Circadian and diel regulation of photosynthesis in the bryophyte *Marchantia*

polymorpha

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Abstract

Circadian rhythms are 24-hour biological cycles that align metabolism, physiology and

development with daily environmental fluctuations. Photosynthetic processes are governed

by the circadian clock in both flowering plants and cyanobacteria, but it is unclear how

extensively this is conserved throughout the green lineage. We investigated the contribution

of circadian regulation to photochemistry in *Marchantia polymorpha*, a liverwort that diverged

from flowering plants early in the evolution of land plants. First, we identified in *M.*

polymorpha the circadian regulation of photosynthetic biochemistry, measured using two

approaches (delayed fluorescence, PAM fluorescence). Second, we identified that light-dark

cycles synchronize the phase of 24 h cycles of photosynthesis in *M. polymorpha*, whereas

the phases of different thalli desynchronize under free running conditions. This might also be

due to masking of the underlying circadian rhythms of photosynthesis by light-dark cycles.

Finally, we used a pharmacological approach to identify that chloroplast translation might be

necessary for clock control of light harvesting in *M. polymorpha*. We infer that the circadian

regulation of photosynthesis are well-conserved amongst terrestrial plants.

Introduction

The rotation of the Earth on its axis causes 24-hour cycles in environmental conditions such

as light and temperature. These diel cycles are thought to have selected for the evolution of

circadian clocks within several Kingdoms of life (Dodd *et al.* 2005, Eelderink-Chen *et al.*

2021, Millar 2016, Ouyang *et al.* 1998, Spoelstra *et al.* 2016). Circadian rhythms are self-

sustaining biological cycles that have a period of about 24 h, which are thought to provide a

biological measure of the time of day. These rhythms coordinate and sequence processes

around the diel cycle, whilst ensuring that responses to environmental cues are appropriate

for the time of day.

Circadian rhythms in plants are generated by a molecular oscillator that is formed from a

series of transcription-translation feedback loops, which has a predominance of negative

feedback steps (Hsu and Harmer 2014). The oscillator is entrained to environmental cues

such as light and temperature, which align the phase of the rhythm with the day/night cycle

so that it provides an accurate biological measure of the time of day. There is circadian

regulation of a variety of key aspects of the physiology of flowering plants, such as stomatal

opening, light harvesting, CO_2 fixation and starch metabolism (Dakhiya and Green 2022,

Dakhiya *et al.* 2017, Dodd *et al.* 2005, Graf *et al.* 2010). Furthermore, the circadian oscillator

contributes to the photoperiodic control of flowering time (Hayama and Coupland 2003, Park

et al. 1999) and the fitness of plants (Dodd *et al.* 2005, Green *et al.* 2002). This means that

circadian regulation has a pervasive influence upon physiology, metabolism, and

development (Hotta *et al.* 2007, Millar 2016), reflecting the close functional relationship

between plants and their fluctuating environments.

Circadian regulation of photosynthesis has been reported in a variety of flowering plants,

including *Arabidopsis thaliana*, *Hordeum vulgare*, *Zea mays*, *Phaseolus vulgaris*, *Kalanchoë*

daigremontiana and *Mesembryanthemum crystallinum* (Dakhiya *et al.* 2017, Davies and

Griffiths 2012, Dodd *et al.* 2003, Dodd *et al.* 2004, Dodd *et al.* 2005, Gould *et al.* 2009,

Hennessey and Field 1991, Wyka *et al.* 2005). Aspects of photosynthetic physiology that are

circadian regulated include the rates of CO_2 assimilation and oxygen evolution, and

photosynthetic light harvesting reported using chlorophyll fluorescence analysis (Dakhiya *et*

al. 2017, Hennessey and Field 1991, Litthauer *et al.* 2015, Schweiger *et al.* 1964). There are

also circadian rhythms in stomatal opening, which are functionally separable from rhythms of

CO_2 assimilation, because rhythms of assimilation continue when the intercellular partial

pressure of CO_2 is held at a constant level to remove the influence of rhythmic stomatal

opening upon assimilation (Hennessey and Field 1991). In addition to flowering plants,

aspects of metabolism associated with photosynthesis are circadian regulated in some

cyanobacteria and algae (Cano-Ramirez *et al.* 2018, Mitsui *et al.* 1986, Samuelsson *et al.*

1983, Schneegurt *et al.* 1994, Schweiger *et al.* 1964). Despite this evidence for the circadian

regulation of photosynthesis, the mechanisms and evolution of this process remain less well

understood.

The pervasiveness of circadian regulation of photosynthesis within flowering plants and

unicellular photosynthetic organisms might suggest that this process is conserved

throughout multicellular plant life, with a potentially ancient evolutionary origin. One

approach to examine this idea is to investigate plant species that diverged from flowering

plants at a relatively early stage of land plant evolution, because this can establish whether

the mechanism might originate from a common ancestor. As a model to investigate this

question, we selected the bryophyte liverwort *Marchantia polymorpha*. *M. polymorpha* is

thought to have diverged from flowering plants about 400 million years ago (Delwiche and

Cooper 2015, Kohchi *et al.* 2021), with liverwort-like plants occurring earlier in the fossil

record than other land plant forms such as vascular plants (Edwards *et al.* 1995). *M.*

polymorpha is a useful model to investigate questions concerning the evolution of circadian

regulation because it has a circadian oscillator that shares some components with flowering

plants (Lagercrantz *et al.* 2021, Linde *et al.* 2017), there is circadian regulation of a subset of

the transcriptome and the position of its thallus lobes (Lagercrantz *et al.* 2021, Lagercrantz

et al. 2020), and it can be used in experimental designs comparable to models such as

Arabidopsis. Whilst circadian rhythms and some aspects of circadian clock structure have

been identified in *M. polymorpha*, it is not yet known whether the regulation of

photosynthesis represents a conserved output from its circadian system.

