# RNA editing enzyme ADAR1 governs the circadian expression of P-glycoprotein in human renal cells by regulating alternative splicing of the *ABCB1* gene

Yuji Omata, Tomoaki Yamauchi, Akito Tsuruta, Naoya Matsunaga, Satoru Koyanagi, Shigehiro Ohdo

### **Supporting information**

## **EXPERIMENTAL PROCEDURES**

Supplementary Table S1 Primer sets for direct sequencing

Supplementary Figure S1 *In silico* prediction of RNA secondary structure of introns of the human *ABCB1* gene by RNAfold.

Supplementary Figure S2 Sequence analysis for exon 8 of the human *PVR* mRNA and intron 27 of the human *ABCB1* pre-mRNA.

Supplementary Figure S3 Electropherograms from direct sequencing of the PCR-amplified product of intron 27 of the human *ABCB1* pre-mRNA.

REFERENCE

#### **EXPERIMENTAL PROCEDURES**

#### **Construction of ADAR1-overexpressing RPTECs**

The ORF sequence of human *ADAR1* with C-terminal GFP-tag was amplified with ADAR1 Human Tagged ORF Clone (RG219761; Origene Technologies, Inc., Rockville, MD) as a template, and subcloned into pLVSIN-CMV Pur Vector (Takara Bio Inc.) using In-Fusion HD Cloning Kit (Takara Bio Inc.) and primer sets as follows: forward 5'-CGGTGAATTCCTCGAGCCGCCGCGATCGCCATG-3', reverse 5'-TAGAACTAGTCTCGATTAAACTCTTTCTTCACCGG-3'. Lentiviral particles were prepared by the Lentiviral High Titer Packaging Mix with pLVSIN series (Takara Bio Inc.) with Lenti-X 293 T cell lines. RPTECs were infected with ADAR1-expressing lentivirus and maintained in a medium containing 5 µg/mL of puromycin (FUJIFILM Wako Pure Chemical Corporation). Overexpression of ADAR1 with GFP-tag was confirmed by Western blotting.

#### **Direct Sequencing**

RNA was extracted using the ReliaPrep RNA Cell Miniprep System (Promega) and treated with DNase I on a column. A total of 500 ng of RNA was used for cDNA synthesis with ReverTra Ace qPCR RT Kit (TOYOBO Co., Ltd.). Genomic DNA was extracted using Wizard Genomic DNA Purification Kit (Promega). Direct sequencing for exon 8 of the PVR gene was conducted as follows. The cDNA and genomic DNA were amplified by the GoTaq Green Master Mix (Promega) with gene-specific primers (Human PVR exon 8 outside; Supplementary Table S1). The PCR-amplified products were used for  $2^{nd}$  amplification as templates with another gene-specific primers (Human *PVR* exon 8 inside; Supplementary Table S1). The PCR-amplified products were analyzed by Sanger sequencing using the same primers used in 2<sup>nd</sup> amplification. Direct sequencing for intron 27 of the ABCB1 gene was conducted as follows. The cDNA and genomic DNA were amplified by the GoTaq Green Master Mix (Promega) with gene-specific primers (Human *ABCB1* intron 27 outside; **Supplementary Table S1**). PCR-amplified products were separated by electrophoresis using 1% agarose gel containing ethidium bromide. PCR-amplified products of intron 27-containing transcripts (maybe both retained intron 27 and pre-mRNA), which were 1,590 bp in length, were extracted and purified using QIAquick Gel Extraction Kit (Qiagen, Hilden, Germany). The purified DNA was amplified by the GoTaq Green Master Mix (Promega) with another gene-specific primers (Human ABCB1 intron 27 inside Forward and outside Reverse; Supplementary Table S1). The PCR-amplified products were analyzed by Sanger sequencing using the same primers used in 2<sup>nd</sup> amplification. Electropherograms were aligned using SnapGene Viewer (https://www.snapgene.com/snapgene-viewer/).

Gene	Primers
Human PVR exon 8 outside	
Forward	5'-CACAGAGCCAGGAATGGAGAGTG-3'
Reverse	5'-ATGTTCCAACCACCAGACAGAGG-3'
Human PVR exon 8 inside	
Forward	5'-GGCTAAAACACTCCACACCCTC-3'
Reverse	5'- ACAAACAGCAGAGCAGCAGA-3'
Human ABCB1 intron 27 outside	
Forward	5'-AGCTCGTGCCCTTGTTAGAC-3'
Reverse	5'-GGCGGTGAGCAATCACAATG-3'
Human ABCB1 intron 27 inside	
Forward	5'-AIGGAGICICGCICIGICG-3'

# Supplementary Table S1 Primer sets for direct sequencing



**Supplementary Figure S1** *In silico* **prediction of RNA secondary structure of introns of the human** *ABCB1* **gene by RNAfold.** The minimum free energy structure with base-pair probabilities calculated to have the lowest value of free energy. Base-pair probabilities are shown by a color spectrum. The RNA editing sites registered in REDIportal are indicated by arrows.



