



HKG *epi* THERAPEUTICS Ltd.
Harnessing the epigenome

HKG *epi* THERAPEUTICS Ltd.

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HKG *epi*THERAPEUTICS Ltd.



CLIA ACCREDITED





- **Disclaimer: Forward-Looking Statements**

- The forecasts and projections presented in this business plan are forward-looking statements. Please note that actual results may vary and are subject to inherent uncertainties and risks. We caution readers not to place undue reliance on these forward-looking statements as they are based on current expectations and assumptions. The company undertakes no obligation to update or revise these statements.

Executive Summary: HKG *epi* Therapeutics



❖ **Mission and vision**

Pioneering early disease detection through epigenetics.



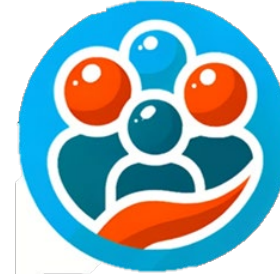
❖ **Unique Technology**

Advanced DNA methylation platform for accurate diagnoses.



❖ **Market Strategy**

Targeting key healthcare sectors globally with innovative solutions..



❖ **Expert Team**

Led by epigenetics pioneers committed to healthcare advancement...



❖ **Financial Snapshot**

Leadership committed to healthcare advancement...



Company mission

- Developing a platform for Early Detection of Disease
- Clinical Application of Epigenetics
- Building a global *epi*Health ecosystem; maintaining health and preventing disease through a cloud-driven, global platform of clinical epigenetic labs and one stop shop for epigenetic tests
- Our vision: *improve lives by detecting and preventing illnesses before they become a major health issue.*

OUR COMPANY I

- ✓ Company Name: **HKG epiTherapeutics Ltd**
- ✓ Date of Establishment: August 1 2016
- ✓ Employee Number: ~20 in HKG, Montreal and Israel and 14 in Shenzhen.
- ✓ Mission: To develop tools for early detection of cancer and promoting healthy aging
- ✓ Core Competitiveness:
 - ✓ All global IP protected by PCT and exclusively licensed or assigned to company
 - ✓ Low cost
 - ✓ High-throughput by automation (R&D with financially supported by the Innovation and Technology Department of HKG SAR)
 - ✓ Resale agreement for epiAging in Germany, HK, UK, Thailand
 - ✓ Global research collaborations and reach: Canada, USA, Israel, Brazil, Bangladesh, India Hong Kong and China.
 - ✓ CAP accredited and CLIA accredited lab in HK science park which was just audited and reaccredited.
- ✓ Goal: To develop a global world-leading innovative multi-cancer early screening platform



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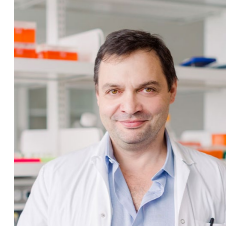
WORLD CLASS LEADERSHIP

Epigenetic pioneer is leading the company

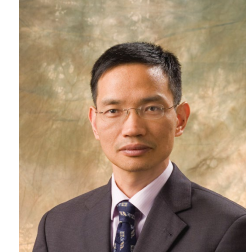


Moshe Szyf, PhD
Glaxo Smith Kline - CIHR chair in pharmacology
James McGill Professor
Department of Pharmacology and Therapeutics at
McGill University, Canada

- Fellow of the Royal society of Canada
- Fellow of the Canadian Academy of Health Sciences
- Pioneer in the field of epigenetics
- Inventor of the first patent in the field
- Founder of the field of behavioral and psychiatric epigenetics
- Published ~300 papers in the area
- 2009 Karolinska Research Lectures at Nobel Forum
- Ted Talk: How early life experience is written into DNA
- Distinguished lecture FDA 2010
- NIMH directors Innovation speaker series 2012
- Quebec Scientist of the year 2009
- UK Royal Society of Art lecture 2013



David Cheishvili, Ph.D
Chief Innovation officer



Ye Xiang, Ph.D
Director and member of
the board of directors



Levan Cheishvili
Chief Technology officer



Carlos Luiz, MD
Medical Director



Charlie Ban
CEO EpiEcho



Sharon Luk, Ph.D
Director of clinical
Services



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Corporate Structure

Board of Directors

HKG epiTherapeutics Hong Kong
(38% Szyf: 62% 8 investors)

MTL Epitherapeutics Ltd.
(90% HKG epitherapeutics, 10%
David Cheishvili)

INEX Epitherapeutics Ltd.
Singapore
(50% HKG epitherapeutics, 50%
Inex)

