

RT² Profiler PCR Array (Rotor-Gene[®] Format)

Rhesus Macaque Ubiquitination (Ubiquitylation) Pathway

Cat. no. 330231 PAQQ-079ZR

For pathway expression analysis

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

Description

The Rhesus Macaque Ubiquitination or Ubiquitylation Pathway RT² Profiler PCR Array profiles the expression of 84 key genes involved in the regulated degradation of cellular proteins by the ubiquitin-proteasome system. Disruption of the proteasomal degradation pathway has been implicated in a wide range of human diseases, such as cancer, diabetes and cardiovascular and neurodegenerative diseases. The array includes ubiquitin-activating enzymes (E1), ubiquitin-conjugating enzymes (E2), and ubiquitin ligases (E3). These enzymes have also been sub-categorized as regulators of key cellular processes, such as apoptosis, the cell cycle and transcription. A set of controls present on each array enables data analysis using the $\Delta\Delta\text{CT}$ 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 expression of a focused panel of genes involved in ubiquitin degradation pathway with this array.

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

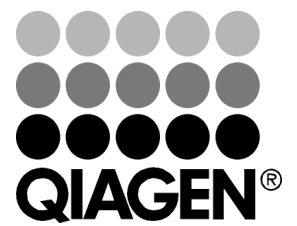
Shipping and storage

RT² 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 .

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.