Here, we used *M. polymorpha* as a tool to investigate three questions concerning the

conservation of circadian regulation of photosynthesis in multicellular plants. First, we

investigated whether there is circadian regulation of photosynthesis in *M. polymorpha*, to

determine whether this process is conserved in a species that diverged from vascular plants

at a relatively early stage of land plant evolution. Second, we investigated whether a

circadian clock in *M. polymorpha* might influence the daily regulation of photosynthesis

under cycles of light and dark, to determine whether circadian regulation might contribute to

the timing of photosynthesis under diel cycles. Finally, we investigated whether

pharmacological approaches might provide insights into the mechanisms that underlie the

circadian rhythms of photosynthesis.

Results

Circadian rhythms of photosynthesis in M. polymorpha

We used two proxies of photosynthetic activity to investigate circadian rhythms of

photosynthesis in *M. polymorpha*. These were delayed chlorophyll fluorescence (DF), and

prompt fluorescence (PF) measured using the pulse amplitude modulation (PAM) method

(Dakhiya and Green 2022, Gould *et al.* 2009, Maxwell and Johnson 2000). When illuminated

leaves are transferred to darkness, the potential for electrons to move down the electron

transport chain is removed, and energy within the system is instead re-emitted as light and

heat. The light emitted within the nanosecond range after the lights turn off is referred to as

prompt fluorescence (Dakhiya and Green 2022, Maxwell and Johnson 2000). Light emitted

within the millisecond to second range following lights-off is referred to as DF (Gould *et al.*

2009, Rees *et al.* 2019, Strehler and Arnold 1951). Both DF and certain features of PF are

circadian-regulated in a range of flowering plants and some algae (Cano-Ramirez *et al.*

2018, Dakhiya and Green 2022, Gould *et al.* 2009, Gyllenstrand *et al.* 2014). One exception

appears to be spruce trees (*Picea abies*), where the DF rhythm damps rapidly in accordance

with a rapidly damping circadian oscillator (Gyllenstrand *et al.* 2014). Features of chlorophyll

fluorescence that oscillate with a circadian rhythm include the apparent quantum yield of

PSII (YII), a measure of non-photochemical quenching of chlorophyll fluorescence within

PSII (NPQ), and the estimated photosynthetic electron transport rate (ETR) (Cano-Ramirez

et al. 2018, Dakhiya and Green 2022, Dakhiya *et al.* 2017). Taken together, rhythms of DF

and PF describe circadian regulation of processes of photochemistry over time, as greater

energy loss as heat (NPQ) and light (DF) indicate lower quantum yields (Murchie and

Lawson 2013).

We investigated whether free-running circadian rhythms of DF occur in *M. polymorpha*. For

these experiments, plants were investigated under conditions of either continuous light,

which was interrupted hourly by darkness for the measurement of DF, or under conditions of

continuous darkness, interrupted hourly by a 5 minute light pulse to allow subsequent DF

measurement. We named these conditions free running light conditions (FRL) and free

running dark conditions (FRD), respectively. We also investigated DF responses under

cycles of 12 h light and 12 h darkness, with the light and dark periods interrupted at hourly

intervals, for 5 minutes, for the measurement of DF. We named these conditions zeitgeber

cycles (ZTC). Rhythmic thalli were identified using Metacycle (Wu *et al.* 2016) ($q <$ 0.001;

period between 18 and 34 hours). Under FRL conditions, 63% of thalli had circadian rhythms

of DF that persisted for 4-5 days, and damped over time (Fig 1a, b; Table 1; Fig. S1;

Metacycle $q <$ 0.001). The mean period of these rhythms under FRL conditions was 25.50 h

± 0.52 h (mean ± s.e.m.). Under FRD conditions, 78% of thalli had circadian rhythms of DF

(Table 1; Metacycle $q <$ 0.001). However, there was considerable variability between the

rhythms of individual thalli, so the mean DF signal from all rhythmic thalli had little overt

rhythmicity (Fig. 1c, d; Fig. S2). In contrast, under zeitgeber cycles, 100% of thalli were

rhythmic (Metacycle $q <$ 0.001), and the rhythms had better phase synchronization between

individual thalli (Fig. 1e, f; Fig. S3).

Rapid damping of a rhythm over time (Fig. 1a, red line) can suggest the desynchronization of

rhythms of individual replicates (Paajanen *et al.* 2021). We hypothesized that such

desynchronization might occur for DF in *M. polymorpha* under our conditions. This is

supported by the considerable range of phases across replicate thalli under both FRL and

FRD conditions (Fig. 1g, h), which contrasts zeitgeber cycle conditions where the phase of

many replicates clustered to the end of the subjective day (6 – 10 h; Fig. 1i). To investigate

this further, we calculated the synchrony of phase between individual replicates, using the

order parameter *r* within the Kuramoto phase oscillator model (Kuramoto 1984, Muranaka

and Oyama 2016). This identified greater phase synchrony under zeitgeber cycles compared

with free running conditions (Fig. 1j), with phase synchrony weakening over time under

zeitgeber cycles. We were interested in the level of phase synchrony of DF in *M. polymorpha*

compared with other species, so also calculated also phase synchrony for DF in

Arabidopsis, *Triticum aestivum* and *Brassica napus* from published data (Rees *et al.* 2019,

Rees *et al.* 2021) (Dataset S1). This identified very high phase synchrony for replicates of

Arabidopsis (r = 0.92) and *B. napus* (r = 0.96) under FRL conditions, compared with

relatively low phase synchrony for both *T. aestivum* under FRL conditions (r = 0.33) and *B.*

napus under FRD conditions (r = 0.47) (Dataset S1).

