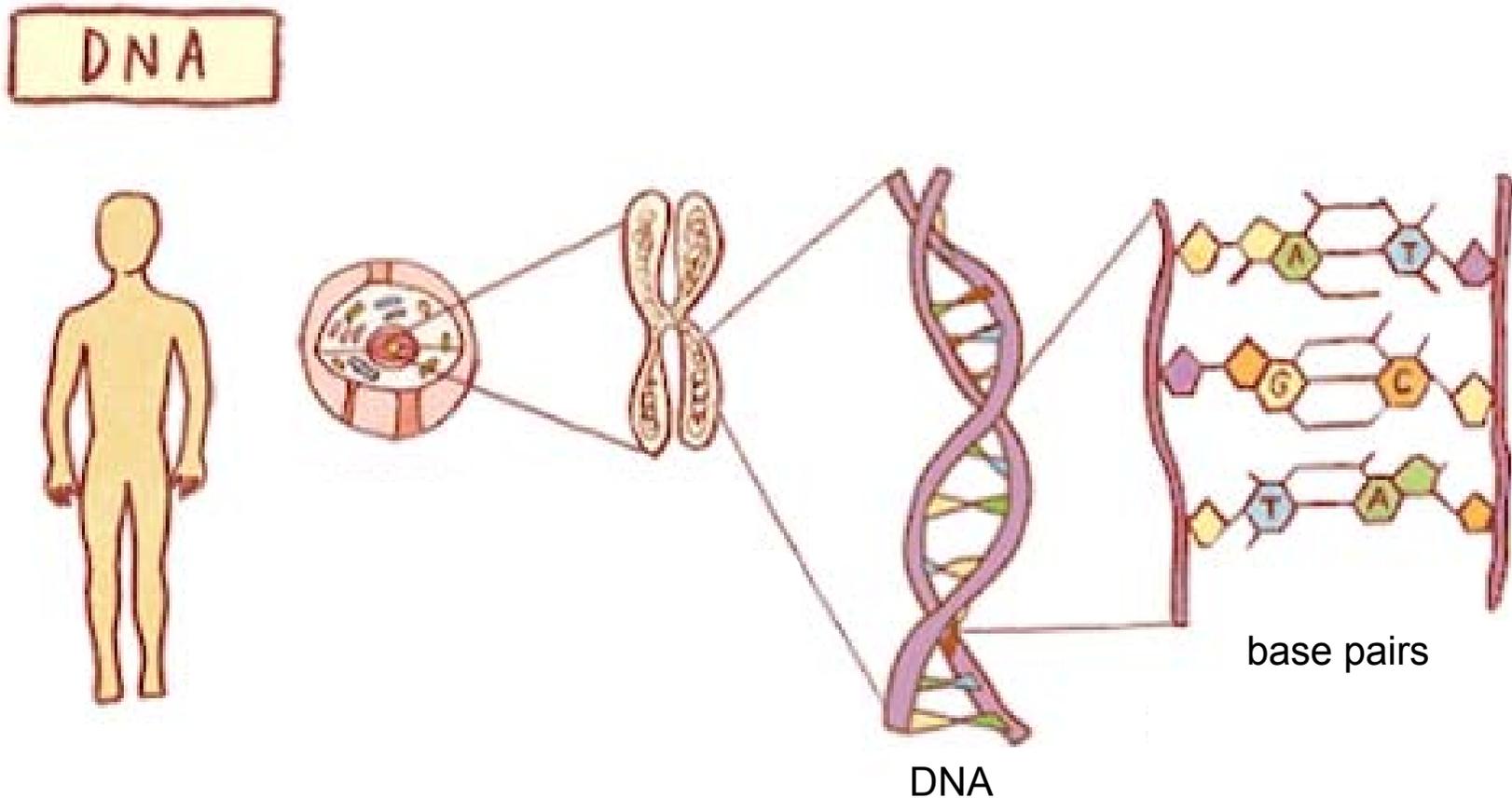
## ***Cancer Genomic Medicine***

*Precision medicine in the diagnosis and treatment of cancer*

# Cancer

Cancer is a disease caused by  
genetic abnormalities

# About DNA



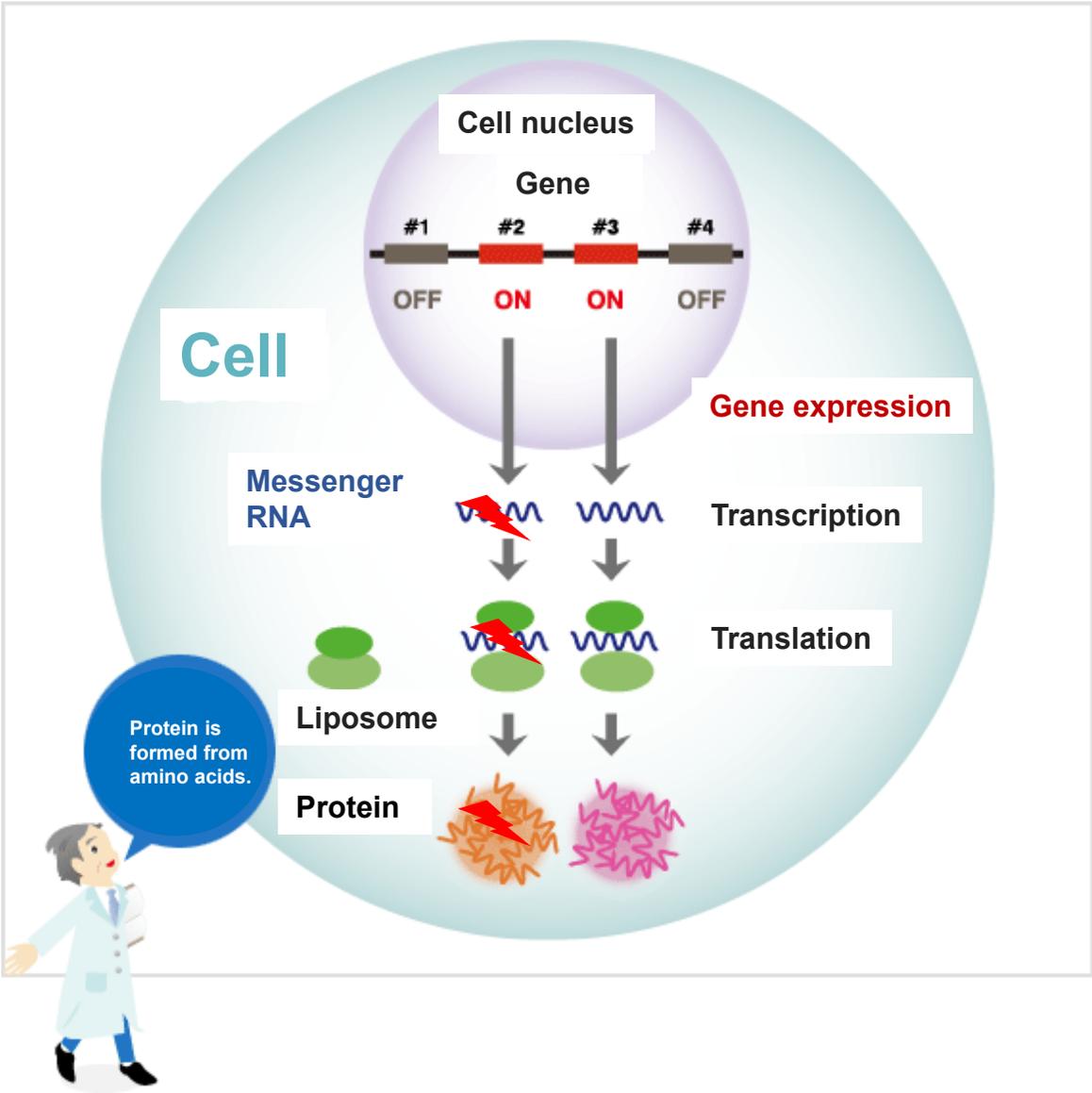
DNA is a biological molecule that regulates expression and transfer of genetic information

In almost all living things, DNA is the carrier and is the genetic information encoded by its base sequence.

Structure and base of DNA

source : *Easy biotechnology and genome*, Chugai Pharmaceutical website (Japanese only, as of June 28, 2019)  
<https://www.chugai-pharm.co.jp/ptn/bio/genome/genomep08.html>

# Protein Synthesis from Gene Transcription



# Gene Sequence Abnormalities

Baseline sequence

Normal

5' – CTA GCC CAA TTA CAT -3'

# Gene Sequence Abnormalities

Baseline sequence

Normal            5' – CTA GCC CAA TTA CAT -3'

Point mutation    5' – CTA ACC CAA TTA CAT -3'

# Gene Sequence Abnormalities

Baseline sequence

Normal

5' – CTA GCC CAA TTA CAT -3'

Deletion

5' – CTA --- CAA TTA CAT -3'

# Gene Sequence Abnormalities

Baseline sequence

Normal

5' – CTA GCC CAA TTA CAT -3'

Insertion

5' – CTA GCC **G**CA ATT ACA T-3'

# Gene Sequence Abnormalities

Baseline sequence

Normal

5' – CTA GCC CAA TTA CAT -3'

Amplification

5' – CTA **GCC GCC** CAA TTA CAT -3'

# Gene Sequence Abnormalities

Baseline sequence

Normal            5' – CTA GCC CAA TTA CAT -3'

Point mutation    5' – CTA **A**CC CAA TTA CAT -3'

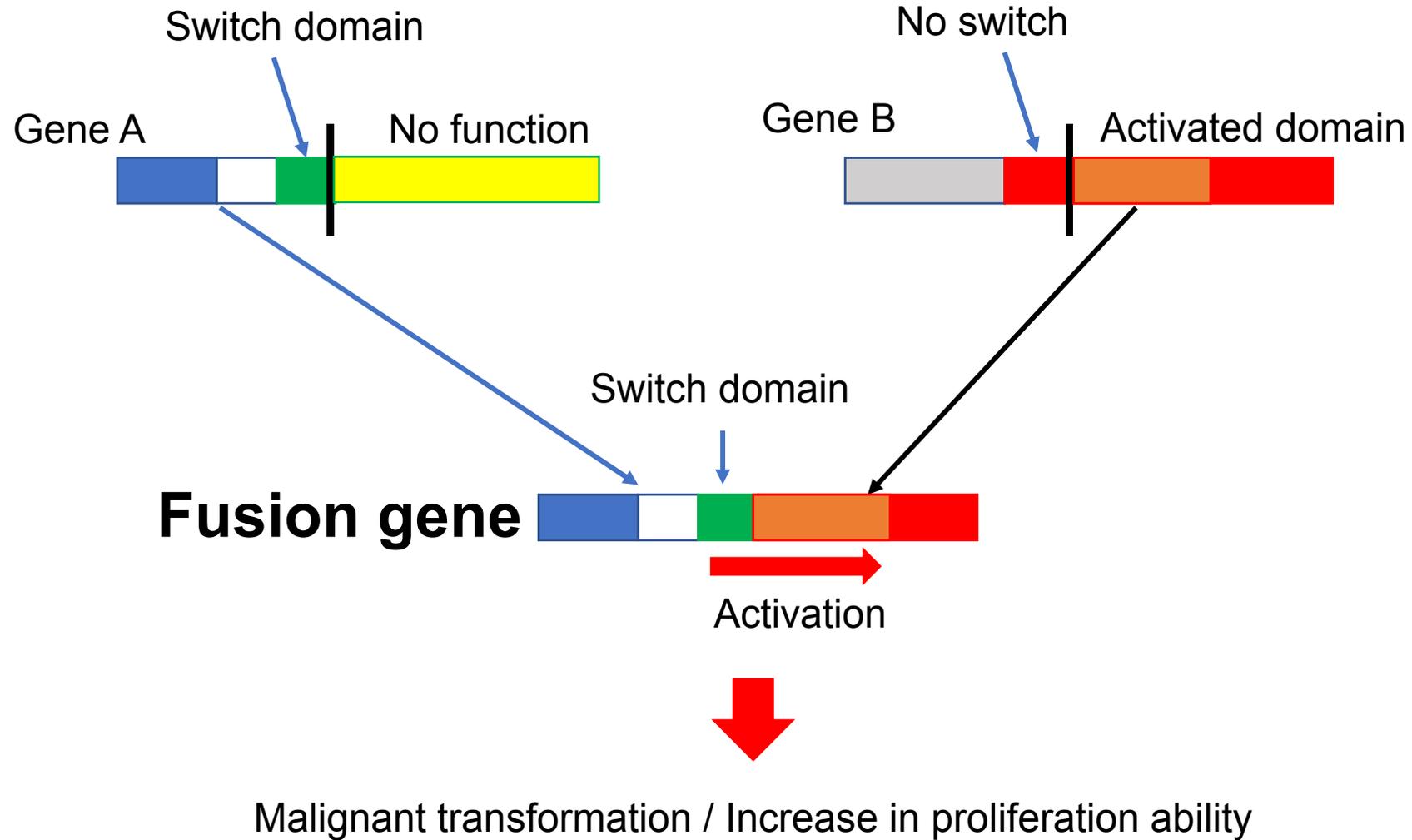
Deletion            5' – CTA --- CAA TTA CAT -3'

Insertion            5' – CTA GCC **G**CA ATT ACA T-3'

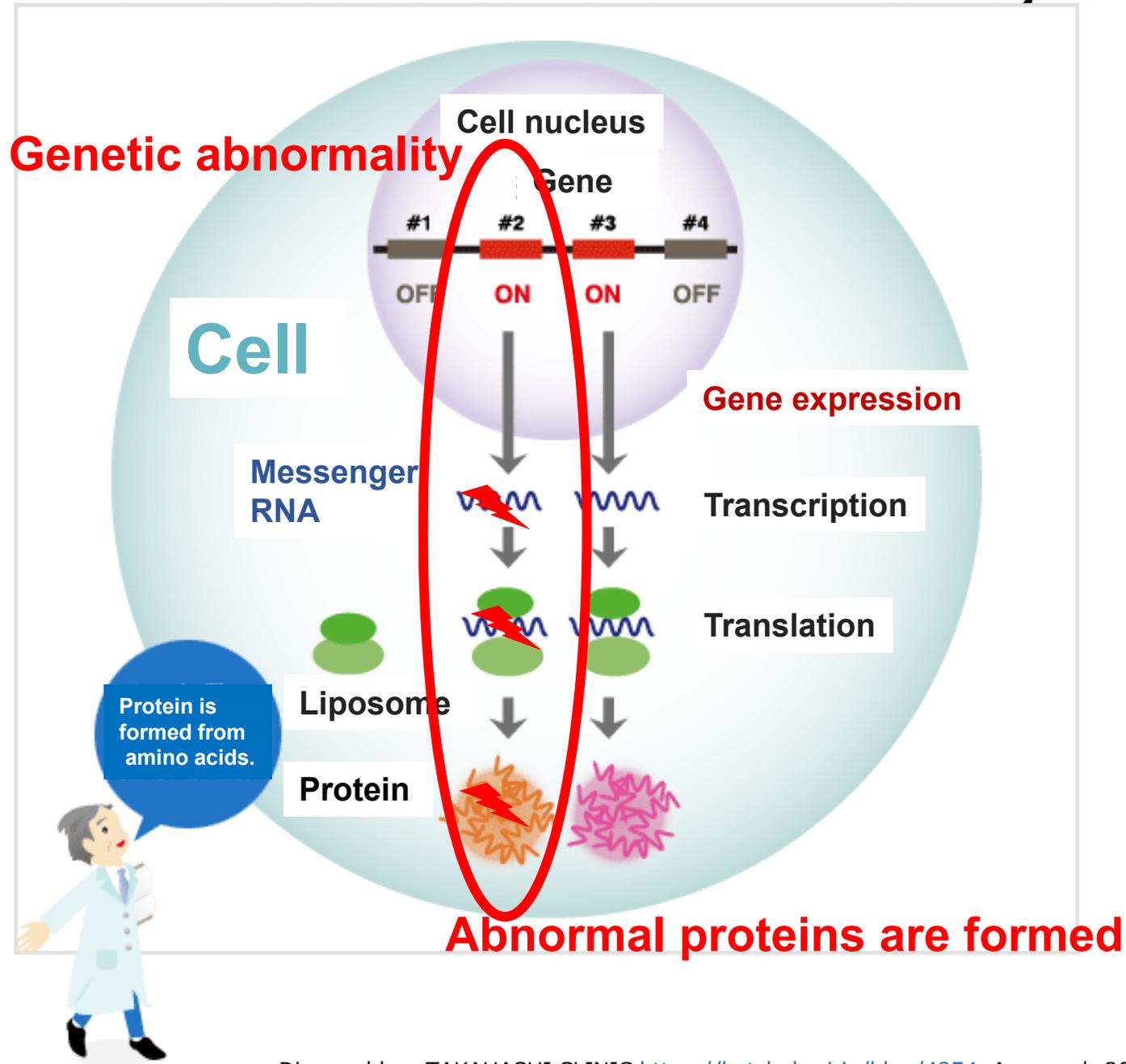
Amplification      5' – CTA **GCC GCC** CAA TTA CAT -3'

# Abnormal Structure of Gene

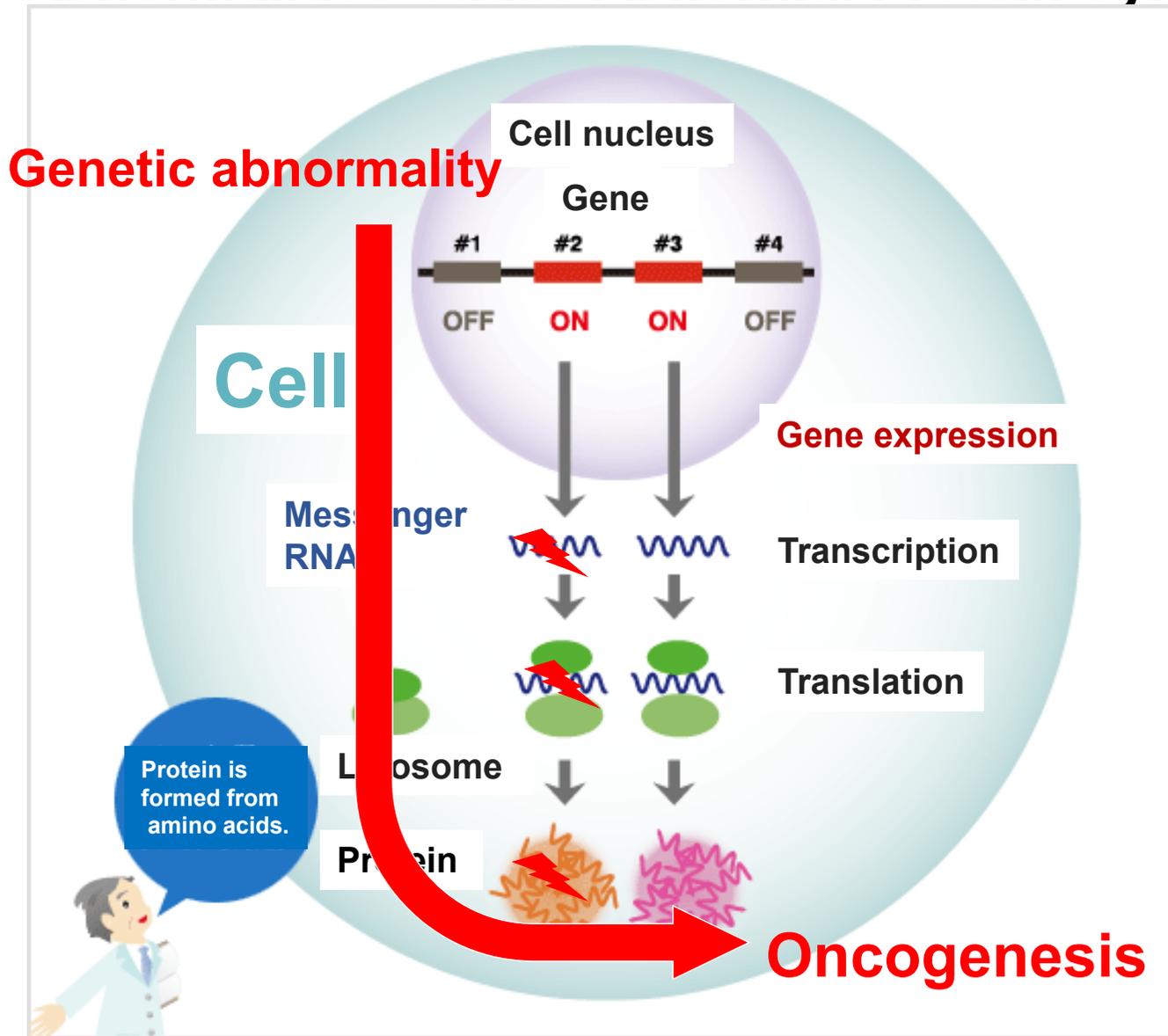
[Conceptual illustration]



