

Uncovering Photoperiodic Regulation Mechanisms in *Neurospora crassa*

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Abstract

Organisms adapt to seasonal changes driven by Earth's rotation around the sun, with photoperiod influencing their biological rhythms. While photoperiodism has been more extensively studied in plants, the mechanisms underlying seasonal responses in fungi remain largely unexplored. This study explores novel mechanisms and identifies genetic components involved in photoperiodic regulation in the model fungus *Neurospora crassa*, focusing on how it adapts reproductive cycles to changes in day length. We tested whether fungi share a general photoperiodic mechanism with flowering plants, which we termed the signal accumulation model. We also tested two alternative hypotheses: the balance model and the gating model. To examine these, we developed a systematic assay to monitor protoperithecia (female organ) formation under varying photoperiods, referred to as the Protoperithecia Assay (PPA), using protoperithecia development as a proxy for the organism's photoperiodic recognition. Interestingly, the data support a balance model, where day length is measured through a dynamic equilibrium between light and dark phases, rather than through signal accumulation. We also observed varying degrees of system sensitivity depending on the time of day, lending support to the gating model. To identify genetic components involved in this fungal day-length sensing, we screened approximately 200 *N. crassa* mutants targeting genes involved in signal transduction, under different photoperiodic conditions. Understanding this unique fungal mechanism may provide broader insights into how non-photosynthetic organisms use ambient light as a seasonal cue.

