

# RT<sup>2</sup> Profiler PCR Array (Rotor-Gene® Format)

## Mouse Amino Acid Metabolism II

Cat. no. 330231 PAMM-130ZR

For pathway expression analysis

| Format                                       | For use with the following real-time cyclers |
|--|--|
| RT <sup>2</sup> Profiler PCR Array, Format R | Rotor-Gene Q, other Rotor-Gene cyclers       |

### Description

The Mouse Amino Acid Metabolism II RT<sup>2</sup> Profiler PCR Array profiles the expression of 84 key genes important in amino acid biosynthesis and degradation. Of the 20 amino acids required for protein synthesis, mammals synthesize the non-essential amino acids in vivo and must obtain the other essential amino acids from their diet or intestinal flora. The interrelated metabolism of amino acids involves key signaling molecules, vitamins and cofactors. Slight alterations in the expression of these metabolic genes impose potentially adverse consequences on mammalian metabolism. For example, the metabolism of histidine forms histamine, a metabolite central to allergic reactions and vasodilation. Expression levels of the enzyme involved in this reaction, DDC, may be related to allergic sensitivities in affected individuals. Therefore, analysis of genes involved in the biosynthesis and degradation of amino acids unlocks the potential to enhance our understanding of basic biological pathways as well as nutritional status in patients with metabolic disorders or nutritional deprivation. This array includes genes important for the metabolism of alanine, asparagine, aspartic acid, histidine, isoleucine, lysine, phenylalanine, serine, glycine, threonine, tyrosine, and valine. Using real-time PCR, you can easily and reliably analyze the expression of a focused panel of genes involved in amino acid metabolism with this array.

For further details, consult the *RT<sup>2</sup> Profiler PCR Array Handbook*.

### Shipping and storage

RT<sup>2</sup> Profiler PCR Arrays in the Rotor-Gene format are shipped at ambient temperature, on dry ice, or blue ice packs depending on destination and accompanying products.

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

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**Note:** Ensure that you have the correct RT<sup>2</sup> Profiler PCR Array format for your real-time cycler (see table above).

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



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Sample & Assay Technologies

## Array layout

The 96 real-time assays in the Rotor-Gene format are located in wells 1–96 of the Rotor-Disc™ (plate A1–A12=Rotor-Disc 1–12, plate B1–B12=Rotor-Disc 13–24, etc.). To maintain data analysis compatibility, wells 97–100 do not contain real-time assays but will contain master mix to account for weight balance.