# Genetic Abnormalities Cause Abnormal Protein Synthesis

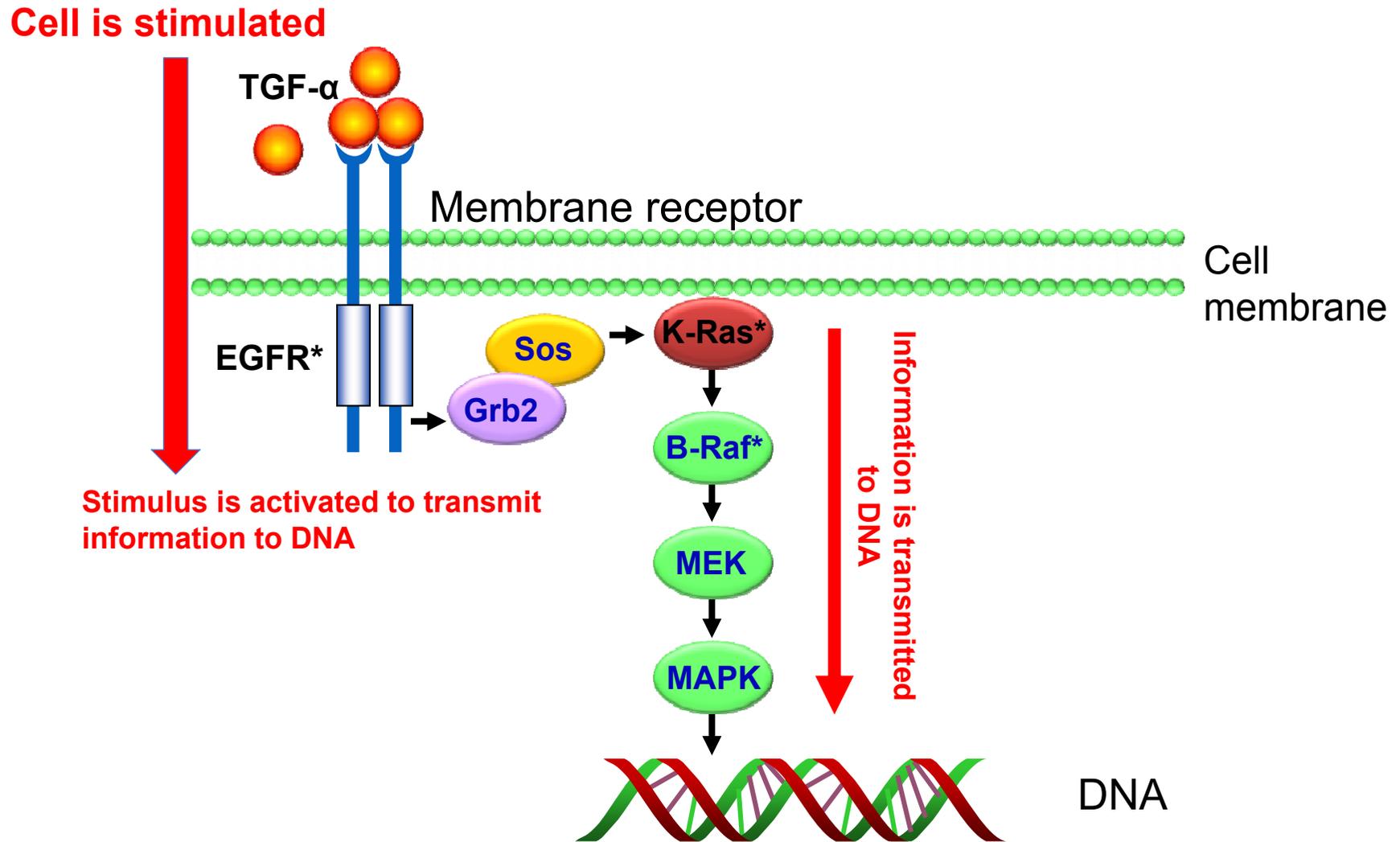


# Genetic Abnormalities Cause Abnormal Protein Synthesis



**Abnormal proteins may induce oncogenesis**

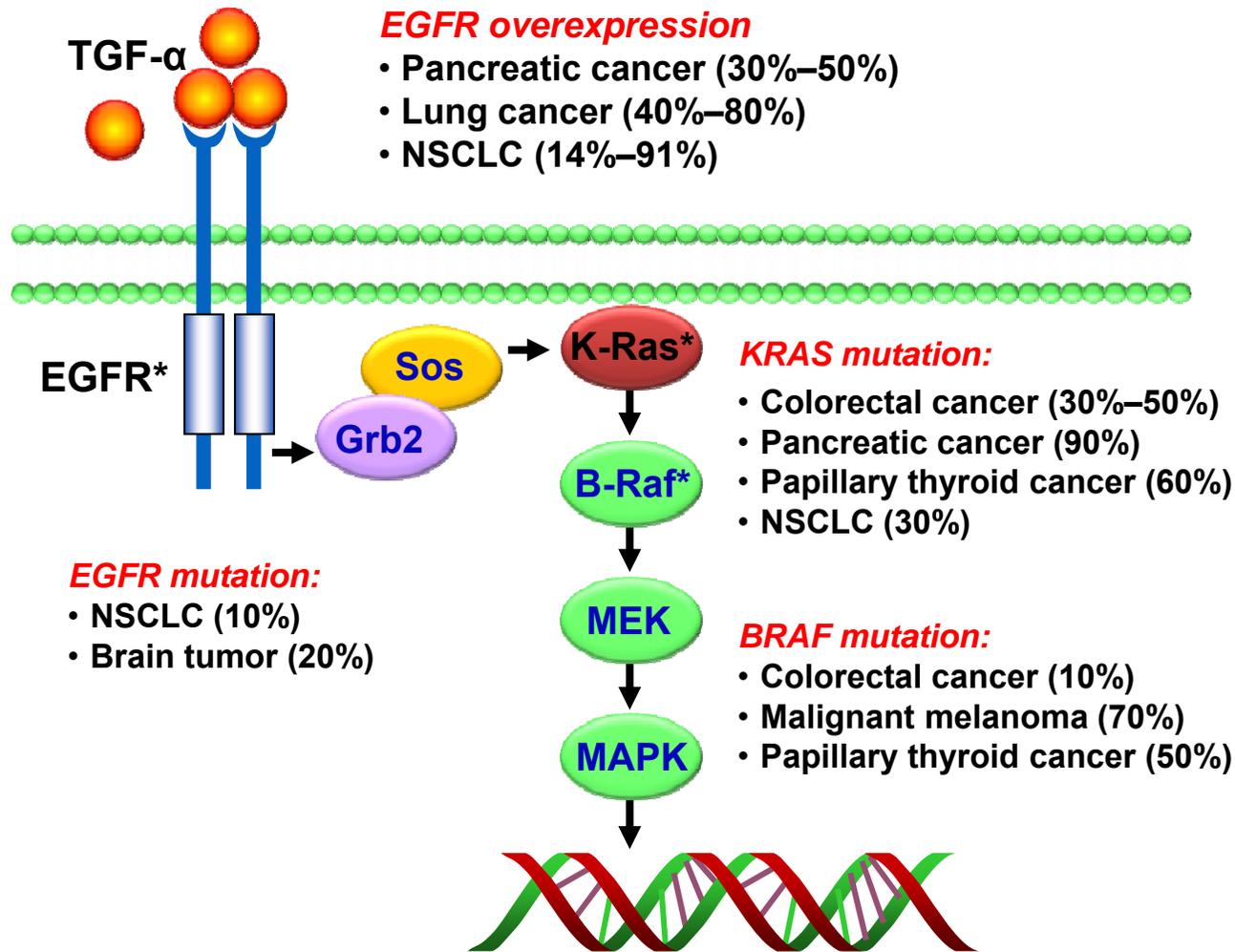
# Example: Cell Signaling



Targeting the Raf-MEK-ERK mitogen-activated protein kinase cascade for the treatment of cancer.

Roberts PJ, Der CJ., Oncogene 2007

# Genetic Abnormalities Cause Abnormal Cell Signaling = Oncogenesis



NSCLC: Non-small cell lung cancer

Targeting the Raf-MEK-ERK mitogen-activated protein kinase cascade for the treatment of cancer.

Roberts PJ, Der CJ., Oncogene 2007

**Gene (genome) abnormalities  
cause cancer**



**Gene (genome) abnormalities need  
to be examined**

# **An Example of Personalized Medicine**

Testing of cancer gene abnormalities  
in individual patients to explore  
therapeutic options

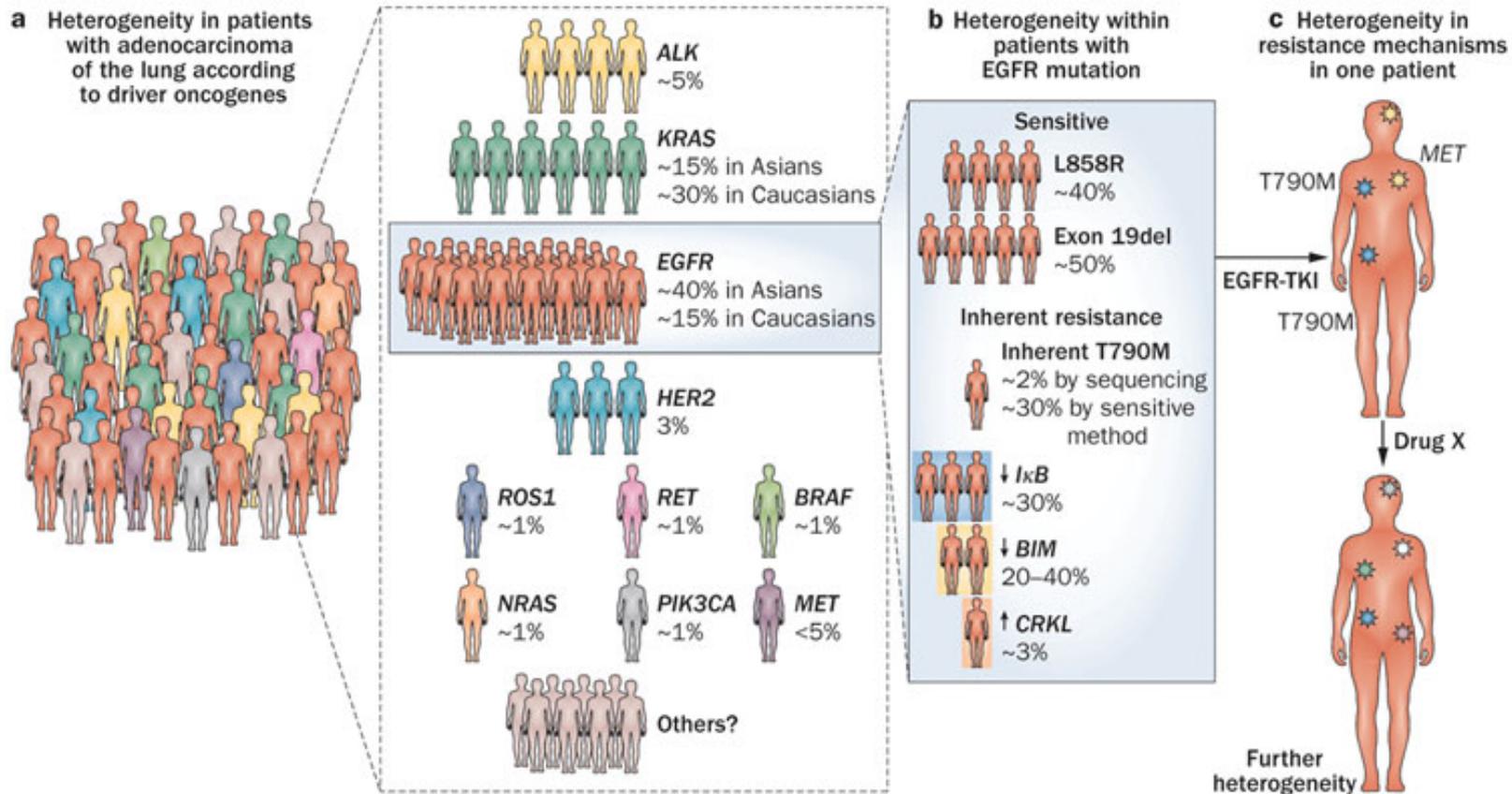
# Testing Methods for Genetic Abnormalities

Under conventional approaches, individual genes are examined for abnormalities and drugs are administered to treat diseases caused by these abnormalities.

Molecular targeted drug therapy

# Why is Genomic Medicine Necessary in the Field of Oncology?

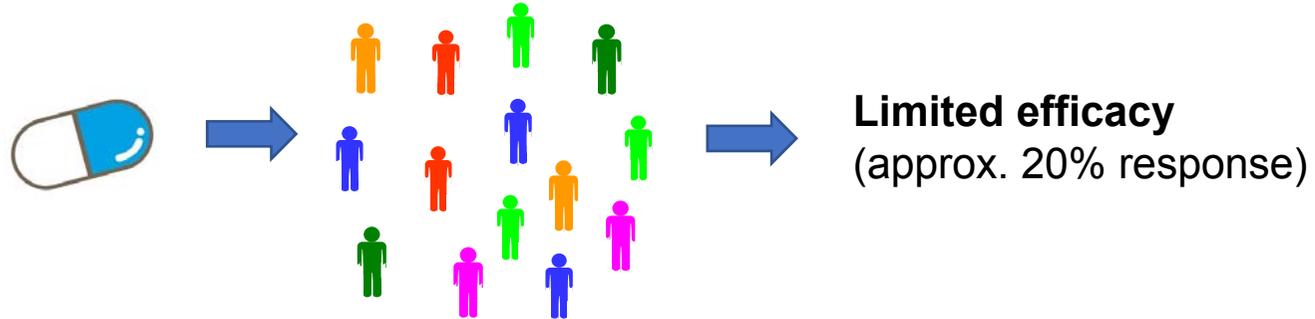
Cancers are caused by abnormal gene functions. For instance, lung cancers caused by different genetic abnormalities will be treated with different therapies.



The treatment of cancer has already entered the age of anticancer drug therapy based on genetic testing, and “cancer genomic medicine” will yield even greater therapeutic advancement.

