

# **Supplemental Data**

## **Manuscript title:**

**The Circadian Clock Gene Bmal1 Controls Intestinal Exporter MRP2 and**

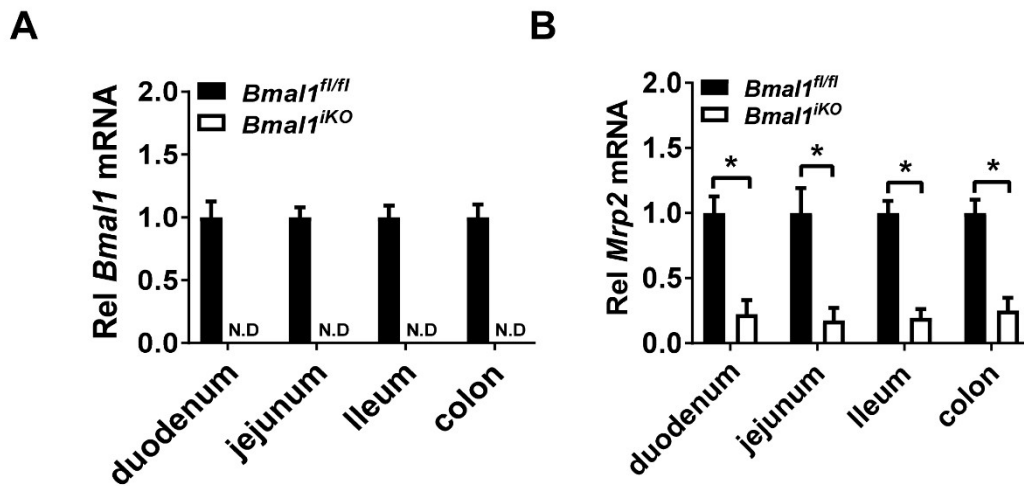
**Drug Disposition**

## **Authors:**

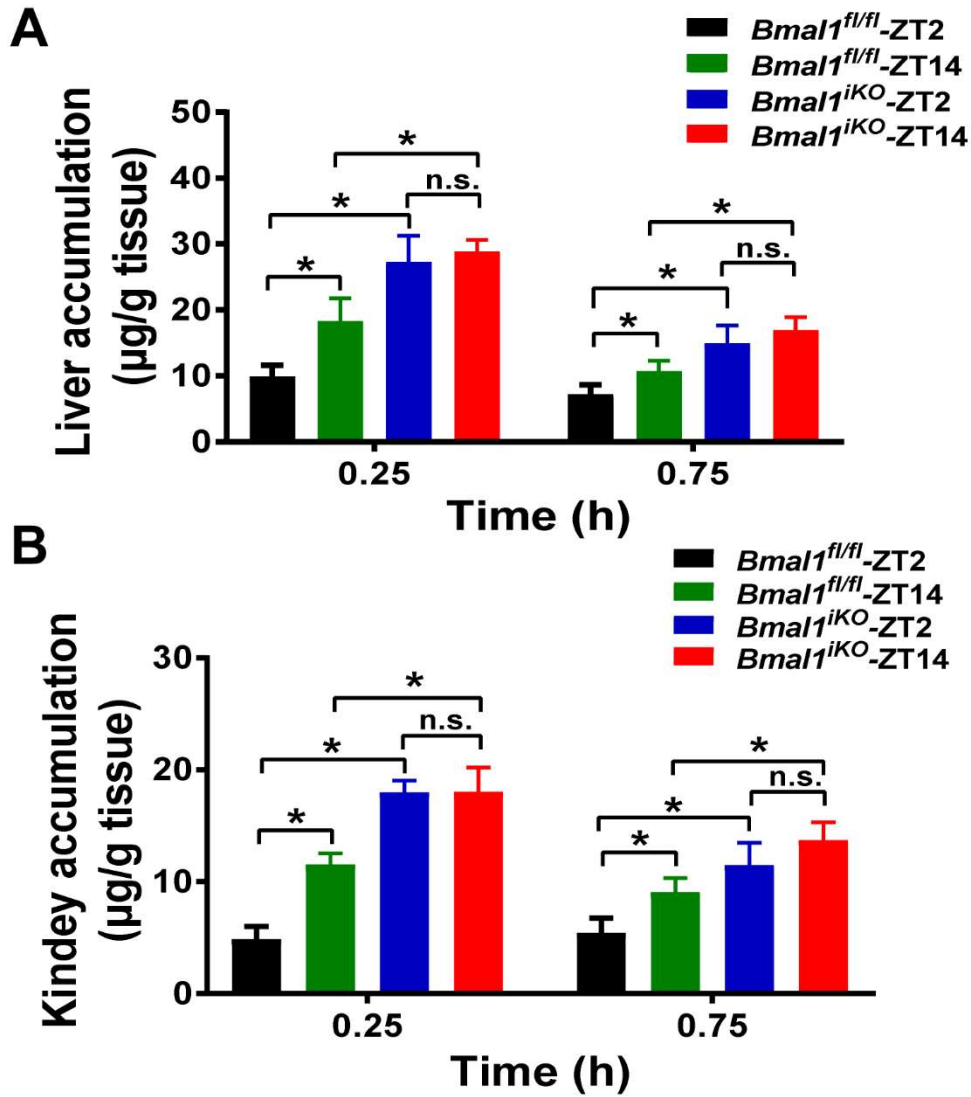
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**Supplementary Table 1.** Oligonucleotides used in this study.