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: RT² Profiler PCR Array

| Position | UniGene | GenBank | Symbol | Description |
|----------|-----------|--------------|--------------|--|
| A01 | Mmu.1688 | XM_001112301 | ANAPC11 | Anaphase-promoting complex subunit 11-like |
| A02 | Mmu.19914 | XM_001090195 | ANAPC2 | Anaphase promoting complex subunit 2 |
| A03 | Mmu.21936 | XM_001087688 | ATG7 | ATG7 autophagy related 7 homolog (<i>S. cerevisiae</i>) |
| A04 | Mmu.26908 | XM_001084740 | BARD1 | BRCA1 associated RING domain 1 |
| A05 | Mmu.17471 | NM_001114949 | BRCA1 | Breast cancer 1, early onset |
| A06 | Mmu.21786 | XM_001097957 | BRCC3 | BRCA1/BRCA2-containing complex, subunit 3 |
| A07 | Mmu.28657 | XM_002805789 | BTRC | Beta-transducin repeat containing E3 ubiquitin protein ligase |
| A08 | Mmu.30036 | XM_001104812 | CBL | Cas-Br-M (murine) ecotropic retroviral transforming sequence |
| A09 | Mmu.9717 | XM_002803512 | CUL1 | Cullin 1 |
| A10 | Mmu.22177 | XM_001087735 | CUL2 | Cullin 2 |
| A11 | Mmu.22552 | XM_001109745 | CUL3 | Cullin 3 |
| A12 | Mmu.3258 | XM_001086195 | CUL4B | Cullin 4B |
| B01 | Mmu.26452 | XM_001103587 | CUL5 | Cullin 5 |
| B02 | Mmu.23329 | XM_001088003 | CUL7 | Cullin 7 |
| B03 | Mmu.21140 | XM_001082958 | DDB1 | Damage-specific DNA binding protein 1, 127kDa |
| B04 | Mmu.23533 | XM_002808316 | DZIP3 | E3 ubiquitin-protein ligase DZIP3-like |
| B05 | Mmu.28476 | XM_001082360 | FBXO3 | F-box protein 3 |
| B06 | Mmu.20609 | XM_001086503 | FBXO31 | F-box protein 31 |
| B07 | Mmu.2644 | XM_001088308 | FBXO4 | F-box protein 4 |
| B08 | Mmu.26121 | XM_001084872 | FBXW10 | F-box and WD repeat domain containing 10 |
| B09 | Mmu.25629 | XM_001108547 | FBXW9 | F-box and WD repeat domain containing 9 |
| B10 | Mmu.32347 | XM_001096775 | HECW1 | HECT, C2 and WW domain containing E3 ubiquitin protein ligase 1 |
| B11 | Mmu.11124 | XM_001088987 | HUWE1 | HECT, UBA and WWE domain containing 1 |
| B12 | Mmu.28561 | XM_002800991 | LOC100425016 | Ubiquitin-conjugating enzyme E2 R1-like |
| C01 | Mmu.9813 | XM_002804538 | LOC694667 | Ubiquitin-conjugating enzyme E2 D2-like |
| C02 | Mmu.29784 | XM_002808017 | LOC697155 | E3 ubiquitin-protein ligase HECW2-like |
| C03 | Mmu.28952 | XM_002808385 | LOC698984 | Cullin-9-like |
| C04 | Mmu.922 | XM_001089338 | LOC701036 | Ubiquitin-conjugating enzyme E2 E1-like |
| C05 | N/A | XM_001089803 | LOC701507 | Ubiquitin-conjugating enzyme E2 R2-like |
| C06 | Mmu.24263 | XM_001090152 | LOC701870 | Von Hippel-Lindau disease tumor suppressor-like |
| C07 | Mmu.19744 | XM_001091222 | LOC702912 | Protein ariadne-1 homolog |
| C08 | N/A | XM_001094634 | LOC703550 | S-phase kinase-associated protein 1-like |
| C09 | Mmu.23669 | XM_001099588 | LOC710899 | E3 ubiquitin-protein ligase parkin-like |
| C10 | Mmu.446 | XM_002808152 | LOC714302 | Cullin-4A-like |
| C11 | Mmu.26182 | XM_001117702 | LOC721528 | Ubiquitin-conjugating enzyme E2 G1-like |
| C12 | Mmu.11534 | XM_001089783 | MARCH5 | Membrane-associated ring finger (C3HC4) 5 |
| D01 | Mmu.31140 | NM_001266402 | MDM2 | Mdm2 p53 binding protein homolog (mouse) |
| D02 | Mmu.4788 | XM_001092086 | MIB1 | Mindbomb homolog 1 (<i>Drosophila</i>) |
| D03 | Mmu.27453 | XM_001093903 | MOCS3 | Molybdenum cofactor synthesis 3 |
| D04 | Mmu.22231 | XM_001096167 | MUL1 | Mitochondrial E3 ubiquitin protein ligase 1 |
| D05 | Mmu.23495 | XM_001085059 | NAE1 | NEDD8 activating enzyme E1 subunit 1 |
| D06 | N/A | XM_001098135 | NEDD8 | Neural precursor cell expressed, developmentally down-regulated 8 |
| D07 | Mmu.22117 | XM_001104796 | RFWD2 | E3 ubiquitin-protein ligase RFWD2-like |
| D08 | Mmu.24222 | XM_001106313 | RNF123 | Ring finger protein 123 |
| D09 | Mmu.19873 | XM_001084777 | RNF148 | Ring finger protein 148 |
| D10 | Mmu.672 | XM_001109956 | SAE1 | SUMO1 activating enzyme subunit 1 |
| D11 | Mmu.22434 | XM_001093834 | SKP2 | S-phase kinase-associated protein 2 (p45) |
| D12 | Mmu.32101 | NM_001257631 | SMURF1 | SMAD specific E3 ubiquitin protein ligase 1 |
| E01 | Mmu.25983 | XM_001109913 | SMURF2 | SMAD specific E3 ubiquitin protein ligase 2 |
| E02 | Mmu.10137 | XM_001086501 | STUB1 | STIP1 homology and U-box containing protein 1, E3 ubiquitin protein ligase |
| E03 | Mmu.19284 | XM_001114254 | SYVN1 | Synovial apoptosis inhibitor 1, synoviolin |
| E04 | Mmu.18639 | NM_001194059 | TMEM189 | Transmembrane protein 189 |
| E05 | Mmu.3286 | NM_001047151 | TP53 | Tumor protein p53 |
| E06 | Mmu.2813 | XM_001092372 | UBA1 | Ubiquitin-like modifier activating enzyme 1 |
| E07 | Mmu.13107 | XM_001091974 | UBA2 | Ubiquitin-like modifier activating enzyme 2 |
| E08 | Mmu.32389 | XM_001088171 | UBA3 | Ubiquitin-like modifier activating enzyme 3 |

