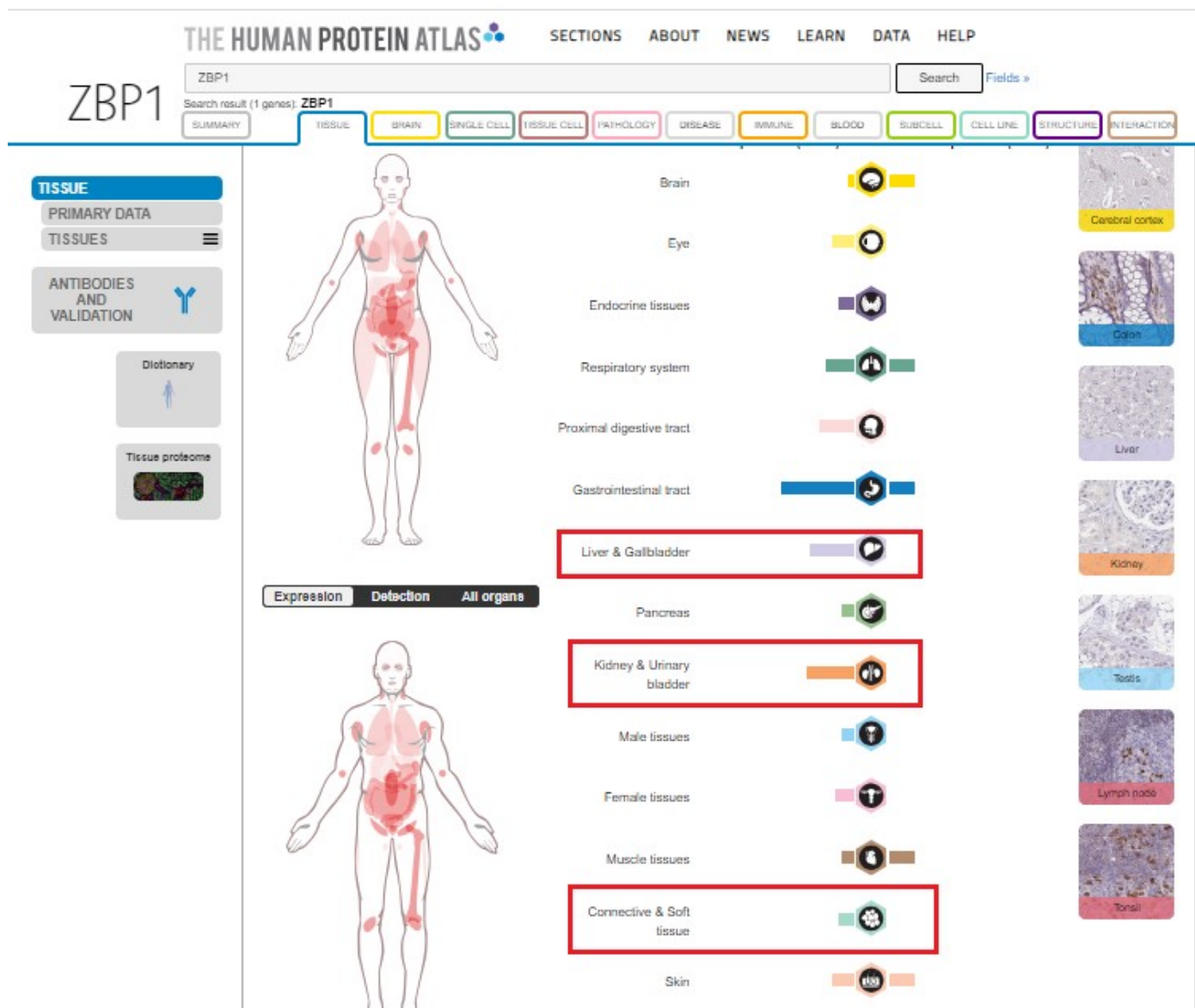


Supplementary information:

Supplementary Figures:

S1 Figures: Validation of the significant expression of the candidate mRNAs (ZBP1, STING1, and DDX58=RIG-1) in the T2DM-related tissues using THE HUMAN PROTEIN ATLAS database <https://www.proteinatlas.org/>, Jan 2022.



