

# GeneQuery<sup>TM</sup> Human Renal Mesangial Cell Biology qPCR Array Kit (GQH-RMB)

Catalog #GK093

#### **Product Description**

ScienCell's GeneQuery<sup>TM</sup> Human Renal Mesangial Cell Biology qPCR Array Kit (GQH-RMB) is designed to facilitate gene expression profiling of 88 key genes involved in essential mesangial cell biology and functions involved in glomerular physiology and pathophysiology. Brief examples of how included genes may be grouped according to their functions are shown below:

- Renal mesangial cell markers: ACTA1, FN1, PDGFRB, TNS1, ITGA8
- **Mesangial matrix assembly:** SERPINB7, Collagen type 4/5, Laminin α/β/γ, SDC2, HSPG2, CSPG5, NID1, ITGA8
- **Growth factor modulation:** IGF1, PTN, MFGE8, GDF10, CTGF, NGF, PSEN2, SH3KBP1, ADAM12, AFAP1L2, SHC3
- Vasculogenesis: VCAM1, ANGPT2, HOPX, MYH11, NRP2, HDAC9, GJA5, ADAMTS1, FHL2, PLN, CTGF
- Endocytosis: CLTA, CLTB, CLTC, AP2A1, AP2A2, AP2B1, AP2M1, AP2S1, SMAD2, SMAD3, ZFYVE9, TGFB receptors
- Transmembrane transporter activity: KCNMB1, GJA5, CACNA2D1, ABCC9, P2RX1, KCNRG, TRPCs, KCNQ5, PIEZO2
- Genes implicated in
  - o **Mesangial proliferative glomerulonephritis:** SERPINB7, IL6, CD79A, CCL2, FGF2, EDN1, CCR5, GAS6, AXL
  - o **Diffuse mesangial sclerosis:** ACTN4, TRPC6, INF2, WT1, PLCE1, PAX2, ARHGDIA, LAMB2, LAMC1

GeneQuery<sup>TM</sup> qPCR array kits are qPCR ready in a 96-well plate format, with each well containing one primer set that can specifically recognize and efficiently amplify a target gene's cDNA. The carefully designed primers ensure that: (i) the optimal annealing temperature in qPCR analysis is 65°C (with 2 mM Mg<sup>2+</sup>, and no DMSO); (ii) the primer set recognizes all known transcript variants of target gene, unless otherwise indicated; and (iii) only one gene is amplified. Each primer set has been validated by qPCR with melt curve analysis, and gel electrophoresis.

#### GeneQuery<sup>TM</sup> qPCR Array Kit Controls

Each GeneQuery<sup>TM</sup> plate contains eight controls (Figure 1).

- Five target housekeeping genes (β-actin, GAPDH, LDHA, NONO, and PPIH), which enable normalization of data.
- The Genomic DNA (gDNA) Control (GDC) detects possible gDNA contamination in the cDNA samples. It contains a primer set targeting a non-transcribed region of the genome.
- Positive PCR Control (PPC) tests whether samples contain inhibitors or other factors that may negatively affect gene expression results. The PPC consists of a predispensed

- synthetic DNA template and a primer set that can amplify it. The sequence of the DNA template is not present in the human genome, and thus tests the efficiency of the polymerase chain reaction itself.
- The No Template Control (NTC) is strongly recommended, and can be used to monitor the DNA contamination introduced during the workflow such as reagents, tips, and the lab bench.

**Kit Components** 

Component	Quantity	Storage
GeneQuery <sup>™</sup> array plate with lyophilized primers	1	4°C or -20°C
Optical PCR plate seal	1	RT
Nuclease-free H <sub>2</sub> O	2 mL	4°C

**Additional Materials Required (Materials Not Included in Kit)** 

Component	Recommended		
Reverse transcriptase	MultiScribe Reverse Transcriptase (Life Tech, Cat. #4311235)		
cDNA template	Customers' samples		
qPCR master mix	FastStart Essential DNA Green Master (Roche, Cat. #06402712001)		

#### **Quality Control**

All the primer sets are validated by qPCR with melt curve analysis. The PCR products are analyzed by gel electrophoresis. Single band amplification is confirmed for each set of primers.