To investigate the robustness of the relatively desynchronized rhythms in *M. polymorpha* we

used the parameter of relative amplitude error (RAE), which is derived from analysis of the

data by fast Fourier transform-non linear least squares method (FFT-NLLS). RAE provides a

measure of the robustness of the oscillation, representing the ratio of the amplitude error to

the amplitude, with a value of 0 indicating a robust and non-damping rhythm, and a value of

1 indicating an absence of rhythmicity, or very noisy data. RAE of < 0.6 is sometimes

considered indicative of rhythmicity (Alabadí *et al.* 2001, Alabadí *et al.* 2002, Para *et al.*

2007). FFT-NLLS analysis suggested relatively robust rhythms under zeitgeber cycles, and

lower rhythmic robustness under FRL and FRD conditions (Fig. 1k). The mean RAE of

rhythms of DF for thalli under FRL conditions was 0.46 ± 0.01, which suggests a lower

robustness of the rhythm compared to rhythms of DF in *Arabidopsis* seedlings under the

same conditions (RAE = 0.31 – 0.44, depending on accession) (Rees *et al.* 2021). Taken

together, this suggests that whilst a proportion of thalli were rhythmic under FRL and FRD

conditions (Table 1), the rhythm of DF varied considerably between thalli and had lower

rhythmic robustness compared with under zeitgeber cycles (Fig. 1k). This also supports the

notion that light input masks or maintains the daily rhythm of DF in *M. polymorpha*.

To extend our DF analysis, we also measured circadian rhythms of photosynthetic activity in

M. polymorpha using the PAM chlorophyll fluorescence method. In this experiment, plants

were maintained under conditions of constant blue actinic light, with alterations in these light

conditions every two hours to measure fluorescence parameters (Cano-Ramirez *et al.* 2018).

Under these conditions within the PAM instrument, there were circadian rhythms in the

effective quantum yield of PSII (Y(II)) (Fig. 2a, b) and a measure of non-photochemical

quenching of chlorophyll fluorescence within PSII (NPQ) (Fig. 2c, d). These rhythms had a

period between 27 h and 28 h (28.10 h ± 0.15 h (Y(II)), 27.55 h ± 0.38 h (NPQ)) in *M.*

polymorpha (Fig. 2b, d). These period estimates are longer compared with the period

estimates obtained using DF (Fig. 1h). This might be caused by the different light conditions

used with the two methods, because the period of some circadian-regulated processes in *M.*

polymorpha depends upon the light intensity and spectrum (Linde *et al.* 2017). Alternatively,

this might reflect differences in the preparation of thalli for the two methods, since thalli were

grown directly from gemmae for DF imaging, whereas for PAM experiments the thalli were

cut into squares to prevent hyponasty. Y(II) and NPQ oscillated with opposing phases (Fig.

2a, c), which is consistent with other reports (Cano-Ramirez *et al.* 2018), and expected

because these measures represent competing fates for light energy absorbed by leaves.

Together with the measurements of DF, we conclude that free-running rhythms of several

photosynthetic outputs occur in *M. polymorpha*. The DF data suggest that free-running

rhythms are less robust and more variable under FRD conditions, but remain detectable in a

proportion of thalli (Table 1). Under zeitgeber cycles, the robustness of individual rhythms

(Fig. 1e, f, k) and relatively high phase synchrony between replicate thalli (Fig. 1j) suggests

that under these conditions, the rhythm of DF is influenced strongly by the zeitgeber cycle.

This indicates that under zeitgeber cycles, the underlying circadian oscillation (Fig. 1e) is

either driven by the light condition, or the underlying circadian rhythm is masked by the light

condition. This is consistent with the notion that the light/dark cycle has a strong effect upon

the diel regulation of the *M. polymorpha* circadian oscillator and its physiological outputs

(Linde *et al.* 2017).

Contribution of circadian regulation in M. polymorpha to the timing of rhythms of delayed

fluorescence under zeitgeber cycles

Because we identified that the zeitgeber cycle strongly influences the daily oscillation of DF

in *M. polymorpha* (Fig. 1a-f), we wished to investigate the extent of clock control of the

phase of delayed fluorescence oscillations in *M. polymorpha* under zeitgeber cycles. To

achieve this, we used an experimental approach whereby the phase of the rhythmic process

was monitored under a range of zeitgeber cycle lengths (known as T cycles; e.g. T24 = 12 h

light / 12 h dark; T28 = 14 h light / 14 h dark, and so on). This can identify effects of circadian

regulation upon the timing of a process under diel cycles, because the circadian oscillator

can influence the phase of the process, producing systematic changes in the phase of either

the entire waveform (Aschoff 1960, Eelderink-Chen *et al.* 2021, Roenneberg *et al.* 2005), or

parts of the rhythm (Dodd *et al.* 2014). Using the zeitgeber cycles of DF described earlier,

we identified oscillations of DF under diel cycles of T20 to T28 (Fig. 3a-d). The period length

of DF matched the T cycle length closely (20.09 h ± 0.06 h (T=20), 22.44 h ± 0.07 h (T=22),

24.36 h ± 0.07 h (T=24) and 28.17 h ± 0.06 h (T=28), measured using FFT-NLLS in

Biodare2). When the phase was normalized to the length of the T cycle, we found that the T

cycle length had a significant influence upon the phase, such that the phase was relatively

late under the longest T cycle (Fig. 3e). In general, a circadian oscillator would be expected

to cause relatively earlier phases under longer T cycles (Aschoff 1960), including in plants

(Dodd *et al.* 2014). This did not occur within our experiment (Fig. 3e), where instead the

phase followed the zeitgeber cycle length (Fig. 3a-e). This suggests that the effect of the

zeitgeber cycle masked the underlying effects of clock control under these conditions, and

that under zeitgeber cycles the effect of circadian regulation upon this aspect of

photosynthetic physiology is not detectable.