STING1

STING1

Search

Fields »

Search result (2 genes): **STING1** | STEEP1

SUMMARY

TISSUE

BRAIN

SINGLE CELL

TISSUE CELL

PATHOLOGY

DISEASE

IMMUNE

BLOOD

SUBCELL

CELL LINE

STRUCTURE

INTERACTION

TISSUE

PRIMARY DATA

TISSUES

ANTIBODIES AND VALIDATION

Dictionary

Tissue proteome

RNA AND PROTEIN EXPRESSION SUMMARY



Expression Detection All organs



RNA expression (nTPM)¹

Protein expression (score)²



RIGI

DDX58 Search Fields »

Search result (2 genes): RIGI | ARL16

- SUMMARY
- TISSUE
- BRAIN
- SINGLE CELL
- TISSUE CELL
- PATHOLOGY
- DISEASE
- IMMUNE
- BLOOD
- SUBCELL
- CELL LINE
- STRUCTURE
- INTERACTION

TISSUE

PRIMARY DATA

TISSUES

ANTIBODIES AND VALIDATION

Dictionary

Tissue proteome

RNA AND PROTEIN EXPRESSION SUMMARY



Expression Detection All organs



RNA expression (nTPM)

Protein expression (score)

Tissue	RNA expression (nTPM)	Protein expression (score)	Image
Brain	Low	Low	Cerebral cortex
Eye	Low	Low	Colon
Endocrine tissues	Low	Low	Liver
Respiratory system	Low	Low	Kidney
Proximal digestive tract	Low	Low	Testis
Gastrointestinal tract	Low	Low	Lymph node
Liver & Gallbladder	High	High	
Pancreas	Low	Low	
Kidney & Urinary bladder	High	High	
Male tissues	Low	Low	
Female tissues	Low	Low	
Muscle tissues	Low	Low	
Connective & Soft tissue	Low	Low	

S2 Figures: Validation of the implication of the candidate mRNAs (ZBP1, STING1, and DDX58=RIG-1) in the development of diabetes, and kidney and liver diseases using DisGeNET database (<https://www.disgenet.org/home/>), the Expression Atlas database (<https://www.ebi.ac.uk/gxa/home>) and diseases database (<https://diseases.jensenlab.org/Search>), Jan 2022.

DISEASES

Disease-gene associations mined from literature

Search
Downloads
About

ZBP1 disease associations

ZBP1 [ENSP00000360215]

Z-DNA-binding protein 1. Participates in the detection by the host's innate immune system of DNA from viral, bacterial or even host origin. Plays a role in host defense against tumors and pathogens. Acts as a cytoplasmic DNA sensor which, when activated, induces the recruitment of TBK1 and IRF3 to its C-terminal region and activates the downstream interferon regulatory factor (IRF) and NF-kappa B transcription factors, leading to type-I interferon production. ZBP1-induced NF-kappaB activation probably involves the recruitment of the RHIM-containing kinases RIPK1 and RIPK3 (By similarity).

Synonyms: ZBP1, ZBP1p, hZBP1, HOY8D1, Q9H171 ...

Linkouts: STRING Pharos UniProt OMIM

Text mining

< Prev | Next >

Name	Z-score	Confidence
Adult respiratory distress syndrome	2.3	*****
Human immunodeficiency virus infectious disease	2.2	*****
Listeriosis	2.1	*****
Cowpox	2.1	*****
Deconvulsating disease	2.1	*****
Diabetes	2.1	*****
Parasitosis	2.1	*****
Liver disease	2.1	*****
Kidney disease	2.1	*****
Chickenpox	2.0	*****

