



Figure S1. Comparison of locomotor activity between two control lines: *Bmal1^{fl/fl}* and *Nkx2.1-Cre;Bmal1^{+/fl}* and *Nkx2.1-Cre;Bmal1^{fl/fl}* (*Nkx2.1-Bmal1^{-/-}*). (A, B) Daily profiles of locomotor activity in LD or DD shown with absolute counts (A) or expressed as percentage of daily total (B). (C, D) Mean free-running period and amplitude (Qp values of periodogram) in DD. Values for activity in DD were calculated for data on Days 5–19 in DD. (E) Activity during 12-h light phase (Light), 12-h dark phase (Dark), 24-h day in LD (Total), and a circadian period in DD (Total/DD). Values are mean \pm SEM. $n = 6$ for *Bmal1^{fl/fl}*, $n = 8$ for *Nkx2.1-Cre;Bmal1^{+/fl}*, $n = 11$ for *Nkx2.1-Cre;Bmal1^{fl/fl}* mice. $F(2,22) = 0.919$, $p = 0.414$ (C); $F(2,22) = 12.41$, $p < 0.001$ (D); $F(2,22) = 14.07$, $p < 0.001$ (E-Light); $F(2,22) = 19.51$, $p < 0.001$ (E-Dark); $F(2,22) = 27.09$, $p < 0.001$ (E-Total); $F(2,22) = 34.20$, $p < 0.001$ (E-Total/DD) by an one-way repeated measures ANOVA; *, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$ as compared to *Bmal1^{fl/fl}* by a Tukey-HSD post-hoc test. The p values calculated by a two-way repeated measures ANOVA for the comparison of Light and Dark in (E) are: the effect of genotype, $F(2,22) = 24.38$, $p < 0.001$; the effect of time, $F(1,22) = 397.22$, $p < 0.001$; and the interaction between genotype and time, $F(2,22) = 8.50$, $p = 0.002$.