

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

Human Drug Metabolism: Phase II Enzymes

Cat. no. 330231 PAHS-069ZA

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 Drug Metabolism Phase II Enzymes RT² Profiler PCR Array contains 84 genes involved in the enzymatic processes of drug biotransformation. Phase II drug metabolism enzymes catalyze the conjugation of lipophilic compounds with hydrophilic functional groups or moieties to form water-soluble conjugates that can be cleared from cells and from the body. This array represents genes encoding the seven major classes of phase II drug metabolism enzymes catalyzing such reactions as glutathione conjugation, glucuronidation, sulfation, methylation, amino acid conjugation, epoxidation, and esterification. Using real-time PCR, you can easily and reliably analyze expression of a focused panel of drug metabolism enzymes 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 | AANAT | ACSL1 | ACSL3 | ACSL4 | ACSM1 | ACSM2B | ACSM3 | AGXT | AS3MT | ASMT | BAAT | CCBL1 |
| B | CES1 | CES2 | CES3 | CES5A | CHST7 | COMT | DDOST | EPHX1 | EPHX2 | GALNT1 | GALNT4 | GAMT |
| C | GCNT1 | GLYAT | GNMT | GSTA1 | GSTA3 | GSTA4 | GSTA5 | GSTK1 | GSTM2 | GSTM3 | GSTM4 | GSTM5 |
| D | GSTO1 | GSTO2 | GSTP1 | GSTT1 | HNMT | INMT | MGAT1 | MGAT2 | MGST1 | MGST2 | MGST3 | NAA20 |
| E | NAT1 | NAT2 | NNMT | NQO1 | NQO2 | PNMT | POMGNT1 | PTGES | SAT1 | SULT1A1 | SULT1A2 | SULT1B1 |
| F | SULT1C2 | SULT1C3 | SULT1C4 | SULT1E1 | SULT2A1 | SULT2B1 | SULT4A1 | SULT6B1 | TPMT | TST | UGCG | UGT1A1 |
| G | UGT1A4 | UGT1A9 | UGT2A1 | UGT2A3 | UGT2B10 | UGT2B17 | UGT2B28 | UGT2B4 | UGT2B7 | UGT3A1 | UGT8 | XDH |
| H | ACTB | B2M | GAPDH | HPRT1 | RPLPO | HGDC | RTC | RTC | PPC | PPC | PPC | PPC |