Supplementary Figure S2 Sequence analysis for exon 8 of the human *PVR* mRNA and intron 27 of the human *ABCB1* pre-mRNA. *A*, Expression of endogenous and exogenous ADAR1 protein in mock-transduced, ADAR1-KD, and ADAR1-overexpressing RPTECs. The protein levels of  $\beta$ -ACTIN are shown as loading controls. *B*, Upper panels show schematic image of procedure of amplification of exon 8 region of the human *PVR* gene for direct sequencing. Lower panels show schematic image of procedure of amplification of intron 27 region of *ABCB1* pre-mRNA for direct sequencing. *C*, Electropherograms around A-to-I editing site of the human *PVR* gene exon 8, which is a well-known A-to-I editing site (1). This region was used as positive control for ADAR1-mediated A-to-I editing. The sequence data are obtained from mock-transduced, ADAR1-KD, and ADAR1-overexpressing RPTECs.

1644 01 00 01 01 01 01 01 01 01 01 01 01 01	
	mock
	ADAR1-KD
a An man A man	ADAR1 overexpress
site1 site2 site3 site4 site5 site6 +759 GACATTGTCTCTACAAAAAAAAAAAAAAAAAAAAAAAGCTGGTGGGAACTTGTGGGAACTTGTAGTCCCAGCTACTCAGAAGGCTGAGGGAACAAAAGAACGCTGAGGTGGGAACTTGTAGTCCCAGCTACTCAGAAGGCTGAGGGAACAAAAAAAA	853
Manhan han han han han han han han han ha	genomic DNA
Martin and Ma	mock
Manhan	ADAR1-KD
site7 site8 site9 site10 site11 site12 site14 site15 site16 site17 site18 site19 site20	ADAR1 overexpress
+854 TO A GO CO A GO A O G TTO A GO C TO CA OTA A A CO A TTO TA CO CO TO CO CO CO TO CO A GO A A TO A GO CO CO TO TTO A A TA A A A A A A A A A A A A	+928
	genomic DNA
Allanda and Allanda and Allanda and Allanda and Allanda and Allanda allanda and Allanda Allanda and Allanda and	mock
	ADAR1-KD
site21 site22 site23 site24	overexpress
41097 TERRATIC TO CONTRACTOR T	+1184 genomic DNA
M.M.M.M.M.M.M.M.M.M.M.M.M.M.M.M.M.M.M.	+1184 genomic DNA mock
	+1184 genomic DNA mock ADAR1-KD
	genomic DNA mock ADAR1-KD ADAR1 overexpress
+1097 /	+1184 genomic DNA mock ADAR1-KD ADAR1 overexpress
+1185	+1184 genomic DNA mock ADAR1-KD ADAR1 overexpress +1272 genomic DNA
	+1184 genomic DNA mock ADAR1-KD ADAR1 overexpress +1272 genomic DNA mock
	+1184 genomic DNA mock ADAR1-KD ADAR1 overexpress +1272 genomic DNA mock ADAR1-KD
	+1184 genomic DNA mock ADAR1-KD ADAR1-KD +1272 genomic DNA mock ADAR1-KD ADAR1 overexpress
+1097 ************************************	+1184 genomic DNA mock ADAR1-KD ADAR1-KD +1272 genomic DNA mock ADAR1-KD ADAR1-KD ADAR1-KD
	+1184 genomic DNA mock ADAR1-KD ADAR1-KD ADAR1 overexpress mock ADAR1-KD ADAR1-KD ADAR1 overexpress +1359 genomic DNA
	+1184 genomic DNA mock ADAR1-KD ADAR1-KD ADAR1 overexpress 41272 genomic DNA ADAR1-KD ADAR1-KD ADAR1 41359 genomic DNA mock
	+1184 genomic DNA mock ADAR1-KD ADAR1-KD +1272 genomic DNA mock ADAR1-KD ADAR1-KD appendic DNA mock ADAR1-KD

Supplementary Figure S3 Electropherograms from direct sequencing of the PCR-amplified product of intron 27 of the human *ABCB1* pre-mRNA. The sequence data are obtained from pre-mRNA extracted from mock-transduced, ADAR1-KD, and ADAR1-overexpressing RPTECs. The RNA editing sites registered in REDIportal are indicated by arrows.

#### REFERENCE

 Hong, H., An, O., Chan, T. H. M., Ng, V. H. E., Kwok, H. S., Lin, J. S., Qi, L., Han, J., Tay, D. J. T., Tang, S. J., Yang, H., Song, Y., Bellido Molias, F., Tenen, D. G., and Chen, L. (2018) Bidirectional regulation of adenosine-to-inosine (A-to-I) RNA editing by DEAH box helicase 9 (DHX9) in cancer. *Nucleic Acids Res.* 46, 7953–7969