



# <section-header>

Discover and validate miRNA expression profiles in a single step.





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# ID3EAL<sup>™</sup> PanoramiR microRNA Knowledge Panel

# Meaningful insights with minimal resource investments.

# Thomas # CHM02D121

The ID3EAL PanoramiR panel is an effective approach to screening for miRNAs important in biological processes, disease mechanisms, or biomarker discovery for disease diagnosis and subtyping.

PanoramiR targets 376 miRNAs expertly curated from miRBase 22, HMDD 3.2 and TCGA. This is further cross-referenced with high-impact publications and in-house research data.



#### **Achieve More with Less**

- **Expertly curated disease-relevant** miRNA panel generates meaningful insights.
- ✓ Highly sensitive and robust ID3EAL™ RT-qPCR based technology enables detection of lowly expressed miRNAs with minimal sample input.
- ✓ **Validated** multiplexed primer pools prevents resource wastage.
- Relevance to over 20 diseases including cancers, dementias, cardiovascular disease, liver diseases and more.
- Relevance to various sample types including plasma, serum and solid tissues.

Category	Examples	microRNA Targets
Cardiovascular	Cardiovascular Disease (e.g. stroke, coronary artery diseases, etc)	140
Metabolic	Diabetes, Liver Cancer	358
Neurological	Alzheimer's Disease, Dementia, Depression, CNS Cancer	114
Respiratory	Lung Disease, Lung Cancer	165
Urological	Kidney Disease, Bladder Cancer	51
Reproductive	Cervical Cancer, Prostrate Cancer	319
Gastrointestinal	Stomach Cancer, Colorectal Cancer	224
Hematological	Leukemia, Lymphoma	150

#### **Disease Categories**

References

Key principles of miRNA involvement in human diseases. Discoveries (Craiova). 2014 Oct-Dec; 2(4): e34.

# Publishable Results In Hours

#### Fast: RNA-to Ct in 2 or 3.5 hours

The proprietary ID3EAL<sup>™</sup> RT-qPCR technology enables results from RNA samples in just 2 hours or up to 3.5 hours with the pre-amplification step to enhance detection with limiting amounts of starting material.

#### **Convenient: Great User Experience**

- Ready-to-use qPCR plate with pre-spotted primers
- Reagents pre-aliquoted into experimentally relevant proportions
- Wet-lab validated assays
- Spike-in RNAs included
- Easy-to-interpret fold-change data from data analysis excel template.



# VItrasensitive and Robust miRNA Detection

Achieve unparalleled sensitivity and specificity with wet-lab validated sequence-specific ID3EAL<sup>™</sup> RT and qPCR forward and reverse primers.



ID3EAL<sup>™</sup> assay primers are optimized by MiRXES' proprietary algorithm and wetlab validated with synthetic miRNA templates and RNA from biological samples. Manufactured in ISO 13485 facility.

# ID3EAL™ miRNA Knowledge Panel Profiling Service

The ID3EAL<sup>™</sup> miRNA Knowledge Panel Profiling Service enables researchers to overcome time, resource and expertise challenges and generate high quality miRNA data for publishing in high impact journals.

Delivering Your microRNA Discoveries



# ID3EAL<sup>™</sup> miRNA Knowledge Panel Profiling Service

## **Delivering Your microRNA Discoveries**

The ID3EAL<sup>™</sup> miRNA Knowledge Panel Profiling Service enables researchers to overcome time, resource and expertise challenges and generate high quality miRNA data for publishing in high impact journals.

- Experienced Scientists with Expert Know-How
- High Quality, Reliable and Reproducible Data
- ✓ Wide Dynamic Coverage of Over 7 logs
- Stringent Quality Controls
- Low Sample Input Requirement
- Publication-Ready Bioinformatics Analyses
- Data Confidentiality

**Proprietary pre-analytical specimen quality** check addresses largest source of variability **in laboratory testing.** 

# From Sample to Publication-Ready Results

Researchers may opt for selected service modules or engage our complete service workflow.







# miRNA Expression Analyses by a MiRXES Partner Service Lab\* using the ID3EAL<sup>™</sup> miRNA Knowledge Panels.

# PCA analysis of miRNA profiles show stratification into distinct sample types and groups

miRNA expression in cultured HCT116 cells and their spent cell culture media (CCM). Principle component analysis (PCA) of miRNA expression data showed clustering of biological replicates of the respective sample types and groups.

\*The experiments and analyses were conducted by a MiRXES partner service lab, the National University of Singapore Yong Loo Lin School of Medicine ncRNA Core.



# Clustering and Volcano plot analyses of blood cancer subtypes by miRNA profiling on serum or platelet-poor plasma (PPP) samples

Unsupervised hierarchical clustering of miRNA expression data stratified uterine cancer samples into various molecular subtypes. Volcano plots of miRNA expression in serum and PPP samples from the same patient revealed identical dysregulated miRNAs.







#### ID3EAL<sup>™</sup> miRNA Profiling Panels

**ParanomiR Panel:** 376 disease-associated miRNAs selected from expert curation of miRNA databases, published literature and in-house research data.

**Biofluid Panel:** 176 miRNAs selected from in-house research of more than 20,000 human serum, plasma and urine specimens in both healthy and pathological conditions.

**Cancer Panel:** 352 miRNAs strongly associated with various cancer types and which regulate key onco- and tumor suppressor genes and pathways.

## **Standard Analyses**

Raw Data Cleaning • Normalization •
Fold-change and p-values • Clustering,
Principal Component Analysis • Volcano
Plots • Gene Ontology on request

## People, Experiences & Expertise

MiRXES scientists and collaborators from top academic and medical institutes have evaluated over 50,000 clinical samples in 30 disease areas and applications using the proprietary ID3EAL<sup>™</sup> technology.

## **Ordering Information:**





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