miProfile™ Human Gastric Cancer miRNA qPCR Array

For focused group profiling of human gastric cancer related miRNA expression

Cat. No. QM008-A (1 x 96-well plate, Format A)
Cat. No. QM008-B (1 x 96-well plate, Format B)
Cat. No. QM008-C (1 x 96-well plate, Format C)
Cat. No. QM008-D (1 x 96-well plate, Format D)
Cat. No. QM008-E (1 x 96-well plate, Format E)

Available as 1 set or 6 sets. Each set contains 80 unique miRNA primers deposited in one 96-well plate.

Introduction

The miProfile human gastric cancer miRNA qPCR array profiles 80 aberrantly expressed miRNAs most relevant to gastric cancer. Gastric cancer is also called stomach cancer, referring to cancer arising from any part of the stomach. Although Helicobacter pylori infection is the main risk factor in about 80% or more of gastric cancers, the molecular pathway underlying Helicobacter pylori infection leading to the development of gastric cancers remains unclear. Better knowledge of changes in gene expression during proliferation and metastasis may lead to improvements in the treatment of advanced gastric cancer. Aberrant miRNA expression has also been frequently reported in gastric cancer, indicating that there is a close correlation between miRNAs and human malignancy, suggesting that miRNA is a novel diagnostic biomarker for gastric cancer detection and should be assessed for its clinical applications in monitoring disease progression. This array contains 80 miRNAs which were reported to be involved specifically in gastric cancer, providing researchers a convenient way to study the regulation of miRNA expression associated with human gastric cancer.

- QM008 plate 01: 80 unique miRNA PCR primer pairs

Shipping and storage condition

Shipped at room temperate
Stable for at least 6 months when stored at -20°C

Array format

GeneCopoeia provides five qPCR array formats (A, B, C, D, and E) suitable for use with the following real-time cyclers.

Important note: Upon receiving, please check to make sure that the correct array format was ordered to ensure the compatibility with your qPCR instrument.

<table>
<thead>
<tr>
<th>Plate format</th>
<th>Instrument provider</th>
<th>qPCR instrument model</th>
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<tbody>
<tr>
<td>A (96-well)</td>
<td>Applied Biosystems</td>
<td>5700, 7000, 7300, 7500, 7700, 7900HT (Standard 96-well block), ViiA™7 (Standard 96-well block)</td>
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<tr>
<td>B (96-well)</td>
<td>Applied Biosystems</td>
<td>7500 (Fast block), 7900HT (Fast block), StepOnePlus™, ViiA™7 (Fast block)</td>
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<td>C (96-well)</td>
<td>Bio-Rad Laboratories</td>
<td>iCycler iQ®, MyiQ™, iQ™5</td>
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<td>D (96-well)</td>
<td>Bio-Rad Laboratories</td>
<td>CFX96™, DNA Engine Opticon™, DNA Engine Opticon 2™, Chromo4™</td>
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<tr>
<td>E (96-well)</td>
<td>Roche Applied Science</td>
<td>LightCycler® 480 (96-well block)</td>
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Quality control

1. Each miRNA-specific primer in the miProfile miRNA qPCR array has been experimentally validated to yield a single dissociation curve peak and to generate a single amplicon of the correct size for the targeted miRNA.

2. The positive PCR controls (PCR) have been verified to amplify a single amplicon of the correct size with Ct values around 20±2.

3. The Spike-in reverse transcription controls (RT) have been verified to amplify a single amplicon of the correct size with Ct values around 20±3.

4. R² > 0.99 was observed for high inter/intra-array reproducibility.

Materials required but not provided

All-in-One™ miRNA First-Strand cDNA Synthesis Kit
All-in-One™ qPCR Mix
Total RNA extraction kit (RNAzol® RT RNA extraction reagent is recommended)
DNase/RNase free tips, PCR reaction tubes, 1.5 ml microcentrifuge tubes
5 ml and 10 ml graduated pipettes, beakers, flasks, and cylinders
10 μl to 1,000 μl adjustable single channel micropipettes with disposable tips
5 μl to 20 μl adjustable multichannel micropipette, disposable tips, and reservoir
qPCR instrument, compatible with gene qPCR arrays ordered

Array layout

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<td>HK4</td>
<td>HK5</td>
<td>HK6</td>
<td>RT</td>
<td>RT</td>
<td>PCR</td>
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- **miRNA primer pairs:** Wells 1-84 are designated wells for pre-deposited miRNA primer pairs.
- **NC:** Negative controls, which only have the pre-deposited reverse universal primers.
- **HK1-6:** Six pre-deposited housekeeping snRNAs primer pairs, which can be used as endogenous positive controls as well as for array normalization.
- **RT:** Three replicates of spike-in reverse transcription controls, which can be used to monitor the efficiency of the RT reaction. These pre-deposited primer pairs specifically amplify the cDNA template reversed transcribed from the spike-in exogenous RNA in the sample.
- **PCR:** Three replicates of positive PCR controls, which are used to verify the PCR efficiency by amplifying the pre-deposited DNA template with its specific pre-deposited primer pairs.

**miRNA primer list**

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