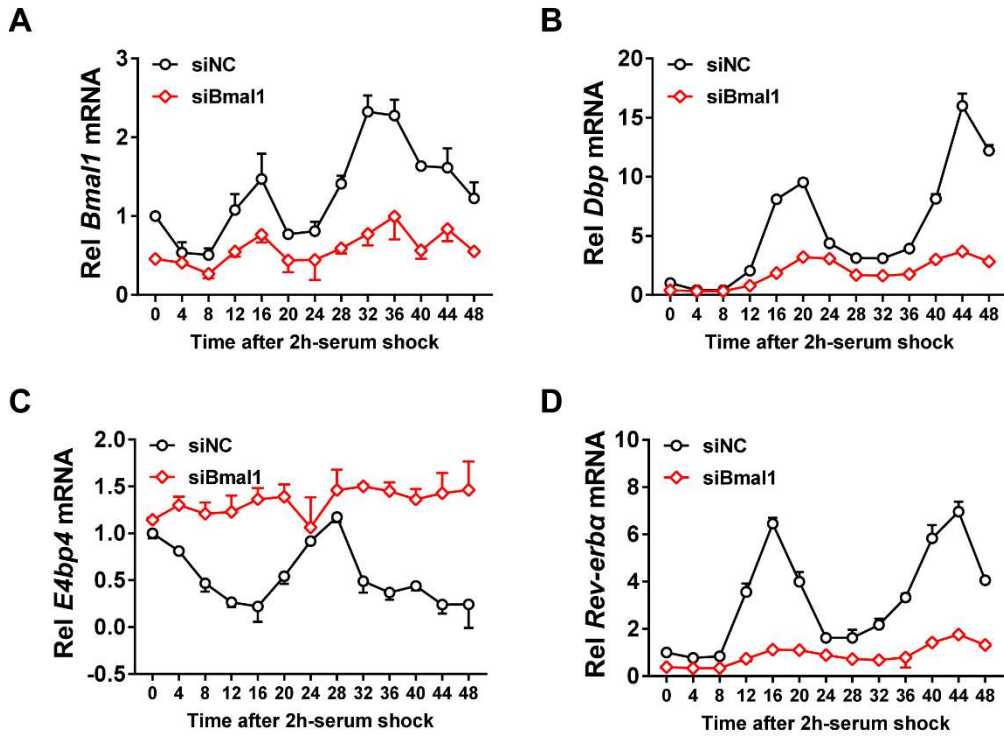
	<b>Forward (5'-3' sequence)</b>	<b>Reverse (3'-5' sequence)</b>
<b>qRT-PCR</b>		
mBmal1	CTCCAGGAGGCAAGAAGATTC	ATAGTCCAGTGGAAGGAATG
mE4bp4	CTTTCAGGACTACCAGACATCCAA	GATGCAACTTCCGGCTACCA
mRev- <i>crba</i>	TTTTTCGCCGGAGCATCCAA	ATCTCGGCAAGCATCCGTTG
mDbp	ACATCTAGGGACACACCCAGTC	AAGTCTCATGGCCTGGAATG
mHlf	GTGACGACTCCTGCTCCA	CAGCCATACTCCATTCCAAC
mTef	GCATCGCCACCATCTTCTT	GCTGCCACATTGTTCTTCTT
mMrp2	GTGTGGATTCCCTTGGGCTTT	CACAACGAACACCTGCTTGG
mAbcc1	CATGTGGACGTGTTTCGA	CACCATCATCCCTGTAATC
mAbcc3	CTGGGTCCCCTGCATCTAC	GCCGTCTTGAGCCTGGATAAC
mSlc19a1	GCCACAGAACCCCTCCTTA	GCATAGTTGAGCCCGACA
mSlc46a1	TCGGGTCTTTCGTGCTTG	GGCTCCTCACCCTCTCA
mFolr1	CCTGGAATGAAGAAAGCA	AAATACAACCCACCTACTCG
Humbs	CCGAGCCAAGGACCAGGATA	CCGAGCCAAGGACCAGGATA
BMAL1	AAATCGCTTTGAGGTGAC	CTTTCGTTTGGCGTTGC
MRP2	CCATAGCTTCATTCCTGAGTAGC	TCAGAGGACGCTTGTAGCCTT
GAPDH	CATGAGAAGTATGACAACAGCCT	AGTCCTTCCACGATACCAAAGT
<b>EMSA</b>		
Mrp2	CTCACTGGGATGACATAGCATTTCATCT	AGATGAATGCTATGTCATCCCAGTGAG
Mrp2(mutant)	CTGACTGGGCTAGCGTCGTATTCATCT	AGATGAATACGACGCTAGCCCAGTCAG
<b>CHIP</b>		
Mrp2	ACTTCCGACCTGCCAACT	CCATCCTCACTGCCCAAT
Mrp2(distal)	ACTTTGCTCTGGGTGTCT	AGGGTTAAGGAAGGGTTT
<b>siRNA</b>		
siE4bp4	GCACAAGCUUCGGAUUAAT	UUUAAUCCGAAGCUUGUGCTT
siRev- <i>crba</i>	CUUCGUUGUUAACGUGAATT	UUCACGUUGAACAAACGAAGTT
siBmal1	GCUCUUUCUUCUGUAGAAUTT	AUUCUACAGAAGAAAGAGCTT
siBMAL1	CCGAGGGAAGAUACUCUUUTT	AAAGAGUAUCUCCCCUCGGTT
siNC	UUCUCCGAACGUGUCACGUTT	ACGUGACACGUUCGGAGAATT



