燃石医学
Burning Rock Dx

Burning Rock Biotech Limited

3Q2023 results

Nasdaq and LSE: BNR

30 Nov 2023

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Our value-building blocks

Extending leadership in NGS-based precision oncology from late-stage to earlier stage patients

New Businesses

Large market potential

At early commercial phase

Early Detection

Asymptomatic population

MRD[1]

Early-stage oncology patients

Biopharma

Global CDx[2] partners for pivotal trials of targeted drugs. Pharma R&D

Common Infrastructure

Accelerating growth of new businesses

- Strong brand to support new product launches & attract talent
- Broad industry network and synergy across different business units
- Large volumes supporting lower cost & faster innovation

Developed Business

Commercial phase

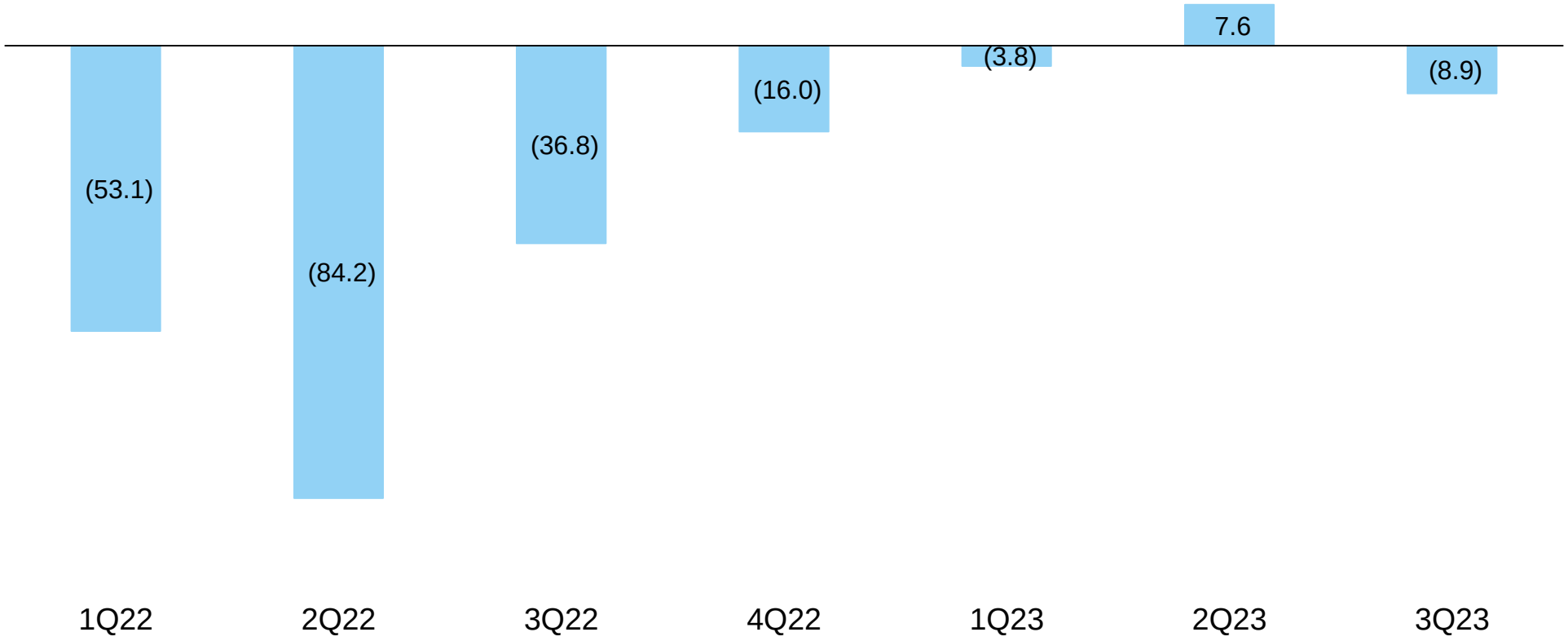
Therapy Selection

Late-stage oncology patients

Notes:
[1] Minimal residual disease of solid tumors
[2] Companion diagnostics

Significant progress towards breakeven

Non-GAAP gross profit *minus* non-GAAP SG&A, excluding R&D* (RMB millions)



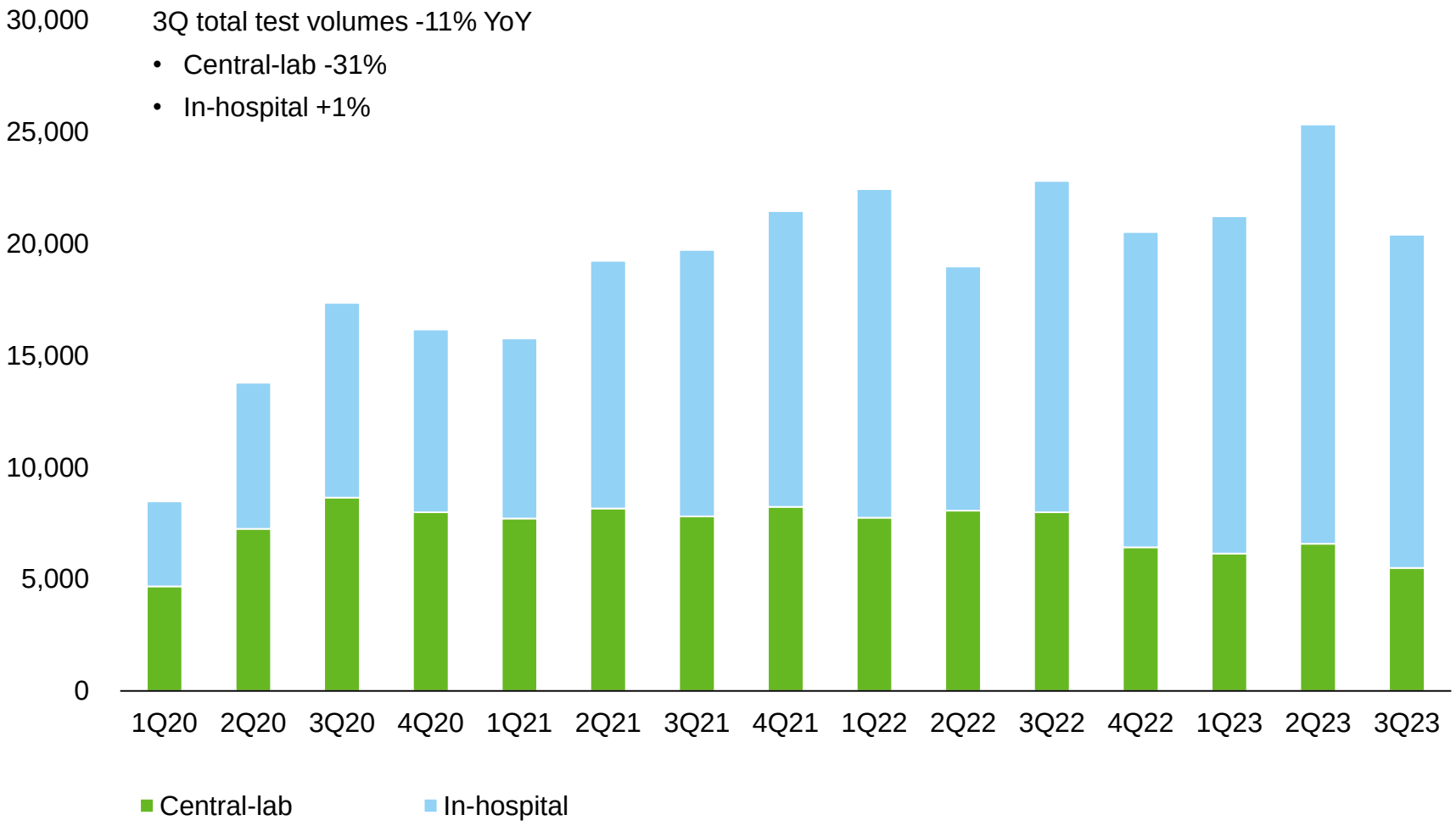
Notes:
* Non-GAAP gross profit, refers to gross profit excluding depreciation and amortization. Non-GAAP SG&A refers to selling and marketing expenses and general and administrative expenses, both excluding their respective share-based compensation and depreciation and amortization.

3Q2023 progress

Corporate	• Significantly narrowed losses and cash outflows vs. 3Q2022 • Execution towards profitability well underway
Therapy selection	• Continued strength in in-hospital channel, despite industry volatility • In-hospital revenues +10% YoY
MRD	• Strong clinical validation publication, with the MEDAL study on lung cancer published in Cancer Cell
Biopharma	• Steady growth, with revenues +31% YoY • Growing backlog, with strong project wins, e.g. entered into CDx contracts with Boehringer Ingelheim
Early detection	• Over C™ MCDBT received Breakthrough designation from China's National Medical Products Administration (NMPA). It's the only early detection test globally that has received breakthrough designation from both the FDA and the NMPA

Quarterly Test volumes

Volumes by channel[1]



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

Financials

RMB millions	2021	2022	1Q22	2Q22	3Q22	4Q22	1Q23	2Q23	3Q23	3Q23 YoY	3Q23 QoQ
Revenue	**507.9**	**563.1**	**135.5**	**130.8**	**154.6**	**142.2**	**142.6**	**146.2**	**127.6**	**-17%**	**-13%**
Central lab	319.4	314.8	74.2	78.6	90.0	72.0	61.8	66.2	53.5	-41%	-19%
In-hospital[1]	165.1	175.3	49.0	34.2	49.6	42.5	51.6	53.8	54.5	10%	1%
Pharma	23.4	73.0	12.3	18.0	15.0	27.7	29.2	26.2	19.6	31%	-25%
Non-GAAP Gross profit[2]	**368.2**	**411.0**	**92.7**	**90.9**	**117.0**	**110.4**	**107.9**	**109.4**	**95.1**	**-19%**	**-13%**
Total opex	**1,161.2**	**1,360.5**	**350.4**	**348.1**	**343.3**	**318.7**	**287.2**	**236.1**	**264.7**	**-23%**	**12%**
R&D[3]	324.1	344.4	100.9	77.7	88.7	77.1	74.0	73.1	64.2	-28%	-12%
S&M[3]	283.4	350.6	84.6	100.3	85.4	80.3	60.5	64.7	56.8	-33%	-12%
G&A[3]	228.8	250.5	61.2	74.8	68.4	46.1	51.2	37.1	47.2	-31%	27%
SBC	280.8	325.1	79.8	76.7	77.4	91.2	77.8	37.2	72.7		
D&A	44.1	89.9	23.9	18.6	23.4	24.0	23.7	24.0	23.8		
Non-GAAP GP – non-GAAP SG&A	**(144.0)**	**(190.1)**	**(53.1)**	**(84.2)**	**(36.8)**	**(16.0)**	**(3.8)**	**7.6**	**(8.9)**		
Operating profit	**(797.1)**	**(980.3)**	**(262.8)**	**(265.5)**	**(234.6)**	**(217.4)**	**(188.5)**	**(135.7)**	**(178.8)**		
Net operating cash flows	**(477.9)**	**(456.9)**	**(144.4)**	**(109.3)**	**(135.5)**	**(67.7)**	**(113.1)**	**(79.2)**	**(47.4)**		
Non-GAAP GP margin[2]	72.5%	73.0%	68.4%	69.5%	75.7%	77.6%	75.7%	74.8%	74.5%		
Opex[3] / revenue	165%	168%	182%	193%	157%	143%	130%	120%	132%		
S&M[3] / revenue	56%	62%	62%	77%	55%	56%	42%	44%	45%		

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)

Strong cash position to fund operations for the next 3 years

Operating loss and cash outflow reduction executing better vs. plan

3Q23 quarterly net operating cash outflow at RMB47m

RMBm	2022	1Q-3Q 2023	2023E[1]	2024E[1]
Operating cash outflow[2]	457	240		
Capex[3]	75	9		
Sum	532	249	c.400	c.200
Cash balance at period-end[4]	925	637		

Estimate assumptions

- Cash spend to focus on early detection clinical studies, the bulk of which will run through 2023 and drop off in 2024
- Commercial business to breakeven during 2023 (no further upside assumed in 2024 estimate)

Notes:
[1] Based on management's current estimate and subject to change
[2] Net cash used in operating activities
[3] Purchase and prepayment of property and equipment and intangible assets, issuance of convertible loan, out of investing cashflows
[4] Consists of Cash and cash equivalents of approximately RMB636.3m, restricted cash of approximately RMB0.5m as of the end of 3Q2023