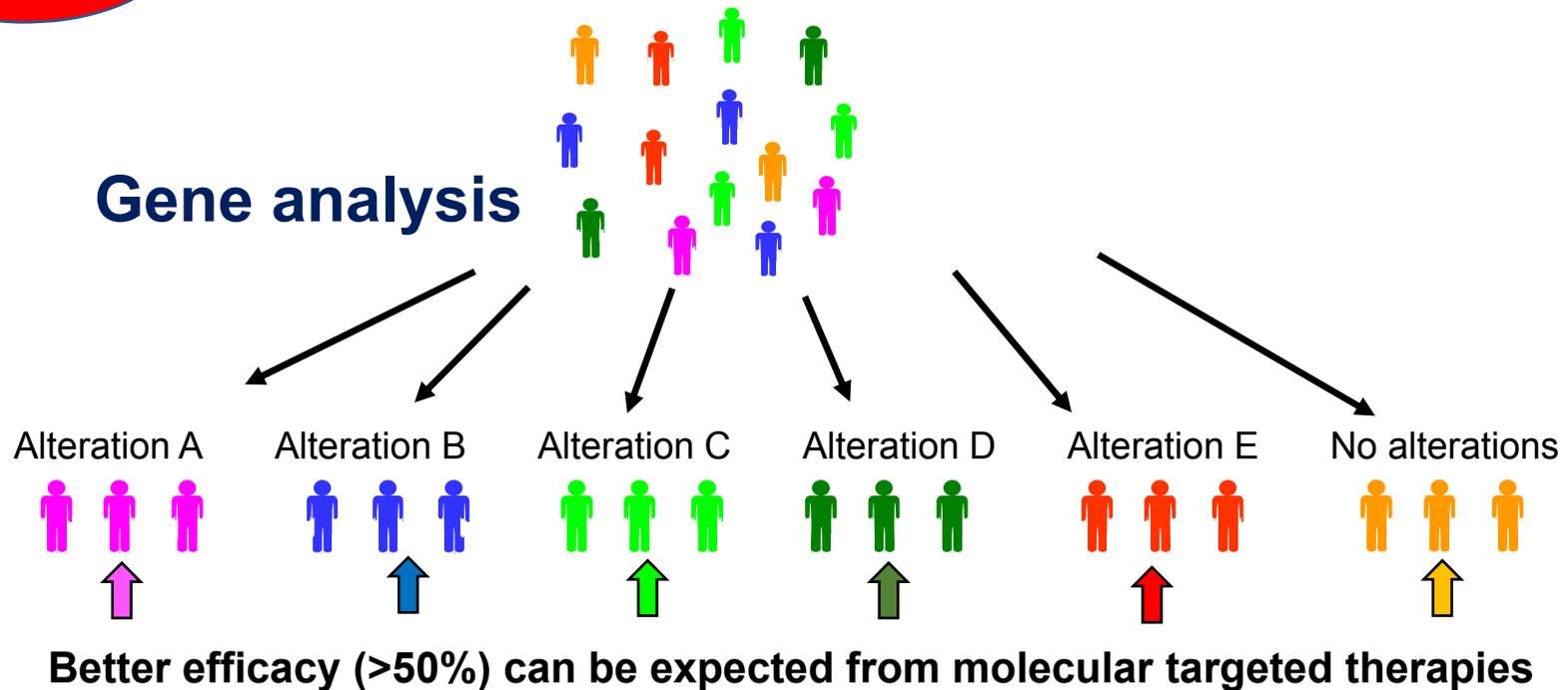
**Before**

**Use one drug for same cancer types**

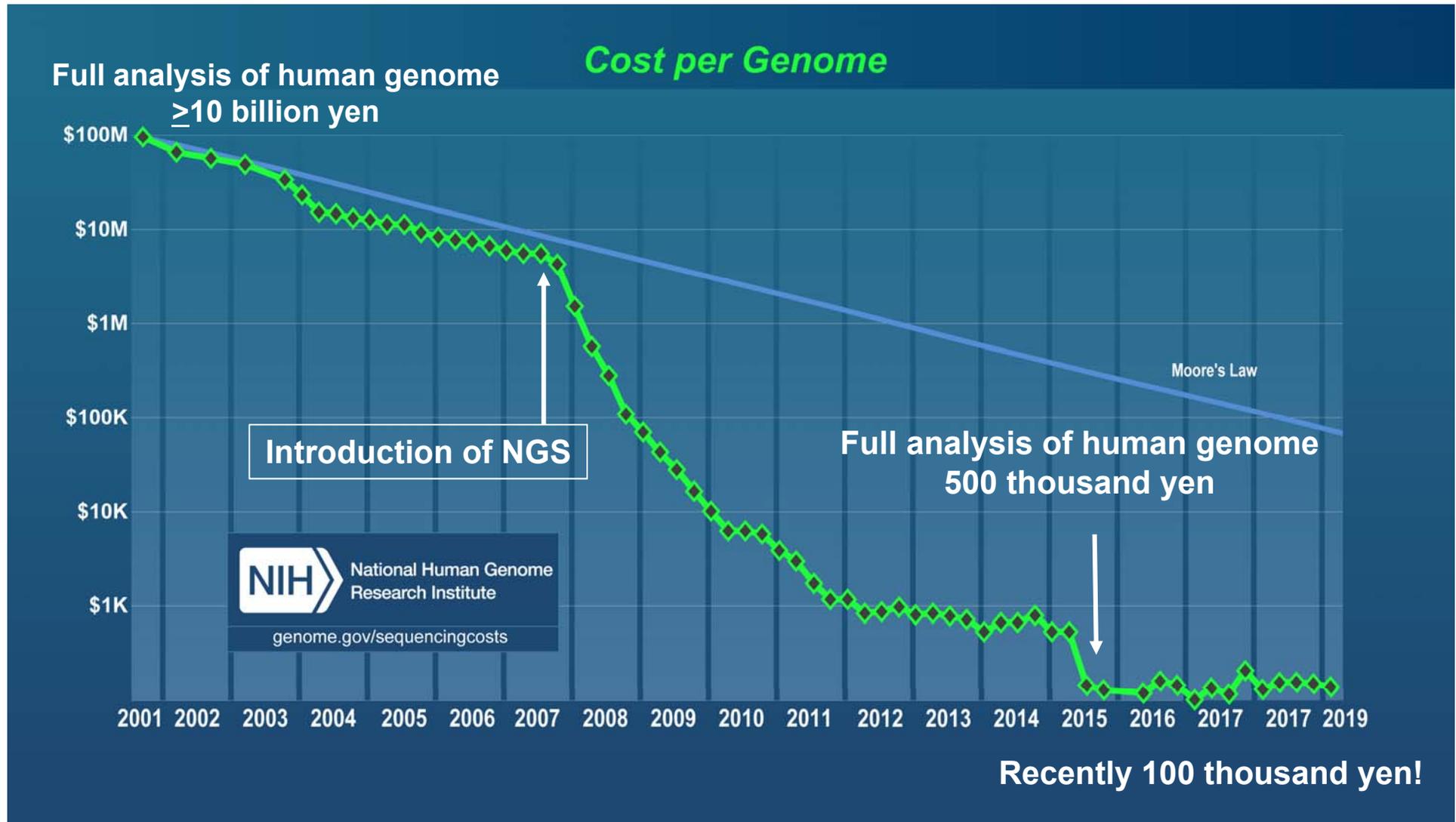


**Now**

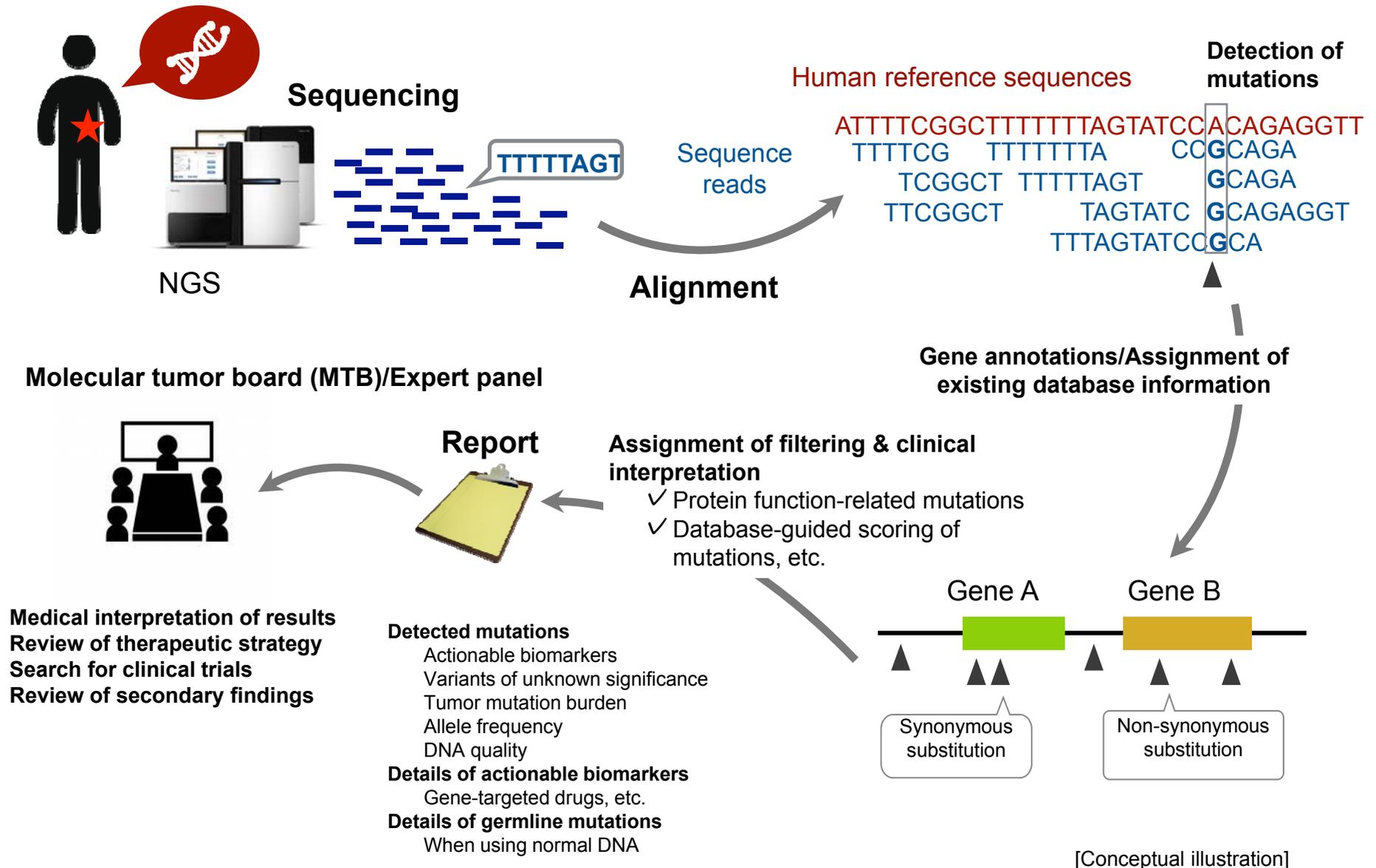
**Use drugs that best suit individuals regardless of cancer types based on their genetic features.**



# Advances in Gene Analysis Technologies



# Flow of Genomic Medicine Using NGS



**Main NGS-based genomic profiling tests with  
regulatory approval & national health insurance coverage  
(NHI) in Japan**

**(Reviewed by Committee on Medical Devices and *in vitro*  
Diagnostics on Dec 13, 2018;**

**Reviewed by Central Social Insurance Medical Council  
[Chuikyo] on May 30, 2019)**

OncoGuide™ NCC Oncopanel System (Sysmex)

FoundationOne® CDx Cancer Genomic Profile  
(Chugai/Roche)

## OncoGuide NCC Oncopanel System

The OncoGuide NCC Oncopanel System is a combination of medical devices comprising a template DNA preparation reagent and analysis program. It is used to output gene mutation data to support the development of therapeutic strategies based on comprehensive genomic profiling of 114-cancer related genes identified in patients with solid cancers.

## FoundationOne CDx Cancer Genomic Profile

FoundationOne CDx Cancer Genomic Profile is an analysis program that outputs data on gene mutations to support the development of treatment strategies and the assessment of drug indications based on comprehensive genomic profiles of 324 cancer-related genes identified in patients with solid cancers.

# FoundationOne CDx Cancer Genomic Profile Can Also be Used as a Companion Diagnostic

Alterations	Cancer type	Relevant drugs
<i>EGFR</i> exon 19 deletions and <i>EGFR</i> exon 21 L858R alterations	Non-small cell lung cancer	afatinib dimaleate, erlotinib hydrochloride, gefitinib, osimertinib mesylate
<i>EGFR</i> exon 20 T790M alterations		osimertinib mesylate
<i>ALK</i> fusion genes		alectinib hydrochloride, crizotinib, ceritinib
<i>BRAF</i> V600E and V600K alterations	Malignant melanoma	dabrafenib mesylate, trametinib dimethyl sulfoxide, vemurafenib
<i>ERBB2</i> copy number alterations ( <i>HER2</i> gene amplification positive)	Breast cancer	trastuzumab (genetical recombination)
<i>KRAS/NRAS</i> wild-type	Colorectal cancer	cetuximab (genetical recombination), panitumumab (genetical recombination)
<i>NTRK1/2/3</i> fusion gene	Solid tumors	entrectinib

Source: Chugai Obtains Approval of FoundationOne CDx Cancer Genomic Profile as a Companion Diagnostic for Rozlytrek  
[https://www.chugai-pharm.co.jp/english/news/detail/20190627120000\\_628.html](https://www.chugai-pharm.co.jp/english/news/detail/20190627120000_628.html) (accessed on June 28, 2019)

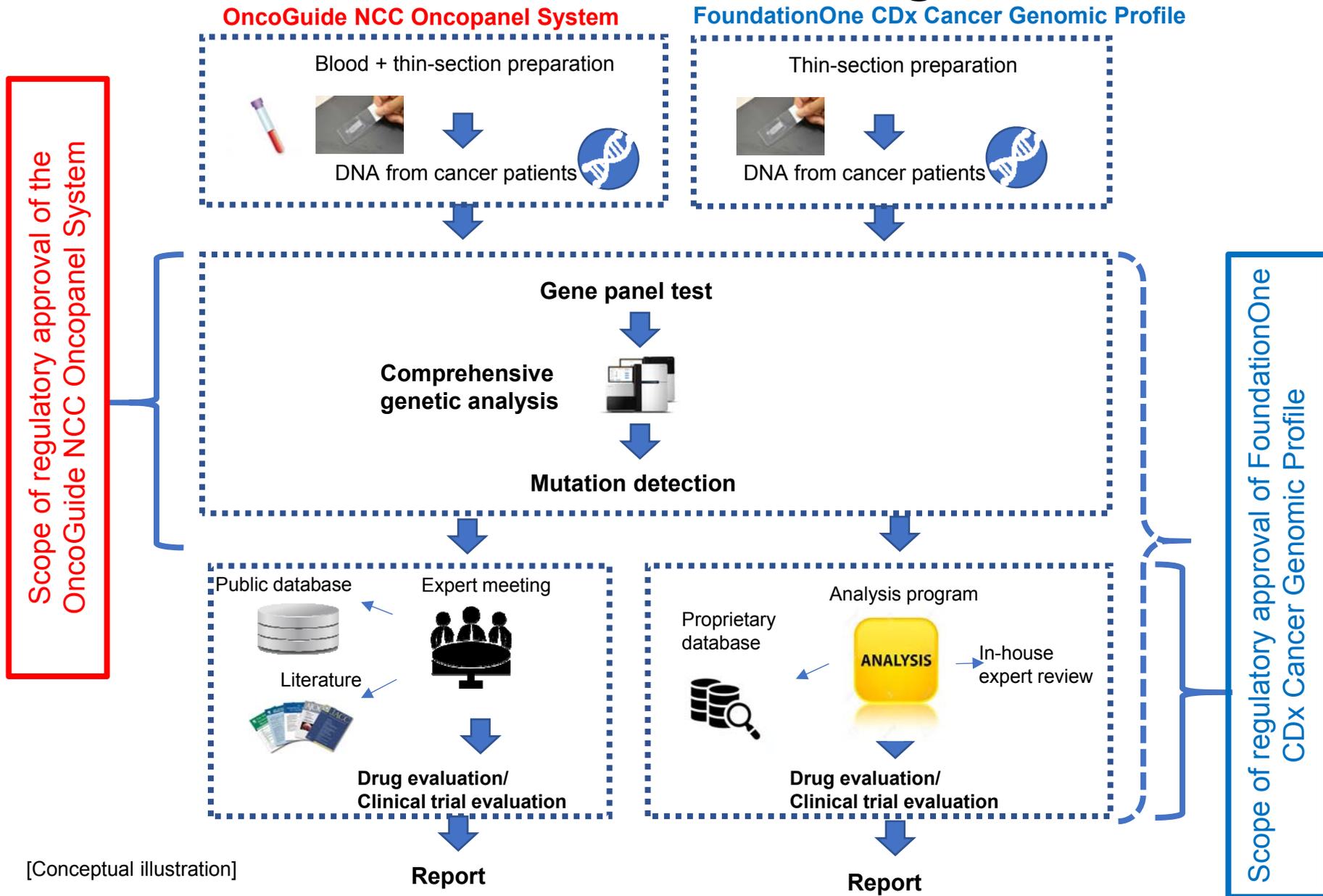
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The OncoGuide NCC Oncopanel System is a **combination of medical devices** comprising a template DNA preparation reagent and analysis program. It is used to output gene mutation data to support the development of therapeutic strategies based on comprehensive genomic profiling of 114-cancer related genes identified in patients with solid cancers.

## FoundationOne CDx Cancer Genomic Profile

FoundationOne CDx Cancer Genomic Profile is an **analysis program** that outputs data on gene mutations to support the development of treatment strategies and the assessment of drug indications based on comprehensive genomic profiles of 324 cancer-related genes identified in patients with solid cancers.

# Difference in Scope of Regulatory Approval of Genomic Medicine using NGS



[Conceptual illustration]

# An Example of C-CAT Survey Results

作成中

## C-CAT調査結果



### 1 基本項目

#### 1-1 患者

登録ID	xxxxxx	匿名化患者ID	ABC123456	検査ID	123456789
年齢	65	性別	男性	がん種	Breast Cancer

#### 1-2 医療機関

連携病院	テスト大学病院	拠点病院	テスト病院	中核拠点病院	テスト病院
------	---------	------	-------	--------	-------

#### 1-3 検査

検体採取日	2016/2/1	腫瘍細胞割合	70%	組織名	肺
パネル名	NCC oncopanel	測定日	2018/4/5		

### 2 検査結果

概要

▲ 薬剤への到達性の指標をご参照ください。

検出変異数	国内承認薬	国内臨床試験中	国内適用外承認	海外臨床試験中	FDA承認薬
体細胞変異: 9 生殖細胞系列変異: 3	5	2	2	2	7

塩基置換、挿入、欠失(DNA)

No.	マーカー	変化	アレル頻度	エビデンスタイプ	臨床的意義	エビデンスレベル	薬剤	薬剤への到達性	QCI
1	EGFR	p.L858R	0.288 (553/1918)	Predictive	Sensitive	A	Geftinib	国内承認 FDA承認	Tier 1A Pathogenic
				Predictive	Sensitive	A	Erlotinib	国内承認 FDA承認	
				Predictive	Sensitive	A	Afatinib	国内承認 FDA承認	
				Predictive	Sensitive	A	Osmertinib	国内承認 FDA承認	
				Oncogenicity	Likely Oncogenic	F			
2	MSH2	p.S900*	0.530 (1459/2752)	Predisposing	Pathogenic	F		Tier 1A Pathogenic	
				Oncogenicity	Likely Oncogenic	F			
3	FGFR3	N718S	0.474 (832/1754)						
4	STK11	F354L	0.502 (722/1437)						
5	APC	D900V	0.500 (150/300)						
6	BRCA1	Y856H	0.47 (94/200)						

作成日: 2018年11月2日  
レポートバージョン: 1.0

作成中

遺伝子再構成(DNA)、構造異型(DNA)

No.	マーカー	変異種類	サイト/バンド	エビデンスタイプ	臨床的意義	エビデンスレベル	薬剤	薬剤ステータス	QCI
7	CCDC6-RET	Rearrangement	chr10:61,657,343 chr10:43,611,194	Predictive	Sensitive	C	Vandetanib	国内適応外	Tier 1A Pathogenic
				Predictive	Sensitive	C	Lenvatinib	国内適応外	
				Predictive	Sensitive	D	Nintedanib		
		Oncogenicity	Oncogenic	F					

コピー数変化 (T/N実施検査のみ)

No.	マーカー	コピー数変化	サイト/バンド	エビデンスタイプ	臨床的意義	エビデンスレベル	薬剤	薬剤ステータス	QCI
8	MYC	19.5	chr8:127,736,069-127,741,434	Oncogenicity	Oncogenic	F			

融合遺伝子(RNA)やエクソスキッピング(RNA)

No.	マーカー	融合遺伝子	サイト/バンド	リード数	エビデンスタイプ	臨床的意義	エビデンスレベル	薬剤	薬剤ステータス	QCI
9	MYB	MYB-NFIB (15/16 - 11/12)	6q23.3-9q23	540 (98.5%)	Predisposing	Pathogenic	F			

Tumor Mutational Burden (T/N実施検査のみ)

Tumor Mutational Burden
6 (Mut/Mb)

生殖細胞系列変異 (T/N実施検査のみ)

▲ 非がん部組織由来のDNA解析結果を表示しております。

No.	マーカー	変化	アレル頻度	エビデンスタイプ	臨床的意義 ClinVar ID	エビデンスレベル	薬剤	薬剤ステータス	QCI
1	PTEN	C136Y	0.500 (150/300)	Predisposing	Pathogenic RCV000169797.1				Pathogenic
2	BRCA1	Y856H	0.47 (94/200)	Predisposing	Likely benign RCV000148383.1				Benign
3	BRCA1	E1559K	-	Predisposing	Pathogenic				Tier 1A Pathogenic
				Predictive	Sensitive	A	Olaparib	国内承認 FDA承認	
				Predictive	Sensitive	D	Rucaparib	国内試験中 FDA承認	
				Predictive	Sensitive	B	Talazoparib	国内試験中 FDA承認	
							Niraparib	FDA承認	

### 3 候補となる臨床試験一覧

※下記の治験・臨床試験については、詳細な適格基準・除外基準に合致しているかが、患者登録受付中であるかが、「実施機関 (連絡先)」への確認が必要となります。

Rucaparibを用いた国内臨床試験

臨床試験DB	試験ID	試験名称	Phase	対象マーカー	データ更新日	実施機関 (連絡先)
Clinicaltrials	NCT03499444	A Study of Rucaparib in Japanese Patients With a Previouslytreated Solid Tumor	P2	BRCA1 E1559K	2018/4/5	ClovisOncology (clovistrials@emergingmed.com)

from Information meeting for Core hospitals on 2019/02/05

作成日: 2018年11月2日  
レポートバージョン: 1.0

# Characteristics of FoundationOne CDx Cancer Genomic Profile

- Comprehensive detection/analysis of mutations\*<sup>1</sup> in 324 oncogenes

\*1: substitutions, insertion/deletion alterations, copy number alterations, and rearrangements

- Companion diagnostics

- MSI\*<sup>2</sup> determination/TMB\*<sup>3</sup> score calculation

\* 2 : Microsatellite Instability      \* 3 : Tumor Mutational Burden

- Analysis result report provides expert review and assignment of clinical significance (annotations)

- Tested and analyzed with abundant experience in FMI

# Analysis Report of FoundationOne CDx

## APPROVED THERAPEUTIC OPTION IN JAPAN

CDx Associated Findings		FDA-APPROVED THERAPEUTIC OPTIONS	
GENOMIC FINDINGS DETECTED			
EGFR L858R		Giotinif® (Afinib®)	
		Heskin® (Gefitinib®)	
		Tacveva® (Erlotinib®)	
TP53M1		Tagrisso® (Osimertinib)	

OTHER ALTERNATIVE BIOMARKERS IDENTIFIED	
Tumor Mutational Burden: 11 Mut/Mb <sup>†</sup>	ARRIS® Q204 <sup>†</sup>
CDKN2A/B loss <sup>†</sup>	TP53 R273P
EGFR amplification <sup>†</sup>	

(1) CDx results with identified gene mutations and corresponding drugs

Results should be used for diagnosis and finding applicable anticancer drugs (companion diagnostics) based on identified gene mutations.