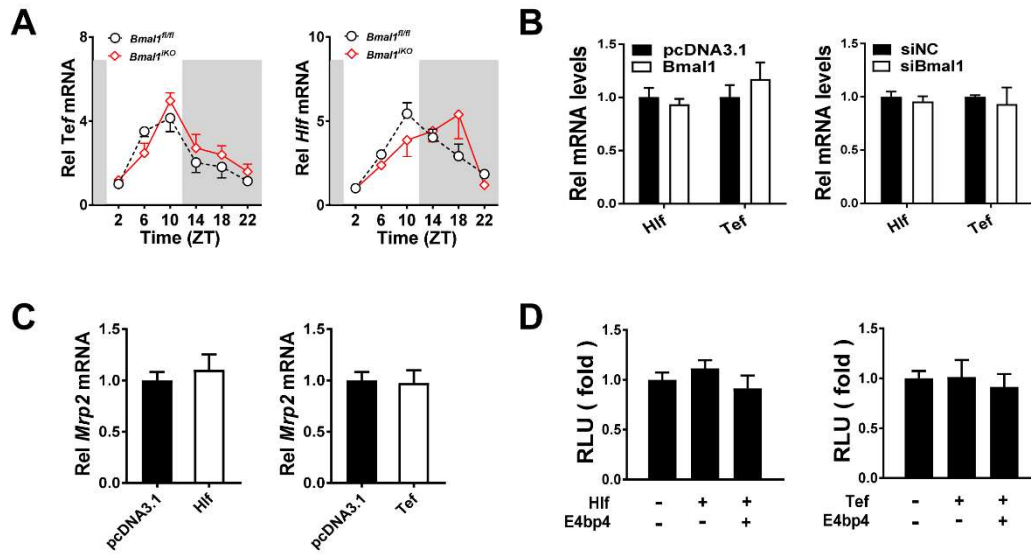
**Supplementary Figure 1. (A)** qRT-PCR analysis of *Bmal1* in the duodenum, jejunum, ileum and colon of *Bmal1<sup>fl/fl</sup>* and *Bmal1<sup>iKO</sup>* mice. **(B)** qRT-PCR analysis of *Mrp2* in the duodenum, jejunum, ileum and colon of *Bmal1<sup>fl/fl</sup>* and *Bmal1<sup>iKO</sup>* mice. Data are mean  $\pm$  SD ( $n = 6$ ). Statistics were performed with Student's t test ( $*p < 0.05$ ).