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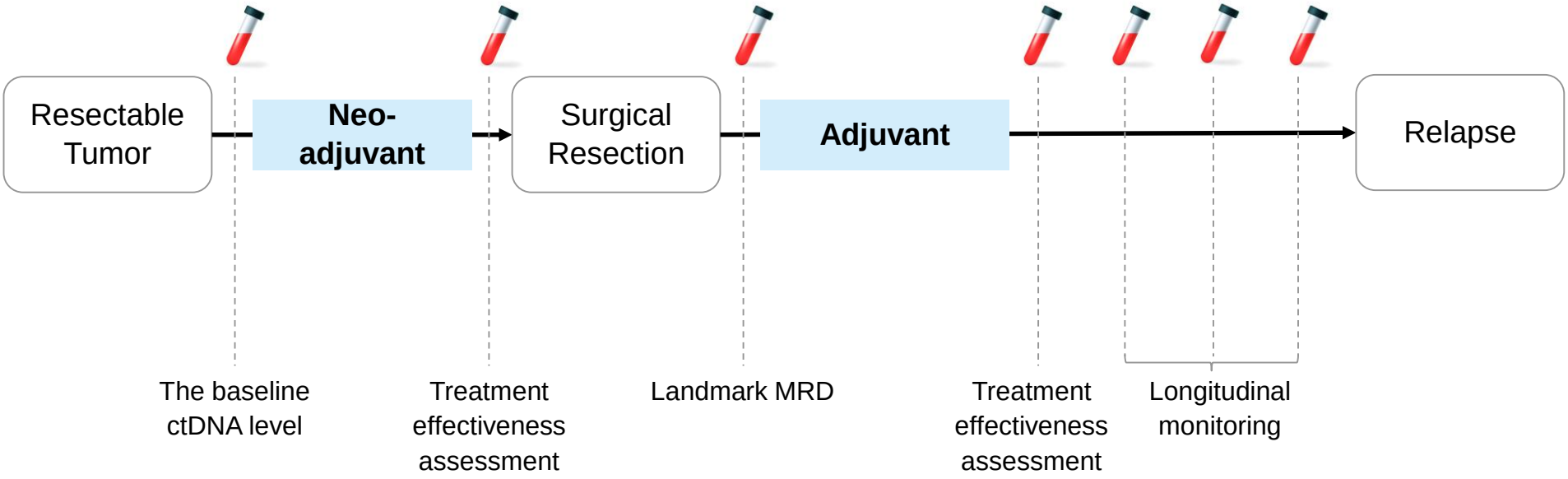
Minimal Residual Disease (MRD)

Burning Rock's MRD clinical publications

Covering adjuvant and relapse settings in lung, colorectal, gastric and other cancers

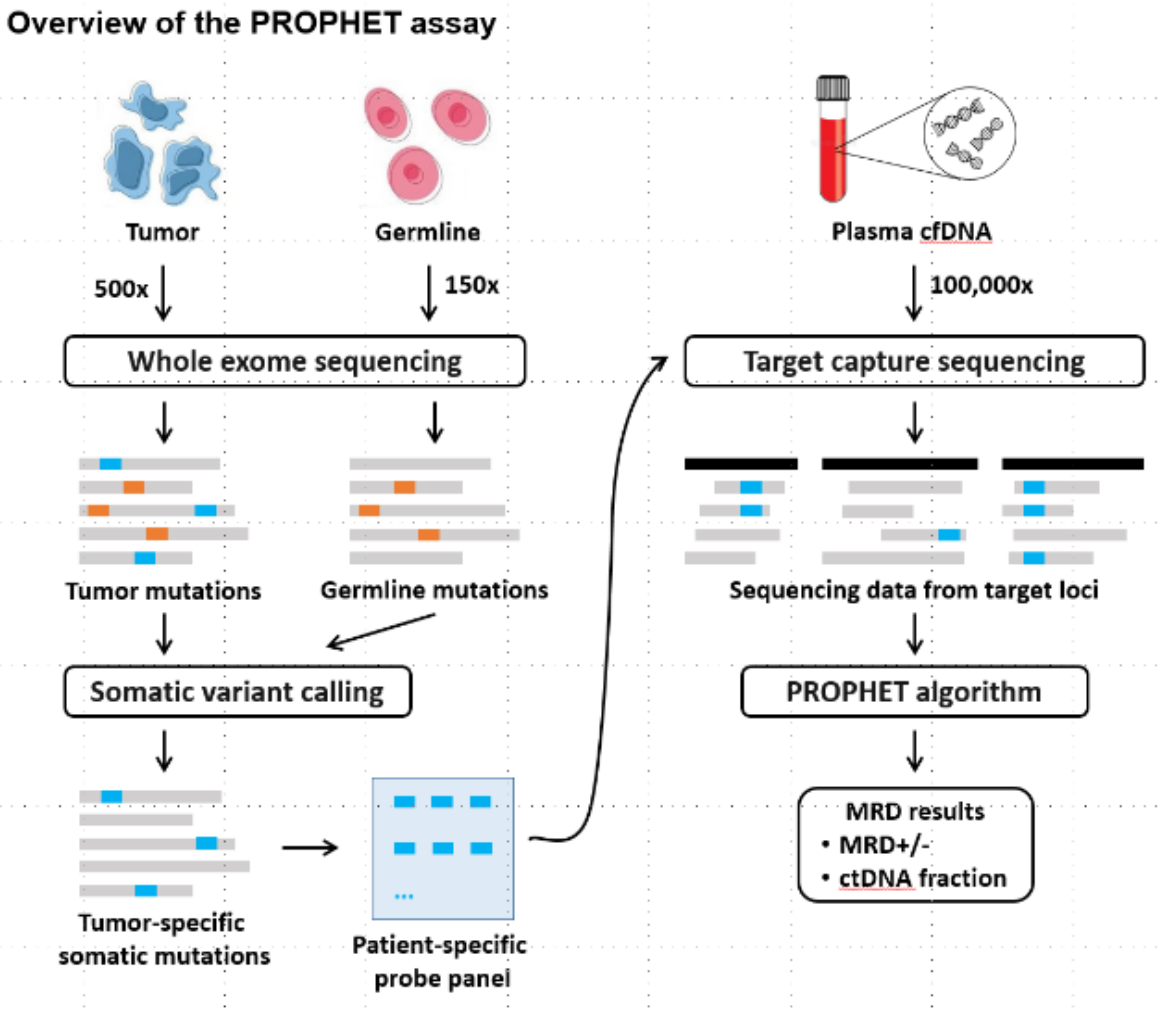


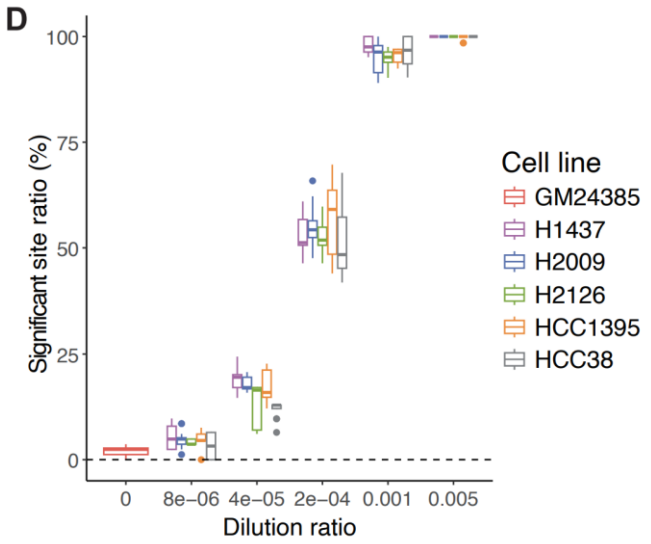
Non-small-cell lung cancer	Baseline, landmark and longitudinal monitoring timepoints completed AACR 2022 Abstract 5916, AACR 2023 Abstract 1039, MEDAL study publication	
Colorectal cancer	Baseline and landmark timepoints AACR 2022 Abstract 5917, ASCO GI 2023 Abstract 213	Longitudinal monitoring ongoing
Gastric cancer	Baseline and landmark timepoints AACR 2023 Abstract 6682	
Pancreatic cancer	Baseline, landmark and longitudinal monitoring timepoints completed ASCO GI 2023 Abstract 744	
Biliary tract cancer	Baseline and landmark timepoints AACR 2023 Abstract 6682	

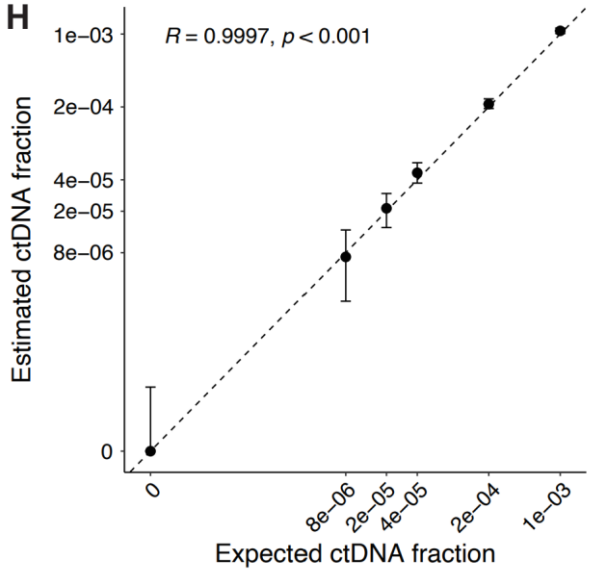
Cancer Cell

Overview of brPROPHET™

An ultrasensitive and quantitative MRD assay







Source: Chen et al., Individualized tumor-informed circulating tumor DNA analysis for postoperative monitoring of non-small cell lung cancer, Cancer Cell, Sep 2023

MEDAL study

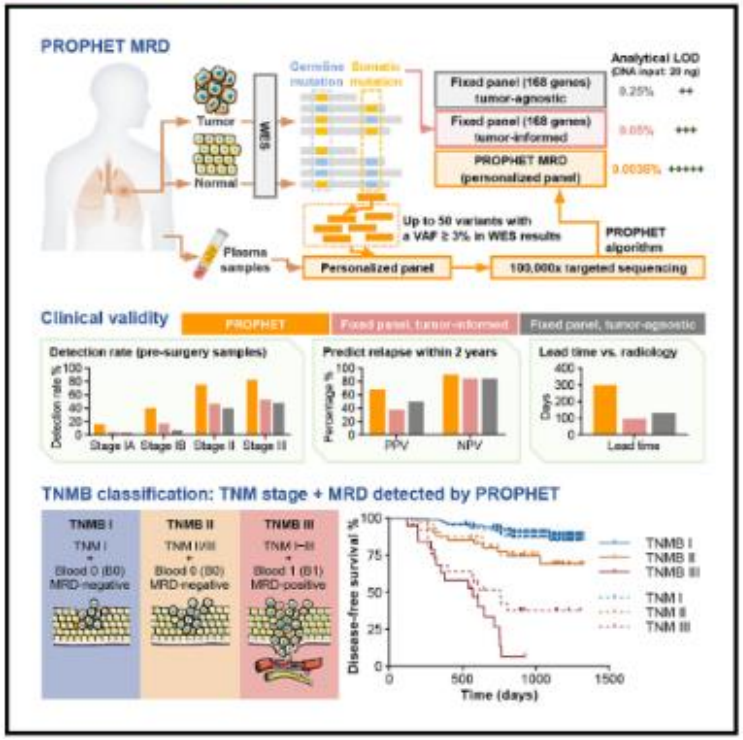
Personalized MRD using brPROPHET™ on non-small cell lung cancer (NSCLC)

Cancer Cell

Article

Individualized tumor-informed circulating tumor DNA analysis for postoperative monitoring of non-small cell lung cancer

Graphical abstract



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Correspondence

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In brief

Chen et al. introduced personalized Patient-specific pROgnostic and Potential tHErapeutic marker Tracking (PROPHET) for detecting molecular residual disease (MRD) in NSCLC, featuring a notably low limit of detection (LOD). It exhibits elevated sensitivity and extended lead time than radiologically confirmed recurrence. It also facilitates prognostic accuracy and postoperative treatment evaluation.

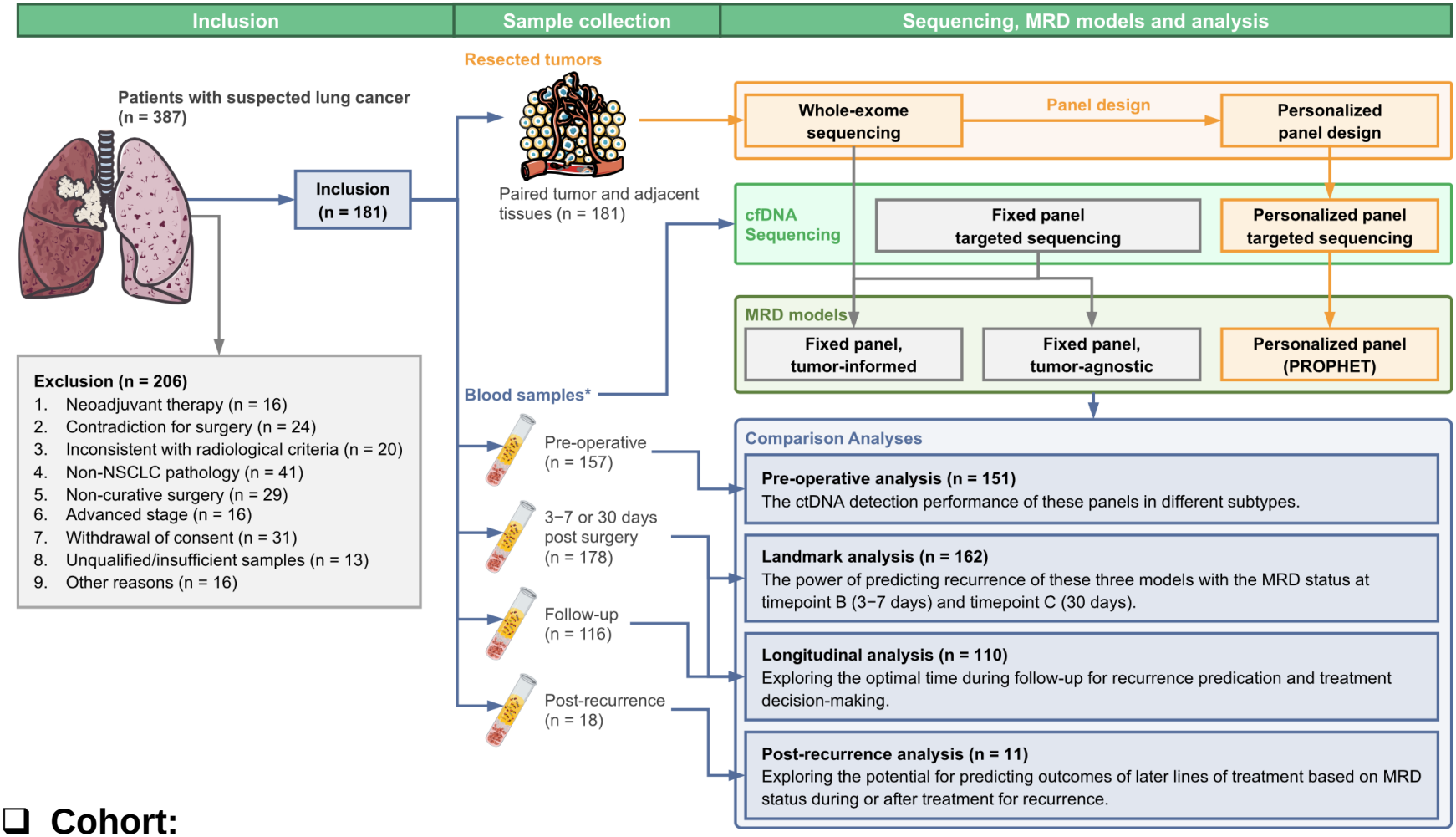
Highlights

- PROPHET outperforms fixed-panel MRD assays in head-to-head comparison in NSCLC
- TNMB stage, integrating landmark ctDNA MRD and TNM, improves prognosis prediction
- PROPHET illustrates a median lead time of 299 days to radiological recurrence
- Post-relapse ctDNA status facilitates decision on later lines of treatment

