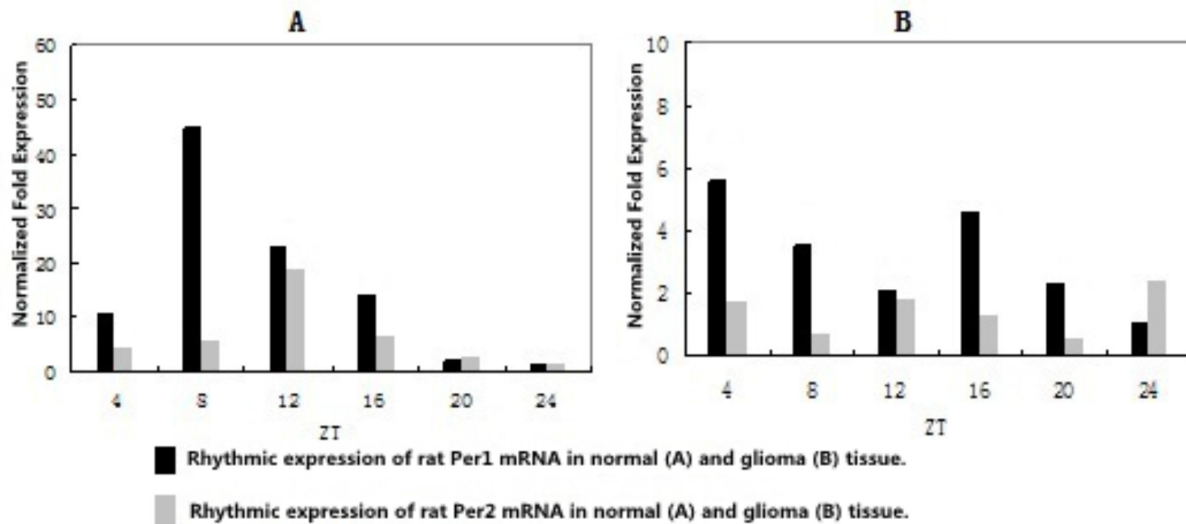


SUPPLEMENTARY FIGURE



Supplementary Figure S1: Rhythmic expression of rat *Per1* and *Per2* mRNA in normal (A) and glioma (B) tissue. Total RNA was extracted from the SCN, and real-time RT-PCR was used to determine *Per1* and *Per2* mRNA levels. Relative expression of each mRNA was normalized to the corresponding level of β -actin mRNA. Data are the mean and standard deviation of three independent experiments.