RT² Profiler PCR Array (Rotor-Gene® Format) Mouse Glucose Metabolism

Cat. no. 330231 PAMM-006ZR

For pathway expression analysis

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

Description

The Mouse Glucose Metabolism RT² Profiler PCR Array profiles the expression of 84 key genes involved in the regulation and enzymatic pathways of glucose and glycogen metabolism. Glycolysis, the TCA cycle and the pentose phosphate pathways break down glucose from carbohydrates into the metabolites necessary for energy production, and gluconeogenesis stores excess energy as glucose. Cells, particularly in skeletal muscle and the liver, store excess glucose as the polysaccharide glycogen, and quickly metabolize it again when necessary. Changes in glucose metabolism gene expression are common in cancerous tissues. Specifically, tumors often show decreased oxidative phosphorylation, even in the presence of sufficient oxygen, due to enhanced transcription of glycolytic genes and/or reduced transcription of TCA cycle genes. In addition, the pathological consequences of diabetes and obesity involve gene expression changes in glucose metabolic pathways. In one notable example, PCK1 overexpression in mice leads to obesity. Using real-time PCR, you can easily and reliably analyze the expression of a focused panel of genes involved in glucose metabolism 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.



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

A01 A02 A03 A04 A05 A06 A07 A08	Mm.282039 Mm.331547 Mm.154581 Mm.237099	NM_134037 NM_007386 NM_080633	Acly Aco1	ATP citrate lyase	
A03 A04 A05 A06 A07 A08	Mm.154581 Mm.237099	_	Aco1	Aconitase 1	
A03 A04 A05 A06 A07 A08	Mm.154581 Mm.237099	_		Aconitase 1	
A05 A06 A07 A08	Mm.237099	1 14/41 U0U033	Aco2	Aconitase 2, mitochondrial	
A06 A07 A08		NM 001081326	Agl	Amylo-1,6-glucosidase, 4-alpha-glucanotransferase	
A06 A07 A08	Mm.275831	NM 007438	Aldoa	Aldolase A, fructose-bisphosphate	
A07 A08	Mm.482116	NM 144903	Aldob	Aldolase B, fructose-bisphosphate	
A08	Mm.7729	NM 009657	Aldoc	Aldolase C, fructose-bisphosphate	
	Mm.282863	NM 007563	Bpgm	2,3-bisphosphoglycerate mutase	
	Mm.58836	NM_026444	Cs	Citrate synthase	
A10	Mm.285076	NM_145614	Dlat	Dihydrolipoamide S-acetyltransferase (E2 component of pyruvate dehydrogenas complex)	
A11	Mm.3131	NM_007861	Dld	Dihydrolipoamide dehydrogenase	
A12	Mm.296221	NM_030225	Dlst	Dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-glutarate complex)	
B01	Mm.70666	NM_023119	Eno1	Enolase 1, alpha non-neuron	
B02	Mm.3913	NM_013509	Eno2	Enolase 2, gamma neuronal	
B03	Mm.251322	NM_007933	Eno3	Enolase 3, beta muscle	
B04	Mm.423078	NM_019395	Fbp1	Fructose bisphosphatase 1	
B05	Mm.391871	NM_007994	Fbp2	Fructose bisphosphatase 2	
B06	Mm.41502	NM 010209	Fh1	Fumarate hydratase 1	
B07	Mm.18064	NM 008061	G6pc	Glucose-6-phosphatase, catalytic	
B08	Mm.22385	NM 175935	G6pc3	Glucose 6 phosphatase, catalytic, 3	
B09	Mm.27210	NM 008062	G6pdx	Glucose-6-phosphate dehydrogenase X-linked	
B10	Mm.29098	NM 176963	Galm	Galactose mutarotase	
B11	Mm.436562	NM 008085	Gapdhs	Glyceraldehyde-3-phosphate dehydrogenase, spermatogenic	
B12	Mm.396102	NM 028803	Gbe1	Glucan (1,4-alpha-), branching enzyme 1	
C01	Mm.220358	NM 010292	Gck	Glucokinase	
C02	Mm.589	NM 008155	Gpi1	Glucose phosphate isomerase 1	
C03	Mm.476745	NM 001031667	Gsk3a	Glycogen synthase kinase 3 alpha	
C04	Mm.394930	NM 019827	Gsk3b	Glycogen synthase kinase 3 beta	
C05	Mm.275654	NM 030678	Gys1	Glycogen synthase 1, muscle	
C06	Mm.275975	NM 145572	Gys2	Glycogen synthase 2	
C07	Mm.22183	NM 173371	H6pd	Hexose-6-phosphate dehydrogenase (glucose 1-dehydrogenase)	
C08	Mm.255848	NM 013820	Hk2	Hexokinase 2	
C09	Mm.267479	NM 001033245	Hk3	Hexokinase 3	
C10	Mm.9925	NM 010497	ldh1	Isocitrate dehydrogenase 1 (NADP+), soluble	
C11	Mm.246432	NM 173011	ldh2	Isocitrate dehydrogenase 2 (NADP+), mitochondrial	
C12	Mm.279195	NM 029573	Idh3a	Isocitrate dehydrogenase 3 (NAD+) alpha	
D01	Mm.29590	NM 130884	Idh3b	Isocitrate dehydrogenase 3 (NAD+) beta	
D02	Mm.14825	NM 008323	ldh3g	Isocitrate dehydrogenase 3 (NAD+), gamma	
D03	Mm.212703	NM_008618	Mdh1	Malate dehydrogenase 1, NAD (soluble)	
D04	Mm.30494	NM 029696	Mdh1b	Malate dehydrogenase 1B, NAD (soluble)	
D05	Mm.297096	NM 008617	Mdh2	Malate dehydrogenase 2, NAD (mitochondrial)	
D06	Mm.276348	NM 010956	Ogdh	Oxoglutarate dehydrogenase (lipoamide)	
D07	Mm.266867	NM 011044	Pck1	Phosphoenolpyruvate carboxykinase 1, cytosolic	
D08	Mm.29856	NM 028994	Pck2	Phosphoenolpyruvate carboxykinase 2 (mitochondrial)	
D09	Mm.1845	NM 008797	Pcx	Pyruvate carboxylase	
D10	Mm.34775	NM 008810	Pdha1	Pyruvate dehydrogenase E1 alpha 1	
D10	Mm.301527	NM 024221	Pdhb	Pyruvate dehydrogenase (lipoamide) beta	
D12	Mm.34411	NM 172665	Pdk1	Pyruvate dehydrogenase kinase, isoenzyme 1	
E01	Mm.29768	NM_133667	Pdk2	Pyruvate dehydrogenase kinase, isoenzyme 1	
E02	Mm.12775	NM 145630	Pdk3	Pyruvate dehydrogenase kinase, isoenzyme 2	
E03	Mm.235547	NM 013743	Pdk4	Pyruvate dehydrogenase kinase, isoenzyme 4	
	Mm.235347 Mm.290750	NM 001024606	Pdp2	Pyruvate dehyrogenase phosphatase catalytic subunit 2	
F04	Mm.370024	NM 198308	Pdpr	Pyruvate dehydrogenase phosphatase regulatory subunit	
E04		14/41 170000	iubi		
E04 E05 E06	Mm.269649	NM 008826	Pfkl	Phosphofructokinase, liver, B-type	