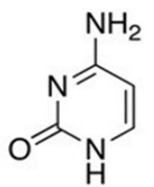
EpiEcho WOFE Beijing China
(100% HKG epitherapeutics)

VIE structure

Epishen (Shenzhen)

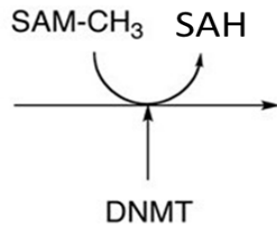
What is DNA methylation? Why is it the next big thing in medicine?

unmethylated

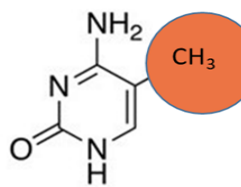


Cytosine

ON



Methylated

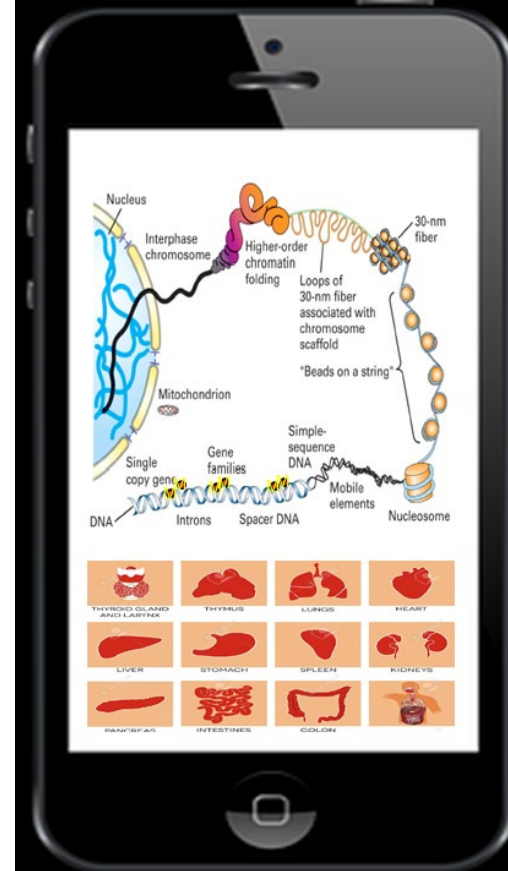


5-Methylcytosine

OFF

DNA methylation is the software of the genome which is written during embryonal development

The DNA methylation software is “bugged” in disease. We can detect these “bugs” using our methods.