Chen et al., 2023, Cancer Cell *41*, 1–14
October 9, 2023 © 2023 Published by Elsevier Inc.
https://doi.org/10.1016/j.ccell.2023.08.010

Source: Chen et al., Individualized tumor-informed circulating tumor DNA analysis for postoperative monitoring of non-small cell lung cancer, Cancer Cell, Sep 2023

Study design



- **Cohort:**
  - 181 patients enrolled Stage I (63%), II (19%), and III (18%)
- **Sampling Time:**
  - Tumor and adjacent paired tissue collected at surgery
  - Blood samples collected at Pre-operative, 3 days, and 30 days post-surgery
  - Median Follow-up Time: 30 months

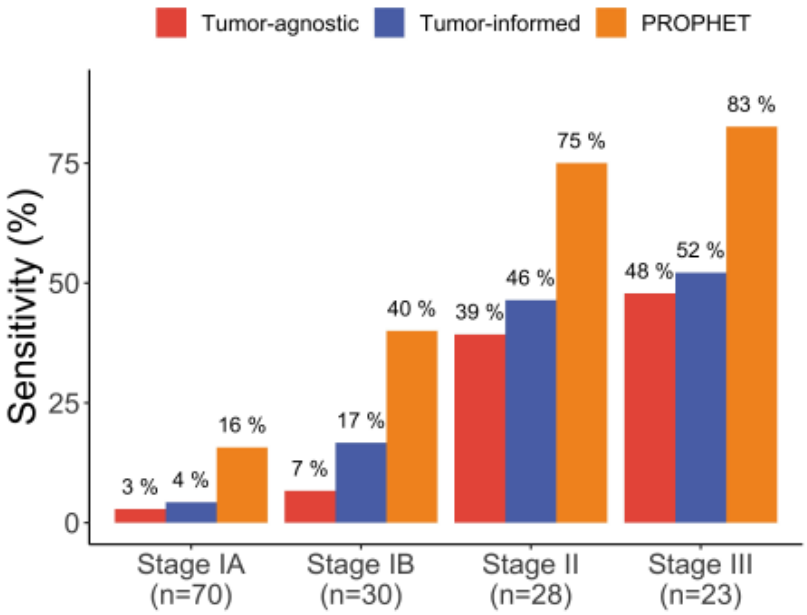
Source: Chen et al., Individualized tumor-informed circulating tumor DNA analysis for postoperative monitoring of non-small cell lung cancer, Cancer Cell, Sep 2023

brPROPHET™ demonstrates superior sensitivity in ctDNA detection

Clinical validation with pre-operative plasma samples

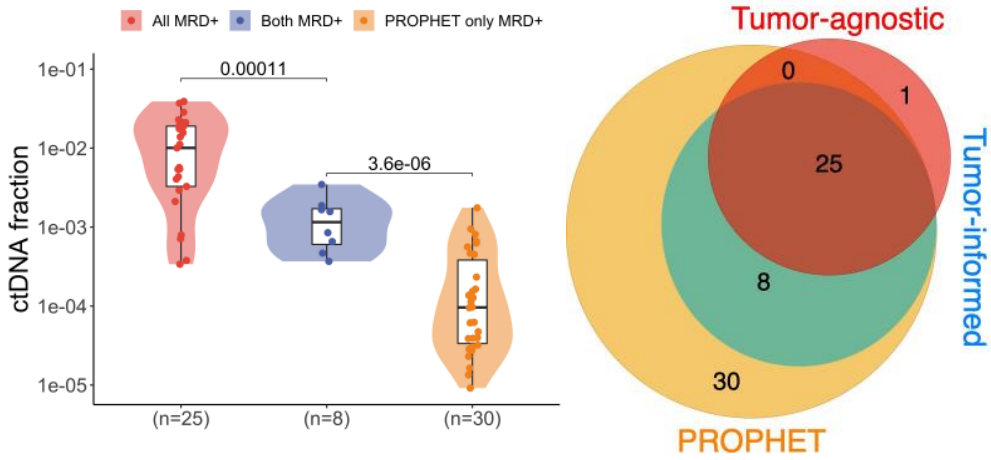
(a)

Sensitivity of pre-operative plasma



(b)

ctDNA fraction distribution of MRD (+) samples detected by different methods

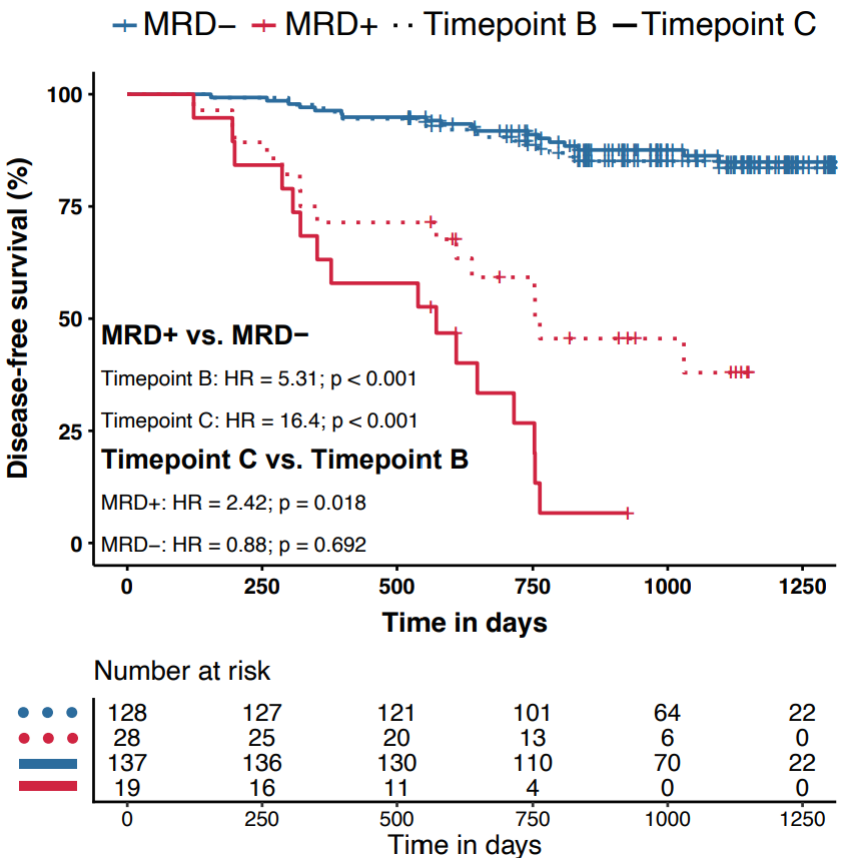


- For pre-operative plasma from patients with different clinical stages, brPROPHET has a higher sensitivity than the other two methods
- The median ctDNA fraction of the 30 patients detected by PROPHET alone was significantly lower than the 25 patients detected by all three MRD assays

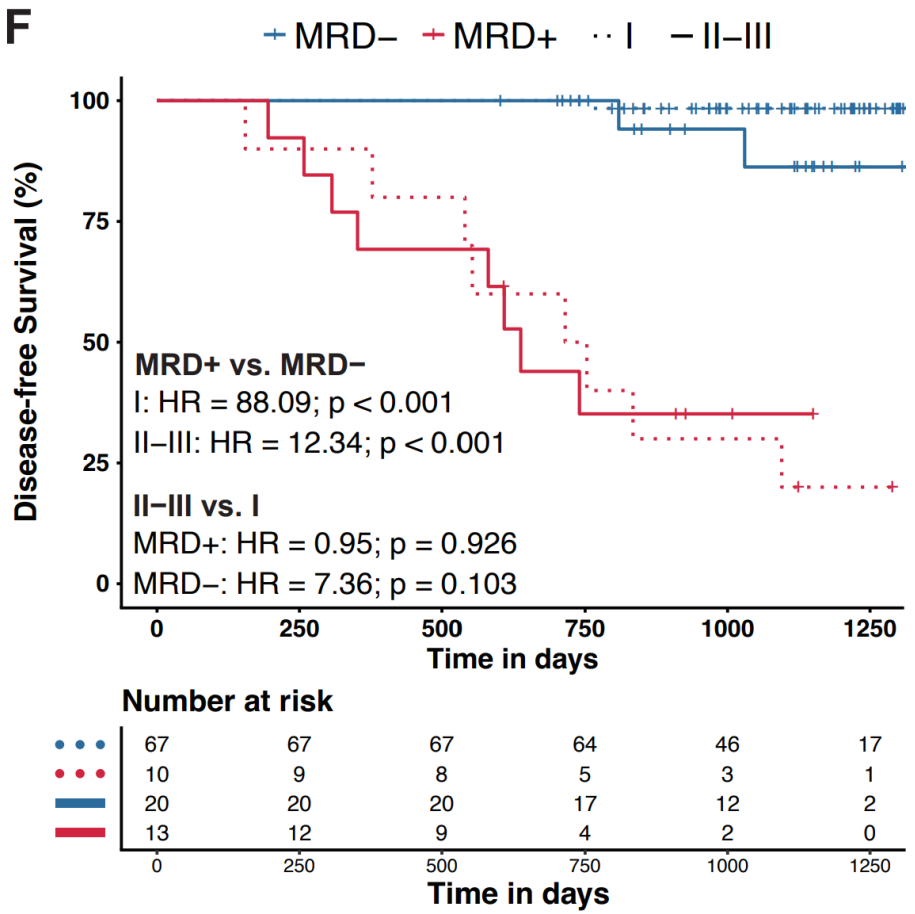
The patient-specific brPROPHET has a higher sensitivity than the two fixed panel detection methods

Source: Chen et al., Individualized tumor-informed circulating tumor DNA analysis for postoperative monitoring of non-small cell lung cancer, Cancer Cell, Sep 2023

brPROPHET™ shows strong prognostic value in post-surgery NSCLC patients



Prognostic analysis at Landmark time points



Longitudinal MRD analysis

Source: Chen et al., Individualized tumor-informed circulating tumor DNA analysis for postoperative monitoring of non-small cell lung cancer, Cancer Cell, Sep 2023

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Early detection

Burning Rock's multi-cancer early detection technology

Competitive technology

Methylation + machine learning to overcome challenges of low ctDNA abundance

nature biomedical engineering

ARTICLES

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

Check for updates

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

Multi-cancer validation data

ELSEVIER

Annals of Oncology

Available online 26 February 2023

In Press, Journal Pre-proof What's this?

