



Burning Rock Biotech Limited

2Q2022 results

BNR US Equity
31 Aug 2022

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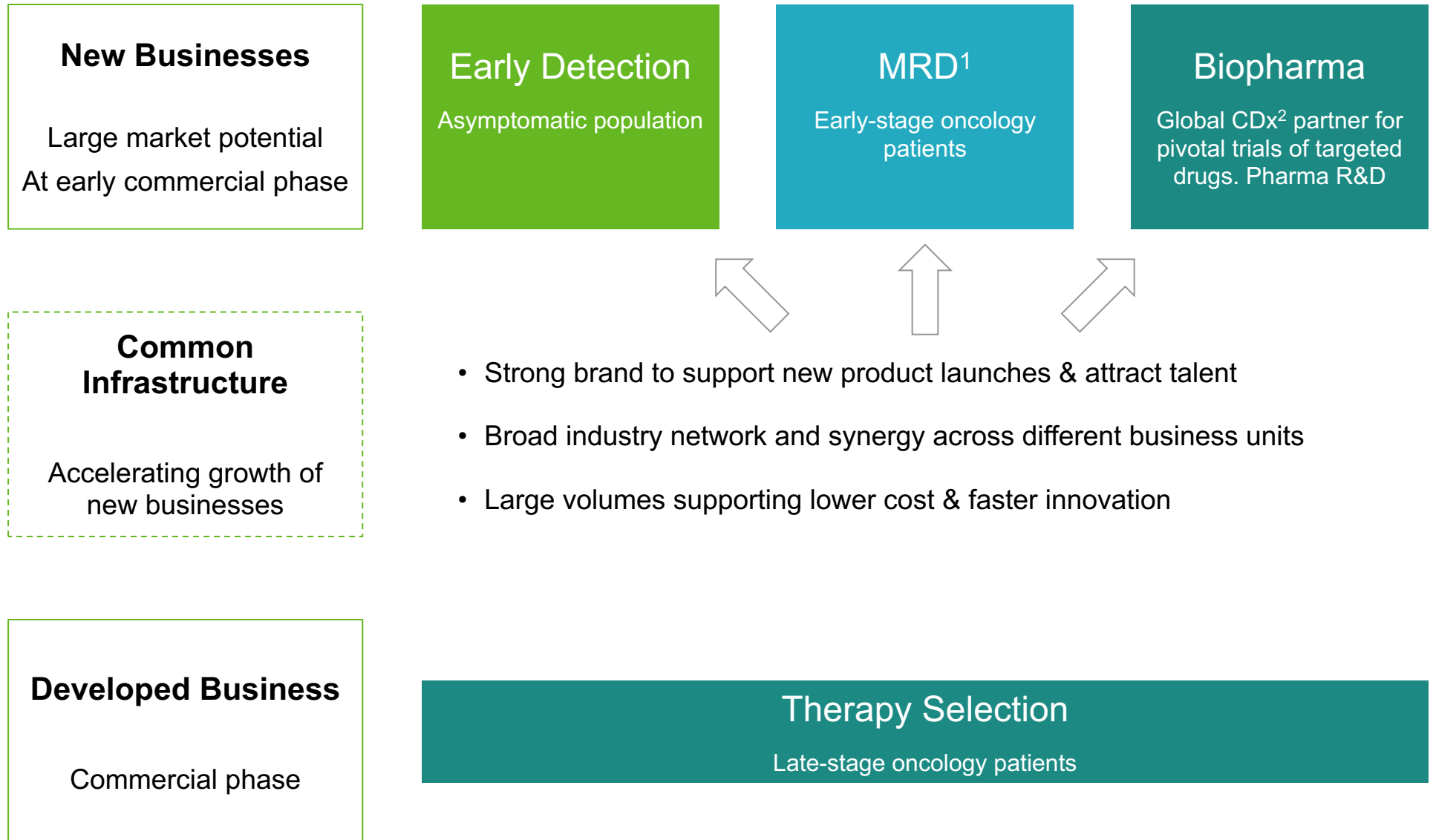
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Our value-building blocks

Extending leadership in NGS-based precision oncology from late-stage to earlier stage patients, opening up addressable market



Notes:

¹ Minimal residual disease of solid tumors

² Companion diagnostics

Objectives by segment

Continued topline growth with higher operating efficiency and reduced cash spend

Therapy selection

- Positive operating profitability in 2023
Through accelerated transition towards the profitable in-hospital channel and reduced opex in central-lab

MRD

- Multi-year, high double digit revenue growth, driving next leg of growth
Greenfield category, no gold standard from older technologies (e.g. PCR)
Indication expansion from NSCLC¹ to CRC², breast and other cancer types via additional clinical studies
Higher product entry barrier of *personalized* MRD test vs. *fixed-panel* products in therapy selection

Biopharma

- High double digit growth
Continued build-up of project backlog, leveraging Burning Rock's strength in quality and product performance
Already profitable due to high sales efficiency

Early detection

- Product – more cancer types, better performance
Incorporate additional signal sources, enrich machine-learning model through large (over 10k+ subjects) studies
- Regulatory – establish approval pathway
Dialogues with the NMPA and additional clinical studies to translate clear unmet need to proof of clinical utility
- Commercial – build first wave of seed customers
Working with a few large hospitals to build blood-based multi-cancer early detection into health check-up routines

Notes:

¹ Non-small cell lung cancer

² Colorectal cancer

Recent progress

+3% YoY revenue growth in 2Q despite severe Covid impact, driven by in-hospital growth outside of Covid impacted regions, MRD contribution and pharma revenues

Therapy selection

- Continued execution of growth via in-hospital – severe Covid impact in Shanghai and Beijing, but other regions combined grew +60% YoY in 2Q (by test volume)
- Team and opex optimization

MRD

- Strong commercial ramp post launch in Mar 2022, following data read-out on NSCLC¹ and CRC² at AACR
- Work underway to launch 2 interventional studies in 2022. Full MEDAL study data on NSCLC¹ expected to be released in 4Q22

Biopharma

- Revenue grew by triple digit YoY to RMB18m, contributing to 14% of overall revenues (up from 5% in 2Q21)
- Strong backlog build, with newly contracted project value +49% YoY to RMB158m during 7M22

Early detection

- Data release – PROMISE study (2,035 participants) for 9-cancer test completed, reading out at ESMO in Sep
- Clinical programs – PREVENT study launched (12,500 participants), China's first multi-cancer prospective interventional study

Notes:

¹ Non-small cell lung cancer

² Colorectal cancer

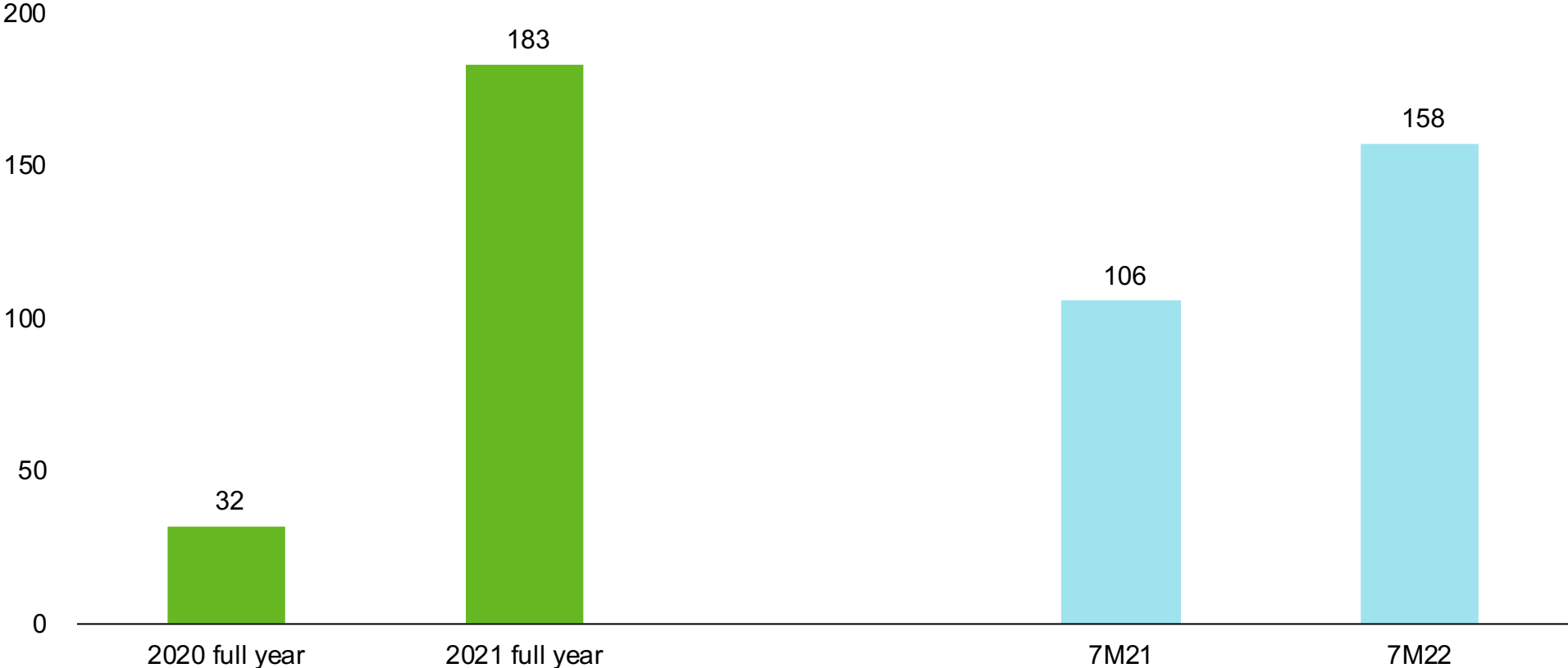
Biopharma services

Rapid backlog build-up continues

1H22 pharma revenues +207% YoY (to RMB18m, contributing to 14% of overall revenues)

Newly contracted pharma projects

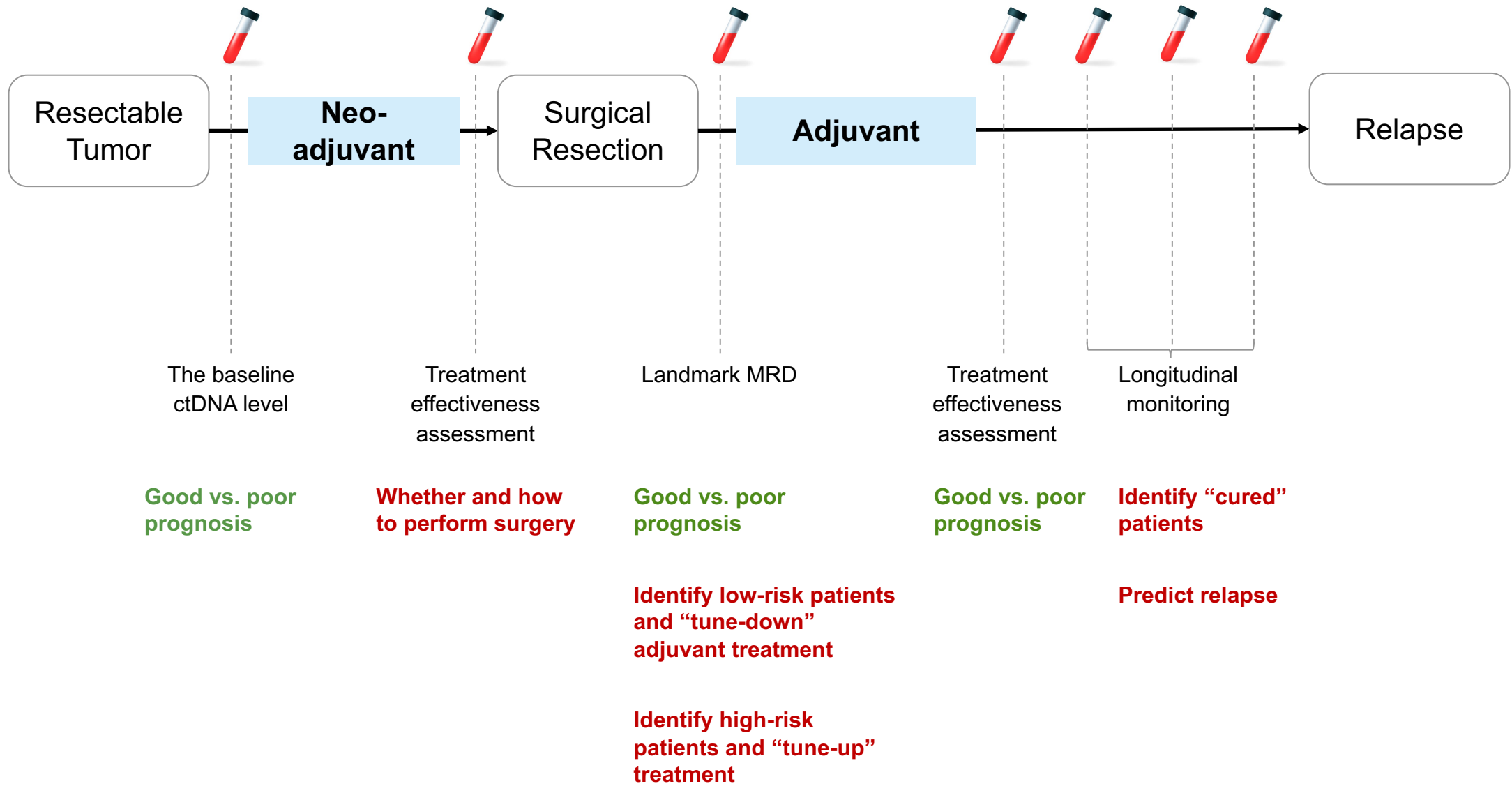
Contract value of new projects (RMB millions)





