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Effect of metformin exposure on growth and photosynthetic performance in the unicellular freshwater chlorophyte, *Chlorella vulgaris*

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Abstract

Many pharmaceuticals have negative effects on biota when released into the environment. For example, recent work has shown that the commonly prescribed antidiabetic drug, metformin (N,N-dimethylbiguanide), has endocrine disrupting effects on fish. However, effects of metformin on aquatic primary producers are poorly known. We exposed cultured isolates of a freshwater chlorophyte, Chlorella vulgaris, to a range of metformin concentrations (0-767.9 mg L⁻¹) to test the hypothesis that exposure negatively affects photosynthesis and growth. A cessation of growth, increase in non-photochemical quenching (NPQ, NPQ_{max}), and reduced electron transport rate (ETR) were observed 24 h after exposure to a metformin concentration of 767.8 mg L⁻¹ (4.6 mM). By 48 h, photosynthetic efficiency of photosystem II (F_v/F_m), α , the initial slope of the ETR-irradiance curve, and E_k (minimum irradiance required to saturate photosynthesis) were reduced. At a lower concentration (76.8 mg L⁻¹), negative effects on photosynthesis (increase in NPQ, decrease in ETR) were delayed, occurring between 72 and 96 h. No negative effects on photosynthesis were observed at an exposure concentration of 1.5 mg L⁻¹. It is likely that metformin impairs photosynthesis either through downstream effects from inhibition of complex I of the electron transport chain or via activation of the enzyme, SnRK1 (sucrose non-fermenting-related kinase 1), which acts as a cellular energy regulator in plants and algae and is an ortholog of the mammalian target of metformin, AMPK (5' adenosine monophosphate-activated protein kinase).

Introduction

Pharmaceuticals and personal care products (PPCPs) comprise a diverse class of chemical compounds that have gained attention as chemicals of emerging concern (CECs) due to their widespread detection in aquatic environments [1–3] and because of their bioactive properties [4–6]. Although the ecological effects of many active pharmaceutical ingredients (APIs) are poorly known, several have been implicated in acute [7–8] and chronic toxicity [9–11] to biota. Direct toxic effects of APIs on microalgae have been observed, with effects varying according to the species and drug [12–16].



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The biguanide drug, metformin (*N*,*N*-dimethylbiguanide), is the most commonly prescribed oral medication used to treat type II diabetes [17,18]. Since the incidence of type II diabetes is on the rise worldwide [19], there are increasing reports of metformin detections in wastewater effluent and surface waters of North America [1,20, 21] and Europe [22–24]. The widespread detection of metformin is due in part to its unique biochemistry and mode of action. Although the mechanisms are not completely understood [25–27], metformin counteracts hyperglycemia through activation of AMP-activated protein kinase (AMPK) [28] by inhibition of complex I of the mitochondrial electron transport chain [29]. AMPK belongs to a class of energy sensing enzymes responsible for maintaining energy homeostasis; specifically, AMPK promotes energy releasing (catabolic) processes and downregulates energy depleting (anabolic) processes (e.g., synthesis of fatty acids, proteins) in response to an increase in adenosine monophosphate (AMP) relative to adenosine triphosphate (ATP) levels (i.e., adenylate charge), and increases catabolic processes (e.g., glycolysis) [30]. An increase in adenylate charge signals a decrease in gluconeogenesis and an increase in glycolysis [26, 28]. Metformin remains chemically unaltered during this process [31].

In the laboratory, fathead minnows exposed to metformin at levels observed in wastewater effluent (40 μ g L⁻¹) showed significant upregulation of mRNA encoding the production of vitel-logenin [32], with stronger estrogenic effects seen in juvenile fish compared to adults [33]. This suggests that metformin may have negative effects on aquatic ecosystems, but effects on the lower aquatic food web are poorly known. Given its role as an AMPK-activator, however, metformin likely influences cell metabolism broadly among eukaryotes and would be therefore be expected to have widespread effects on the ecosystem. Limited data on the influence of metformin on phytoplankton suggest that lethal effects would not be expected at typical environmental levels concentrations $< ~80 \ \mu$ g L⁻¹ (with the highest values observed downstream of wastewater treatment plants [20, 24]) [12]; however, more subtle effects may occur, from direct effects such as suppression of photosynthesis or changes in species composition [13] due to chronic exposure, to indirect effects, such as the preferential elimination of grazers and trophic cascades [34].