(2) Non-CDx results with identified gene mutations and biomarkers that are not used for companion diagnostics

Identified gene mutations and biomarkers that are not used for companion diagnostics are listed. Results should be used for diagnosis and support for treatment decision

\* : You can review mutation data file and download/print the Analysis results reports at Chugai FMI portal site. Printed Analysis result report will be sent to medical institutions from Health inspection stations. Contents of both reports are identical.

- [Precautions]**
- In case of confirmation of the Analysis results reports, please refer to the latest package insert of FoundationOne CDx Cancer Genomic Profile.
  - "APPROVED THERAPEUTIC OPTION IN JAPAN" is prepared based on approved drug information at the time of first approval of FoundationOne CDx Cancer Genomic Profile in Japan. In case of determining therapeutic drugs, please refer to the latest package inserts of the drugs.

APPROVED THERAPEUTIC OPTION IN JAPAN provides CDx results with identified gene mutations and corresponding drugs, Non-CDx results with other identified gene alterations, testing results of microsatellite instability and tumor mutation burden.

Source: Chugai website (accessed on June 28, 2019)

<https://chugai-pharm.jp/content/dam/chugai/product/f1t/cdx/report-fullmock/doc/F1CDx-Report-Fullmock.pdf>

# Analysis Report of FoundationOne CDx Cancer Genomic Profile

## PROFESSIONAL SERVICES

The image shows a screenshot of a FoundationOne CDx Cancer Genomic Profile report. The report is titled "FOUNDATION ONE CDx" and "Lung Adenocarcinoma". It includes sections for "Marker Findings", "Genomic Findings", "Tumor Mutation Burden", "Microsatellite Instability", and "Copy Number Alterations". A table lists therapeutic drugs with their indications for different cancer types and biomarkers.

**(1) Results of gene mutation and biomarker detection**

**(2) Therapeutic drugs with an indication for the cancer type with detected gene mutations and biomarkers**

**(3) Therapeutic drugs with an indication for other cancer types with detected gene mutations and biomarkers**

**(4) Ongoing clinical trial information for the treatment of cancers with detected gene mutations and biomarkers**

**(5) Therapeutic drugs with an indication for the cancer type with detected gene mutations, but with no expected effectiveness due to tolerance etc.**

Explanations for detected gene mutations and biomarkers, therapeutic drugs with expected efficacy and clinical trials are described in following pages. Explanations for VUS (Variant of Unknown Significance) and FoundationOne CDx Cancer Genomic Profile are described in Appendices.

\* : You can review mutation data file and download/print the Analysis result report on Chugai FMI portal site. Printed Analysis result report will be sent to medical institutions from Health inspection stations. Contents of both reports are identical.

**[Precautions]**

- In case of confirmation of the Analysis result report, please refer to the latest package insert of FoundationOne CDx Cancer Genomic Profile.
- "PROFESSIONAL SERVICES" is not approved by the Ministry of Health, Labour and Welfare.

PROFESSIONAL SERVICES provide information such as detected gene mutation and MSI/TMB (biomarker), therapeutic drugs with expected clinical efficacy, clinical studies and tolerance etc.

# **Cancer genomic medicine in Japan**

# Draft Selection Criteria for Core Hospitals for Cancer Genomic Medicine

資料5

がんゲノム医療中核拠点病院(案)等の  
指定要件(案)

厚生労働省健康局  
がん・疾病対策課

# Report of Cancer Genomic Medicine Promotion Consortium Council (excerpt)

Requirements for medical institutions to implement cancer genomic medicine:

- (1) A system to conduct panel tests (including outsourcing to external organizations)
- (2) A group of experts to offer medical interpretation of panel test results (including collaboration with other organizations in some areas of medical care)
- (3) Ability to provide expert genetic counseling for patients with hereditary and other cancers
- (4) Access to a certain number of patients who are eligible for panel testing, etc.
- (5) Ability to collect and manage panel test results and clinical information in a secured manner, and to register necessary information with Center for Cancer Genomics and Advanced Therapeutics.
- (6) A system for rapid freezing and storage of surgical specimens and other biological specimens
- (7) An appropriate framework for and experiences in conducting clinical studies including those for advanced medical care, investigator-initiated clinical trials, and global clinical trials.
- (8) Comprehensive and accessible contact services for patients and other stakeholders regarding use of medical information, clinical trial information and etc.

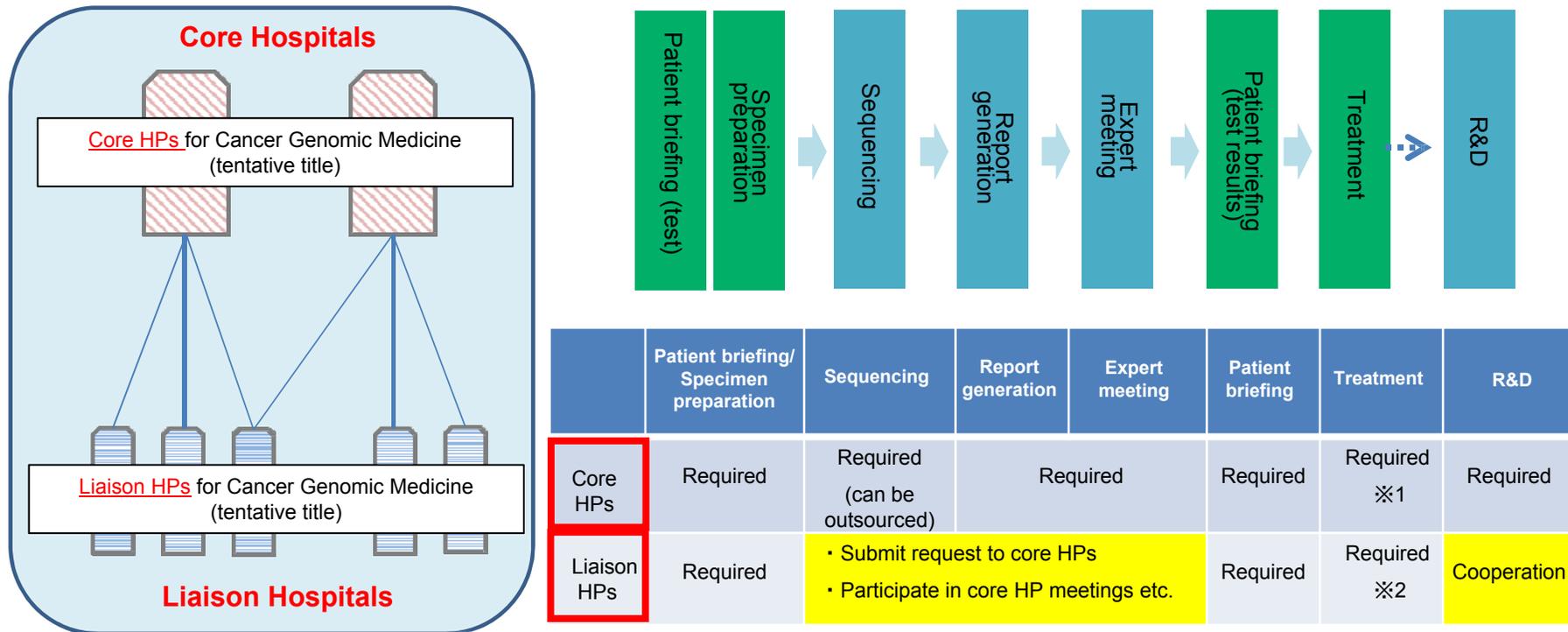
# Committee on Designation of Core Hospital for Cancer Genomic Medicine (2018/2/14)

The committee selected the following medical institutions.

Minister of Health, Labour and Welfare will designate Core Hospital for Cancer Genomic Medicine based on today's discussion.

#	Prefecture	Core hospitals for cancer genomic medicine
1	Hokkaido	Hokkaido University Hospital
2	Miyagi	Tohoku University Hospital
3	Chiba	National Cancer Center (NCC) Hospital East
4	Tokyo	Keio University Hospital
5	Tokyo	University of Tokyo Hospital
6	Tokyo	NCC Hospital
7	Aichi	Nagoya University Hospital
8	Kyoto	Kyoto University Hospital
9	Osaka	Osaka University Hospital
10	Okayama	Okayama University Hospital
11	Fukuoka	Kyushu University Hospital

# Conceptual Illustration of the System for Delivering Cancer Genomic Medicines and Required Functions (draft)



※1 In order to secure access to non-approved & off-label drugs, the Core Hospital for Cancer Genomic Medicine (tentative title) must establish a system capable of taking the initiative in conducting clinical trials (including investigator-initiated trials) and advanced medical care.

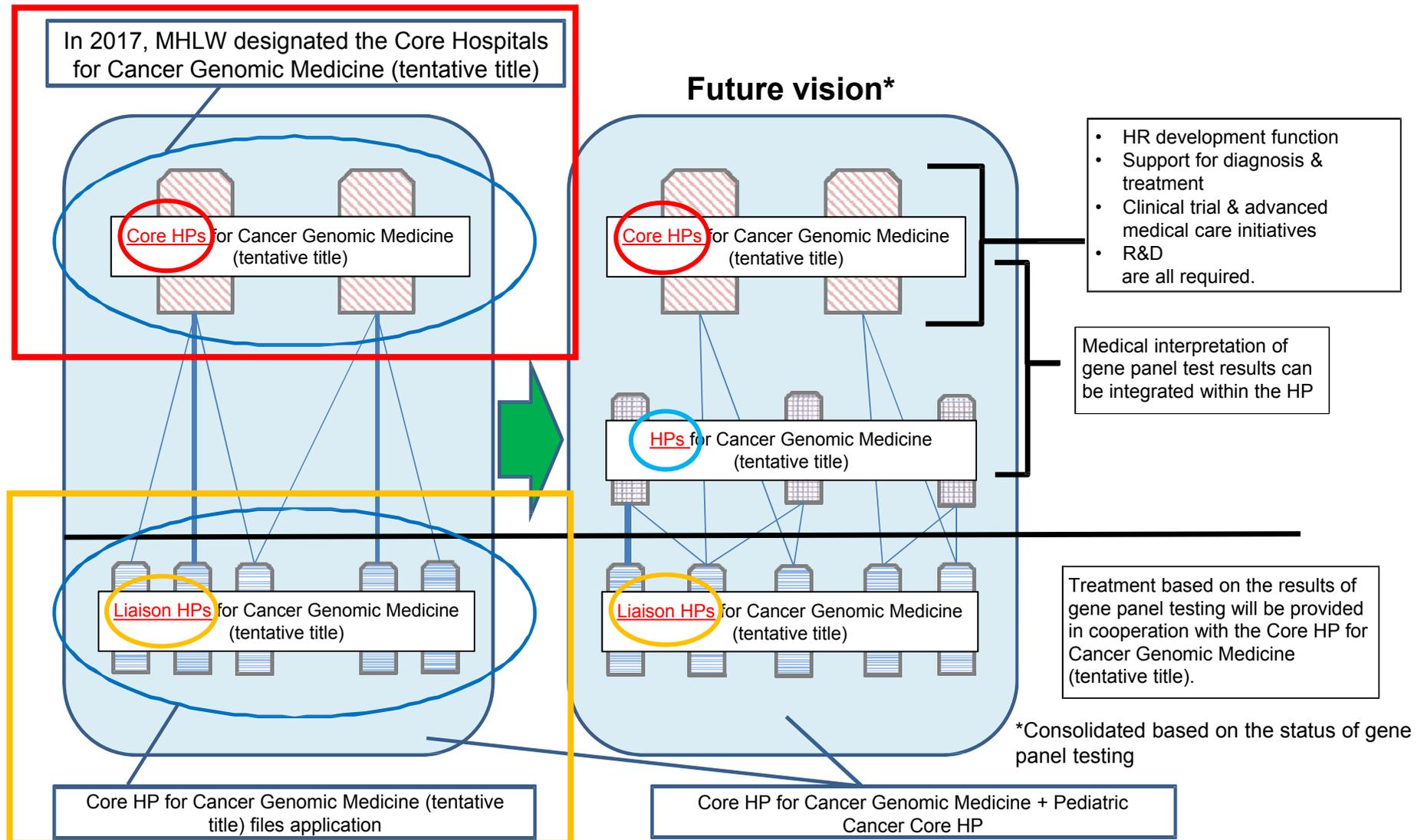
※2 The Liaison Hospital for Cancer Genomic Medicine (tentative title) must establish a system to facilitate participation in the above-mentioned clinical trials and advanced medical care.

## Number of Liaison Hospitals Affiliated with Each Core Hospital

Core hospitals for cancer genomic medicine	Apr 2018	Oct 2018	Apr 2019
Hokkaido University Hospital	2	4	4
Tohoku University Hospital	6	8	8
National Cancer Center (NCC) Hospital East	6	8	9
Keio University Hospital	24	35	36
National Cancer Center Hospital	9	20	23
University of Tokyo Hospital	14	16	19
Nagoya University Hospital	14	18	21
Kyoto University Hospital	20	23	28
Osaka University Hospital	8	13	15
Okayama University Hospital	16	21	29
Kyushu University Hospital	13	16	19
<b>No. of liaison hospitals</b> <small>*Some liaison hospitals are affiliated with multiple core hospitals.</small>	<b>100</b>	<b>135</b>	<b>156</b>

System for Delivering Cancer Genomic Medicine: Cancer and Disease Control Division,  
Health Services Bureau, Ministry of Health, Labour and Welfare (MHLW)

**Future vision of the system for delivering cancer genomic medicine (draft)**



Source: The 2<sup>nd</sup> Sub-working group meeting on the designated requirements of core HPs for Cancer Genomic Medicine (tentative title), Document #4 (2017/09/11) <https://www.mhlw.go.jp/file/05-Shingikai-10901000-Kenkoukyoku-Soumuka/0000177033.pdf>

Accessed: 2019/06/25

# Outline of the Working Group Meeting on the Designated Requirements of Core HPs for Cancer Genomic Medicine (Document in Japanese Only)

資料 1

## 「がんゲノム医療中核拠点病院等の指定要件に関するワーキンググループ」 開催要綱

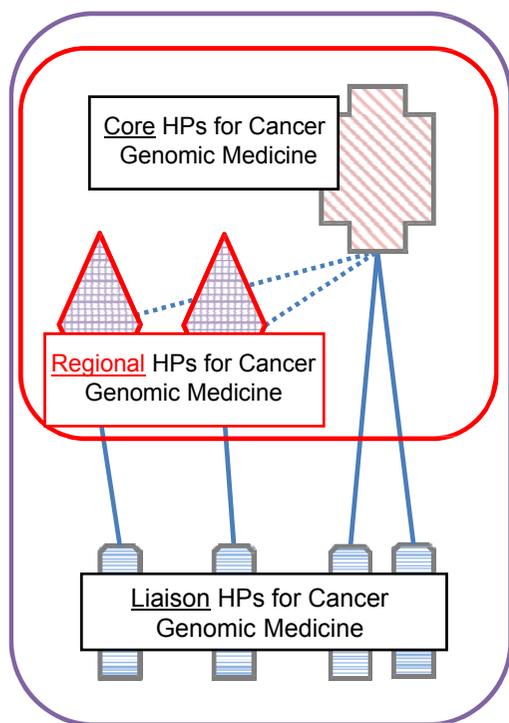
### 1. 趣旨

政府としては、平成 30 年（2018）3 月に閣議決定された第 3 期がん対策推進基本計画に基づき、ゲノム医療を必要とするがん患者が、全国どこにいても、がんゲノム医療を受けられる体制整備を進めてきた。

平成 29（2017）年 12 月に「がんゲノム医療中核拠点病院等の整備について「がんゲノム医療中核拠点病院等の整備に関する指針（平成 29 年 12 月 25 日健発 1225 第 3 号厚生労働省健康局長通知の別添）」を発出し、平成 30（2018）年 2 月に、がんゲノム医療を牽引する高度な機能を有する医療機関として「がんゲノム医療中核拠点病院（以下、中核拠点病院）」を全国に 11 箇所指定し、中核拠点病院と連携して、がんゲノム医療を提供する「がんゲノム医療連携病院」を平成 31（2019）年 4 月までに 156 箇所公表してきた。

今般、がんゲノム医療提供体制をさらに充実させるため、「がんゲノム医療中核拠点病院等の整備に関する指針」を見直すとともに、自施設でがんゲノム医療を完結できる医療機関として「がんゲノム医療拠点病院」の指定要件を策定するため、本ワーキンググループを設置し、検討結果を「がん診療提供体制のあり方に関する検討会」に報告することとする。

# Future functions of Core, Regional & Liaison Hospitals for Cancer Genomic Medicine (draft)



The Core HP for Cancer Genomic Medicine submits the application for accreditation of the Liaison HP for Cancer Genomic Medicine\*