Normal

Cancer

Normal

Cardiovascular disease

Normal

Metabolic disorders

Normal

Obesity

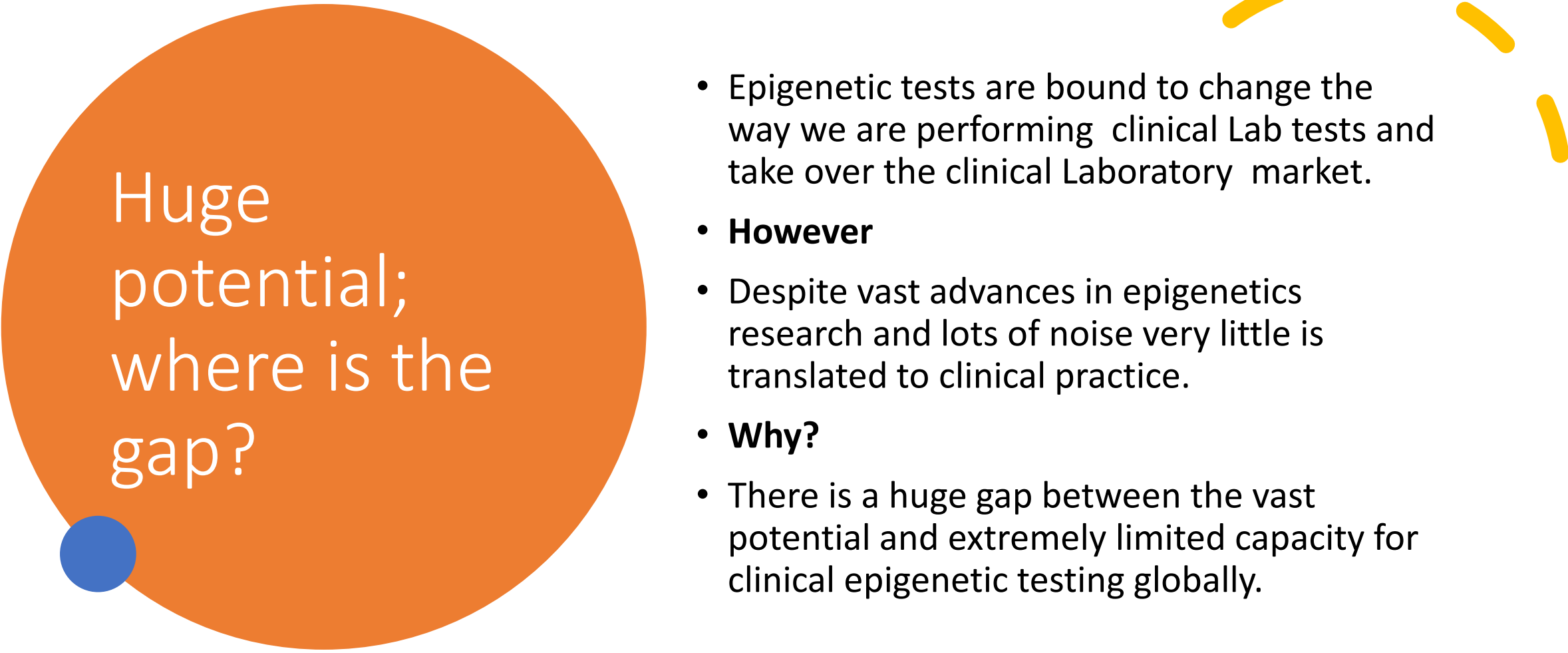
Normal

Mental health disorder

Normal aging

Accelerated aging

DNA methylation is the code of
our biological state, our
exposures and experiences, we
know how to read this code and
be informed by it.



Huge
potential;
where is the
gap?

- Epigenetic tests are bound to change the way we are performing clinical Lab tests and take over the clinical Laboratory market.
- **However**
- Despite vast advances in epigenetics research and lots of noise very little is translated to clinical practice.
- **Why?**
- There is a huge gap between the vast potential and extremely limited capacity for clinical epigenetic testing globally.

The next big thing



A high throughput, robust global platform for delivery of an evolving landscape of clinical epigenetic tests.



A clinical epigenetic lab blue-print which is the focal node for a health building and maintenance ecosystem.



A clinical lab for the 22nd century that revolutionizes clinical laboratory medicine.



Our plan

Focus

- Focus not on a test, but on a unified platform for many diverse tests

Translate

- Translate epigenetic discoveries to a panel of clinically validated and accessible tests through a unified platform for, discovery, research and development.

Develop

- Develop a scalable automated cloud-based platform for high throughput-one shop delivery of clinical epigenic tests.

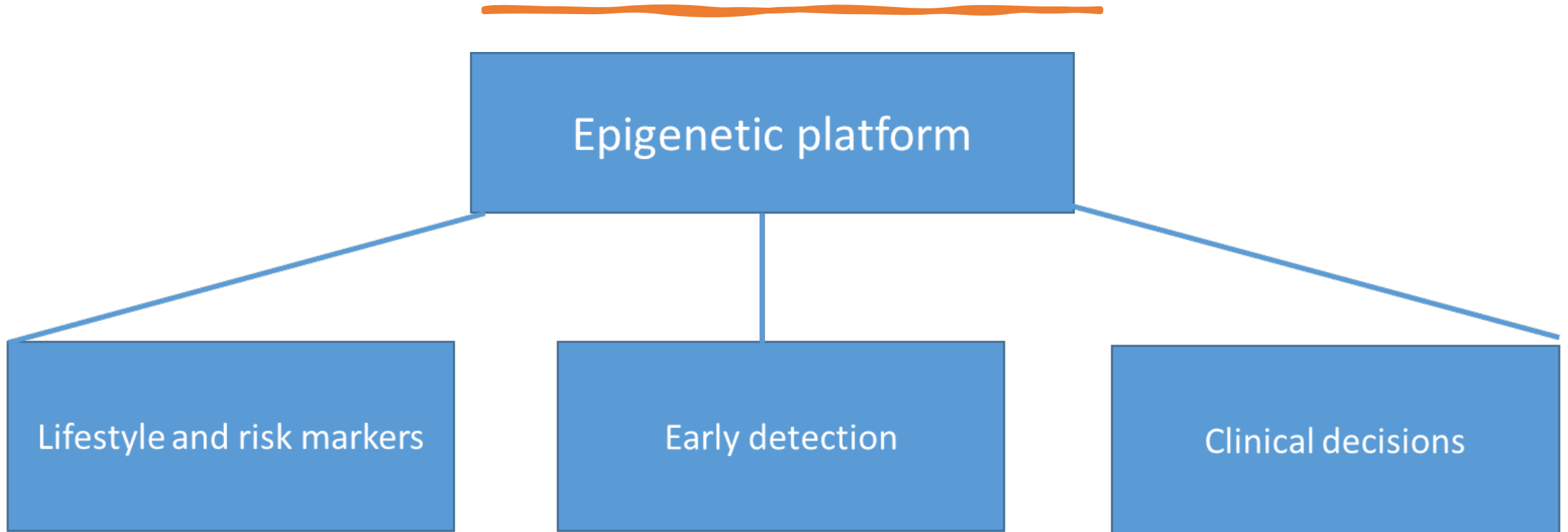
Create

- Create a blueprint of the clinical epigenetic lab of the 22nd century that would be implemented globally