The objective of this study was to determine the effects of metformin exposure on photosynthesis and growth in the freshwater chlorophyte, *Chlorella vulgaris*, isolated from the Columbia River, USA. This organism was chosen because it is a cosmopolitan freshwater taxa and its response to exposure should be broadly relevant. Moreover, as a chlorophyte, it shares similarities to land plants in terms of its photosynthetic machinery owing to a common evolutionary pathway [35]. Since photosynthesis is a highly regulated process in algae that enables cells to acclimate to fluctuating light environments, it was expected that responses in photosynthetic characteristics would be observed before changes in growth. The data will aid in characterizing toxicological effects of metformin exposure on AMPK/SnRK1/SNF1-containing eukaryotes in aquatic environments.

Materials and methods

Chemicals and materials

All glassware used in media preparation and culturing was cleaned according to the USGS National Water-Quality Assessment (NAWQA) protocols to minimize contamination from trace organic and inorganic constituents [36].

Metformin standards (as metformin hydrochloride) for the analysis of concentrations in growth media (0–1000 mg L⁻¹) were prepared by dilutions of a primary metformin stock solution (10 mg mL⁻¹ dissolved in mineral water; Toronto Research Chemicals). Mobile phase solvents included 0.1% UPLC-grade formic acid in HPLC-grade water (Sigma-Aldrich; solvent A), UPLC-grade 0.1% formic acid in HPLC-grade acetonitrile (Sigma-Aldrich; solvent B).

Aliquots of 0.2 μ m-filtered water were dispensed into 1.5 mL (12 x 32 mm) amber borosilicate glass SUN-SRI standard opening auto-sampler vials (Thermo Scientific). Internal standard (IS) solutions of deuterated metformin (Toronto Research Chemicals) were prepared from frozen stock solutions (1 mg mL⁻¹ in Milli-Q water) diluted to yield a working solution (of 998 ng mL⁻¹). Internal standard (10 μ g L⁻¹) was added to each sample and standard.

Experimental design

Isolates of *C. vulgaris* from the Columbia River (1.8-5.1 µm in diameter) were grown in triplicate batch cultures in 250 mL glass Erlenmeyer flasks in WC medium [37] at 18 °C under a 12:12 light:dark cycle in a walk-in environmental chamber equipped with full-spectrum lighting (~191±18 µmol photons m⁻² s⁻¹). Cell densities were determined at each time point using a Beckman Coulter Model Z2 particle counter (Beckman Coulter, Indianapolis, IN) after dilution with Isoton II phosphate-buffered saline (Beckman Coulter, Indianapolis, IN). Cells were kept in suspension by continuous gentle shaking (80 rpm) to prevent negative effects of high biomass such as CO₂ limitation of growth [38].

Triplicate cultures were grown to mid-exponential phase before receiving one of three target metformin concentrations: 1.5 mg L^{-1} (9.1 μ M), 76.8 mg L^{-1} (0.46 mM), and 767.8 mg L^{-1} (4.6 mM), plus controls (0 mg L⁻¹; no metformin added). These concentrations were selected to provide a broad range for the study of effects on photosynthesis. In addition, although the concentrations were much higher than would be observed in nature, the density of cells in culture was also orders of magnitude higher than in natural samples; thus, the average (or nominal) exposure by any given cell in the population could be considered to be approximately similar to environmental concentrations (e.g., the nominal concentration was the quotient of added concentration/cell density multiplied by the average cell density observed in a river sample [~4000 cells mL⁻¹], yielding cell-specific exposure values of 11.5 ng L⁻¹, 80 μ g L⁻¹, and 500 µg L⁻¹). This was done to alleviate potential suppression of toxicity that can occur when high cell densities are used in bioassays [39]. A 1 mL pooled sample (i.e., 0.33 mL from each of three replicates) was taken at 0, 5, and 96 h to measure dissolved metformin concentrations in the medium to confirm the quantities added. The samples were filtered and frozen at -20°C prior to analysis by LC-MS/MS. An additional flask was included in each treatment and control to record pH over the course of the experiment. The initial pH was 7.3 ±0.017.

