

# Moving Early Diagnostics to Global Patients

- Advancing equitable cancer care through innovation

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SeekIn

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Amsterdam

# Cancer

## - The emperor of all maladies

- A huge burden globally and will stay with human forever
- Significant progresses in the past 100 years especially in the last two decades
- Deep knowledge and improved outcome
- Precision cancer management has been taking place in top academic centers
- Expenditure becomes another burden on the public healthcare systems and the families
- Accessibility, affordability and equity are huge challenges

# The perspectives

Advancing equitable cancer care through innovation

- Scalable
- Cost-effective
- Convenience

Prevention, Diagnosis and Treatment

# Blood tests for getting right treatments at earlier time



Indicated subjects: Healthy population

Post-op patients of radical resection

Late-stage patients

Launch date: Nov. 2018

May 2019

Aug. 2020



One-Size-Fits-All

# SeekIn has dedicated to figure out solutions to key challenges

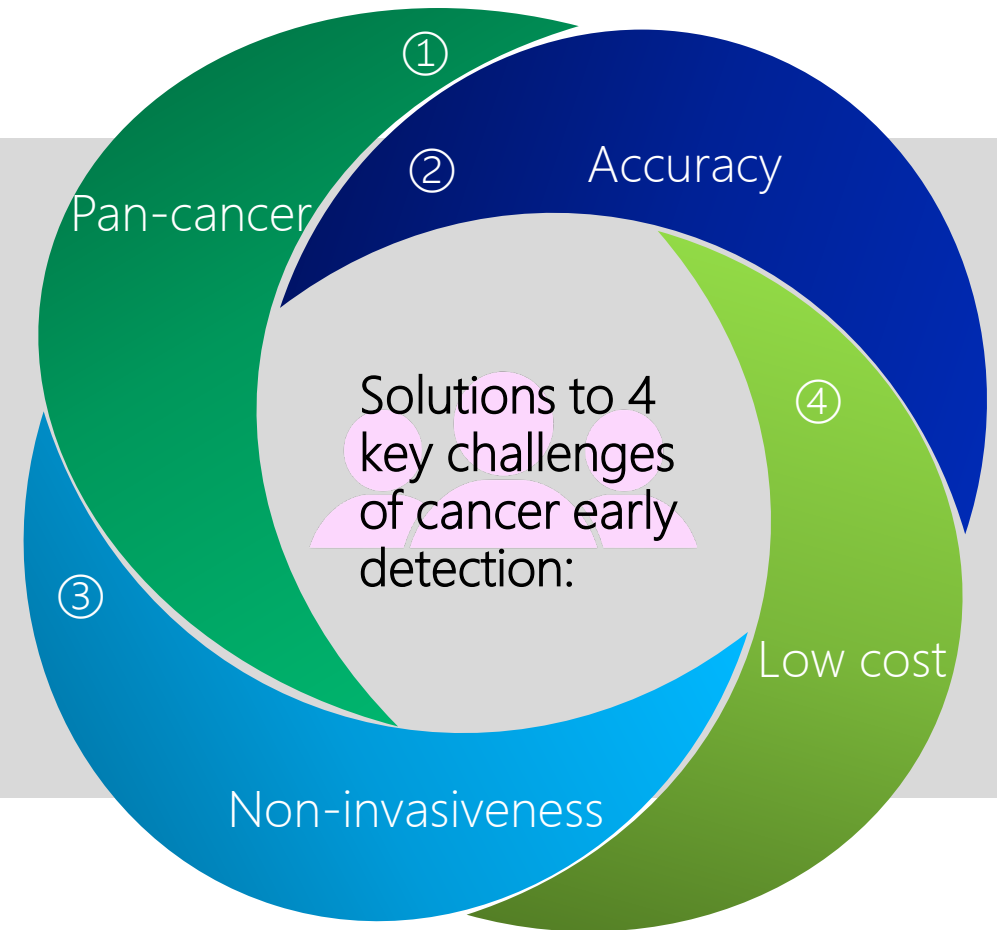
## Mission

- ▶ Detecting cancer early when it is still curable

## Vision

- ▶ Leading novel screening approaches to reduce cancer mortality by 15% in screened population by 2030

## Holy Grail



seek and you will find – Matthew 7:7

# "top-down" approach

## Conventional

4-5 cancer types  
High-risk population  
Low accuracy  
Invasive

## Single cancer type

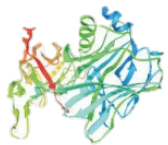
One cancer type  
High-risk population  
Complementary to SOC  
screening  
Noninvasive

## Multi-cancer types

>5 cancer types  
Average-risk population  
Bottom-up approach  
High cost

## Pan-cancer

Pan-cancer  
Average-risk population  
Top-bottom approach  
Cost-effective



### Level 0

CT, B ultrasound,  
Cytology, Protein markers



### Level 1

<5 markers



### Level 2

$10^6$  markers



Marker A+B+C+D

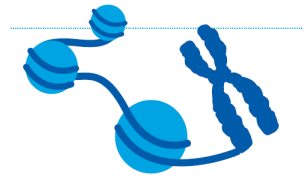
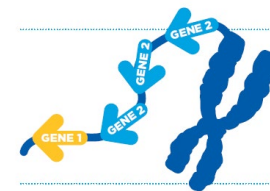
"Bottom-up":  
Add up the markers of individual  
cancer types



SeekInCare®

### Level 3

Whole genome  
3 billion bases




"top-down":  
Utilize common features for all  
cancer types

## Detection of asymptomatic cancers by shallow genome sequencing and 7 protein markers in ~2 million pregnant women (2016-2017)

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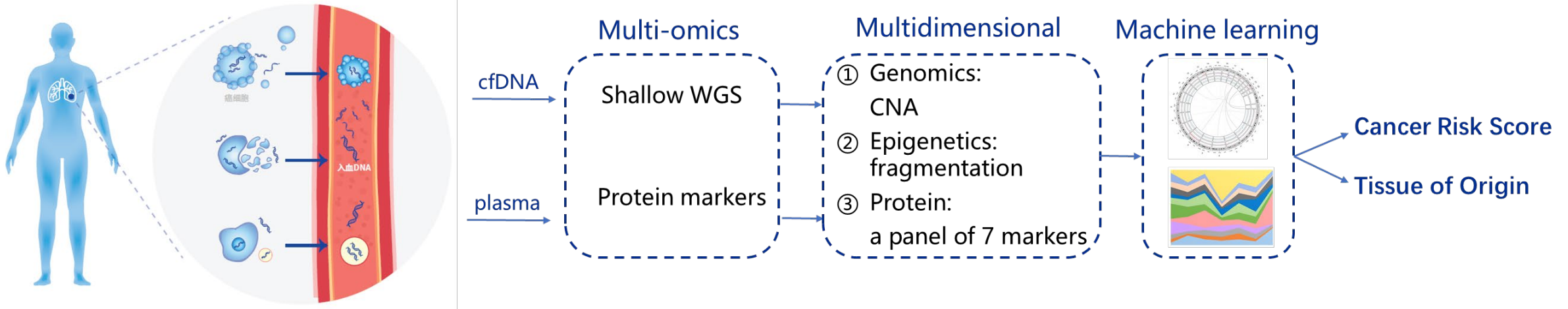
### Identifying occult maternal malignancies from 1.93 million pregnant women undergoing noninvasive prenatal screening tests

[Xing Ji MD](#), [Jia Li PhD](#), ... [Mao Mao MD, PhD](#)  [+ Show authors](#)

[Genetics in Medicine](#) 21, [2293–2302](#) (2019) | [Cite this article](#)

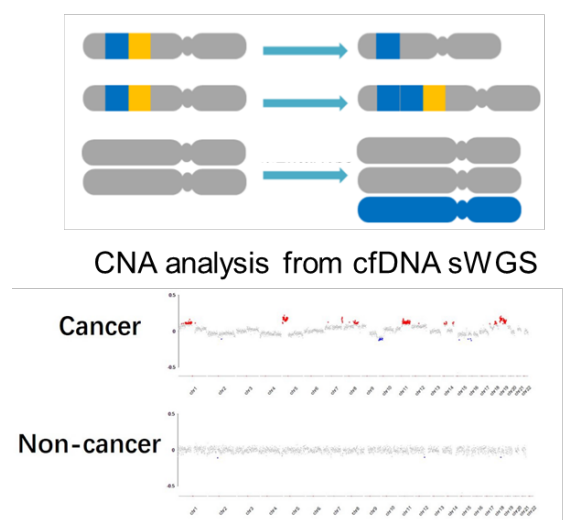
seek and you will find – Matthew 7:7

# Technical edge

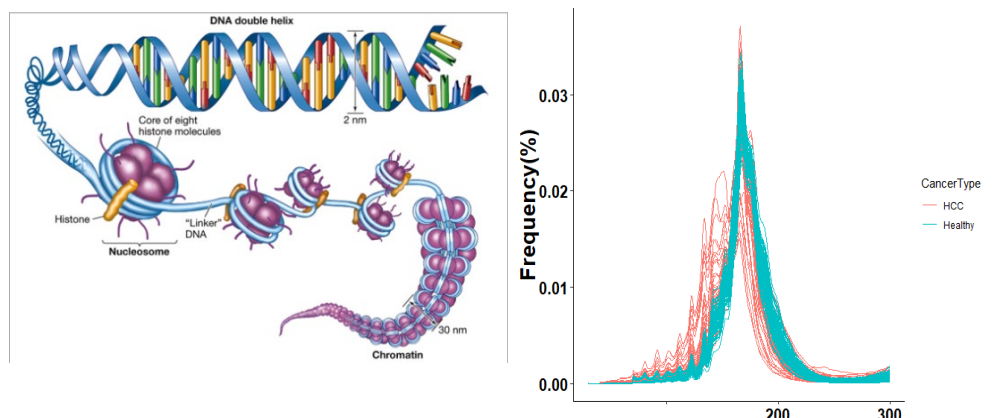


We capture the cancer genomic landscape via a panoramic view by shallow WGS. Thus cancer hallmarks such as CNA and fragment size in conjunction with protein biomarkers can be utilized to refine the MCRC model.

## ① Copy number aberration (CNA)

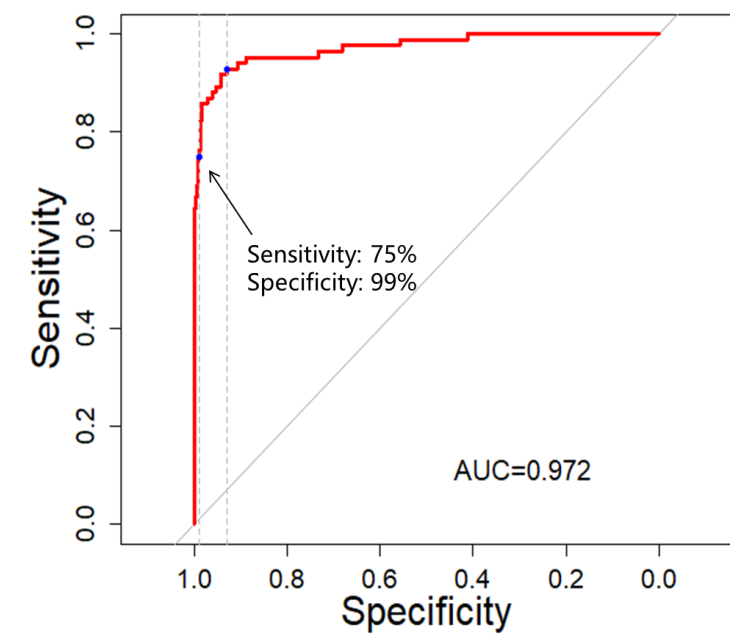


## ② Fragmentation



## ③ Protein markers

AFP, CEA, CA199, CA125, CA153, CA211, CA724





思康宁®  
SeekInCare®

**the first-in-class blood-based pan-cancer early detection test**

CE Marking



A panoramic view of cancer genomics landscape + protein markers



Big data + AI

**A blood-based test generating CRS and locating TOO**

comprehensive

Can detect molecular cancer signals in all cancer types

accurate

More accurate than other NGS-based costly tests  
TOO can provide feasible clinical workup suggestions

cost-effective

Technical edge ensures controllable cost

non-invasive

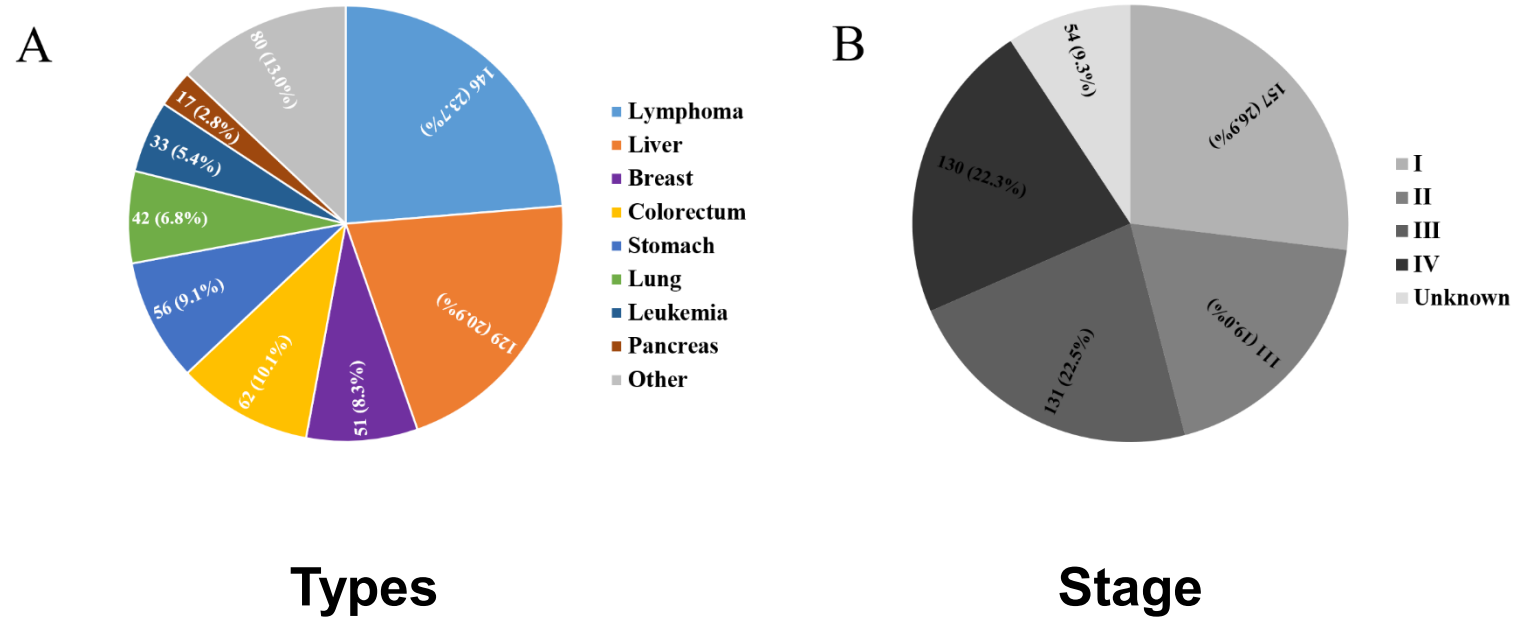
All the test needs is only 10 mL peripheral blood from one blooddraw

"An inborn pan-cancer early detection test"

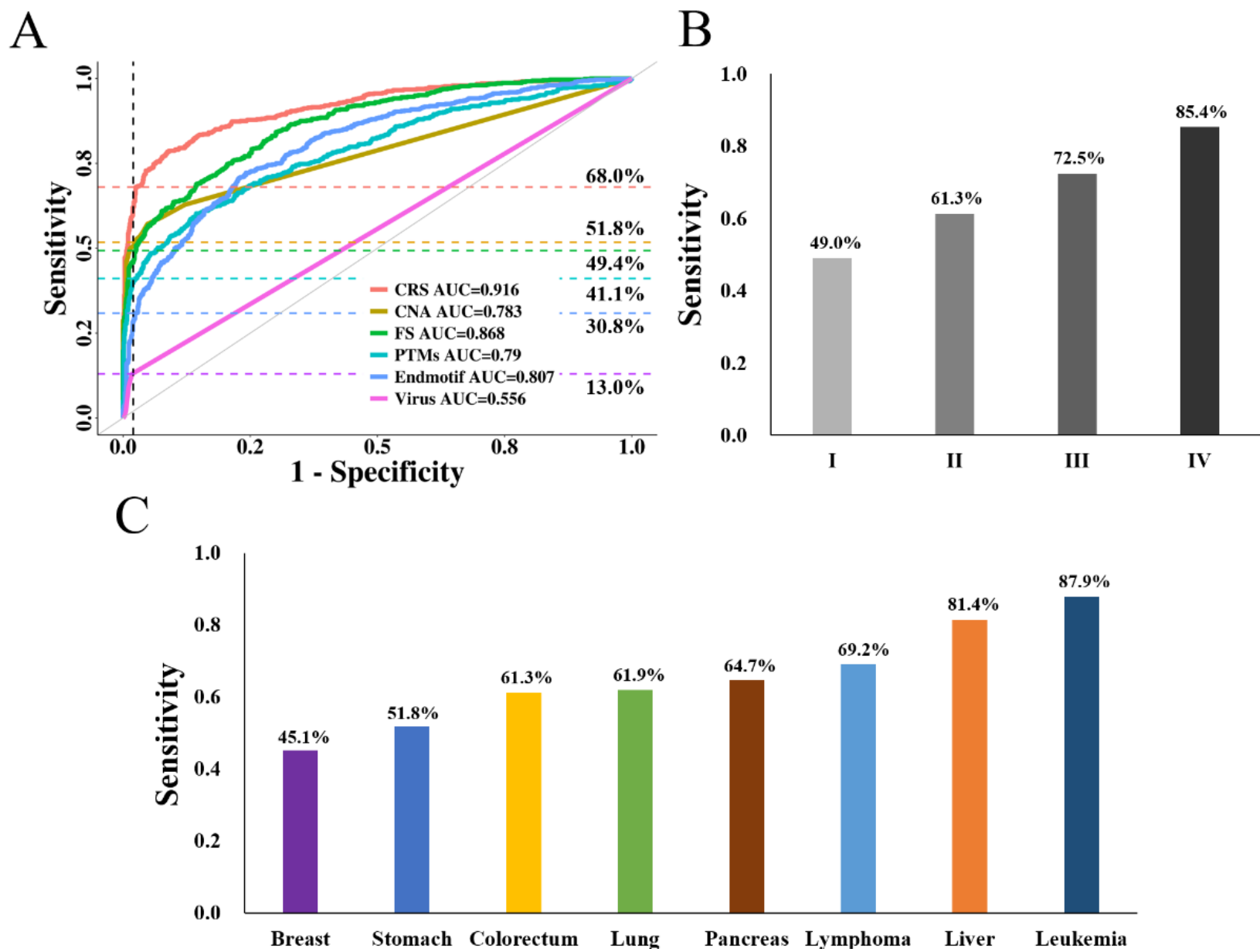
Methodology paper: Meng Z, et al. Non-invasive detection of hepatocellular carcinoma with circulating tumor DNA features and AFP. J Mol Diagn. 2021 Sep;23(9):1174-1184.