Original Article

Unintrusive multi-cancer detection by circulating cell-free DNA methylation sequencing (THUNDER): development and independent validation studies

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

ESMO 2022

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

Regulatory breakthrough

FDA breakthrough device designation granted

国家药品监督管理局 National Medical Products Administration China NMPA breakthrough designation granted

Product development roadmap

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)

6-cancer
2018 – 2020

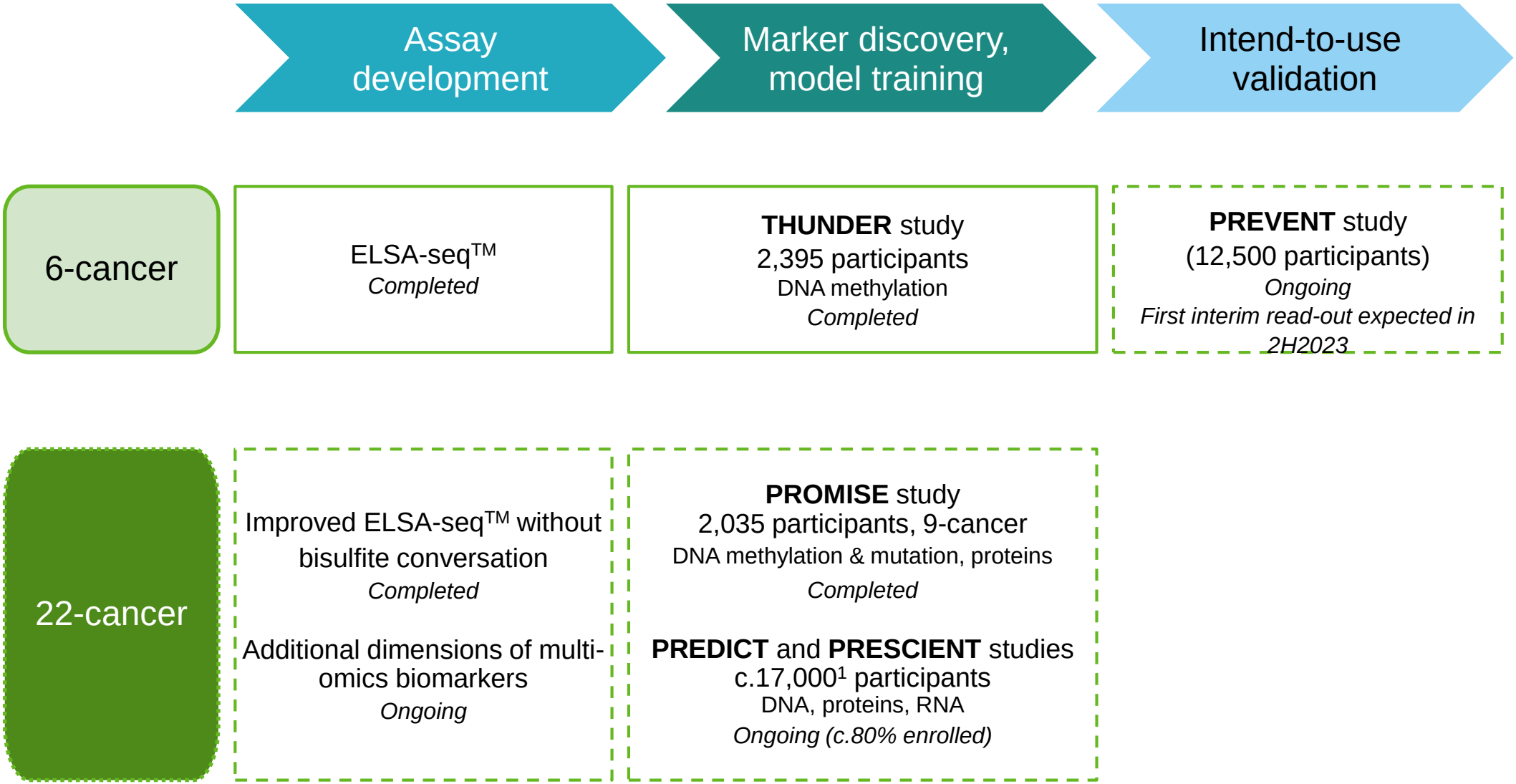
- Lung, CRC, HCC, Ovarian, Pancreatic, Esophageal
- THUNDER study (N=2,385) completed, 98.9% specificity and 69.1% sensitivity, top-2 predicted origin accuracy of 91.7% (independent validation cohort)
- PREVENT study (prospective and interventional, IU population) ongoing

22-cancer
2019 – Ongoing

- Covering 88% of China’s cancer incidence
- PROMISE study (N=2,035) completed, improved performance vs. 6-cancer test
- PREDICT and PRESCIENT studies ongoing

Clinical programs

One of the largest datasets globally, prospectively enrolled, across a large number of cancer types / stages



Note：

[1] Total number of subjects for Predict and Prescient studies.

Running the largest clinical programs in China supported by top physicians

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 basis[1]
  - Ranked top 5 in the 2019 China's general hospital rankings[2]

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 Sciences[3]
  - 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

- Fellow of the Chinese Academy of Sciences
- President of CHCAMS

Prof. Jie Wang

- Head of the Dept. of Medicine, CHCAMS

PREVENT

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

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Appendix 1

Early detection

Product Development Roadmap

Proof-of-concept
2016 – 2019

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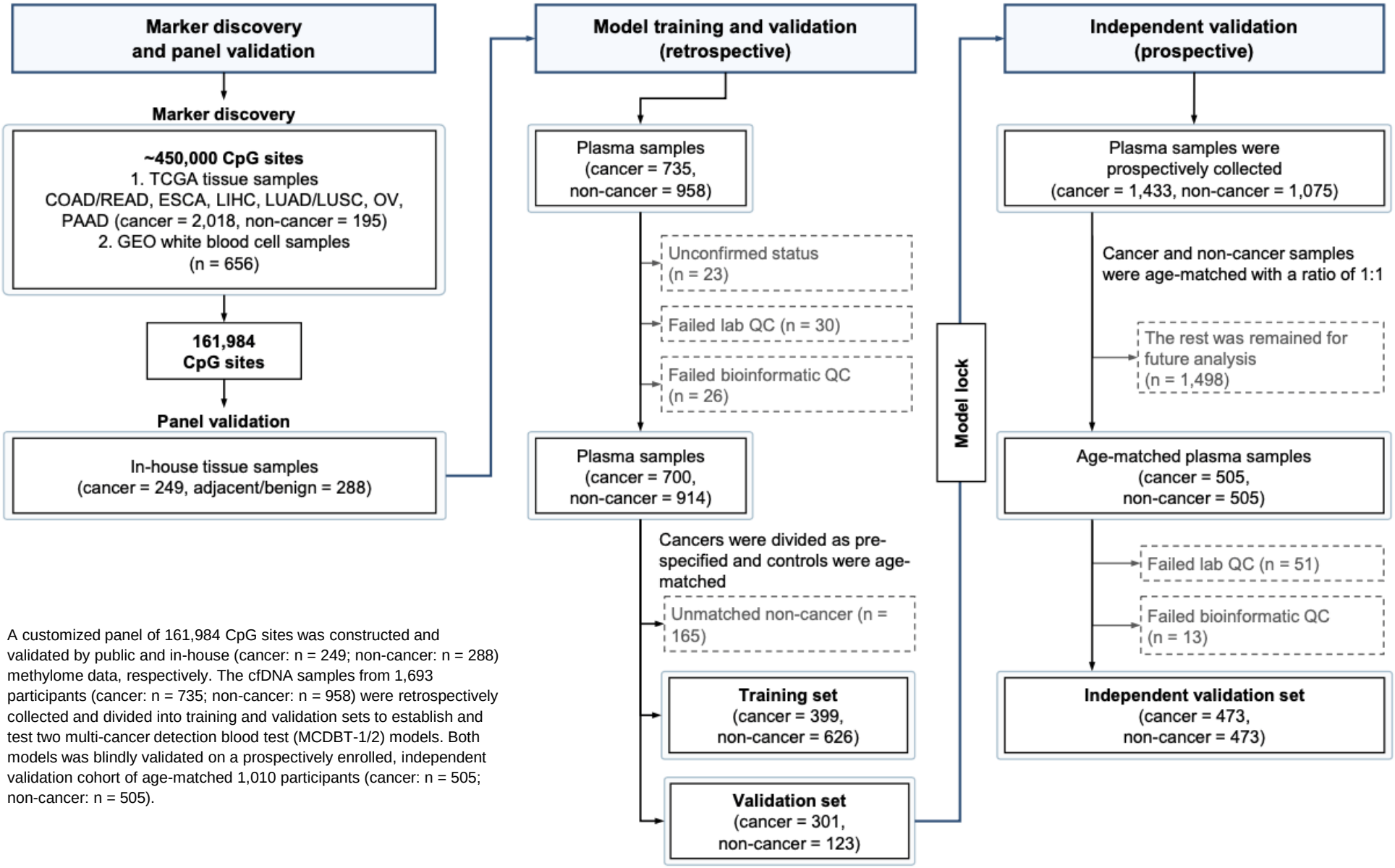
6-cancer test marker discovery and model training

The THUNDER study, 2395 participants



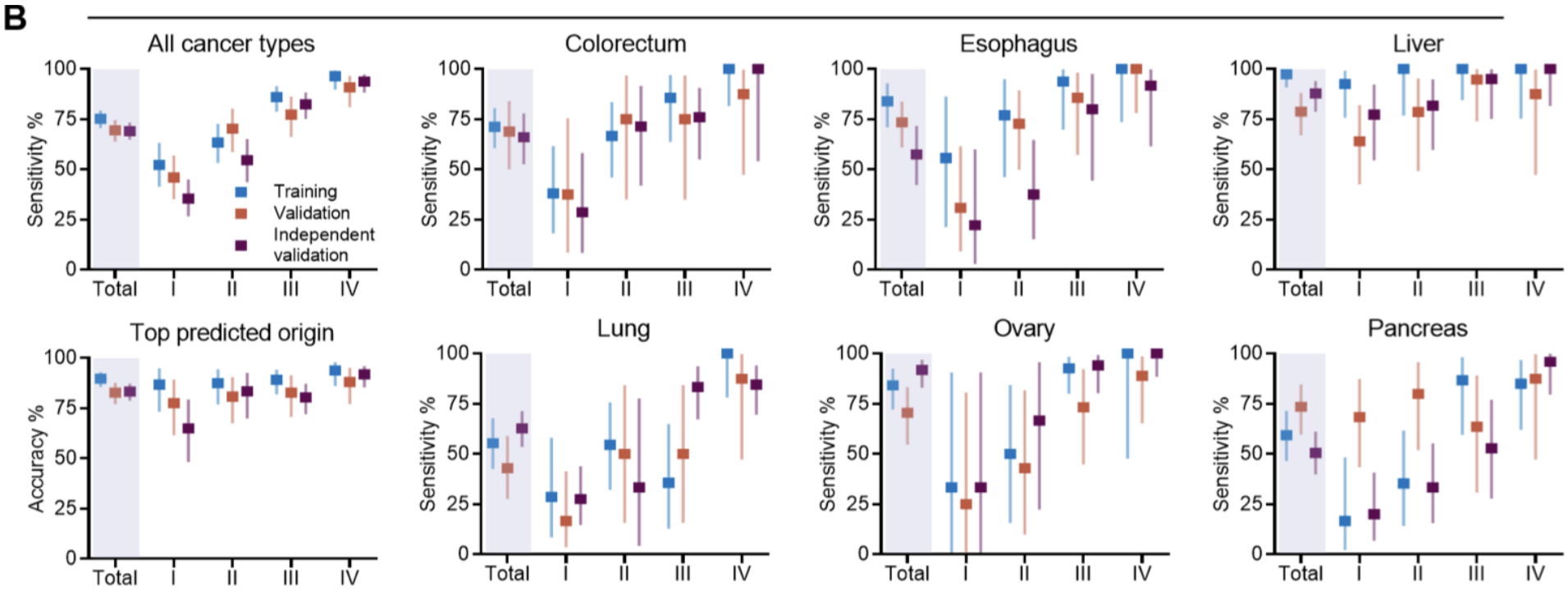
A customized panel of 161,984 CpG sites was constructed and validated by public and in-house (cancer: n = 249; non-cancer: n = 288) methylome data, respectively. The cfDNA samples from 1,693 participants (cancer: n = 735; non-cancer: n = 958) were retrospectively collected and divided into training and validation sets to establish and test two multi-cancer detection blood test (MCDBT-1/2) models. Both models was blindly validated on a prospectively enrolled, independent validation cohort of age-matched 1,010 participants (cancer: n = 505; non-cancer: n = 505).

Source: Gao et al., Unintrusive multi-cancer detection by circulating cell-free DNA methylation sequencing (THUNDER): development and independent validation studies, ASCO 2022

6-cancer test, detection-of-cancer performance in case-control cohorts

The THUNDER study

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.



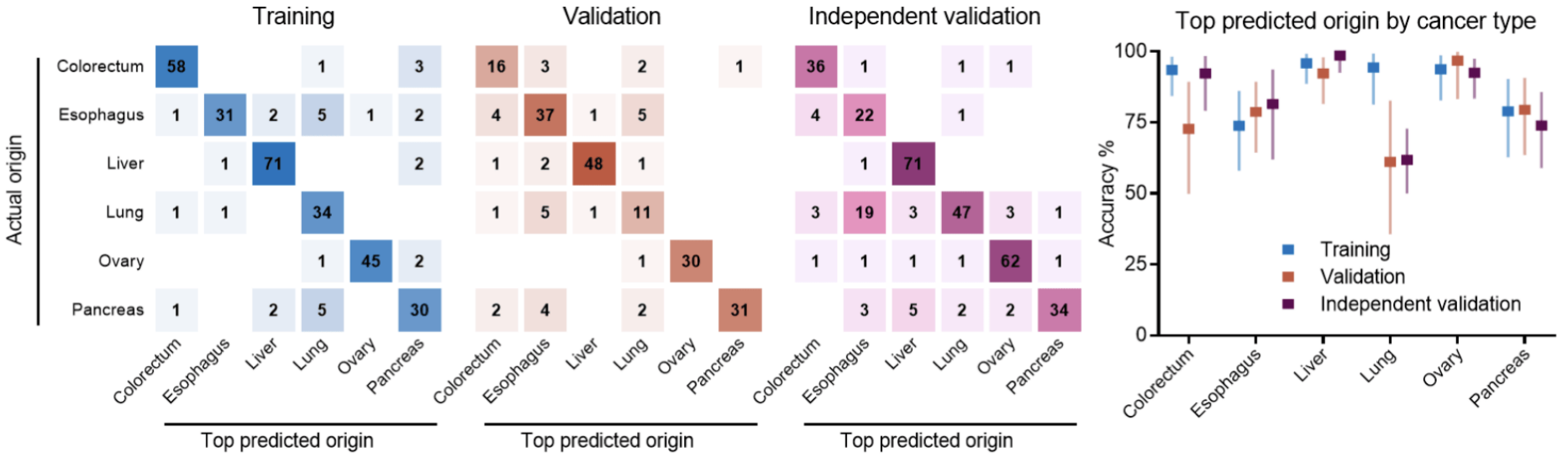
Data set	Specificity (%)	Sensitivity (%)	Accuracy of top predicted origin (%)	Accuracy of top two predicted origins (%)
Training set	99.7 (98.9-100.0)	75.2 (70.6-79.4)	89.7 (85.7-92.9)	94.7 (91.5-96.9)
Validation set	100.0 (97.0-100.0)	69.4 (63.9-74.6)	82.8 (77.0-87.6)	89.4 (84.5-93.3)
Independent validation set	**98.9 (97.6-99.7)**	**69.1 (64.8-73.3)**	**83.2 (78.7-87.1)**	**91.7 (88.2-94.5)**

Source: Gao et al., Unintrusive multi-cancer detection by circulating cell-free DNA methylation sequencing (THUNDER): development and independent validation studies, ASCO 2022

6-cancer test, top-predicted-origin performance in case-control cohorts

The THUNDER study

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.