Integrate

- Integrate epigenetic testing into a platform of a multitiered epihealth ecosystem.

Epigenetic markers portfolio



Products and Production Laboratory Overview



Global Operations and Expertise in Epigenetic Testing: HKG epiTHERAPEUTICS Ltd

Innovation and Data
Analysis
(Montreal, Canada)

Clinical Management
and Regulatory
Oversight
Sanford, FL. USA

Computation and
Cloud Management
(Israel)

Test Production and
R & D
(Hong Kong science
park)

China Test
Production Lab
Shenzhen

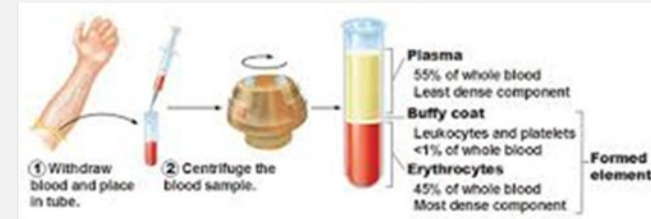
Products : Ready to market test Production system and lab launched in 2021



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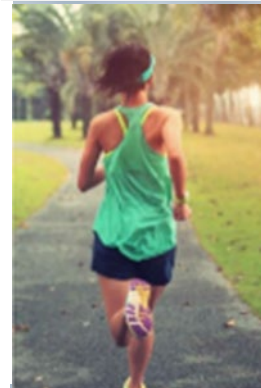


Early Detection of Liver Cancer



Early detection product pipeline including panCancer product.

Early Detection of Cervical Cancer



Detection of Biological Age



*epi*Smoke
*epi*Age (Saliva)



*epi*Cervix (Pap smears)

Our regulatory strategy is via LDT path; The lab in HKSTP is CLIA and CAP accredited and CLIA registered; our plan is to copy this platform in strategic global locations



COLLEGE of AMERICAN
PATHOLOGISTS

CERTIFICATE OF ACCREDITATION

**HKG epiTherapeutics Ltd
Laboratory
Hong Kong, China
Carlos J. Ruiz, MD**

CAP Number: 9143982
AU-ID: 2142729

The organization named above meets all applicable standards for accreditation and is hereby accredited by the College of American Pathologists' Laboratory Accreditation Program. Reinspection should occur prior to **December 27, 2023** to maintain accreditation.

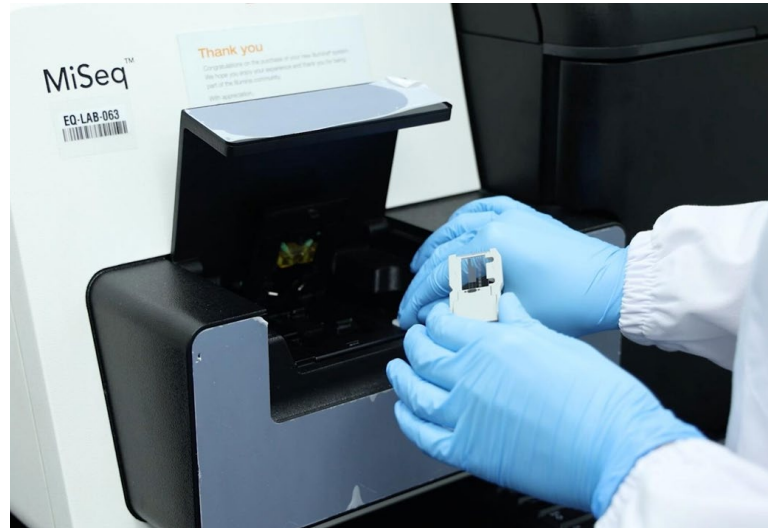
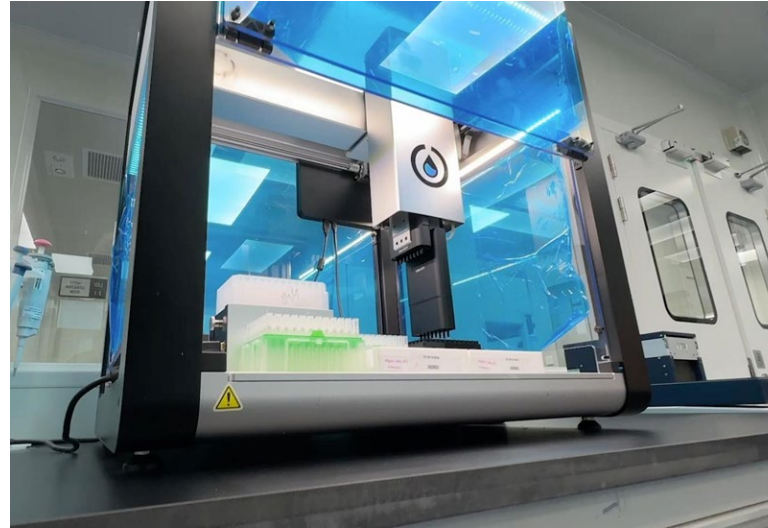
Accreditation does not automatically survive a change in director, ownership, or location and assumes that all interim requirements are met.