MRD

MRD test plays a role at multiple timepoints throughout the treatment journey



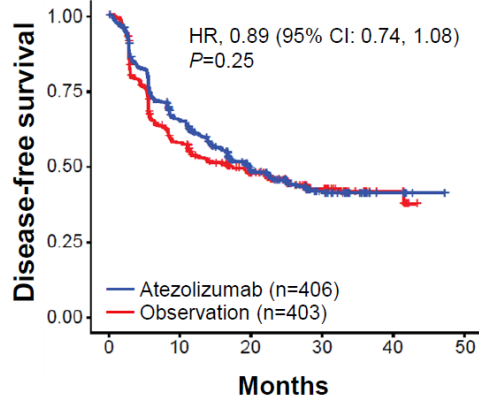
Nice-to-have prognosis

Actionable diagnosis that drives treatment choice

How do MRD studies advance utility

Example 1: IMvigor010, enrich the high-risk group and "tune-up" adjuvant treatment

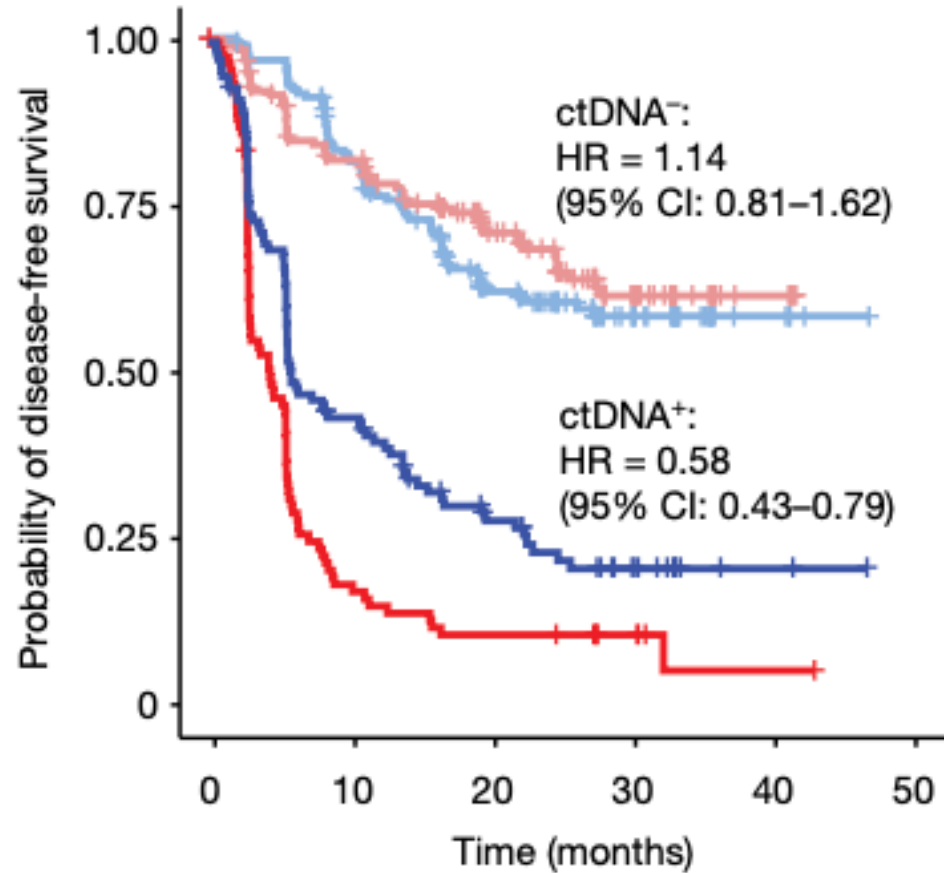
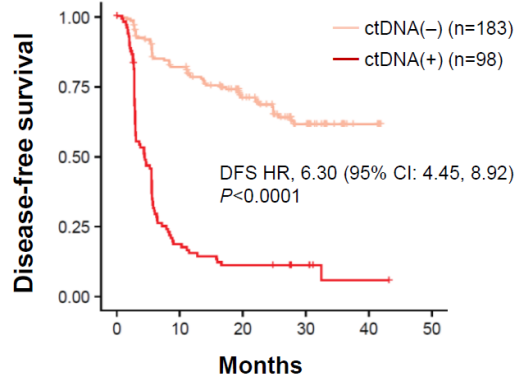
ITT



Atezo, MIUC Adjuvant Therapy
"All comers" demonstrated NO efficacy
TMB/PD-L1 showed NO prediction



Observation arm



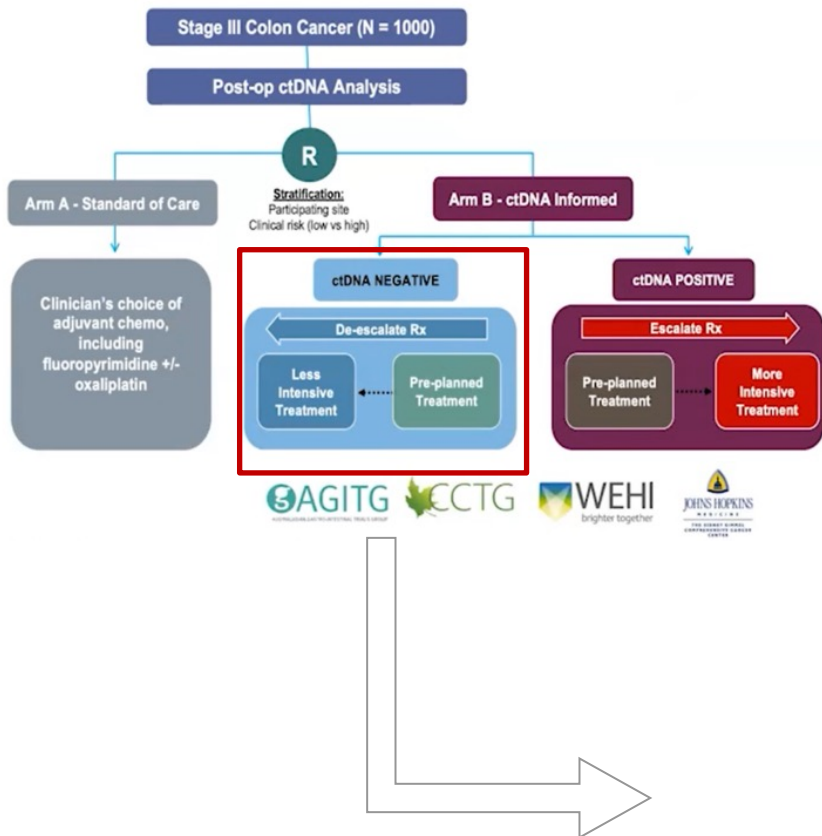
Indeed, only baseline MRD+ pts showed benefit

Landmark MRD+ pts (39%) had worse prognosis
Maybe only those patients can benefit?

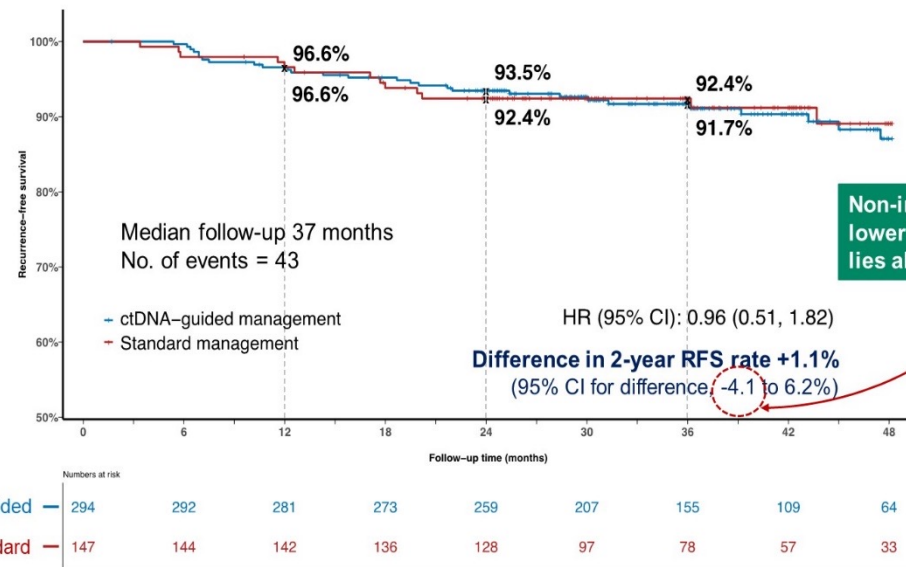
How do MRD studies advance utility

Example 2: Dynamic, identify low-risk patients and “tune-down” adjuvant treatment

DYNAMIC-III



ctDNA-guided vs standard



The ctDNA-guided MRD- group had fewer patients with adjuvant chemotherapy than the standard group (15% vs. 29%) with non-inferiority of 3-year RFS (92.4% vs 91.7%)

MRD clinical adoption through physician consensus

Chinese oncologists developing consensus on MRD applications in solid tumors, e.g. lung cancer

第18届中国肺癌高峰论坛 ——肺癌分子(微小)残留病灶(MRD)的检测和临床应用共识

共识一：MRD的概念

- 肺癌分子残留病变，指的是经过治疗后，传统影像学(包括PET/CT)或实验室方法不能发现，但通过液体活检发现的癌来源分子异常，代表着肺癌的持续存在和临床进展可能；
- 肺癌分子异常：指的是在外周血可稳定检测出丰度 $\geq 0.02\%$ 的ctDNA，包括肺癌驱动基因或其他的 I / II类基因变异。

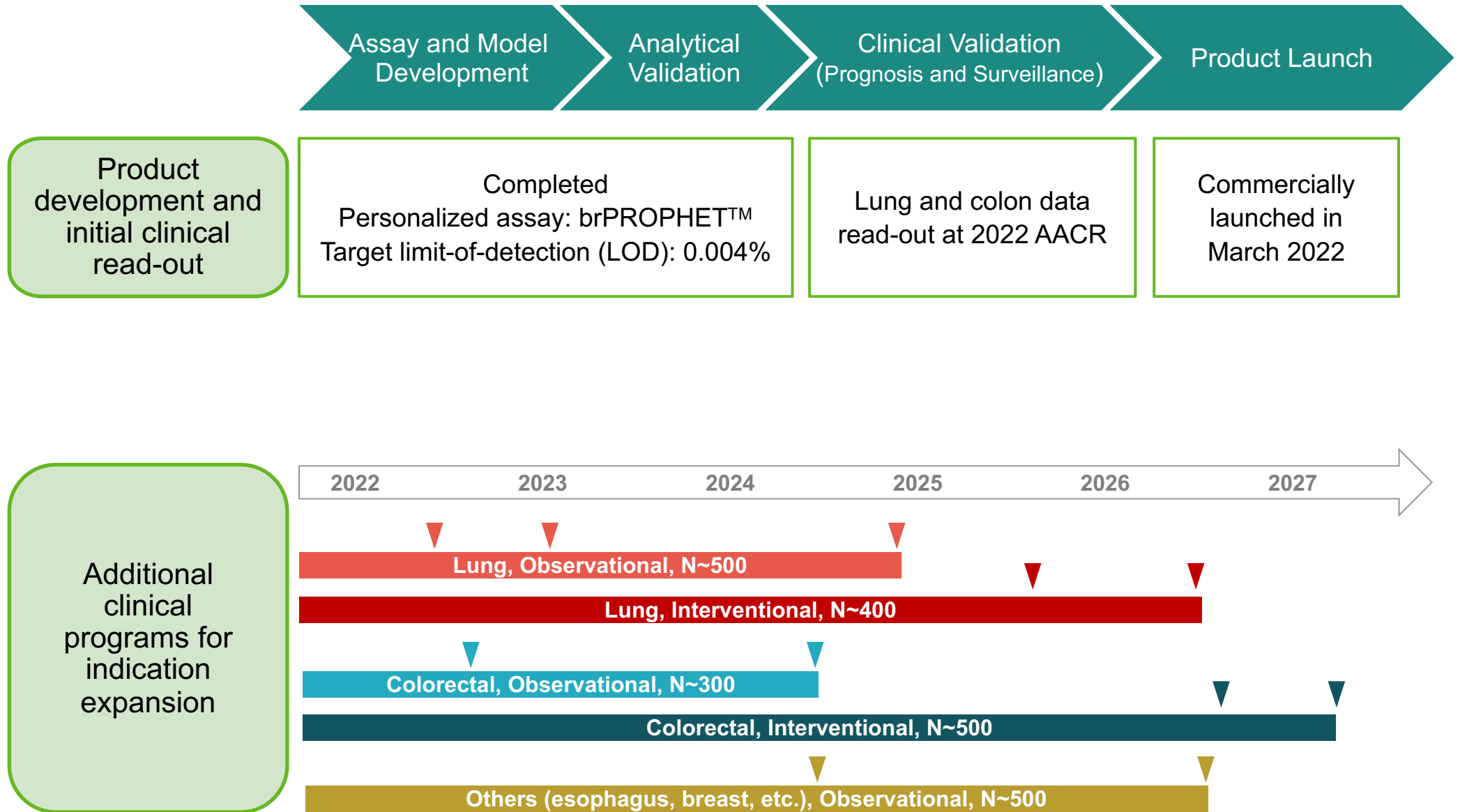
共识二：MRD检测的基本技术要求

- MRD检测的基本技术，包括Tumor-informed assays(个体化定制)和 Tumor agnostic assays(NGS panel和多组学技术)，目前均处在探索阶段，需要前瞻性研究确定其敏感性、特异性和预测价值；
- 采用二代测序技术(NGS)，所选的多基因 panel中必须覆盖患者 I / II类基因变异，基本技术标准是可稳定检出丰度 $\geq 0.02\%$ 的ctDNA；
- 驱动基因阳性的非小细胞肺癌，MRD的分子panel应包括该驱动基因；
- MRD评估报告中必须包括cfDNA丰度， ctDNA丰度，所检测基因VAF值；
- 需要建立针对免疫治疗的MRD标准。

