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Comparative sensitivity to the fungicide tebuconazole of biofilm and plankton microbial communities in freshwater ecosystems

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Highlights

- Biofilm and plankton microbial communities were exposed to Tebuconazole (TBZ).
- TBZ had effects on biofilm function including respiration and photosynthetic activity.
- Biofilm communities from polluted sites exhibited induced tolerance to TBZ.
- Plankton communities responded to TBZ by increasing bacterial cell densities.
- TBZ toxicity depends on the nature of microbial communities and pre-exposure to TBZ.

Abstract

Stream and lake ecosystems in agricultural watersheds are exposed to fungicide inputs that can threaten the structure and functioning of aquatic microbial communities. This research analyzes the impact of the triazole fungicide tebuconazole (TBZ) on natural biofilm and plankton microbial communities from sites presenting different degrees of agricultural contamination. Biofilm and plankton communities from less-polluted (LP) and polluted (P) sites were exposed to nominal concentrations of 0 (control), 2 and 20 $\mu\text{g TBZ L}^{-1}$ in 3-week microcosm experiments. Descriptors of microbial community structure (bacterial density and chlorophyll-a concentration) and function (bacterial respiration and production and photosynthesis) were analyzed to chart the effects of TBZ

and the kinetics of TBZ attenuation in water during the experiments. The results showed TBZ-induced effects on biofilm function (inhibition of substrate-induced respiration and photosynthetic activity), especially in LP-site communities, whereas plankton communities experienced a transitory stimulation of bacterial densities in communities from both LP and P sites. TBZ attenuation was stronger in biofilm (60–75%) than plankton (15–18%) experiments, probably due to greater adsorption on biofilms. The differences between biofilm and plankton responses to TBZ were likely explained by differences in community structure (presence of extracellular polymeric substances (EPS) matrix) and microbial composition. Biofilm communities also exhibited different sensitivity levels according to their in-field pre-exposure to fungicide, with P-site communities demonstrating adaptation capacities to TBZ. This study indicates that TBZ toxicity to non-targeted aquatic microbial communities essentially composed by microalgae and bacteria was moderate, and that its effects varied between stream and lake microbial communities.

Keywords

Algae; Bacteria; Fungicide; Photosynthetic efficiency; Adaptation; Substrate-induced respiration

1. Introduction

Freshwater ecosystems in agricultural watersheds are widely exposed to multiple pesticidecontamination events, including fungicides which pose major ecotoxicological threats to aquatic microbial community biodiversity and function (e.g. Montuelle et al., 2010 and Rasmussen et al., 2012a). The intensive use of modern azole fungicides like fluconazole, propiconazole and tebuconazole (TBZ) (Richardson, 2009) has prompted studies aiming to understand their effects on aquatic detritivores and microbial decomposers (Bundschuh et al., 2011 and Rasmussen et al., 2012b). TBZ has been found in numerous freshwater ecosystems in Europe, including rivers and lakes, and is reported as relatively persistent even after wastewater treatment (Berenzen et al., 2005 and Kahle et al., 2008).

The main aquatic microbial communities potentially threatened by TBZ are fungi. TBZ affects fungal activity-related processes such as litter decomposition (Rasmussen et al., 2012b and Bundschuh et al., 2011) and inhibits ergosterol biosynthesis, which hampers fungal mycelium growth (Copping and Hewitt, 1998). Exposing submerged leaves to TBZ led to lower fungal biomass accumulation and reduced the ability of microorganisms to decompose cellulose and hemicellulose compounds (Artigas et al., 2012). Important bacterial groups such as nitrogen-fixing and nitrifying bacteria can also be directly or indirectly affected by fungicides (Johnsen et al., 2001). Specifically, TBZ has been reported to stimulate ammonifying bacteria populations at early exposure in soils and nitrifying bacteria in later exposure phases (Chen et al., 2001, Cycoń et al., 2006 and Muñoz-Leoz et al., 2011). This stimulating effect on bacterial groups could be associated with increased levels of nutrients and energy sources released from dead fungal hyphae killed by TBZ (Cycoń et al., 2006) or by chemicals present in the fungicide formulation becoming available to bacteria (Muñoz-Leoz et al., 2011). Although such results suggest that similar effects will be found in aquatic ecosystems, with consequences on microbial processes, only a few studies have explored the effects of TBZ on aquatic microbial communities other than fungi in biofilms and plankton (Tlili et al., 2011a and Artigas et al., 2012).

Microbial community exposure to TBZ varies between stream and lake ecosystems. Pesticide mobilization in agricultural watersheds mostly depends on drift, run-off and drainage processes from the watershed (Kolpin et al., 1996), and often coincides with fungicide applications in conjunction with cultural practices (Rabiet et al., 2010). However, the arrival of fungicides into surface water

depends on watershed topography (especially field slope), soil characteristics (i.e. organic carbon-water binding coefficient), and the half-life of the molecule (49–610 days for TBZ in aerobic soils; Strickland et al., 2004). Once in the aquatic ecosystem, hydrological conditions in lotic ecosystems mostly generate peak exposure scenarios (lasting a few hours to days) at high contaminant concentrations, whereas lentic ecosystems establish a more chronic exposure scenario (several months) at lower contaminant concentration (Simmons and Wallschlager, 2005). Consistently with this pattern, TBZ concentrations recorded in rivers (often at $\mu\text{g L}^{-1}$) exceed those found in lakes (often at ng L^{-1} ; Richardson, 2009, Rabiet et al., 2010 and Kahle et al., 2008). However, TBZ concentrations in rivers are highly variable over time (Rabiet et al., 2010) and chronic exposure scenarios to low fungicide concentrations can also be detected (i.e. Reilly et al., 2012). Knowing that level and duration of exposure of microbial communities to the pesticide are important factors determining the adaptation processes of microorganisms (Le Jeune et al., 2007 and Pesce et al., 2009), microbial communities from stream biofilms and lakes are likely to display contrasted susceptibility to this fungicide.