Position	UniGene	GenBank	Symbol	Description	
E08	Mm.336205	NM_008828	Pgk1	Phosphoglycerate kinase 1	
E09	Mm.717	NM_031190	Pgk2	Phosphoglycerate kinase 2	
E10	Mm.2325	NM_025700	Pgm1	Phosphoglucomutase 1	
E11	Mm.217764	NM_028132	Pgm2	Phosphoglucomutase 2	
E12	Mm.390201	NM_028352	Pgm3	Phosphoglucomutase 3	
F01	Mm.212889	NM_173021	Phka1	Phosphorylase kinase alpha 1	
F02	Mm.237296	NM_199446	Phkb	Phosphorylase kinase beta	
F03	Mm.3159	NM_011079	Phkg1	Phosphorylase kinase gamma 1	
F04	Mm.274473	NM_026888	Phkg2	Phosphorylase kinase, gamma 2 (testis)	
F05	Mm.383180	NM_013631	Pklr	Pyruvate kinase liver and red blood cell	
F06	Mm.287178	NM_021463	Prps1	Phosphoribosyl pyrophosphate synthetase 1	
F07	Mm.79179	NM_029294	Prps111	Phosphoribosyl pyrophosphate synthetase 1-like 1	
F08	Mm.272955	NM_026662	Prps2	Phosphoribosyl pyrophosphate synthetase 2	
F09	Mm.256926	NM_133198	Pygl	Liver glycogen phosphorylase	
F10	Mm.27806	NM_011224	Pygm	Muscle glycogen phosphorylase	
F11	Mm.22519	NM_153196	Rbks	Ribokinase	
F12	Mm.240912	NM_025683	Rpe	Ribulose-5-phosphate-3-epimerase	
G01	Mm.17905	NM_009075	Rpia	Ribose 5-phosphate isomerase A	
G02	Mm.158231	NM_023281	Sdha	Succinate dehydrogenase complex, subunit A, flavoprotein (Fp)	
G03	Mm.246965	NM_023374	Sdhb	Succinate dehydrogenase complex, subunit B, iron sulfur (Ip)	
G04	Mm.198138	NM_025321	Sdhc	Succinate dehydrogenase complex, subunit C, integral membrane protein	
G05	Mm.10406	NM_025848	Sdhd	Succinate dehydrogenase complex, subunit D, integral membrane protein	
G06	Mm.38951	NM_011506	Sucla2	Succinate-Coenzyme A ligase, ADP-forming, beta subunit	
G07	Mm.29845	NM_019879	Suclg1	Succinate-CoA ligase, GDP-forming, alpha subunit	
G08	Mm.371585	NM_011507	Suclg2	Succinate-Coenzyme A ligase, GDP-forming, beta subunit	
G09	Mm.29182	NM_011528	Taldo 1	Transaldolase 1	
G10	Mm.290692	NM_009388	Tkt	Transketolase	
G11	Mm.4222	NM_009415	Tpi1	Triosephosphate isomerase 1	
G12	Mm.28877	NM_139297	Ugp2	UDP-glucose pyrophosphorylase 2	
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, 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.

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 or can be requested from QIAGEN Technical Services or your local distributor.

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