

RT² Profiler PCR Array (96-Well Format and 384-Well [4 x 96] Format)

Human Induced Pluripotent Stem Cells

Cat. no. 330231 PAHS-092ZA

For pathway expression analysis

| Format | For use with the following real-time cyclers |
|---|--|
| RT ² Profiler PCR Array, Format A | Applied Biosystems® models 5700, 7000, 7300, 7500, 7700, 7900HT, ViiA™ 7 (96-well block); Bio-Rad® models iCycler®, iQ™ 5, MyiQ™, MyiQ2; Bio-Rad/MJ Research Chromo4™; Eppendorf® Mastercycler® ep realplex models 2, 2s, 4, 4s; Stratagene® models Mx3005P®, Mx3000P®; Takara TP-800 |
| RT ² Profiler PCR Array, Format C | Applied Biosystems models 7500 (Fast block), 7900HT (Fast block), StepOnePlus™, ViiA 7 (Fast block) |
| RT ² Profiler PCR Array, Format D | Bio-Rad CFX96™; Bio-Rad/MJ Research models DNA Engine Opticon®, DNA Engine Opticon 2; Stratagene Mx4000® |
| RT ² Profiler PCR Array, Format E | Applied Biosystems models 7900HT (384-well block), ViiA 7 (384-well block); Bio-Rad CFX384™ |
| RT ² Profiler PCR Array, Format F | Roche® LightCycler® 480 (96-well block) |
| RT ² Profiler PCR Array, Format G | Roche LightCycler 480 (384-well block) |
| RT ² Profiler PCR Array, Format H | Fluidigm® BioMark™ |



Sample & Assay Technologies

Description

The Human Induced Pluripotent Stem Cells RT² Profiler PCR Array profiles the expression of 84 key genes involved in induced pluripotent stem cell (iPSC) research. iPSCs hold the promise to provide treatments for a multitude of diseases by converting adult somatic cells into pluripotent cells that are able to differentiate into any one of a variety of cell types, avoiding the ethics of embryonic stem cell (ESC) use. The process starts by transfecting or transducing somatic cells (such as keratinocytes) with constructs ectopically expressing a combination of specific transcription factors (KLF4, MYC, POU5F1, and/or SOX2). These transcription factors reprogram or induce the somatic cells to "dedifferentiate", losing markers of the original cell type and gaining markers of pluripotent cells. Often, a combination of additional transcription factors (such as ESRRB, LIN28A, NANOG, MYCN, and/or NR5A2) increases induction efficiency. To control the procedure, the expression of multiple gene classes included on this array must be monitored simultaneously: representative parental cell line genes, the ectopically expressed transcription factors, markers of iPSCs, and markers of the redifferentiation into ectoderm, endoderm, and mesoderm. ESCs and iPSCs have proven to not be functionally identical. Therefore, this array also analyzes genes highly expressed in both cell types to help distinguish them and better understand their differences. Because the expression of typical housekeeping or reference genes often proves inconsistent in these types of studies, the array includes another gene (NAT1) used in iPSC gene expression for data normalization if needed. A set of controls present on each array enables data analysis using the $\Delta\Delta CT$ method of relative quantification and assessment of reverse transcription performance, genomic DNA contamination, and PCR performance. Using real-time PCR, research studies can easily and reliably analyze the expression of a focused panel of genes involved in the induced pluripotent stem cell dedifferentiation and redifferentiation processes with this array.

For further details, consult the *RT² Profiler PCR Array Handbook*.

Shipping and storage

RT² Profiler PCR Arrays in formats A, C, D, E, F, and G are shipped at ambient temperature, on dry ice, or blue ice packs depending on destination and accompanying products. RT² Profiler PCR Arrays in format H are shipped on dry ice or blue ice packs.

For long term storage, keep plates at -20°C.

Note: Ensure that you have the correct RT² Profiler PCR Array format for your real-time cycler (see table above).

Note: Open the package and store the products appropriately immediately on receipt.