Abrogation of circadian rhythms of photosynthesis in M. polymorpha by a potential inhibitor

of chloroplast translation

Whilst the canonical circadian oscillator underlies the daily timing of photosynthesis (Dodd *et*

al. 2004, Dodd *et al.* 2005), the mechanisms by which the circadian oscillator regulates

photosynthesis remain poorly understood. One approach to identify potential cellular

processes associated with circadian rhythms of photosynthesis is to use a pharmacological

strategy, because several chemicals are known to inhibit specific processes within

chloroplasts. We combined a pharmacological approach with the monitoring of circadian

rhythms of chlorophyll fluorescence to identify candidate processes associated with the

circadian regulation of photosynthesis.

First, we tested whether an inhibitor of chloroplast translation can attenuate circadian

rhythms of chlorophyll fluorescence in *M. polymorpha*. Lincomycin is an antibiotic that binds

to the 50S subunit of bacteria-like ribosomes (Chang and Weisblum 1967) and is thought to

inhibit chloroplast but not mitochondrial protein synthesis (Sullivan and Gray 1999, Zhao *et*

al. 2018). We supplemented the growth media with a range of lincomycin concentrations (25

µg ml^{-1}, 50 µg ml^{-1} and 100 µg ml^{-1}, and a vehicle control), and monitored rhythms of

photosynthesis using the PAM fluorescence method (Fig. 4a-l; Fig. S4a-d). There was a

dose-dependent effect of lincomycin on the amplitude of rhythms of Y(II) and NPQ under

free running light conditions (Fig. 4f, l; Fig. S4b, d), such that the amplitudes of these

parameters were reduced by greater lincomycin concentrations (Fig. 4f, l; Fig. S4b, d)).

Furthermore, the RAE of circadian rhythms of Y(II) was significantly greater in the presence

of lincomycin compared with the vehicle control (Fig. 4e; Fig. S4a). The effect of lincomycin

on the RAE of rhythms of NPQ was less consistent across two experiments (Fig. 4k; Fig.

S4c), suggesting that lincomycin might have a greater effect upon the rhythmic robustness of

Y(II) compared with NPQ. These measures of photosynthesis are ratiometric, so the

decreased amplitude is unlikely to be caused by a lower fluorescence signal in the presence

of lincomycin. Whilst we cannot exclude the possibility that lincomycin in *M. polymorpha*

affects processes other than plastid translation, our data suggest that chloroplast translation

might be required to maintain circadian rhythms of photosynthetic light harvesting.

Next, we tested the hypothesis that an inhibitor of chloroplast transcription can attenuate

circadian rhythms of photosynthesis in *M. polymorpha.* In Arabidopsis, mutants of a

circadian-regulated sigma submit of plastid encoded plastid RNA polymerase (PEP) known

as SIG5 can alter the phase of circadian rhythms of delayed fluorescence (Noordally *et al.*

2013). Therefore, we reasoned that chloroplast transcription by PEP might contribute to

circadian rhythms of photosynthesis in *M. polymorpha*. To test this, we supplemented the

growth media with rifampicin, which is thought to inhibit chloroplast transcription by one form

of PEP (Pfannschmidt and Link 1997, Surzycki 1969). We measured rhythms of Y(II) and

NPQ, under free running light conditions, in the presence of several concentrations of

rifampicin (Fig. 4m-x; Fig. S4e-h). Across two experiments, under these experimental

conditions, there were no consistent differences caused by the range of rifampicin

concentrations tested upon the circadian rhythm of the apparent quantum yield of PSII or

NPQ (Fig. 4q, r, w, x; Fig. S4e, f, g, h).

Discussion

Here, we present evidence for the circadian regulation of two proxies for photosynthetic

activity in *M. polymorpha*. Our experiments cannot prove that circadian rhythms of

photosynthesis occurred in early land plants, but the presence of these rhythms in both

bryophytes and angiosperms suggests that they might be well-conserved. Given that

reporters of photosynthetic activity are also circadian regulated in aquatic photosynthetic

organisms such as cyanobacteria, uni- and multicellular algae (Cano-Ramirez *et al.* 2018,

Hastings *et al.* 1961, Okada *et al.* 1978, Schneegurt *et al.* 1994, Schweiger *et al.* 1964,

Sorek and Levy 2012, Sorek *et al.* 2013, Sweeney and Haxo 1961), our study suggests that

the circadian regulation of photosynthesis was conserved from aquatic ancestors during the

terrestrialization of plants. One interpretation is that circadian regulation of photosynthesis

confers a selective advantage to all photosynthetic life. The nature of the advantage and

degree of selection pressure for this might depend upon the ecological niche occupied by

the organism (Hellweger *et al.* 2020), particularly because certain species (e.g. *Picea abies*)

appear to lack these rhythms (Gyllenstrand *et al.* 2014).