Background

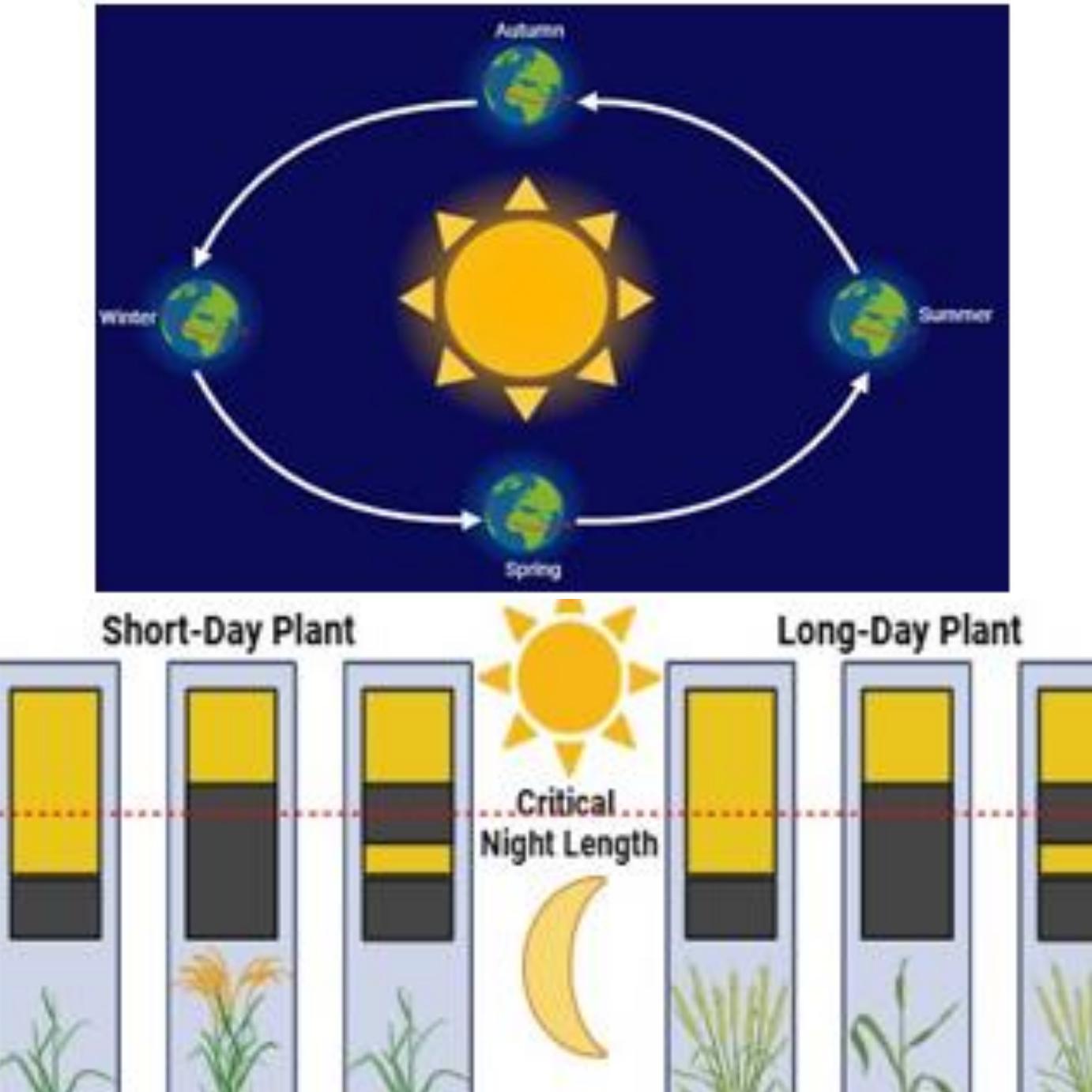


Figure 1. Photoperiodic Regulation. A. Earth's Rotation Around the Sun. B. The Photoperiodic recognition in fungi. Mechanism in Short-Day and Long-Day Plants