Array layout (96-well)

For 384-well 4 x 96 PCR arrays, genes are present in a staggered format. Refer to the *RT² Profiler PCR Array Handbook* for layout.

| | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
|---|--------|--------|---------|-------|--------|--------|--------|--------|--------|-------|-------|-------|
| A | ACTC1 | AICDA | ALDH1A1 | ALDH2 | ALPL | APC | BGLAP | BMP2 | BRIX1 | CCNA2 | CCNE1 | CD34 |
| B | CD9 | CDC42 | CDH1 | CDH2 | CDK1 | COL1A1 | COL2A1 | COL9A1 | DNMT3B | DPPA2 | DPPA3 | EMX2 |
| C | EP300 | ESRRB | FABP7 | FGF2 | FGF4 | FGFR1 | FOXA2 | FOXD3 | GABRB3 | GATA2 | GATA4 | GDF3 |
| D | GJA1 | GJB2 | GRB7 | HAND1 | HDAC2 | HNF4A | HSPA9 | KAT2A | KAT7 | KAT8 | KLF4 | KRT15 |
| E | LEFTY1 | LEFTY2 | LIN28A | MESP1 | MYBL2 | MYC | MYCN | NANOG | NAT1 | NCAM1 | NES | NODAL |
| F | NR5A2 | NUMB | OLIG2 | OTX2 | PARD6A | PAX6 | PECAM1 | PODXL | POU5F1 | REST | RUNX1 | RUNX2 |
| G | SOX15 | SOX17 | SOX2 | SYCP3 | TBX3 | TCF3 | TDGF1 | TERT | TP53 | TUBB3 | UTF1 | ZFP42 |
| H | ACTB | B2M | GAPDH | HPRT1 | RPPL0 | HGDC | RTC | RTC | PPC | PPC | PPC | PPC |

Gene table: RT² Profiler PCR Array

| Position | UniGene | GenBank | Symbol | Description |
|----------|-----------|-----------|---------|--|
| A01 | Hs.118127 | NM_005159 | ACTC1 | Actin, alpha, cardiac muscle 1 |
| A02 | Hs.149342 | NM_020661 | AICDA | Activation-induced cytidine deaminase |
| A03 | Hs.76392 | NM_000689 | ALDH1A1 | Aldehyde dehydrogenase 1 family, member A1 |
| A04 | Hs.604551 | NM_000690 | ALDH2 | Aldehyde dehydrogenase 2 family (mitochondrial) |
| A05 | Hs.75431 | NM_000478 | ALPL | Alkaline phosphatase, liver/bone/kidney |
| A06 | Hs.158932 | NM_000038 | APC | Adenomatous polyposis coli |
| A07 | Hs.654541 | NM_199173 | BGLAP | Bone gamma-carboxyglutamate (gla) protein |
| A08 | Hs.73853 | NM_001200 | BMP2 | Bone morphogenetic protein 2 |
| A09 | Hs.718510 | NM_018321 | BRIX1 | BRX1, biogenesis of ribosomes, homolog (S. cerevisiae) |
| A10 | Hs.58974 | NM_001237 | CCNA2 | Cyclin A2 |
| A11 | Hs.244723 | NM_001238 | CCNE1 | Cyclin E1 |
| A12 | Hs.374990 | NM_001773 | CD34 | CD34 molecule |
| B01 | Hs.114286 | NM_001769 | CD9 | CD9 molecule |
| B02 | Hs.467637 | NM_001791 | CDC42 | Cell division cycle 42 (GTP binding protein, 25kDa) |
| B03 | Hs.461086 | NM_004360 | CDH1 | Cadherin 1, type 1, E-cadherin (epithelial) |
| B04 | Hs.464829 | NM_001792 | CDH2 | Cadherin 2, type 1, N-cadherin (neuronal) |
| B05 | Hs.732435 | NM_001786 | CDK1 | Cyclin-dependent kinase 1 |
| B06 | Hs.172928 | NM_000088 | COL1A1 | Collagen, type I, alpha 1 |
| B07 | Hs.408182 | NM_001844 | COL2A1 | Collagen, type II, alpha 1 |
| B08 | Hs.590892 | NM_001851 | COL9A1 | Collagen, type IX, alpha 1 |
| B09 | Hs.643024 | NM_006892 | DNMT3B | DNA (cytosine-5)-methyltransferase 3 beta |
| B10 | Hs.351113 | NM_138815 | DPPA2 | Developmental pluripotency associated 2 |
| B11 | Hs.131358 | NM_199286 | DPPA3 | Developmental pluripotency associated 3 |
| B12 | Hs.202095 | NM_004098 | EMX2 | Empty spiracles homeobox 2 |
| C01 | Hs.517517 | NM_001429 | EP300 | E1A binding protein p300 |
| C02 | Hs.435845 | NM_004452 | ESRRB | Estrogen-related receptor beta |
| C03 | Hs.26770 | NM_001446 | FABP7 | Fatty acid binding protein 7, brain |
| C04 | Hs.284244 | NM_002006 | FGF2 | Fibroblast growth factor 2 (basic) |
| C05 | Hs.1755 | NM_002007 | FGF4 | Fibroblast growth factor 4 |
| C06 | Hs.264887 | NM_015850 | FGFR1 | Fibroblast growth factor receptor 1 |
| C07 | Hs.155651 | NM_021784 | FOXA2 | Forkhead box A2 |
| C08 | Hs.546573 | NM_012183 | FOXD3 | Forkhead box D3 |
| C09 | Hs.302352 | NM_000814 | GABRB3 | Gamma-aminobutyric acid (GABA) A receptor, beta 3 |
| C10 | Hs.367725 | NM_032638 | GATA2 | GATA binding protein 2 |
| C11 | Hs.243987 | NM_002052 | GATA4 | GATA binding protein 4 |
| C12 | Hs.86232 | NM_020634 | GDF3 | Growth differentiation factor 3 |
| D01 | Hs.700699 | NM_000165 | GJA1 | Gap junction protein, alpha 1, 43kDa |
| D02 | Hs.524894 | NM_004004 | GJB2 | Gap junction protein, beta 2, 26kDa |
| D03 | Hs.86859 | NM_005310 | GRB7 | Growth factor receptor-bound protein 7 |
| D04 | Hs.152531 | NM_004821 | HAND1 | Heart and neural crest derivatives expressed 1 |
| D05 | Hs.3352 | NM_001527 | HDAC2 | Histone deacetylase 2 |
| D06 | Hs.116462 | NM_178849 | HNF4A | Hepatocyte nuclear factor 4, alpha |
| D07 | Hs.184233 | NM_004134 | HSPA9 | Heat shock 70kDa protein 9 (mortalin) |
| D08 | Hs.463045 | NM_021078 | KAT2A | K(lysine) acetyltransferase 2A |
| D09 | Hs.21907 | NM_007067 | KAT7 | K(lysine) acetyltransferase 7 |