Expression Atlas

Gene expression across species and biological conditions

Query single cell expression

To Single Cell Expression Atlas >

Home
Browse experiments
Download
Release notes
FAQ
Help
Licence
About
Support

Results for ZBP1 (symbol) AND diabetes mellitus

Baseline expression
Differential expression

Filter your results

Kingdom

Animals

Species

Homo sapiens

Mus musculus

Experiment type

Microarray 1-colour mRNA differential

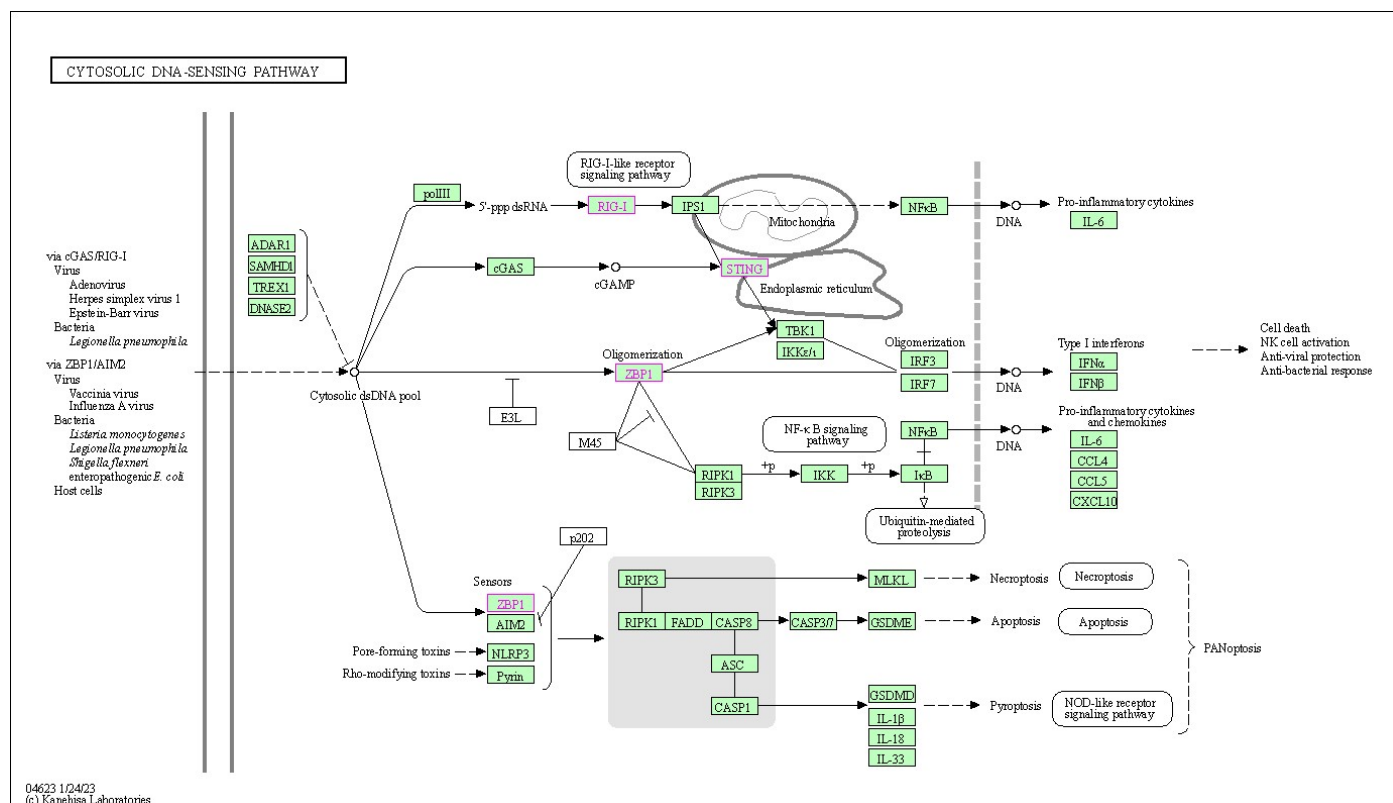
RNA-seq mRNA differential

Display log₂-fold change

Download results

Log ₂ -fold change	Species	Gene name	Comparison	Experimental variables	Experiment name
1.1		<u>Zbp1</u>	'control; streptozotocin-induced diabetes' vs 'control; control'	diet, phenotype	<u>Hepatic gene expression in streptozotocin-induced diabetic mice fed a phloridzin diet</u>
1.1		<u>ZBP1</u>	'advanced' vs 'early' in 'steatosis; liver disease; Hepatitis C virus genotype 3'	clinical information, disease, disease staging, infect	<u>RNA-seq of hepatic biopsies taken from patients with chronic liver disease presenting with different aetiologies (HCV, FLD) and fibrosis stages.</u>