# The 2<sup>nd</sup> validation study

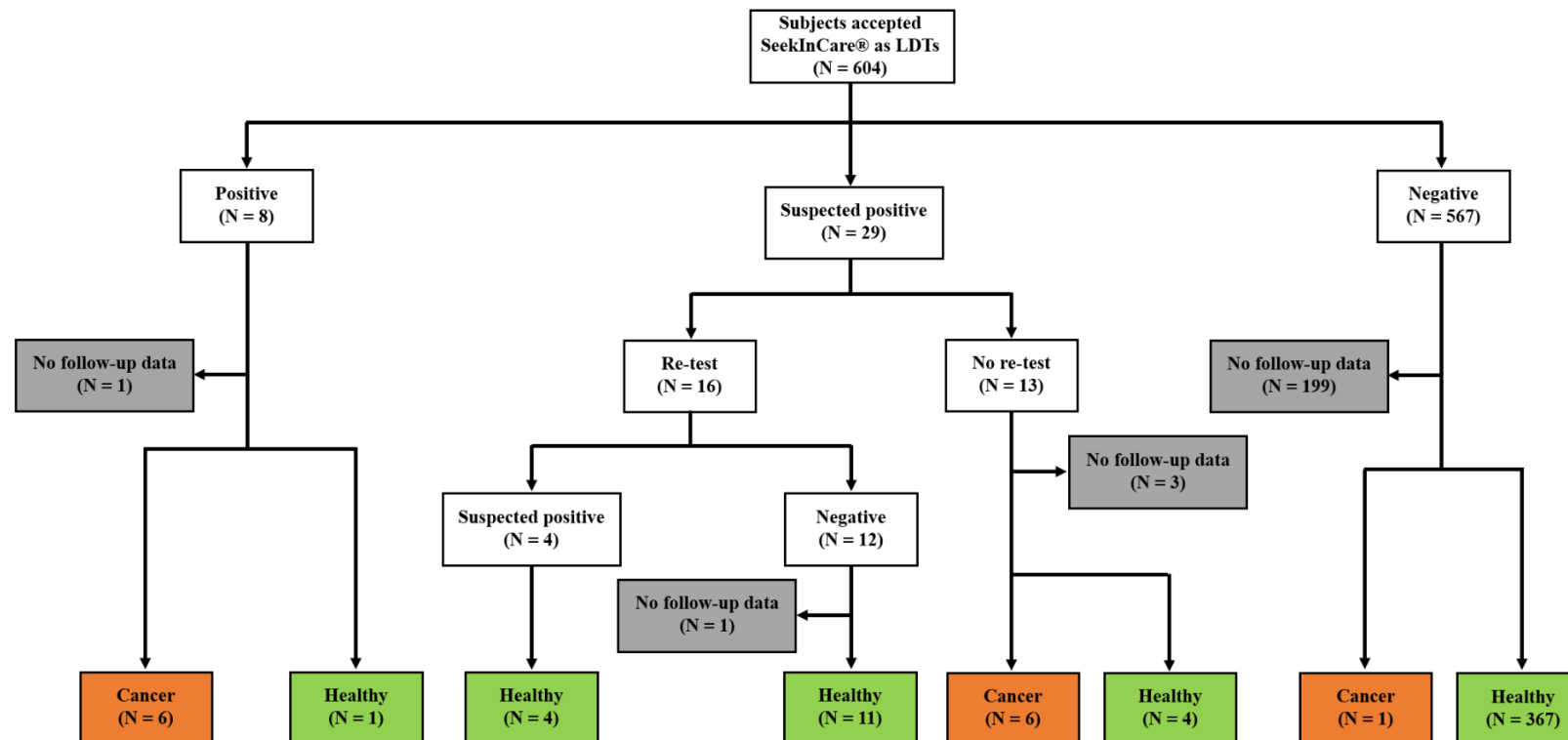
616 stage I-IV cancer patients that cover eight common cancers and the other types and 898 healthy subjects.



# Performance of SeekInCare® in the 2<sup>nd</sup> validation study



# Performance of SeekInCare® in the real world



92.3% sensitivity, 97.8% specificity, 48% PPV and 99.8% NPV

# Low-cost cancer detection: OncoSeek®

思康永®  
OncoSeek®

CE Marking



OncoSeek®, a combination of 7 clinically validated plasma tumor markers (AFP, CA125, CA15-3, CA19-9, CA72-4, CEA, and CYFRA21-1), using a new multivariate OncoSeek algorithm combined with big data and AI, performs early detection for 12 common cancers by calculating the probability of cancer (POC) and tracing the cancer tissue of origin (TOO).

Bladder

Breast

Cervix

Colorectum

Esophagus

Gallbladder

Liver

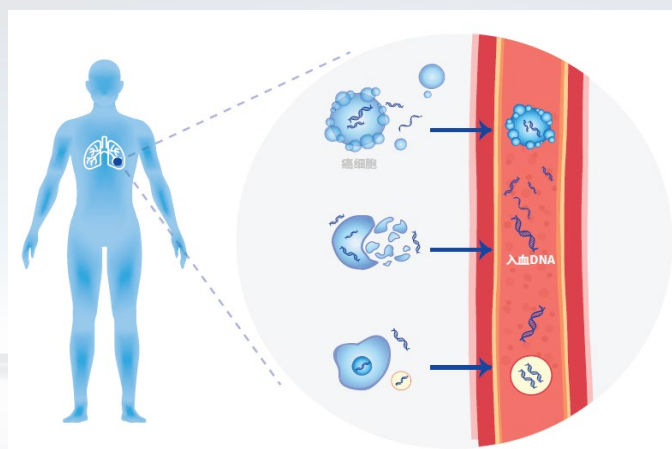
Lung

Ovary

Pancreas

Stomach

Testis



plasma

**Bladder:** CEA

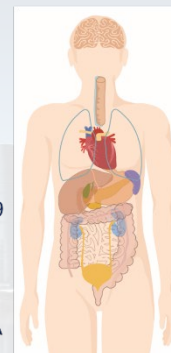
**Breast:** CA15-3, CEA

**Cervix:** CEA

**Colorectum:** CEA, CA19-9

**Esophagus:** CEA

**Gallbladder:** CA19-9, CEA



**Liver:** AFP, CA19-9, CEA

**Lung:** CEA, CYFRA21-1

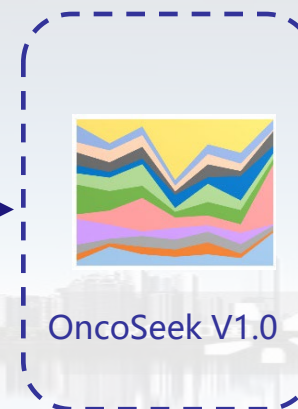
**Ovary:** CA125

**Stomach:** CEA, CA19-9, CA72-4

**Pancreas:** CEA, CA19-9

**Testis:** AFP

Protein marker panel

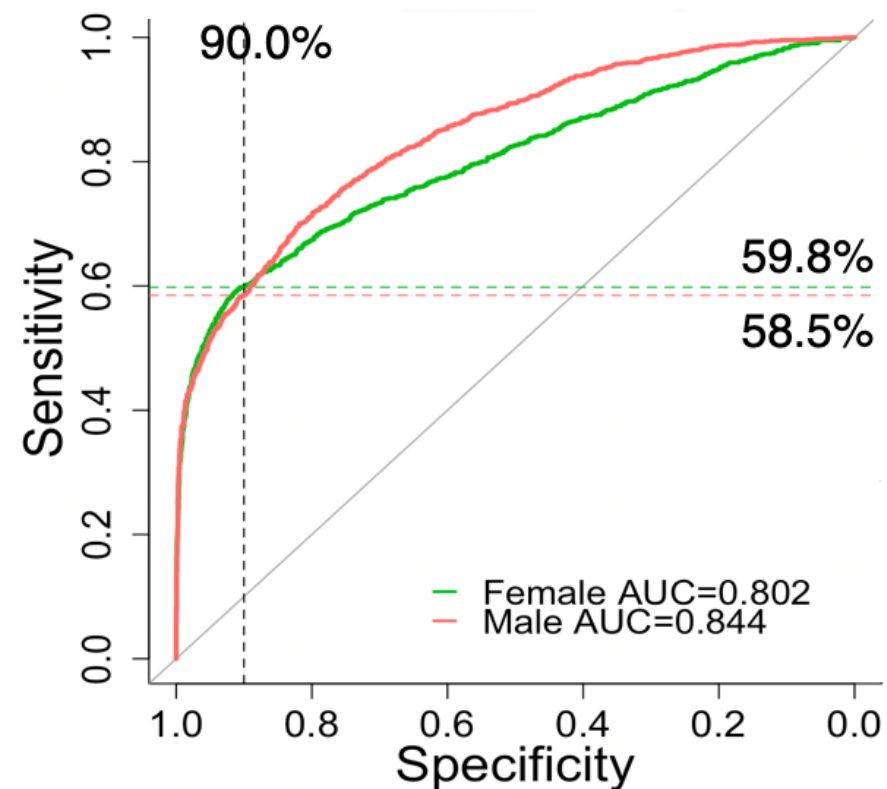


Machine learning

POC

TOO

# Low-cost cancer detection: OncoSeek®



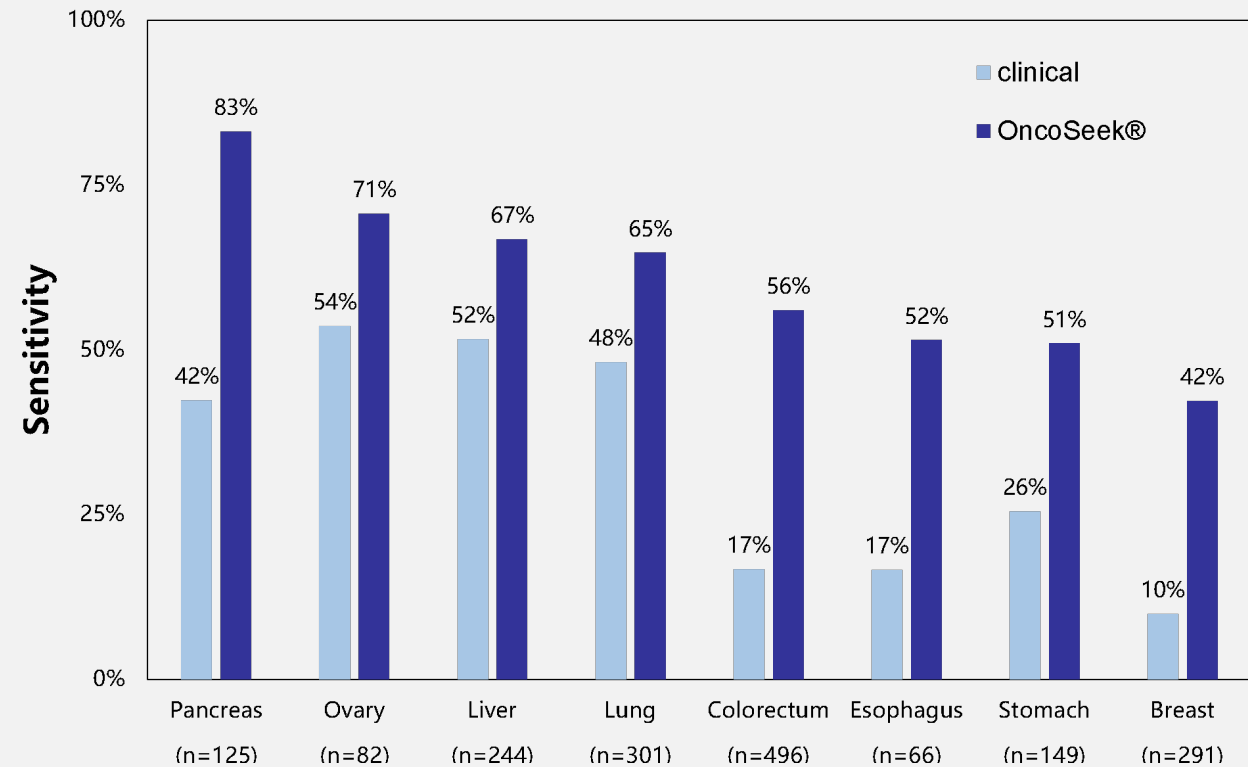
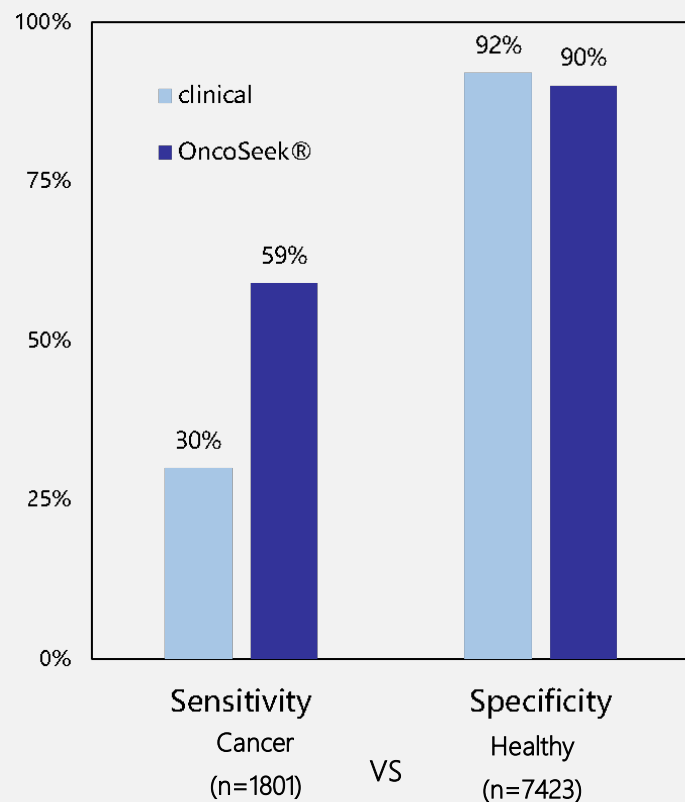
	Female		Male	
	Cancer	Healthy	Cancer	Healthy
Predicted cancer	518	341	547	400
Predicted noncancer	348	3075	388	3607
Sensitivity % (95% CI)	59.8 (56.5—63.1)		58.5 (55.3—61.7)	
Specificity % (95% CI)	90.0 (89.0—91.0)		90.0 (89.0—90.9)	
PPV % (95% CI)	60.3 (56.9—63.6)		57.8 (54.5—60.9)	
NPV % (95% CI)	89.8 (88.8—90.8)		90.3 (89.3—91.2)	



- ✓ Using US FDA/CE/NMPA approved reagents and instrument (Roche cobas e411) to improve the accuracy of cancer detection
- ✓ Based on the analysis of ~10,000 blood samples, the sensitivity of OncoSeek was ~60% at 90% specificity, and the AUC was greater than 0.80

# Low-cost cancer detection: OncoSeek®

Based on the analysis of nearly 10,000 samples, with specificity at ~90%, the sensitivity of OncoSeek® (59%) for cancer detection is two times more than that of using current clinical method.



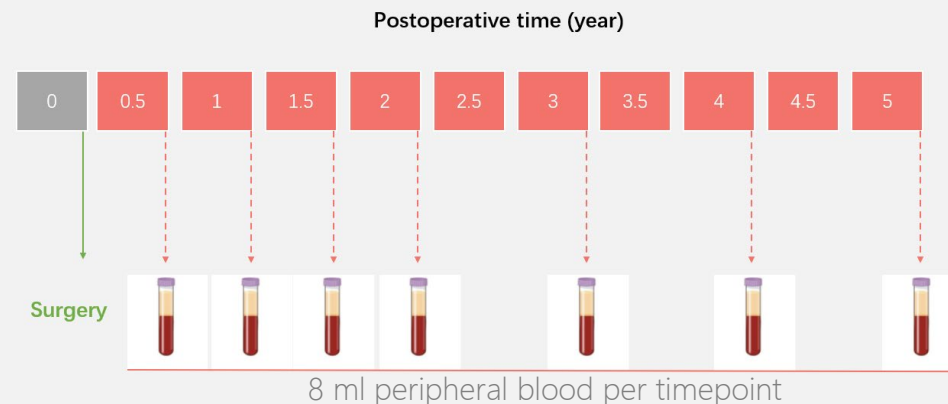
# Recurrence monitoring - SeekInCure™

思康安®  
SeekInCure™

CE Marking



SeekInCure™ is applied to monitoring cancer recurrence and risk evaluation for post-op cancer patients. By monitoring the variation of cancer genome in cell-free DNA (cfDNA) of post-op cancer patients continually, combined with plasma protein markers, equipping big data and artificial intelligence, it generates cancer recurrence index (CRI) .



