

Supplemental Information

**Circadian clock gene Bmal1 regulates bilirubin detoxification: a potential
mechanism of feedback control of hyperbilirubinemia**

Supplementary Table 1. Mouse primer sequences for quantitative real-time PCR (qPCR)

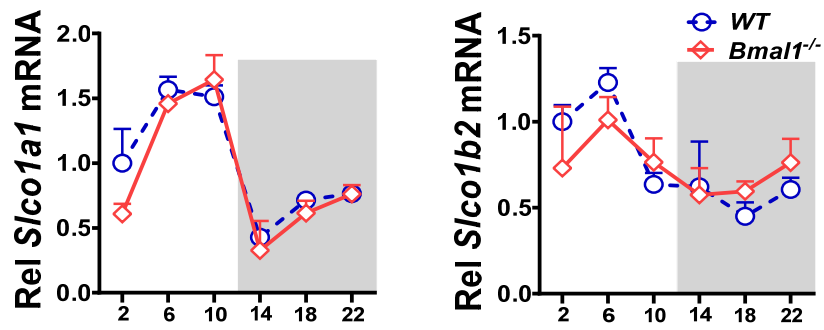
Gene	Forward (5'-3' sequence)	Reverse (5'-3' sequence)
<i>Bmal1</i>	CTCCAGGAGGCAAGAAGATTC	ATAGTCCAGTGAAGGAATG
<i>Ugt1a1</i>	CACTGGCTGAGTATGCTTGG	CTTCTGGAATGGCACAGGGAA
<i>Mrp2</i>	AGCCGACCTGCCTCTTTGTT	CAAGAGCCAAAGAAAGCCCAA
<i>Mrp3</i>	CTGCCCCAGACGGACTTTAT	TCTGGTGCGTAGTTTCGGAG
<i>Sloc1a1</i>	CAGGGGCATGCAGGATGTAT	GTCTGGAGAGTGGATGTCGC
<i>Sloc1b2</i>	CACCGACCAAAGCTGATTGG	AGAAATGTGGCAACGCAGTC
<i>Homx1</i>	CCTTCTGGTATGGGCCTCAC	CCTCGTGGAGACGCTTTACA
<i>Blvra</i>	ACGAGAACTTGGGAGCCTTG	AGGCGACATCAACCTCTTGG
<i>E4bp4</i>	CTTTCAGGACTACCAGACATCCAA	GATGCAACTTCCGGCTACCA
<i>Pck1</i>	ATGAAAGGCCGCACCATGTA	GCACAGATATGCCCATCCGA
<i>G6pase</i>	TGGTAGCCCTGTCTTTCTTTG	TTCCAGCATTACACTTTCCT
<i>Cyp4a14</i>	CCCAAAGGTATCACAGCCACAA	CAGCAATTCAAAGCGGAGCAG
<i>Ppib</i>	CTCTCGGAGCGCAATATGAAG	GCAAAGGAAGACGACGGAG

Supplementary Table 2. Primer sequences for ChIP assays.