S3 Figure: The visualization of the candidate mRNAs (ZBP1, STING1, and DDX58=RIG-1) in the cGAS-STING pathway through KEGG pathway database (<https://www.genome.jp/kegg/>, Jan 2022)



S4 Figures: Validation of the implication of the selected biochemical effectors proteins (LC3B=MAP1LC3B, and TNF) in the development of diabetes, and kidney and liver diseases using DisGeNET database (<https://www.disgenet.org/home/>), Jan 2022.

Summary of GDAs | Evidences for GDAs | Summary of VDAs | Evidences for VDAs

MAP1LC3B, microtubule associated protein 1 light chain 3 beta, 81631

N. diseases: 167; N. variants: 1

Source: ALL

Results per page 200

Filter within current results:

Disease ^	Type ⇅	Disease Class ⇅	Semantic Type ⇅	N. genes _d ⇅	N. SNPs _d ⇅	Score _{gda} ⇅	EL _{gda} ⇅	EI _{gda} ⇅	N. PMIDs ⇅	N. SNPs _{gda} ⇅	First Ref. ⇅	Last Ref. ⇅
Diabetes Mellitus	group	Nutritional and Metab...	Disease or Syndr...	2803	824	0.010	None	1.000	1		2019	2019
Diabetic Retinopathy	disease	Eye Diseases; Endoc...	Disease or Syndr...	645	213	0.020	None	1.000	2		2017	2019

MAP1LC3B, microtubule associated protein 1 light chain 3 beta, 81631

N. diseases: 167; N. variants: 1

Source: ALL

Results per page 200

Filter within current results: liver

Disease ^	Type ⇅	Disease Class ⇅	Semantic Type ⇅	N. genes _d ⇅	N. SNPs _d ⇅	Score _{gda} ⇅	EL _{gda} ⇅	EI _{gda} ⇅	N. PMIDs ⇅	N. SNPs _{gda} ⇅	First Ref. ⇅	Last Ref. ⇅
Adult Liver Carcinoma	disease	Digestive System Dis...	Neoplastic Process	1377	72	0.010	None	1.000	1		2018	2018
Chemical and Drug Induced Liver Injury	disease	Digestive System Dis...	Disease or Syndr...	461	38	0.300	None	1.000	1		2014	2014
Chemically-Induced Liver Toxicity	disease	Digestive System Dis...	Disease or Syndr...	412		0.300	None	1.000	1		2014	2014
Drug-Induced Acute Liver Injury	disease	Digestive System Dis...	Disease or Syndr...	413		0.300	None	1.000	1		2014	2014
Drug-Induced Liver...	phenotype	Digestive System Dis...	Disease or Syndr...	537	29	0.300	None	1.000	1		2014	2014
Liver and Intrahepa...	disease	Digestive System Dis...	Neoplastic Process	1395	73	0.010	None	1.000	1		2018	2018
Liver carcinoma	disease	Digestive System Dis...	Neoplastic Process	5725	942	0.060	None	1.000	6		2014	2018

MAP1LC3B, microtubule associated protein 1 light chain 3 beta, 81631

N. diseases: 167; N. variants: 1

Source: ALL

Results per page 200

Filter within current results: renal

Disease ^	Type ⇅	Disease Class ⇅	Semantic Type ⇅	N. genes _d ⇅	N. SNPs _d ⇅	Score _{gda} ⇅	EL _{gda} ⇅	EI _{gda} ⇅	N. PMIDs ⇅	N. SNPs _{gda} ⇅	First Ref. ⇅	Last Ref. ⇅
Conventional (Clear Cell) Renal Cell Carcinoma	disease	Neoplasms; Female ...	Neoplastic Process	2346	222	0.040	None	1.000	4		2012	2019
Renal Cell Carcino...	disease	Neoplasms; Female ...	Neoplastic Process	2084	288	0.020	None	1.000	2		2019	2019

