RT² Profiler PCR Array (Rotor-Gene® Format) Human Fatty Acid Metabolism

Cat. no. 330231 PAHS-007ZR

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 Human Fatty Acid Metabolism RT² Profiler PCR Array profiles the expression of 84 key genes involved in the regulation and enzymatic pathways of fatty acid metabolism. Cells, particularly in skeletal muscle and adipose tissue, primarily store energy as triacylglycerols and, when needed, break them down again into glycerol and fatty acids for activation and transport into the mitochondria. The process of â-oxidation then metabolizes these activated fatty acids yielding acetyl-CoA, the initial metabolite necessary for the TCA cycle and ketogenesis. During resting states, cells store excess energy by re-synthesizing fatty acids in a process tightly regulated by hormones. Alterations in the expression of genes involved in fatty acid metabolism, such as CRAT, often associate with metabolic syndrome and insulin resistance. These two syndromes are risk factors for multiple diseases including diabetes and obesity as well as other prevalent health problems such as cardiovascular disease. Using real-time PCR, you can easily and reliably analyze the expression of a focused panel of genes involved in fatty acid 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.



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	Hs.706758	NM_001607	ACAA1	Acetyl-CoA acyltransferase 1
A02	Hs.200136	NM 006111	ACAA2	Acetyl-CoA acyltransferase 2
A03	Hs.331141	NM 025247	ACAD10	Acyl-CoA dehydrogenase family, member 10
A04	Hs.441378	NM 032169	ACAD11	Acyl-CoA dehydrogenase family, member 11
A05	Hs.567482	NM 014049	ACAD9	Acyl-CoA dehydrogenase family, member 9
A05	Hs.471277	NM 001608	ACAD	Acyl-CoA denydrogenase, long chain
A00 A07	Hs.471277 Hs.445040	NM 000016	ACADL	
		-		Acyl-CoA dehydrogenase, C-4 to C-12 straight chain
A08	Hs.507076	NM_000017	ACADS	Acyl-CoA dehydrogenase, C-2 to C-3 short chain
A09	Hs.81934	NM_001609	ACADSB	Acyl-CoA dehydrogenase, short/branched chain
A10	Hs.437178	NM_000018	ACADVL	Acyl-CoA dehydrogenase, very long chain
A11	Hs.232375	NM_000019	ACAT1	Acetyl-CoA acetyltransferase 1
A12	Hs.571037	NM_005891	ACAT2	Acetyl-CoA acetyltransferase 2
B01	Hs.568046	NM_001037161	ACOT1	Acyl-CoA thioesterase 1
B02	Hs.591756	NM_130767	ACOT12	Acyl-CoA thioesterase 12
B03	Hs.122038	NM_001037162	ACOT6	Acyl-CoA thioesterase 6
B04	Hs.126137	NM_181866	ACOT7	Acyl-CoA thioesterase 7
B05	Hs.444776	NM 005469	ACOT8	Acyl-CoA thioesterase 8
B06	Hs.298885	NM 001033583	ACOT9	Acyl-CoA thioesterase 9
B07	Hs.464137	NM 004035	ACOX1	Acyl-CoA oxidase 1, palmitoyl
B08	Hs.444959	NM 003500	ACOX2	Acyl-CoA oxidase 2, branched chain
B09	Hs.479122	NM 003501	ACOX3	Acyl-CoA oxidase 3, pristanoyl
B10	Hs.655760	NM 015162	ACSBG1	Acyl-CoA synthetase bubblegum family member 1
B10 B11	Hs.465720	NM 030924	ACSBG2	Acyl-CoA synthetase bubblegum family member 2
B12		-		, , , , , , , , , , , , , , , , , , , ,
	Hs.406678	NM_001995	ACSL1	Acyl-CoA synthetase long-chain family member 1
C01	Hs.655772	NM_004457	ACSL3	Acyl-CoA synthetase long-chain family member 3
C02	Hs.268785	NM_004458	ACSL4	Acyl-CoA synthetase long-chain family member 4
C03	Hs.11638	NM_016234	ACSL5	Acyl-CoA synthetase long-chain family member 5
C04	Hs.14945	NM_001009185	ACSL6	Acyl-CoA synthetase long-chain family member 6
C05	Hs.706754	NM_005622	ACSM3	Acyl-CoA synthetase medium-chain family member 3
C06	Hs.450804	NM_001080454	ACSM4	Acyl-CoA synthetase medium-chain family member 4
C07	Hs.659606	NM_017888	ACSM5	Acyl-CoA synthetase medium-chain family member 5
C08	Hs.632733	NM_000690	ALDH2	Aldehyde dehydrogenase 2 family (mitochondrial)
C09	Hs.274539	NM 004051	BDH1	3-hydroxybutyrate dehydrogenase, type 1
C10	Hs.124696	NM 020139	BDH2	3-hydroxybutyrate dehydrogenase, type 2
C11	Hs.503043	NM 001876	CPT1A	Carnitine palmitoyltransferase 1A (liver)
C12	Hs.439777	NM 004377	CPT1B	Carnitine palmitoyltransferase 1B (muscle)
D01	Hs.112195	NM 152359	CPT1C	Carnitine palmitoyltransferase 1C
D02	Hs.705379	NM 000098	CPT2	Carnitine palmitoyltransferase 2
D02	Hs.12068	NM 000755	CRAT	Carnitine O-acetyltransferase
D03	Hs.12008	NM 021151	CROT	Carnifine O-acelylinansferase
D04 D05	Hs.125039 Hs.492212	-	DECR1	,
		NM_001359		2,4-diencyl CoA reductase 1, mitochondrial
D06	Hs.628831	NM_020664	DECR2	2,4-dienoyl CoA reductase 2, peroxisomal
D07	Hs.76394	NM_004092	ECHS1	Enoyl CoA hydratase, short chain, 1, mitochondrial
D08	Hs.15250	NM_006117	ECI2	Enoyl-CoA delta isomerase 2
D09	Hs.429879	NM_001966	EHHADH	Enoyl-CoA, hydratase/3-hydroxyacyl CoA dehydrogenase
D10	Hs.380135	NM_001443	FABP1	Fatty acid binding protein 1, liver
D11	Hs.282265	NM_000134	FABP2	Fatty acid binding protein 2, intestinal
D12	Hs.657242	NM_004102	FABP3	Fatty acid binding protein 3, muscle and heart (mammary-derived growth inhibitor)
E01	Hs.391561	NM 001442	FABP4	Fatty acid binding protein 4, adipocyte
E02	Hs.408061	NM 001444	FABP5	Fatty acid binding protein 5 (psoriasis-associated)
E03	Hs.519719	NM 001445	FABP6	Fatty acid binding protein 6, ileal
E04	Hs.26770	NM 001445	FABP7	Fatty acid binding protein 7, brain
E04 E05	Hs.26770 Hs.83190	NM 004104	FABP7	Fatty acid synthase
		-	GCDH	
E06	Hs.532699	NM_000159		Glutaryl-CoA dehydrogenase
E07	Hs.1466	NM_000167	GK	Glycerol kinase
E08	Hs.98008	NM_033214	GK2	Glycerol kinase 2