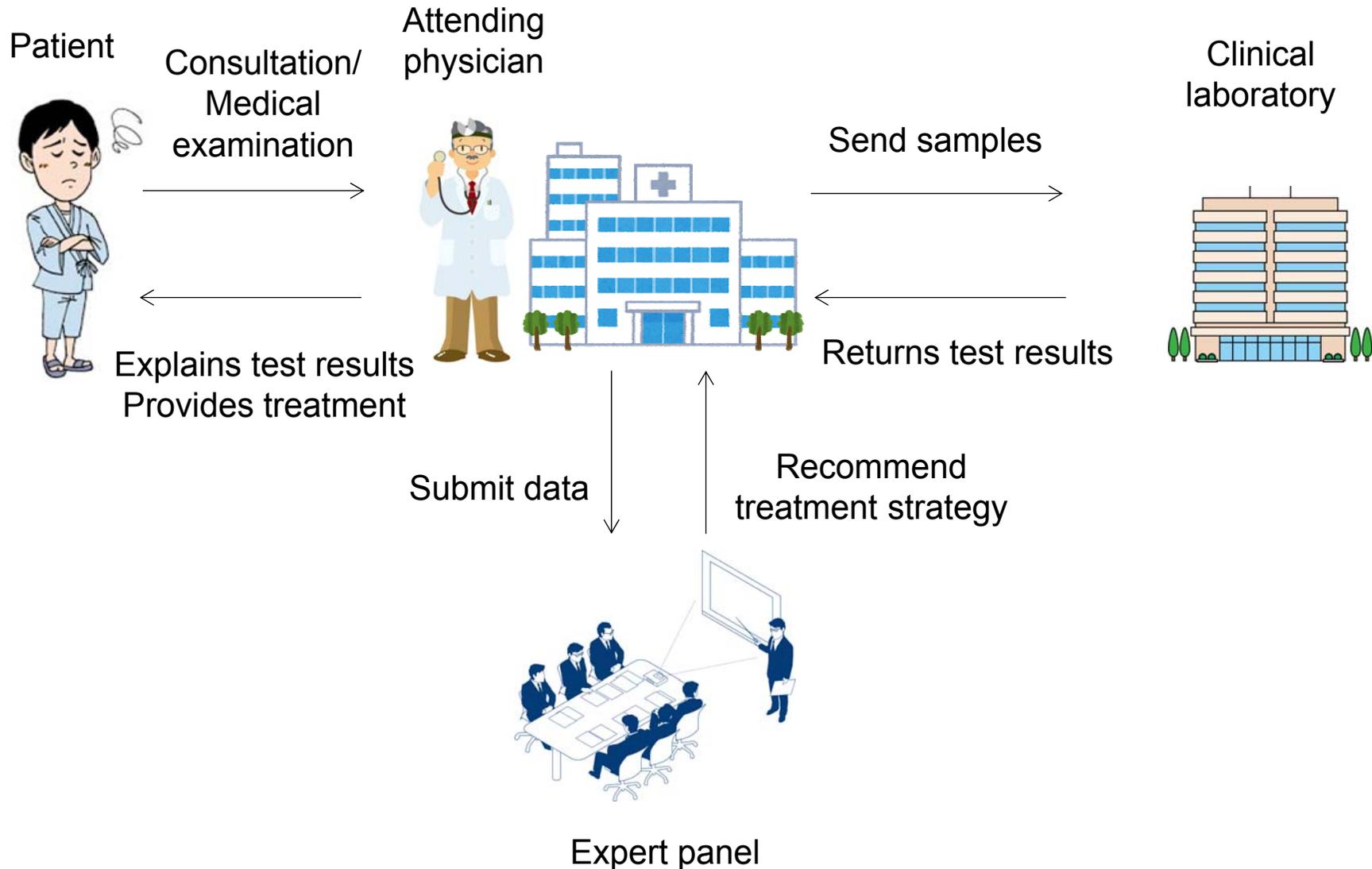
	Patient briefing (test) Specimen preparation	Sequencing	Expert panel	Report generation	Patient briefing (test results)	Treatment	Advanced medical care, clinical trial, R&D	HR development
	Patient briefing/ Specimen preparation	Sequencing	Expert meeting	Report generation	Patient briefing	Treatment	R&D	HR development
Core HPs	Required	Can be outsourced	Required		Required	Required	Required	Required
Regional HPs (proposed)	Required	Can be outsourced	Required		Required	Required	Work together	Work together
Liaison HPs	Required	Can be outsourced	Participate in Core or Regional HP meetings, etc.		Required	Required	Work together	Work together

\*After confirming that the developmental requirements stipulated in the above-mentioned guidance have been satisfied, the Core HP or Regional HP for Cancer Genomic Medicine will submit an application to the MHLW Minister seeking accreditation of medical institutions with which it collaborates as Liaison HPs for Cancer Genomic Medicine.

Thereafter, the Core HP or Regional HP for Cancer Genomic Medicine will submit approximately one application **each year** to the Minister of MHLW to seek accreditation of additional Liaison HPs.

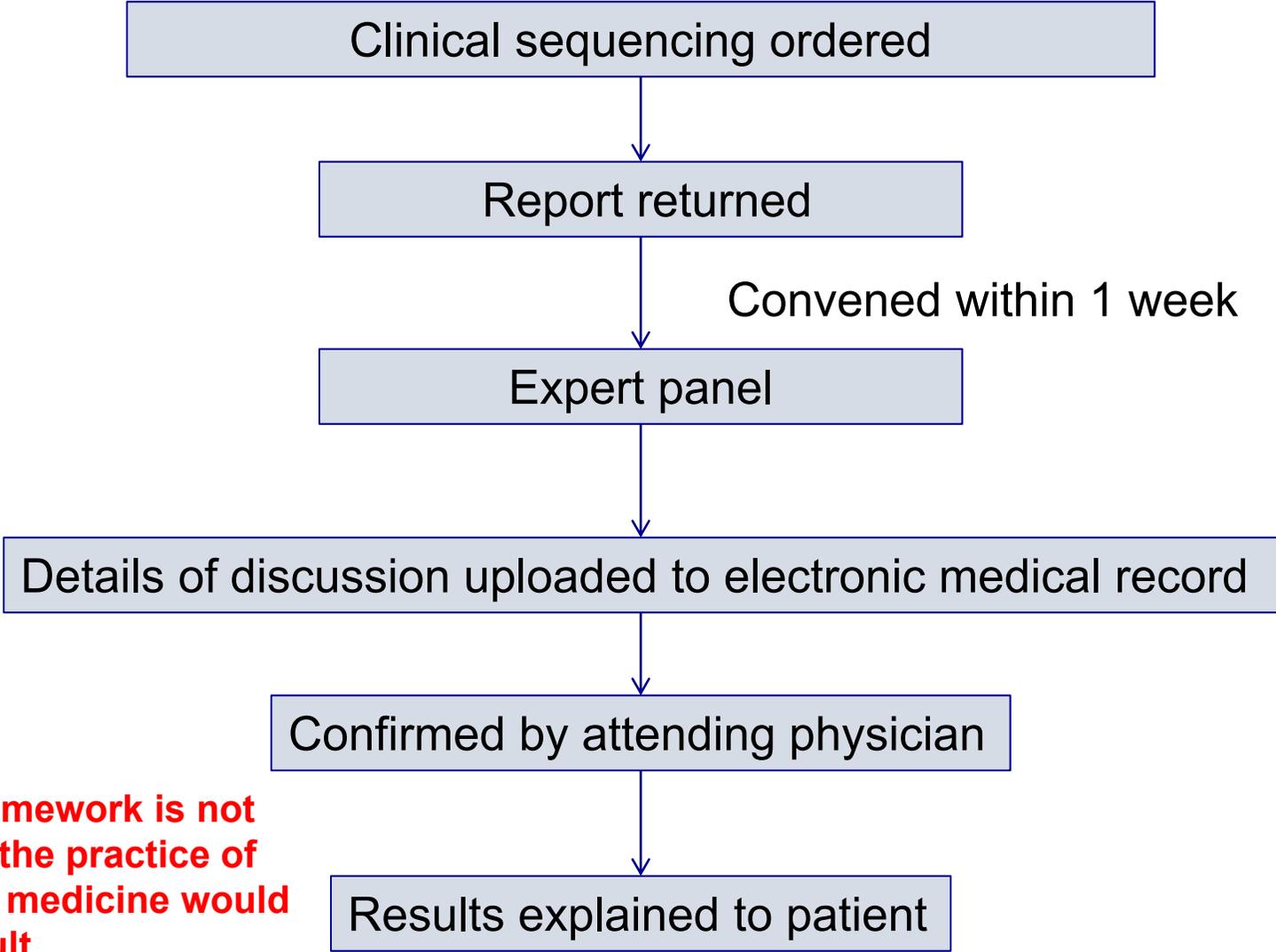
# Ideal Concept for Delivering Cancer Genomic Medicine

All aspects should be integrated within a single hospital



# Convening the expert panel

# Flowchart for returning reports to attending physicians & explaining results to patients

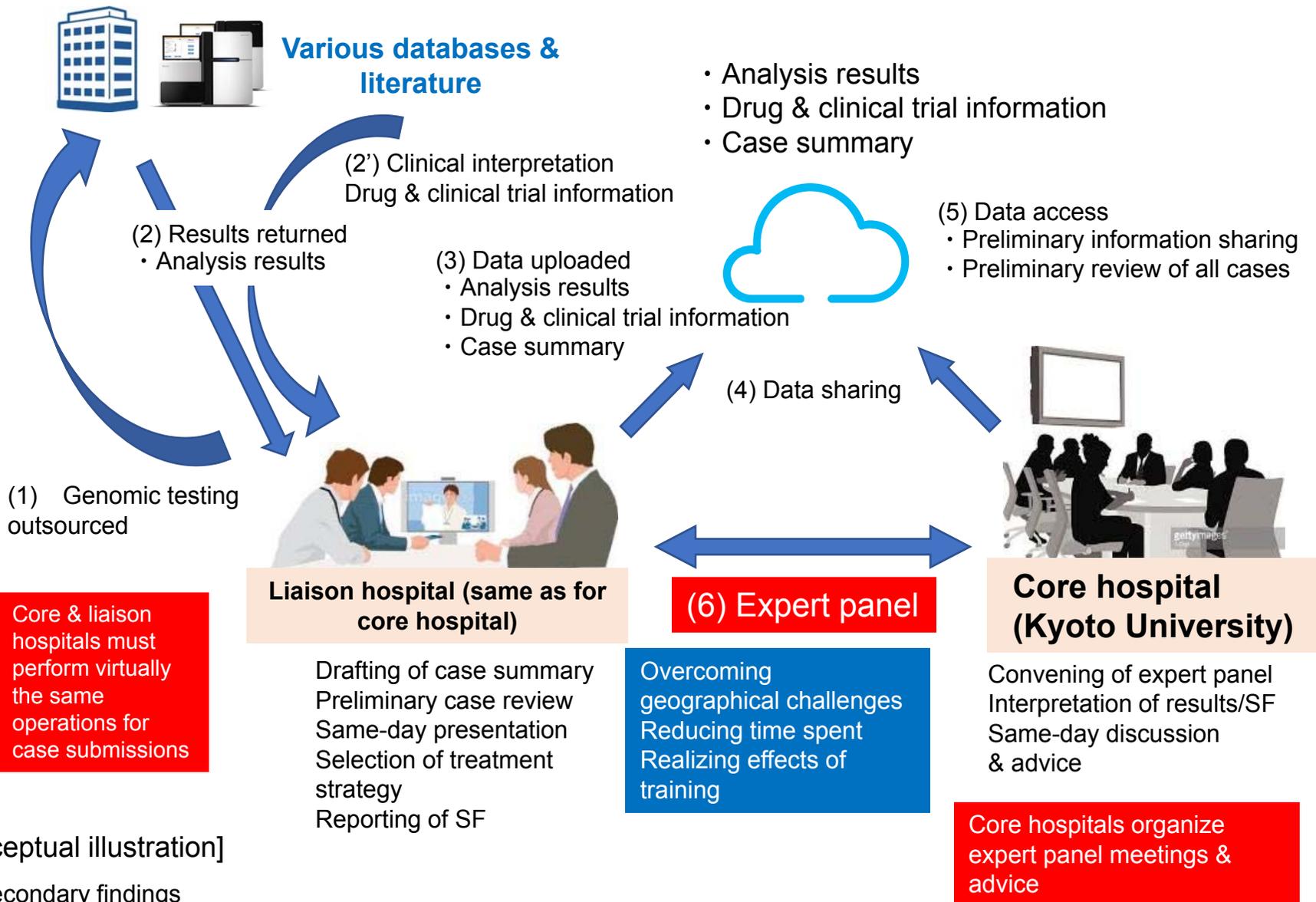


**If this framework is not present, the practice of genomic medicine would be difficult.**

# Practical Example of Expert Panel Operation via Web-based System

Clinical laboratory

Cloud-based data storage contracted by core hospital



[Conceptual illustration]

SF: Secondary findings





# Expert Panel Structure & Performance

- Panel members

Specialization	Specialization
Clinical oncologist (clinical department)	Accredited genetic counselor
Surgical oncologist (surgery department)	Bioinformatician (NGS analyst)
Pathologist	Biobank personnel
Radiotherapist	Genomic medicine personnel
Clinical geneticist	CRC
Nurse	Postgraduate student in basic medical sciences
Pharmacist	Postgraduate student in genetic counseling
Clinical laboratory technician	Participation is also open to interested individuals

Expert panel members are required to sign a confidentiality agreement, and attendance is confirmed.

- Time & date of expert panel meetings:

From 5:30pm every Tuesday @Common Conference Room, 1F, Kyoto University Hospital Cancer Center

- Meeting format:

Web conference via WebEx: Liaison Hospitals attend (it is allowed to join the conference without cases)

# Procedure in Preparation for Expert Panel Meeting

	Fri	Sat	Sun	Mon	Tue
<b>Core hospitals</b>	<ul style="list-style-type: none"> <li>• Check liaison hospital report(s)</li> <li>• Notify liaison hospital coordinating physician and web conference supervisor of scheduled meeting</li> <li>• Also notify core hospital participants of scheduled meeting</li> <li>• Report whether any cases will be reviewed (by 9:00 am on the day of meeting)</li> </ul>			<ul style="list-style-type: none"> <li>• Distribute case summaries (for preliminary reviews)</li> </ul>	<ul style="list-style-type: none"> <li>• Send web conference invitation</li> <li>• Prepare web conference connection (from 3:00 pm)</li> <li>• Prepare web conference meeting (from 5:00 pm)</li> <li>• Convene web-based expert panel meeting (5:30 pm – around 7:00 pm )</li> </ul>
<b>Liaison hospitals</b>	<ul style="list-style-type: none"> <li>• Report if any case reviews are scheduled (by 9:00 am on the day of meeting)</li> <li>• Upload analysis report of case review(s) to Box</li> </ul>			<ul style="list-style-type: none"> <li>• Upload presentation summary to Box by morning of meeting</li> </ul>	<ul style="list-style-type: none"> <li>• Check web conference connection (from 3:00 pm)</li> <li>• Prepare web conference meeting (from 5:00 pm)</li> <li>• Participate in meeting via web conferencing</li> </ul>

Draft case summary  
 Review secondary findings  
 Discussion of relevant literature

## Challenges

- (1) Burden on web conference supervisor
- (2) Burden of drafting summaries
- (3) Limit on number of cases that can be reviewed (limit of 7-8 cases)

# Standardizing Summaries for Web-based Expert Panel Meetings

## **Case outline**

Age, sex, clinical diagnosis, histopathological diagnosis, family history, past & current medical history, representative images, etc.

## **Sequencing results**

Positive biomarker, VUS, allele frequency, tumor mutational burden (TMB), any secondary findings (Y/N), etc.

## **Drug & clinical trial information on positive biomarker**

## **Interpretation of secondary findings**

## **Summary of results**

## **Discussion of relevant literature**

Drug & clinical trial evidence on positive biomarker, etc.

\*Due to data storage limitations, summaries must not exceed 15 PowerPoint slides and/or 5 MB in size.

VUS: Variant of unknown significance

# **Cancer Clinical Sequencing in Practice**

**Cancer clinical sequencing at KUH  
Data from OncoPrime, a non-covered  
medical treatment**

# Target Patients

---

- Cancers of unknown primary origin
- Rare cancers
- Recurrent advanced cancers refractory to standard-of-care therapy

# List of Genes Sequenced in OncoPrime

■ Gene mutations (SNV, insertion, deletion) (all exons of 210 genes)

■ Translocations (17 genes)

■ Genes with FDA-approved drugs

ABL	BLM	CRLF2	ESR1	H3F3A	MAP3K1	NFE2L2	PIK3R2	SETD2	TP53
ABL2	BRAF	CSF1R	EZH2	HNF1A	MAPK1	NOTCH1	PIK3R5	SF3B1	TP63
ACVR1B	BRCA1	CTNNA1	FAM123B	HRAS	MDM2	NOTCH2	PMS1	SMAD2	TP73
AKT1	BRCA2	CTNNB1	FANCA	IDH1	MDM4	NOTCH3	PMS2	SMAD3	TPMT
AKT2	BTK	CYP1A2	FBXW7	IDH2	MED12	NOTCH4	PPP2R1A	SMAD4	TRAF7
AKT3	CARD11	CYP2C19	FGFR1	IGF1R	MEN1	NPM1	PRDM1	SMARCA4	TSC1
ALK	CASP8	CYP2C9	FGFR2	IGF2R	MET	NRAS	PTCH1	SMARCB1	TSC2
APC	CBL	CYP2D6	FGFR3	IKZF1	MITF	NTRK1	PTCH2	SMO	TSHR
AR	CCND1	DAXX	FGFR4	IL7R	MLH1	NTRK2	PTEN	SOCS1	TYMS
ARAF	CCND2	DDR2	FLT1	INSR	MLL	NTRK3	PTPN11	SRC	U2AF1
ARID1A	CCND3	DNMT3A	FLT3	JAK1	MPL	PALB2	RAD50	SRSF2	UGT1A1
ARID1B	CCNE1	DPYD	FLT4	JAK2	MRE11A	PARP1	RAD51	STAG2	VHL
ASXL1	CDC73	EGFR	FOXL2	JAK3	MSH2	PAX5	RAF1	STAT1	VKORC1
ATM	CDH1	EP300	G6PD	KDM6A	MSH6	PBRM1	RB1	STAT3	WRN
ATR	CDK4	ERBB2	GATA1	KDR	MTHFR	PDGFRA	RET	STK11	WT1
ATRX	CDK6	ERBB3	GATA2	KIT	MTOR	PDGFRB	RICTOR	SUFU	XPC
AURKA	CDKN2A	ERBB4	GATA3	KLF4	MYC	PDK1	RNF43	TERT	XRCC1
AURKB	CDKN2B	ERCC1	GLI1	KRAS	MYCN	PGR	ROS1	TET2	
AXIN1	CEBPA	ERCC2	GNA11	MAML1	MYD88	PHF6	RPTOR	TGFBR2	
BAP1	CHEK1	ERCC3	GNAQ	MAP2K1	NBN	PIK3CA	RSPO2	TNFAIP3	
BCL2	CHEK2	ERG	GNAS	MAP2K2	NF1	PIK3CG	RSPO3	TOP1	
BCOR	CREBBP	ERRFI1	GRIN2A	MAP2K4	NF2	PIK3R1	RUNX1	TOP2A	