TNF, tumor necrosis factor, 7124

1 - 200 of 2724 results

Download Share

N. diseases: 2724; N. variants: 31

Add/Remove filter

Source: ALL

Results per page 200

Filter within current results: diab

Disease	Type	Disease Class	Semantic Type	N. genes _d	N. SNPs _d	Score _{gda}	EL _{gda}	EI _{gda}	N. PMIDs	N. SNPs _{gda}	First Ref.	Last Ref.
Diabetes Mellitus	disease	Nutritional and Metab	Disease or Syndr	3134	2672	0.600	None	0.950	179	7	1995	2020
Diabetes Mellitus	disease	Nutritional and Metab	Experimental Mo	522		0.500	None	1.000	10		2002	2014
Diabetes Mellitus, I...	disoco	Nutritional and Metab...	Disease or Syndr...	1675	954	0.400	None	0.950	101	2	1989	2020
Diabetic Nephropathy	disoco	Female Urogenital Dis...	Disease or Syndr...	1189	238	0.300	None	0.957	47	3	1997	2020

TNF, tumor necrosis factor, 7124

1 - 200 of 2724 results

Download Share

N. diseases: 2724; N. variants: 31

Add/Remove filter

Source: ALL

Results per page 200

Filter within current results: kid

Disease	Type	Disease Class	Semantic Type	N. genes _d	N. SNPs _d	Score _{gda}	EL _{gda}	EI _{gda}	N. PMIDs	N. SNPs _{gda}	First Ref.	Last Ref.
Kidney Failure, Acute	disease	Female Urogenital Di...	Disease or Syndr...	826	32	0.400	None	1.000	29	1	2002	2020
Kidney Failure, Chr...	disease	Female Urogenital Di...	Disease or Syndr...	827	425	0.400	None	0.960	25		1994	2019

TNF, tumor necrosis factor, 7124

1 - 200 of 2724 results

Download Share

N. diseases: 2724; N. variants: 31

Add/Remove filter

Source: ALL

Results per page 200

Filter within current results: liver

Disease	Type	Disease Class	Semantic Type	N. genes _d	N. SNPs _d	Score _{gda}	EL _{gda}	EI _{gda}	N. PMIDs	N. SNPs _{gda}	First Ref.	Last Ref.
Liver Failure, Acute	disease	Digestive System Dis...	Disease or Syndr...	282	21	0.600	None	1.000	27		1998	2019
Chemical and Drug Induced Liver Injury	disease	Digestive System Dis...	Disease or Syndr...	461	38	0.500	None	1.000	12		1995	2019
Liver Cirrhosis, Experimental	disease	Pathological Conditio...	Experimental Mo...	870		0.500	None	1.000	2		2005	2014
Liver carcinoma	disease	Digestive System Dis...	Neoplastic Process	5725	942	0.400	None	0.979	146	2	1990	2020
Liver diseases	group	Digestive System Dis...	Disease or Syndr...	1019	100	0.400	None	1.000	30		1991	2019
Fatty Liver	disease	Digestive System Dis...	Disease or Syndr...	875	35	0.400	None	0.950	20		2006	2020
Liver neoplasms	group	Digestive System Dis...	Neoplastic Process	1424	7	0.400	None	1.000	16		1990	2019
Malignant neoplas...	disease	Digestive System Dis...	Neoplastic Process	1649	88	0.400	None	1.000	13		2008	2019

Activate Windows
Go to Settings to activate Windows.

S5 Figure: Validation of the interaction between the three key mRNAs and two effector proteins with the retrieved miR-1976 from mirwalk3 (<http://mirwalk.umm.uni-heidelberg.de/>, Jan 2022)