| Position | UniGene | GenBank | Symbol | Description |
|----------|-----------|--------------|-----------|---|
| E09 | Mmu.28903 | XM_002804103 | UBA6 | Ubiquitin-like modifier activating enzyme 6 |
| E10 | Mmu.20021 | XM_001105981 | UBA7 | Ubiquitin-like modifier activating enzyme 7 |
| E11 | Mmu.839 | XM_001082047 | UBE2A | Ubiquitin-conjugating enzyme E2A |
| E12 | Mmu.2430 | NM_001194433 | UBE2B | Ubiquitin-conjugating enzyme E2B |
| F01 | N/A | XM_001083155 | UBE2C | Ubiquitin-conjugating enzyme E2C |
| F02 | Mmu.890 | XM_001097762 | UBE2D1 | Ubiquitin-conjugating enzyme E2D 1 |
| F03 | Mmu.4221 | XM_001109666 | UBE2D3 | Ubiquitin-conjugating enzyme E2 D3-like |
| F04 | Mmu.18684 | XM_001089978 | UBE2E2 | Ubiquitin-conjugating enzyme E2E 2 |
| F05 | Mmu.1876 | XM_001100742 | UBE2E3 | Ubiquitin-conjugating enzyme E2 E3-like |
| F06 | Mmu.13402 | XM_001085096 | UBE2F | Ubiquitin-conjugating enzyme E2F (putative) |
| F07 | Mmu.19159 | XM_001101137 | UBE2G2 | Ubiquitin-conjugating enzyme E2G 2 |
| F08 | Mmu.995 | XM_001093712 | UBE2H | Ubiquitin-conjugating enzyme E2H |
| F09 | Mmu.10576 | XM_001118584 | UBE2I | Ubiquitin-conjugating enzyme E2I |
| F10 | Mmu.1281 | XM_001096217 | UBE2J1 | Ubiquitin-conjugating enzyme E2, J1, U |
| F11 | Mmu.24789 | XM_001092754 | UBE2J2 | Ubiquitin-conjugating enzyme E2 J2-like |
| F12 | Mmu.4081 | XM_001092929 | UBE2K | Ubiquitin-conjugating enzyme E2K |
| G01 | Mmu.1675 | XM_001087183 | UBE2L3 | Ubiquitin-conjugating enzyme E2L 3 |
| G02 | Mmu.19225 | XM_001099932 | UBE2M | Ubiquitin-conjugating enzyme E2M |
| G03 | Mmu.2752 | XM_001111833 | UBE2Q1 | Ubiquitin-conjugating enzyme E2Q family member 1 |
| G04 | Mmu.20815 | XM_001087564 | UBE2S | Ubiquitin-conjugating enzyme E2S |
| G05 | Mmu.21550 | XM_002808256 | UBE2T | Ubiquitin-conjugating enzyme E2 T-like |
| G06 | Mmu.20584 | XM_001085749 | UBE2W | Ubiquitin-conjugating enzyme E2W (putative) |
| G07 | Mmu.26065 | XM_001088335 | UBE2Z | Ubiquitin-conjugating enzyme E2 Z-like |
| G08 | Mmu.12182 | XM_001108510 | UBE3A | Ubiquitin protein ligase E3A |
| G09 | Mmu.24744 | XM_001101836 | UBE4B | Ubiquitination factor E4B |
| G10 | Mmu.2057 | XM_001105721 | UBR1 | Ubiquitin protein ligase E3 component n-recognin 1 |
| G11 | Mmu.20747 | XM_001088074 | UBR2 | Ubiquitin protein ligase E3 component n-recognin 2 |
| G12 | Mmu.1001 | XM_001083173 | WWP1 | WW domain containing E3 ubiquitin protein ligase 1 |
| H01 | Mmu.11089 | NM_001033084 | ACTB | Actin, beta |
| H02 | Mmu.5037 | NM_001047137 | B2M | Beta-2-microglobulin |
| H03 | Mmu.3145 | XM_001105471 | GAPDH | Glyceraldehyde-3-phosphate dehydrogenase |
| H04 | Mmu.12316 | XM_001097691 | LOC709186 | Hypoxanthine-guanine phosphoribosyltransferase-like |
| H05 | Mmu.2512 | XM_001115079 | RPL13A | Ribosomal protein L13A |
| H06 | N/A | SA_00125 | QGDC | Rhesus Macaque 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 ROX™ 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.

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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