## Characteristics

Precise

Personalized  
surveillance

Non-invasive

Blood-based

Broad-spectrum

Primary/Metastatic  
cancerous lesions

Multi-omics

sWGS+  
protein markers

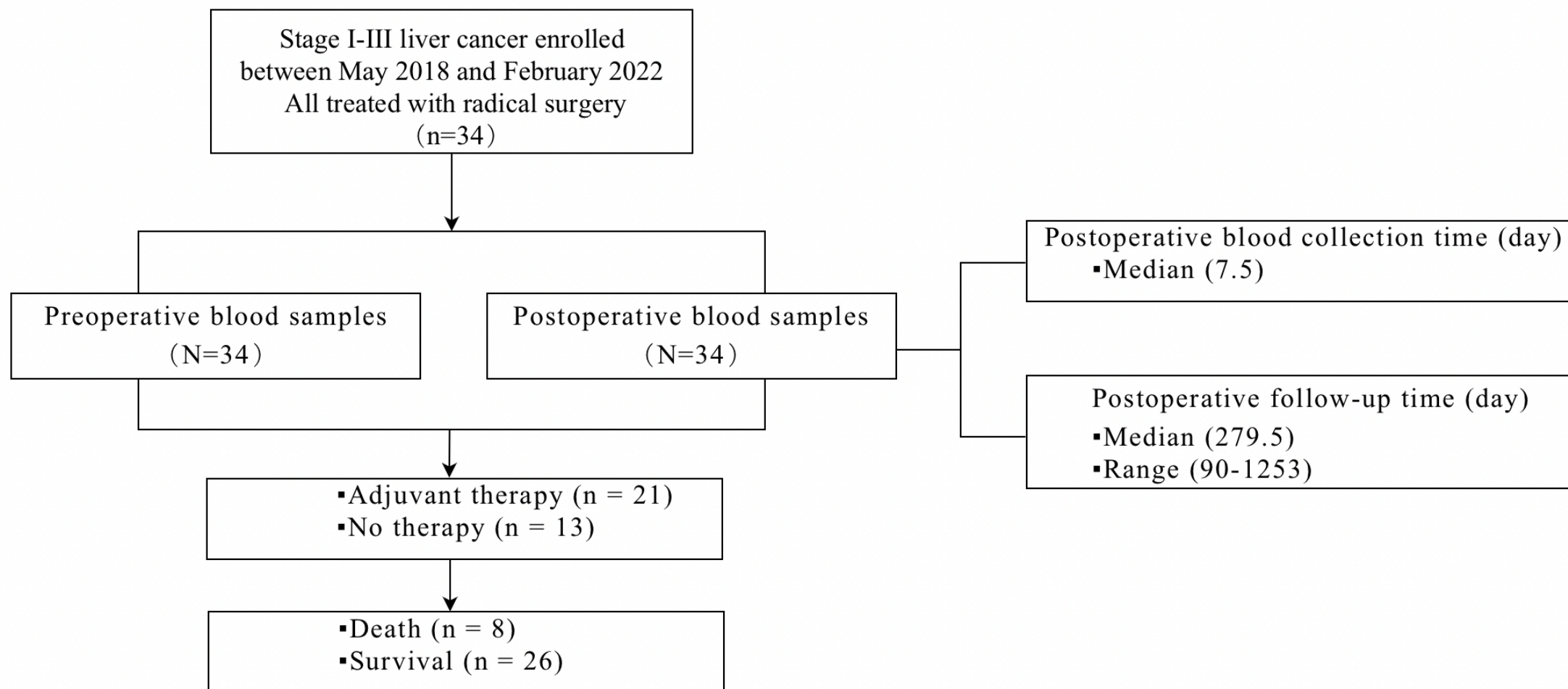
Cost-effective

Affordable

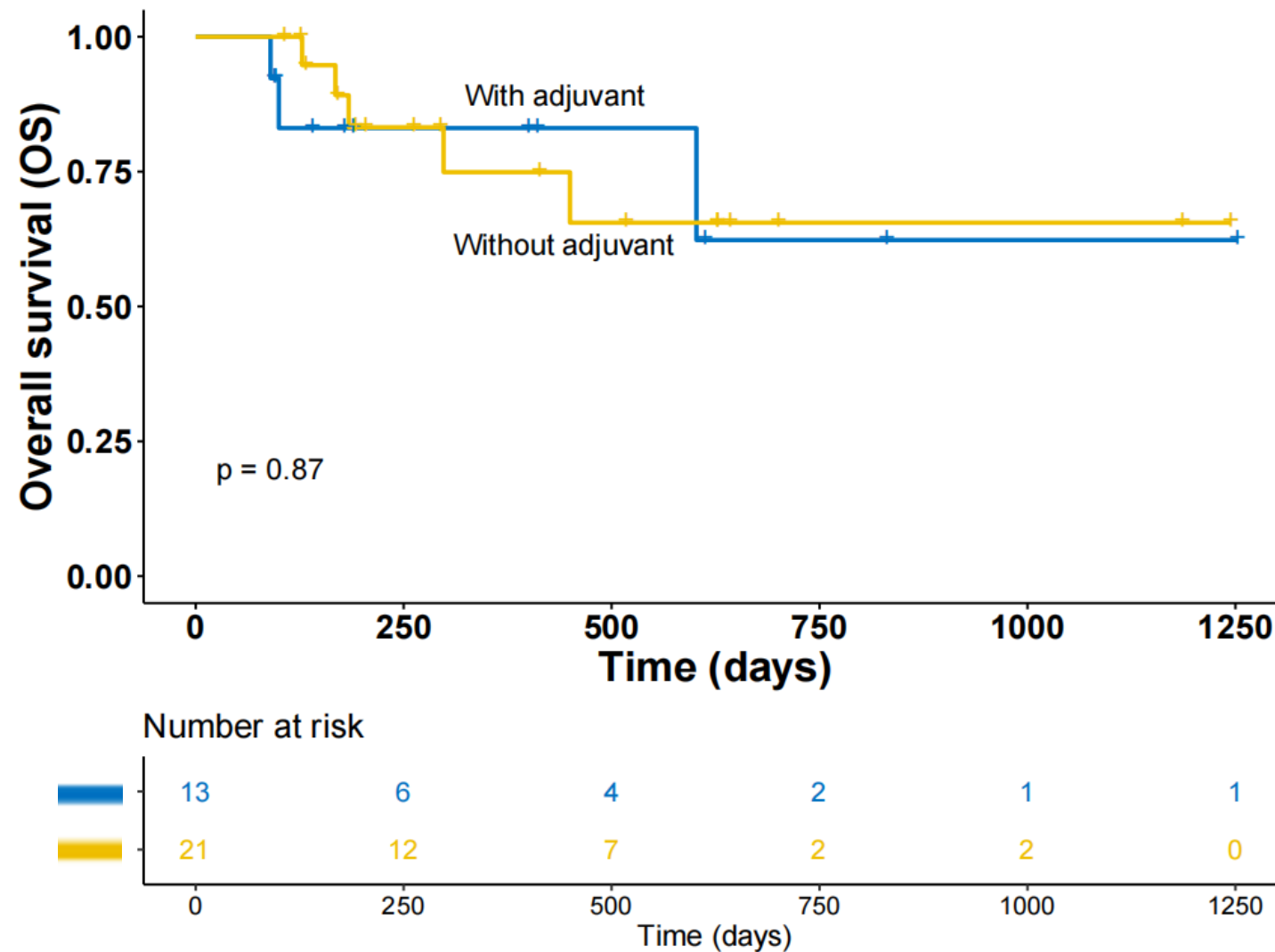


# SeekInCure™ – HCC prospective MRD study

Prospective HCC MRD study (2018.05 ~ 2022.02): enrolled HCC radical surgery patients.  
8ml blood sample collected at preoperative (treatment-naïve) and postoperative timepoints.

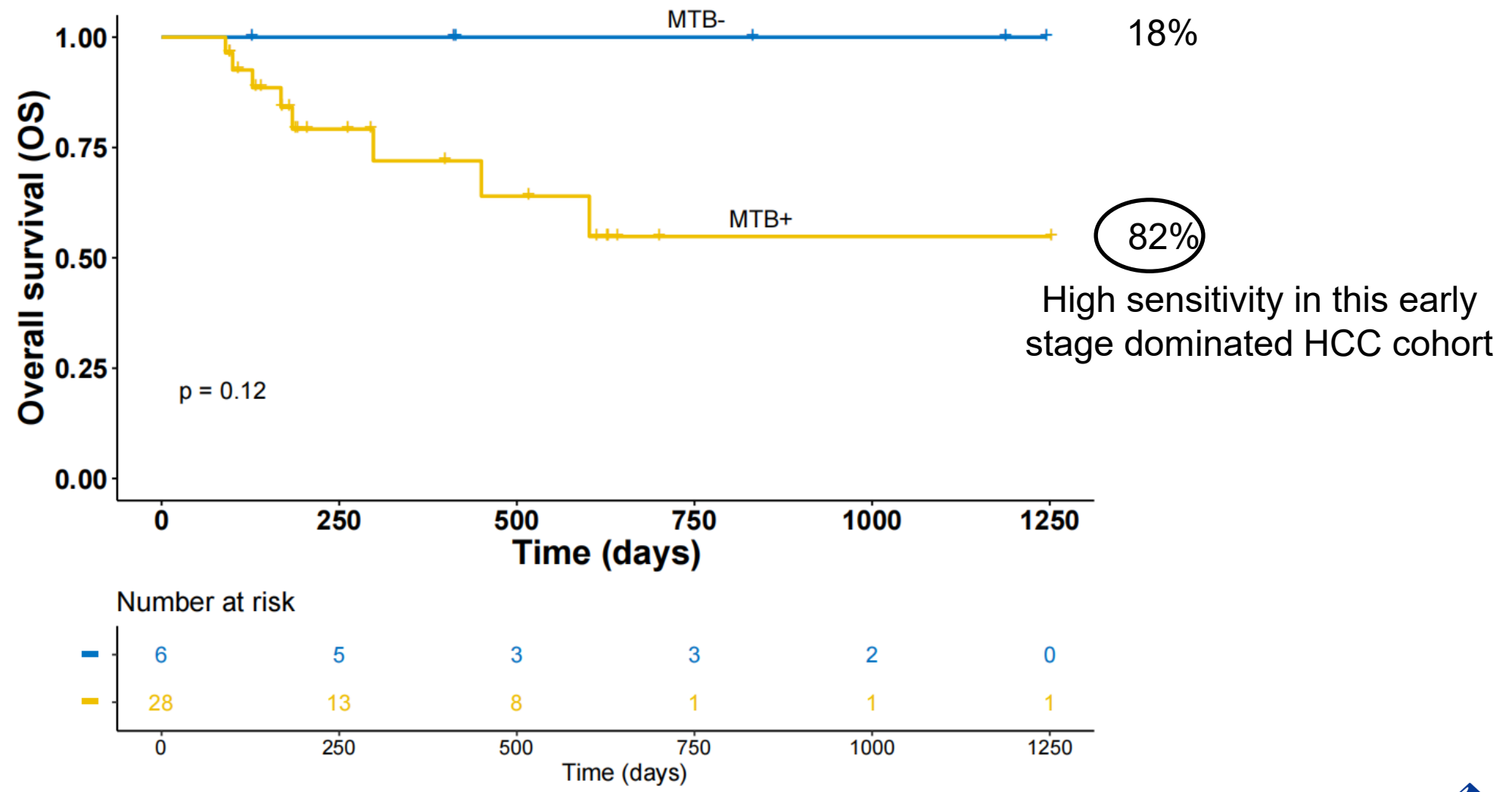


# Adjuvant has no influence on survival



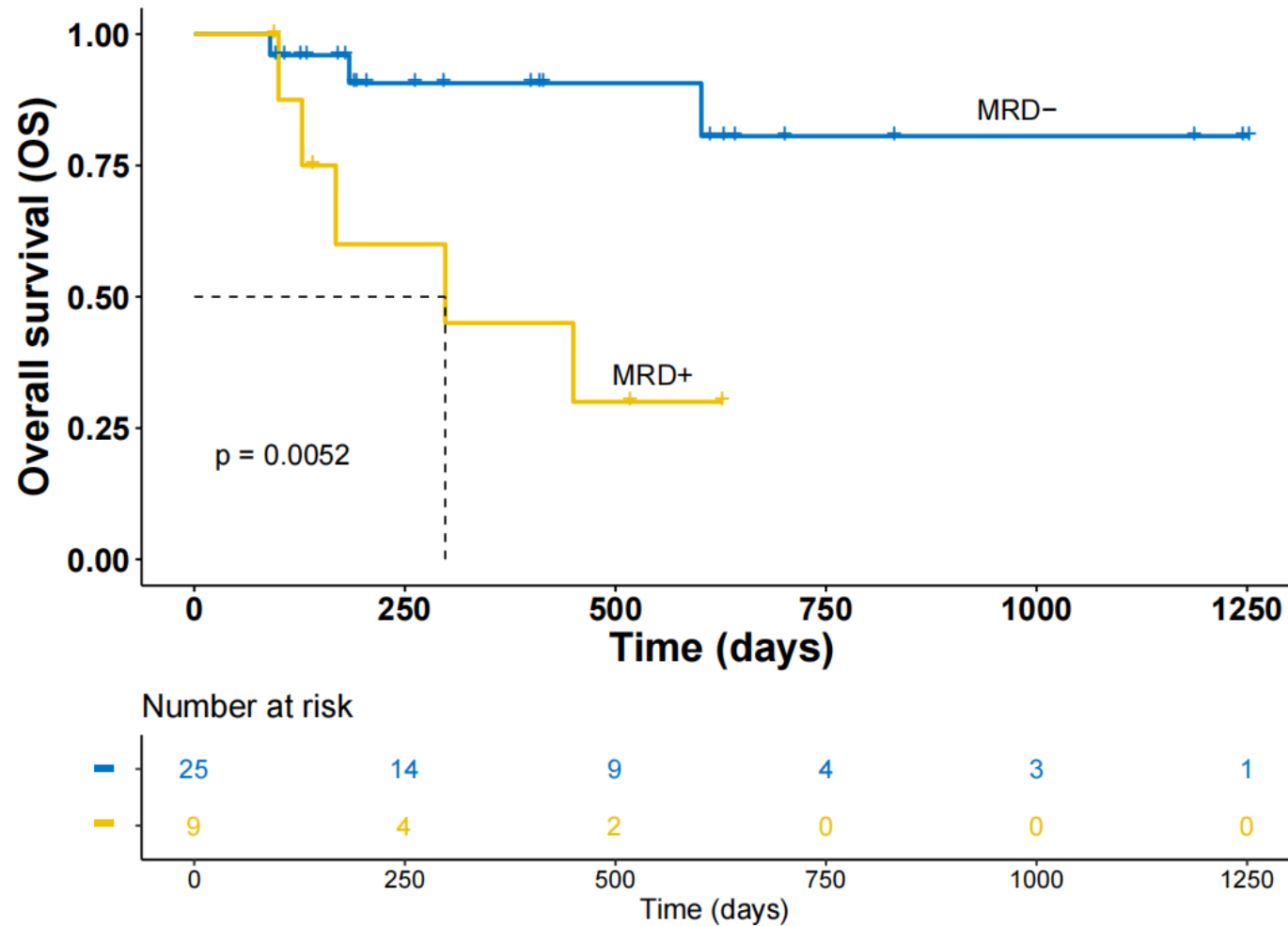
To be treated or not to be?

## SeekInCure MTB detection for prognosis



MTB: molecular tumor burden. MTB+/-: molecular signal of tumor is detectable/undetectable

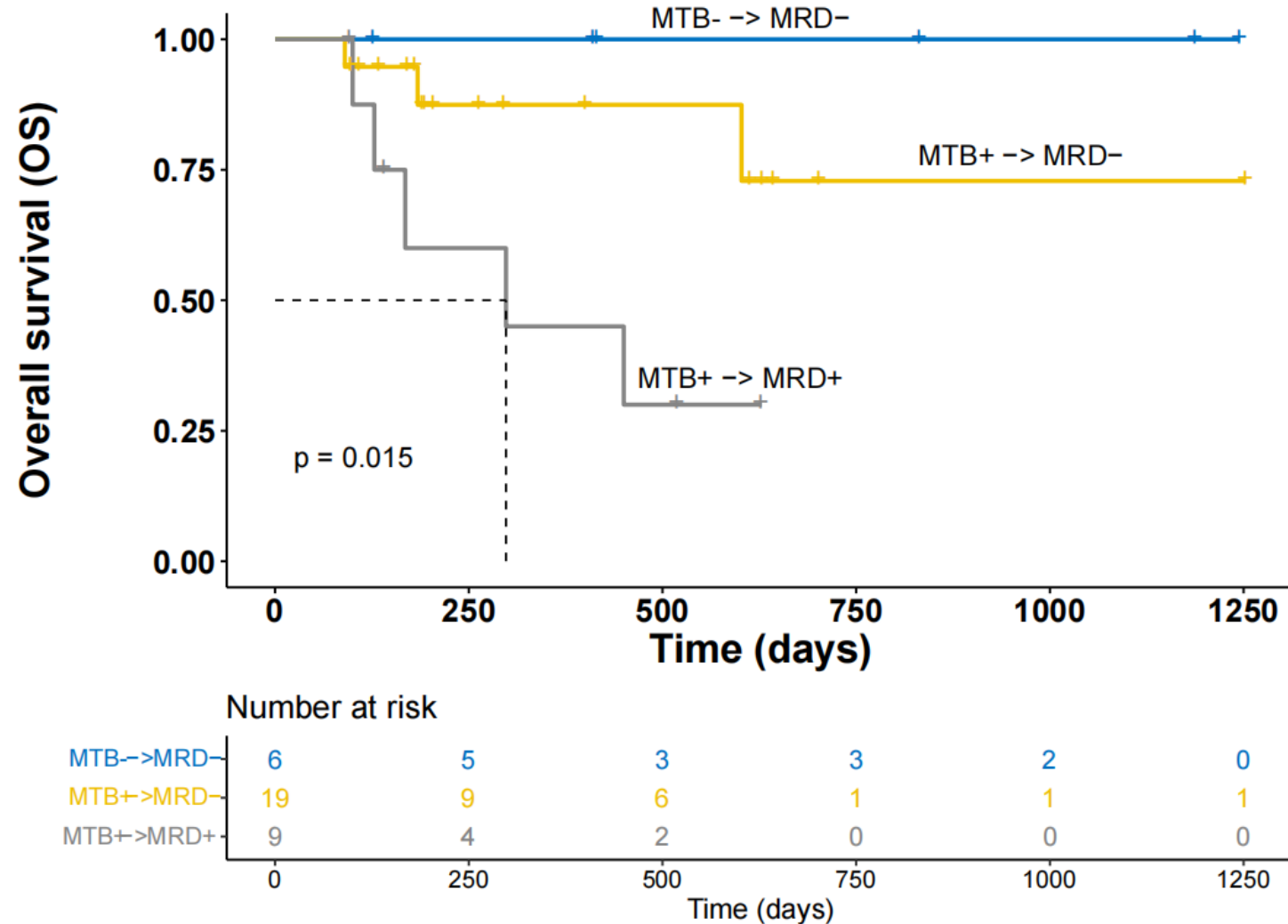
## SeekInCure MRD detection for prognosis



After surgery, patients with MRD+ have shorter survival time.  
Median overall survival time is less than 1 year.