miRWalk

HOME FAQ RESOURCES ABOUT

hsa-miR-1976 GeneSymbol 0.9 3UTR set filter

miRNA	RefseqID	GeneSymbol	Duplex	Score	Position	Binding Site	Au	Me	N Pairings	Targetscan	Mirdb	Mirtarbase
hsa-miR-1976	NM_001367258	STING1	details	1.00	3UTR	1667,1693	0.43	-5.481	18	—	—	—
hsa-miR-1976	NM_198282	STING1	details	1.00	3UTR	1894,1920	0.43	-5.481	18	—	—	—
hsa-miR-1976	NM_014314	DDX58	details	0.92	3UTR	4343,4393	0.43	-4.171	15	—	—	—
hsa-miR-1976	NM_014314	DDX58	details	1.00	3UTR	4002,4020	0.47	-12.956	14	—	—	—
hsa-miR-1976	NM_022818	MAP1LC3B	details	0.92	3UTR	1298,1314	0.59	-6.675	11	—	—	—
hsa-miR-1976	NM_001160418	ZBP1	details	0.92	3UTR	1627,1656	0.41	-5.495	18	—	—	—
hsa-miR-1976	NM_001301738	STING1	details	1.00	3UTR	1707,1733	0.43	-5.481	18	—	—	—

miRWalk

HOME FAQ RESOURCES ABOUT

hsa-miR-1976 GeneSymbol 0.9 5UTR set filter

miRNA	RefseqID	GeneSymbol	Duplex	Score	Position	Binding Site	Au	Me	N Pairings	Targetscan	Mirdb	Mirtarbase
hsa-miR-1976	NM_000594	TNF	details	1.00	5UTR	14,33	0.43	-9.268	16	—	—	—

S6 Figure: The association of miR-1976 with cGAS-STING-related pathways through DIANA tools mirPath 3 (<http://www.microrna.gr/miRPathv3>, Jan 2022)