Cell density was determined in each treatment every 24 h (i.e., 0, 24 h, 48 h, 72 h, and 96 h). Photosynthetic performance was determined at shorter time intervals (0, 1, 3, and 5 h) following metformin addition, and then every 24 h for 96 h, since changes in photosynthesis are more likely to be observed over shorter time intervals than are changes in growth. Photosynthetic performance was characterized using a Water-PAM (Pulse-Amplitude-Modulation) chlorophyll fluorometer (Heinz Walz, Germany) according to parameters shown in Table 1. All samples taken for PAM fluorometry were diluted to 3 mL with WC media and darkadapted for 30 min to fully oxidize Photosystem II and maximize fluorescence potential upon light saturation. Dark-adapted samples were exposed to 10 s of far-red light just before PAM measurements to process remaining inter-system electrons through excitation of Photosystem I and oxidation of the plastoquinone pool [40–42]. A 4-min light curve (where fluorescence of cells exposed to a saturating light pulse is measured at increasing levels of actinic light) was recorded for each dark-adapted sample.

Light saturation was determined by the maximum fluorescence (F_m) resulting from the initial saturating pulse and subsequent fluorescence peaks (F_m) associated with the stepwise actinic pulses. Maximum quantum yield of Photosystem II (Φ PSII_{max}, or F_v/F_m), relative electron transport rate (ETR), light intensity at saturation (E_k) , and non-photochemical quenching (NPQ) were calculated and considered indicators of photosynthetic performance [43, 44].



Notation	Determination method	Description
Fm F'm	Direct measurement	Fluorescence when PSII reaction centers are closed and plastoquinone pool is reduced
Fo	Direct measurement	Fluorescence when PSII reaction centers are open and plastoquinone pool is oxidized
$\Phi PSII_{max}$	$\frac{F_v}{F_m}$ or $\frac{F_m-F_o}{F_m}$	Efficiency of dark-adapted PSII to absorb light
ΦPSII	$rac{F'_v}{F'_m}$ or $rac{F'_m-F}{F'_m}$	Efficiency of PSII in the presence of light
ETR	ΦPSII x PAR ^a x 0.5 x 0.84	Rate of electron movement in the photosynthetic ETC ^b
α	Initial slope of light curve	Efficiency of electron transport in the photosynthetic ETC
Ek	$\frac{\text{ETR}_{max}}{\alpha}$	Irradiance at onset of light saturation
NPQ	$\frac{F_m - F'_m}{F'_m}$	Excitation dissipated by heat
	Notation Fm F_o ΦPSIImax ΦPSII ETR α Ek NPQ	$\begin{array}{c c} \textbf{Notation} & \textbf{Determination method} \\ \hline & & \\ F_m \\ F_{m}' \\ \hline & \\ \hline & \\ \hline & \\ F_o \\ \hline & \\ \hline & \\ F_o \\ \hline & \\ \hline & \\ \hline & \\ F_o \\ \hline & \\ \hline \\ \hline$

Table 1. Photosynthetic parameters measured or calculated by PAM fluorometer.