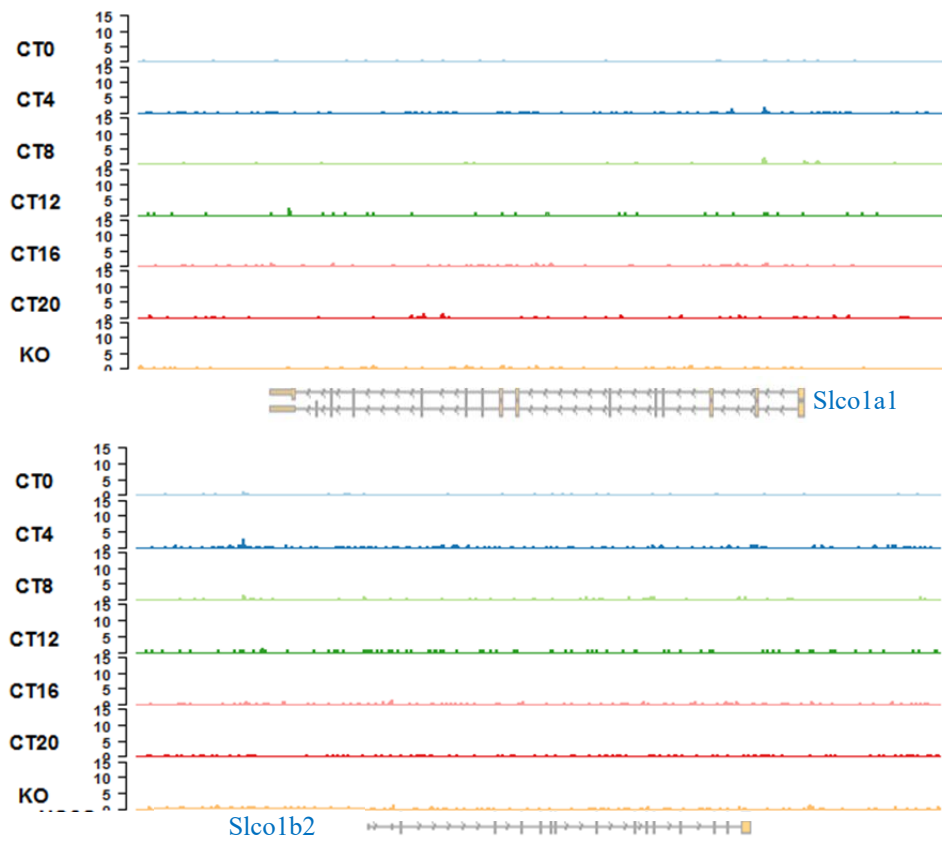
Gene	Forward (5'-3' sequence)	Reverse (5'-3' sequence)
Ugt1a1-E-box	TCATGGGGCACAATCAATCA	ACGGAAGAAGCAGAAGCAAAC
Ugt1a1-nonspecific region	GCTGCCATCTTGAAACCCAG	AATCTCGGGTCCTTATGCTTGC
Mrp2-E-box-1	ACATCTCTGTGAACTCTTAACCG	TAATGCAGTTGGCAGGTCGG
Mrp2-E-box-2	GAGAAGTGATAAGCCTCCAGT	GAATGCTATGTCATCCCAGTGAGC
Mrp2-nonspecific region	GTTTCGCTATCACATTTCA	CTCATTGCCTTCACTTCA