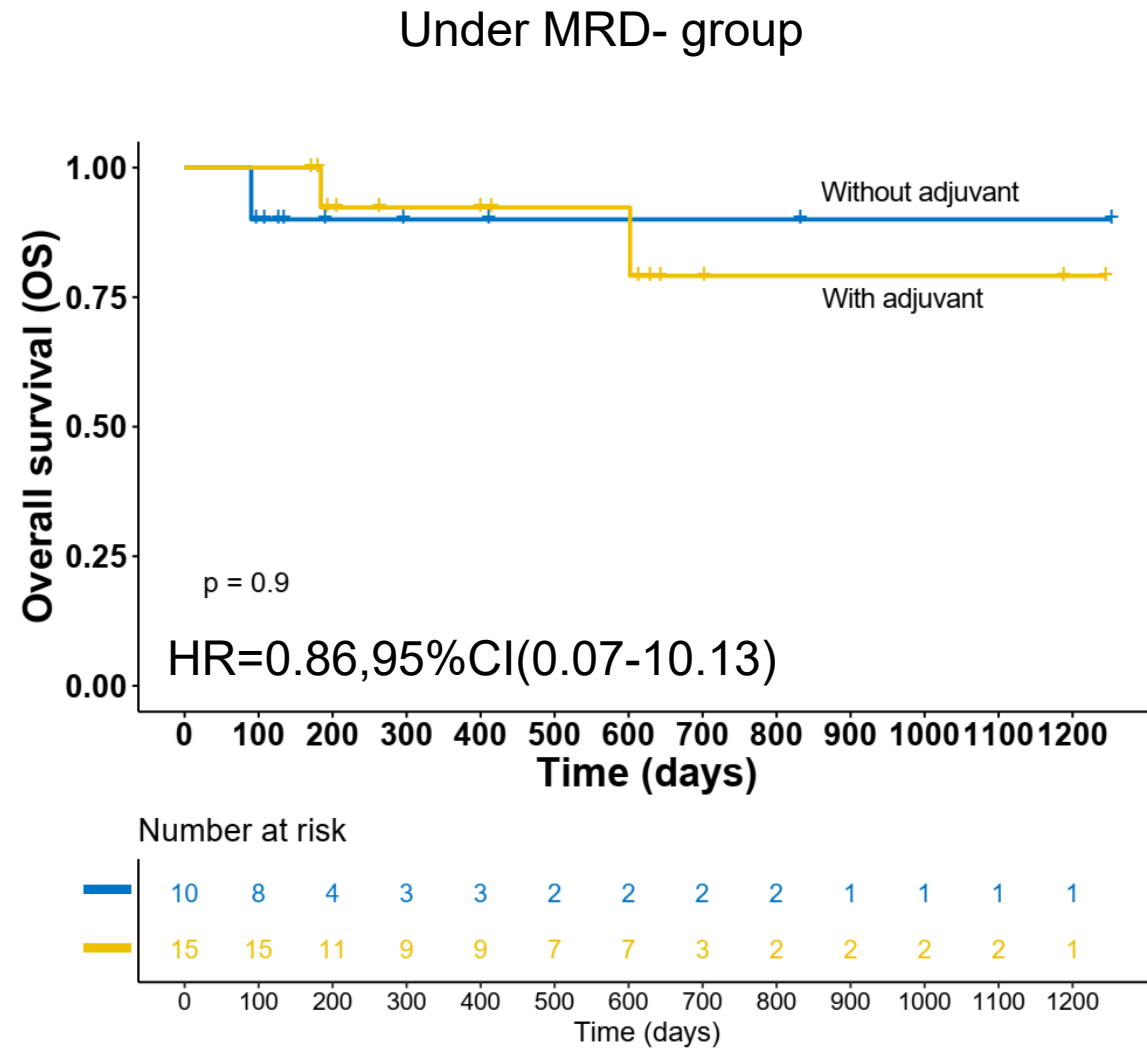
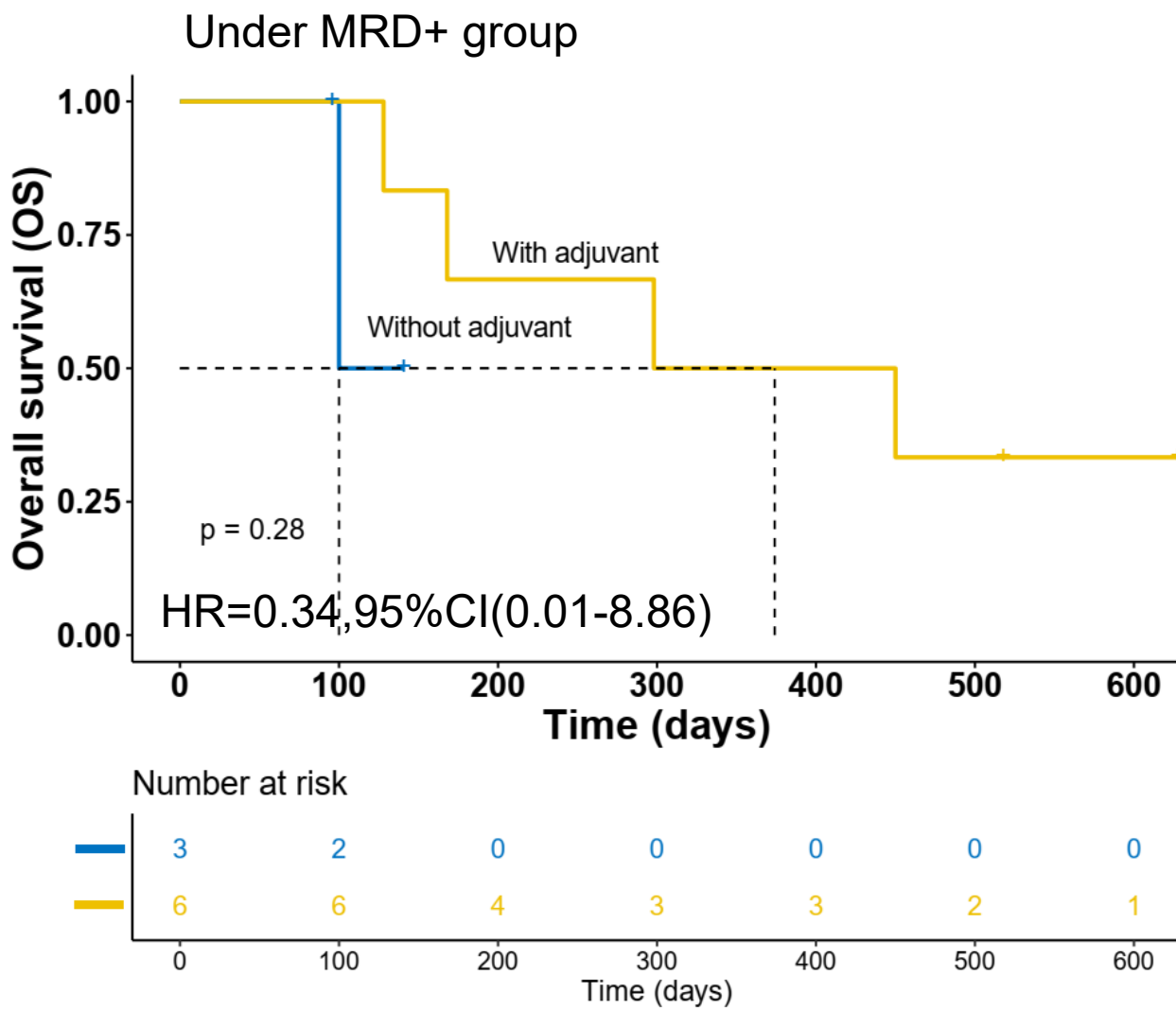
# Pre-/post- operative changes and outcome

Patients who are MTB- and MRD- have very good OS



31% MTB+ patients remained positive (i.e. MRD+) after surgery

# SeekInCure™ – with/without adjuvant



Among HCC MRD- group, adjuvant has no influence on survival

- SeekInCure is a cost-effective pan cancer MRD test that does not require cancer tissue analysis.
- Both preoperative MTB and postoperative MRD values are prognostic.
- Double negative (MTB- and MRD-) patients have a very favorable outcome (100% OS).
- Double positive (MTB+ and MRD+) patients have the worst outcome (median overall survival time is less than 1 year).
- Adjuvant has no influence on survival, especially in MRD- patients.

# Response monitoring - SeekInClarity™

思康明®

SeekInClarity™

CE Marking



✓

Pan-cancer

✓

Pan-indication

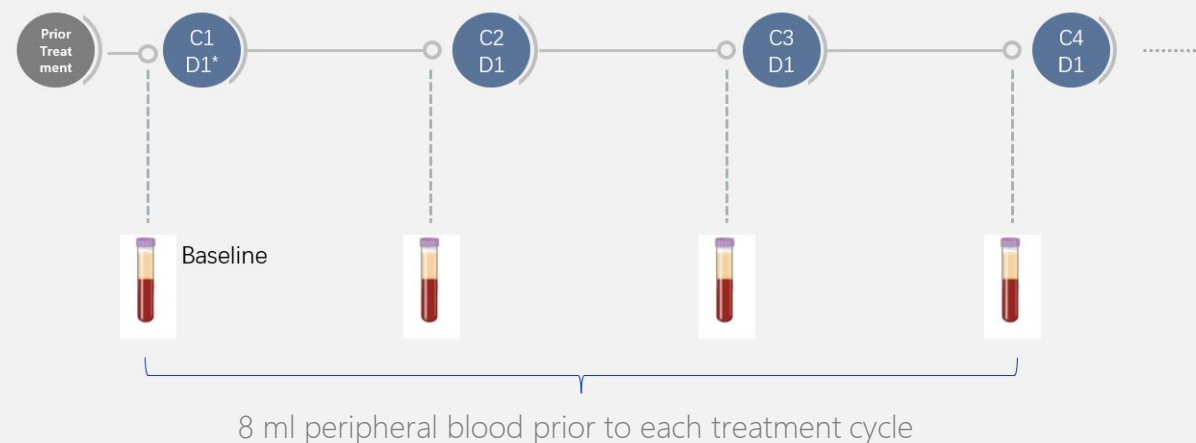
✓

Multi-omics

✓

Prompt monitoring

SeekInClarity™ is the first-in-class blood-based pan-cancer pan-indication treatment response monitoring test, which uses a cutting edge multivariable molecular tumor burden (MTB) algorithm. Molecular response index (MRI) model maps the panoramic cancer genome by shallow whole genome sequencing (sWGS) data, evaluates the copy number aberration (CNA), fragment size (FS) and 7 plasma protein markers, to predict tumor burden and therapeutic efficacy of the late-stage cancer patients during treatment including chemotherapy, target therapy and immunotherapy or combination therapies.

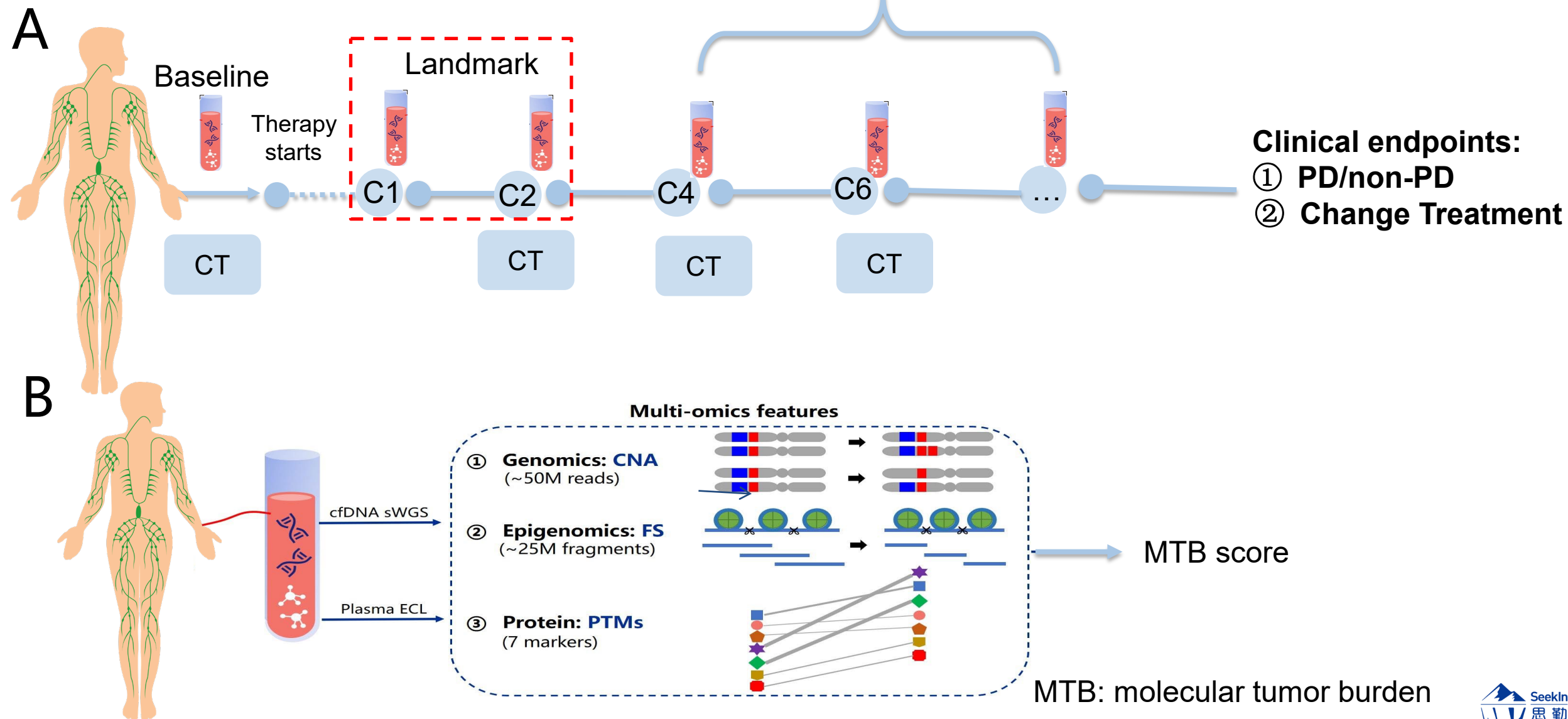


\*: Cycle 1 Day 1, the 1<sup>st</sup> day of each treatment cycle



# SeekInClarity™ for lymphoma treatment response

Longitudinal: analyzed every 2 cycles within 2 years



# Clinical summary of lymphoma patients

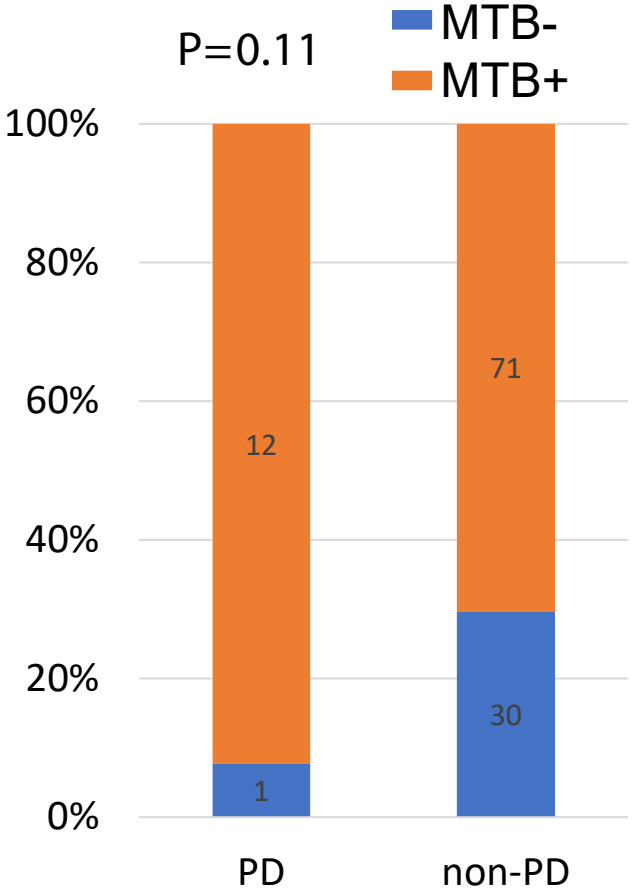
Characteristics	No. (%)
No. of patients	154
Median age, years (range)	55 (13, 81)
Sex	
Female	82 (53.2)
Male	72 (46.8)
Subtype	
DLBCL	64 (41.6)
NK/TCL	20 (13)
HL	14 (9.0)
FL	11 (7.1)
AITL	8 (5.2)
Others	37 (11.7)
Stage	
I	27 (17.5)
II	30 (19.5)
III	17 (11.0)
IV	73 (47.4)
NaN	7 (4.5)

Regimens	
R-CHOP	27 (21.8)
R-CDOP	17 (13.7)
PD-1	9 (7.3)
BR	9 (7.3)
DDGP	9 (7.3)
AVD	6 (4.8)
CHOP	5 (4.0)
CDOP	4 (3.2)
Other regimens	32 (25.8)
NaN	6 (4.8)