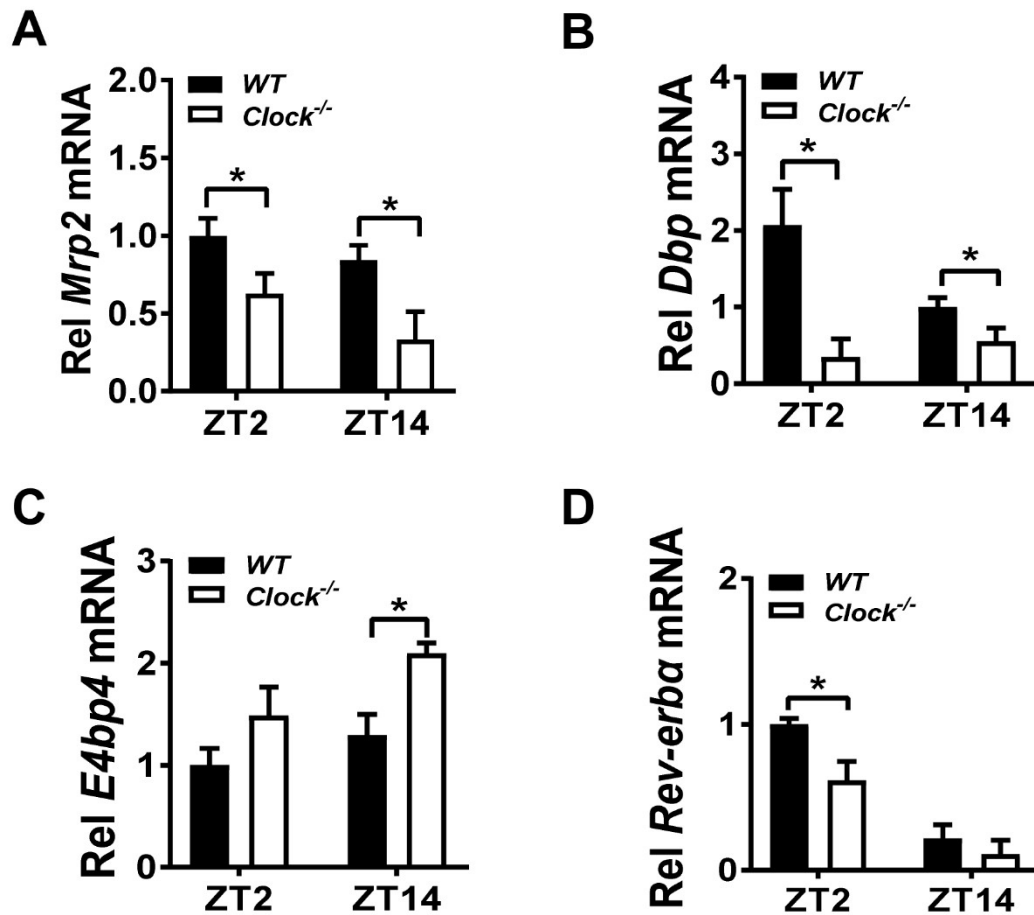
**Supplementary Figure 2. (A)** The liver accumulation of MTX at 0.25 and 0.75 h post oral administration (50 mg/kg) at dosing time of ZT2 and ZT14 in *Bmal1<sup>fl/fl</sup>* and *Bmal1<sup>iKO</sup>* mice. **(B)** The kidney accumulation of MTX at 0.25 and 0.75 h post oral administration (50 mg/kg) at dosing time of ZT2 and ZT14 in *Bmal1<sup>fl/fl</sup>* and *Bmal1<sup>iKO</sup>* mice. Data are mean  $\pm$  SD ( $n = 5$ ). Statistics were performed with Two-way ANOVA and Bonferroni post hoc test ( $*p < 0.05$ ).