#### **Product Use**

GQH-RMB is for research use only. It is not approved for human or animal use, or for application in clinical or *in vitro* diagnostic procedures.

#### **Shipping and Storage**

The product is shipped at ambient temperature. Upon receipt, the plate should be stored at 4°C and is good for up to 12 months. For long-term storage (>1 year), store the plate at -20°C in a manual defrost freezer.

**Note:** The primers in each well are lyophilized.

- 1. Prior to use, allow plates to warm to room temperature.
- 2. Briefly centrifuge at 1,500x g for 1 minute before slowly peeling off the seal.
- 3. Prepare 20 µl PCR reactions for one well as shown in Table 1.

Table 1

cDNA template	0.2 – 250 ng
2x qPCR master mix	10 μ1
Nuclease-free H <sub>2</sub> O	variable
Total volume	20 μl

*Important: Only* use polymerases with hot-start capability to prevent possible primer-dimer formation. *Only* use nuclease-free reagents in PCR amplification.

4. Add the mixture of 2x qPCR master mix, cDNA template, and nuclease-free H<sub>2</sub>O to each well containing the lyophilized primers. Seal the plate with the provided optical PCR plate seal.

Important: In NTC control well, do NOT add cDNA template. Add 2x qPCR master mix and nuclease-free H2O only.

- 5. Briefly centrifuge the plates at 1,500x g for 1 minute at room temperature. For maximum reliability, replicates are strongly recommended (minimum of 3).
- 6. For PCR program setup, please refer to the instructions of the master mix of the user's choice. We recommend a typical 3-step qPCR protocol for a 200nt amplicon:

Three-step cycling protocol

Step	Temperature	Time	Number of cycles
Initial denaturation	95°C	10 min	1
Denaturation	95°C	20 sec	
Annealing	65°C	20 sec	40
Extension	72°C	20 sec	40
Data acquisition	Plat	e read	
Recommended	Melting curve analysis		1
Hold	4°C	Indefinite	1

7. (Optional) Load the PCR products on 1.5% agarose gel and perform electrophoresis to confirm the single band amplification in each well.

Figure 1. Layout of GeneQuery<sup>TM</sup> qPCR array kit controls.

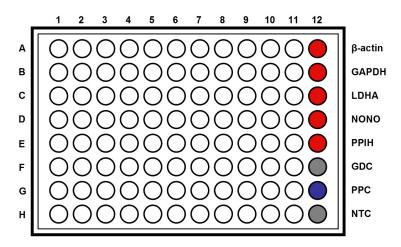


Table 2. Interpretation of control results:

Controls	Results	Interpretation	Suggestions
Housekeeping gene controls	Variability of a housekeeping gene's Cq value	The expression of the housekeeping gene is variable in samples; cycling program is incorrect	Choose a constantly expressed target, or analyze expression levels of multiple housekeeping genes; use correct cycling program and make sure that all cycle parameters have been correctly entered
gDNA Control (GDC)	Cq ≥ 35	No gDNA detected	N/A
	Cq < 35	The sample is contaminated with gDNA	Perform DNase digestion during RNA purification step
Positive PCR Control (PPC)	Cq > 30; or The Cq variations > 2 between qPCR Arrays.	Poor PCR performance; possible PCR inhibitor in reactions; cycling program incorrect	Eliminate inhibitor by purifying samples; use correct cycling program and make sure that all cycle parameters have been correctly entered
No Template Control (NTC)	Positive	DNA contamination in workflow	Eliminate sources of DNA contamination (reagents, plastics, etc.)

Figure 2. A typical amplification curve showing the amplification of a qPCR product.