## Gene table: RT<sup>2</sup> Profiler PCR Array

| Position | UniGene   | GenBank      | Symbol  | Description  |
|----------|-----------|--------------|---------|--|
| A01      | Mm.35020  | NM_011834    | Aadat   | Aminoadipate aminotransferase  |
| A02      | Mm.33970  | NM_026276    | Aasdhpt | Aminoadipate-semialdehyde dehydrogenase-phosphopantetheinyl transferase  |
| A03      | Mm.18651  | NM_013930    | Aass    | Aminoadipate-semialdehyde synthase   |
| A04      | Mm.259315 | NM_172961    | Abat    | 4-aminobutyrate aminotransferase   |
| A05      | Mm.213898 | NM_029638    | Abp1    | Amiloride binding protein 1 (amine oxidase, copper-containing)   |
| A06      | Mm.10530  | NM_007382    | Acadm   | Acyl-Coenzyme A dehydrogenase, medium chain  |
| A07      | Mm.18759  | NM_007383    | Acads   | Acyl-Coenzyme A dehydrogenase, short chain   |
| A08      | Mm.334274 | NM_025826    | Acadsb  | Acyl-Coenzyme A dehydrogenase, short/branched chain  |
| A09      | Mm.439711 | NM_009338    | Acat2   | Acetyl-Coenzyme A acetyltransferase 2  |
| A10      | Mm.3874   | NM_007410    | Adh5    | Alcohol dehydrogenase 5 (class III), chi polypeptide   |
| A11      | Mm.38151  | NM_009634    | Adsl    | Adenylosuccinate lyase   |
| A12      | Mm.338021 | NM_007422    | Adss    | Adenylosuccinate synthetase, non muscle  |
| B01      | Mm.7457   | NM_016702    | Agxt    | Alanine-glyoxylate aminotransferase  |
| B02      | Mm.290578 | NM_020559    | Alas1   | Aminolevulinic acid synthase 1   |
| B03      | Mm.284446 | NM_009656    | Aldh2   | Aldehyde dehydrogenase 2, mitochondrial  |
| B04      | Mm.109341 | NM_026316    | Aldh3b1 | Aldehyde dehydrogenase 3 family, member B1   |
| B05      | Mm.393311 | NM_172532    | Aldh5a1 | Aldehyde dehydrogenase family 5, subfamily A1  |
| B06      | Mm.247510 | NM_134042    | Aldh6a1 | Aldehyde dehydrogenase family 6, subfamily A1  |
| B07      | Mm.244518 | NM_027908    | Amdhd1  | Amidohydrolase domain containing 1   |
| B08      | Mm.390225 | NM_001013814 | Amt     | Aminomethyltransferase   |
| B09      | Mm.67281  | NM_009675    | Aoc3    | Amine oxidase, copper containing 3   |
| B10      | Mm.130752 | NM_138679    | Ash1l   | Ash1 (absent, small, or homeotic)-like (Drosophila)  |
| B11      | Mm.2942   | NM_012055    | Asns    | Asparagine synthetase  |
| B12      | Mm.293574 | NM_023113    | Aspa    | Aspartoacylase   |
| C01      | Mm.27335  | NM_130452    | Bbox1   | Butyrobetaine (gamma), 2-oxoglutarate dioxygenase 1 (gamma-butyrobetaine hydroxylase)  |
| C02      | Mm.4606   | NM_007532    | Bcat1   | Branched chain aminotransferase 1, cytosolic   |
| C03      | Mm.25848  | NM_007533    | Bckdha  | Branched chain ketoacid dehydrogenase E1, alpha polypeptide  |
| C04      | Mm.329582 | NM_016668    | Bhmt    | Betaine-homocysteine methyltransferase   |
| C05      | Mm.259916 | NM_172264    | Chdh    | Choline dehydrogenase  |
| C06      | Mm.23278  | NM_177450    | Cndp1   | Carnosine dipeptidase 1 (metallopeptidase M20 family)  |
| C07      | Mm.100940 | NM_007744    | Comt    | Catechol-O-methyltransferase   |
| C08      | Mm.20115  | NM_010018    | Dao     | D-amino acid oxidase   |
| C09      | Mm.167781 | NM_138942    | Dbh     | Dopamine beta hydroxylase  |
| C10      | Mm.3636   | NM_010022    | Dbt     | Dihydrolipoamide branched chain transacylase E2  |
| C11      | Mm.12906  | NM_016672    | Ddc     | Dopa decarboxylase   |
| C12      | Mm.3131   | NM_007861    | Dld     | Dihydrolipoamide dehydrogenase   |
| D01      | Mm.296221 | NM_030225    | Dlst    | Dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-glutarate complex)   |
| D02      | Mm.21789  | NM_028772    | Dmgdh   | Dimethylglycine dehydrogenase precursor  |
| D03      | Mm.24452  | NM_053119    | Echs1   | Enoyl Coenzyme A hydratase, short chain, 1, mitochondrial  |
| D04      | Mm.3798   | NM_010176    | Fah     | Fumarylacetoacetate hydrolase  |
| D05      | Mm.36278  | NM_080845    | Ftcd    | Formiminotransferase cyclodeaminase  |
| D06      | Mm.4784   | NM_008078    | Gad2    | Glutamic acid decarboxylase 2  |
| D07      | Mm.237085 | NM_013847    | Gcat    | Glycine C-acetyltransferase (2-amino-3-ketobutyrate-coenzyme A ligase)   |
| D08      | Mm.2475   | NM_008097    | Gcdh    | Glutaryl-Coenzyme A dehydrogenase  |
| D09      | Mm.274852 | NM_138595    | Gldc    | Glycine decarboxylase  |
| D10      | Mm.29395  | NM_010321    | Gnmt    | Glycine N-methyltransferase  |
| D11      | Mm.19039  | NM_010324    | Got1    | Glutamate oxaloacetate transaminase 1, soluble   |
| D12      | Mm.30130  | NM_182805    | Gpt     | Glutamic pyruvic transaminase, soluble   |
| E01      | Mm.260164 | NM_008212    | Hadh    | Hydroxyacyl-Coenzyme A dehydrogenase   |
| E02      | Mm.291463 | NM_145558    | Hadhb   | Hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit |
| E03      | Mm.18603  | NM_008230    | Hdc     | Histidine decarboxylase  |
| E04      | Mm.157442 | NM_013547    | Hgd     | Homogentisate 1, 2-dioxygenase   |
| E05      | Mm.286458 | NM_145567    | Hibadh  | 3-hydroxyisobutyrate dehydrogenase   |
| E06      | Mm.222063 | NM_146108    | Hibch   | 3-hydroxyisobutyryl-Coenzyme A hydrolase   |