Gene table: RT² Profiler PCR Array

| Position | UniGene | GenBank | Symbol | Description |
|----------|-----------|-----------|--------|--|
| A01 | Hs.431417 | NM_001088 | AANAT | Aralkylamine N-acetyltransferase |
| A02 | Hs.406678 | NM_001995 | ACSL1 | Acyl-CoA synthetase long-chain family member 1 |
| A03 | Hs.655772 | NM_004457 | ACSL3 | Acyl-CoA synthetase long-chain family member 3 |
| A04 | Hs.268785 | NM_004458 | ACSL4 | Acyl-CoA synthetase long-chain family member 4 |
| A05 | Hs.306812 | NM_052956 | ACSM1 | Acyl-CoA synthetase medium-chain family member 1 |
| A06 | Hs.567879 | NM_182617 | ACSM2B | Acyl-CoA synthetase medium-chain family member 2B |
| A07 | Hs.706754 | NM_005622 | ACSM3 | Acyl-CoA synthetase medium-chain family member 3 |
| A08 | Hs.144567 | NM_000030 | AGXT | Alanine-glyoxylate aminotransferase |
| A09 | Hs.123461 | NM_020682 | AS3MT | Arsenic (+3 oxidation state) methyltransferase |
| A10 | Hs.522572 | NM_004043 | ASMT | Acetylserotonin O-methyltransferase |
| A11 | Hs.284712 | NM_001701 | BAAT | Bile acid CoA: amino acid N-acyltransferase (glycine N-choloyltransferase) |
| A12 | Hs.495250 | NM_004059 | CCBL1 | Cysteine conjugate-beta lyase, cytoplasmic |
| B01 | Hs.558865 | NM_001266 | CES1 | Carboxylesterase 1 |
| B02 | Hs.282975 | NM_198061 | CES2 | Carboxylesterase 2 |
| B03 | Hs.268700 | NM_024922 | CES3 | Carboxylesterase 3 |
| B04 | Hs.350800 | NM_145024 | CES5A | Carboxylesterase 5A |
| B05 | Hs.138155 | NM_019886 | CHST7 | Carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 7 |
| B06 | Hs.370408 | NM_000754 | COMT | Catechol-O-methyltransferase |
| B07 | Hs.523145 | NM_005216 | DDOST | Dolichyl-diphosphooligosaccharide--protein glycosyltransferase |
| B08 | Hs.89649 | NM_000120 | EPHX1 | Epoxide hydrolase 1, microsomal (xenobiotic) |
| B09 | Hs.212088 | NM_001979 | EPHX2 | Epoxide hydrolase 2, cytoplasmic |
| B10 | Hs.514806 | NM_020474 | GALNT1 | UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 1 (GalNAc-T1) |
| B11 | Hs.25130 | NM_003774 | GALNT4 | UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 4 (GalNAc-T4) |
| B12 | Hs.81131 | NM_000156 | GAMT | Guanidinoacetate N-methyltransferase |
| C01 | Hs.521568 | NM_001490 | GCNT1 | Glucosaminyl (N-acetyl) transferase 1, core 2 |
| C02 | Hs.145384 | NM_005838 | GLYAT | Glycine-N-acyltransferase |
| C03 | Hs.144914 | NM_018960 | GNMT | Glycine N-methyltransferase |
| C04 | Hs.446309 | NM_145740 | GSTA1 | Glutathione S-transferase alpha 1 |
| C05 | Hs.102484 | NM_000847 | GSTA3 | Glutathione S-transferase alpha 3 |
| C06 | Hs.485557 | NM_001512 | GSTA4 | Glutathione S-transferase alpha 4 |
| C07 | Hs.646984 | NM_153699 | GSTA5 | Glutathione S-transferase alpha 5 |
| C08 | Hs.390667 | NM_015917 | GSTK1 | Glutathione S-transferase kappa 1 |
| C09 | Hs.279837 | NM_000848 | GSTM2 | Glutathione S-transferase mu 2 (muscle) |
| C10 | Hs.2006 | NM_000849 | GSTM3 | Glutathione S-transferase mu 3 (brain) |
| C11 | Hs.348387 | NM_000850 | GSTM4 | Glutathione S-transferase mu 4 |
| C12 | Hs.75652 | NM_000851 | GSTM5 | Glutathione S-transferase mu 5 |
| D01 | Hs.190028 | NM_004832 | GSTO1 | Glutathione S-transferase omega 1 |
| D02 | Hs.203634 | NM_183239 | GSTO2 | Glutathione S-transferase omega 2 |
| D03 | Hs.523836 | NM_000852 | GSTP1 | Glutathione S-transferase pi 1 |
| D04 | Hs.268573 | NM_000853 | GSTT1 | Glutathione S-transferase theta 1 |
| D05 | Hs.42151 | NM_006895 | HNMT | Histamine N-methyltransferase |
| D06 | Hs.632629 | NM_006774 | INMT | Indolethylamine N-methyltransferase |
| D07 | Hs.519818 | NM_002406 | MGAT1 | Mannosyl (alpha-1,3-)glycoprotein beta-1,2-N-acetylglucosaminyltransferase |