Burning Rock development plans

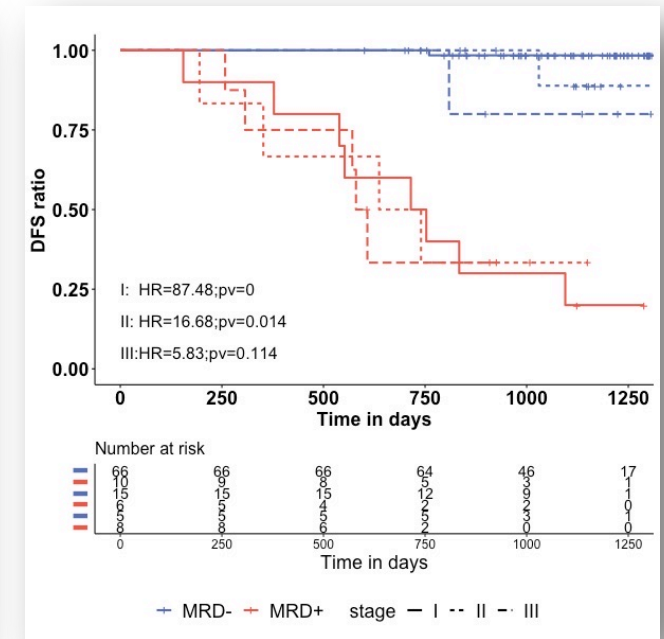
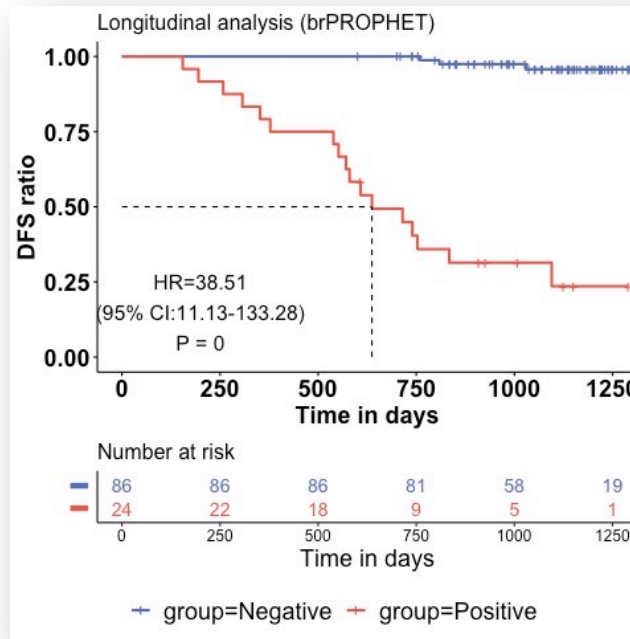
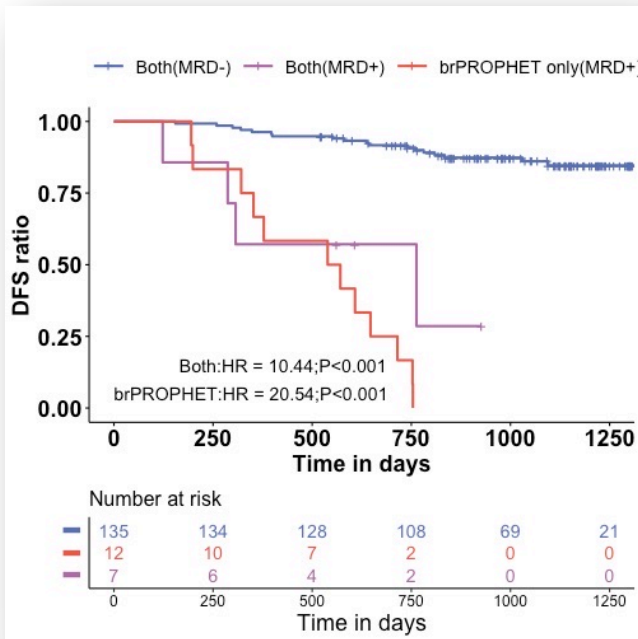
Personalized approach (brPROPHET™) demonstrating strong analytical performance

Additional clinical studies to expand indications



MRD clinical validation data readout

NSCLC – MEDAL study



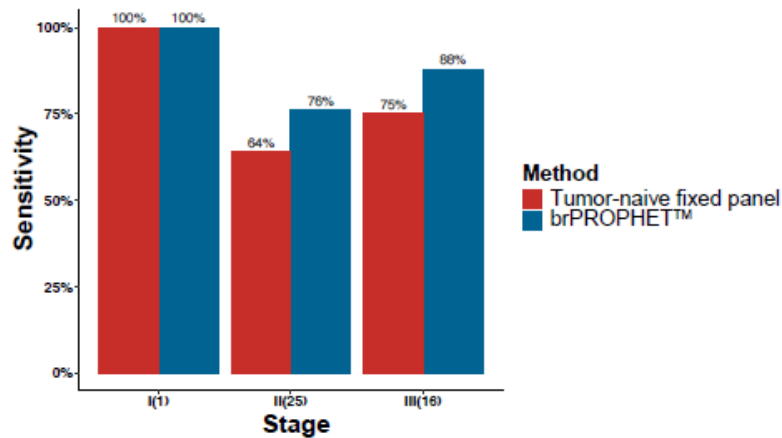
- *brPROPHET identified 2.7 times more true high-risk patients than the fixed panel approach at the landmark time point*
- *Longitudinally MRD negative patients has near-perfect prognosis with median of 3-year follow-up*
- *The prognosis differentiation holds true for patients with different clinical stage*

MRD clinical validation data readout

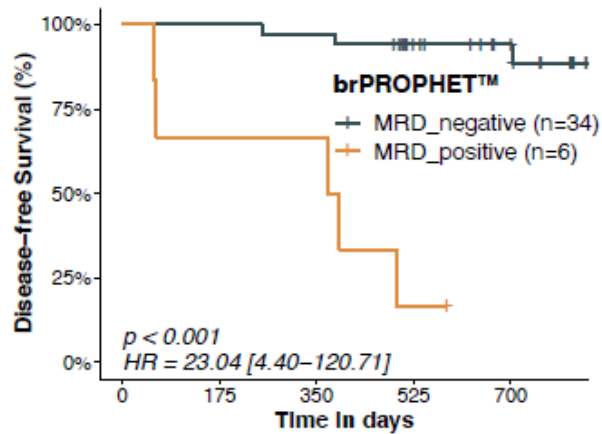
CRC

Session OPO.PR02.01 - Clinical Prevention, Early Detection, and Interception

5917 - Patient-specific tumor-informed circulating tumor DNA (ctDNA) analysis for postoperative monitoring of patients with stages I-III colorectal cancer (CRC)

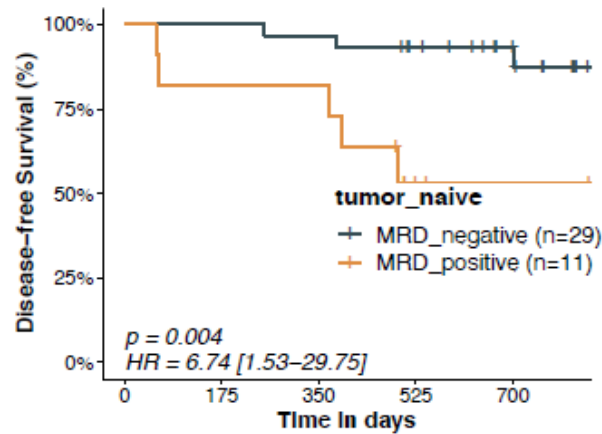


brPROPHET™ demonstrated superior sensitivity and specificity to fixed panel in pre-operative ctDNA detection and post-operative MRD calling among relapsed patients



Number at risk

—	34	34	33	25	17
—	6	4	4	1	0



Number at risk

—	29	29	28	24	16
—	11	9	9	2	1



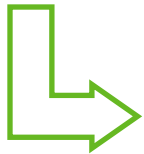
Early detection

Product development since 2016

Demonstrated high specificity and tissue-of-origin detection capability

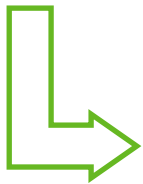
Proof-of-concept
2016 – 2019

- Proof of concept on our methylation based, machine learning aided technology platform
- Results published on *Nature Biomedical Engineering*, “Ultrasensitive detection of circulating tumour DNA via deep methylation sequencing aided by machine learning”



3-cancer
2017 – 2020

- Lung, Colorectal Cancer (CRC), Hepatocellular Carcinoma (HCC)
- 95.1% specificity and 80.8% sensitivity¹



Product development complete

6-cancer
2018 – Nov 2020

- Lung, CRC, HCC, Ovarian, Pancreatic, Esophageal
- 98.9% specificity and 69.1% sensitivity²



Product development in progress

9-cancer
2019 – Ongoing

- Additional cancer types: Gastric, Biliary Tract, Head & Neck
- PROMISE study concluded, reading out at ESMO in Sep 2022
- Ongoing PREDICT study

22-cancer³
2020 – Ongoing

- BR-22 covers 88% of China’s cancer incidence
- Ongoing PRESCIENT study

Notes:

¹ Training and validation cohorts combined, 490 cancer samples, 226 control samples. Sample size is aggregated through a series of case-control studies. 95.1% specificity (95% CI 91.2-97.4) and 80.8% sensitivity (95% CI 77.0-84.1)

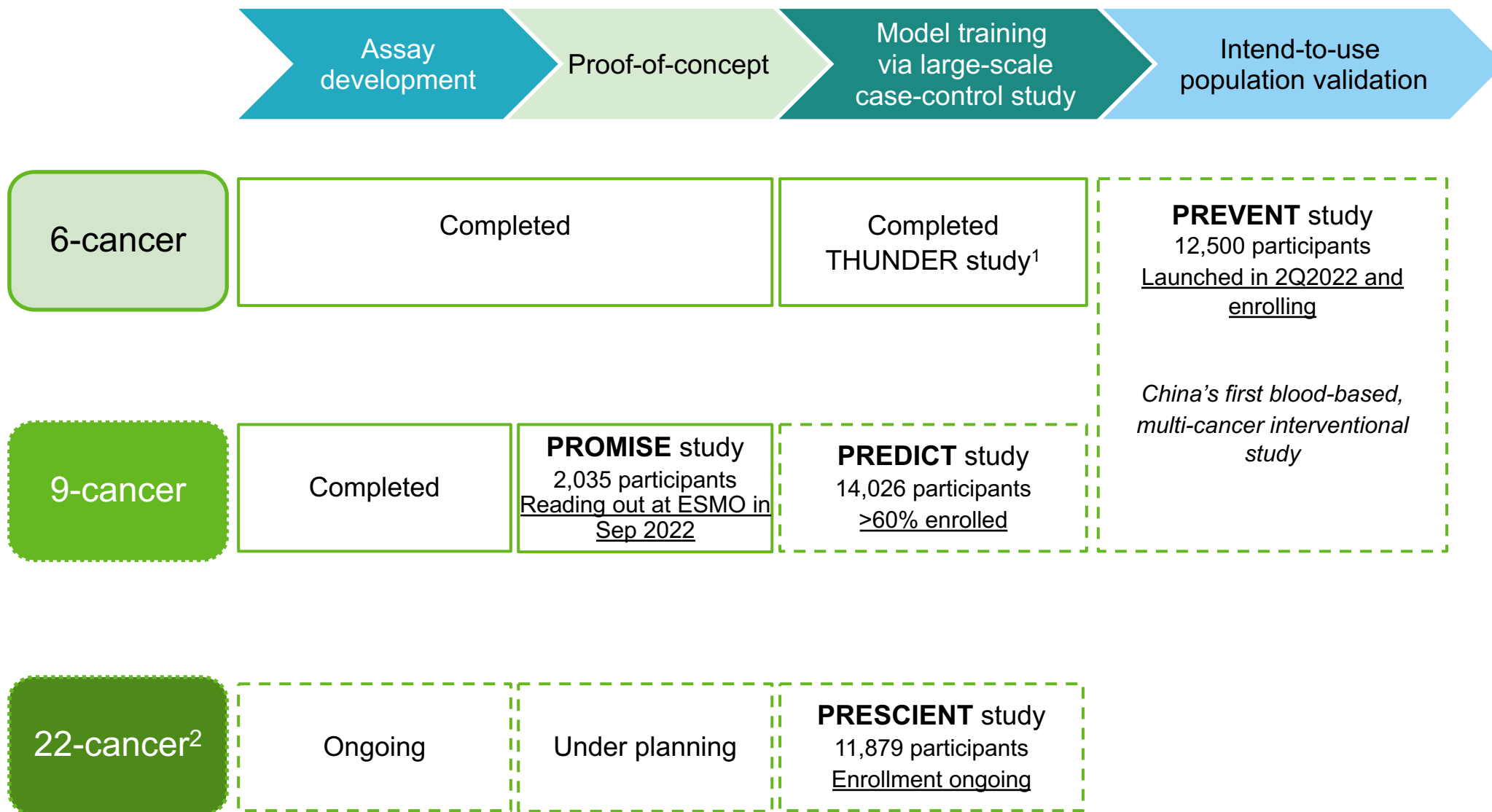
² Unintrusive multi-cancer detection by circulating cell-free DNA methylation sequencing (THUNDER); development and independent validation studies, ASCO 2022. Further details in Appendix 1.

³ Final number of cancer types subject to development progress

Clinical programs

9-cancer development first read-out in Sep (PROMISE study)

China's first interventional study for multi-cancer launched in 2Q (PREVENT study)



Notes:

¹ THUNDER series of studies. Latest results presented at ASCO 2022, Unintrusive multi-cancer detection by circulating cell-free DNA methylation sequencing (THUNDER): development and independent validation studies

² Final number of cancer types subject to development progress

Burning Rock's early detection technology

Globally competitive technology with multi-cancer validation

Competitive technology

Methylation + machine learning to overcome challenges of low ctDNA abundance, leading to feasibility of multi-cancer early detection

Multi-cancer validation data

nature
biomedical engineering

ARTICLES

<https://doi.org/10.1038/s41551-021-00746-5>



Ultrasensitive detection of circulating tumour DNA via deep methylation sequencing aided by machine learning

VIRTUAL 2020 ESMO ASIA

Early detection and localization of multiple cancers using a blood-based methylation assay (ELSA-seq)

AACR 2022

Session OPO.CL11.01 - Biomarkers

5116 - Analytical performance of ELSA-seq, a blood-based test for early detection of multiple cancers

Session OPO.CL11.01 - Biomarkers

5109 - Development of cfDNA reference standards for methylation-sequencing tests

ASCO 2022

Clinical validation of a multicancer detection blood test by circulating cell-free DNA (cfDNA) methylation sequencing: The THUNDER study.