Methods and Materials

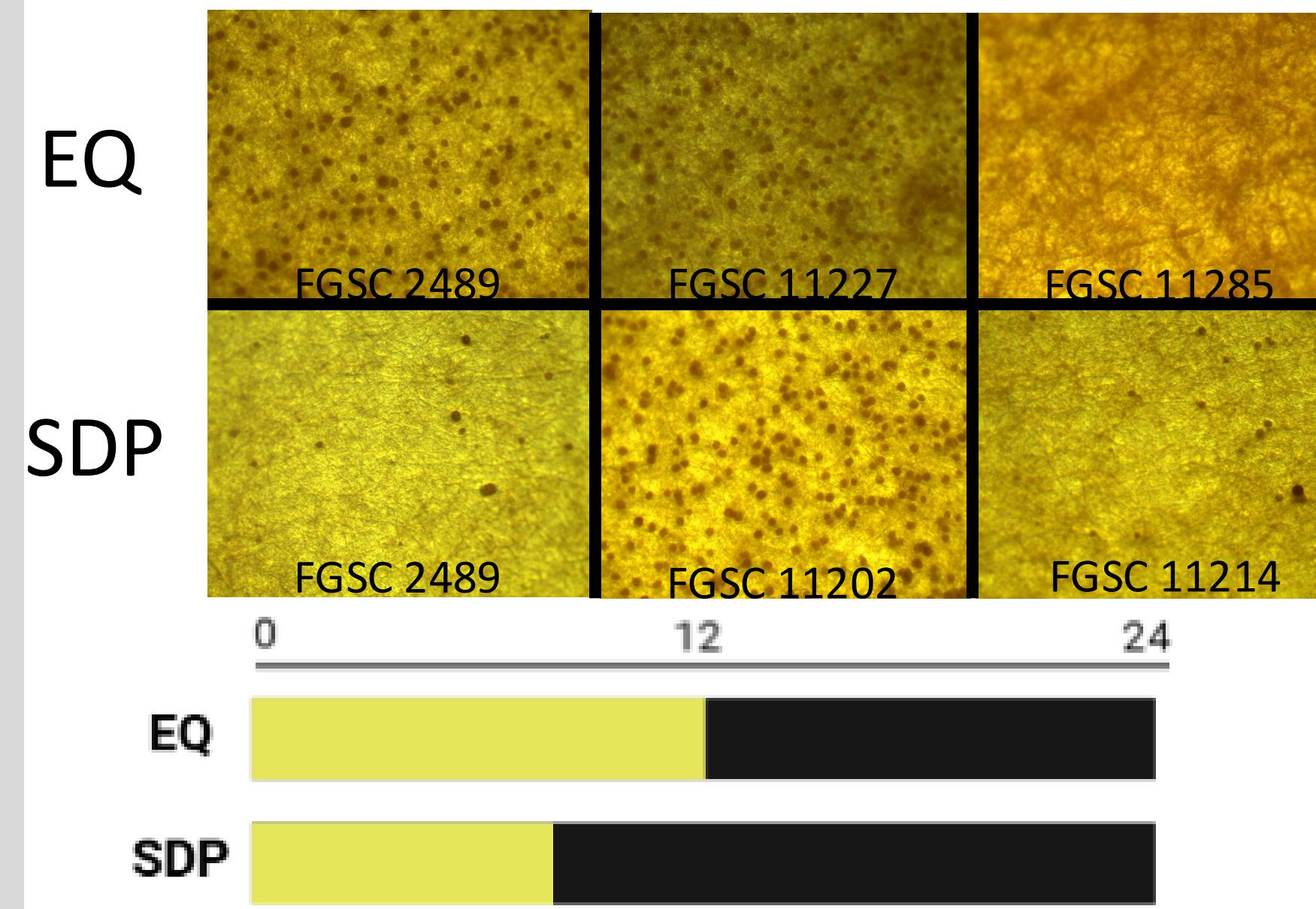


Figure 5. Light Conditions Model

Results