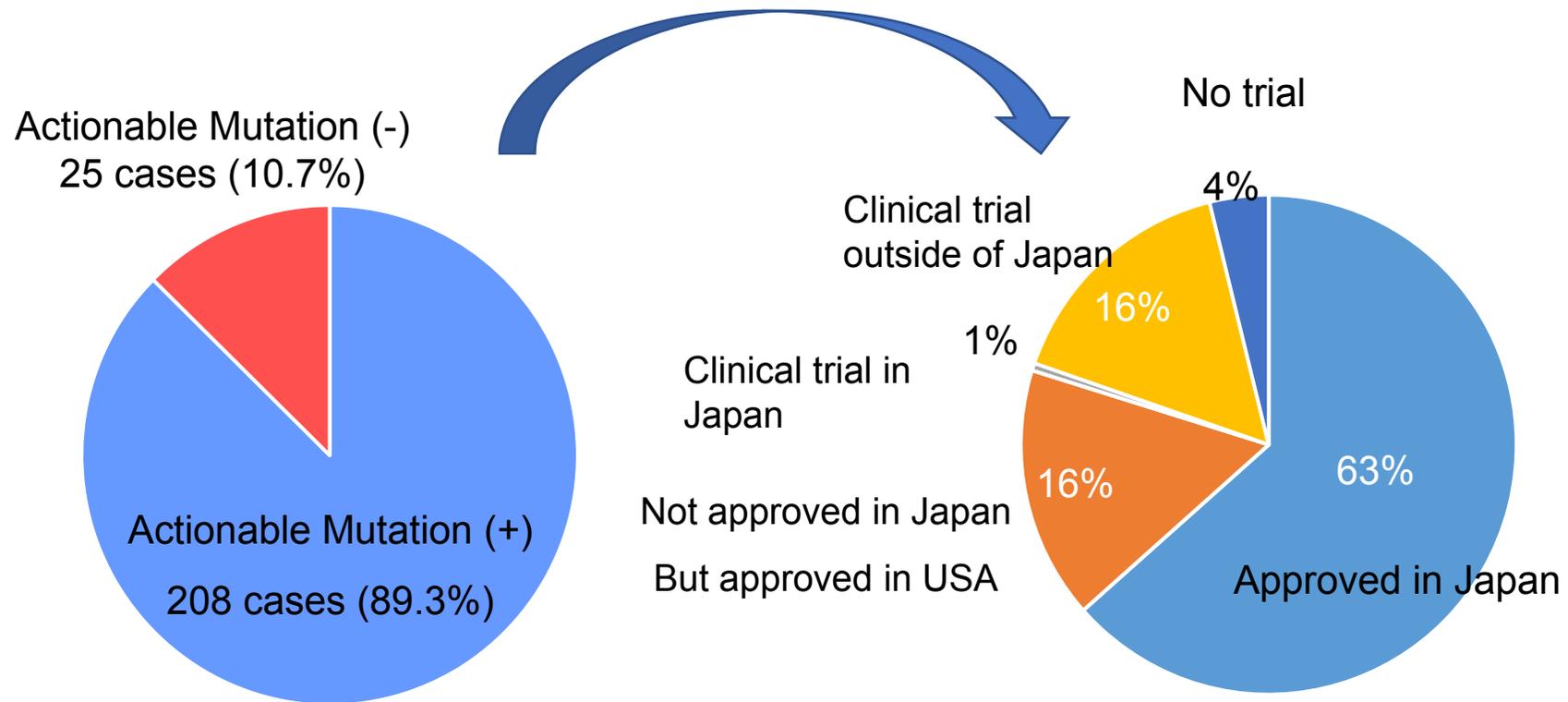
ALK *
BCR
ETV4
MLL *
RARA
BRAF *
EGFR *
ETV6
PDGFRB *
ROS1 *
ETV5
ETV1
EWSR1
RAF1 *
TMPRSS2
PDGFRA *
RET *

\*Genes marked with an asterisk are included in both lists.



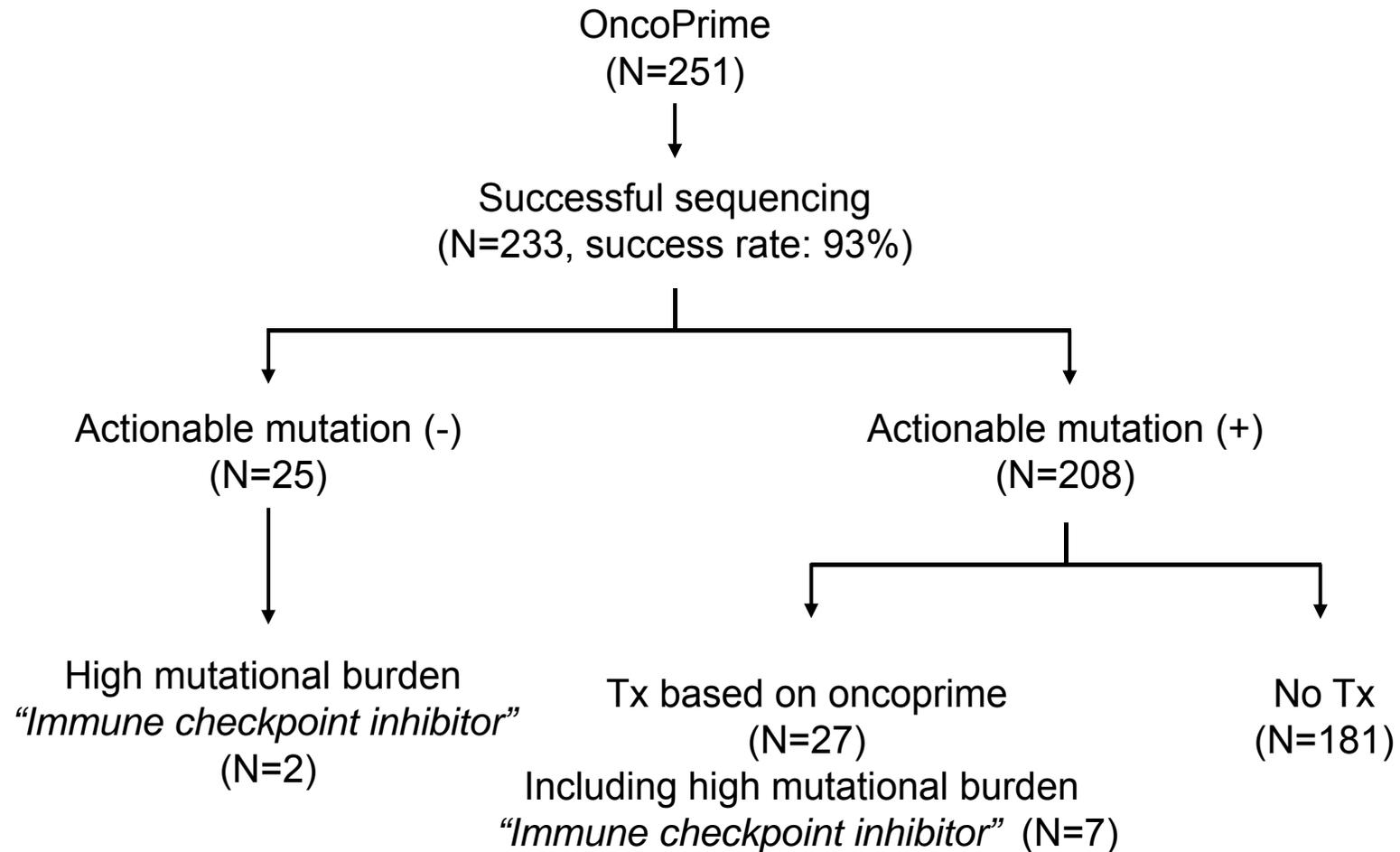
# Actionable Mutation and Druggable Mutation

2015 April ~ 2018 September (n=251)



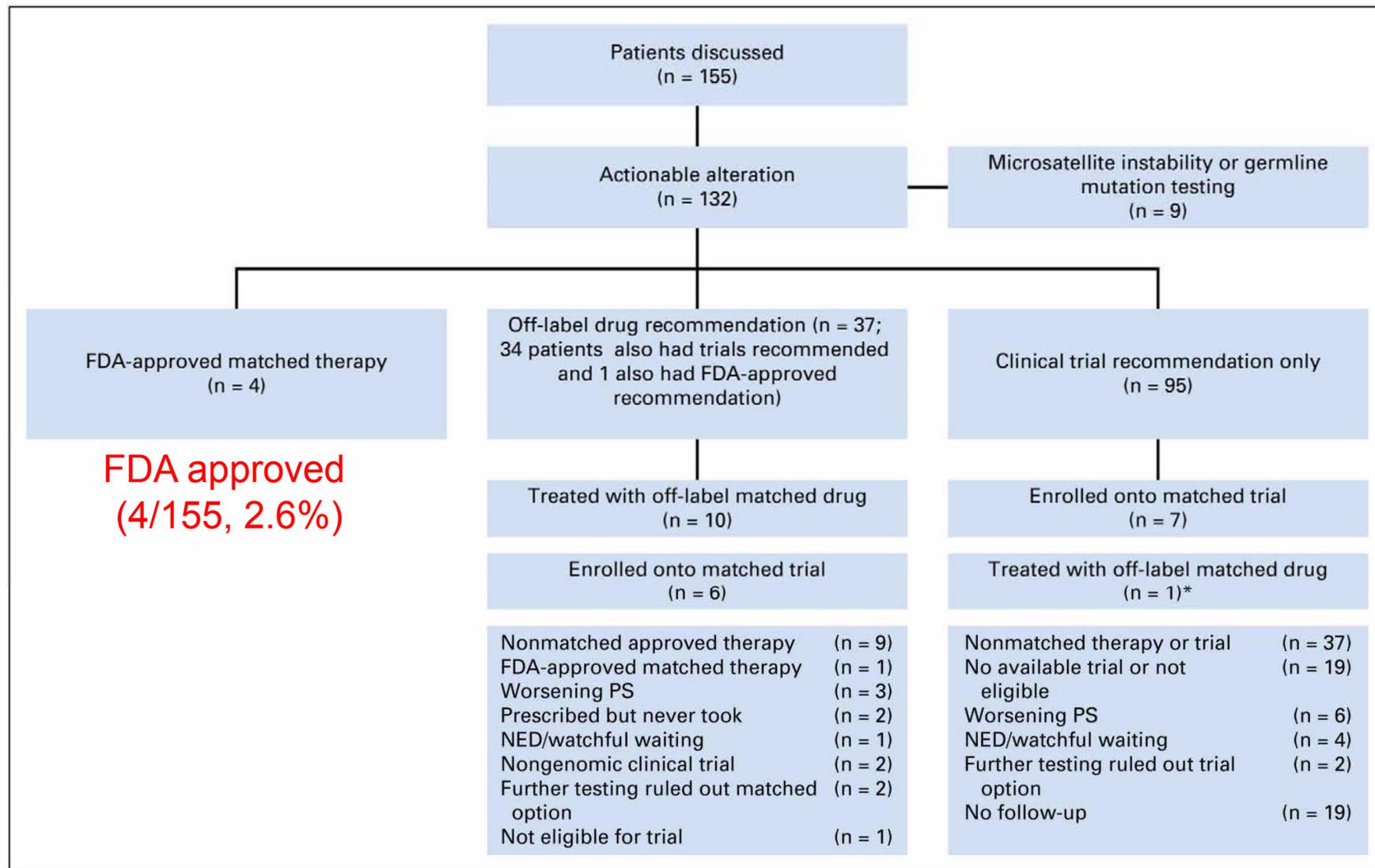
Most of them were OFF-label

# Clinical Flow after Sequencing



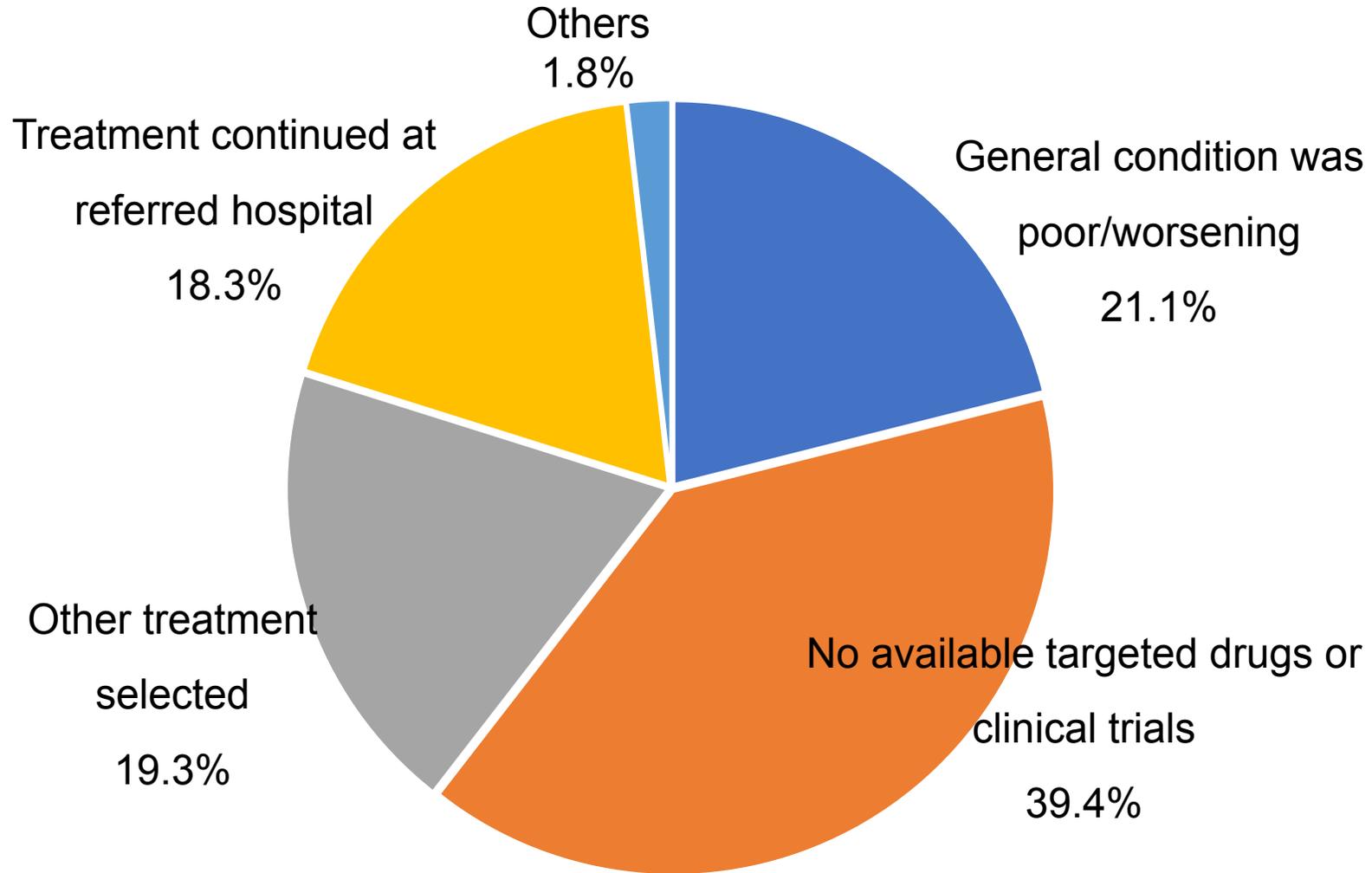
Pts administered Tx = 12.4% (29/233)

# (Reference) Personalized Medicine in the Oncology Clinic: Implementation and Outcomes of the Johns Hopkins Molecular Tumor Board



**Off-label (37/155, 24%)**

# Reasons why Therapies could not be Selected Based on Test Results



# Responders to Treatment

# Precision Cancer Medicine in Practice

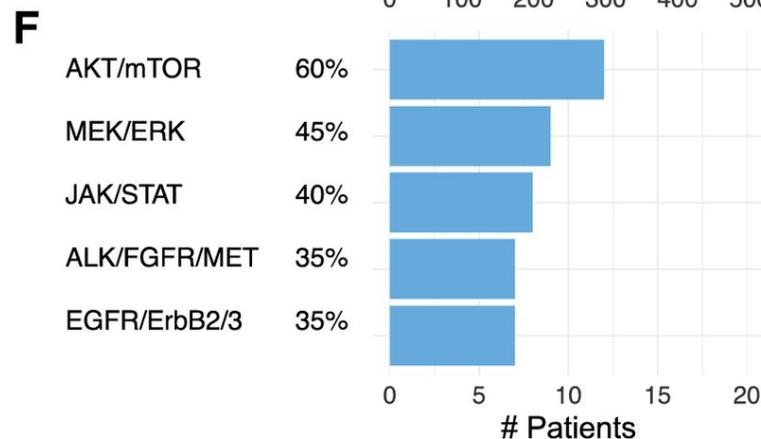
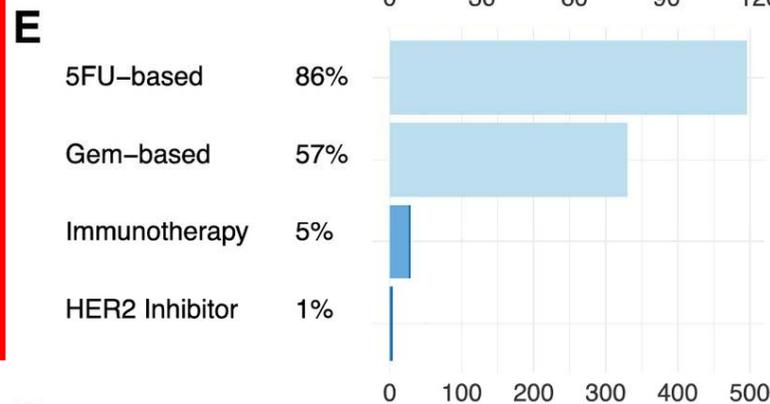
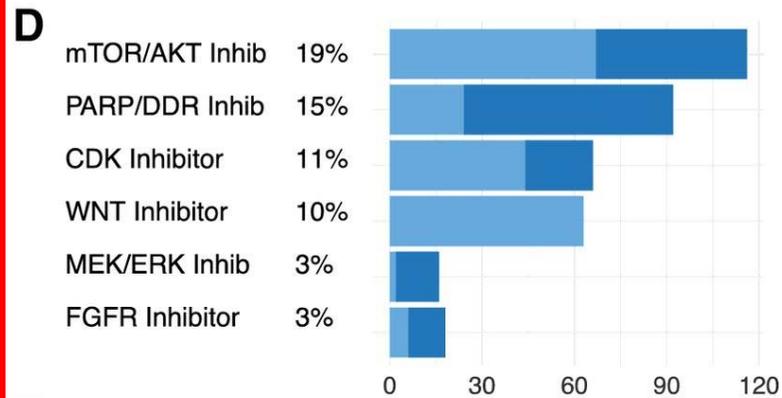
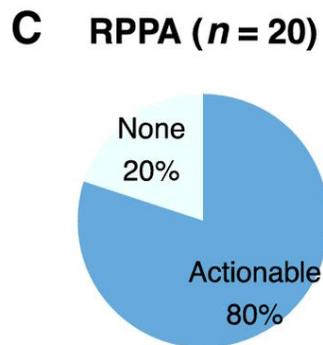
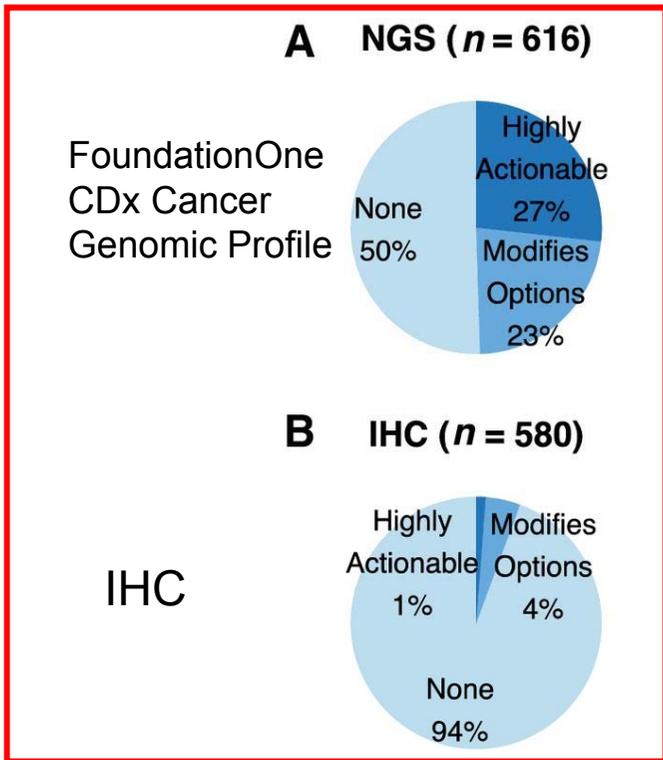
Treatment A won in a randomized clinical trial.  
Let's use this drug.