ESMO 2022

The performance of a multi-cancer early detection model based on liquid biopsy of multi-omics biomarkers: A proof of concept study (PROMISE study)

Leadership in multi-cancer early detection

First-in-class, high entry-barrier, multi-year effort

Challenges

BNR position

1

Technology

Low amount of cancer signal

in the circulating bloodstream, much more challenging vs. tissue

Proprietary chemistry and algorithm

- On par with global leader, competitive sensitivity in earlier stages for certain cancers
- Multi-year lead vs. China peers (most showing liver-cancer and colon-cancer data only)

2

Clinical

Large, multi-year studies required

from case-control to intend-to-use population, from observational to interventional (e.g. CCGA study: 15,254 participants, 8,584 with cancer, 6,670 without cancer)

Sponsorship from top physicians

- Catching up with global leader, to improve specificity and tissue-of-origin performance through large clinical studies
- Multi-year lead in China as the only company with studies over 10,000+ subject scale launched

3

Regulatory

First-in-class in nature

with no established regulatory pathway

Leading regulatory capability in China

- Exploring possible pathway, leveraging experience through the country's first NGS kit approval by the NMPA

4

Commercial

Unprecedented product

Multi-pronged approach

- Initially working with hospital health check-up departments, leveraging synergy from in-hospital therapy selection business

Leadership from top-tier principal investigators key to clinical success

Also drives increasing recognition on multi-cancer early detection among clinicians

PREDICT



- Leading site: Shanghai Zhongshan Hospital
 - One of China's largest comprehensive academic hospitals
 - Performs c.104,000 operations and serves c.169,000 inpatients and over 4,236,000 outpatients on an annual basis¹
 - Ranked top 5 in the 2019 China's general hospital rankings²

Principal Investigator: Prof. Jia Fan



- Fellow of the Chinese Academy of Sciences
- President of Shanghai Zhongshan Hospital

PRESCIENT



- Leading site: Cancer Hospital of the Chinese Academy of Medical Sciences³
 - The first and top cancer-specialist hospital in China
 - The National Clinical Center for Cancer Research, the National Center for Quality Control on Standardized Cancer Treatment and Diagnosis, the National Clinical Center for Drug Research

Principal Investigators

Prof. Jie He



Prof. Jie Wang



- Fellow of the Chinese Academy of Sciences
- President of CHCAMS
- Head of the Dept. of Medicine, CHCAMS

PREVENT



四川大学华西医学中心
WEST CHINA MEDICAL CENTER OF SICHUAN UNIVERSITY

- Leading site: West China Hospital
 - One of the largest hospitals in China, performed 196,000 surgeries and 7.8 million out-patient services in 2021
 - Ranked #2 in the Fudan Best Hospital in China Rankings (2009-2020)

Principal Investigator: Prof. Weiming Li



- President of West China Hospital

Notes:
¹ Based on 2018 statistics
² <http://rank.cn-healthcare.com/rank/general-best>
³ CHCAMS

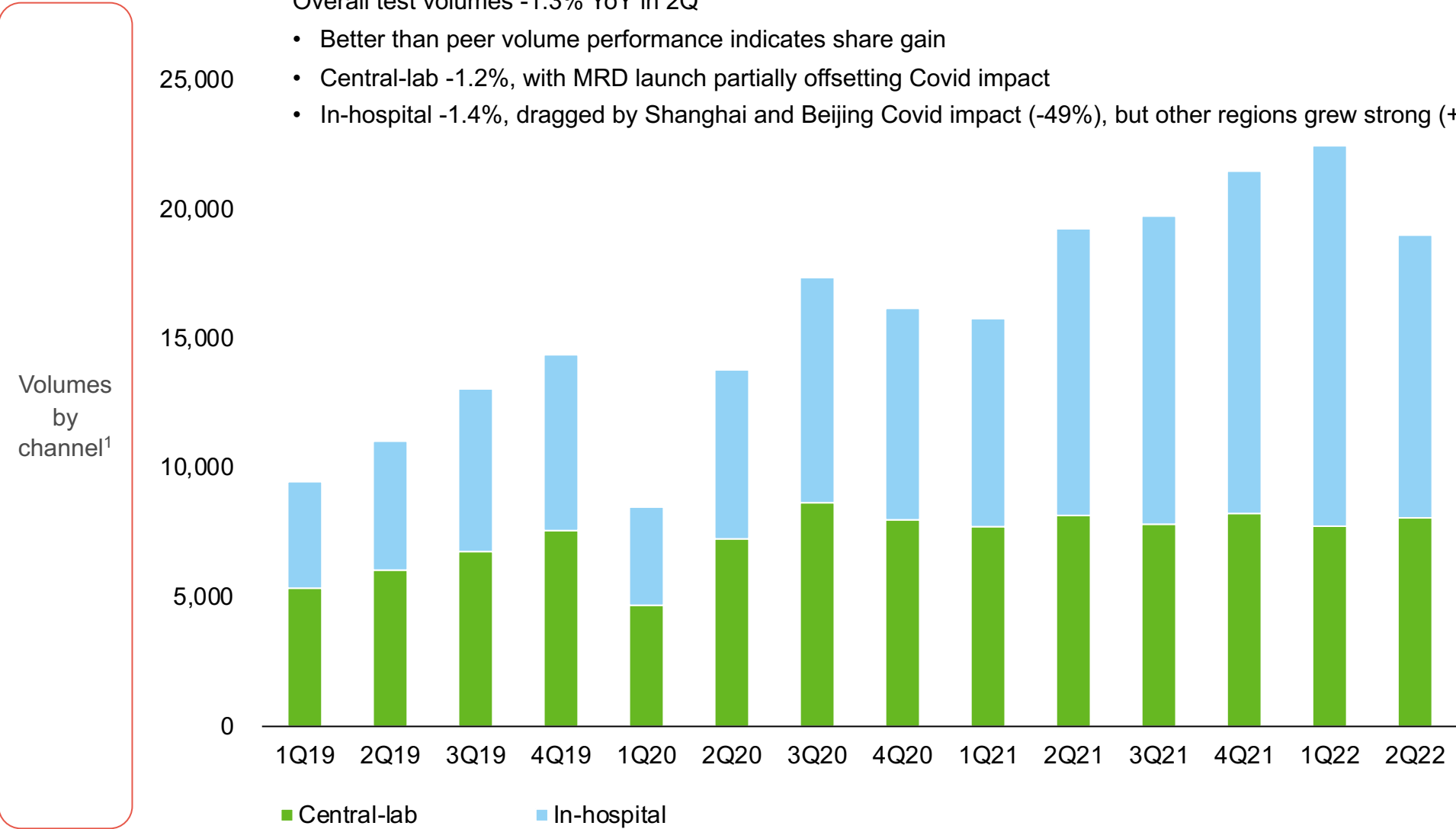


Financials

In-hospital and MRD driving growth uplift, but patient volumes negatively impacted by Covid related restrictions

Overall test volumes -1.3% YoY in 2Q

- Better than peer volume performance indicates share gain
- Central-lab -1.2%, with MRD launch partially offsetting Covid impact
- In-hospital -1.4%, dragged by Shanghai and Beijing Covid impact (-49%), but other regions grew strong (+60%)



Notes:

¹ Central-lab (LDT) volumes represented by the number of patients tested. In-hospital (IVD) volumes represented by the number of testing kits shipped to partner hospitals

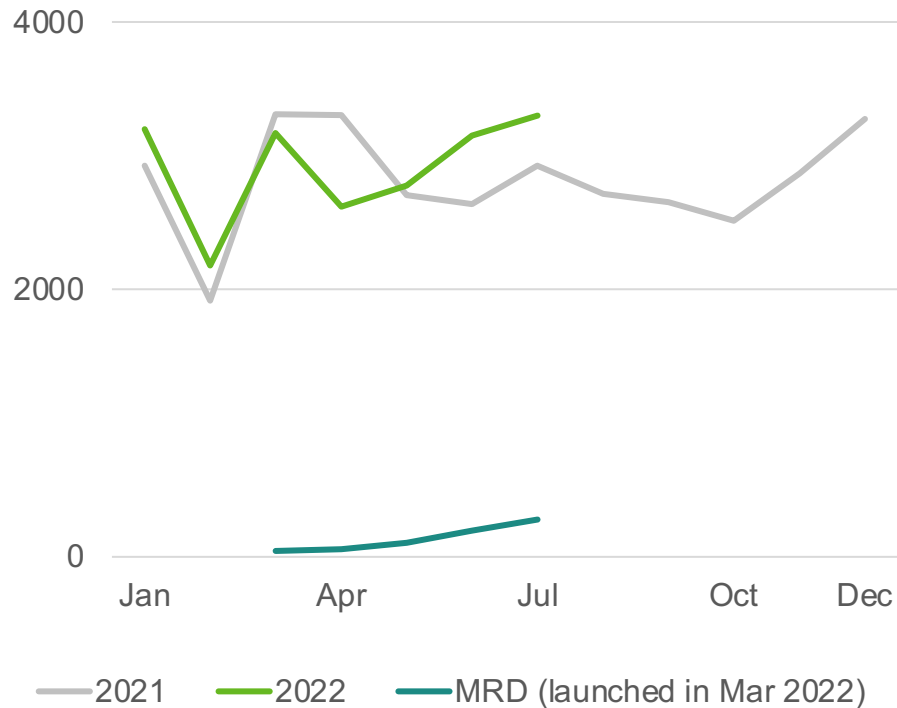
Latest trends

Continued secular growth of in-hospital penetration and MRD, despite Covid disruptions

Overall volume growth turned from -11% in Jun to +6% in Jul

Central-lab volumes

Jul +13% YoY, improved vs. 2Q's -1%, driven by MRD



In-hospital volumes

Shanghai, Beijing not yet fully recovered from Covid, strong growth in other regions

Volume growth YoY	Jan 2022	Feb	Mar	Apr	May	Jun	Jul
All regions	+149%	+95%	+36%	+60%	-11%	-28%	+0.2%
Shanghai and Beijing	+147%	+60%	+35%	+43%	-68%	-79%	-25%
Other regions	+152%	+154%	+37%	+78%	+56%	+51%	+41%

Notes:

Central-lab (LDT) volumes represented by the number of patients tested. In-hospital (IVD) volumes represented by the number of testing kits shipped to partner hospitals
 MRD – minimal residual disease testing

Financials

Opex to trend down over time

RMB1.15bn / USD172m cash and investments on balance as of Jun 2022

RMB millions	2021	19 YoY	20 YoY	21 YoY	1Q21	2Q21	3Q21	4Q21	1Q22	2Q22	2Q22 YoY	2Q22 QoQ	2022 Guide
Revenue	507.9	83%	13%	18%	106.6	127.3	126.6	147.3	135.5	130.8	3%	-3%	620
Central lab	319.4	71%	8%	7%	74.6	80.0	78.8	86.0	74.2	78.6	-2%	6%	
In-hospital ¹	165.1	164%	34%	40%	29.0	40.5	43.7	51.9	49.0	34.2	-16%	-30%	
Pharma	23.4	25%	(17%)	59%	3.1	6.8	4.1	9.4	12.3	18.0	165%	46%	
Non-GAAP Gross profit²	368.2				77.1	90.7	93.0	107.4	92.7	90.9	0%	-2%	
Total opex	1,161.2	49%	64%	60%	248.8	292.3	262.7	357.5	350.4	348.1	19%	-1%	
R&D ³	324.1				52.7	84.3	73.5	113.6	100.9	77.7	-8%	-23%	
S&M ³	283.4				50.0	62.7	72.1	98.6	84.6	100.3	60%	19%	
G&A ³	228.8				52.6	51.1	51.7	73.4	61.2	74.8	46%	22%	
SBC	280.8				84.4	83.0	53.3	60.2	79.8	76.7			
D&A	44.1				9.1	11.2	12.1	11.7	23.9	18.6			
Operating profit	(797.1)				(171.9)	(202.0)	(171.1)	(252.1)	(262.8)	(265.5)			
Net operating cash flows	(477.9)				(113.1)	(119.0)	(133.4)	(112.3)	(144.4)	(109.3)			
Non-GAAP GP margin ²	72.5%				72.4%	71.2%	73.4%	72.9%	68.4%	69.5%			
Opex ³ / revenue	165%				146%	156%	156%	194%	182%	193%			
S&M ³ / revenue	56%				47%	49%	57%	67%	62%	77%			

Notes:

¹ Within in-hospital segment, over 95% revenues are kit revenues, which are recurring in nature; the remaining are instrument revenues. In-hospital primarily through direct-sales model

² Non-GAAP gross margin, which is defined as gross margin excluding depreciation and amortization (D&A)

³ Excluding share based compensation (SBC) and depreciation and amortization (D&A)

Appendix 1

Early detection

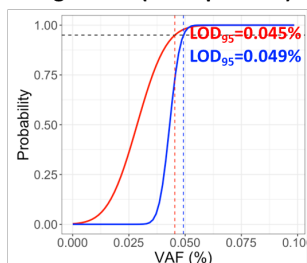
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Session OPO.CL11.01 - Biomarkers

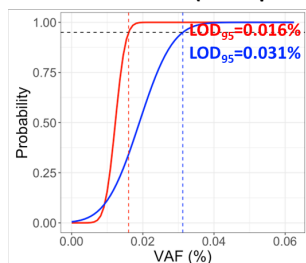
5116 - Analytical performance of ELSA-seq, a blood-based test for early detection of multiple cancers

- Analytical sensitivity.** The limit of detection with 95% probability (LOD_{95}) was established using 5ng DNA, the lowest claimed input amounts. Two models were assessed with a fixed training specificity at 95% (MCDBT-1) and 99% (MCDBT-2), respectively. Among six tested cancer types, the LOD_{95} was estimated down to 0.02% with respect to VAF.