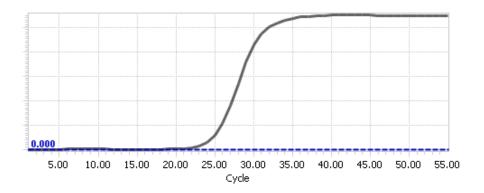
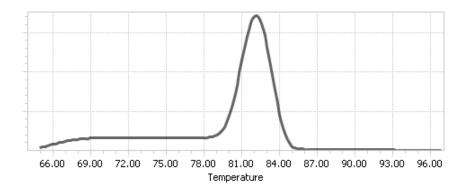


Figure 3. A typical melting peak of a qPCR product.



#### Quantification Method: Comparative $\Delta\Delta$ Cq (Quantification Cycle Value) Method

1. **Note:** Please refer to your qPCR instrument's data analysis software for data analysis. The method provided here serves as guidance for quick manual calculations.

You can use one or more housekeeping genes as a reference to normalize samples.

*Important:* We highly recommend using all 5 housekeeping genes included in this kit,  $\beta$ -actin, GAPDH, LDHA, NONO, and PPIH.

2. For a single housekeeping gene,  $\Delta$ Cq (ref) is the quantification cycle number change for that housekeeping gene (HKG) between an experimental sample and control sample.

$$\Delta$$
Cq (ref) = Cq (HKG, experimental sample) - Cq (HKG, control sample)

When using multiple housekeeping genes as a reference, we recommend normalizing using the geometric mean [1] of the expression level change, which is the same as normalizing using the arithmetic mean of  $\Delta Cq$  of the selected housekeeping genes.

 $\Delta$ Cq (ref) = average ( $\Delta$ Cq (HKG1),  $\Delta$ Cq (HKG2),.....,  $\Delta$ Cq (HKG n)) (n is the number of housekeeping genes selected)

If using all 5 housekeeping genes included in this kit,  $\beta$ -actin, GAPDH, LDHA, NONO, and PPIH, use the following formula:

$$\Delta$$
Cq (ref) = ( $\Delta$ Cq( $\beta$ -actin)+ $\Delta$ Cq(GAPDH)+ $\Delta$ Cq(LDHA)+ $\Delta$ Cq(NONO)+ $\Delta$ Cq(PPIH)) /5

*Note:*  $\Delta$ Cq (HKG) = Cq (HKG, experimental sample) - Cq (HKG, control sample), and  $\Delta$ Cq (HKG) value can be positive, 0, or negative.

3. For any of your genes of interest (GOI),

$$\Delta$$
Cq (GOI) = Cq (GOI, experimental sample) - Cq (GOI, control sample)

$$\Delta\Delta$$
Cq =  $\Delta$ Cq (GOI) -  $\Delta$ Cq (ref)

Normalized GOI expression level fold change =  $2^{-\Delta\Delta Cq}$ 

#### References

[1] Vandesompele J, De Preter K, Pattyn F, Poppe B, Van Roy N, De Paepe A, Speleman F. (2002) "Accurate normalization of real-time quantitative RT-PCR data by geometric averaging of multiple internal control genes." *Genome Biol.* 3(7): 1-12.

#### Example: Comparative ΔΔCq (Quantification Cycle Value) Method

Table 3. Cq (Quantification Cycle) values of 2 genes-of-interest and 5 housekeeping genes obtained for experimental and control samples.

	Genes o	f Interest		House	keeping G	enes	
Samples	GOI1	GOI2	β-actin	GAPDH	LDHA	NONO	PPIH
Experimental	21.61	22.19	17.16	17.84	20.12	19.64	26.40
Control	33.13	26.47	18.20	18.48	20.57	19.50	26.55

$$\Delta$$
Cq (ref) = ( $\Delta$ Cq( $\beta$ -actin)+ $\Delta$ Cq(GAPDH)+ $\Delta$ Cq(LDHA)+ $\Delta$ Cq(NONO)+ $\Delta$ Cq(PPIH)) /5 = ((17.16-18.20)+(17.84-18.48)+(20.12-20.57)+(19.64-19.50)+(26.40-26.55))/5 = -0.43

$$\Delta$$
Cq (GOI1) = 21.61 - 33.13  
= -11.52

$$\Delta$$
Cq (GOI2) = 22.19 - 26.47  
= -4.28

$$\Delta\Delta$$
Cq (GOI1) =  $\Delta$ Cq (GOI1) -  $\Delta$ Cq (ref)  
= -11.52 - (-0.43)  
= -11.09

$$\Delta\Delta$$
Cq (GOI2) =  $\Delta$ Cq (GOI2) -  $\Delta$ Cq (ref)  
= -4.28 - (-0.43)  
= -3.85