Our experiments identified that under free running light conditions, the period of the rhythm

of delayed fluorescence was not uniform between replicate thalli, with this variation leading

to phase desynchrony (Fig. 1a-d). In comparison to *M. polymorpha*, the circadian rhythm of

DF in Arabidopsis and other flowering plants is relatively stable, with a period of about 24 h

and individual replicates relatively synchronized (Dataset S1) (Gould *et al.* 2009, Rees *et al.*

2021). Other reporters for circadian rhythms in *M. polymorpha* are also less robust than in

flowering plants. For example, rhythms of some promoter-luciferase reporters damp quite

rapidly and present a range of periods across replicates, with this difference in robustness

evident when circadian rhythms in *M. polymorpha* and Arabidopsis are compared

(Lagercrantz *et al.* 2021, Linde *et al.* 2017) (Dataset S1). This desynchronization of replicate

thalli is reminiscent of the phase desynchronization that occurs between the rhythms of

individual cells in the leaves of *Lemna gibba* under free running conditions, compared with

the maintenance of phase synchrony in *L. gibba* under zeitgeber cycles (Muranaka and

Oyama 2016).

One potential explanation for this difference between circadian rhythms in *M. polymorpha*

and flowering plants might relate to circadian oscillator structure (Linde *et al.* 2017). The

circadian oscillator in *M. polymorpha* has been proposed to include a smaller number of

components than the oscillator of Arabidopsis, with greater levels of oscillator complexity

thought to confer robustness to perturbation (Shalit-Kaneh *et al.* 2018, Troein *et al.* 2009).

Therefore, the less robust circadian oscillations of photosynthesis that we report might arise

from lower levels of redundancy within a less complex circadian clock in *M. polymorpha*

compared with Arabidopsis. An alternative explanation for this difference might relate to

differences in the strength of coupling of circadian rhythms between cells and across tissues

in Arabidopsis and *M. polymorpha*. Although there is weak intercellular coupling of circadian

rhythms in Arabidopsis (Wenden *et al.* 2012), differences between the circadian period of

individual cells means that the rhythms of cell populations become progressively more

desynchronized over time in the absence of intercellular coupling, leading to an apparent

damping of the rhythm (Gould *et al.* 2018, Muranaka and Oyama 2016, Paajanen *et al.*

2021). Therefore, rhythms might damp more rapidly in *M. polymorpha* if intercellular coupling

of circadian rhythms is weaker compared with Arabidopsis, especially when combined with a

potentially less robust circadian oscillator within each cell. The idea of relatively unstable

circadian rhythms in the bryophytes is also supported by DF data from *Physcomitrium*

patens, which is arrhythmic under constant light and has a noisy rhythm under low light

conditions (Gyllenstrand *et al.* 2014). It will be interesting in future to determine whether

there is simply phase desynchronization between individual replicate thalli of *M. polymorpha*,

or patchy or complete intercellular desynchronization within individual thalli.

The thallus of *M. polymorpha* lacks stomatal pores for gas exchange and, instead, has air

pores in its surface that are formed by a channel passing through a cylindrical structure of 16

cells (Jones and Dolan 2017, Shimamura 2016). Unlike the stomata of flowering plants, the

air pores of *M. polymorpha* are thought to be static structures that do not adjust in response

to environmental cues. Therefore, circadian rhythms of the reporters of photosynthesis

described here might be due entirely to the circadian regulation of the biochemical reactions

of photosynthesis, rather than changes in the rate of CO_2 and O_2 exchange with the

atmosphere. Nevertheless, we cannot rule out alterations in conductivity to gas exchange

through diel or circadian changes in cell turgor or potential "motor cells" within liverwort air

pores (Walker and Pennington 1939). Such changes might influence the measures of

photosynthesis that we used, because the presence of the pores and air spaces within the

thallus appears to reduce its resistance to CO_2 diffusion (Green and Snelgar 1982).

The reduction in robustness of rhythms of Y(II) caused by lincomycin could suggest that

chloroplast translation is necessary for circadian rhythms of this reporter of photosynthesis.

Lincomycin inhibits the synthesis of PSII D1, PSII D2 and RbcS proteins, and the expression

of photosynthesis-associated nuclear encoded genes (PhAnGs) in angiosperms (Bachmann

et al. 2004, Karpinska *et al.* 2017, Mulo *et al.* 2003). Perhaps inhibiting the expression of

these proteins prevents the replacement of PSII components, with ensuing impacts upon

circadian cycles of light harvesting. The absence of any effect of rifampicin on circadian

rhythms of Y(II) or NPQ, contrasting the effect of lincomycin, could suggest that rhythms

of these parameters involve genes transcribed by nuclear encoded plastid RNA

polymerase (NEP) rather than rifampicin-sensitive PEP, there is compensation by NEP

for reduced PEP activity, or specific roles for the rifampicin sensitive- and insensitive

forms of PEP (Pfannschmidt and Link 1994). Whilst this suggests that protein synthesis in

chloroplasts is necessary for the circadian regulation of photosynthesis, it does not identify

that this is a rate-limiting regulatory step that underlies the clock control of the process. It

also supports the idea that *M. polymorpha* represents an informative model for investigation

of mechanisms underlying the circadian regulation of physiology, particularly considering the

range of gene-editing tools available.