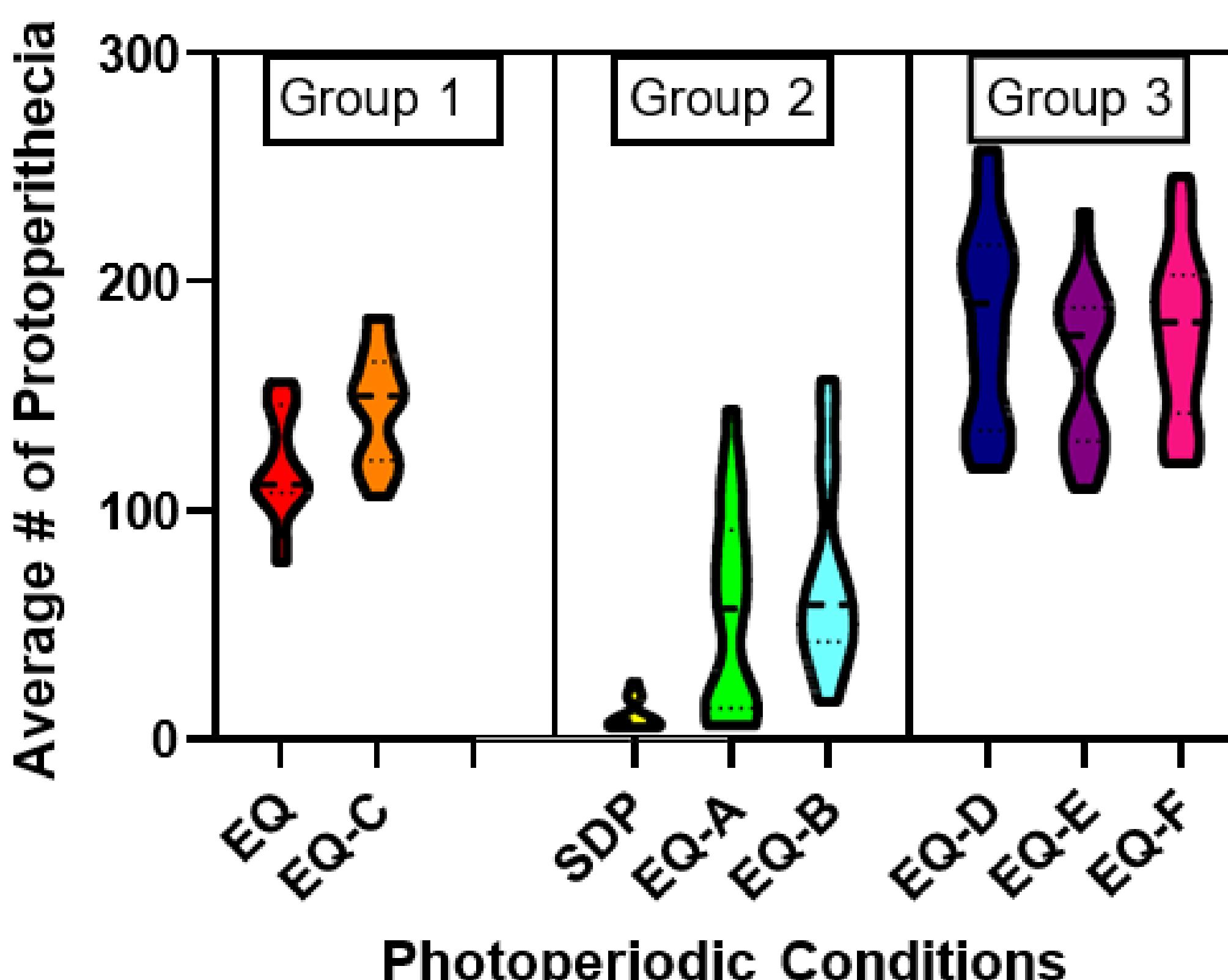
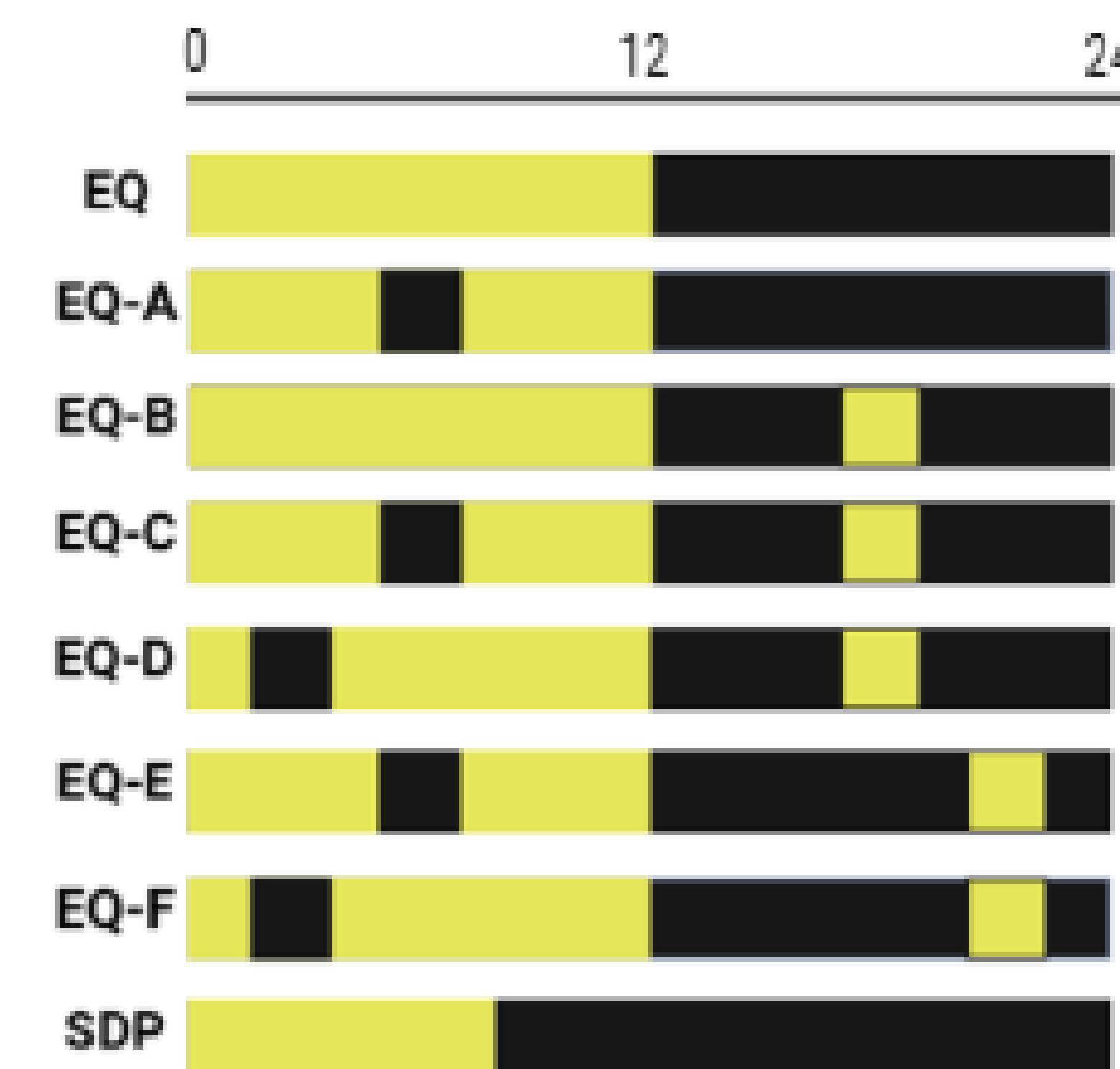
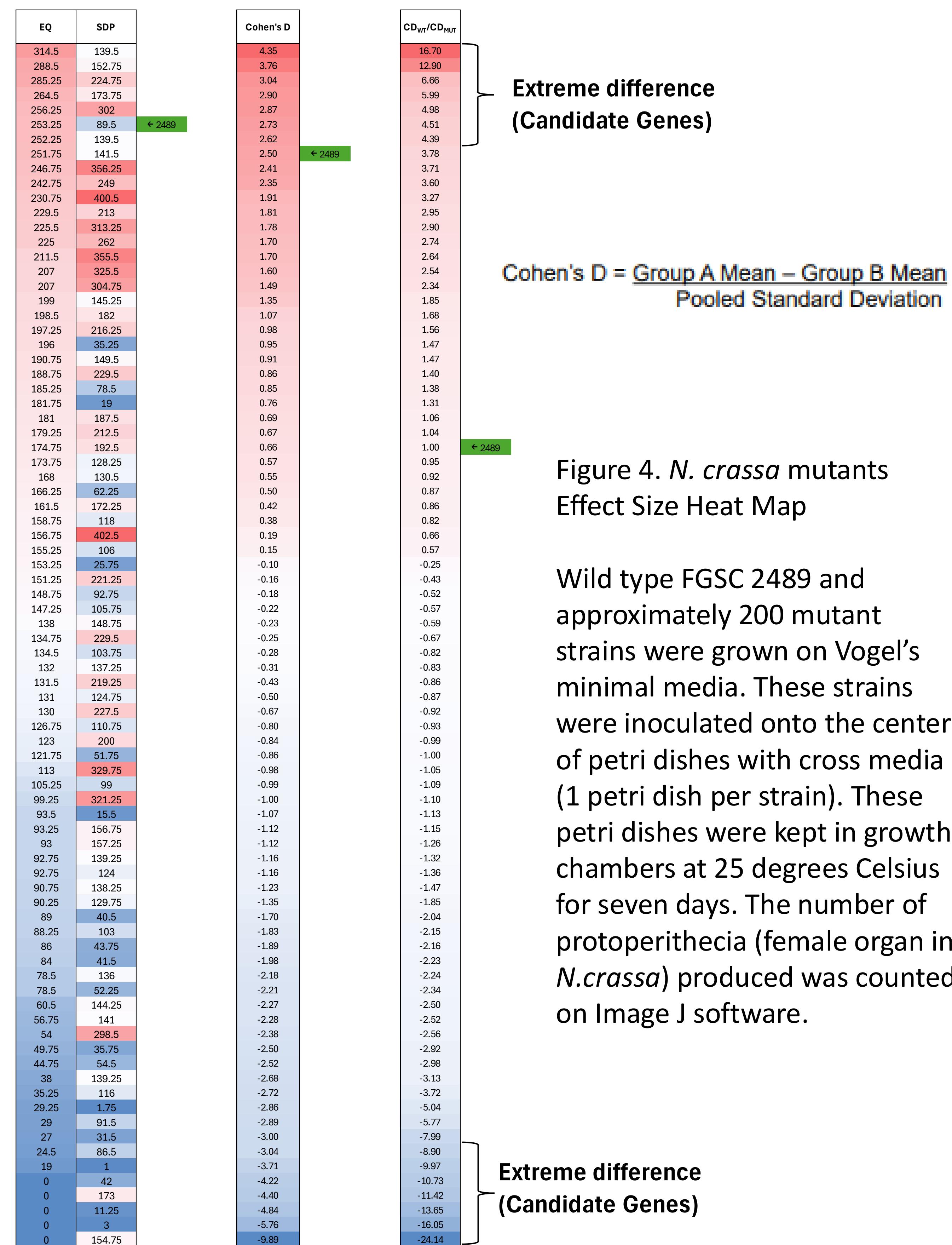