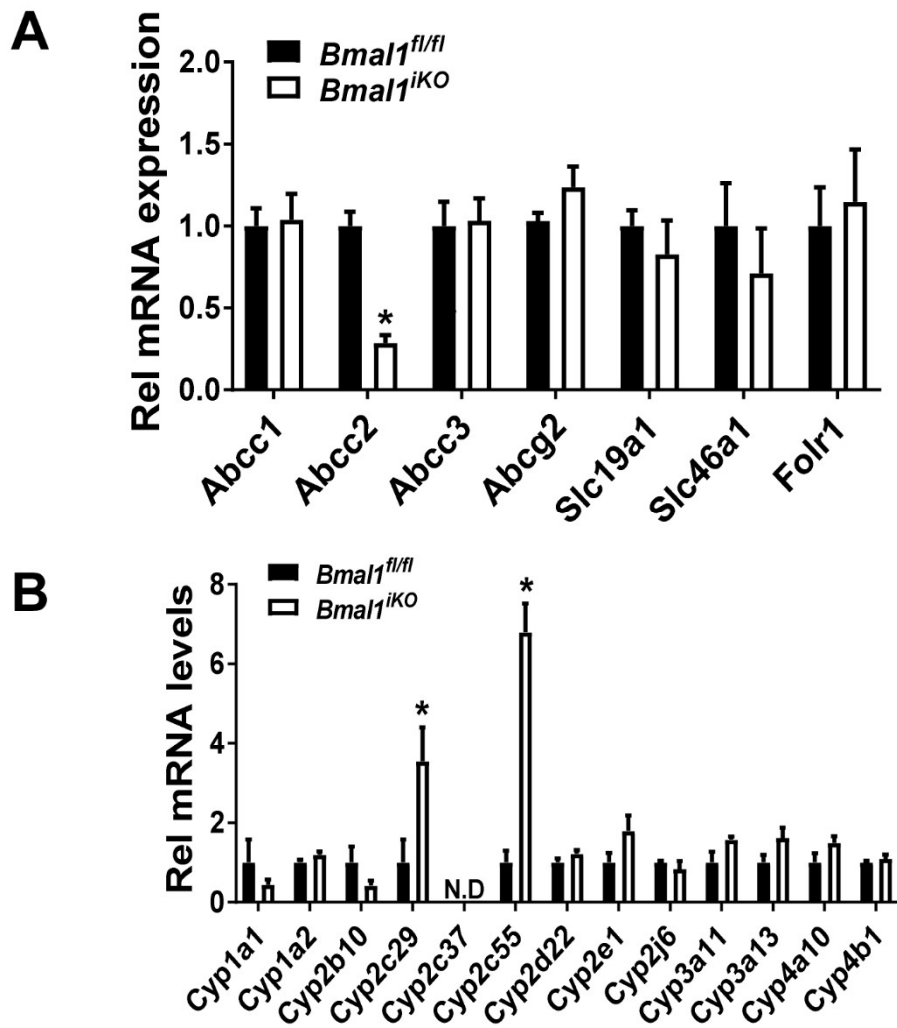
**Supplementary Figure 3.** (A) qRT-PCR analysis of Bmal1 in serum-shocked CT26 cells treated with siNC or siBmal1. Data are mean  $\pm$  SD ( $n = 6$ ). (B) qRT-PCR analysis of Dbp in serum-shocked CT26 cells treated with siNC or siBmal1. Data are mean  $\pm$  SD ( $n = 6$ ). (C) qRT-PCR analysis of E4bp4 in serum-shocked CT26 cells treated with siNC or siBmal1. Data are mean  $\pm$  SD ( $n = 6$ ). (D) qRT-PCR analysis of Rev-erba in serum-shocked CT26 cells treated with siNC or siBmal1. Data are mean  $\pm$  SD ( $n = 6$ ). Statistics were performed with Two-way ANOVA ( $*p < 0.05$ ).