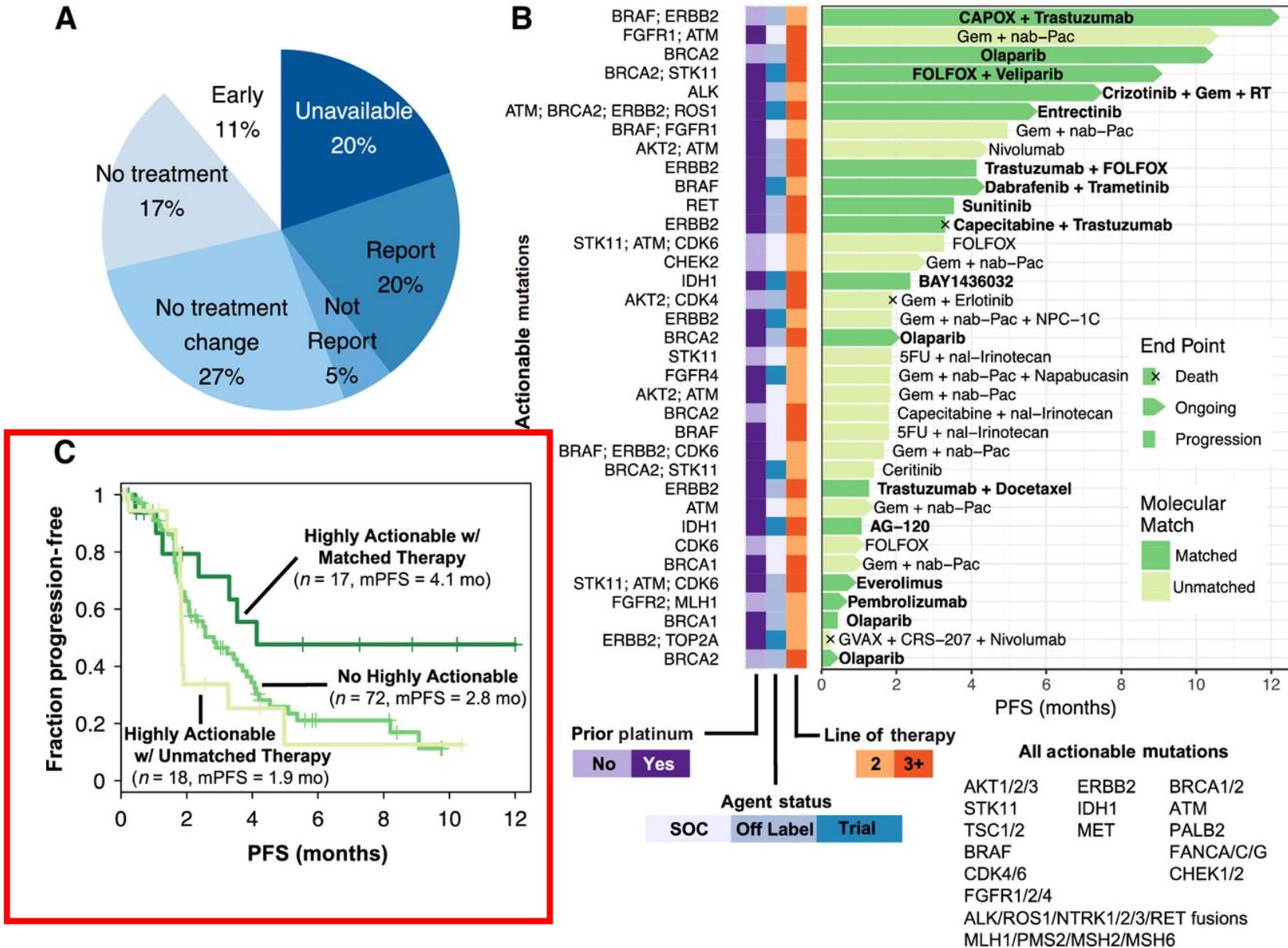
Your cancer was caused by a mutation in  
gene XX. So, let's start a treatment  
targeting this mutation.

# Comparison of Actionable Biomarkers in Pancreatic Cancer Patients

(99 % of results come from FoundationOne CDx Cancer Genomic Profile)

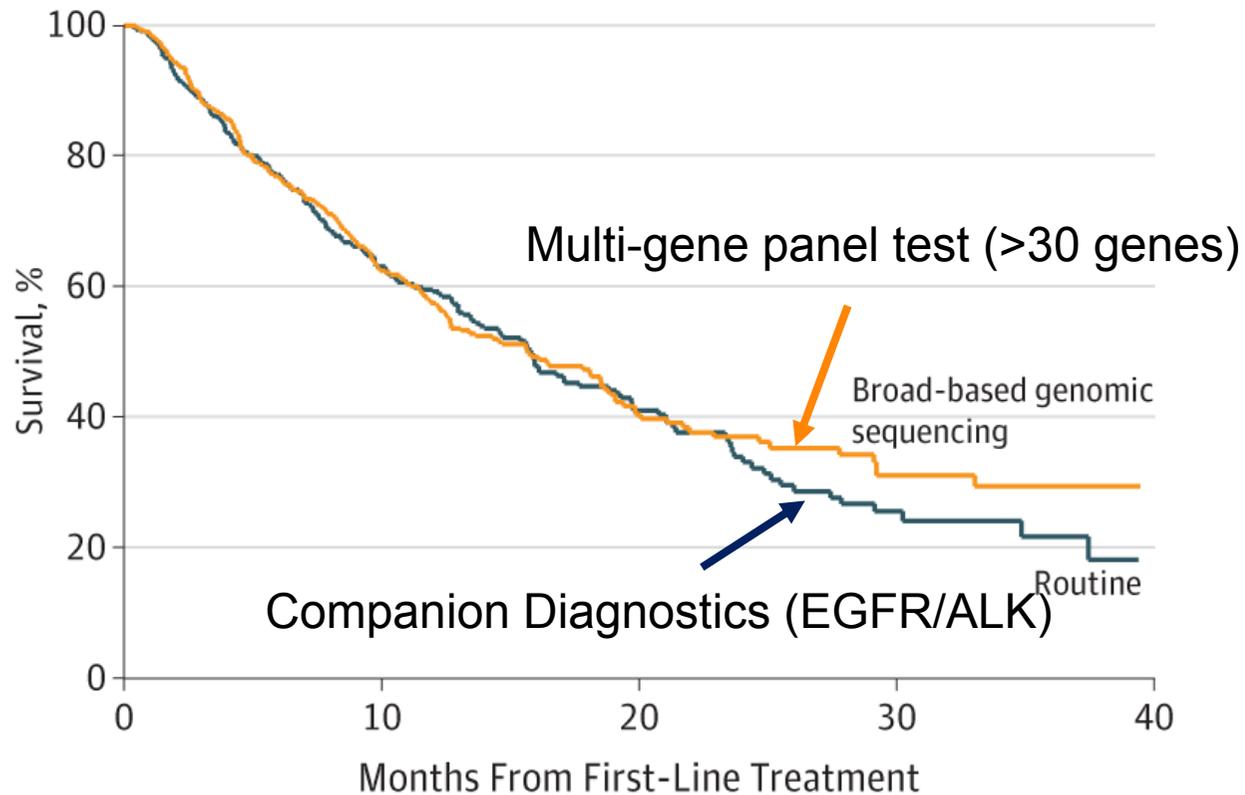


# Treatment Effect on Pancreatic Cancer Patients with Actionable Mutations



# Comprehensive Panel Testing and CDx Shows No Difference in Prognosis in NSCLC

From: Association of Broad-Based Genomic Sequencing With Survival Among Patients With Advanced Non-Small Cell Lung Cancer in the Community Oncology Setting



No. at risk						
Testing method						
Broad-based genomic sequencing	515	195	64	23	8	
Routine	513	192	66	18	<5	

# **Challenges to Cancer Genomic Medicine in Japan**

# Patients Who Can Use Genomic Testing under the NHI Scheme

Patients with solid tumors for which there are no standard therapies, and patients with locally advanced or metastatic solid tumors who have completed standard therapies (including those who are expected to complete standard therapies)

Patients for whom their treating physician judges to be suitable for chemotherapy based on the functional status of organs, their overall condition and etc, in accordance with relevant chemotherapy guidelines

**In the US, all patients with stage 3 or 4 cancer can use FoundationOne CDx under CMS coverage.**

The 415th Central Social Insurance Medical Council (General Meeting)  
<https://www.mhlw.go.jp/content/12404000/000513115.pdf> Accessed: 2019/7/3

Centers for Medicare & Medicaid Services Press release on March 16, 2018  
<https://www.cms.gov/newsroom/press-releases/cms-finalizes-coverage-next-generation-sequencing-tests-ensuring-enhanced-access-cancer-patients> Accessed: 2019/7/3

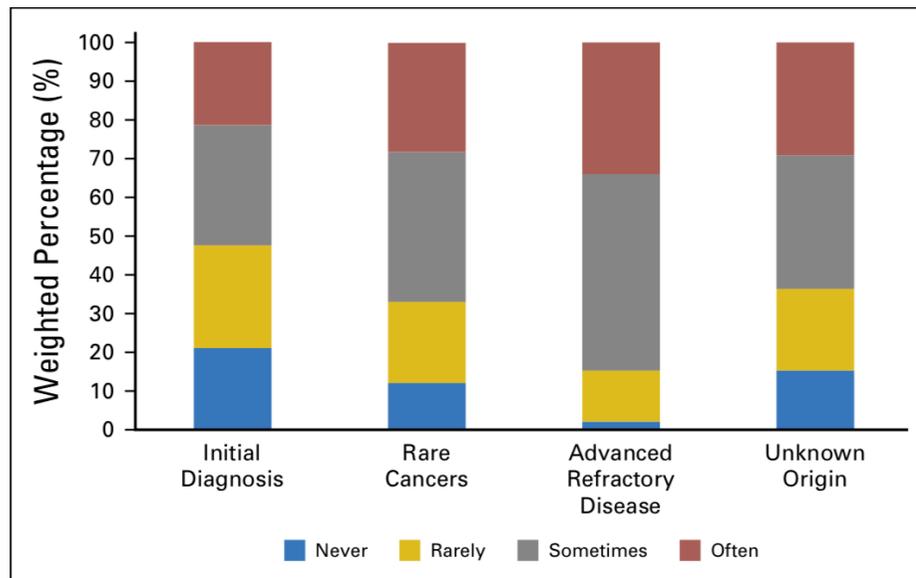
## Use of Next-Generation Sequencing Tests to Guide Cancer Treatment: Results From a Nationally Representative Survey of Oncologists in the United States

Research period

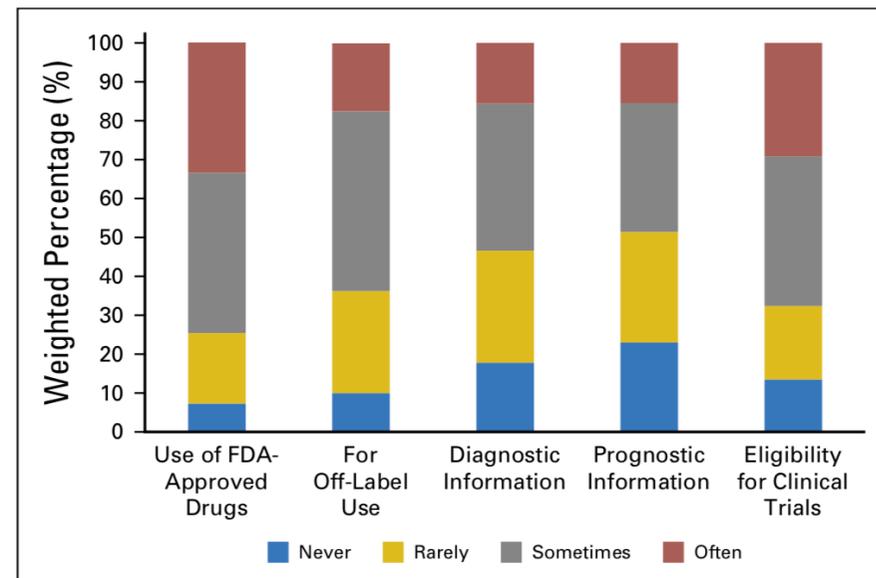
- one year, 2017
- conducted by email
- N=1,281

(cooperation rate=38%)

### When to use NGS?



### What to use NGS for?

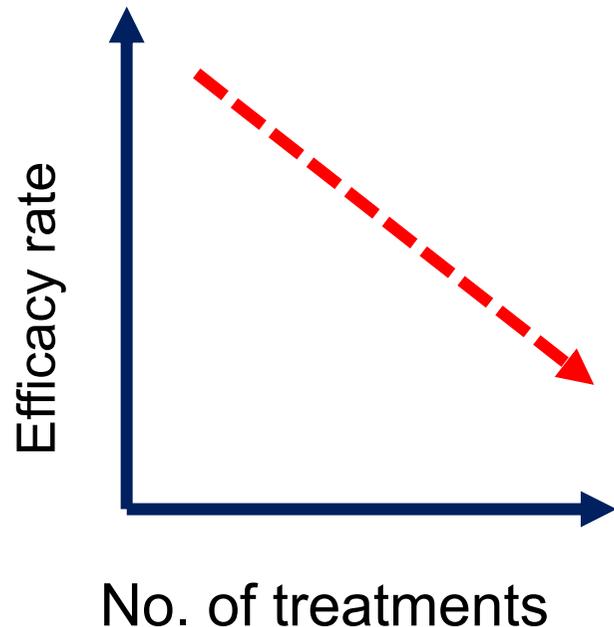


75.6% of oncologists who answered use NGS to determine treatment plan

# Eligibility for genomic testing in Japan

1. Cancers of unknown primary origin and Rare cancers
2. Cancers refractory to standard-of-care therapy

[Conceptual illustration]

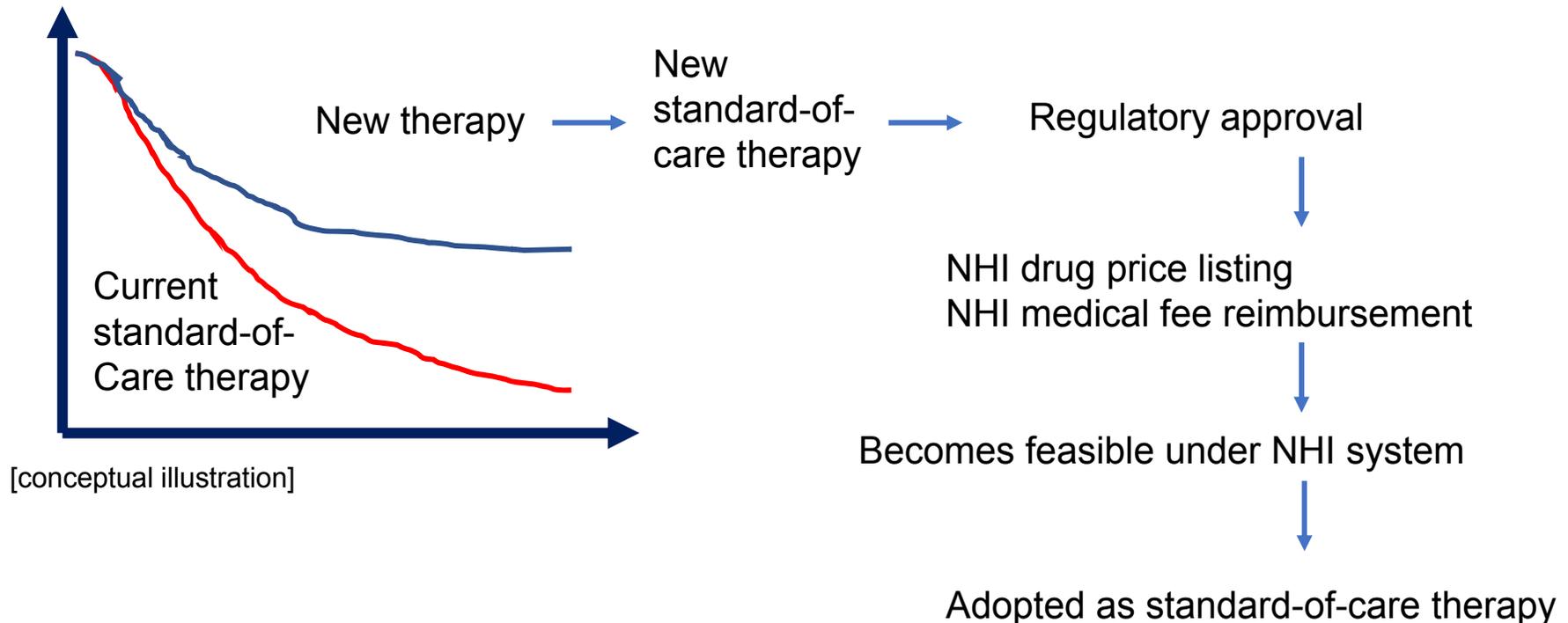


When a treatment is repeated, its effectiveness generally decreases (development of resistance).

The decision to administer a drug that is expected to be effective at the initial stage should therefore be viewed as a logical one.