| Position | UniGene | GenBank | Symbol | Description |
|-----------------|----------------|----------------|---------------|--|
| D08 | Hs.93338 | NM_002408 | MGAT2 | Mannosyl (alpha-1,6-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase |
| D09 | Hs.389700 | NM_020300 | MGST1 | Microsomal glutathione S-transferase 1 |
| D10 | Hs.81874 | NM_002413 | MGST2 | Microsomal glutathione S-transferase 2 |
| D11 | Hs.191734 | NM_004528 | MGST3 | Microsomal glutathione S-transferase 3 |
| D12 | Hs.368783 | NM_016100 | NAA20 | N(alpha)-acetyltransferase 20, NatB catalytic subunit |
| E01 | Hs.591847 | NM_000662 | NAT1 | N-acetyltransferase 1 (arylamine N-acetyltransferase) |
| E02 | Hs.2 | NM_000015 | NAT2 | N-acetyltransferase 2 (arylamine N-acetyltransferase) |
| E03 | Hs.503911 | NM_006169 | NNMT | Nicotinamide N-methyltransferase |
| E04 | Hs.406515 | NM_000903 | NQO1 | NAD(P)H dehydrogenase, quinone 1 |
| E05 | Hs.533050 | NM_000904 | NQO2 | NAD(P)H dehydrogenase, quinone 2 |
| E06 | Hs.1892 | NM_002686 | PNMT | Phenylethanolamine N-methyltransferase |
| E07 | Hs.525134 | NM_017739 | POMGNT1 | Protein O-linked mannose beta1,2-N-acetylglucosaminyltransferase |
| E08 | Hs.146688 | NM_004878 | PTGES | Prostaglandin E synthase |
| E09 | Hs.28491 | NM_002970 | SAT1 | Spermidine/spermine N1-acetyltransferase 1 |
| E10 | Hs.567342 | NM_001055 | SULT1A1 | Sulfotransferase family, cytosolic, 1A, phenol-preferring, member 1 |
| E11 | Hs.546304 | NM_177528 | SULT1A2 | Sulfotransferase family, cytosolic, 1A, phenol-preferring, member 2 |
| E12 | Hs.129742 | NM_014465 | SULT1B1 | Sulfotransferase family, cytosolic, 1B, member 1 |
| F01 | Hs.436123 | NM_001056 | SULT1C2 | Sulfotransferase family, cytosolic, 1C, member 2 |
| F02 | Hs.535156 | NM_001008743 | SULT1C3 | Sulfotransferase family, cytosolic, 1C, member 3 |
| F03 | Hs.312644 | NM_006588 | SULT1C4 | Sulfotransferase family, cytosolic, 1C, member 4 |
| F04 | Hs.479898 | NM_005420 | SULT1E1 | Sulfotransferase family 1E, estrogen-preferring, member 1 |
| F05 | Hs.515835 | NM_003167 | SULT2A1 | Sulfotransferase family, cytosolic, 2A, dehydroepiandrosterone (DHEA)-preferring, member 1 |
| F06 | Hs.369331 | NM_004605 | SULT2B1 | Sulfotransferase family, cytosolic, 2B, member 1 |
| F07 | Hs.189810 | NM_014351 | SULT4A1 | Sulfotransferase family 4A, member 1 |
| F08 | Hs.631892 | NM_001032377 | SULT6B1 | Sulfotransferase family, cytosolic, 6B, member 1 |
| F09 | Hs.444319 | NM_000367 | TPMT | Thiopurine S-methyltransferase |
| F10 | Hs.474783 | NM_003312 | TST | Thiosulfate sulfurtransferase (rhodanese) |
| F11 | Hs.304249 | NM_003358 | UGCG | UDP-glucose ceramide glucosyltransferase |
| F12 | Hs.554822 | NM_000463 | UGT1A1 | UDP glucuronosyltransferase 1 family, polypeptide A1 |
| G01 | Hs.554822 | NM_007120 | UGT1A4 | UDP glucuronosyltransferase 1 family, polypeptide A4 |
| G02 | Hs.554822 | NM_021027 | UGT1A9 | UDP glucuronosyltransferase 1 family, polypeptide A9 |
| G03 | Hs.225950 | NM_006798 | UGT2A1 | UDP glucuronosyltransferase 2 family, polypeptide A1, complex locus |
| G04 | Hs.122583 | NM_024743 | UGT2A3 | UDP glucuronosyltransferase 2 family, polypeptide A3 |
| G05 | Hs.201634 | NM_001075 | UGT2B10 | UDP glucuronosyltransferase 2 family, polypeptide B10 |
| G06 | Hs.575083 | NM_001077 | UGT2B17 | UDP glucuronosyltransferase 2 family, polypeptide B17 |
| G07 | Hs.653154 | NM_053039 | UGT2B28 | UDP glucuronosyltransferase 2 family, polypeptide B28 |
| G08 | Hs.285887 | NM_021139 | UGT2B4 | UDP glucuronosyltransferase 2 family, polypeptide B4 |
| G09 | Hs.654424 | NM_001074 | UGT2B7 | UDP glucuronosyltransferase 2 family, polypeptide B7 |
| G10 | Hs.254699 | NM_152404 | UGT3A1 | UDP glycosyltransferase 3 family, polypeptide A1 |
| G11 | Hs.144197 | NM_003360 | UGT8 | UDP glycosyltransferase 8 |
| G12 | Hs.250 | NM_000379 | XDH | Xanthine dehydrogenase |
| 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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