**Supplementary Figure 4. (A)** qRT-PCR analysis of *Tef* and *Hlf* mRNA in small intestine of *Bmal1<sup>fl/fl</sup>* and *Bmal1<sup>iKO</sup>* mice. **(B)** Overexpression or knockdown of *Bmal1* in CT26 cells had no effect on *Tef* and *Hlf* mRNA expression. **(C)** Overexpression of *Tef* or *Hlf* in CT26 cells had no effect on *Mrp2* mRNA expression. **(D)** Overexpression of *Tef* or *Hlf* had no effect on *Mrp2* promoter activity. NIH3T3 cells were transfected with *Mrp2*-Luc reporter and *Tef* or *Hlf* expression plasmids. After 24 h treatment, luciferase reporter activities were measured. Data are mean  $\pm$  SD ( $n = 6$ ).



**Supplementary Figure 5. (A)** Intestinal *Mrp2* mRNA levels in WT and *Clock*<sup>-/-</sup> mice at ZT2 and ZT14. **(B)** Intestinal *Dbp* mRNA levels in WT and *Clock*<sup>-/-</sup> mice at ZT2 and ZT14. **(C)** Intestinal *E4bp4* mRNA levels in WT and *Clock*<sup>-/-</sup> mice at ZT2 and ZT14. **(D)** Intestinal *Rev-erba* mRNA levels in WT and *Clock*<sup>-/-</sup> mice at ZT2 and ZT14. Data are mean ± SD ( $n = 6$ ). Statistics were performed with Two-way ANOVA and Bonferroni post hoc test ( $*p < 0.05$ ).



**Supplementary Figure 6.** (A) The mRNA levels of transporters involved in intestinal disposition of MTX in small intestine of *Bmal1<sup>fl/fl</sup>* and *Bmal1<sup>iKO</sup>* mice. (B) Intestinal mRNA levels of CYPs in *Bmal1<sup>fl/fl</sup>* and *Bmal1<sup>iKO</sup>* mice. Data are mean  $\pm$  SD ( $n = 6$ ). Statistics were performed with Student's t test ( $*p < 0.05$ ).