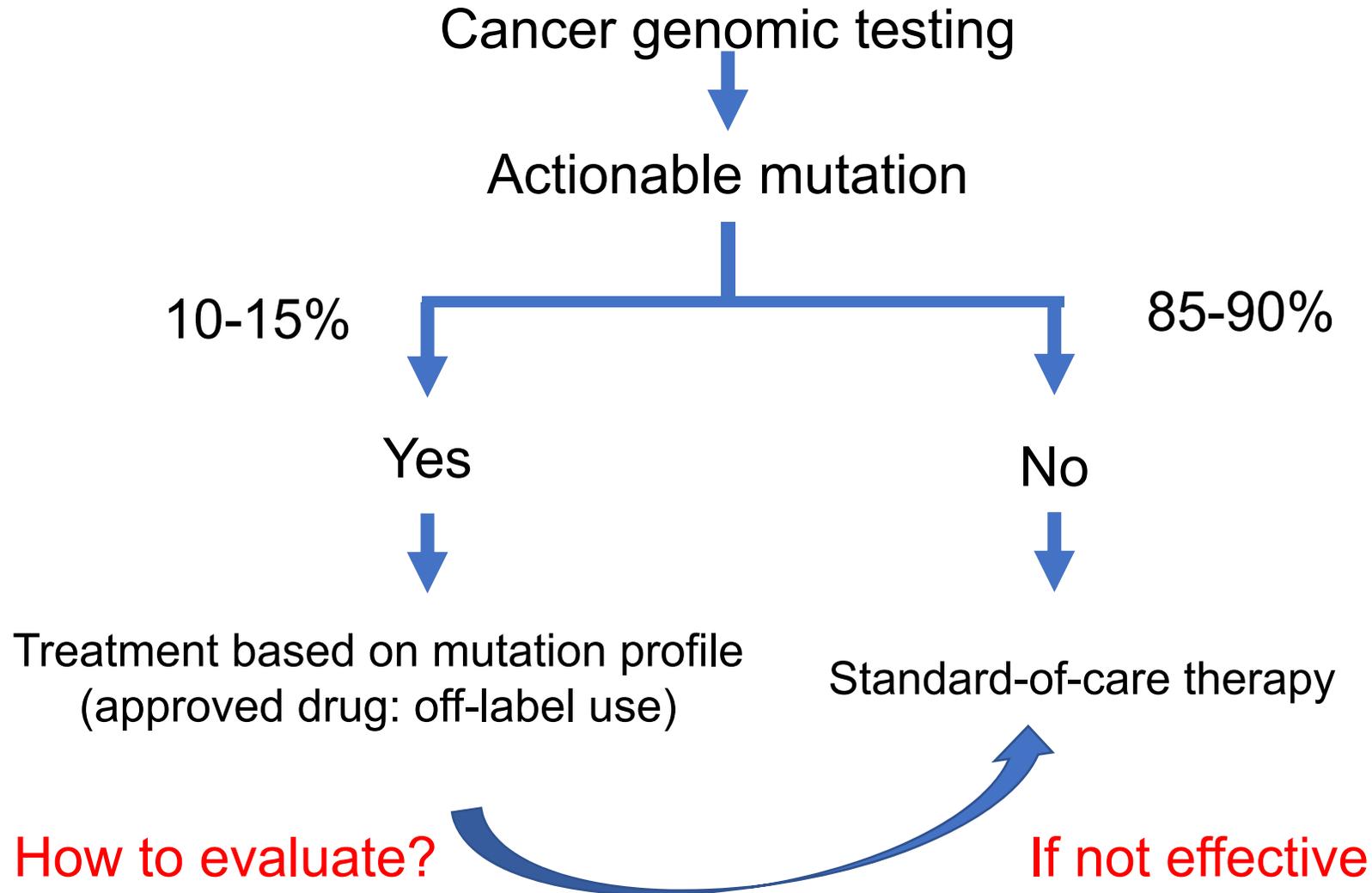
# So is it acceptable to perform genomic testing before initial treatment?

Standard-of-care therapy is determined by the results of large-scale Phase III clinical trials.



Even if a drug candidate is identified in genomic testing, there is no guarantee that it will become readily available

# Flow of Genomic Testing Used for Initial Treatment



# Eligibility for genomic testing in Japan

1. Cancers of unknown primary origin and Rare cancers
2. Cancers refractory to standard-of-care therapy

Is it possible to treat patients with targeted drugs under NHI coverage if genomic testing identified actionable alterations for patients whose cancers is unknown of its primary origin or for patients with rare cancers?

Eg.) Genomic testing identified ALK gene fusion in patients with cancer of unknown primary. Then, is it possible to treat patients under the diagnosis of ALK positive lung cancer?

Answer is NO.

Is it possible to treat such a patient based on MoA with detailed record of his symptoms?

# Self-contradiction in Precision Cancer Medicine

Precision cancer medicine is a treatment tailored to each individual



The results of our genomic testing have identified drugs that are a potential match for each individual!



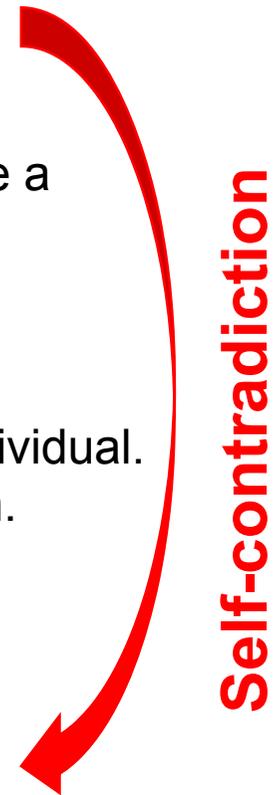
But we don't know if this drug is a good match for this particular individual.  
And we don't know if it will be effective in this particular organ.



We'll evaluate the drug in a clinical trial.  
It may have different effects on different organs



The question of whether a drug will be effective in a particular individual can only be answered by evaluating the drug in that individual.



# **Issues on Access to Treatment**

# Treatment-related Issue: How can Patients be Treated with Off-label Drugs?

	Patient briefing/ Specimen preparation	Sequencing	Expert meeting	Report drafting	Patient briefing	Treatment	R&D
Core hospital	Required	Required (Can be outsourced)	Required		Required	Required ※1	Required
Liaison hospital	Required	<ul style="list-style-type: none"> <li>Ask for Core Hospitals</li> <li>Participate in Core or Regional Hospital meetings, etc</li> </ul>			Required	Required ※2	Cooperation

MHLW guidance states that the use of off-label drugs based on genomic testing is anticipated in clinical trials and advanced medical treatments...

But do clinical trials & advanced medical treatments constitute precision medicine?

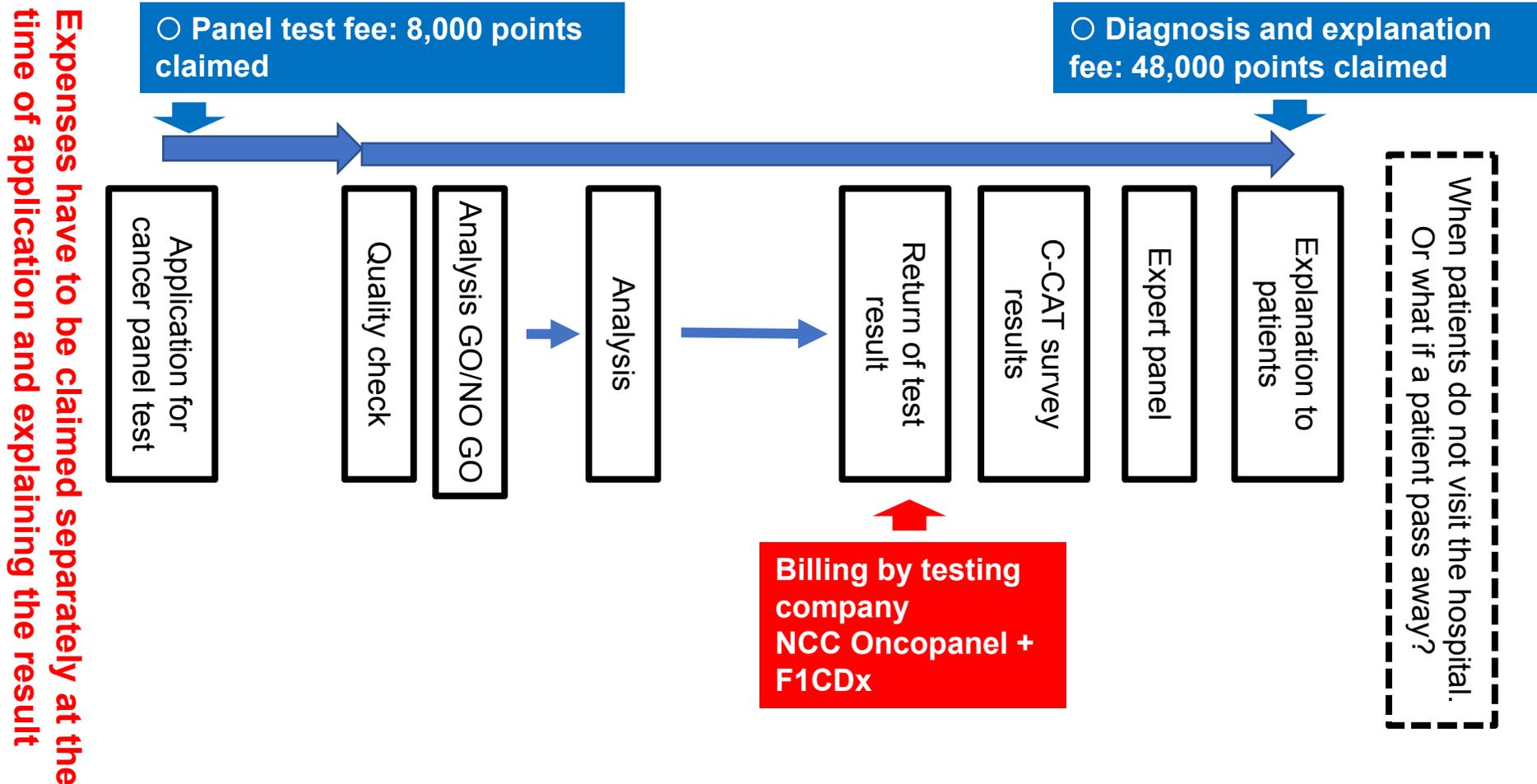
If such treatment is to be administered in a clinical study, it should be done in a 'basket trial' to obtain an additional indication for an approved drug.

# Issue in Treatment: How to Treat?

- Dealing with off-label use of drugs
  - ✓ For the time being, apply for off-label use on a case-by-case basis
  - ✓ In the future, it will be necessary to leverage programs such as compassionate use & 'patient-requested medical care' (although there are considerable obstacles because the usage would have to be based on clinical trials)
  - ✓ **Introduction of single patient IND**
- What about off-patent drugs or drugs without data protection period?
  - ✓ Companies would not be willing to conduct clinical trials for these drugs
- Should beneficiaries pay?
  - ✓ Dealing with test and treatment costs involving life insurance companies
  - ✓ Expansion of private insurance including non-covered medical treatments

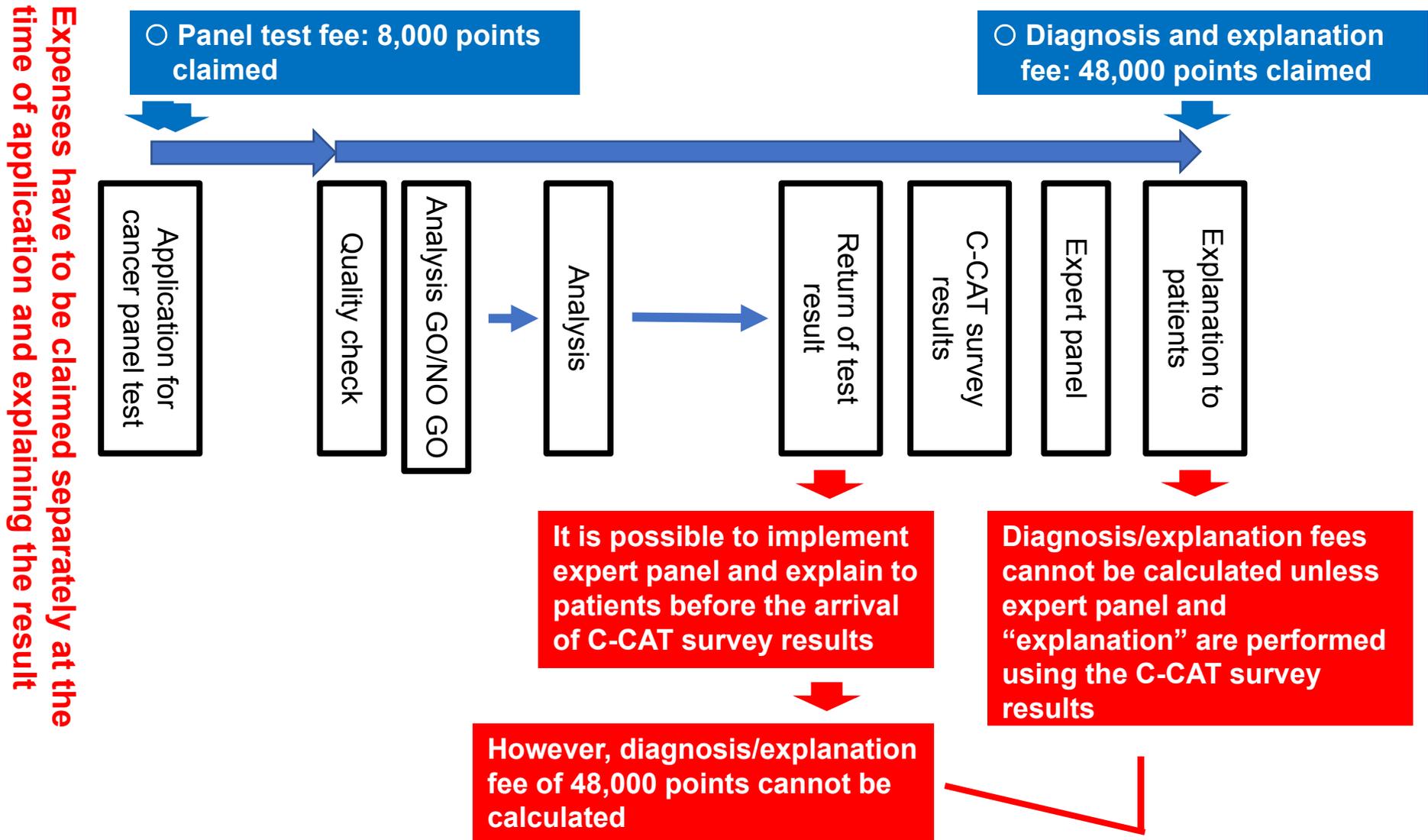
# **Timing Issues on Calculation of Medical Fees**

# Timing Issues on Calculation of Medical Fees



**It is important to select patients whose condition is unlikely to worsen after the tests**

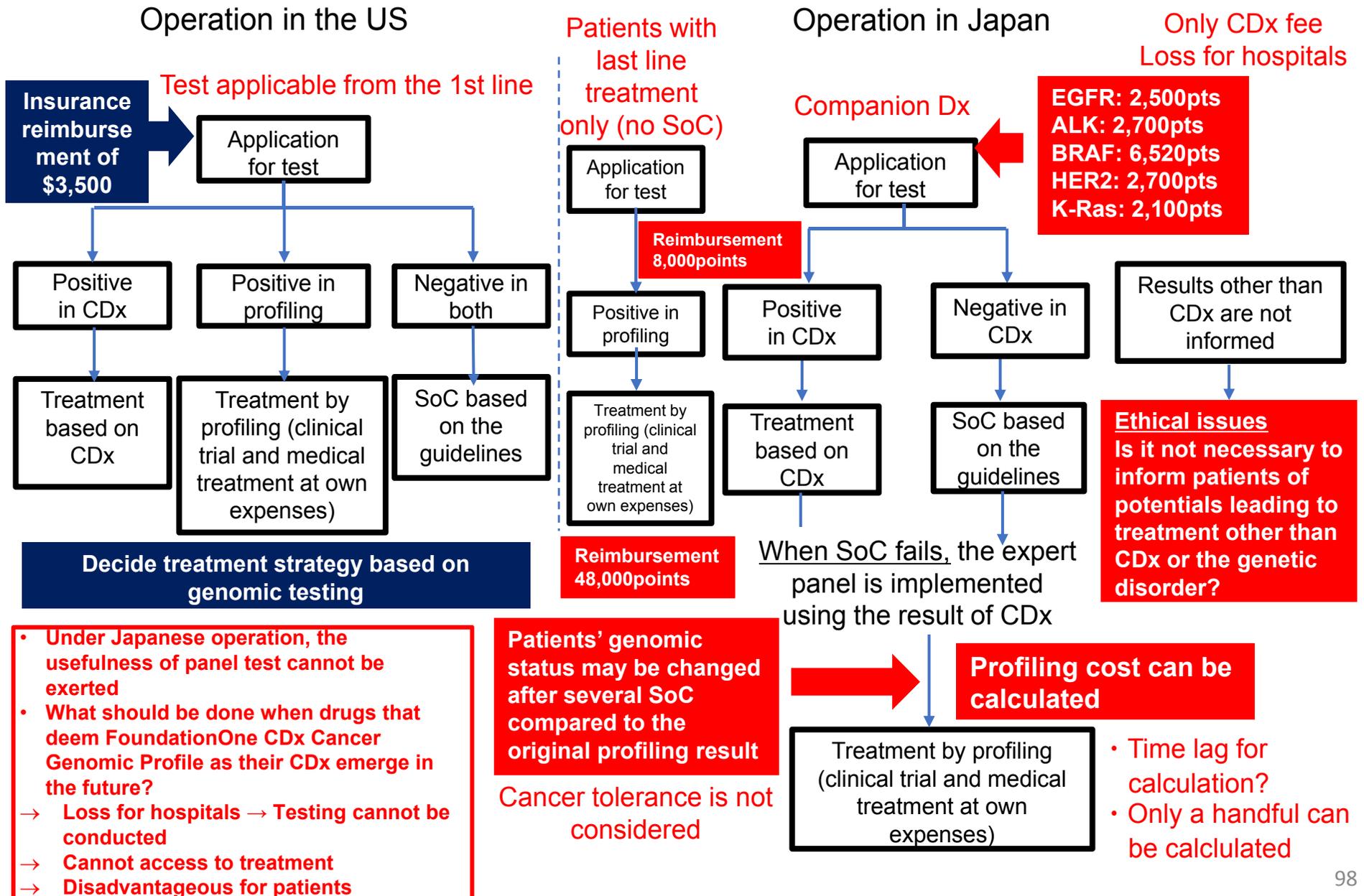
# Timing Issues on Calculation of Medical Fees



Should expert panel be implemented twice? 96

# **Challenges in the Use of FoundationOne CDx Cancer Genomic Profile as Companion Diagnostics**

# Timing issues with FoundationOne CDx Cancer Genomic Profile as Companion Diagnostics



# Take Home Messages

1. FoundationOne CDx is now available under the National Health Insurance scheme in Japan.
2. The time of Precision Medicine has come.
3. Further efforts are needed to bring an optimized treatment to patients faster.

**Thank you.**

# Disclaimer

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