| Position | UniGene | GenBank | Symbol | Description |
|-----------------|----------------|----------------|---------------|--|
| D10 | Hs.533803 | NM_032188 | KAT8 | K(lysine) acetyltransferase 8 |
| D11 | Hs.376206 | NM_004235 | KLF4 | Kruppel-like factor 4 (gut) |
| D12 | Hs.654570 | NM_002275 | KRT15 | Keratin 15 |
| E01 | Hs.656214 | NM_020997 | LEFTY1 | Left-right determination factor 1 |
| E02 | Hs.520187 | NM_003240 | LEFTY2 | Left-right determination factor 2 |
| E03 | Hs.86154 | NM_024674 | LIN28A | Lin-28 homolog A (C. elegans) |
| E04 | Hs.447531 | NM_018670 | MESP1 | Mesoderm posterior 1 homolog (mouse) |
| E05 | Hs.179718 | NM_002466 | MYBL2 | V-myb myeloblastosis viral oncogene homolog (avian)-like 2 |
| E06 | Hs.202453 | NM_002467 | MYC | V-myc myelocytomatosis viral oncogene homolog (avian) |
| E07 | Hs.25960 | NM_005378 | MYCN | V-myc myelocytomatosis viral related oncogene, neuroblastoma derived (avian) |
| E08 | Hs.635882 | NM_024865 | NANOG | Nanog homeobox |
| E09 | Hs.591847 | NM_000662 | NAT1 | N-acetyltransferase 1 (arylamine N-acetyltransferase) |
| E10 | Hs.503878 | NM_000615 | NCAM1 | Neural cell adhesion molecule 1 |
| E11 | Hs.527971 | NM_006617 | NES | Nestin |
| E12 | Hs.370414 | NM_018055 | NODAL | Nodal homolog (mouse) |
| F01 | Hs.33446 | NM_003822 | NR5A2 | Nuclear receptor subfamily 5, group A, member 2 |
| F02 | Hs.654609 | NM_003744 | NUMB | Numb homolog (Drosophila) |
| F03 | Hs.176977 | NM_005806 | OLIG2 | Oligodendrocyte lineage transcription factor 2 |
| F04 | Hs.288655 | NM_021728 | OTX2 | Orthodontic homeobox 2 |
| F05 | Hs.112933 | NM_016948 | PARD6A | Par-6 partitioning defective 6 homolog alpha (C. elegans) |
| F06 | Hs.270303 | NM_000280 | PAX6 | Paired box 6 |
| F07 | Hs.514412 | NM_000442 | PECAM1 | Platelet/endothelial cell adhesion molecule |
| F08 | Hs.744213 | NM_005397 | PODXL | Podocalyxin-like |
| F09 | Hs.249184 | NM_002701 | POU5F1 | POU class 5 homeobox 1 |
| F10 | Hs.307836 | NM_005612 | REST | RE1-silencing transcription factor |
| F11 | Hs.149261 | NM_001754 | RUNX1 | Runt-related transcription factor 1 |
| F12 | Hs.535845 | NM_004348 | RUNX2 | Runt-related transcription factor 2 |
| G01 | Hs.95582 | NM_006942 | SOX15 | SRY (sex determining region Y)-box 15 |
| G02 | Hs.98367 | NM_022454 | SOX17 | SRY (sex determining region Y)-box 17 |
| G03 | Hs.518438 | NM_003106 | SOX2 | SRY (sex determining region Y)-box 2 |
| G04 | Hs.506504 | NM_153694 | SYCP3 | Synaptonemal complex protein 3 |
| G05 | Hs.744016 | NM_016569 | TBX3 | T-box 3 |
| G06 | Hs.371282 | NM_003200 | TCF3 | Transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47) |
| G07 | Hs.385870 | NM_003212 | TDGF1 | Teratocarcinoma-derived growth factor 1 |
| G08 | Hs.492203 | NM_198253 | TERT | Telomerase reverse transcriptase |
| G09 | Hs.437460 | NM_000546 | TP53 | Tumor protein p53 |
| G10 | Hs.511743 | NM_006086 | TUBB3 | Tubulin, beta 3 |
| G11 | Hs.458406 | NM_003577 | UTF1 | Undifferentiated embryonic cell transcription factor 1 |
| G12 | Hs.335787 | NM_174900 | ZFP42 | Zinc finger protein 42 homolog (mouse) |
| H01 | Hs.520640 | NM_001101 | ACTB | Actin, beta |
| H02 | Hs.534255 | NM_004048 | B2M | Beta-2-microglobulin |
| H03 | Hs.592355 | NM_002046 | GAPDH | Glyceraldehyde-3-phosphate dehydrogenase |
| H04 | Hs.412707 | NM_000194 | HPRT1 | Hypoxanthine phosphoribosyltransferase 1 |
| H05 | Hs.546285 | NM_001002 | RPLP0 | Ribosomal protein, large, P0 |
| H06 | N/A | SA_00105 | HGDC | Human Genomic DNA Contamination |
| H07 | N/A | SA_00104 | RTC | Reverse Transcription Control |
| H08 | N/A | SA_00104 | RTC | Reverse Transcription Control |
| H09 | N/A | SA_00104 | RTC | Reverse Transcription Control |
| H10 | N/A | SA_00103 | PPC | Positive PCR Control |
| H11 | N/A | SA_00103 | PPC | Positive PCR Control |
| H12 | N/A | SA_00103 | PPC | Positive PCR Control |