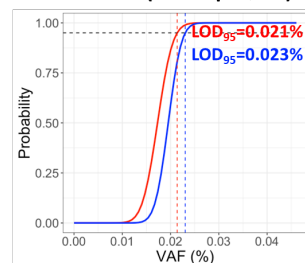
Lung cancer (EGFR:p.L858R)



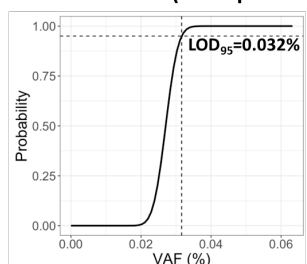
Colorectal cancer (EGFR:p.G719S)



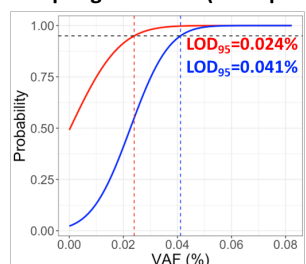
Liver cancer (NRAS:p.Q61L)



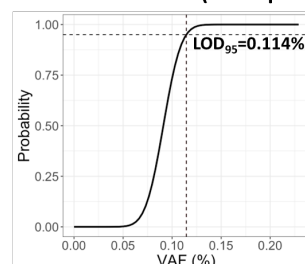
Ovarian cancer (EGFR:p.R215Q)



Esophageal cancer (TP53:p.H214R)



Pancreatic cancer (KRAS:p.G12R)

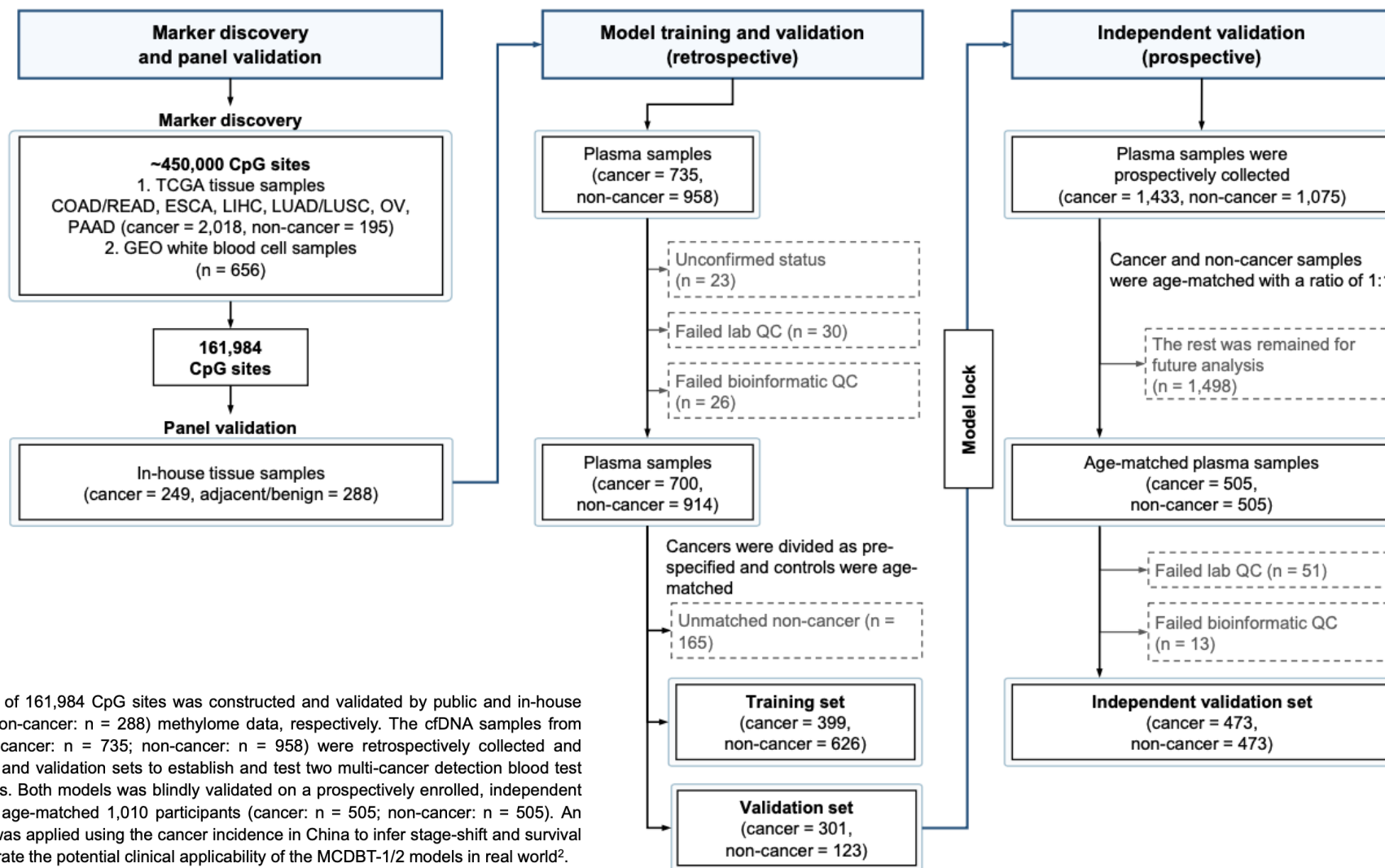


Full analytical validation study was conducted on ELSA-seq. LoD was demonstrated to be between 0.02% and 0.11% across different cancer types.

Figure 3: The LOD_{95} for 6 cancer types using two prediction models. Probit fit of DOC accuracy versus VAF using cell line dilution series. The red and blue curves represent MCDBT-1 and MCDBT-2 results, respectively. The black curves indicate the same results obtained by both models. The dotted lines indicate the LOD_{95} for each model.

ASCO 2022 – Thunder study read-out of the 6-cancer test Cohort

Fig 1. Flow chart.

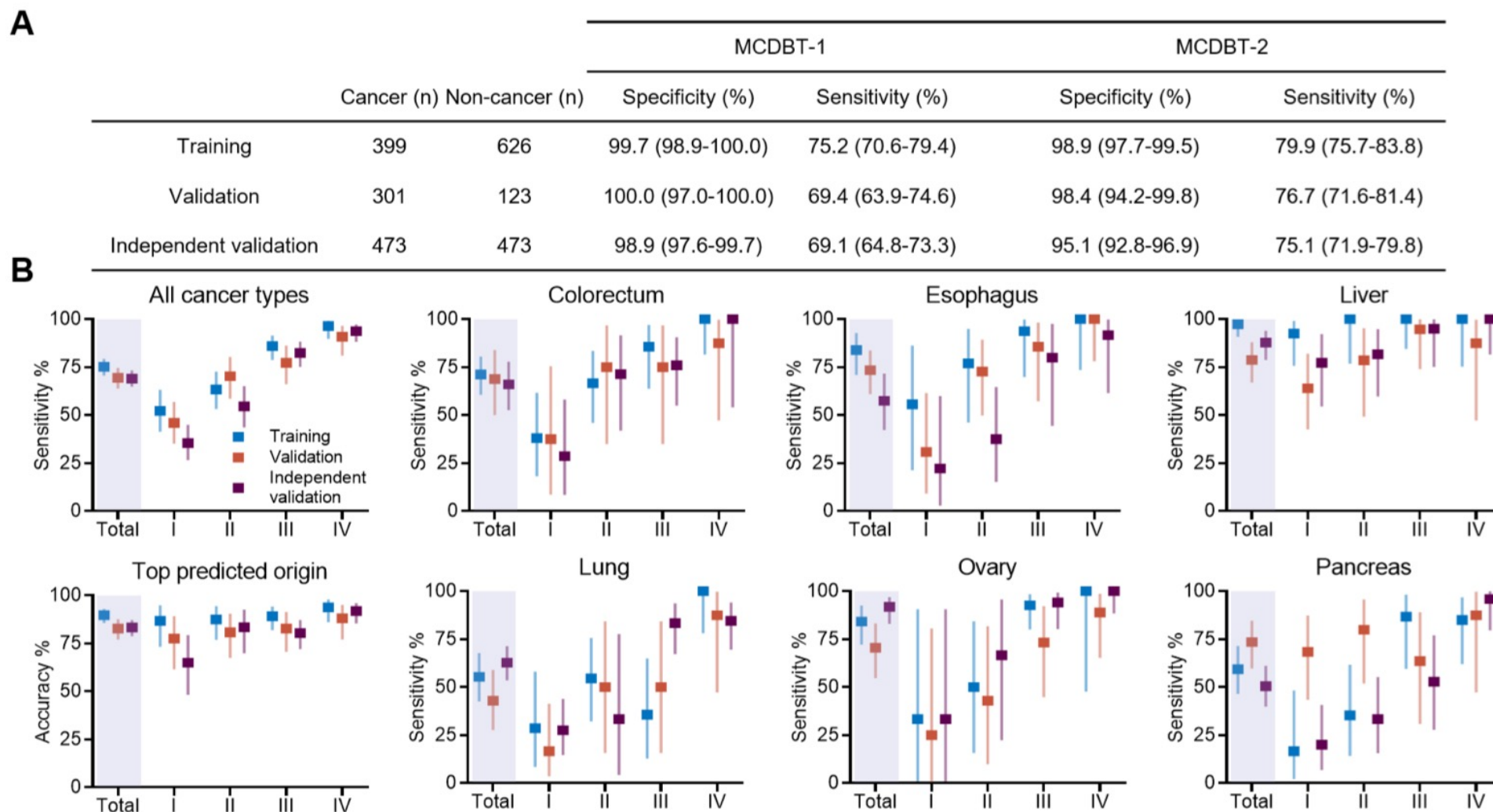


A customized panel of 161,984 CpG sites was constructed and validated by public and in-house (cancer: n = 249; non-cancer: n = 288) methylome data, respectively. The cfDNA samples from 1,693 participants (cancer: n = 735; non-cancer: n = 958) were retrospectively collected and divided into training and validation sets to establish and test two multi-cancer detection blood test (MCDBT-1/2) models. Both models were blindly validated on a prospectively enrolled, independent validation cohort of age-matched 1,010 participants (cancer: n = 505; non-cancer: n = 505). An interception model was applied using the cancer incidence in China to infer stage-shift and survival benefits to demonstrate the potential clinical applicability of the MCDBT-1/2 models in real world².

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Clinical performance on cancer detection

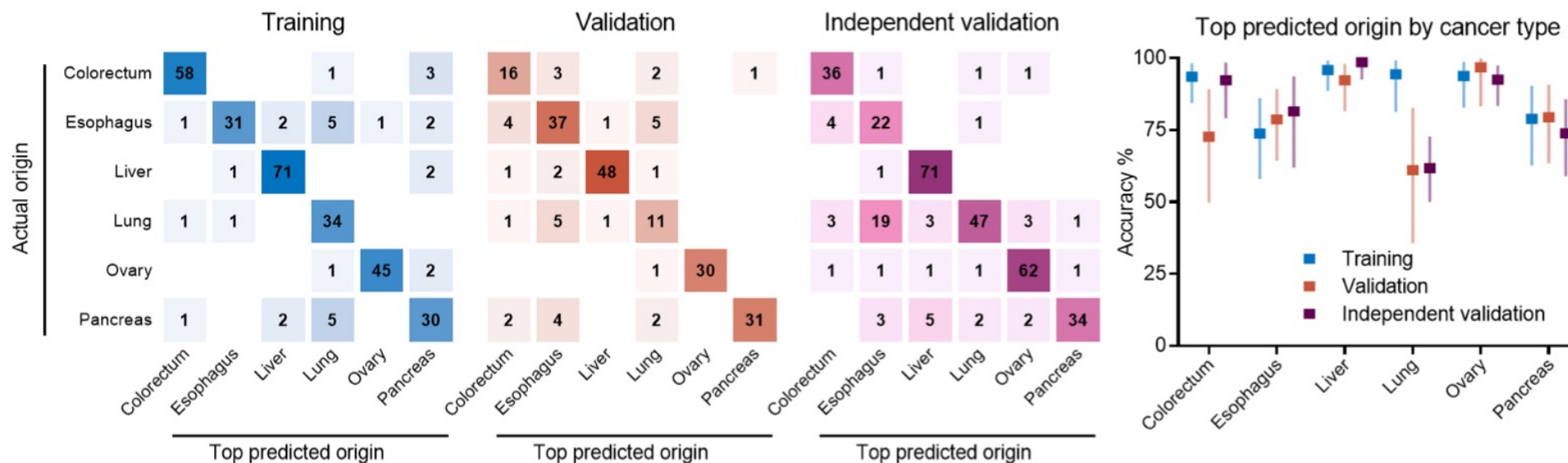
Fig 3. Performance of the MCDBT-1/2 models. A. Sensitivity, specificity, accuracy of top predicted origin, and accuracy of top two predicted origins. **B.** The overall sensitivity, accuracy of top predicted origin, and sensitivity stratified by cancer types reported by tumor stage.



ASCO 2022 – Thunder study read-out of the 6-cancer test

Clinical performance on tissue of origin

Fig 4. Top predicted origin for the MCBDT-1 model. Confusion matrices representing the predicted origin in the training, the validation, and the independent validation sets.