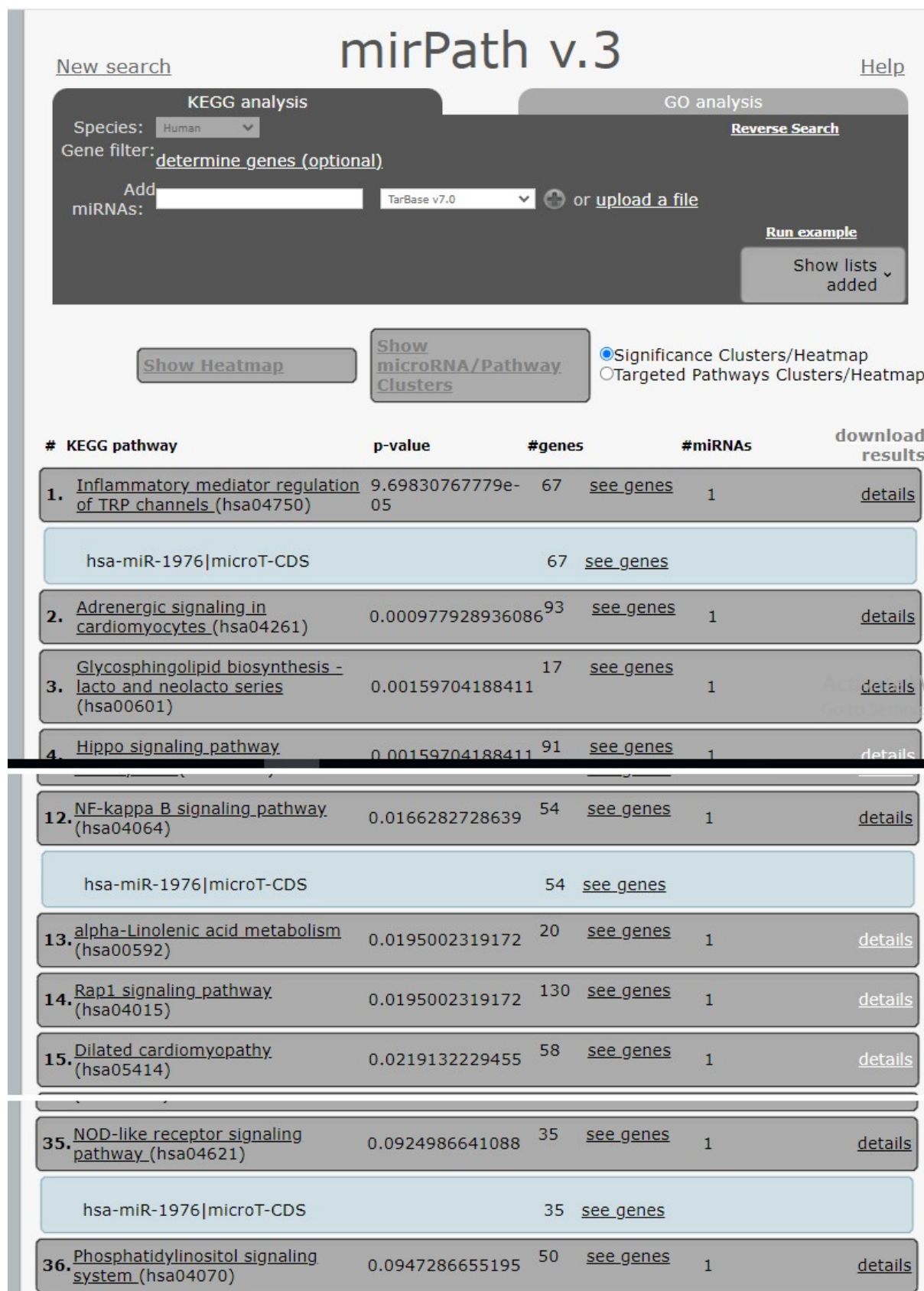


Figure S7: Retrieval of AC074117.2 lncRNA from DIANA Tools database (<https://diana.e-ce.uth.gr/lncbasev3>, Jan 2022)

miRNA: Location: [Clear All](#)

[Execute Search](#) [Clear All](#) [Download](#)

Interactions: 38, Experiments 1 (low: 0, high: 1) Cell lines: 1, Tissues: 1, Publications: 1

Filters [Reset](#)

- Tissue: [Select Tissues](#)
- Cell Type: [Select Cell Types](#)
- Method: [Select Methods](#)
- Validated As: [Select Validated as](#)
- Validation Type: [Select Validation Types](#)
- miRNA Conf. Level: [Select Confidence Levels](#)
- Biotypes: [Select Biotypes](#)
- Species: [Select Species](#)
- lncRNA Source: [Select lncRNA Sources](#)
- Variants: [All](#) [With](#) [Without](#)

Page: [First](#) [Previous](#) [1](#) [2](#) [3](#) [4](#) [Next](#) [Last](#)

Gene name	miRNA name	Experiments	Publications	Cell Lines	Tissues	miRNA Conf.	DIANA Links
AC007619.2	hsa-miR-1976	1	1	1	1	Low	lncRNA Expr UCSC mT TB InP mP
AC012123.2	hsa-miR-1976	1	1	1	1	Low	lncRNA Expr UCSC mT TB InP mP
AC012531.1	hsa-miR-1976	1	1	1	1	Low	lncRNA Expr UCSC mT TB InP mP
AC013404.1	hsa-miR-1976	1	1	1	1	Low	lncRNA Expr UCSC mT TB InP mP
AC074117.2	hsa-miR-1976	1	1	1	1	Low	lncRNA Expr UCSC mT TB InP mP

Publication	Methods	Tissue	Cell Type	Tested Cell Line	Category	Experimental Condition
Karginov FV et al. 2013	IP	Kidney	293S	NA	Embryonic/Fetal	Active Windows

Figure S8: lncRNA AC074117.2 / miR-1976 interaction using IntaRNA 2.0 tool.



Supplementary Tables:

Table S4: Primers used for qPCR.

Primer name	Gene ID	Cat.no.	Tm	Product size
<i>TMEM173 (STING1)</i>	NM_198282	249900 Gene Globe ID: QT00055440	70 °C	125 bp
<i>ZBP1</i>	NM_001160418	249900 Gene Globe ID: QT00029631	70 °C	83 bp
<i>DDX58</i>	NM_014314	249900 Gene Globe ID: QT00040509	70 °C	114 bp
<i>GAPDH</i>	NM_002046	249900 Gene Globe ID: QT00079247	70°C	95 bp
<i>miR_1976</i>	MIMAT0009451	339350 Gene Globe ID: ZP00000388	75 °C	85 bp
<i>AC074117.2</i>	ENST00000417130	Gene Globe ID: UPFH0284997	58.8 °C	82 bp