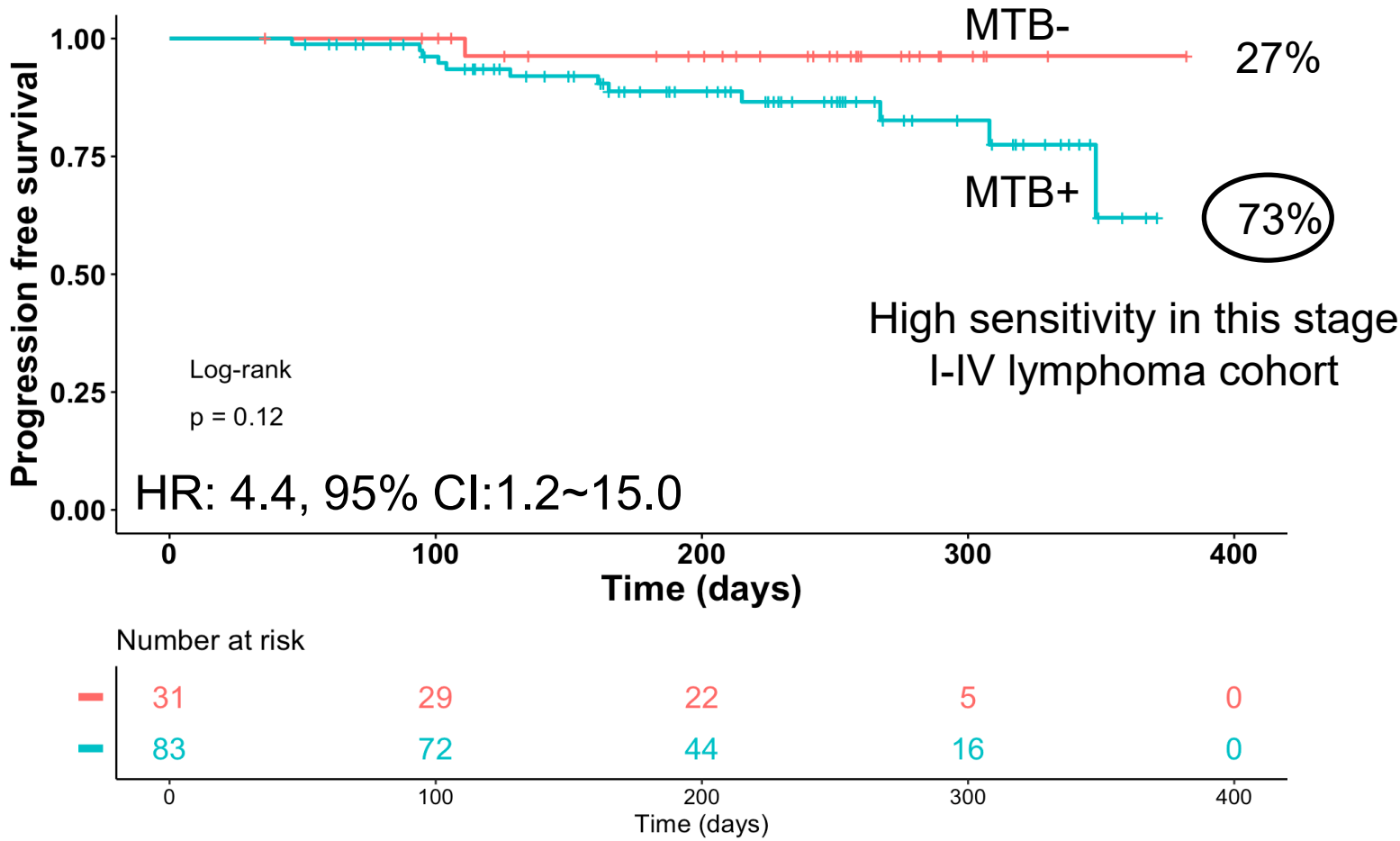
114 patients were included in the interim analysis

# SeekInClarity™ analysis at baseline

Correlation between MTB status at baseline and clinical outcome (PFS)



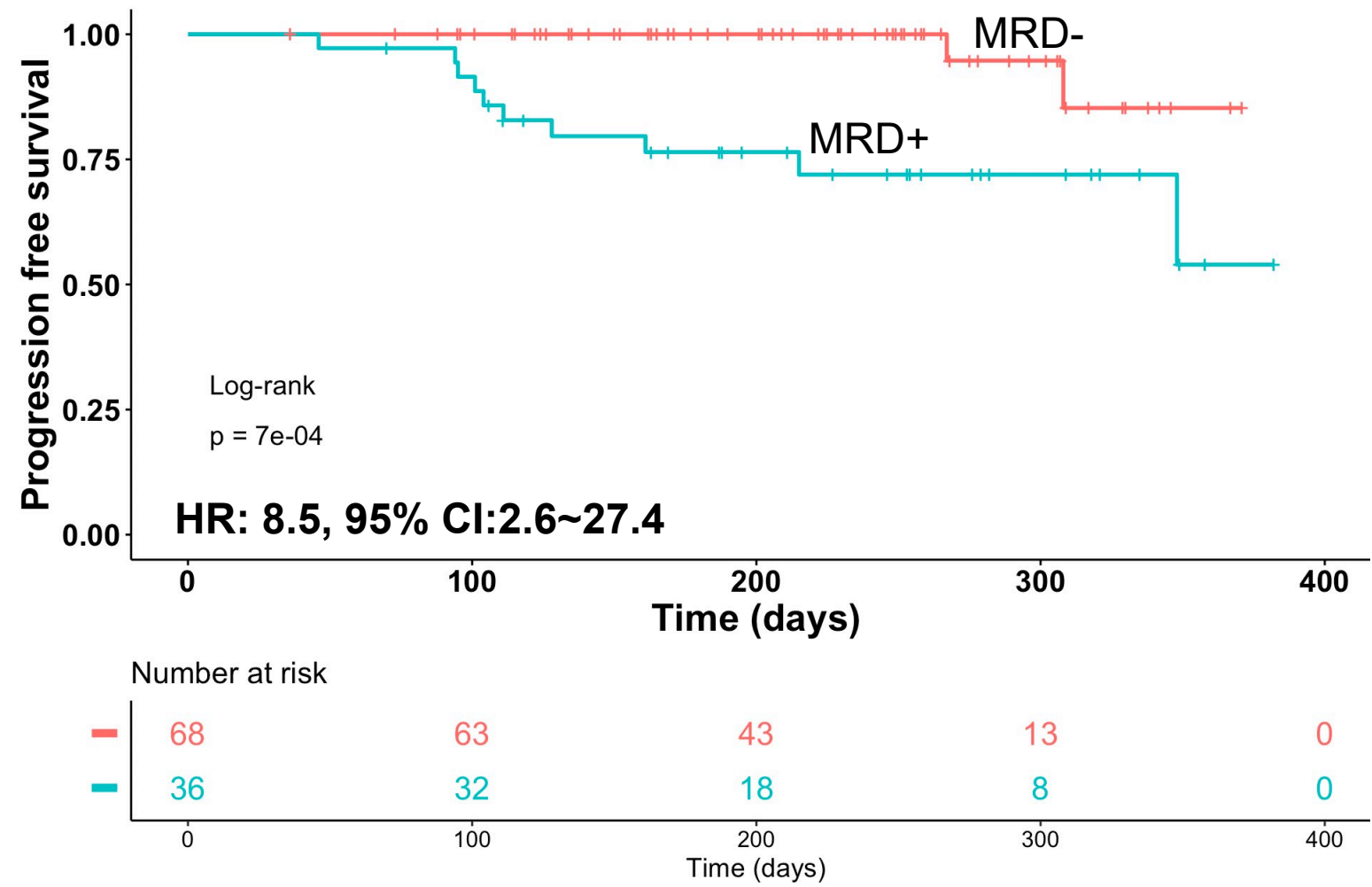
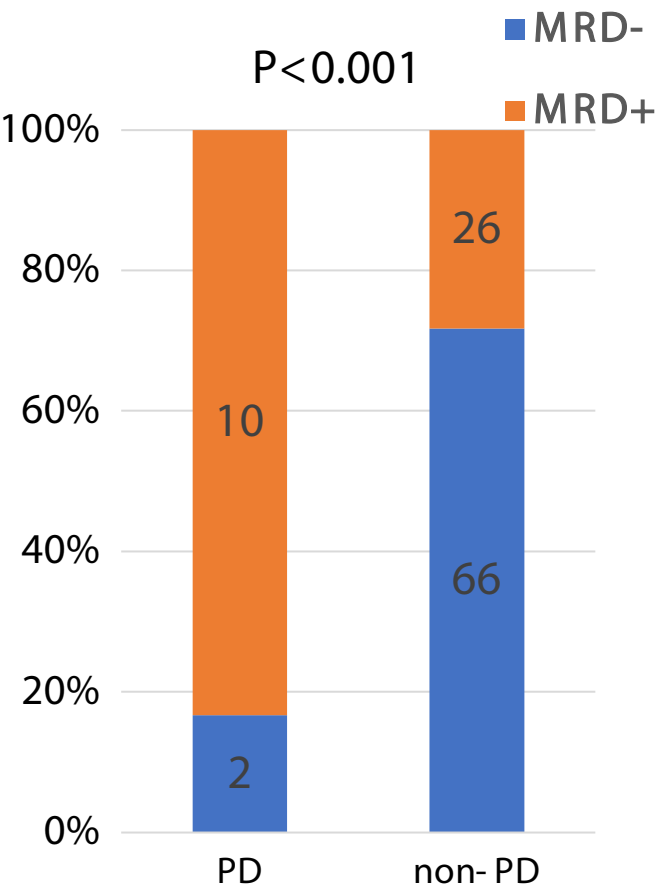
PD: Progressive disease, i.e. non-responders  
Non-PD: responders, including SD, PR, CR



MTB: molecular tumor burden  
MTB+/-: molecular signal of tumor is detectable/undetectable

# SeekInClarity™ analysis at landmark

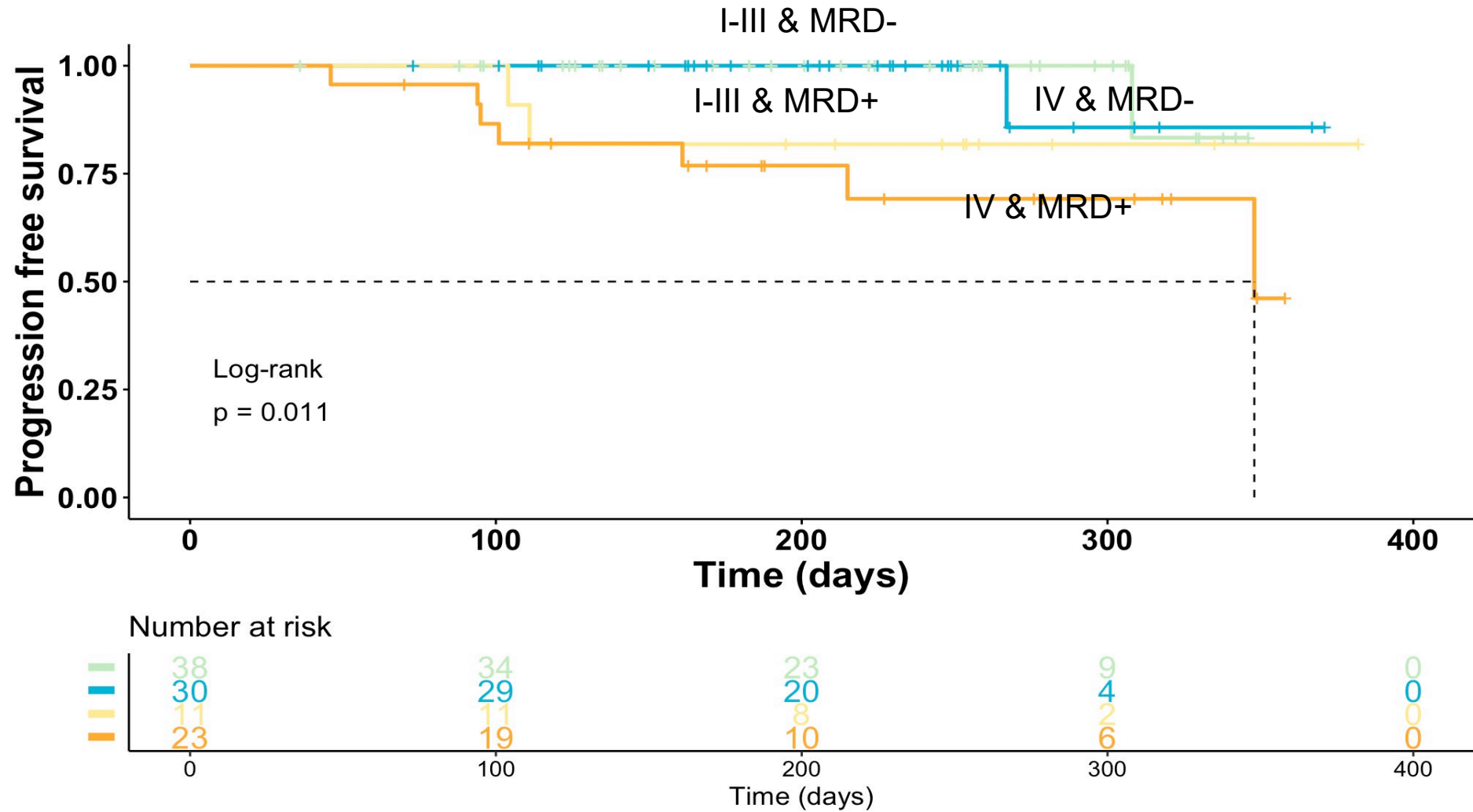
MRD status at landmark is prognostic on treatment effectiveness



\*Samples collected after 2 cycles were defined as landmark, except 4 out of 104 patients used 1 cycle samples instead.

# SeekInClarity™ vs tumor stage

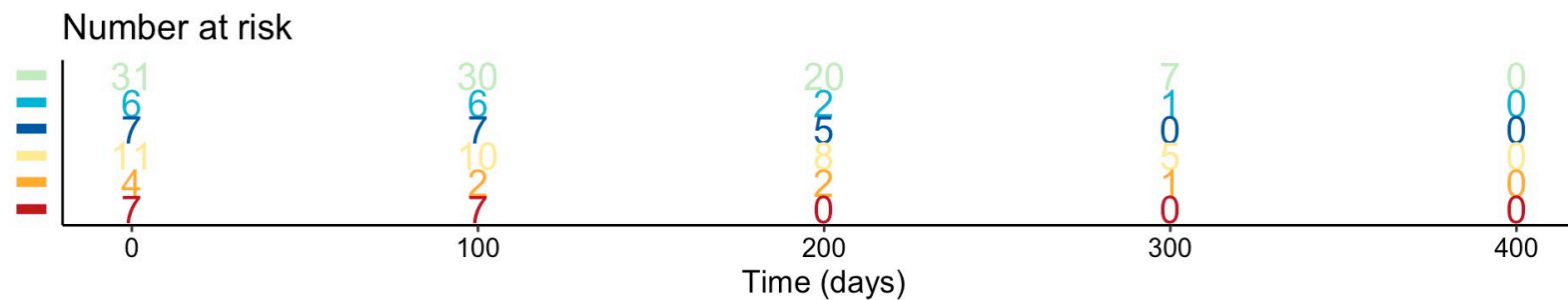
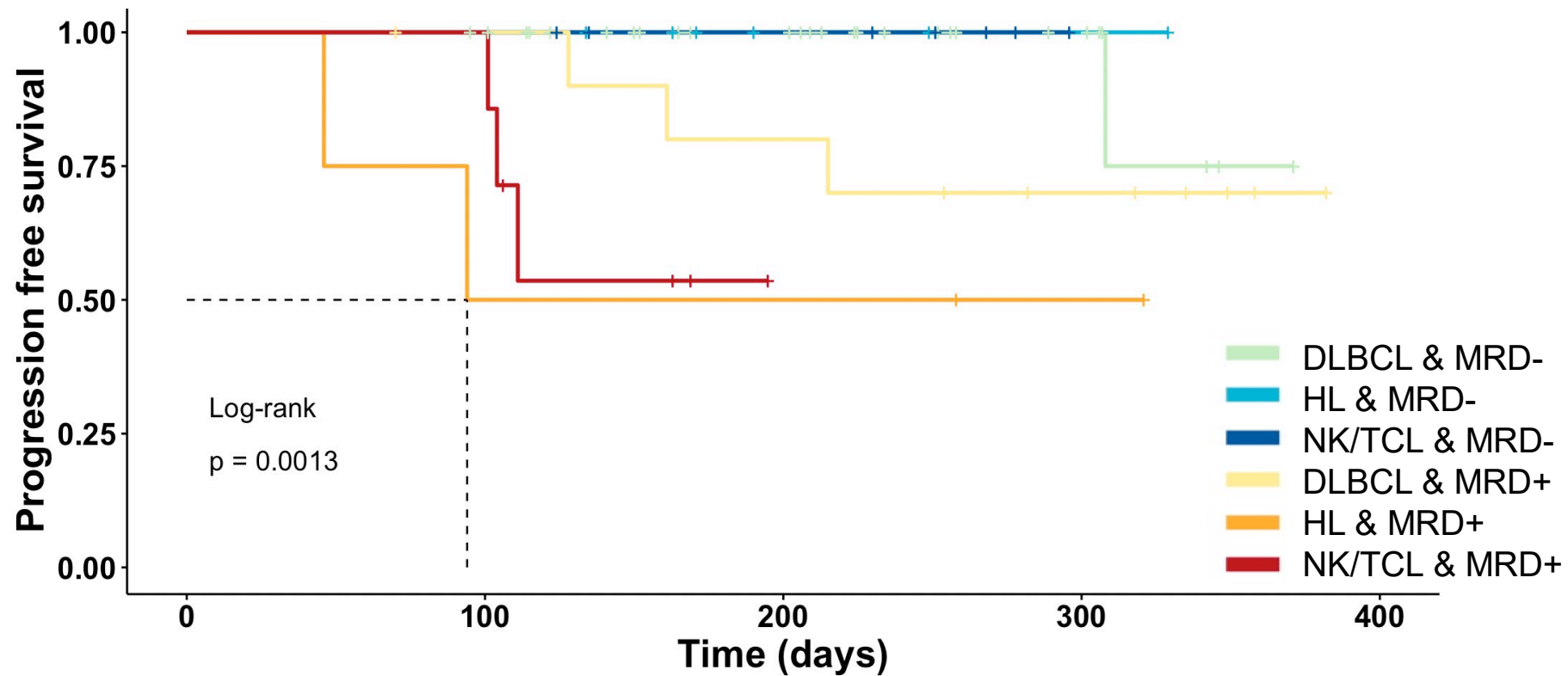
MRD status is more prognostic than tumor staging