Whilst the apparent reduction of robustness of circadian rhythms in *M. polymorpha* might

increase the susceptibility of rhythms in *M. polymorpha* to environmental perturbation (Linde

et al. 2017), it is alternatively possible that a circadian oscillator with greater variability might

confer selective advantages. For example, plasticity of circadian regulation of the daily

photosynthetic cycle of CAM plants, and of carbohydrate metabolism in Arabidopsis, are

both thought to allow better alignment between metabolic processes and daily cycles of

energy availability (Borland *et al.* 2011, Dodd *et al.* 2002, Graf *et al.* 2010, Webb *et al.* 2019).

In comparison, very little is known about diel cycles of metabolism in *M. polymorpha*, or how

circadian regulation might confer fitness to *M. polymorpha*. To provide a better context for

understanding these findings, it might be valuable in future to investigate the regulation and

adaptive functions of the circadian system in *M. polymorpha*, or in naturally occurring

populations of *M. polymorpha*.

Materials and Methods

Plant material and growth conditions

Tissue of *Marchantia polymorpha* subsp. *ruderalis* (Tak-1 background) was cultivated under

sterile conditions, using half-strength Gamborg B5 media containing 0.8% (w/v) agar at pH

5.8 (Solly et al. 2017). Cultivation occurred in Panasonic MLR-352 growth chambers, under

cycles of 12 h light / 12 h darkness, at 19 °C, under 100 µmol photons m^{-2} s^{-1} of white light

(light spectrum in Fig. S5a). For inhibitor experiments, lincomycin (Alfa Aesar) or rifampicin

(Melford Laboratories Ltd) were added for the PAM chlorophyll fluorescence experiments at

concentrations of 25, 50 and 100 µg/ml, with a dimethylsulfoxide (DMSO) vehicle control

where necessary.

For DF experiments, gemmae (Shimamura 2016) were positioned on nutrient agar and

cultivated for 14 days prior to the start of the experiment, and entrained under 12 h light /

12 h dark cycles for this entire period. For PAM fluorescence experiments, pieces of 21-day

old thallus cultivated under 12 h light / 12 h dark cycles were cut into square sections under

sterile conditions and allowed to regenerate for 6 days, during which they were entrained

under 12 h light / 12 h dark cycles. Preparation of the thallus as square shapes prevented

hyponasty during these experiments. For both DF and PAM experiments, plants were

entrained under conditions described above prior to imaging.

PAM chlorophyll fluorescence measurement

Circadian time-series of chlorophyll fluorescence measurements were obtained using an

IMAGING-PAM M series chlorophyll fluorescence imaging system (Walz GmbH, Effeltrich,

Germany), using a protocol described previously (Cano-Ramirez *et al.* 2018). Briefly, rapid

light curves (RLCs) were obtained at 2 hour intervals, with the plants maintained under

constant blue light at other times during the timecourse. Each RLC was acquired over a

period of 6 minutes, which was preceded by a 20 minute period of dark adaptation. For the

acquisition of each RLC, a saturating light pulse (845 µmol m^{-2} s^{-1}) was applied, and followed

by an increase in actinic light across 16 steps, at 20 second intervals, to a maximum of 789

µmol m^{-2} s^{-1}. After acquiring the RLC, samples were held under continuous blue light (15

µmol m^{-2} s^{-1}; light spectrum in Fig. S5b) until the next measurement cycle. PAM chlorophyll

fluorescence parameters were measured at 108 µmol m^{-2} s^{-1} actinic light on the RLC, which

was similar to the growth and entrainment conditions (100 µmol m^{-2} s^{-1}).

Measurement of delayed fluorescence

Delayed fluorescence analysis was performed as described previously (Gould *et al.* 2009,

Rees *et al.* 2019). This involved image capture with Retiga LUMO cameras (QImaging,

Canada), fitted with Xenon 0.95/25 mm lenses (Scheneider-Kreuznach, Germany). Plants

were illuminated with mixed red/blue LED arrays (approx. 60 µmol m^{-2} s^{-1}; light spectrum in

Fig. S5c) under the control of the µManager Software (v1.4.19, Open Imaging) and an

Arduino Uno microcontroller board. The camera and LEDs were mounted in growth

chambers maintained at 19 °C (Sanyo MIR-553). The camera settings for all experiments

were Binning = 4, Gain = 1, and Readout-Rate=0.650195 MHz, 16 bits. Three control scripts

were used for DF measurement in *Marchantia polymorpha*. The free running light condition

script was a regime of 59 minutes of light followed by one minute of image acquisition, the

free running dark condition script was 54 minutes of darkness followed by five minutes of

light and one minute of image acquisition, and the zeitgeber script was an alternating

combination of FRL and FRD scripts that swapped every half T-cycle. Exposure of plants to

5 minutes of light before image acquisition under both free running dark conditions and the

dark period of zeitgeber cycles was necessary to obtain the DF signal, and might be

perceived by plants as a low fluence response equivalent to an average photosynthetically

active radiation (PAR) of approximately 3 µmol m^{-2} s^{-1} over each hour. All experiments

started at entrained dawn, and finished after seven days of imaging. T-cycle experiments

employed the zeitgeber cycle script, but using daily periods of 20 h (10L/10D), 22 h

(11L/11D) and 28 h (14L/14D), respectively. For analysis, images were imported into FIJI

(version 1.53c) and regions of interest were selected and extracted using the multi-measure

plugin to measure integrated density of each region. Raw data before filtration are provided

in Dataset S2. The DF data were baseline and amplitude (BAMP) de-trended using

Biodare2, and a moving average was used to smooth traces from individual thalli.