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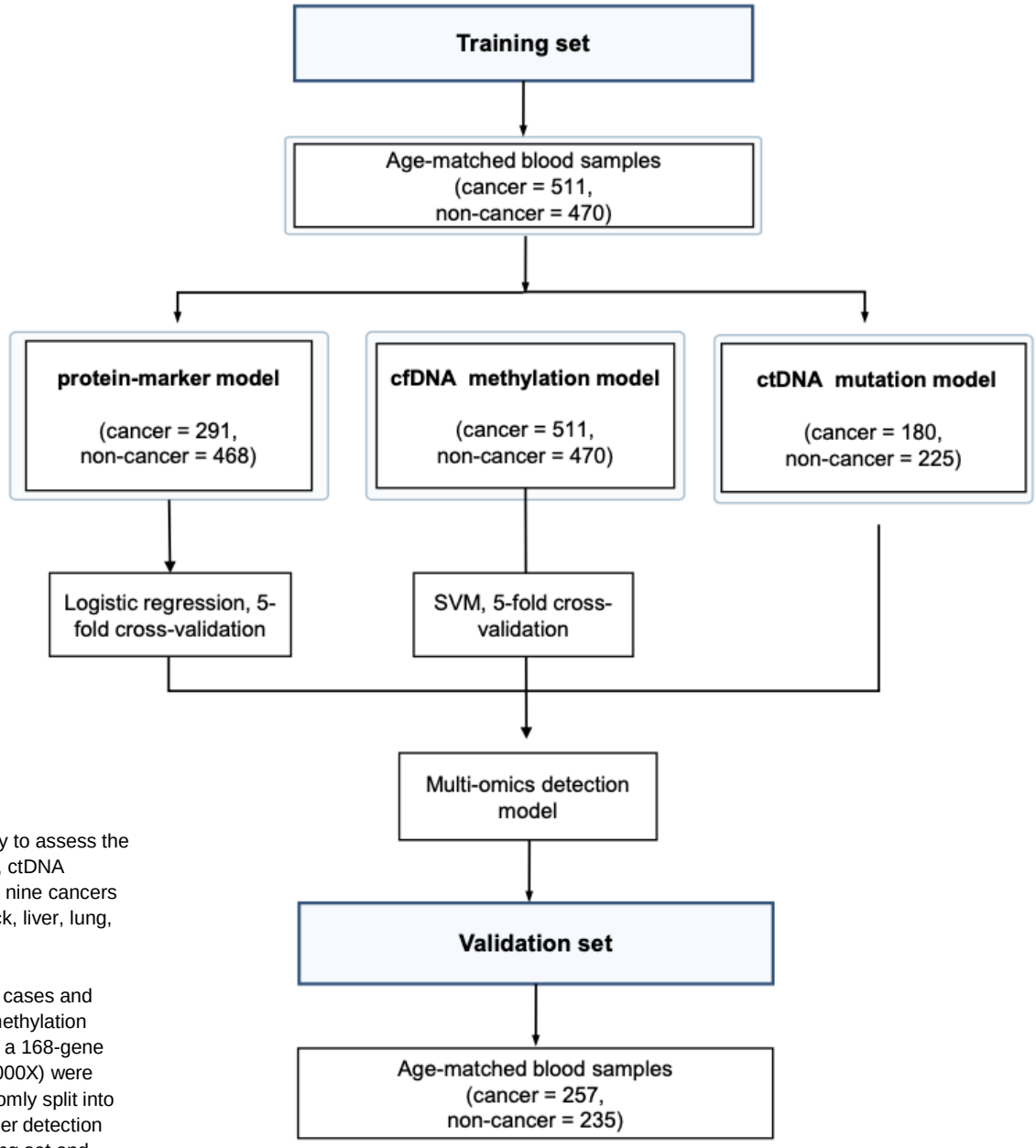
- Lung, CRC, HCC, Ovarian, Pancreatic, Esophageal
- THUNDER study (N=2,385) completed, 98.9% specificity and 69.1% sensitivity, top-2 predicted origin accuracy of 91.7% (independent validation cohort)
- PREVENT study (prospective and interventional, IU population) ongoing

22-cancer
2019 – 2022

- Covering 88% of China’s cancer incidence
- PROMISE study (N=2,035) completed, improved performance vs. 6-cancer test
- PREDICT/PRESCIENT studies ongoing

9-cancer test, multi-omics model

The PROMISE study



PROMISE is a prospective multicenter case-control study to assess the performance of multi-omics including cfDNA methylation, ctDNA mutation and protein biomarkers in the early detection of nine cancers in the biliary tract, colorectum, esophagus, head and neck, liver, lung, ovary, pancreas and stomach.

Blood samples were prospectively collected from cancer cases and non-cancer controls. A targeted cell-free DNA (cfDNA) methylation panel of ~490,000 CpG sites (1,000X) by ELSA-seq and a 168-gene mutation panel (35,000X, matched white blood cells:10,000X) were sequenced. Age-matched cases and controls were randomly split into training (n = 981) and test sets (n = 492). The multi-cancer detection blood test (MCDBT) models were developed in the training set and then validated in the test set.

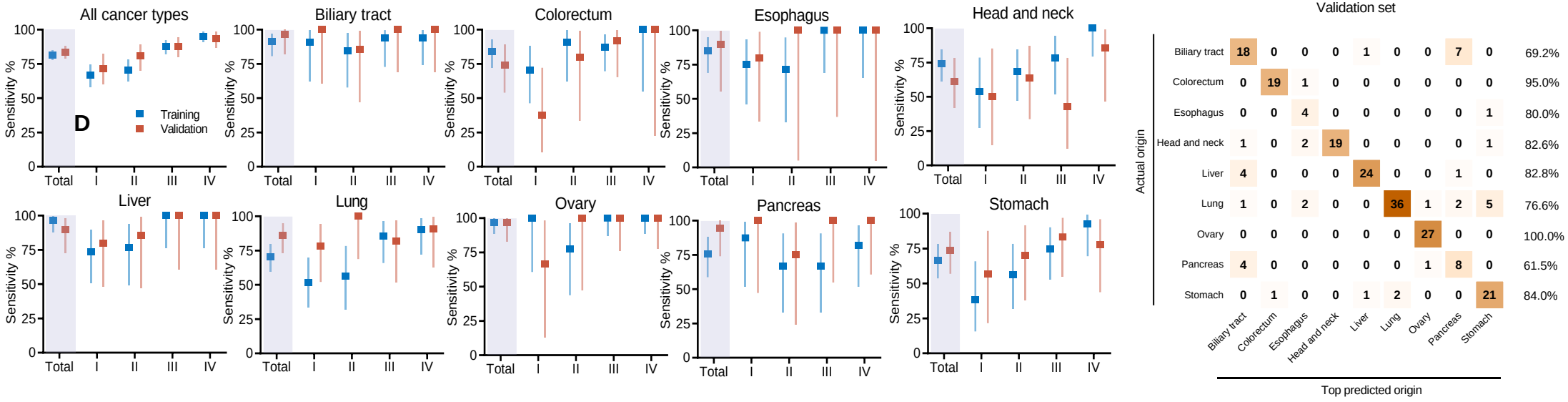
Source: Gao et al., a multi-cancer early detection model based on liquid biopsy of multi-omics biomarkers, ESMO 2022

9-cancer test multi-omics model performance

The PROMISE study

	Cancer (n)	Non-cancer (n)	Specificity (%)	Sensitivity (%)	Accuracy of top predicted origin (%)
Training	470	511	97.9% (96.1%-99.0%)	81.7% (78.1%-84.9%)	86.6% (83.0%-90.0%)
Validation	257	235	98.3% (96.6%-99.4%)	83.7% (79.0%-88.0%)	81.9% (76.0%-87.0%)

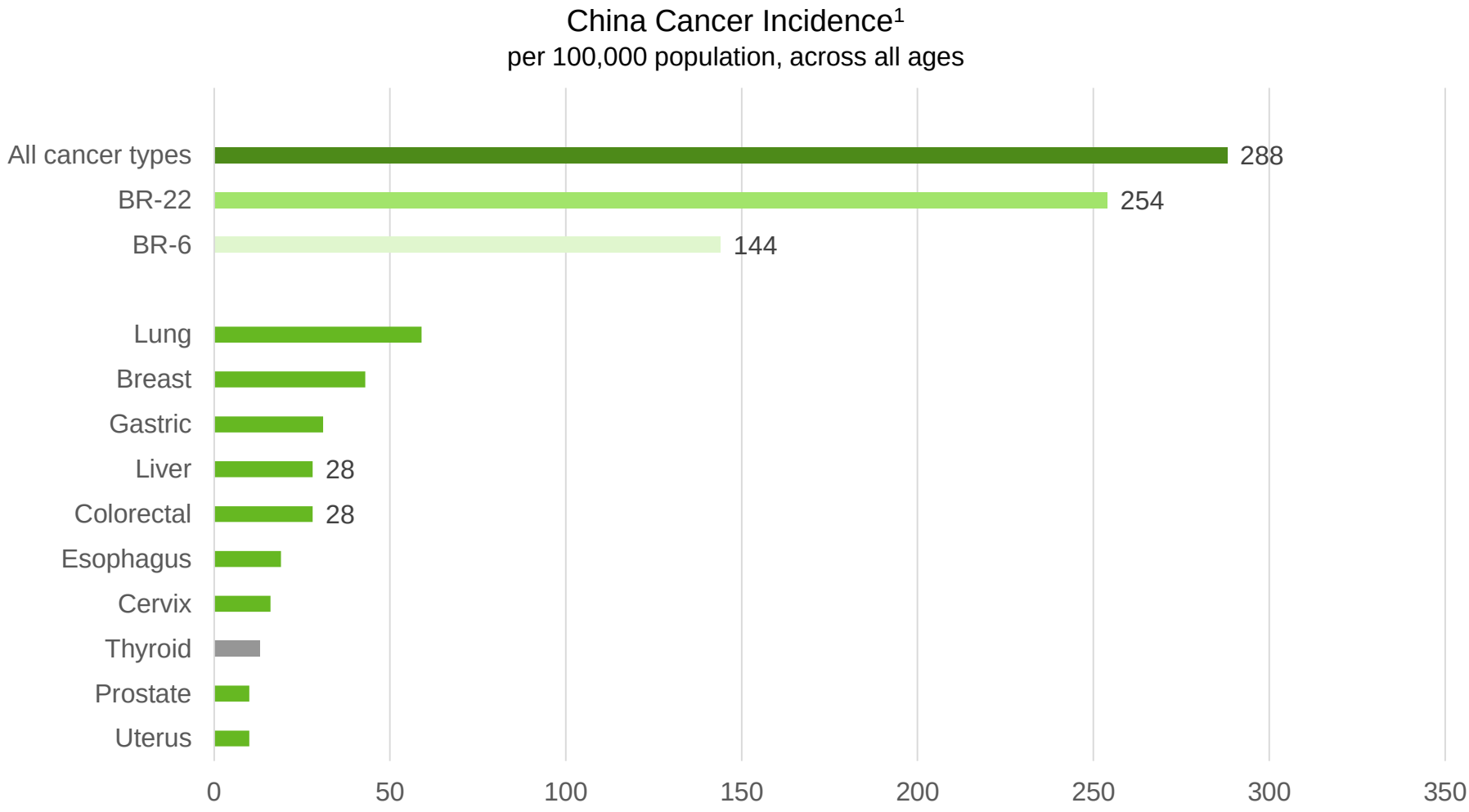
	Multi-omics	Methylation	Mutation	Protein
Specificity (95% CI)	98.3% (96.6%–99.4%)	99.1% (97.3%–99.8%)	99.6% (97.9%–100.0%)	99.6% (98.7%–100.0%)
Sensitivity (95% CI)	83.7% (78.6%–88.0%)	79.0% (73.5%–83.8%)	49.4% (41.9%–57.0%)	47.8% (40.8%–54.9%)



- PROMISE demonstrated 83.7% sensitivity and 98.3% specificity for 9 cancers
- Methylation contributed >90% of the total sensitivity, while protein and mutation collectively provided <10% additional positive detections

Source: Gao et al., a multi-cancer early detection model based on liquid biopsy of multi-omics biomarkers, ESMO 2022