Related products

For optimal performance, RT² Profiler PCR Arrays should be used together with the RT² First Strand Kit for cDNA synthesis and RT2 SYBR® Green qPCR Mastermixes for PCR.

| Product | Contents | Cat. no. |
|--|---|----------|
| RT ² First Strand Kit (12) | Enzymes and reagents for cDNA synthesis | 330401 |
| RT ² SYBR Green qPCR Mastermix (2)* | For 2 x 96 assays in 96-well plates; suitable for use with real-time cyclers that do not require a reference dye, including: Bio-Rad models CFX96, CFX384, DNA Engine Opticon 2; Bio-Rad/MJ Research Chromo4; Roche LightCycler 480 (96-well and 384-well); all other cyclers | 330500 |
| RT ² SYBR Green ROX™ qPCR Mastermix (2)* | For 2 x 96 assays in 96-well plates; suitable for use with the following real-time cyclers: Applied Biosystems models 5700, 7000, 7300, 7500 [Standard and FAST], 7700, 7900HT 96-well block [Standard and FAST] and 384-well block, StepOnePlus; Eppendorf Mastercycler ep realplex models 2, 2S, 4, 4S; Stratagene models Mx3000P, Mx3005P, Mx4000; Takara TP-800 | 330520 |
| RT ² SYBR Green Fluor qPCR Mastermix (2)* | For 2 x 96 assays in 96-well plates; suitable for use with the following real-time cyclers: Bio-Rad models iCycler, iQ5, MyiQ, MyiQ2 | 330510 |

* Larger kit sizes available; please inquire.

RT² Profiler PCR Array products are intended for molecular biology applications. These products are not intended for the diagnosis, prevention, or treatment of a disease.

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