Figure 3. EQ Interruption Balance Model Experiment. Equinox (EQ), short- day (SDP), equinox-A (EQ-A), equinox-B (EQ-B), equinox-C (EQ-C), equinox- D (EQ-D), equinox-E (EQ-E), equinox- F (EQ-F). Group 1 exhibits an optimal production of protoperithecia, Group 2 exhibits a decrease in protoperithecia production, Group 3 exhibits an increase in protoperithecia production, nested one-way ANOVA test, p-value < 0.05, n=4.



Cohen's D = $\frac{\text{Group A Mean} - \text{Group B Mean}}{\text{Pooled Standard Deviation}}$

Figure 4. *N. crassa* mutants Effect Size Heat Map



Extreme difference (Candidate Genes)

Hypothesis

We hypothesize that candidate genes involved in light signal transduction mediate the balance model of photoperiodic regulation in *Neurospora crassa*, and their activity is selectively gated by circadian timing.

Conclusion

In conclusion, our study set out to uncover novel mechanisms by which fungi recognize and respond to photoperiods, with a particular emphasis on the molecular and genetic components underlying this regulation. Unlike plants, fungi appear to rely on distinct pathways for photoperiodic recognition, where the balance between light and dark phases—as well as the timing and duration of interruptions—plays a critical role. To explore these mechanisms, we screened approximately 200 *Neurospora crassa* mutants targeting genes involved in signal transduction and light perception. This approach led to the identification of several promising candidate genes that may contribute to photoperiodic regulation in fungi. These findings suggest that fungi possess unique signaling networks that interpret environmental light cues in ways not yet fully understood. As a future direction, we plan to genotype these candidate strains and perform functional analyses to validate their roles, ultimately deepening our understanding of the genetic architecture behind fungal photoperiodism and its broader implications for circadian biology.

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