

Abstract



Searching for Light-Induced Genes in Hymenopteran Insect – Nasonia vitripennis: Transcription of 13% of the Genes Is Regulated by Light ⁺

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Hymenopteran insects, including bees, wasps, and ants, are crucial species of our ecosystems and fulfill valuable ecological roles in nature, as well as in agricultural and industrial activities. With recent reports on dramatic global declines in population size for both bees and wasps, there is an urgent need to understand their circadian system, since it is essential for orientation, foraging, and anticipation of the environment (Eban-Roth-schild and Bloch, 2011).

A key component in circadian system regulation is light. In *Drosophila*, the light input pathway involves the light-sensitive protein – cryptochrome (CRY) by inducing protein degradation (Katz and Minke, 2009). In mammals, a non-light sensitive CRY is present and the light input pathway is through CREB by inducing *period* (PER) expression (Chauhan et al., 2017). However, for other insect species like hymenopterans, that possess only non-light sensitive CRY, the mechanisms behind circadian light entrainment remain unclear.

We hypothesize that there may be light-driven circadian gene induction in hymenopteran insects, similar to what has been reported for mammals. To investigate this hypothesis, we have subjected *Nasonia* to different light conditions and conducted an RNAsequencing experiment to identify any light-induced gene expression. Our results provide a better understanding of the molecular mechanisms underlying the circadian light input pathway in hymenopterans as well as insight into the possible disruption of circadian entrainment by artificial light in hymenopteran insects.

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