Blue: MRD-  
Orange: MRD+

# SeekInClarity™ vs different subtypes

MRD status is more prognostic than lymphoma subtyping

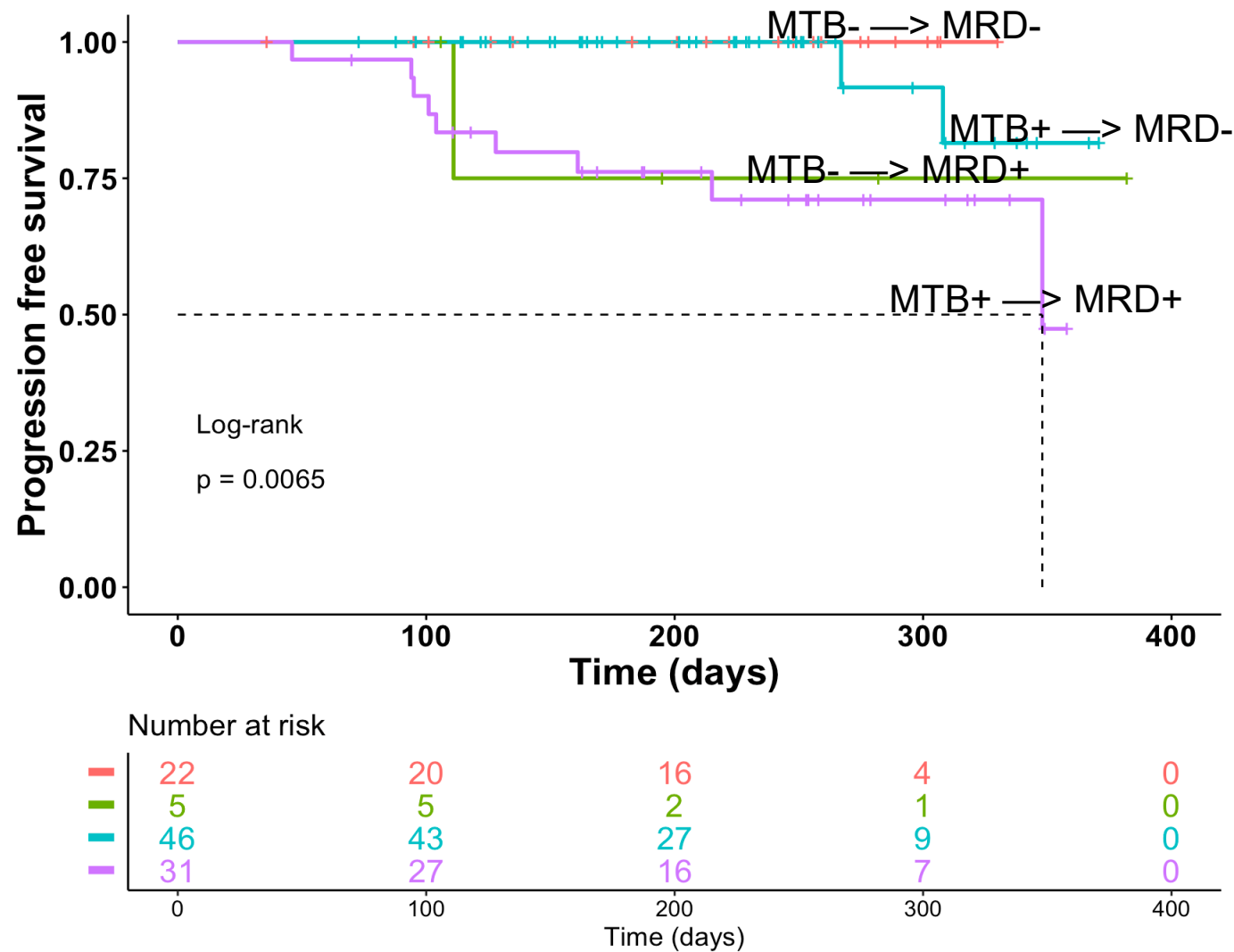


Blue: MRD-  
Orange: MRD+

# MTB status changes before/after treatment

Patients who are MTB negative at both before and after treatment do extremely well

Baseline → landmark

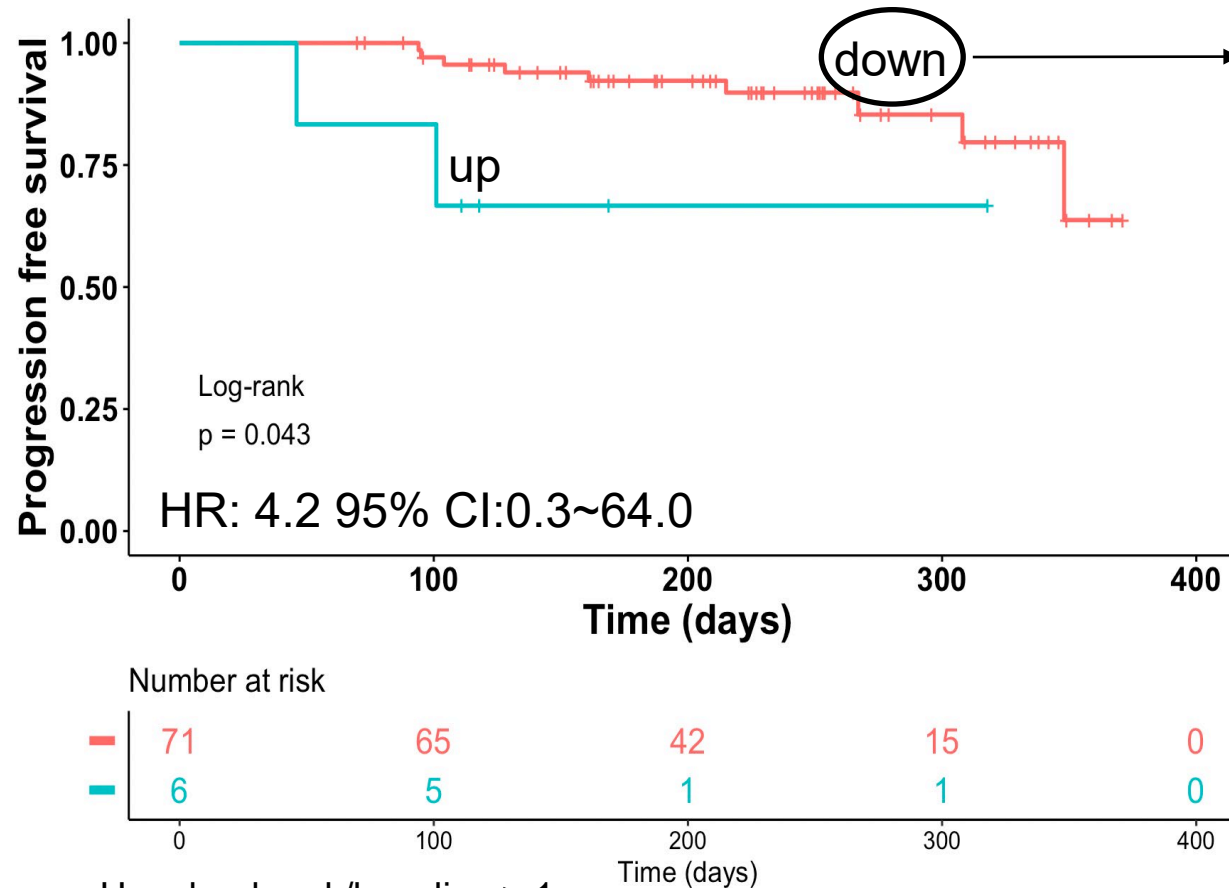


60% TMB+ (baseline) patients can achieve MRD- (ctDNA/PTM clearance) at Landmark

# MTB dynamic changes before and after treatment

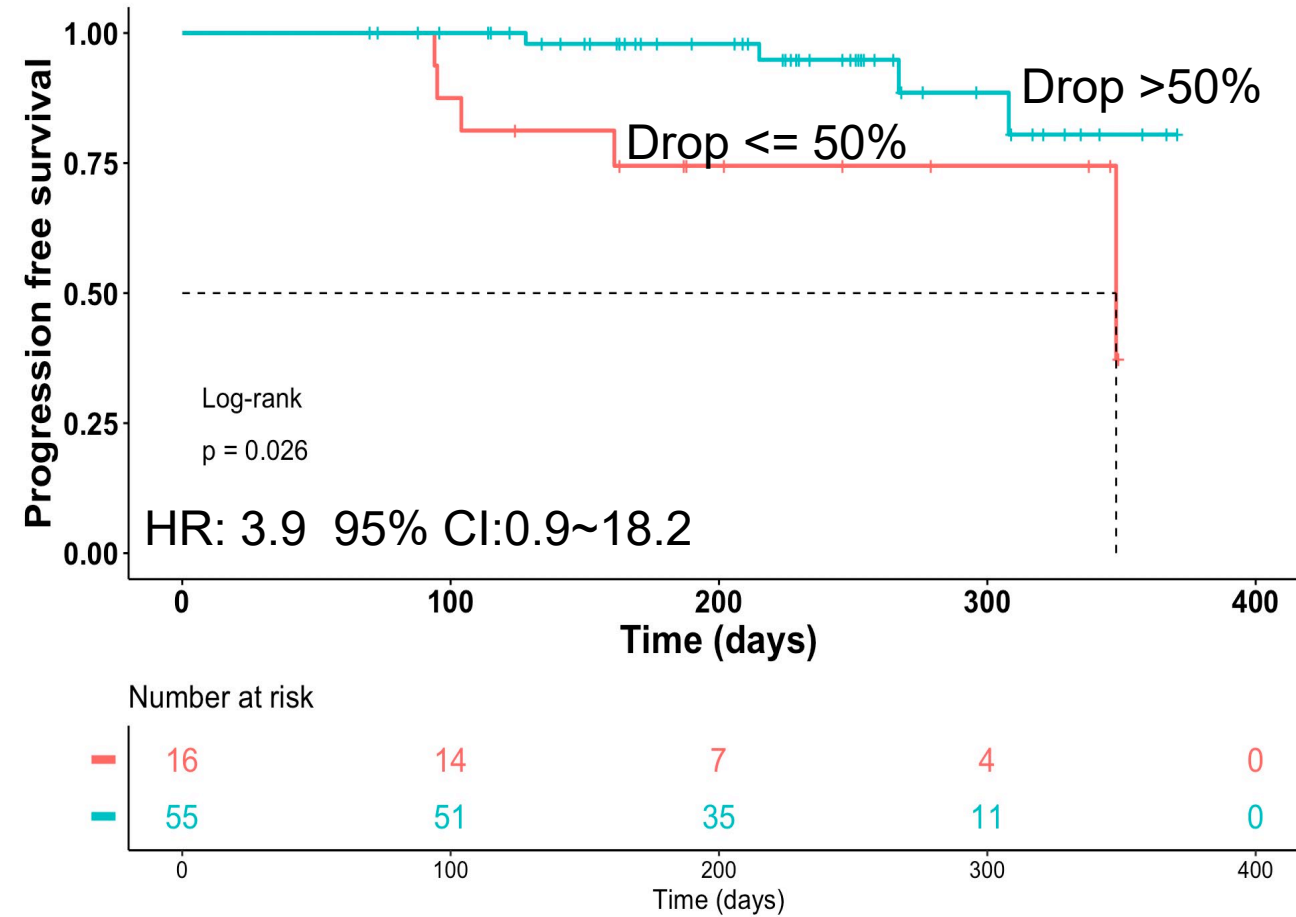
MTB dynamic change and the rate of MTB/ctDNA clearance are correlated with outcome

MTB dynamic changes in baseline + group



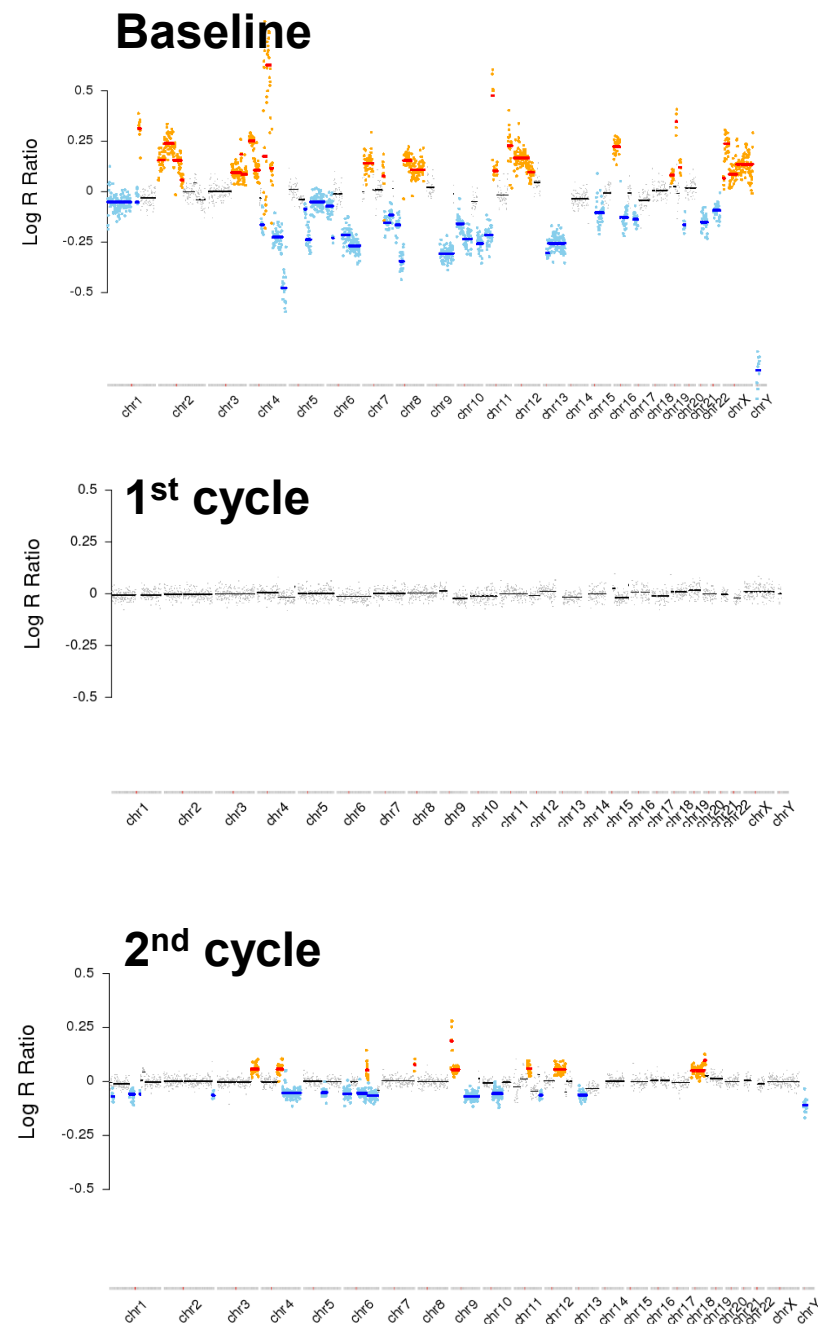
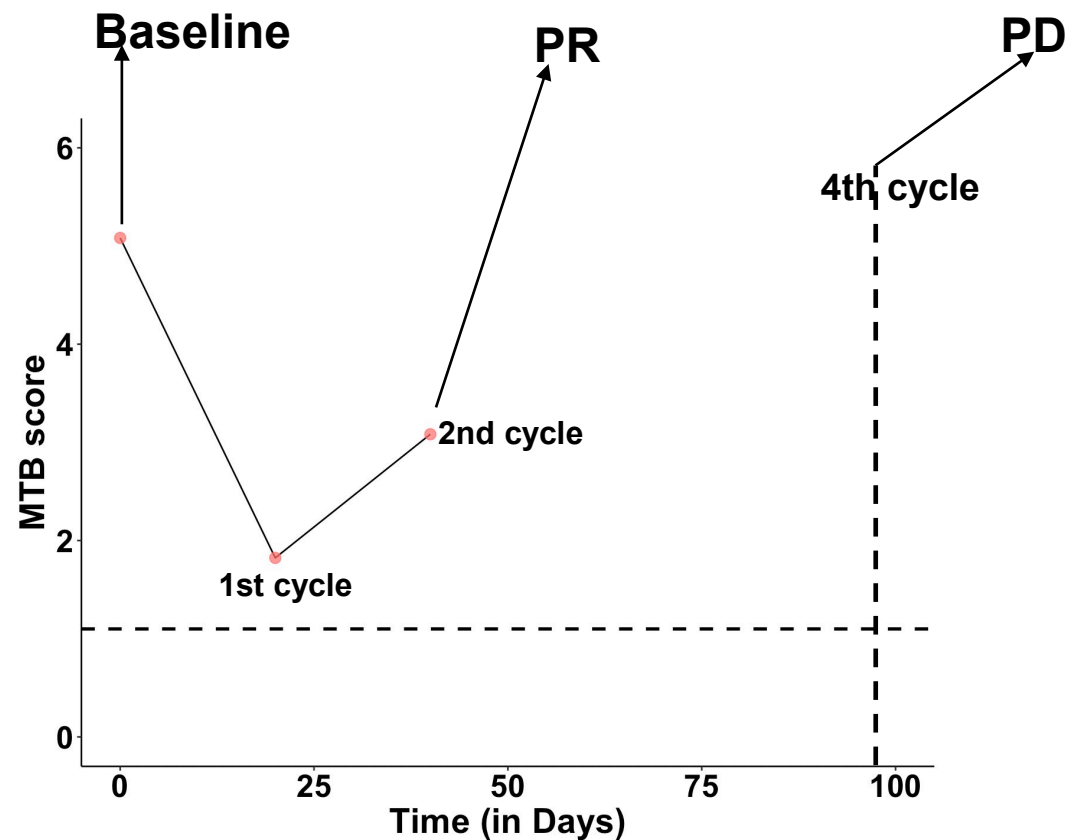
Up : landmark/baseline > 1  
Down: landmark/baseline < 1