| Position | UniGene   | GenBank   | Symbol   | Description   |
|----------|-----------|-----------|----------|---|
| E07      | Mm.33120  | NM_080462 | Hnmt     | Histamine N-methyltransferase                             |
| E08      | Mm.439709 | NM_008277 | Hpd      | 4-hydroxyphenylpyruvic acid dioxygenase                   |
| E09      | Mm.6994   | NM_016763 | Hsd17b10 | Hydroxysteroid (17-beta) dehydrogenase 10                 |
| E10      | Mm.21118  | NM_172015 | Iars     | Isoleucine-tRNA synthetase                                |
| E11      | Mm.21108  | NM_173740 | Maoa     | Monoamine oxidase A                                       |
| E12      | Mm.10093  | NM_028626 | Mcee     | Methylmalonyl CoA epimerase                               |
| F01      | Mm.2326   | NM_010798 | Mif      | Macrophage migration inhibitory factor                    |
| F02      | Mm.259884 | NM_008650 | Mut      | Methylmalonyl-Coenzyme A mutase                           |
| F03      | Mm.276348 | NM_010956 | Ogdh     | Oxoglutarate dehydrogenase (lipoamide)                    |
| F04      | Mm.263539 | NM_008777 | Pah      | Phenylalanine hydroxylase                                 |
| F05      | Mm.23876  | NM_144844 | Pcca     | Propionyl-Coenzyme A carboxylase, alpha polypeptide       |
| F06      | Mm.4223   | NM_008811 | Pdha2    | Pyruvate dehydrogenase E1 alpha 2                         |
| F07      | Mm.16898  | NM_016966 | Phgdh    | 3-phosphoglycerate dehydrogenase                          |
| F08      | Mm.8543   | NM_008952 | Pipox    | Pipecolic acid oxidase                                    |
| F09      | Mm.251003 | NM_011962 | Plod3    | Procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3        |
| F10      | Mm.57030  | NM_008890 | Pnmt     | Phenylethanolamine-N-methyltransferase                    |
| F11      | Mm.186185 | NM_007453 | Prdx6    | Peroxiredoxin 6   |
| F12      | Mm.289936 | NM_177420 | Psat1    | Phosphoserine aminotransferase 1                          |
| G01      | Mm.271784 | NM_133900 | Psph     | Phosphoserine phosphatase                                 |
| G02      | Mm.278467 | NM_138665 | Sardh    | Sarcosine dehydrogenase                                   |
| G03      | Mm.28685  | NM_145565 | Sds      | Serine dehydratase  |
| G04      | Mm.29890  | NM_028230 | Shmt2    | Serine hydroxymethyltransferase 2 (mitochondrial)         |
| G05      | Mm.131443 | NM_013761 | Srr      | Serine racemase   |
| G06      | Mm.1292   | NM_009377 | Th       | Tyrosine hydroxylase                                      |
| G07      | Mm.394228 | NM_138758 | Tmlhe    | Trimethyllysine hydroxylase, epsilon                      |
| G08      | Mm.4991   | NM_009417 | Tpo      | Thyroid peroxidase  |
| G09      | Mm.238127 | NM_011661 | Tyr      | Tyrosinase  |
| G10      | Mm.30438  | NM_031202 | Tyrp1    | Tyrosinase-related protein 1                              |
| G11      | Mm.32002  | NM_175137 | Vars2    | Valyl-tRNA synthetase 2, mitochondrial (putative)         |
| G12      | Mm.439878 | NM_025375 | Wbscr22  | Williams Beuren syndrome chromosome region 22             |
| H01      | Mm.328431 | NM_007393 | Actb     | Actin, beta   |
| H02      | Mm.163    | NM_009735 | B2m      | Beta-2 microglobulin                                      |
| H03      | Mm.343110 | NM_008084 | Gapdh    | Glyceraldehyde-3-phosphate dehydrogenase                  |
| H04      | Mm.3317   | NM_010368 | Gusb     | Glucuronidase, beta                                       |
| H05      | Mm.2180   | NM_008302 | Hsp90ab1 | Heat shock protein 90 alpha (cytosolic), class B member 1 |
| H06      | N/A       | SA_00106  | MGDC     | Mouse 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, RT<sup>2</sup> Profiler PCR Arrays should be used together with the RT<sup>2</sup> First Strand Kit for cDNA synthesis and RT<sup>2</sup> SYBR<sup>®</sup> Green qPCR Mastermixes for PCR.

| Product   | Contents   | Cat. no. |
|---|--|----------|
| RT <sup>2</sup> First Strand Kit (12)                           | Enzymes and reagents for cDNA synthesis  | 330401   |
| RT <sup>2</sup> SYBR Green ROX <sup>™</sup> FAST Mastermix (2)* | For 2 x 96 assays in 96-well plates; suitable for use with the Rotor-Gene Q and other Rotor-Gene cyclers | 330620   |

\* Larger kit sizes available; please inquire.

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RT<sup>2</sup> Profiler PCR Array products are intended for molecular biology applications. These products are not intended for the diagnosis, prevention, or treatment of a disease.

For up-to-date licensing information and product-specific disclaimers, see the respective QIAGEN kit handbook or user manual. QIAGEN kit handbooks and user manuals are available at [www.qiagen.com](http://www.qiagen.com) or can be requested from QIAGEN Technical Services or your local distributor.

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