Burning Rock's 22-cancer test covers 88% of China's cancer incidence



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

Leadership in multi-cancer early detection

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

	Challenges	BNR position
1 Technology	**Low amount of cancer signal** in the circulating bloodstream, much more challenging compared to 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 that has launched studies with over 10,000+ subjects
3 Regulatory	**First-in-class** with no established regulatory pathway	**Leading regulatory capability in China** • Exploring possible pathway, leveraging experience through the country's first NMPA-approved NGS kit
4 Commercial	**Unprecedented product**	**Multi-pronged approach** • Initially working with hospitals' health check-up departments, leveraging synergy from in-hospital therapy selection business

燃石医学
Burning Rock Dx

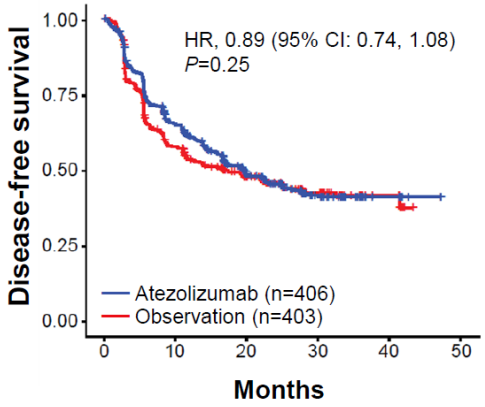
Appendix 2

MRD

How do MRD studies advance utility

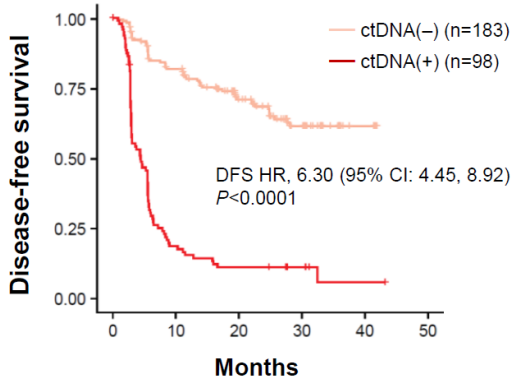
Example 1: IMvigor010, enrich the high-risk group and "tune-up" adjuvant treatment

ITT

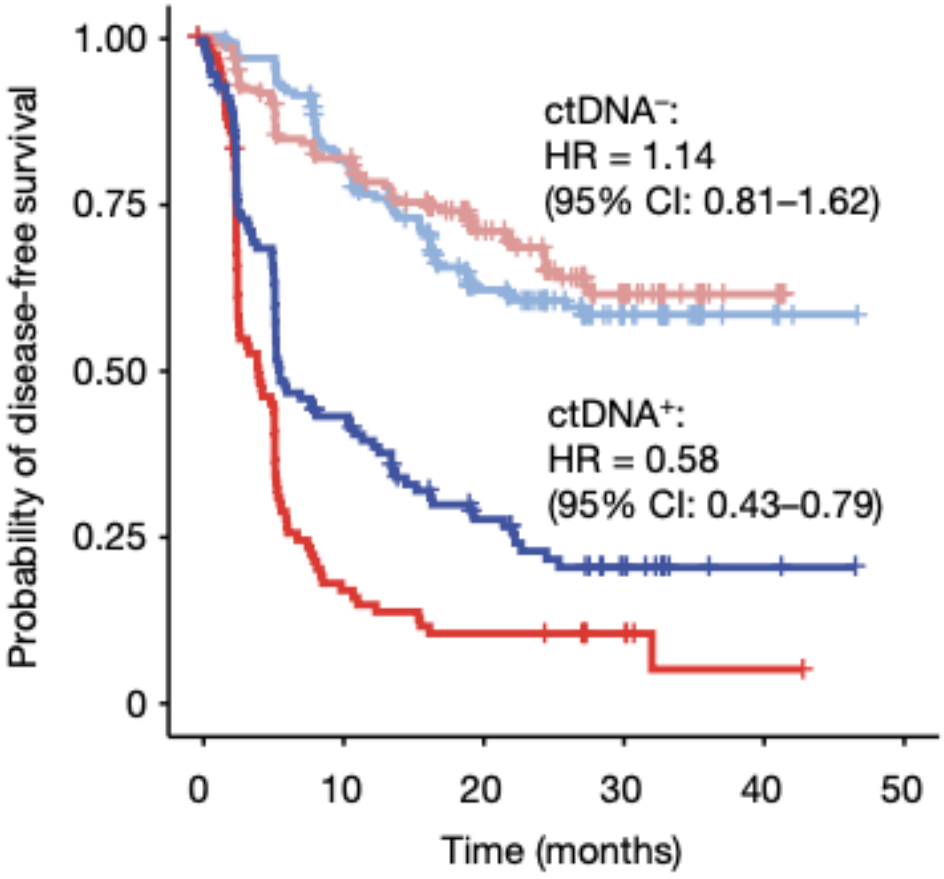


Atezo, MIUC Adjuvant Therapy
"All comers" demonstrated NO efficacy
TMB/PD-L1 showed NO prediction

Observation arm



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

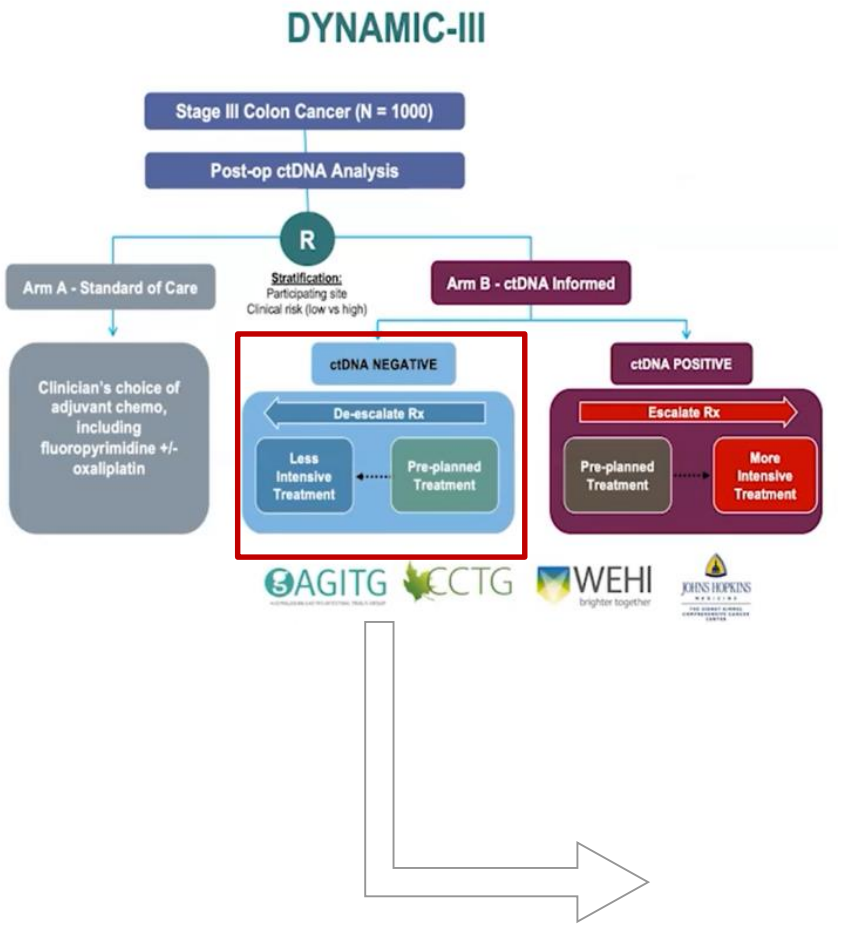


Indeed, only baseline MRD+ pts showed benefit

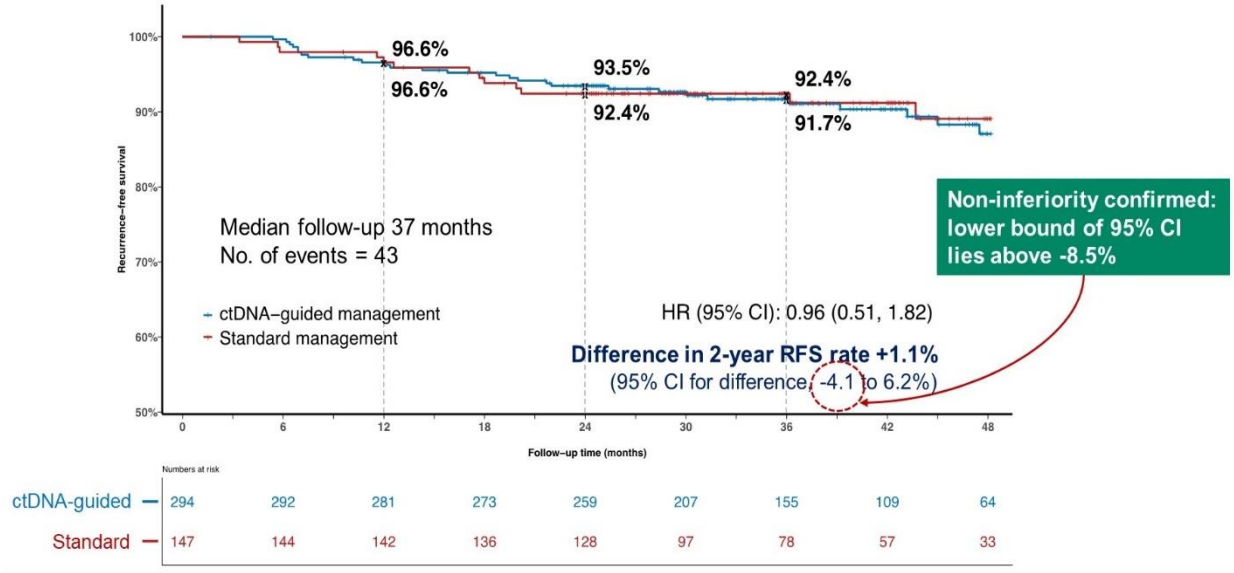
Nature. 2021 Jun 16. doi: 10.1038/s41586-021-03642-9.

How do MRD studies advance utility

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

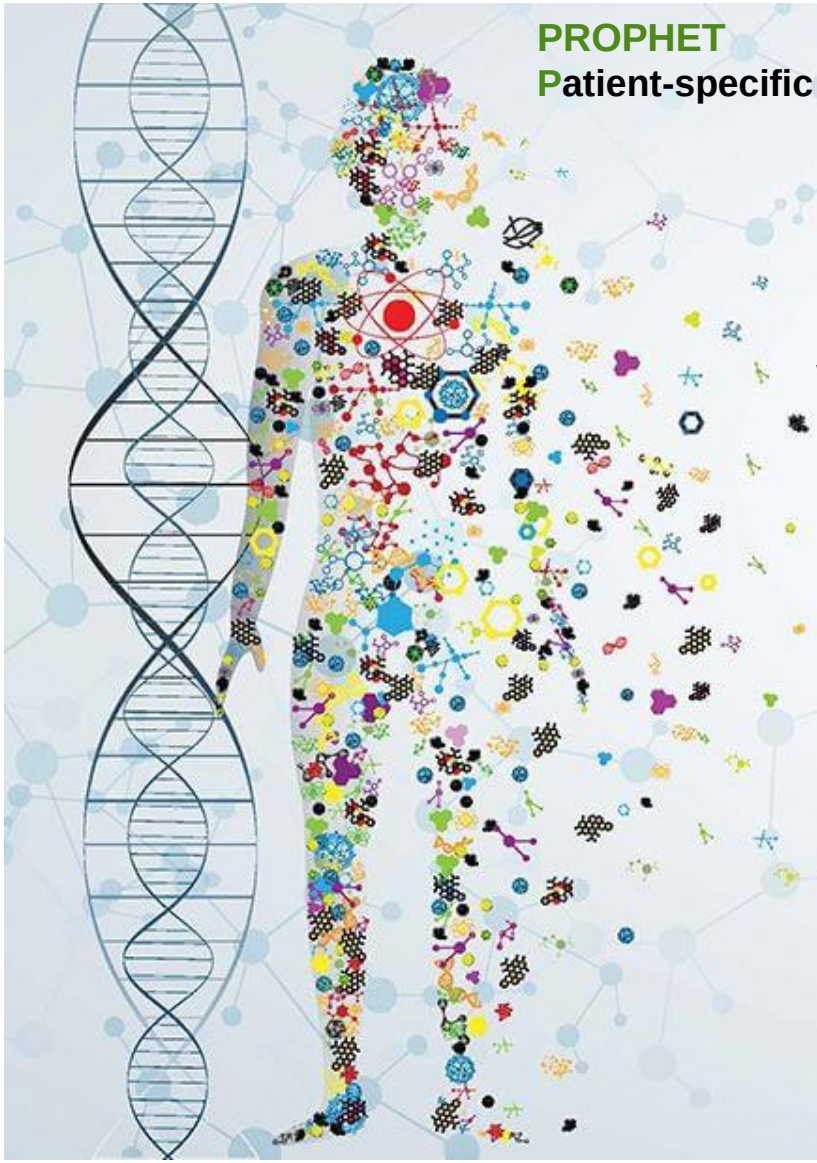


ctDNA-guided vs standard



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%)

brPROPHET™ – Burning Rock's MRD solution



PROPHET
Patient-specific pROgnostic and Potential tHErapeutic marker Tracking

Whole Exome Sequencing Tumor Profiling with add-on region for SV detection

SNV/SV/CNV/MSI/HLA/Therapy Selection

Personalized Panel

Intelligent selection of 50 tracking sites

brPROPHET MRD Assay

100,000x Raw Depth/UMI error correction/Tumor Fraction Estimation

Gastric cancer cohort publication at AACR 2023

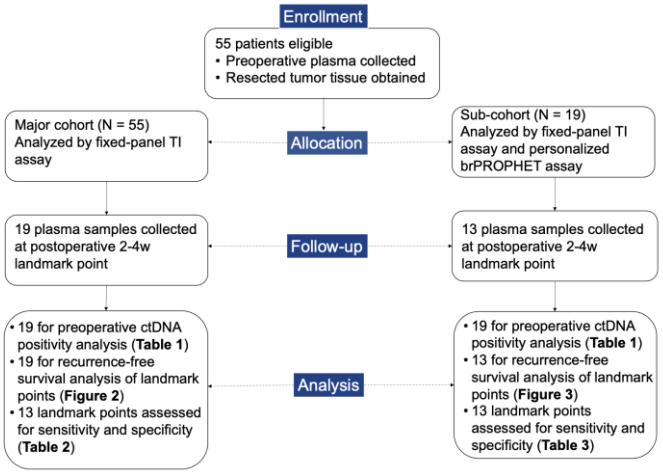
Circulating tumor DNA - based molecular residual disease predicts relapse in patients with resectable gastric cancer