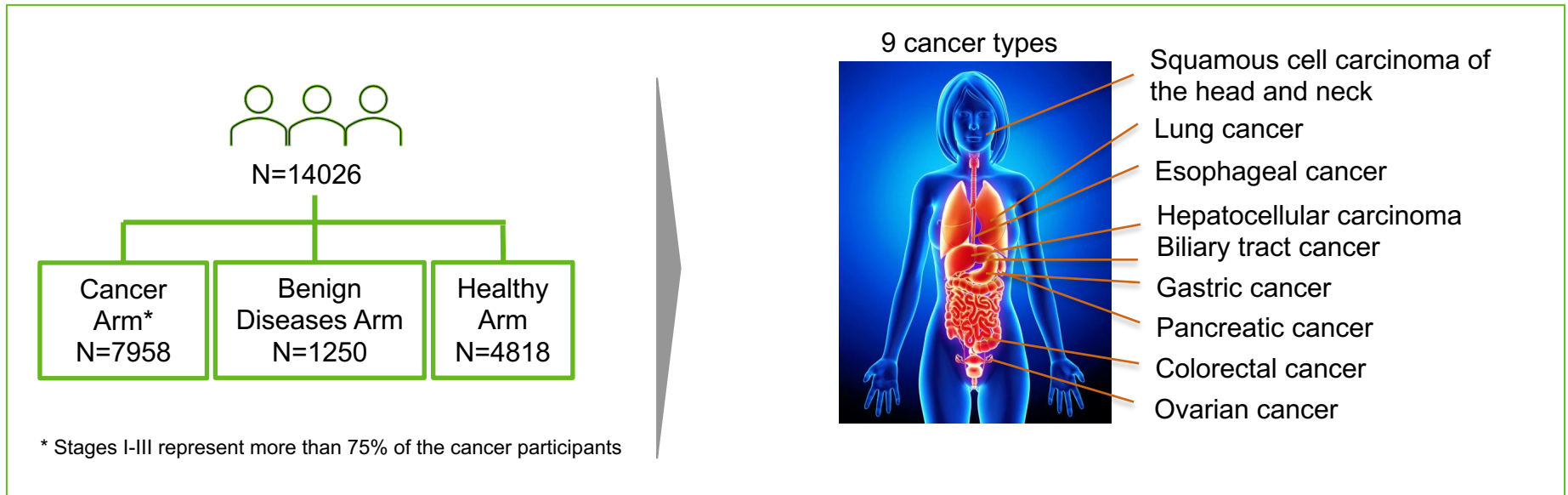


The PREDICT study (NCT04817306)

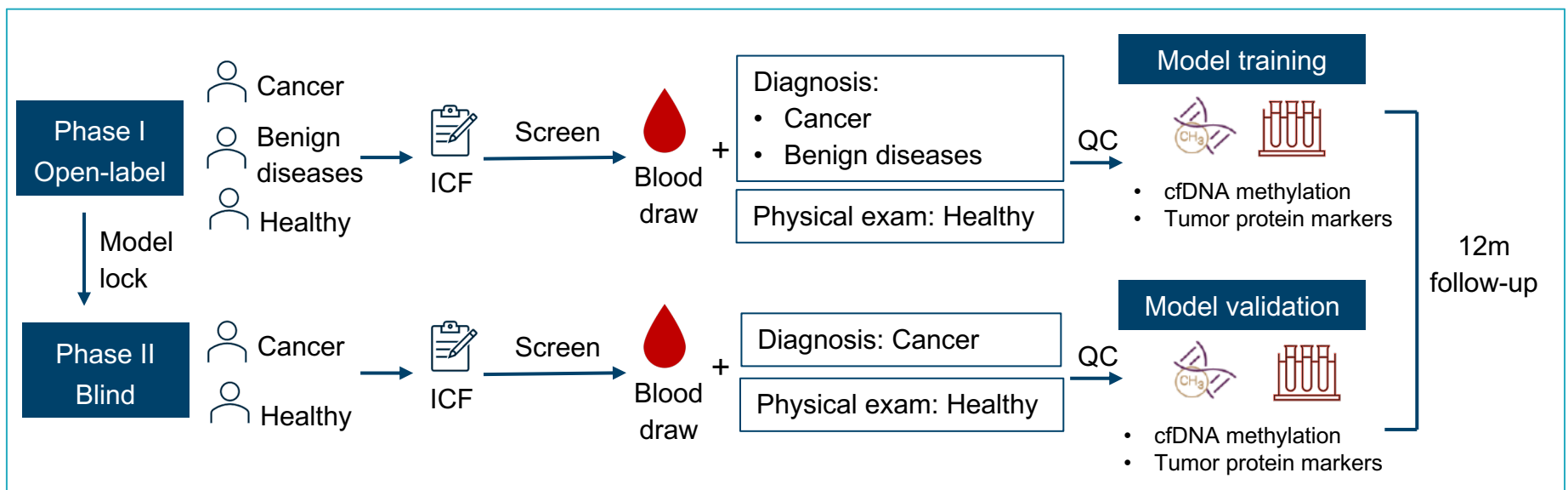
Study design

PREDICT is a *prospective, multi-center, case-control, observational* study for the detection of 9 cancer types through a cell-free DNA (cfDNA) methylation based, machine learning aided model

Participants



Study Design



The PREDICT study (NCT04817306)

National Oncology Conference on Standardized Diagnosis and Treatment, Beijing, 14th-16th May 2021

The image is a composite of three parts related to the PREDICT study. At the top, a banner for the 'National Oncology Conference on Standardized Diagnosis and Treatment' (NCC) in Beijing, May 2021, features logos for the National Cancer Center and other institutions, along with the title '全国肿瘤规范化诊疗工作会议暨肿瘤多学科诊治及转化研究高峰论坛'. Below this is a presentation slide titled '国内率先启动“泛癌种”早筛研究' (Domestic率先启动 "泛癌种" 早筛研究). The slide includes a diagram of 'cfDNA甲基化' (cfDNA methylation) with a human silhouette and DNA helix, and a '多癌种' (Multi-cancer) network diagram connecting 'Lung' and 'Breast' to 'Non-cancer'. A text box describes the study as a prospective, multi-center exploration and validation of an early cancer differentiation diagnosis model based on cfDNA methylation detection. A screenshot of the ClinicalTrials.gov entry for 'Pan-cancer Early Detection project (PREDICT)' is also shown, listing the study identifier NCT04817306 and its status as 'Not yet recruiting'. On the right, a man in a suit is speaking at a podium during the conference, with a screen behind him displaying '多癌种' and a network diagram.

国内率先启动“泛癌种”早筛研究

cfDNA甲基化

多癌种

基于cfDNA甲基化检测的早期癌症鉴别诊断模型在多癌种中的探索及验证：一项前瞻性、多中心研究 (Pan-Cancer Early Detection Project, PREDICT)

研究预计纳入癌症、良性病变及健康受试者；
样本量：14026例

NIH U.S. National Library of Medicine
ClinicalTrials.gov
Pan-cancer Early Detection project (PREDICT)
ClinicalTrials.gov Identifier: NCT04817306
Recruitment Status: Not yet recruiting
First Posted: May 12, 2020
Last Update Posted: May 14, 2020
See Contacts and Locations

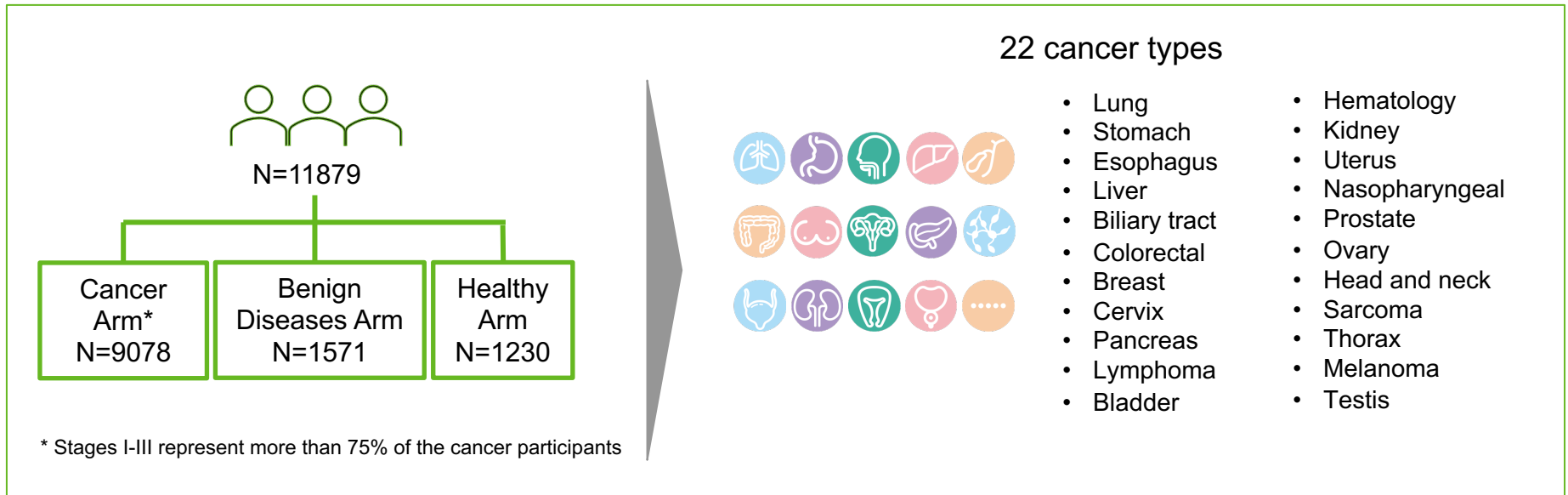
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The PRESCIENT study (NCT04822792)

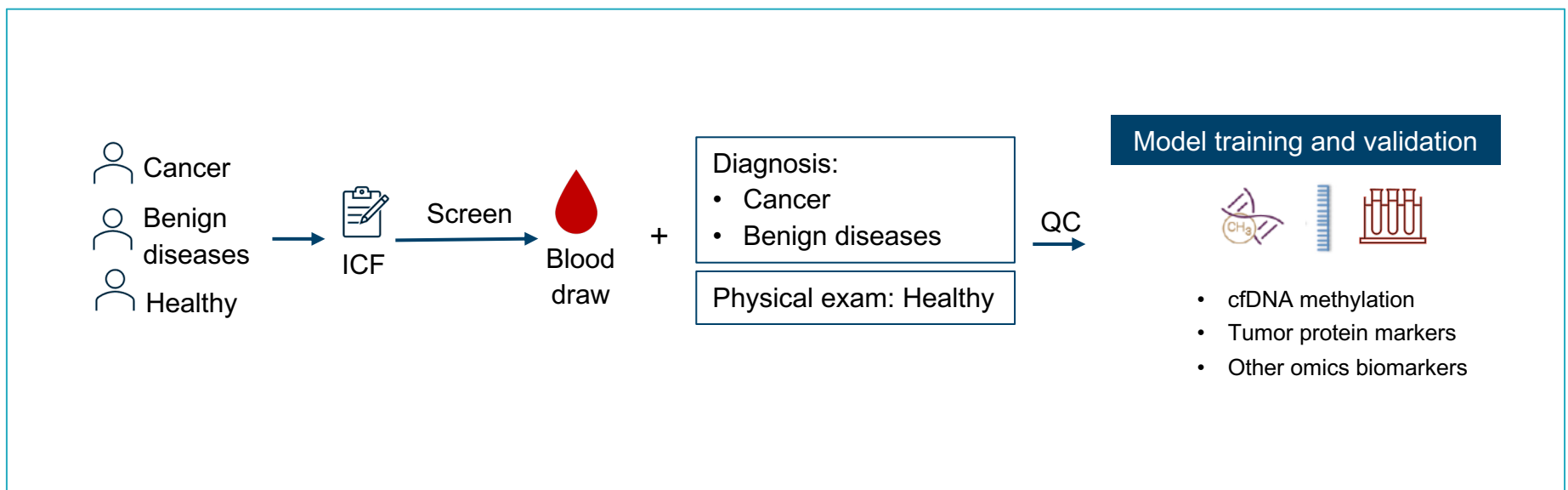
Study design

PRESCIENT is a *prospective, multi-center, case-control, observational* study aimed to train and validate the performance of a multi-omics model in the detection of 22 cancers

Participants

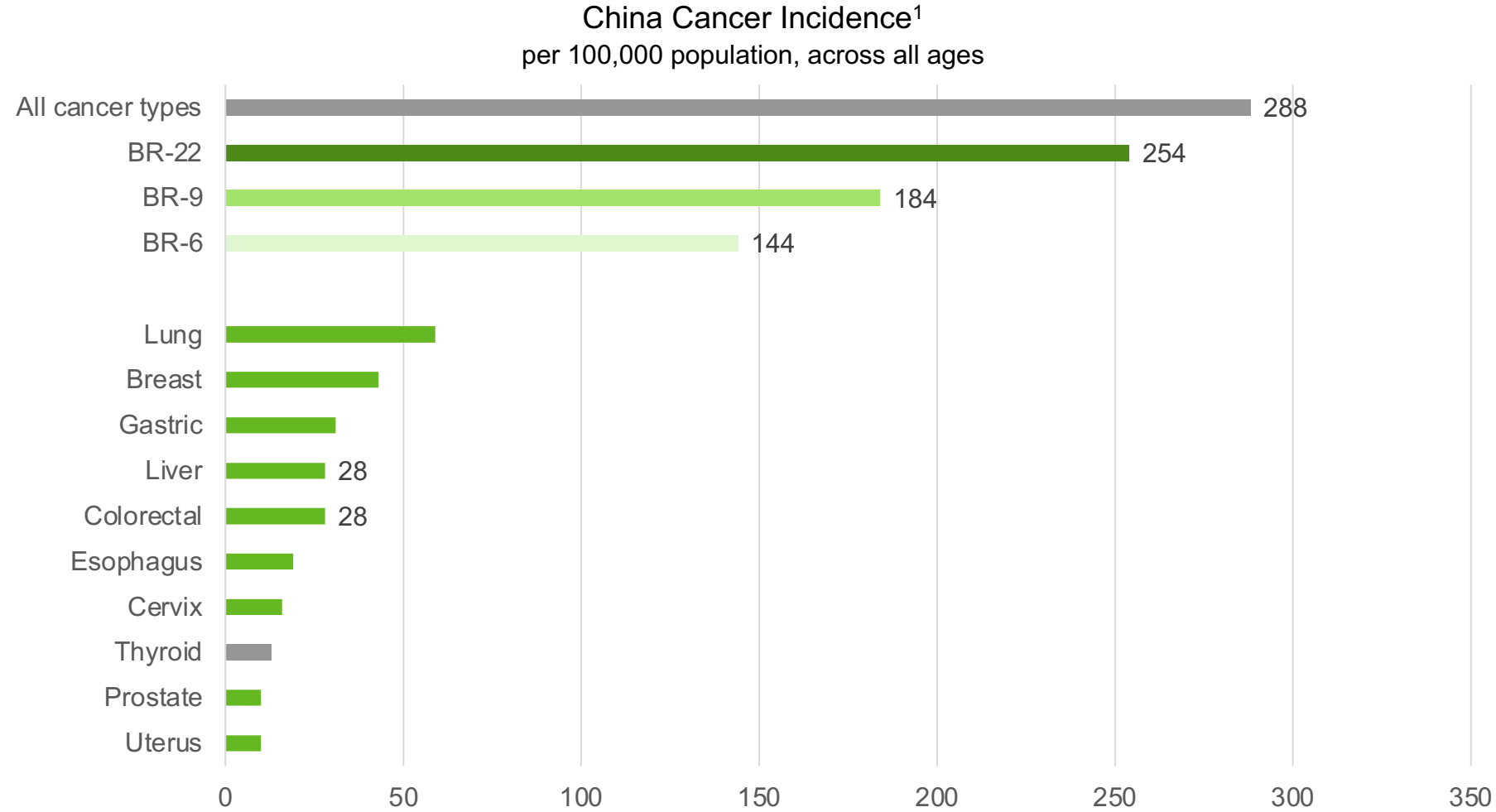


Study Design



Multi vs. single cancer early detection

Multiple times larger TAM



BR-22 covers 88% of China's cancer incidence²

Notes:
¹ Incidence data per "2018 China cancer registry annual report", J He et al., ISBN 978-7-117-28585-8
² Final number of cancer types subject to development progress