Supplementary Table 3. Oligonucleotide sequences for EMSA assays

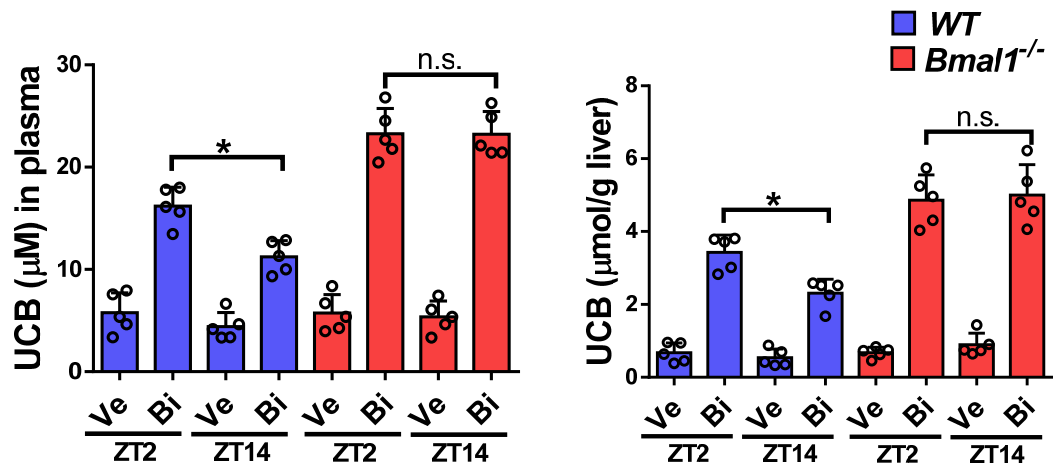
Oligonucleotide	Forward (5'-3' sequence)	Reverse (5'-3' sequence)
Mrp2-E-box1(biotin)	TAAAACCCACATGCAGGGAT	ATCCCTGCATGTGGGGTTTTA
Mrp2-E-box1(cold)	TAAAACCCACATGCAGGGAT	ATCCCTGCATGTGGGGTTTTA
Mrp2-E-box1(mut)	TAAAACCCTGTGCACAGGGAT	ATCCCTGTGCACAGGGTTTTA
Mrp2-E-box2(biotin)	TCCCAGTCACATGTCTGCTCA	TGAGCAGACATGTGACTGGGA
Mrp2-E-box2(cold)	TCCCAGTCACATGTCTGCTCA	TGAGCAGACATGTGACTGGGA
Mrp2-E-box2(mut)	TCCCAGTAGTCTTCTGCTCA	TGAGCAGAAGCACTACTGGGA
Ugt1a1-E-box(biotin)	TTCGGATGTTCCAGCTGCCTTTATCTCC	GGAGATAAAGGCAGCTGGAACATCCGAA
Ugt1a1-E-box(cold)	TTCGGATGTTCCAGCTGCCTTTATCTCC	GGAGATAAAGGCAGCTGGAACATCCGAA
Ugt1a1-E-box(mut)	TTCGGATGTTTCATCGATCCTTTATCTCC	GGAGATAAAGGATCGATGAACATCCGAA



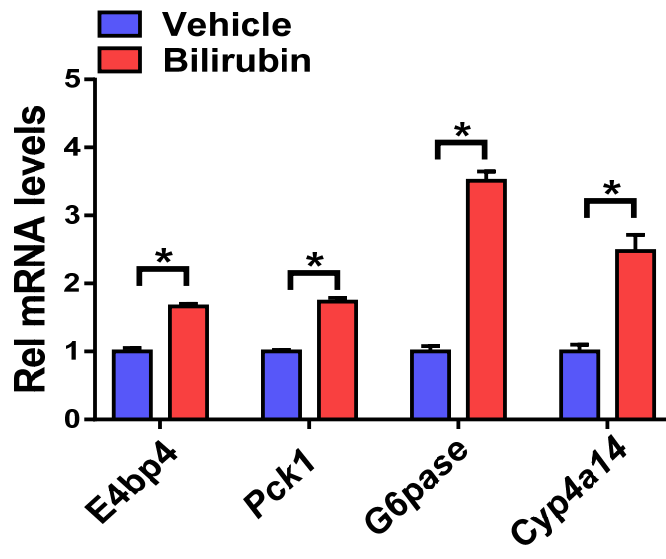
Supplementary Figure 1. Bmal1 ablation does not affect *Slco1a1* and *Slco1b2* expressions. (A) Circadian mRNA levels of *Slco1a1* and *Slco1b2* in the livers of WT and *Bmal1*^{-/-} mice measured by qPCR. Data are mean ± SD (n=5).



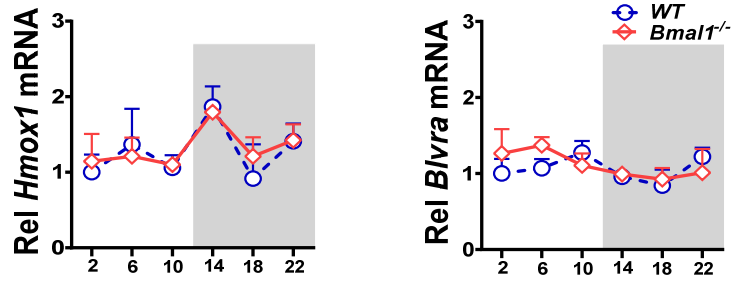
Supplementary Figure 2. ChIP-sequencing for circadian Bmal1 binding to *Slco1a1* and *Slco1b2*. No obvious peaks were found in the data. ChIP-sequencing traces were generated from GSE39977.