MTB dynamic changes in baseline + and down group



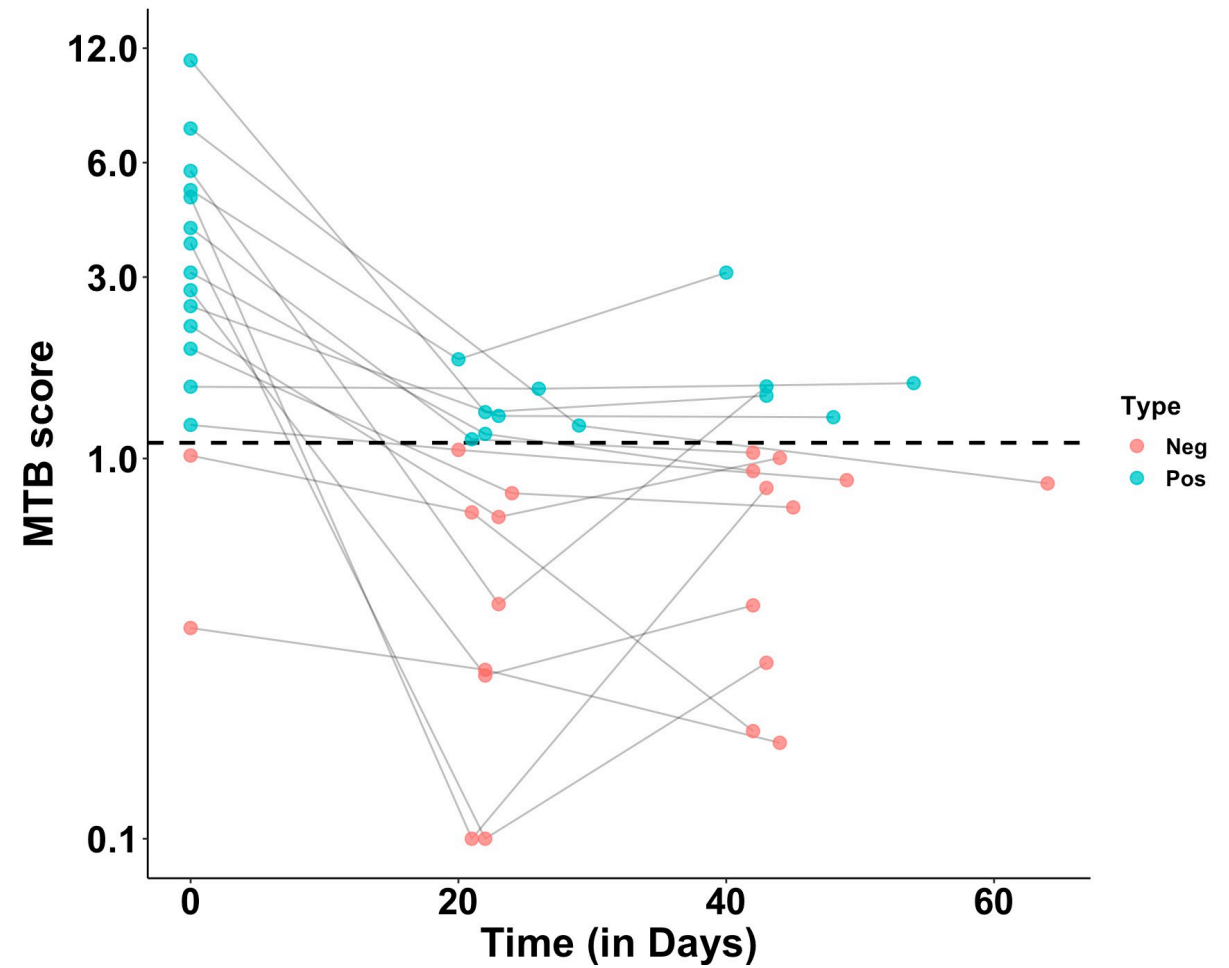
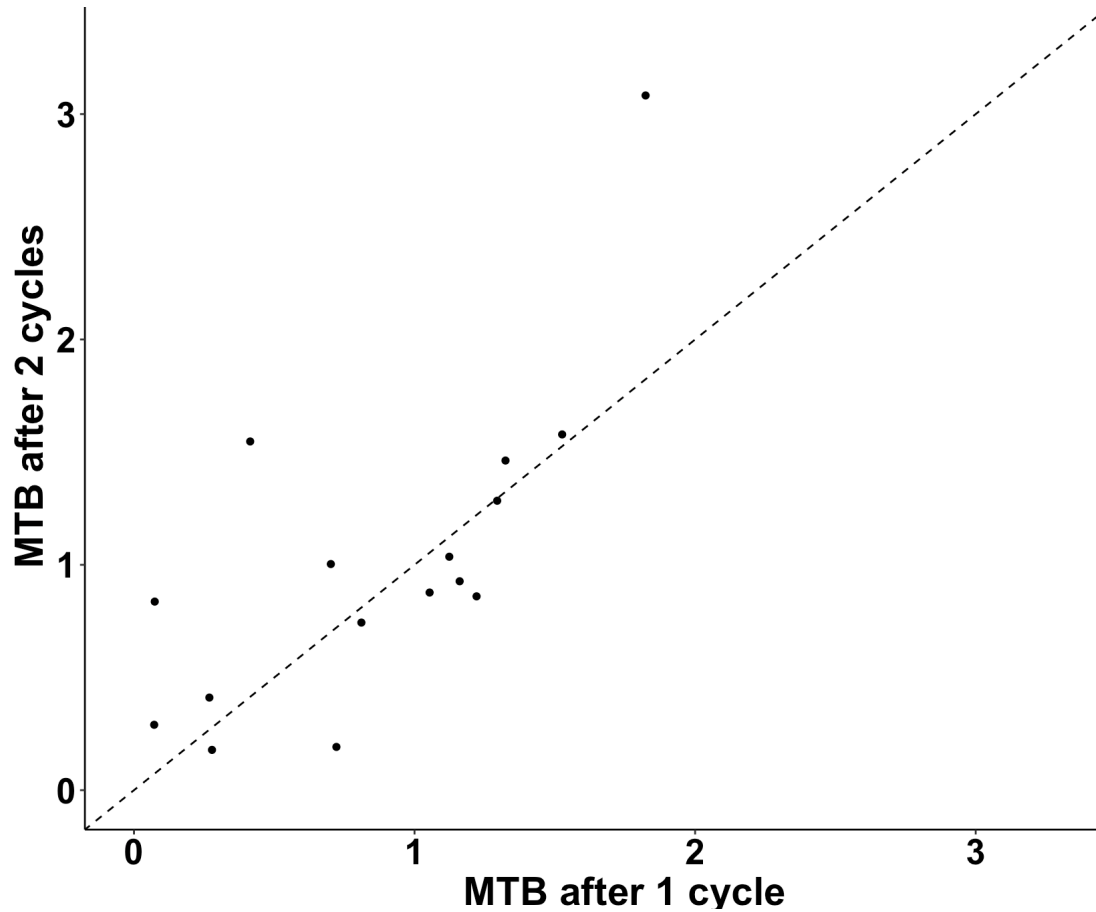


Dynamic change of MTB is prior to imaging (CT) change



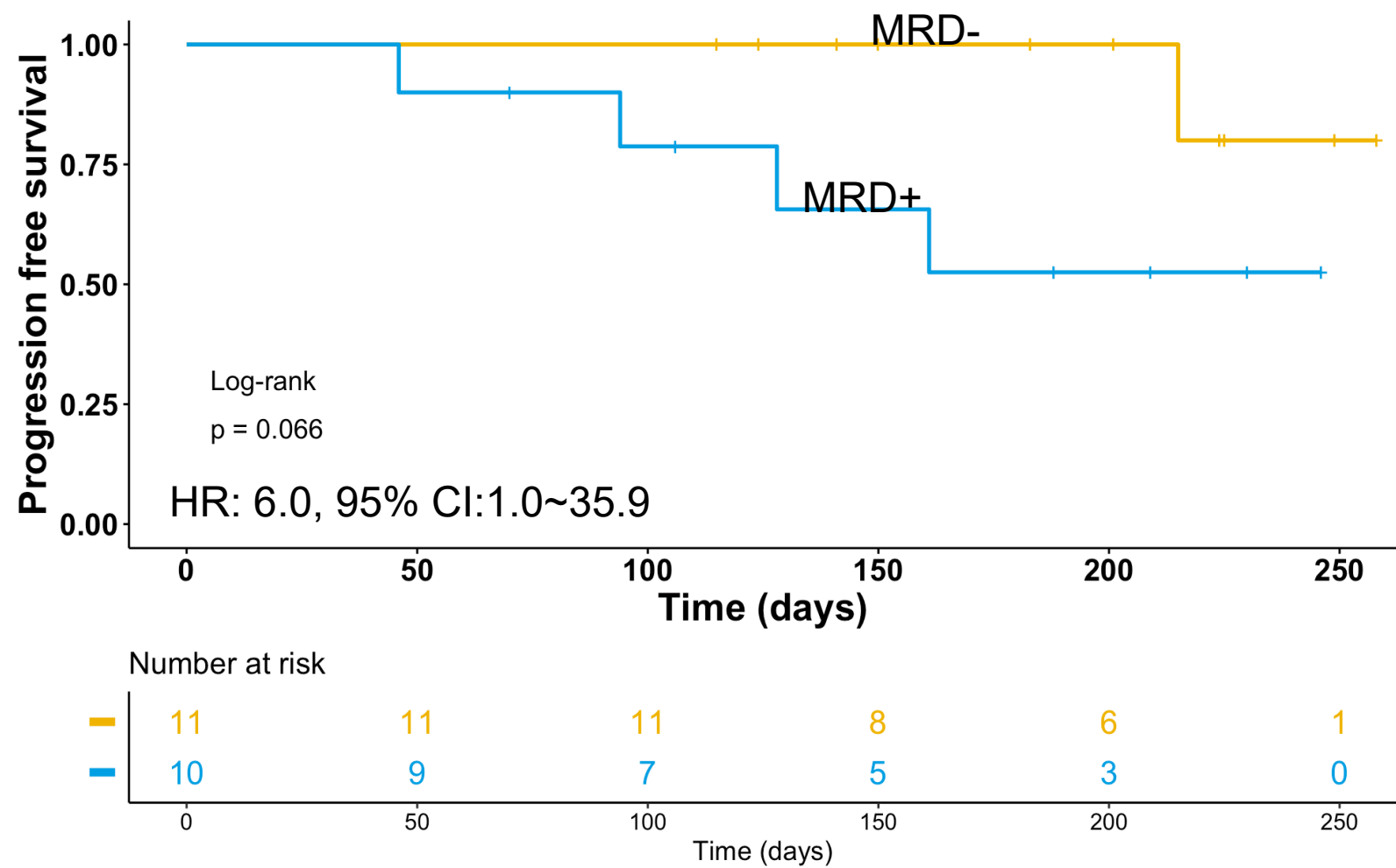
# MTB: 1 vs 2 cycles of treatment

MTB value after 1 cycle treatment has a good concordance with that after 2 cycles treatment



# MRD statue after one cycle treatment is sufficient to predict outcome

Even only after one cycle treatment, SeekInClarity also has prognostic value for therapeutic effectiveness



Quick and quantitative assessment of efficacy is the holy grail of oncology drug clinical development

## SeekInClarity™ – summary

- SeekInClarity is a cost-effective pan cancer/pan indication treatment response monitoring test that does not require cancer tissue analysis.
- Both baseline MTB and landmark MRD values are prognostic.
- Double negative (MTB- and MRD-) patients have a very favorable outcome (100% PFS).
- Double positive (MTB+ and MRD+) patients have the worst outcome.
- 60% baseline positive patients can achieve ctDNA/PTM clearance (MRD-) after 2 cycles of treatment.
- SeekInClarity analysis after 1 cycle of treatment is sufficient to assess response.

# Blood tests for getting right treatments at earlier time

Advancing equitable cancer care through innovation



Indicated subjects: Healthy population

Launch date: Nov. 2018



Post-op patients of radical resection

May 2019



Late-stage patients

Aug. 2020



One-Size-Fits-All

A photograph of two men shaking hands across a wooden table. The man on the left is older, with white hair, wearing a dark suit jacket over a red shirt. The man on the right is younger, with dark hair, wearing a light-colored striped button-down shirt. They are both smiling. In the background, there is a large whiteboard with a camera mounted on top. The entire image is framed by a thin white border.

**SEEKIN & INSPIRE2LIVE**

**GROUND-BREAKING NEW  
TECHNOLOGY FOR MULTI-  
CANCER EARLY DETECTION**

**JOINING FORCES FOR  
THE BENEFIT OF PATIENTS**

#WeAreThePatientsVoiceInCancer  
#IfAboutUsNotWithoutUs

A faint, light blue world map is centered on the slide, showing the continents of North America, South America, Europe, Africa, and Asia. Two white diagonal lines cross the slide: one from the top right to the bottom left, and another from the bottom left to the top right, intersecting near the center.

# Lead the World!

真理至上，以人为本

*seek and you will find*

– Matthew 7:7

# SeekInCare® - Comparison to other cancer early detection tests

	Method	Sensitivity	Specificity	Pan-cancer	Tissue of origin
Plasma tumor markers	chemiluminescence, flow cytometry based fluorescent microsphere...	30%	92%	✓	✓ (<20%)
Specialized cancer detection	varied based on different cancer types	varied based on different cancer types and technologies. Mainly applied for high-risk individuals.		✗	✗
Grail Multi-cancer	cfDNA methylation	51.5%	99%	✓	✓ (89%)
SeekInCare® Pan-cancer	cfDNA CNA & FS Plus PTMs	68%	98%	✓	✓ (67%)



# Comparison with other common technical approaches

Company	Technical methodology	Enrichment	Sequencing	Cost	Genomic coverage	Sensitivity	Specificity
Grail	cfDNA methylation panel	Targeted capture probes	139X NGS	NA	1 M methylation sites	51.5% <sup>1</sup>	99% <sup>1</sup>
Thrive	ctDNA mutation Targeted capture panel <sub>4</sub>	Targeted capture probes /Multiplex PCR based amplicons	3000X NGS	\$ 500	10-1000 genes	70% <sup>2</sup>	99% <sup>2</sup>
Delfi	cfDNA fragmentomics	In silico enrichment of cancer signals (Fragmenomics)	2X WGS	NA	Whole genome (3 B bases)	73% <sup>3</sup>	98% <sup>3</sup>
SeekIn	cfDNA panoramic view + Protein tumor markers	In silico enrichment of cancer signals (CNA and FS)	3X WGS	\$ 200	Whole genome (3 B bases)	75% <sup>4</sup>	99% <sup>4</sup>

1. Liu, M. C. et al. Sensitive and specific multi-cancer detection and localization using methylation signatures in cell-free DNA. Annals of Oncology 31, 745–759 (2020).

2. Cohen, J. D. et al. Detection and localization of surgically resectable cancers with a multi-analyte blood test. Science 359, 926–930 (2018).

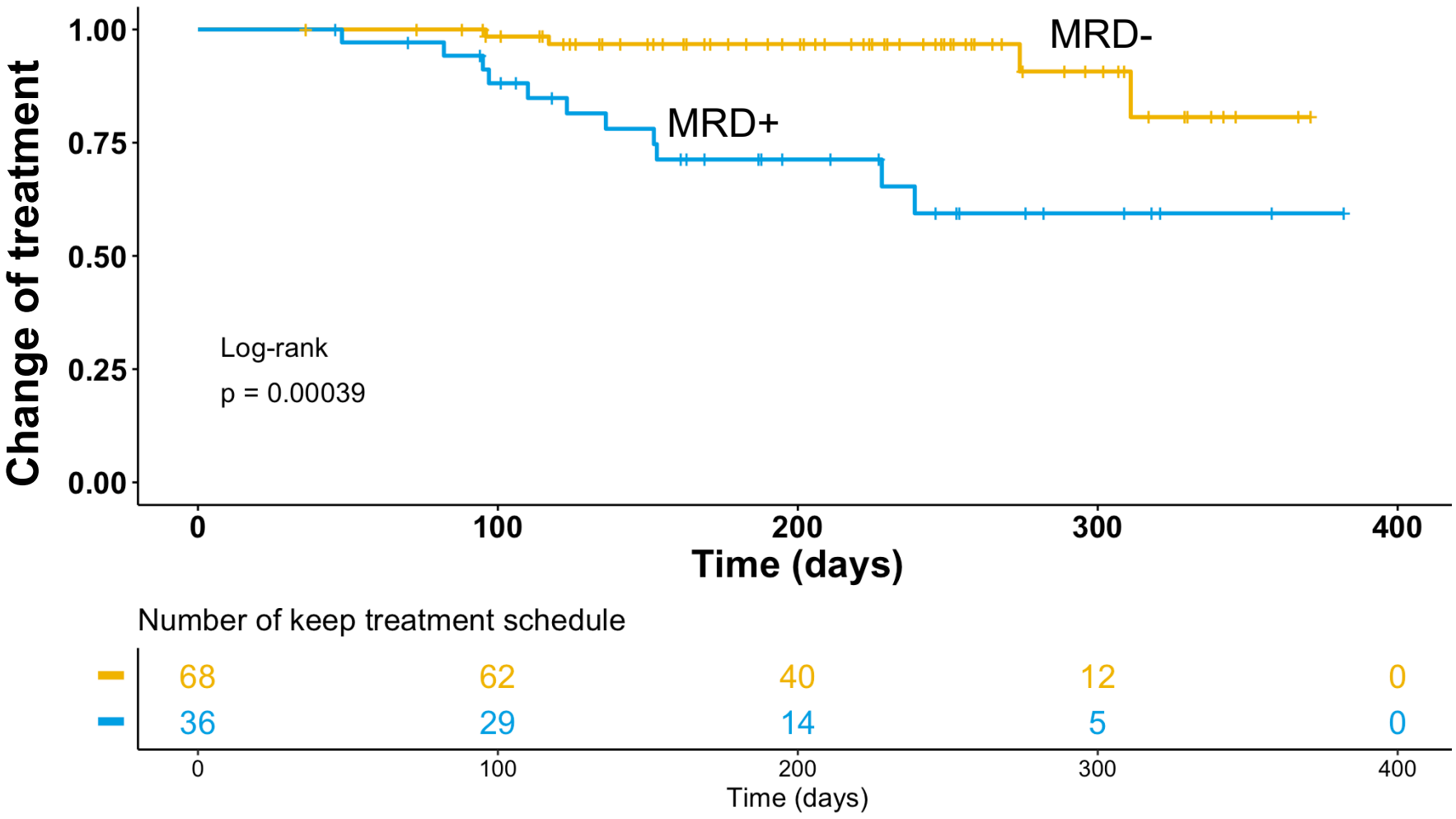
3. Cristiano, S. et al. Genome-wide cell-free DNA fragmentation in patients with cancer. Nature, 570(7761):385-389. (2019).