Normalized GOI1 expression level fold change = 
$$2^{-\Delta\Delta Cq \text{ (GOI1)}}$$
  
=  $2^{11.09}$   
= 2180

Normalized GOI2 expression level fold change = 
$$2^{-\Delta\Delta Cq \text{ (GOI2)}}$$
  
=  $2^{3.85}$   
= 14.4

*Conclusion:* Upon treatment, expression level of GOI1 increased 2,180 fold, and expression level of GOI2 increased 14.4 fold.



# GeneQuery<sup>TM</sup> Human Renal Mesangial Cell Biology qPCR Array Kit (GQH-RMB)

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GeneQuery  $^{TM}$  Human Renal Mesangial Cell Biology qPCR Array Plate Layout\* (8 controls in Bold and Italic)

	1	2	3	4	5	6	7	8	9	10	11	12
A	ABCC9	AP2A1	CCL2	COL5A1	FHL2	IGF1	LAMA2	MFGE8	PDGFRB	SFRP2	TGFBR3	β-actin
В	ACTA1	AP2A2	CCR5	COL5A2	FN1	IL6	LAMA3	MYH11	PIEZO2	SH3KBP1	TNS1	GAPDH
C	ACTN4	AP2B1	CD79A	COL5A3	GAS6	INF2	LAMA4	NGF	PLCE1	SHC3	TRPC1	LDHA
D	ADAM12	AP2M1	CLTA	CSPG5	GDF10	ITGA8	LAMA5	NID1	PLN	SMAD1	TRPC4	NONO
E	ADAMTS1	AP2S1	CLTB	CTGF	GJA5	KCNMB1	LAMB1	NID2	PSEN2	SMAD2	TRPC6	PPIH
$\mathbf{F}$	AFAP1L2	ARHGDIA	CLTC	EDN1	HDAC9	KCNQ5	LAMB2	NRP2	PTN	SMAD3	VCAM1	GDC
G	AGTR1	AXL	COL4A1	ETS1	HOPX	KCNRG	LAMC1	P2RX1	SDC2	TGFBR1	WT1	PPC
H	ANGPT2	CACNA2D1	COL4A2	FGF2	HSPG2	LAMA1	MECOM	PAX2	SERPINB7	TGFBR2	ZFYVE9	NTC

<sup>\*</sup> gene selection may be updated based on new research and development

### Plate type A

Brand	Model	kit catalog #
ABI / Life Tech	ABI 5700	GK093-A
	ABI 7000	GK093-A
	ABI 7300	GK093-A
	ABI 7500	GK093-A
	ABI 7700	GK093-A
	ABI 7900 HT	GK093-A
	QuantStudio	GK093-A
	ViiA 7	GK093-A
Bio-Rad	Chromo4	GK093-A
	iCycler	GK093-A
	iQ5	GK093-A
	MyiQ	GK093-A
	MyiQ2	GK093-A
Eppendorf / Life Tech	Matercycler ep realplex 2	GK093-A
	Matercycler ep realplex 4	GK093-A
Stratagene	MX3000P	GK093-A
J	MX3005P	GK093-A

### Plate type B

Brand	Model	kit catalog #
ABI / Life Tech	ABI 7500 Fast	GK093-B
	ABI 7900 HT Fast	GK093-B
	QuantStudio Fast	GK093-B
	StepOnePlus	GK093-B
	ViiA 7 Fast	GK093-B
Bio-Rad	CFX Connect	GK093-B
	CFX96	GK093-B
	DNA Engine Opticon 2	GK093-B
Stratagene	MX4000	GK093-B

## Plate type C

Brand	Model	kit catalog #
Roche	Lightcycler 96	GK093-C
	Lightcycler 480 (96-well)	GK093-C