

Gene	GeneBank accession number	Primers for qPCR		Product size [bp]	Primers for dsRNA synthesis		Product size [bp]
		Sense	Antisense		Sense	Antisense	
<i>per</i>	EF364034.1	CCGAACGAGGTGGTGG TAAA	GCCAAGGTACGGAACA GACT	99	CGCAACACCTTTGCATC ACA	TTTACCACCACCTCGTT CGG	299
<i>tim</i>	FQ029930.1	GAAGTCAAGCAGGCG TCTA	AACTTGCCTGGAGTCC ACAG	94	GGAGACCTGCTTCGCT AAGAT	CTGCGGTCCAGAAATG AGGT	305
<i>cry2</i>	EF396286.1	CAAGTGTATGCGCAAC TCGC	TTCGGACTTGGTGACG ACTG	98	GAGGTGCGATGTGGGT AACT	TGCAGCAGGAACCTCC ATTT	300
<i>cyc</i>	ON052028	GTGGCCAACAACACGA TCGT	ACCTCCCTCTGCTTGT CATGT	92	GTGGCCAACAACACGA TCGT	TTCTGGTTGAAGACGG CGTC	290
<i>pdp1</i>	ON052027	CCCCTCGAAGACAAG AAAGAT	ACTTGAGATCGGGGTC ATAGGG	107	TCTCAAGTACGCGGAC CTAGAC	CGGGACTTCTTGATCA TTGGCT	300
<i>EF1-A</i>	HQ177154.1	CGTACTCATTGTCGCC GCTG	CGAGTGTGAAAGCGAG CAGA	95	-	-	-
<i>SSU</i>	M32419.1	-	-	-	GCCTCATTCCCTGTTTC AAG	AGCAAGGAACCCATCC ATTT	292

Table S1. Information about the starters used in the experiments. The primers shown on the two columns on the left (orange) were used to study the daily expression profile of selected clock genes—*period* (*per*), *timeless* (*tim*), *cryptochrome 2* (*cry2*), *cycle* (*cyc*), *pdp1* (*PAR domain protein 1*)—and the *EF1-A* (*Elongation factor 1-alpha*) gene, which was used as a housekeeping standard for normalizing qRT-PCR results. The primers shown on the two columns on the right (purple) were used for *in vitro* synthesis (transcription) to generate dsRNAs used to knock out the expression of the same biological clock genes when injected to the larvae; primers for amplifying *SUU* (*ribulose-1,5-bisphosphate carboxylase small subunit*) of cultivated tobacco were used to produce non-specific (plant) dsRNA which was applied to the larvae in the control group. In addition, the GenBank accession numbers for studied genes and the sizes of products obtained from qRT-PCR and *in vitro* transcription (production of dsRNAs) are shown.