^aPAR = photosynthetically active radiation

^bETC = electron transport chain

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Determination of metformin concentrations

Following a 10 μ L injection, metformin was separated using a Shimadzu Prominence HPLC system using two binary pumps (Shimadzu LC-20AD XR Prominence LC pumps) and gradient elution on the reverse-phase column (Synerg Hydro-RP LC Column (250 x 4.6 mm, 4 μ m, 80 Å; Phenomonex, Torrance, CA). A ThermoFisher (Waltham, MA) BetaBasic C8 Javelin guard column (10 x 2.1 mm, 1.5 μ m) minimized column contamination from particulate matter in samples. The analyte was identified, detected, and confirmed using an AB Sciex QTRAP 5500 mass spectrometer (Applied Biosystems/MDS Sciex Instruments, Concord, ON, Canada) and Analyst 1.6.2 software. Samples with metformin concentrations >75 mg L⁻¹ were diluted 1:500 in mineral water to improve peak resolution. The concentration of metformin was calculated from a standard curve of measured area ratios (analyte area/IS area) versus expected concentration ratios (analyte concentration/IS concentration). A single six-point linear standard curve with a 1/x weighting factor was established for each run.

Statistics

Curves generated from ETR data were fit by non-linear regression using Microsoft Excel Solver; curve fitting for growth data was performed using GraphPad Prism (version 7.04, GraphPad Software, La Jolla, California). The resulting models were used to estimate parameters of growth and photosynthesis and differences among treatments and controls over time were tested using two-way repeated-measures ANOVAs performed using Microsoft Excel 2016 or GraphPad Prism.

Cell count data for short-term responses (0-5 h) to metform addition were plotted against time and fitted with an exponential growth equation:

$$N_t = M_o \ e^{\mu t} \tag{1}$$

where N_t is the number of cells at time, t, N_0 is the initial number of cells, and μ is the specific growth rate. Cell count data for the 0–96 h time period were plotted against time and fitted with a Weibull growth curve model (5) [45]. In addition to N_t , N_0 , and μ , the time corresponding to the point of inflection (δ) and N_{max} (the maximum number of cells) are included in the

Weibull model:

$$N_{t} = N_{max} - (N_{max} - N_{0})e^{(-(\mu t)^{\delta})}$$
(2)

Specific growth rates at each exposure level and time point were evaluated for differences relative to controls according to a two-way ANOVA.

ETR data were fitted according to Platt et al. [46] and photosynthetic parameters including the initial slope of the ETR curve (α), slope of the photoinhibition region (β), minimum saturating irradiance (E_k), and ETR_{max} were determined from the fitted data. Resulting models were used to compare parameter values among exposure levels and time points.

$$ETR = ETR_{max}[1 - exp(-\alpha I/ETR_{max})][exp(-\beta I/ETR_{max})]$$
(3)

Maximum quantum yield of PSII (Φ PSII_{max}) and NPQ were compared among exposure levels and time using two-way repeated-measures ANOVAs.

Results

Effects of metformin on cell density and growth

Metformin concentrations remained within 20% of initial spike concentrations during the experiment (data not shown). Metformin reduced the culture density of *C. vulgaris* at exposures of 76.8 mg L⁻¹ (0.46 mM) and 767.8 mg L⁻¹ (4.6 mM) due to a rapid reduction in growth that was not observed in controls (Fig 1). There was a significant interaction between treatment and time (ANOVA, F(27,72) = 17.3, p<0.0001), whereby cell densities were reduced relative to controls by 48 h after an exposure of 767.8 mg L⁻¹ (Tukey HSD). Unlike cell density, specific growth rates during the mid-exponential growth phase did not differ from the controls at any exposure concentration (data not shown), suggesting that growth ceased in response to exposure, rather than slowed.

The culture pH varied from 7.32 \pm 0.017 at T₀ to 7.36 \pm 0.054 at T₇₂ to 7.7 \pm 0.03 at the end of the experiment (T₁₆₆); since pH did not increase beyond 7.7, it is unlikely that the cultures were limited by the availability of carbon (as carbon dioxide or bicarbonate) in the medium [47].

