| Gene | Direction | Primer sequence (5' to 3') | | |
|----------|-----------|----------------------------|--|--|
| Cirbp | Forward | GTGGGCGATGACATTGCC | | |
| | Reverse | ACTGAGGTGCCAGGAGCAAC | | |
| Fgf21 | Forward | ATGGAATGGATGAGATCTAGAGTTGG | | |
| | Reverse | TCTTGGTGGTCATCTGTGTAGAGG | | |
| Pdk4 | Forward | CACATGCTCTTCGAACTCTTCAAG | | |
| | Reverse | TGATTGTAAGGTCTTCTTTTCCCAAG | | |
| Hmgcs2 | Forward | ACCTGCTCACCTGCTCTCAGCAG | | |
| | Reverse | GGTCAGTTTGGTCCACATATTGG | | |
| Bmal1 | Forward | ACGACATAGGACACCTCGCAGA | | |
| | Reverse | CGGGTTCATGAAACTGAACCATC | | |
| Clock | Forward | ACCACAGCAACAGCAACAAC | | |
| | Reverse | GGCTGCTGAACTGAAGGAAG | | |
| Per3 | Forward | TGCGCTGTGTACACAGTGTGC | | |
| | Reverse | CAGTCACATCTGCCTGGTGTG | | |
| Per2 | Forward | CACTCAGGAGTGCATGGAGGAGA | | |
| | Reverse | CTGCTCTTGCACCTTGACCAGGT | | |
| Dbp | Forward | GGAACTGAAGCCTCAACCAAT | | |
| | Reverse | CTCCGGCTCCAGTACTTCTCA | | |
| 18S rRNA | Forward | GTAACCCGTTGAACCCCATT | | |
| | Reverse | CCATCCAATCGGTAGTAGCG | | |

Supplemental Table 1. Primer sequences for quantitative real-time PCR



Supplemental Figure 1. Temporal mRNA expression profiles of *Cirbp* and circadian clock-related genes in the heart of KD mice.

Mice were fed with normal (ND; unfilled circles) or ketogenic (KD; filled circles) diets for 14 days under LD 12:12 (lights on at 0 h) and then total RNA was extracted from the heart after sacrifice. Messenger RNA levels were measured by quantitative RT-PCR. Maximal value for ND mice is expressed as 100%. Values are shown as means \pm SEM (n = 4). Significant differences between groups are indicated as **P* < 0.05, ***P* < 0.01. Horizontal open and solid bars, day and night, respectively.

| | | MESOR | Amplitude | Acrophase (h) |
|-------|----|------------------|----------------|-------------------|
| Bmal1 | ND | 42.7 ± 4.7 | 51.5 ± 6.6 | 0.76 ± 0.49 |
| | KD | 39.2 ± 4.5 | 47.3 ± 6.4 | 18.6 ± 0.51** |
| Clock | ND | 68.6 ± 2.1 | 27.4 ± 3.0 | 1.6 ± 0.41 |
| | KD | $59.7 \pm 3.0^*$ | 29.3 ± 4.3 | 18.3 ± 0.56** |
| Per3 | ND | 52.4 ± 3.0 | 41.7 ± 4.2 | 11.3 ± 0.39 |
| | KD | 49.4 ± 1.3 | 34.1 ± 1.9 | 6.5 ± 0.21 ** |
| Per2 | ND | 44.9 ± 3.9 | 44.5 ± 5.5 | 14.1 ± 0.47 |
| | KD | 84.5 ± 5.1** | 49.4 ± 7.2 | 8.5 ± 0.55** |
| Dbp | ND | 42.8 ± 4.5 | 47.3 ± 6.3 | 10.6 ± 0.51 |
| | KD | 22.1 ± 3.6** | 25.9 ± 5.1* | 4.6 ± 0.75** |

Supplemental Table 2. Cosinor analysis of *Bmal1*, *Clock*, *Per3*, *Per2*, and *Dbp* mRNA expression in the heart of mice fed with a ketogenic diet.

KD, ketogenic diet; ND, normal diet. MESOR, mean statistics of rhythm; Amplitude, one-half the total peak-trough variation; Acrophase, delay from 0:00 (lights on). Data are shown as means \pm SEM (n = 4). Significant differences compared with ND value are indicated as **P* < 0.05, ***P* < 0.01.