Michael Bradley Datto, MD, PhD, FCAP
Chair, Accreditation Committee

Emily Volk, MD, FCAP
President, College of American Pathologists



HKG Epigenetic Test "Production" Laboratory at HKSTP



- Laboratory Construction Tailored for Biochemical Testing Procedures.
- Segregated Spaces for Each Testing Step.
- Automated Testing Process with Cloud-Based Management, Analysis and Reporting.

Recent peer reviewed clinical validations of our tests

nature communications



Article

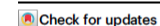
<https://doi.org/10.1038/s41467-023-39055-7>

A high-throughput test enables specific detection of hepatocellular carcinoma

Received: 24 September 2022

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Published online: 07 June 2023



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High-throughput tests for early cancer detection can revolutionize public health and reduce cancer morbidity and mortality. Here we show a DNA methylation signature for hepatocellular carcinoma (HCC) detection in liquid biopsies, distinct from normal tissues and blood profiles. We developed a classifier using four CpG sites, validated in TCGA HCC data. A single F12 gene CpG site effectively differentiates HCC samples from other blood samples, normal tissues, and non-HCC tumors in TCGA and GEO data repositories. The markers were validated in a separate plasma sample dataset from HCC patients and controls. We designed a high-throughput assay using next-generation sequencing and multiplexing techniques, analyzing plasma samples from 554 clinical study participants, including HCC patients, non-HCC cancers, chronic hepatitis B, and healthy controls. HCC detection sensitivity was 84.5% at 95% specificity and 0.94 AUC. Implementing this assay for high-risk individuals could significantly decrease HCC morbidity and mortality.

Received: 27 January 2023 | Revised: 17 July 2023 | Accepted: 21 July 2023

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RESEARCH ARTICLE

Cancer Genetics and Epigenetics



Validation of novel DNA methylation markers in cervical precancer and cancer

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²Gerald Bronfman Department of Oncology, McGill University, Montréal, Quebec, Canada

³HKG Epitherapeutics, Hong Kong

⁴Department of Pharmacology and Therapeutics, McGill University, Montréal, Quebec, Canada

Correspondence

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Funding information

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Abstract

We have recently identified, using a genome-wide approach, new methylation markers which were evaluated among various cervical intraepithelial neoplasia (CIN) grades and cervical cancer. We herein validate the methylated state of these genes in independent study populations, based on histology ascertained outcomes regardless of human papillomavirus status. *CA10*, *DPP10*, *FMN2* and *HAS1* (discovery set: 54 normal, 50 CIN1, 40 CIN2, 42 CIN3) were evaluated by targeted bisulfite next generation sequencing (NGS) (Illumina MiSeq platform) in 258 (training set: 100 normal, 50 CIN1, 50 CIN2, 50 CIN3, 8 cancers) and 373 (validation set: 100 normal, 57 CIN1, 61 CIN2, 53 CIN3, 102 cancers) physician-collected samples (PreservCyt). Using targeted amplification NGS data from the training set for 94 normal and eight cancer samples, we calculated for each gene the median methylation value. These were summed and normalized to compute a four-gene Marker Polygenic Score (MPS). We compared the relationship between MPS and progression from normal through CIN grades and cancer, separately in the training and validation sets, and tested its clinical performance via receiver-operating characteristic curves. MPS increased with increasing CIN grade, and accurately predicted cervical cancer in the

Thank
you



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- For more info:
<https://www.hkgepitherapeutics.com/>
 - <https://www.epi-age.com/>