Influence of metformin on photosynthetic performance

No short-term (<5 h) effects on electron transport rate (ETR)(Fig 2) or photosynthetic parameters (α , Φ PSII_{max}, E_k)(Fig 3) were observed at different exposure levels [ETR_{max}, (F(12,32) = 0.8302, p = 0.6199); α (F(12,32) = 1.84, p = 0.082), Φ PSII_{max} (F(12,32) = 1.09, p = 0.401, and E_k (F(12,32) = 1.405, p = 0.2145)]. Significant variability in β across treatment and time (F(12,32) = 3.224, p = 0.004) appeared to be stochastic, with no systematic differences among any of the treatments when single effects were examined.

By 24 h, ETR_{max} was reduced in the 767.8 mg L⁻¹ exposure (F(12,32) = 38.91, p<0.0001; Tukey HSD, p = 0.0065, p<0.0001) (Fig 2). Further decreases were observed until ETR reached baseline levels at 96 h, with the steepest decline occurring 48–72 h after exposure. At lower exposure levels, reduction of ETR_{max} was delayed or absent. At the 76.8 mg L⁻¹ exposure level, ETR declined between 72 and 96 h (Tukey HSD, p = 0.007), while no negative effects on ETR were observed at the 1.5 mg L⁻¹ exposure.

There were significant interactions between exposure concentration and α (F(12,32) = 40.02, p<0.0001) and exposure concentration and E_k (F(12,32) = 12.62, p<0.0001). Upon exposure to 767.8 mg L⁻¹ metformin, α values were 10-fold lower than controls and other treatments (Tukey HSD, p = 0.0029) (Fig 3). By 24 h, average E_k values in the 767.8 mg L⁻¹ exposure level were significantly lower than in the controls, decreasing from 885 to 263 µmol



Fig 1. Algal growth curves (in log values) at different metformin concentrations. (A) Culture densities in *Chlorella vulgaris* in four triplicate cultures (three treatments and a control); first panel tracks cell densities over 96 h prior to a spike of three concentrations of metformin at 120 h, indicated in the second panel. Short-term changes in cell density were tracked for 5 h (120–129 h). Third panel shows cell densities at 144 h (24 h post-spike), 168 h (48 h post spike), 192 h (72 h post-spike), and 216 h (96 h post spike). (B) Ratio of cell density determined after inoculation relative to initial value (C/C₀, where C₀ = initial concentration at time = 120 h). Error bars represent ± 1 standard deviation.

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photons m⁻² s⁻¹ over 96 h (Tukey HSD, p = 0.0003). Φ PSII_{max} decreased >two-fold (~0.7 to 0.3 µmol photons⁻¹ m² s⁻¹) by 48 h (p = 0.0005). A significant increase in NPQ_{max} was observed between 0 and 48 h at the highest exposure concentration (F(12,32) = 15.6, p<0.0001; Tukey HSD, p<0.0001). NPQ_{max} of the 76.8 mg L⁻¹ exposure also increased with time, reaching ~0.9 by 96 h (Tukey HSD, p<0.0001).

NPQ was elevated at irradiances $>\sim 1500 \mu moL$ photons m⁻² s⁻¹ 24 h and 48 h after exposure to the highest metformin concentration relative to controls and to the other treatments (Fig 4). Between 48 and 72 h post-spike, NPQ declined in the highest exposure treatment, but increased in the 76.8 mg L⁻¹ treatment at the higher irradiances. By 96 h, NPQ was lower in the highest-exposure cultures than the controls, while values exceeded controls in the 76.8 mg L⁻¹ treatment for most irradiance levels (Fig 4).