Data analysis

The DF data were smoothed using a 5 h moving average (Gould *et al.* 2009) and baseline

and amplitude (BAMP) detrended using Biodare2 (Zielinski *et al.* 2014). These data are

provided Fig. 1b, d, f and Fig. S1, S2 and S3 to exemplify variation in the DF rhythms across

all thalli. Data detrending was necessary because rapid thallus growth during the experiment

produced a continuously increasing raw signal. The data were subsequently analyzed using

Metacycle (Wu *et al.* 2016) to identify rhythmic thalli ($q < 0.001$; period range 18 – 34 h).

Data that passed this filtration step were used for calculation of rhythmic features and

presented in Fig. 1a, c, e, g-k and Table 1.

Rhythmic features of the data were quantified using the Fast Fourier Transform-Non-linear

least squares method (FFT-NLLS) algorithm within Biodare2 (Zielinski *et al.* 2014), and also

with Metacycle (Wu *et al.* 2016). Statistical analysis on derived data was conducted using

IBM SPSS Statistics v24 and SigmaPlot v14 for PAM chlorophyll fluorescence and DF data.

Graphs were prepared using R, with a combination of the ggplot2, scales, rhesape2,

cowplot, png, grid, RColorBrewer, wesanderson, colorspace, ggpubr, ggrepel, and

patchwork packages. Heat maps were created using the R package Pheatmap. A 5 h

moving average was applied to DF and PAM fluorescence data before analysis (Gould *et al.*

2009).

For calculation of the change in mean RAE and synchronisation index over time (Fig. 1j, k),

data were smoothed using a moving average, and filtered using Metacycle to identify

rhythmic thalli ($q < 0.001$; period range 18 – 34 h). RAE estimates and phases were re-

calculated using FFT-NLLS across a 48h sliding window (e.g. 24-72 h, 48-96 h, and so on)

Only rhythmic thalli that did not produce error messages within Biodare were included in the

means for each time window (FRL: N= 35-52; FRD: N=47-62; ZTC: N=95-98). The

synchronisation index is Kuramoto's order parameter, r (Kuramoto 1984, Nykamp accessed

2022), which represents the synchrony of phases in any given time window, where r = 1

indicates absolute phase synchrony. Our code for the calculation of the synchronisation

index and mean phases for each time window is available on GitHub

(https://github.com/AHallLab/KuramotoParameters).

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Figure legends

Figure 1. Circadian and diel rhythms of delayed fluorescence of *Marchantia polymorpha*

thalli under (a, b) free running light, (c, d) free running dark, and (e, f) zeitgeber cycle

conditions. (a, c, e) indicate DF signal from thalli categorized as rhythmic after filtration with

Metacycle (*q*-value < 0.001; period between 18 h and 34 h). The mean is the thick coloured

line and individual replicate thalli are fine grey lines. (b, d, f) Heat maps summarize DF

intensity of 50 individual and randomly selected thalli across two experimental repeat

experiments under (b) free running light, (d) free running dark and (f) zeitgeber cycle

conditions. Data in (a, c, e) are smoothed with a 5-hour moving average, baseline and

amplitude de-trended, and normalized to the mean signal. On heat maps, colour values are

scaled per row using Z-score normalization. (g-i) Phase of replicate thalli relative to dawn

(0), calculated using fast Fourier transform-non linear least squares method (FFT-NLLS).

Radial scale shows relative amplitude error (RAE) of each thallus, with RAE=0 at the exterior

and RAE=1 at the centre of the circles, respectively. (j) Comparison of relative

synchronization between replicate thalli under each light condition, using the order

parameter within the Kuramoto phase oscillator model to provide a synchronization index.

Greater values indicate better phase synchrony between replicates. (k) Comparison of RAE

change over time under each light condition. Data derive from 100 individual thalli, across

two independent experimental replicates, for each condition.

Figure 2. Circadian rhythms of modulated chlorophyll fluorescence in thalli of *Marchantia*

polymorpha. Under free running conditions, time-course analysis of (a, b) the apparent

quantum yield of PSII (Y(II)), and (c, d) non-photochemical quenching of PSII fluorescence

(NPQ). (a, c) are the normalized signal level for each chlorophyll fluorescence parameter,

processed using amplitude and baseline detrending and normalized to the mean (using

Biodare2), and shaded grey areas indicate the subjective night. (b, d) compare the estimated

period and relative amplitude error (RAE) of the rhythmic thalli, with period, RAE and the

presence of rhythmicity quantified using FFT-NLLS (using Biodare2). n = 20 independent

thalli.

Figure 3. Analysis of the contribution of circadian regulation to the rhythms of delayed

chlorophyll fluorescence under zeitgeber cycles. DF rhythms were quantified under T cycles

of lengths of (a) 20 h, (b) 22 h, (c) 24 h and (d) 28 h, with the mean intensity plotted. (e) The

phase was estimated from each time-series and normalized to a 24 h T cycle length, to

evaluate the phase relationship between the rhythm of DF and the zeitgeber cycle. A strong

contribution of the circadian clock to the timing of the rhythm under zeitgeber cycles would

be expected to drive a relatively earlier phase under longer T-cycles, which (e) is not the

case. (a-d) Shaded areas indicate dark periods. n = 150 thalli per T cycle (T20: 126 of 150

points rhythmic; T22: 127 of 150 points rhythmic; T24: 122 of 150 points rhythmic T28: 129

of 150 points rhythmic). Phase estimates obtained using FFT-NLLS, with data amplitude and

baseline detrended. Only rhythmic thalli were included in FFT-NLLS analysis. In (e), different

letters indicate significantly different mean phases (one-way ANOVA, where $p < 0.05$ based

on Tukey's post-hoc test).