Multi vs. single cancer early detection in China

Significantly higher technology barrier

Single-cancer test

- Established technology, typically PCR based, with readily available products
 - US – First FDA approved product in 2014 (first submission in 2012)
 - China – NMPA approved products (class-III, including tissue and blood-based) in 2017, 2018, 2019, 2020, 2021, etc
- Small panel, low cost
- Relatively simple genomic data analytics

Multi-cancer test

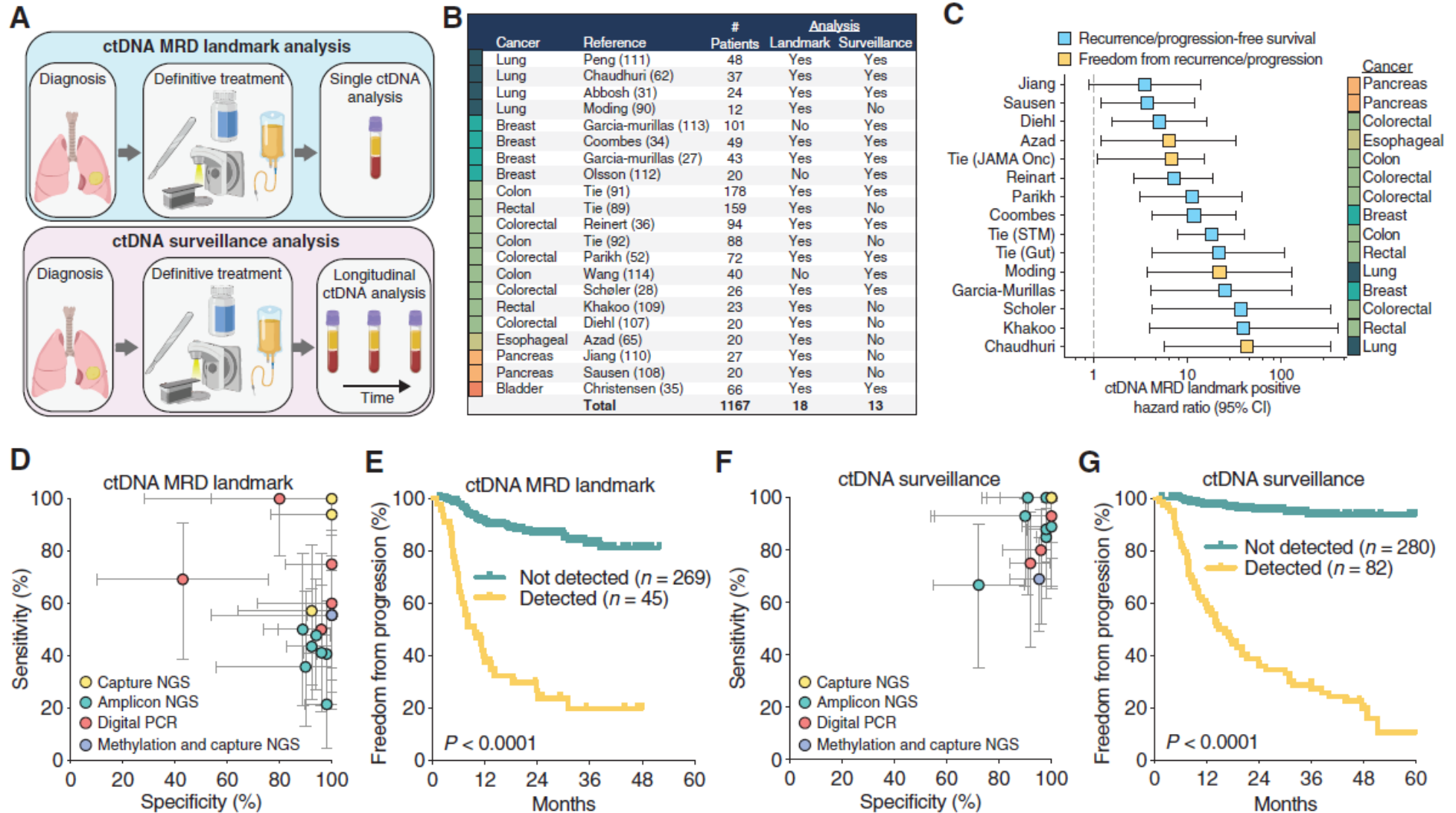
- Biologically, blood-based tests are multi-cancer in nature
- Highly complex technology with product risk
 - Globally, only a small number of innovators have locked-down products going under intended-use validation
- Data as a key factor for development and validation
 - Evolving dataset leads to continuous product improvement and greater validation
- Unprecedented commercial potential
 - Possibility to fundamentally shift oncology landscape from late-stage therapeutics to earlier stage intervention

Appendix 2

MRD

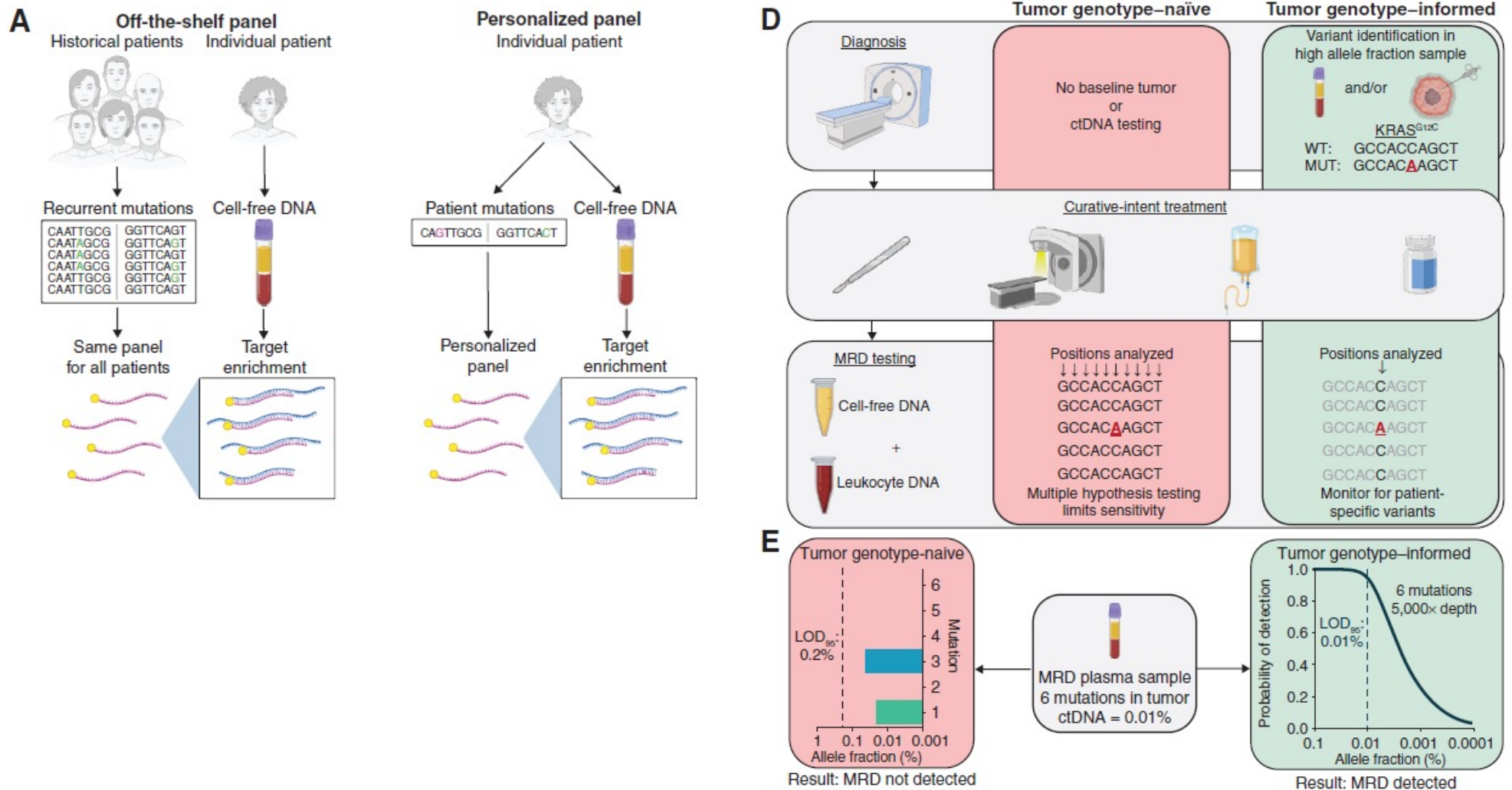
Clinical utilities of MRD in solid tumors

1) risk stratification and regimen selection (landmark analysis), 2) relapse monitoring (surveillance analysis)



Clinical utilities of MRD in solid tumors

Fixed panel vs. personalized panel approaches




Appendix 3

Therapy selection

NMPA approved NGS panels

NMPA approved testing kit by major NGS-focused companies¹

	First NMPA-approved kit	Second NMPA-approved kit
 燃石医学 Burning Rock Dx	EGFR, ALK, BRAF, KRAS Approved in Jul 2018 <u>First approved NGS kit in China</u>	EGFR, KRAS, MET, ERBB2, BRAF, PIK3CA, ALK, ROS1, RET Approved in Mar 2022
Novogene 诺禾	EGFR, KRAS, BRAF, PIK3CA, ALK, ROS1 Approved in Aug 2018	
Geneseeq 世和	EGFR, ALK, ROS1, BRAF, KRAS, ERBB2 Approved in Sep 2018	
BGI 华大	EGFR, KRAS, ALK Approved in Aug 2019	
Gene+ 吉因加	EGFR, KRAS, ALK Approved in Dec 2019	
Genetron 泛生子	EGFR, KRAS, BRAF, ERBB2, PIK3CA, ALK, ROS1, MET Approved in Feb 2020	
Genecast 臻和	KRAS, NRAS, BRAF, PIK3CA Approved in Mar 2021	
3DMed 思路迪		

Highlights on our second NMPA-approved kit

- Only 30ng DNA input required, applicable to small tissue samples
- First NMPA approved NGS kit with CNV² mutation type, with MET exon14 skipping

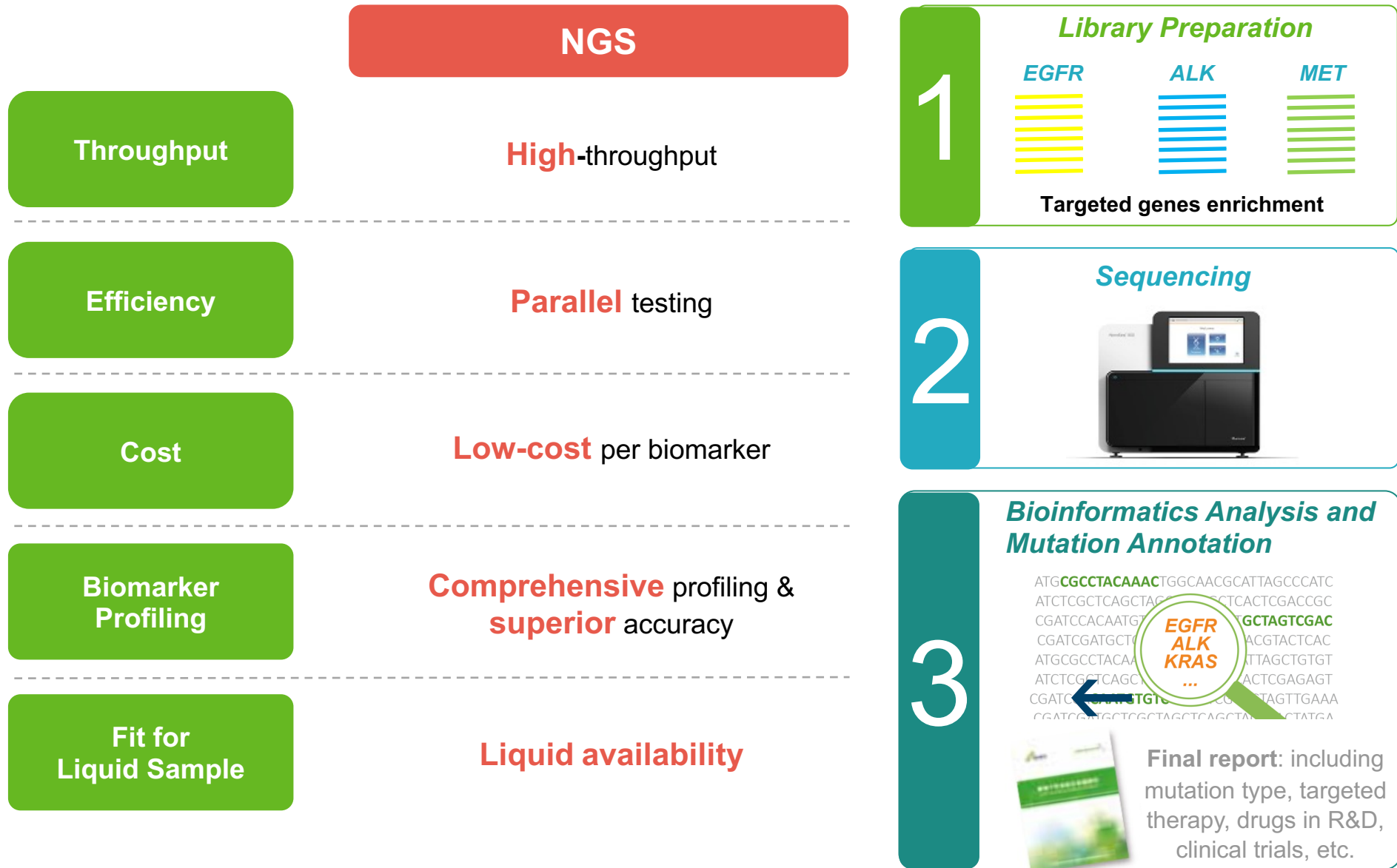
Notes:

¹ Major NGS-focused companies listed. The list is not exhaustive. A total of 13 kits have been approved by the NMPA as of the date of this presentation

² Copy number variation

NGS testing

Diagnostics companies focus on steps 1 and 3



Leading liquid-biopsy product in China, with globally competitive performance

Demonstrated in high-impact analytical validation study

SEQC2
Study
Overview

MAQC/SEQC Consortium Projects – An Overview

- An FDA-led community-wide consortium effort to assess technical performance and application of emerging technologies (e.g., genomics).