4. Mao, M. et al. Development of a blood-based cancer screening assay with a novel multivariate cancer risk score (MCRS) model by integrating shallow WGS data and plasma protein markers. Oral presentation at ASCO Breakthrough 2019.

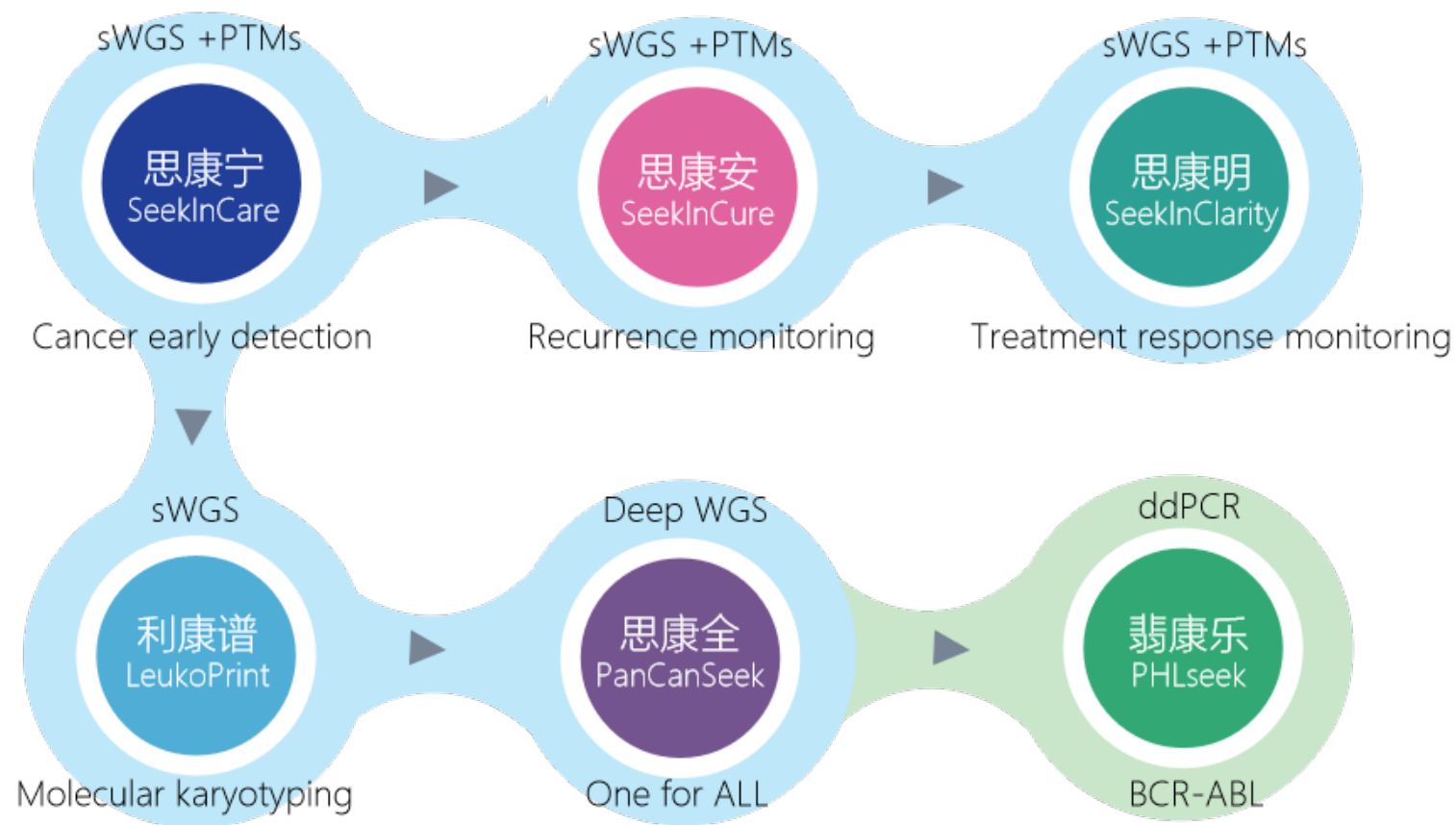


# SeekInClarity™ analysis at landmark

Time to changing the first-line treatment in MRD+ group is significant less than that in MRD- group.



# From Solid Tumor to Leukemia



# LeukoPrint® A new generation of genetic molecular technology



the first-in-class Leukemia sWGS CNA test

CE Marking



Without culturing cells in vitro

High sensitivity and low failure rate

Automatic readout system

By evaluating the copy number aberration (CNA) in Bone marrow cells and/or circulating cell free DNA (cfDNA) via shallow whole genome sequencing (sWGS), in conjunction with conventional cytogenetics, LeukoPrint® makes leukemia diagnosis, molecular subtyping, prognostic stratification and treatment responds monitoring more precise and comprehensive.

- **Sample requirement:** 10ml blood
- **TAT:** 10 work days
- **Results readout:** copy number aberration (CNA)
- **Indicated subjects:** Patients with suspected or confirmed leukemia (MDS/AML/ALL/CLL/MM)

# Comparison to other karyotyping tests

Coverage area
Resolution
Sample Requirement
Readout
Throughput
TAT
Cost

Karyotyping	FISH
23 pairs of chromosomes	Single probe
≥5Mb	≥100kb
High	High
Visual inspection	Visual inspection
Low	Low
7-10d	3-5d
Medium	High



## LeukoPrint®

Whole genome

≥100kb

Low

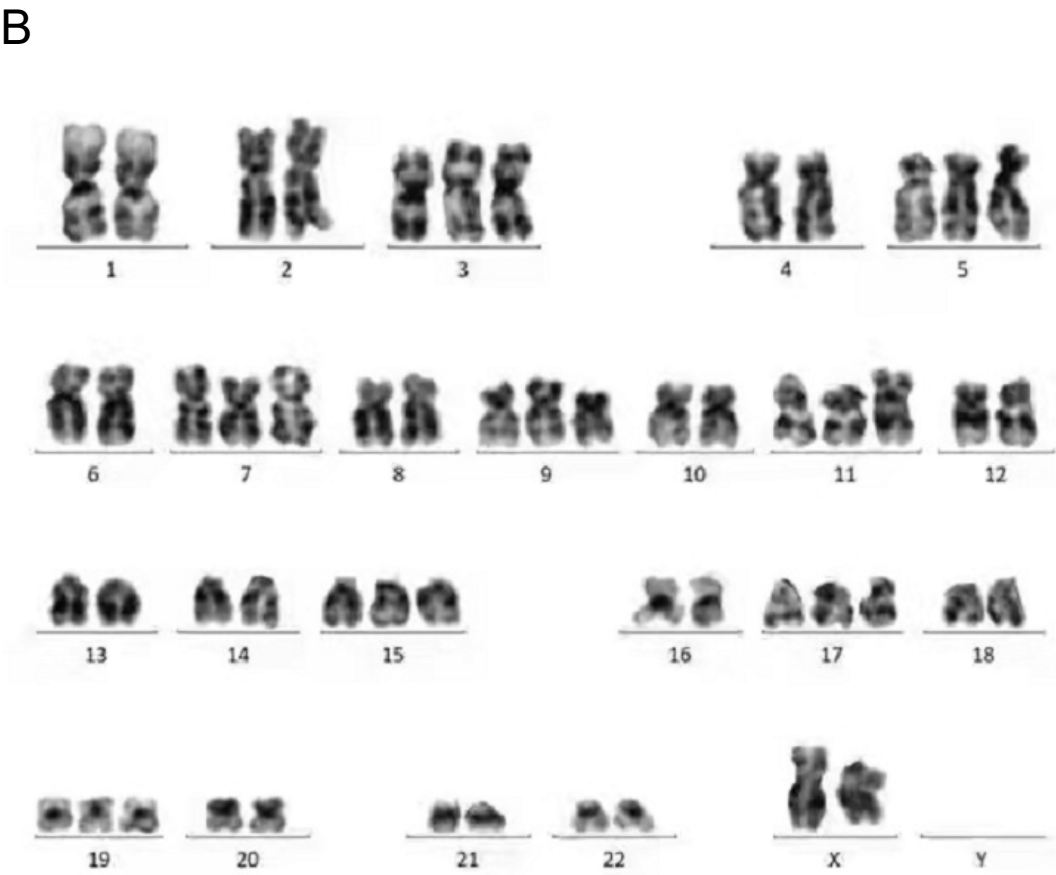
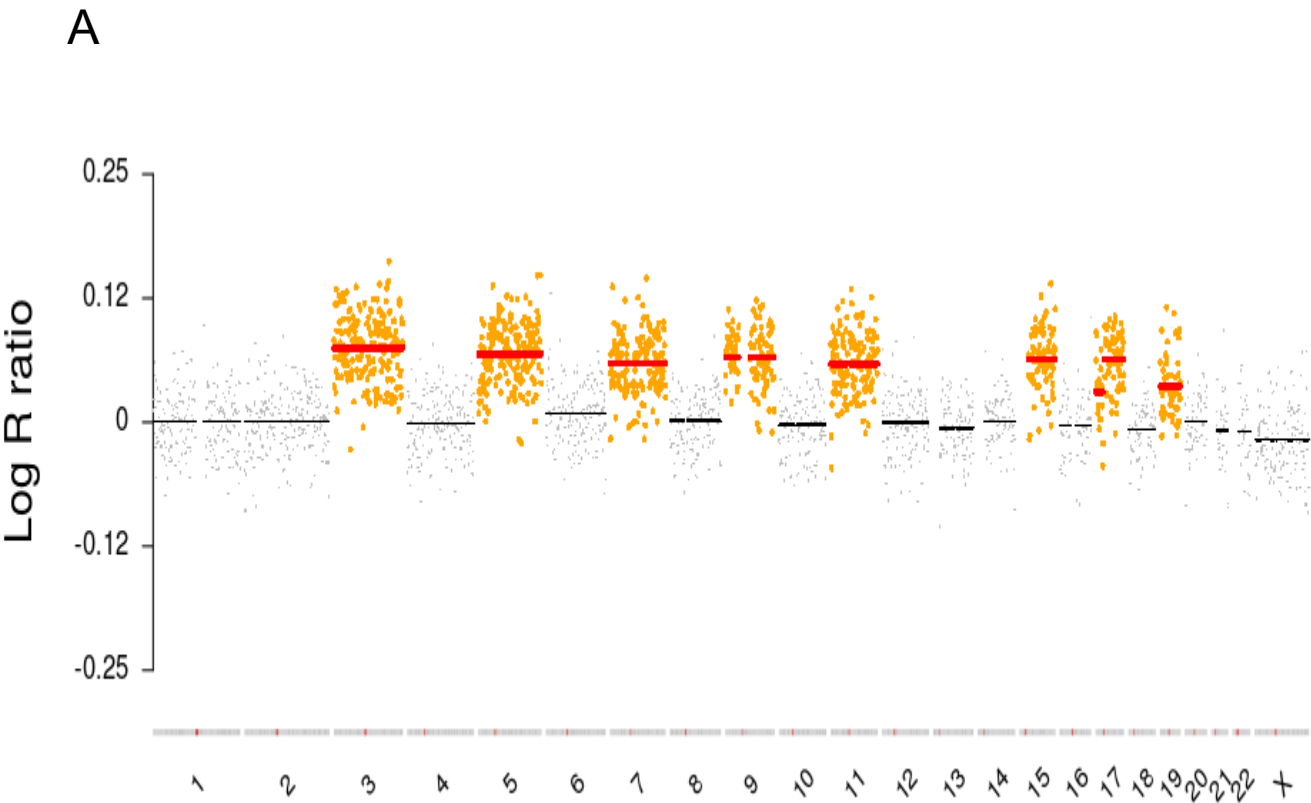
Digital

High

10-12d

Medium

Figure 2



染色体位置标注出来

# PanCanSeek® Integrated Solutions of Cytogenetics and Molecular Diagnosis in Leukemia



PanCanSeek®  
思康全®

CE Marking



**One for All**

01

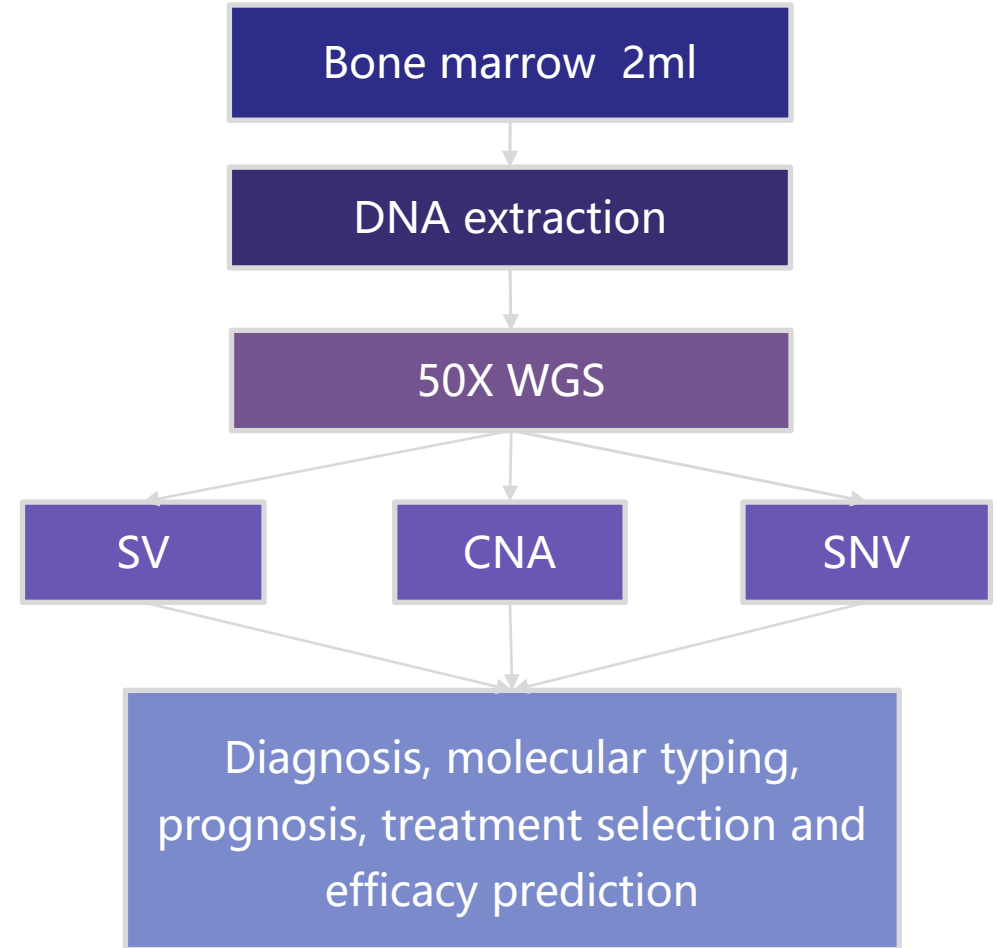
Structural variants (SV)

02

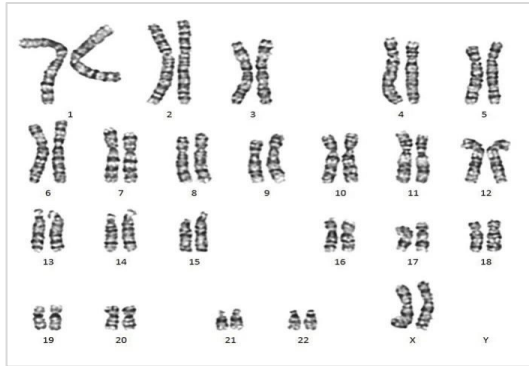
Copy number variation (CNA)

03

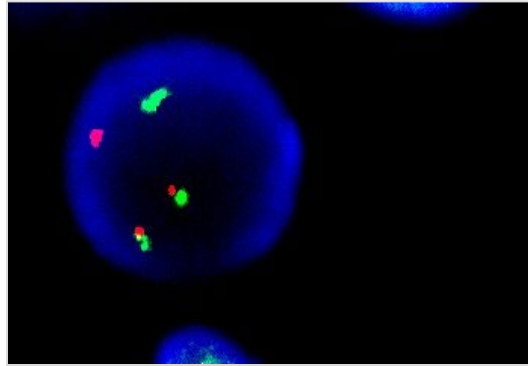
Point mutations (SNV, INDEL)



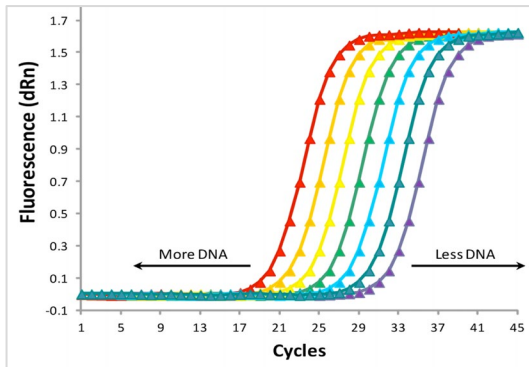
# PanCanSeek® faster, more accurate and more comprehensive



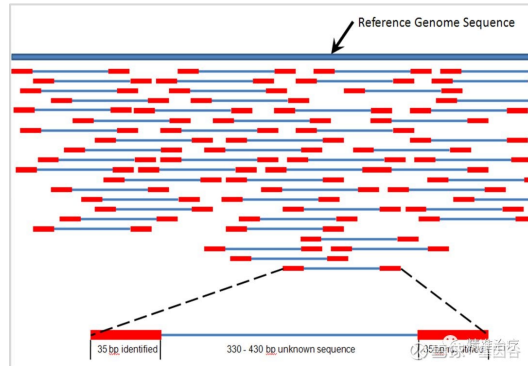
Karyotyping



FISH



qPCR



NGS Panel

思康全®PanCanSeek®



Compared with traditional diagnostic methods, PanCanSeek can complete the detection of all genetic mutations in leukemia patients in one single test - faster, more accurate and more comprehensive



# Product portfolio