Discussion

Growth in *Chlorella vulgaris* ceased 24–48 h after exposure to 767.8 mg L^{-1} (4.6 mM) metformin, while exposure to lower concentrations (i.e., 76.8 mg L^{-1}) led to delayed effects (72–96 h) on photosynthesis and growth, including an increase in non-photochemical quenching



Fig 2. Effects of metformin on electron transport rates (ETR). Parameters were estimated by models of best fit according to Platt (1980) and include electron transport efficiency (α), photoinhibition (β), maximum relative electron transport rate (ETR_{max}), and minimum saturating irradiance (E_k). Symbols indicate different metformin additions: 0 µg L⁻¹ (black circles), 1.5 mg L⁻¹ (open squares), 76.8 mg L⁻¹ (open diamonds), and 767.8 mg L⁻¹ (open triangles).

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Fig 3. Effects of metformin on photosynthetic parameters of *C. vulgaris* over 96 h of exposure for four treatment levels: No addition (Control, open circles); 1.5 mg L⁻¹ (Treatment A, open triangles); 76.8 mg L⁻¹ (Treatment B, open squares); and 767.8 mg L⁻¹ (Treatment C, closed diamonds). Parameters include (A) maximum yield of PSII (Φ PSII_{max}), (B) electron transport efficiency (initial slope of the ETR curve, α), (C) minimum saturating irradiance (E_k), and (D) maximum nonphotochemical quenching (NPQ_{max}).

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Fig 4. Non-photochemical quenching (NPQ) versus photosynthetically active radiation (PAR) after exposure to three levels of metformin (Treatment A = 1.5 mg L^{-1} , Treatment B = 76.8 mg L^{-1} , and Treatment C = 767.8 mg L^{-1}) and a control over 96 h.

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(NPQ), the dissipation of light energy not used to carry out photochemistry. Delayed effects on growth could reflect a shift to the use of internal carbon stores, as occurs under conditions of low or fluctuating irradiance [48] or it could occur if upregulation of membrane transporters is required for metformin to enter mitochondria where it inhibits complex I and activates AMPK [31, 49]. Indeed, metformin is known to accumulate within mitochondria of mamma-lian cells [50, 51] through active transport across the membrane down a chemiosmotic gradient via organic cation transporters (e.g., OCT1–3; [52]). The number of available transporters may thus dictate effective metformin concentrations found within mitochondria, which require time for upregulation and thus lead to delayed effects.

Photosynthetic performance in *C. vulgaris* was impaired 24 h after exposure at the highest concentration (767.8 mg L⁻¹), shown by a ~20% decline in both electron transport rate (ETR-max) and in the minimum irradiance capable of saturating photosynthesis (E_k). These effects indicate a reduced capacity to process light energy, which is also supported by the observed decline in α and F_v/F_m (after 48 h) and most dramatically in elevated NPQ and NPQ_{max} values (after 24 h). The depression in ETR_{max} observed in *C. vulgaris* compared to controls is reminiscent of the response to low light in a number of algal species, where light-limited cells reach a much lower maximum rate of photosynthesis compared to those grown at saturating irradiance [53–55]. A decrease in F_v/F_m is generally interpreted as a signal of physiological stress in unialgal cultures; F_v/F_m is also typically depressed under high light [56]. To date, effects of metformin on photosynthesis have not been shown; the unusual biochemistry of the compound makes it somewhat difficult to predict its physiological target in aquatic microorganisms, since there are no chemical analogues (aside from other biguanide compounds) with known toxic modes of action with which to compare effects as has been done with other pharmaceutical compounds such as fluoxetine [57].

Non-photochemical quenching describes a number of processes involved in energy dissipation and photoprotection in photosynthetic cells [58]. NPQ components-differentiated by relaxation kinetics in response to a saturating pulse of light—include qE, a rapid-response deexcitation mechanism driven by the pH of the chloroplast lumen, qT, quenching that accompanies state transitions (a shift in the location of light harvesting antennae complexes from the readily fluorescent photosystem II to the weakly fluorescent photosystem I), and qI, or photoinhibitory quenching, which is associated with either protein damage or the reversible downregulation of photosystem II, sometimes referred to as qRC (where RC refers to reaction center [59]). Among these, the likely origin of the response by *C. vulgaris* to metformin exposure in this study is qI. The rationale behind this hypothesis is that the timescale of response by qE is short (seconds); the timescale for qT is slower, but still on the order of minutes. On the other hand, quenching that occurs at the reaction center is slower (hours). The fact that changes in NPQ and NPQ_{max} occurred over a period of days in the present study suggests a longer-term loss of capacity to process light energy. In addition, it has been shown that qE deficient mutants of Chlamydomonas reinhardtii are able to acclimate to growth under constant-but not fluctuating—conditions [60]; in contrast, growth was impaired in the present study.