Issues and Study Objectives

- FDA approved several NGS tests with sensitivity for AF ~5%
- Hundreds lab developed tests (LDT): sensitivity ~ 2-10%
- FDA approved ctDNA tests with sensitivity for AF ~0.3%



Evaluating the analytical validity of circulating tumor DNA sequencing assays for precision oncology

- False positive rate estimate through known negatives
- All of them by VAF ranges:
 - 0.1 - 0.5%, 0.5 - 2.5%, >2.5%
 - Finer VAF ranges for sensitivity: 0.1 - 0.2%, 0.2 - 0.3%, 0.3 - 0.5%
- Evaluate the impact of DNA input amount
 - Three levels of input for Ef: 10ng, 25ng, 50ng
- Evaluate the impact of synthetic plasma (DNA extraction)
 - Qubit HS calibration and quantification
 - Calculate extraction yield

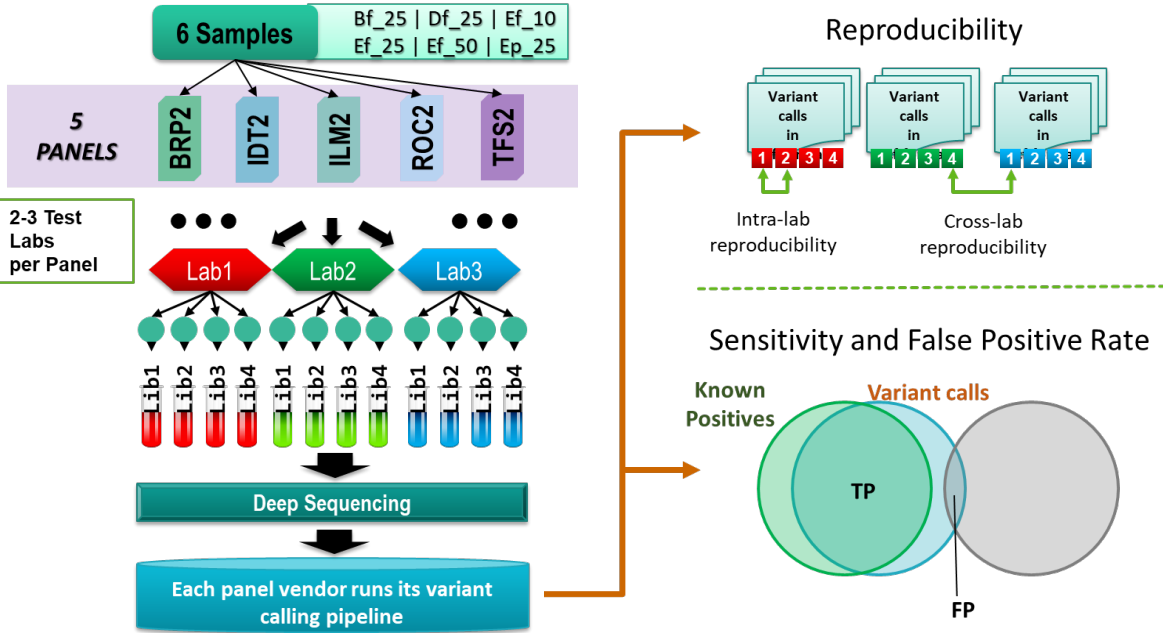
- Enzymatic fragmentation
 - better ligation efficiency
- Gel-based size selection (160bp-180bp) to mimic cfDNA
- 1ng/ul to mimic concentration after DNA extraction from plasma
- Ep: 40ng/ml Ef in synthetic plasma

BRP2: Burning Rock Dx LungPlasma v4
IDT2: IDT xGen Non-Small Cell Lung Cancer
ILM2: Illumina TruSight 170 with UMI
ROC2: Roche AVENIO ctDNA Expanded Kit
TFS2: Thermo Fisher Oncomine Lung cfDNA Assay

Liquid
Biopsy

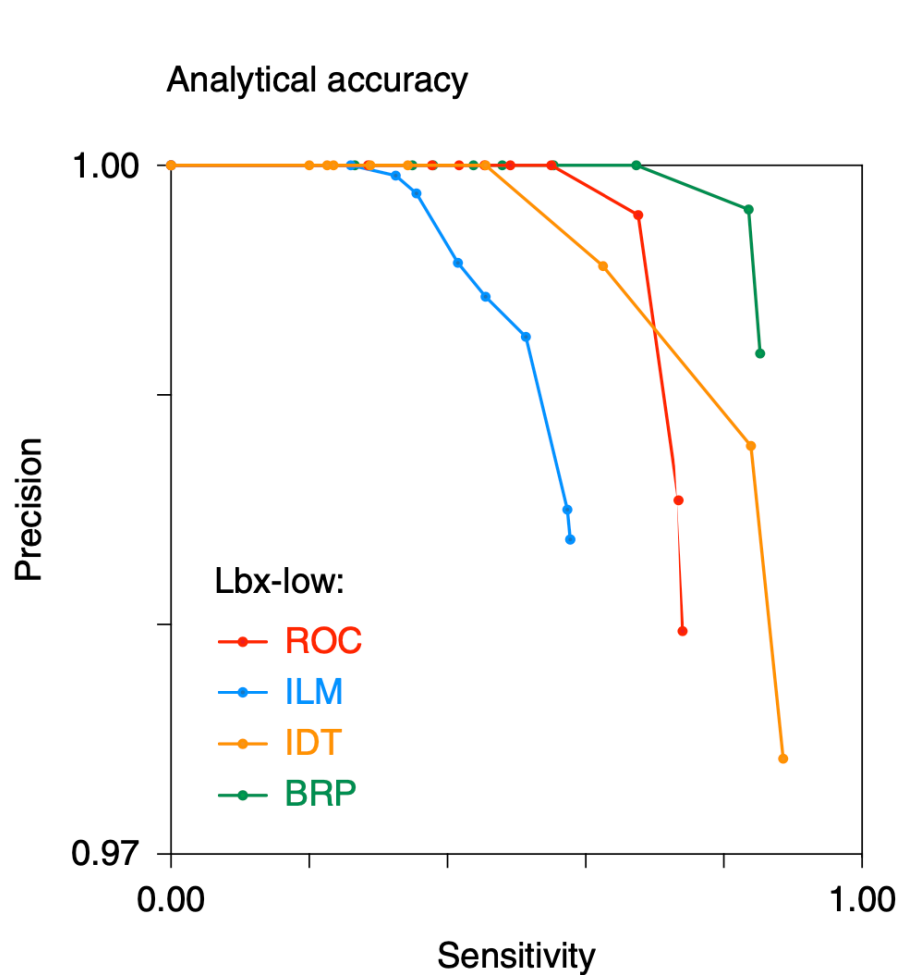
Participating assays and study design

Name	Vendor	ctDNA assay	Sequencing platform	Target genes	Reportable region (kb)	Coding (kb)	CTR (kb)	Negatives (× 1,000)	Variants
ROC	Roche Sequencing Solutions	AVENIO ctDNA (Expanded Kit)	Illumina NextSeq	77	161.7	140.2	103.8	47.1	189
ILM	Illumina	TruSight Tumor 170 + UMI	Illumina NovaSeq	154	501.0	390.1	338.4	133.0	574
IDT	Integrated DNA Technologies	xGen Non-small Cell Lung Cancer	Illumina NovaSeq	24	110.1	93.2	76.5	39.3	130
BRP	Burning Rock Biotech	Lung Plasma v4	Illumina NovaSeq	168	226.9	148.5	125.1	53.4	229
TFS	Thermo Fisher Scientific	Oncomine Lung cfDNA assay	Ion Torrent S5 XL	11	1.9	1.6	1.3	0.8	5



Source: "Evaluating the analytical validity of circulating tumor DNA sequencing assays for precision oncology", Nature Biotechnology, Apr 2021

Overall analytical accuracy and specificity

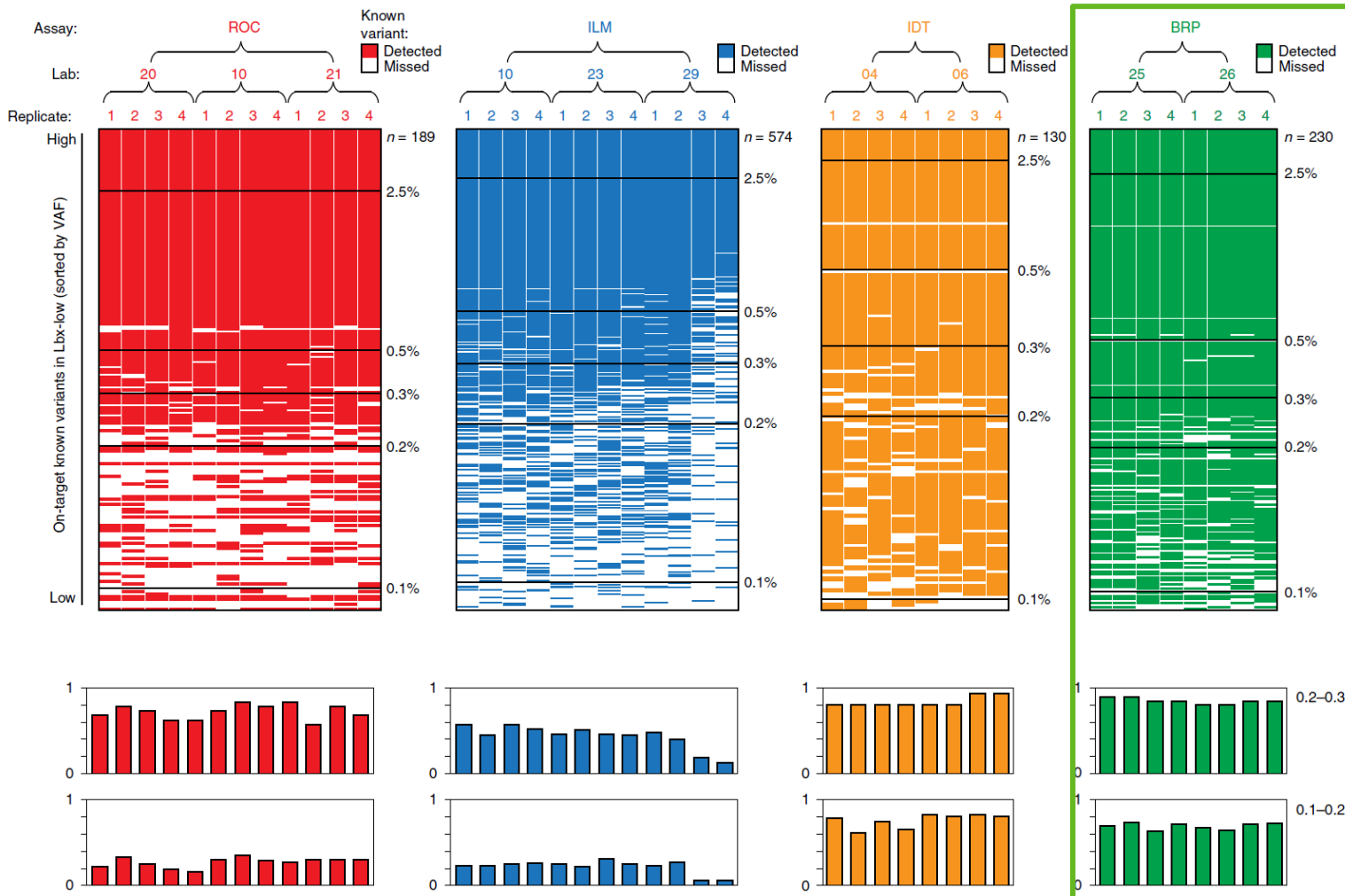


Assay	Known negatives (kb)	FPs per replicate (mean [range])	FP-rate (FP / kb) at specified VAF threshold		
			> 0%	> 0.1%	> 0.5%
ROC	47.1	2.91 [1-6]	0.061	0.044	0.000
ILM	133	5.25 [2-10]	0.039	0.039	0.008
IDT	39.3	2.75 [0-6]	0.070	0.057	0.000
BRP	53.4	1.65 [0-5]	0.030	0.007	0.000

The analytical accuracy was measured by **Precision-Sensitivity** plot (25ng LBx-Low)
 The false positive rates were computed by FP/kb region.
 Once different VAF threshold increases, FP rates dropped further.

“To compare the accuracy of the participating ctDNA assays, we generated precision recall curves, ranking known variants and FPs according to their observed VAFs. **For Lbx-low samples at 25ng input, BRP was the most accurate assay, with roughly equivalent sensitivity but superior precision to IDT** (Fig. 4b and Supplementary Fig. 4c).”

Performance – Sensitivity



- LBx-low (25 ng input) replicates in each participating assay in different expected VAF bin.

“The most sensitive assays (IDT and BRP) achieved sensitivity greater than 0.90 for variants with 0.3–0.5% VAF; however, no assays reached this mark for variants with 0.2–0.3% or 0.1–0.2% VAF (Fig. 4a).”

“The performance characteristics of the assays evaluated here were broadly similar to what has been reported by several ctDNA sequencing providers (based on internal testing) that did not participate in this study. **During validation of the Guardant360 CDx hybrid capture assay, variants were detected with high sensitivity (~94%) at VAF ≥ 0.4%, declining to ~64% among variants with VAF ranging from 0.05% to 0.25%.** **FoundationACT showed ~99% sensitivity for SNVs with VAF > 0.5%, ~95% for 0.25%–0.5% VAF and ~70% for 0.125–0.25% VAF.**”