Position	UniGene	GenBank	Symbol	Description	
E09	Hs.524418	NM_005276	GPD1	Glycerol-3-phosphate dehydrogenase 1 (soluble)	
E10	Hs.512382	NM_000408	GPD2	Glycerol-3-phosphate dehydrogenase 2 (mitochondrial)	
E11 Hs.516032			HADHA	Hydroxyacyl-CoA dehydrogenase/3-ketoacyl-CoA thiolase/enoyl-CoA hydratase	
		NM_000182		(trifunctional protein), alpha subunit	
E12	Hs.533444	NM_000191	HMGCL	3-hydroxymethyl-3-methylglutaryl-CoA lyase	
F01	Hs.397729	NM_002130	HMGCS1	3-hydroxy-3-methylglutaryl-CoA synthase 1 (soluble)	
F02	Hs.59889	NM_005518	HMGCS2	3-hydroxy-3-methylglutaryl-CoA synthase 2 (mitochondrial)	
F03	Hs.656980	NM_005357	LIPE	Lipase, hormone-sensitive	
F04	Hs.180878	NM_000237	LPL	Lipoprotein lipase	
F05	Hs.94949	NM_032601	MCEE	Methylmalonyl CoA epimerase	
F06	Hs.485527	NM_000255	MUT	Methylmalonyl CoA mutase	
F07	Hs.472491	NM_022120	OXCT2	3-oxoacid CoA transferase 2	
F08	Hs.281680	NM_018441	PECR	Peroxisomal trans-2-enoyl-CoA reductase	
F09	Hs.437403	NM_021129	PPA1	Pyrophosphatase (inorganic) 1	
F10	Hs.43322	NM_006251	PRKAA1	Protein kinase, AMP-activated, alpha 1 catalytic subunit	
F11	Hs.437039	NM_006252	PRKAA2	Protein kinase, AMP-activated, alpha 2 catalytic subunit	
F12	Hs.715515	NM_006253	PRKAB1	Protein kinase, AMP-activated, beta 1 non-catalytic subunit	
G01	Hs.50732	NM_005399	PRKAB2	Protein kinase, AMP-activated, beta 2 non-catalytic subunit	
G02	Hs.631630	NM 002730	PRKACA	Protein kinase, cAMP-dependent, catalytic, alpha	
G03	Hs.487325	NM_182948	PRKACB	Protein kinase, cAMP-dependent, catalytic, beta	
G04	Hs.530862	NM_002733	PRKAG1	Protein kinase, AMP-activated, gamma 1 non-catalytic subunit	
G05	Hs.647072	NM_016203	PRKAG2	Protein kinase, AMP-activated, gamma 2 non-catalytic subunit	
G06	Hs.591634	NM_017431	PRKAG3	Protein kinase, AMP-activated, gamma 3 non-catalytic subunit	
G07	Hs.363138	NM_198580	SLC27A1	Solute carrier family 27 (fatty acid transporter), member 1	
G08	Hs.11729	NM_003645	SLC27A2	Solute carrier family 27 (fatty acid transporter), member 2	
G09	Hs.438723	NM_024330	SLC27A3	Solute carrier family 27 (fatty acid transporter), member 3	
G10	Hs.656699	NM_005094	SLC27A4	Solute carrier family 27 (fatty acid transporter), member 4	
G11	Hs.292177	NM_012254	SLC27A5	Solute carrier family 27 (fatty acid transporter), member 5	
G12	Hs.49765	NM_014031	SLC27A6	Solute carrier family 27 (fatty acid transporter), member 6	
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	RPLPO	Ribosomal protein, large, PO	
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		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
RT² 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 <u>www.qiagen.</u> <u>com</u> or can be requested from QIAGEN Technical Services or your local distributor.

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