The increase in NPQ over time in *C. vulgaris* in response to metformin exposure suggests that the cells were less able to process the same amount of light energy over time. qI results from a long-term downregulation of photosystem II, and likely occurs through a combination of photoprotection and photoinhibition [61]. In the present case, the increase in NPQ likely accompanied photoinhibition (and damage to photosystem II) rather than photoprotective quenching, given that a reduction in F_v/F_m was observed. That is, F_o should decrease in direct proportion to F_m in the case of photoprotective quenching, while increased F_o is accompanied by decreased F_m when quenching is achieved through photoinhibition, thus reducing (F_m - F_o)/ F_m , or F_v/F_m [61].

Interestingly, activation of SnRK1 (SNF1-related protein kinase 1)—an ortholog of AMPK, the target of metformin—is also associated with a decrease in the rate of photosynthesis in plants [62, 63]. Although not completely understood, SnRK1 activation occurs under energy deficit conditions, for example in darkness; to conserve energy, energetically demanding processes such as protein synthesis and cell division are inhibited [64, 65]. In contrast, SnRK1 is inhibited by sugars [66]. Recently, genes for SnRK1 were discovered in eukaryotic algae [67], and regulatory proteins involved in SnRK1 signaling pathways have been identified in both green algae and diatoms [68, 69]. Since energy sensing kinases are highly conserved among humans (AMPK), plants (SnRK1), and phytoplankton (SnRK1) [70, 71] the observed effects on growth and photosynthesis in *C. vulgaris* could be due to activation of SnRK1. It is possible that SnRK1 is implicated in stress responses to emerging contaminants such as pharmaceutical by algae more broadly, and therefore identifying factors that activate SnRK1 pathways would be valuable in assessing ecosystem effects of PPCPs.

Alternatively, it is possible that metformin inhibits complex I of the electron transport chain within algal mitochondria, resulting in a suppression of photosynthesis due to a reduction in interactions between organelles, either driven by inhibition of complex I or through activation of SnRK1. SnRK1 has been implicated in energy signaling between organelles [72]; in addition, a significant role for metabolic interactions between the chloroplast and the mitochondrion was demonstrated in mutants lacking complexes I and IV activity in another green alga, *Chlamydomonas*, which showed a 40% reduction in the light-saturated capacity of O_2 evolution [73]. This reduction in the rate of photosynthesis is similar to the ~20% reduction in ETR_{max} in this study, although more dramatic, consistent with the idea that complex I may be a target in eukaryotic phytoplankton. Another example comes from the diatom, *Phaeodacty-lum tricornutum*, where upregulation of alternative oxidase (AOX) activity in the mitochondrion occurred in response to iron-deplete conditions [74] suggesting a possible way to reroute photosynthetic electrons towards respiration rather than photosynthesis [75].