Figure 4. Amplitude of circadian rhythm of chlorophyll fluorescence parameters is reduced

by the inhibitor of chloroplast translation lincomycin. (a-f) Effect of range of concentrations of

lincomycin upon the circadian oscillation of the apparent quantum yield of PSII (Y(II)). (g-l)

Effect of range of concentrations of lincomycin upon the circadian oscillation of non-

photochemical quenching of chlorophyll fluorescence. (m-r) Effect of range of concentrations

of rifampicin upon the circadian oscillation of the apparent quantum yield of PSII (Y(II)). (s-x)

Effect of range of concentrations of rifampicin upon the circadian oscillation of non-

photochemical quenching of chlorophyll fluorescence. (e, f, k, l, q, r, w, x) Relative amplitude

error, period and amplitude were estimated using FFT-NLLS with amplitude and baseline

detrending. (e, k, q, w) Different letters next to data points indicate significant differences in

period (upper case)) and RAE (lower case), with data plotted ± s.e.m. (f, l, r, x). Analyzed

with one-way ANOVA, where $p < 0.05$ based on Tukey's post-hoc test; n = 25 thalli. On time-

series plots, white sections indicate subjective day and grey shading indicates subjective

night.

Figure S1. Example of raw delayed fluorescence data for 64 individual replicate thalli of *M.*

polymorpha, under free running light conditions. Clear areas and shaded areas on the plots

indicate subjective day and night, respectively.

Figure S2. Example of raw delayed fluorescence data for 64 individual replicate thalli of *M.*

polymorpha, under free running dark conditions. Clear areas and shaded areas on the plots

indicate subjective day and night, respectively.

Figure S3. Example of raw delayed fluorescence data for 64 individual replicate thalli of *M.*

polymorpha, under zeitgeber cycles. Clear areas and shaded areas on the plots indicate day

and night, respectively.

Figure S4. Amplitude of circadian rhythm of chlorophyll fluorescence parameters is reduced

by the inhibitor of chloroplast translation lincomycin. This is a second direct repeat of the

experiment in Fig. 4. (a-d) Effect of range of concentrations of lincomycin upon the circadian

oscillation of (a, b) the apparent quantum yield of PSII (Y(II)) and (c, d) non-photochemical

quenching of chlorophyll fluorescence. (e-h) Effect of range of concentrations of rifampicin

upon the circadian oscillation of (e, f) Y(II) and (g, h) non-photochemical quenching of

chlorophyll fluorescence. (a, c, e, g) Different letters next to data points indicate significant

differences in period (upper case) and RAE (lower case), with data plotted ± s.e.m. (b, d, f,

h) Different letters indicate significantly different mean phases. Analyzed with one-way

ANOVA, where $p < 0.05$ based on Tukey's post-hoc test; n = 25 thalli.

Figure S5. Spectra of light conditions used for experiments. Photon flux density spectrum

within (a) growth chambers used for cultivation and entrainment of M. polymorpha, (b)

imaging PAM system actinic light conditions and (c) delayed fluorescence imaging LED

array.

Dataset S1. Comparison of synchrony of circadian rhythms of delayed fluorescence in

variety of species and experimental conditions. This uses using the order parameter within

the Kuramoto phase oscillator model to provide a synchronization index. Data derive from

this paper, (Rees *et al.* 2019, Rees *et al.* 2021).

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Experiment 1

Condition	Number of thalli	Number of rhythmic thalli	Rhythmic thalli (%)
FRL	50	33	66
FRD	50	42	84
ZTC	50	50	100

Experiment 2

Condition	Number of thalli	Number of rhythmic thalli	Rhythmic thalli (%)
FRL	50	30	60
FRD	50	36	72
ZTC	50	50	100

Experiments combined

Condition	Number of thalli	Number of rhythmic thalli	Rhythmic thalli (%)
FRL	100	63	63
FRD	100	78	78
ZTC	100	100	100

Table 1. Comparison of number of *M. polymorpha* thalli having a circadian or diel

rhythm of delayed fluorescence under each light condition, from two independent

experiments. Data were classified as rhythmic when the Metacycle q-value was <

0.001 and period lengths between 18 h and 34 h. The abbreviations FRL, FRD and

ZTC represent free running conditions of light, free running conditions of darkness,

and zeitgeber cycles (light/dark cycles), respectively.

(a) Control treatment
(g) Control treatment
(m) Control treatment
(s) Control treatment

(b) 25 µg ml⁻¹ lincomycin
(h) 25 µg ml⁻¹ lincomycin
(n) 25 µg ml⁻¹ rifampicin
(t) 25 µg ml⁻¹ rifampicin

(c) 50 µg ml⁻¹ lincomycin
(i) 50 µg ml⁻¹ lincomycin
(o) 50 µg ml⁻¹ rifampicin
(u) 50 µg ml⁻¹ rifampicin

(d) 100 µg ml⁻¹ lincomycin
(j) 100 µg ml⁻¹ lincomycin
(p) 100 µg ml⁻¹ rifampicin
(v) 100 µg ml⁻¹ rifampicin

(e) Lincomycin
(k) Lincomycin
(q) Rifampicin
(w) Rifampicin

(f) Lincomycin
(l) Lincomycin
(r) Rifampicin
(x) Rifampicin

- No treatment (NT) - 25 µg ml⁻¹ - 50 µg ml⁻¹ - 100 µg ml⁻¹
- No treatment (NT) - 25 µg ml⁻¹ - 50 µg ml⁻¹ - 100 µg ml⁻¹