Pei Xue[1], Yanfei Shao[1], Xueliang Zhou[1], Haiyan Li[2], Yang Wang[2], Chenyang Wang[2], Hao Zhang[2], Bing Li[2], Shuo Shi[2], Haiwei Du[2], Jing Sun[1]
1. Department of General Surgery, Ruijin Hospital, Shanghai Jiao Tong University, Shanghai, China
2. Burning Rock Biotech, Guangzhou, China

2023 AACR
#1037

AACR American Association for Cancer Research ANNUAL MEETING 2023
APRIL 14-19 • #AACR23

Study cohort



Personalized assay significantly out-performs fixed panels

The ctDNA+ rate of preoperative samples detected by fixed panel and personalized brPROPHET™ assays

ctDNA⁺ rate	Stage I	Stage II	Stage III	Overall
Fixed panel	0% (0/4)	0% (0/3)	58.3% (7/12)	36.8% (7/19)
brPROPHET	100% (4/4)	66.7% (2/3)	91.7% (11/12)	89.5% (17/19)

Postoperative prognosis

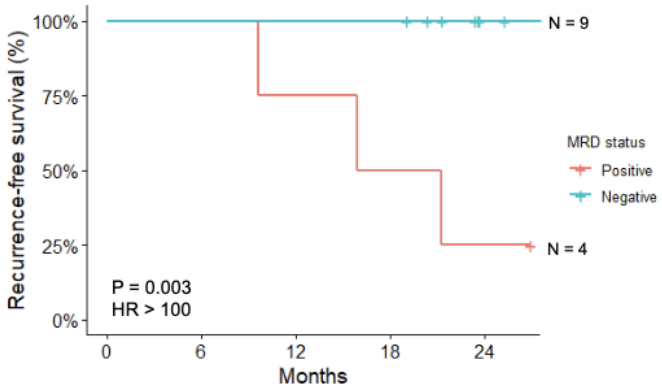


Table 3. The performance of MRD status detected by brPROPHET in predicting tumor recurrence.

MRD status	Tumor recurrence		
	+	-	
+	3	1	PPV=75%
-	0	9	NPV=100%
	Sensitivity=100%	Specificity=90%	

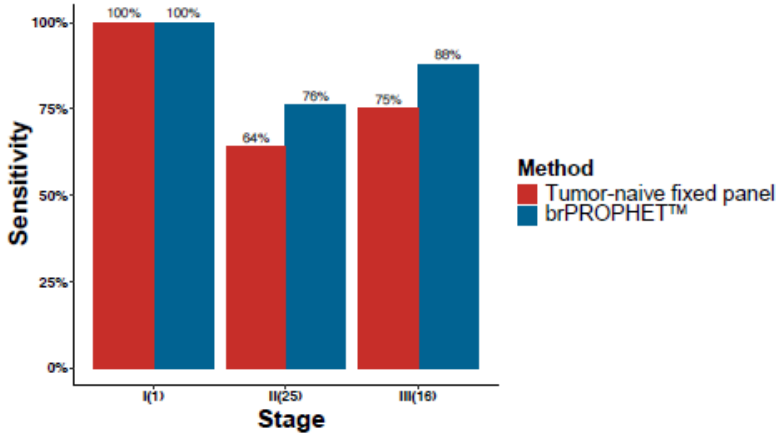
PPV, positive predictive value; NPV, negative predictive value.

Source: Xue et al., Circulating tumor DNA based molecular residual disease predicts relapse in patients with resectable gastric cancer, #1037 AACR 2023

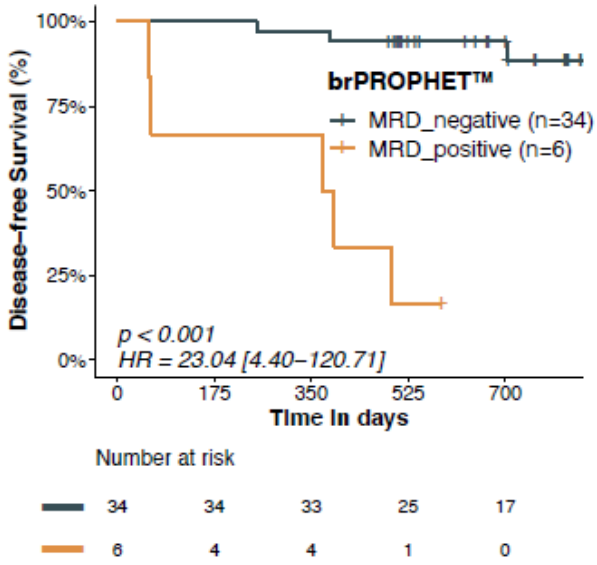
Colorectal cancer cohort publication at AACR 2022

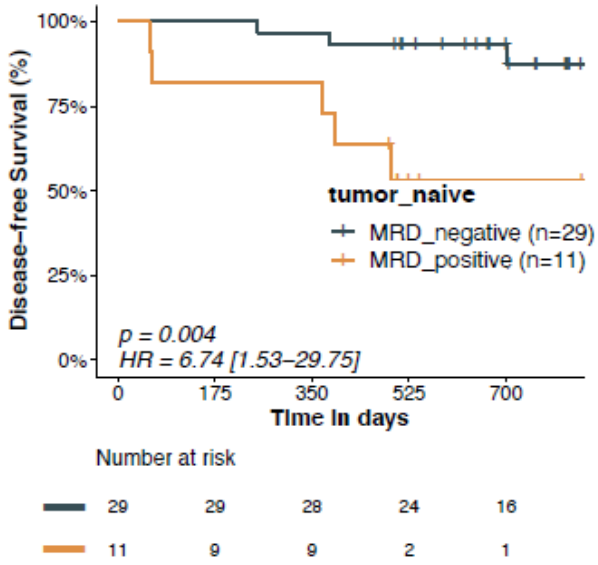
Session OPO.PR02.01 - Clinical Prevention, Early Detection, and Interception

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

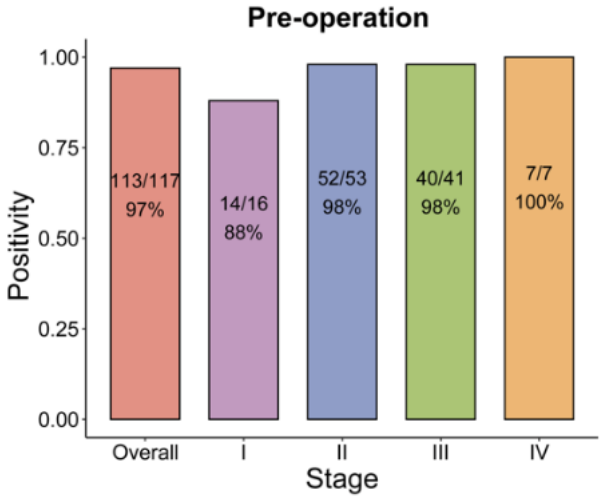




Second colorectal cancer cohort publication at ASCO GI 2023

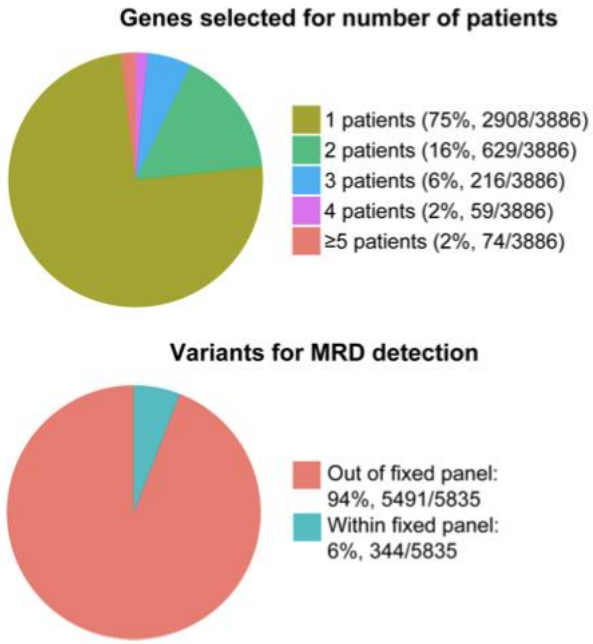
brPROPHET™ has high detection sensitivity

Preoperative ctDNA was detected in 97% (113/117) of the patients with 88% (14/16), 98% (52/53), 98% (40/41), and 100% (7/7) in stage I, II, III and IV, respectively



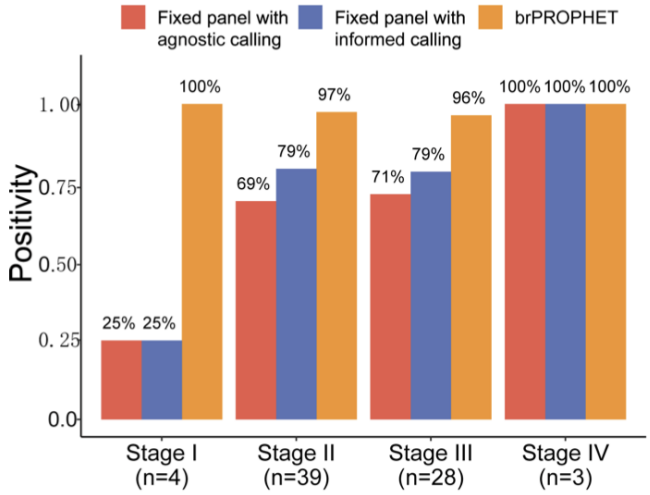
Most mutation variants fall outside of fixed panels

Only 6% of designed variants were included in the fixed panel. 75% of genes selected for panel design were private to a specific patient.



brPROPHET™ significantly out-performs fixed panels

Preoperative ctDNA was detected in 97% (113/117) of the patients with 88% (14/16), 98% (52/53), 98% (40/41), and 100% (7/7) in stage I, II, III and IV, respectively



Patients: A total of 117 patients (stage II/III 53 [45.3%] / 41 [35.0%]) who received surgery were analyzed. A subset of 74 patients were analyzed for comparisons of different methods.
Samples: Tumor tissue samples were collected at the surgery. Plasma samples collected at baseline, landmark 7-day and 1-month, and longitudinal points were analyzed.

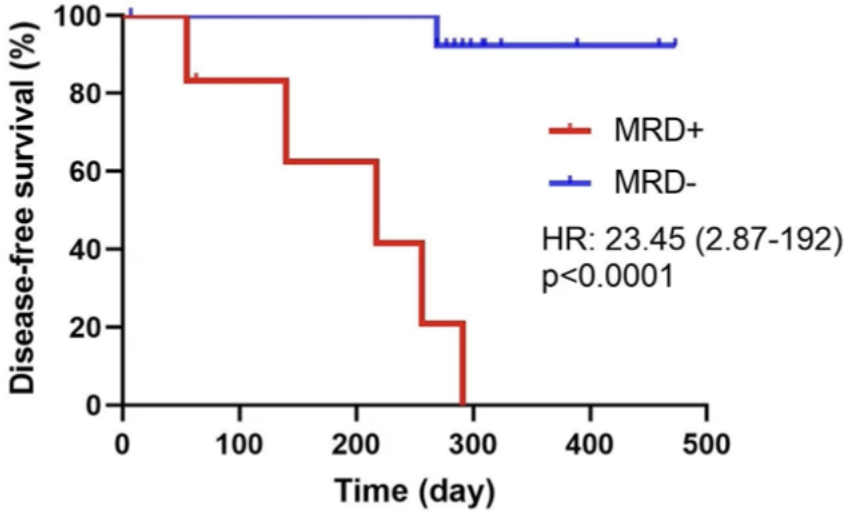
Source: Cao et al., Patient-specific tumor-informed circulating tumor DNA analysis for molecular residual disease detection in surgical patients with stages I-IV colorectal cancer, ASCO GI 2023

Pancreatic cancer cohort publication at ASCO GI 2023

Table 1: ctDNA detection at serial timepoints

	Baseline (Day 0)	Timepoint A (Day 7)	Timepoint B (Day 30)	Timepoint C (During AT)	Follow-ups
Positive	20	2	1	2	4
Negative	0	16	9	12	5
Positive Rate	**100%**	**11.1%**	**10%**	**14.3%**	**44.4%**

Figure 1: Longitudinal MRD detection is associated with shorter disease-free survival



Patients: A total of 20 patients (stage I/II 10 [50.0%] / 9 [45.0%]) were analyzed. 13 (65.0%) patients were treated with adjuvant therapy (AT) after surgery.
Samples: Tumor tissue samples were collected at the surgery. Plasma samples collected at baseline (n=20), landmark 7-day (n=18) and 1-month (n=10), and longitudinal points (n=23) were analyzed. Patients were followed for a median of 302 days.