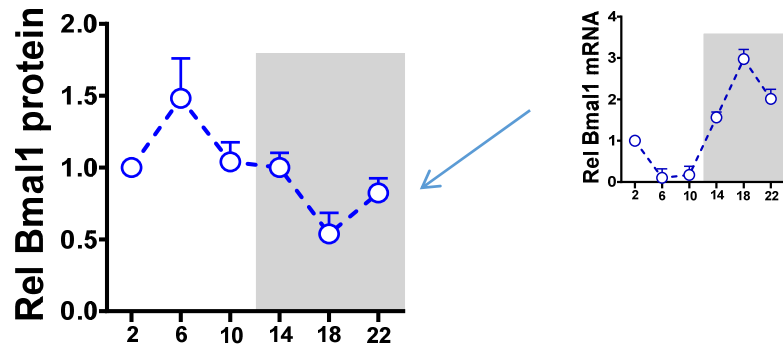
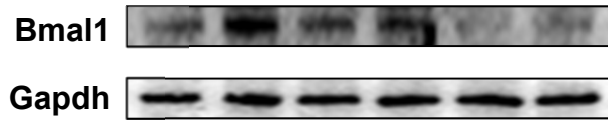
Supplementary Figure 3. Measurements of UCB in plasma and livers of bilirubin or vehicle-treated mice at ZT2 or ZT14. Data are mean \pm SD ($n=5$). * $P < 0.05$ (t test). Ve: vehicle; Bi: bilirubin.



Supplementary Figure 4. qPCR measurements of mRNA levels of *E4bp4*, *Pck1*, *G6pase* and *Cyp4a14* in Hepa1-6 cells. Data are mean \pm SD (n=5). *P < 0.05 (t test).

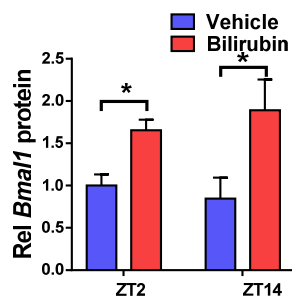


Supplementary Figure 5. *Bmal1* ablation does not affect *Hmox1* and *Blvra* expressions. (A) Circadian mRNA levels of *Hmox1* and *Blvra* in the livers of WT and *Bmal1*^{-/-} mice measured by qPCR. Data are mean \pm SD ($n=5$).

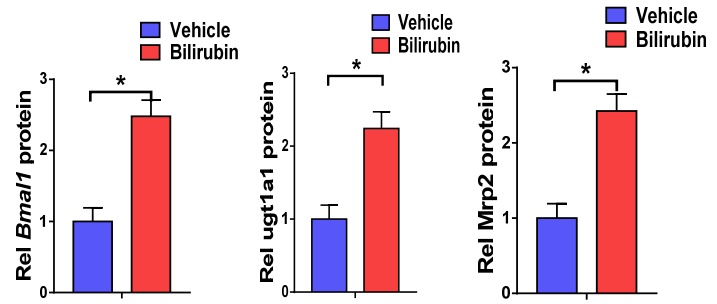


Supplementary Figure 6. Circadian protein (left panel) and mRNA (right panel) expression of Bmal1 in the livers of Wild-type mice.

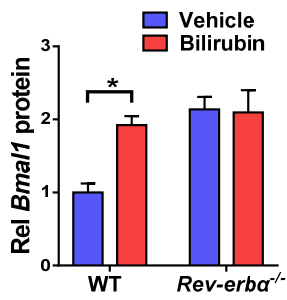
6-B



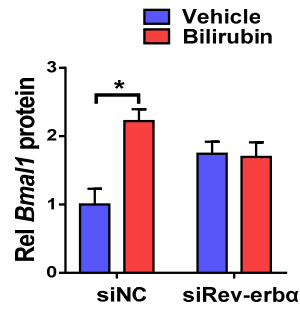
6-E



6-J



6-K



Supplementary Figure 7. Quantification of the blot band density for Figure 6.