The fact that inhibition of complex I by metformin is observed across different phylogenetic groups including the yeast, Pichia pastoris, the bacterium, Escherichia coli, and the bovine heart mitochondria [49, 51] indicates that the biguanide binding site is found in the core subunit that is conserved across taxa, not just in mammalian cells [76]. Although the reported effect concentrations tend to be substantially higher than those found in the extracellular matrix or ambient environment (e.g., IC₅₀ of 22.6 ±4.3 mM in P. pastoris [49]), the 1000-fold accumulation of metformin within the mitochondrial matrix [51, 77] makes even low levels physiologically relevant. Moreover, in the environment, persistent exposure to metformin (for example in slow-moving reservoirs or lakes or else due to constant input fluxes [78]) could potentially lead to bioaccumulation and negative effects as transporters are synthesized. The issue of bioaccumulation is complex, since the multitude of pharmaceutically active ingredients with highly diverse chemistries show different propensities for bioaccumulation or biomagnification in the food web [79]. A recent study in Taihu Lake in China suggested that trophic magnification of several pharmaceuticals including norloxacin, ciproflaxin, and tetracycline are low [80, 81]. Persistent exposure to the antibiotic, ciproflaxin, downstream of a wastewater treatment plant in Kansas, United States, however, was associated with shifts in algal assemblages, a loss of genus richness, decrease in biomass yield, and changes in algal biovolumes [13].

In addition to deleterious effects on biota, exposure to chemical contaminants can lead to physiological acclimation (at sub-lethal concentrations) as well as evolutionary adaptation [82], which enables certain genotypes to survive, owing to the occurrence of beneficial mutations; this phenomenon is referred to as evolutionary rescue [83]. In the face of environmental deterioration, species–including microalgae [84, 85]–can become resistant to xenobiotics,

which likely underlies shifts in algal assemblages observed downstream of wastewater treatment plants [13] as sensitive species are lost and resistant species persist. Evolutionary rescue appears to be strongest when effects are slow, with repeated exposure, and dispersal is modest [83]. Although exposures to low levels of a variety of pharmaceuticals have unknown consequences for aquatic biota [86], evolutionary rescue could be invoked in response to circumstance where risks are high, for example if chemical mixtures, particular life stages, or different physiological stressors render certain genotypes sensitive to exposure in ways that cannot be overcome by short-term physiological acclimation.

The present study is limited by the fact that effects of long-term exposures were not assessed; however, exposure to pharmaceutical compounds and other chemicals of emerging concern can occur over short or long time periods, making studies of both types necessary to evaluate ecological effects of unregulated contaminants. In addition, this study focuses on a single pharmaceutical compound in isolation, exposed to a single target organism; this clearly does not adequately represent natural environmental conditions [81], and therefore the results should be interpreted as such. Nevertheless, our approach was to determine whether exposure to an unusual pharmaceutical compound would elicit effects on photosynthesis in suspended algal populations, and the results shed some light on potential mechanisms.

Finally, recent evidence has shown that different biguanides [including antimalarial drugs, anti-hyperglycemic drugs (metformin, phenformin, and buformin), and drugs for the treatment of cardiovascular disease and cancer; [51]) inhibit complex I to varying degrees. We tested the effects of one biguanide on photosynthetic performance; it is possible that other biguanides could have different effects on complex I or in the activation of SnRK1, which could influence photosynthetic processes differently, meriting further investigation into their toxicological effects. Future work should explore the mechanisms by which SnRK1 activation occurs, as well as responses by SnRK1 to environmental toxicants in order to better understand broader impacts on photosynthetic processes.

Supporting information

S1 Table. Cell concentration data used to calculate experiment growth rates. Cell concentrations determined at each experimental time point and used to calculate specific growth rate. Cells were enumerated using a Coulter counter. Triplicate samples were averaged, and cell densities determined by accounting for dilution of the samples prior to enumeration. (XLSX)

S2 Table. Photosynthetic performance data showing response to metformin additions. Primary data produced by a Pulse Amplitude Modulated fluorometer for experimental time points. The data were collected for 120 h before spiking the cultures with three concentrations of metformin, plus a control. Photosynthetic characteristics were tracked for 1, 3, and 5 h after the spike and each 24 h thereafter (until 96 h after the spike). F = fluorescence; F_m ' = effective maximum fluorescence; F_v/F_m = yield of PSII; ETR = electron transport rate; PAR = photosynthetically active radiation; NPQ = nonphotochemical quenching. Primary ETR and NPQ data from light curves are also included. (CSV)

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