Source: Wang et al., Patient-specific tumor-informed circulating tumor DNA (ctDNA) assay predicts cancer recurrence in patients with resected pancreatic cancer, ASCO GI 2023

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Appendix 3

Therapy selection

NMPA approved NGS panels

NMPA approved testing kits by major NGS-focused companies[1]

	First NMPA-approved kit	Second NMPA-approved kit
燃石医学 Burning Rock Dx	EGFR, ALK, BRAF, KRAS Approved in Jul **2018** First approved NGS kit in China	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

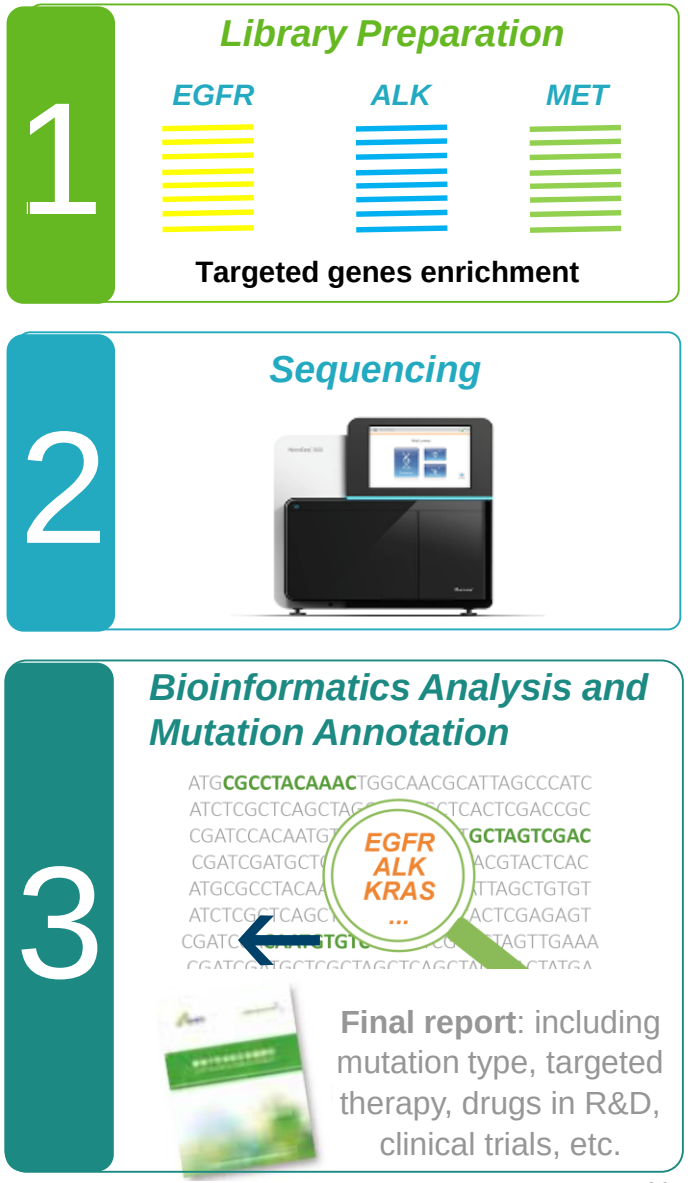
- Only 30ng DNA input required, applicable to small tissue samples
- First NMPA approved NGS kit with CNV[2] mutation type, with MET exon14 skipping

Notes:
[1] 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

	NGS
Throughput	**High**-throughput
Efficiency	**Parallel** testing
Cost	**Low-cost** per biomarker
Biomarker Profiling	**Comprehensive** profiling & **superior** accuracy
Fit for Liquid Sample	**Liquid availability**



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%

of ctDNA

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nature biotechnology

ARTICLES

https://doi.org/10.1038/s41587-021-00857-z

Check for updates

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

Liquid Biopsy

- 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

25 | Ef_10
50 | Ep_25

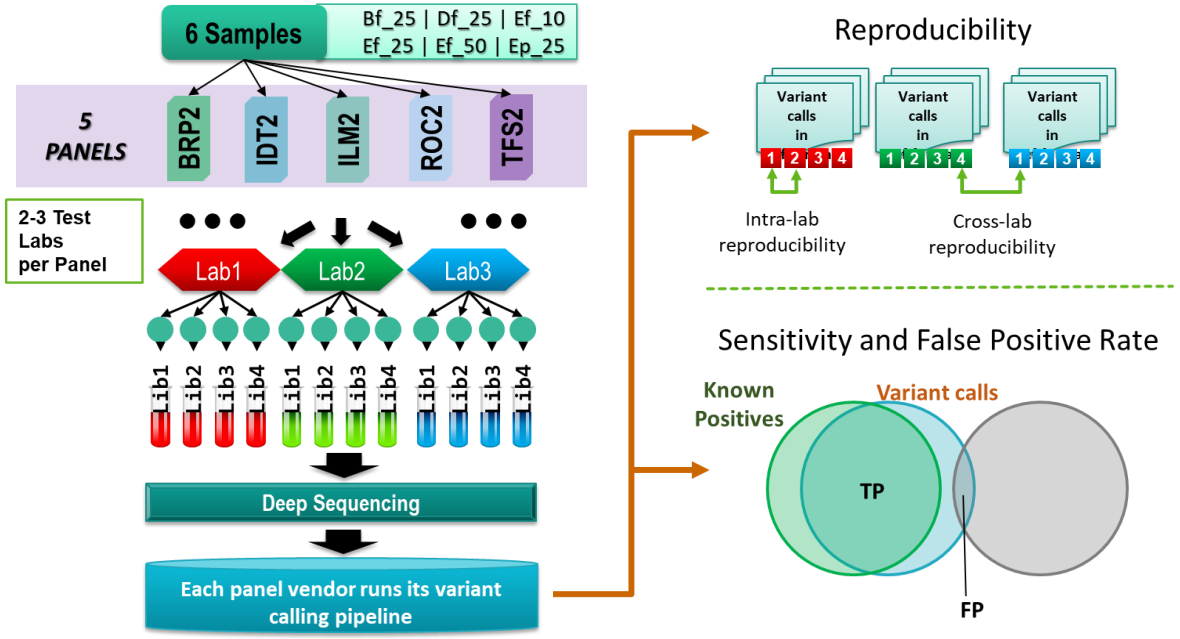
TFS2

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

Source:
Slides from “Establishing the analytical validity of circulating tumor DNA sequencing for precision oncology“, 5th Annual Liquid Biopsy for Precision Oncology Summit, Feb 2021
Further information in Appendix 2

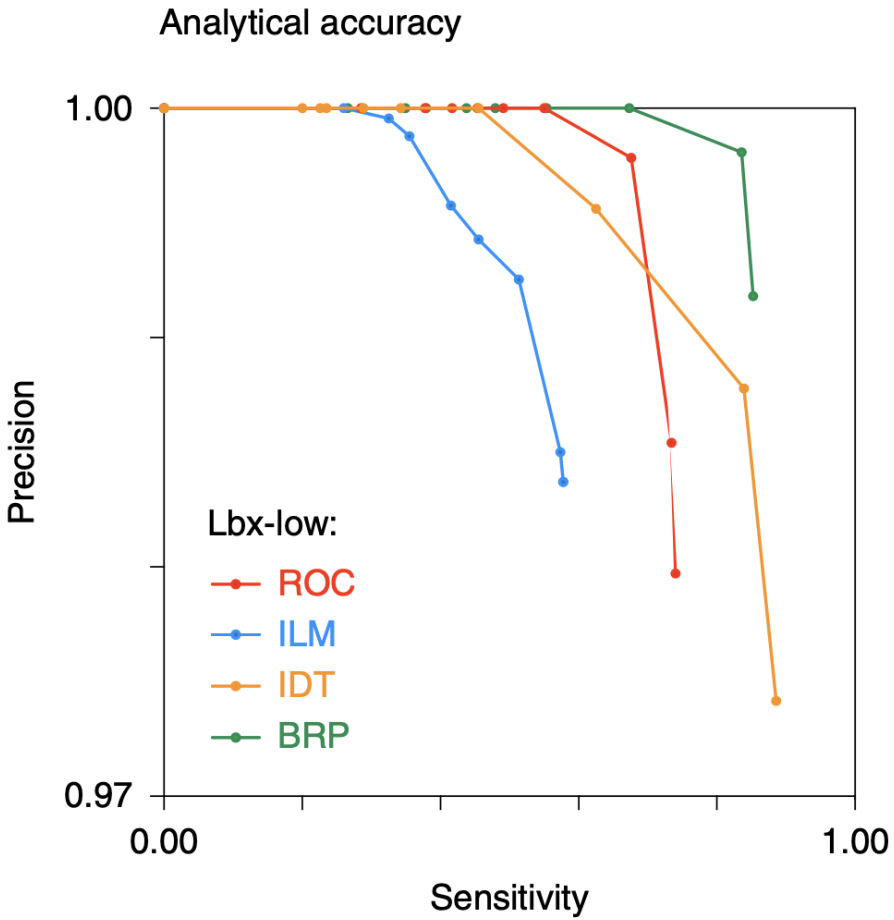
Participating assays and study design

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



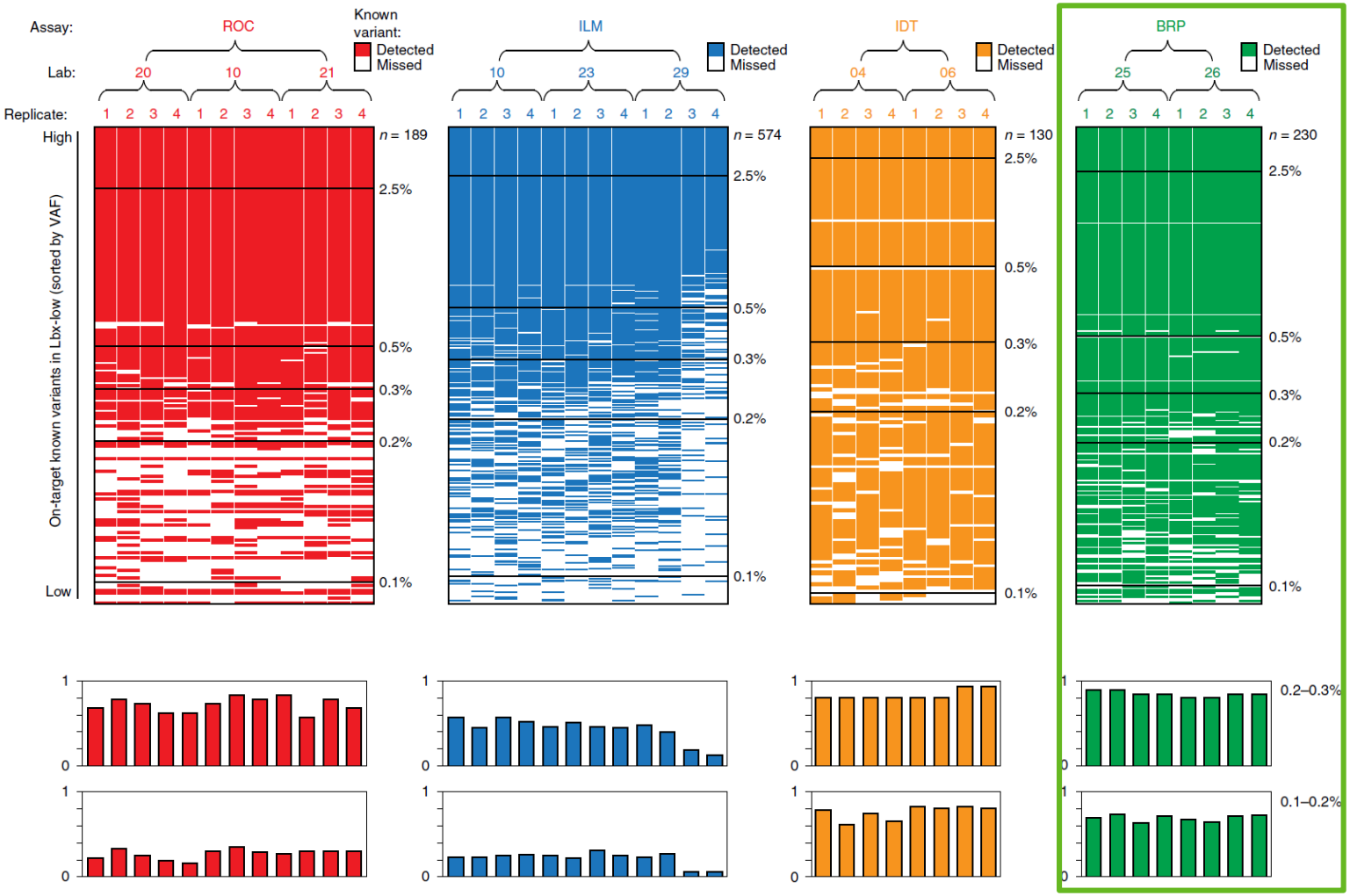
Assay	Known negatives (kb)	FPs per replicate (mean [range])	FP-rate (FP / kb) at specified VAF threshold > 0%	> 0.1%	> 0.5%
ROC	47.1	2.91 [1-6]	0.061	0.044	0.000
ILM	133	5.25 [2-10]	0.039	0.039	0.008
IDT	39.3	2.75 [0-6]	0.070	0.057	0.000
BRP	53.4	1.65 [0-5]	0.030	0.007	0.000

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). "

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

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."

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