Comparative studies assessing pollutant toxicity between lotic (stream) and lentic (lake) ecosystems are rare in the literature (Munawar et al., 1999 and Simmons and Wallschlager, 2005), despite their potential utility for better understanding how the ecological, hydrological and biogeochemical characteristics of each type of ecosystem can modulate pollutant ecotoxicity. In this global framework, we adopted an experimental approach to investigate the effects of TBZ on non-targeted microalgal and bacterial communities living in stream biofilms and lake plankton samples collected in natural ecosystems. For each ecosystem (stream and lake), samples were collected in sites qualified as “less polluted” and “polluted” based on previous in situ measurements in order to assess the influence of pre-exposure to TBZ on their susceptibility to the fungicide. The responses of all these communities to TBZ (at two doses, 2 and 20 $\mu\text{g TBZ L}^{-1}$) were examined using a range of structural and functional descriptors. This experimental scheme was thus designed to test the respective effects of community lifestyle (benthic versus planktonic) and pre-exposure to the pollutant on the responses of non-targeted aquatic microbial communities to TBZ.

2. Materials and methods

2.1. Biological communities

Biofilm and plankton microbial communities from less-polluted (LP) and polluted (P) sites were obtained from stream and lake ecosystems in eastern France. Biofilms were obtained from two sites (upstream in Saint Joseph and downstream in Saint Ennemon) of Morcille River (Beaujolais vineyard region, 46.15°N 4.60°E), a siliceous stream composed by three nested sub-watersheds that experiences an upstream-to-downstream gradient of agricultural contamination (Rabiet et al., 2010 and Montuelle et al., 2010). This pollution gradient is characterized by increases in dissolved organic carbon, phosphate (125% to 300% increase between upstream and downstream sites, respectively) and copper, arsenic, herbicide (diuron and its breakdown products) and fungicide (azoxystrobin, carbendazim, tebuconazole, procymidone and dimetomorph) concentrations (Montuelle et al., 2010). Previous surveys have shown that TBZ concentrations ranged from 0.001–0.1 $\mu\text{g L}^{-1}$ at the upstream LP site of Saint Joseph to 0.01–4 $\mu\text{g L}^{-1}$ at the downstream P site of Saint Ennemon (Margoum C., pers. comm.). Artificial glass substrata (20 cm² tiles) were left to colonize for one month at both study sites. Biofilm colonization was conducted during late summer, when the highest TBZ contamination peaks had been detected at the P site (Montuelle et al., 2010). The resulting biofilms from LP and P sites were collected, transported cold (4 °C) to the laboratory, and placed in indoor stream channels for TBZ experiments.

Plankton communities were obtained from two different peri-alpine lakes experiencing contrasting degrees of agricultural contamination — one less polluted (Lake Aiguebelette, mean TBZ concentration $< 0.001 \mu\text{g L}^{-1}$) and one polluted (Lake Geneva, mean TBZ concentration = $0.001 \mu\text{g L}^{-1}$). Lake Aiguebelette (LP site) is situated in a small forested watershed (70 km²) bearing low anthropogenic pressure, whereas Lake Geneva (P site) has a much larger watershed (7395 km²) bearing high pressure due to urbanization and agriculture. Samples (400 L) from each lake were collected from the 0–50 cm layer and filtered through a 50 μm pore-size filter to remove zooplankton. Samples were collected at a central part in each lake (Lake Aiguebelette: 5°47'56", 45°33'02", Lake Geneva: 6°35'34", 46°27'22"). Samples with plankton communities from LP and P sites were transported in plastic containers, stored overnight at 4 °C, and used for subsequent TBZ experiments in indoor microcosms.

2.2. Laboratory experiments

The experiments were performed in glass indoor stream channels ($L \times W \times D = 63 \text{ cm} \times 11 \text{ cm} \times 4 \text{ cm}$) for biofilms and in transparent polycarbonate bottles (20 L) for plankton. Each type of microbial community (LP-site biofilm, P-site biofilm, LP-site plankton, P-site plankton) was subjected to three different treatments consisting of control (non-contaminated), low-TBZ dose ($2 \mu\text{g L}^{-1}$) and high-TBZ dose ($20 \mu\text{g L}^{-1}$) exposure.

For biofilms, each microbial community (LP and P sites) and treatment was tested in three independent channel replicates (18 channels in total), each connected to a separate 5 L reservoir (Pyrex, USA) by means of an aquarium pump (NEWA MJ 750). Water temperature was set to 20 °C and flow conditions were set to 0.19 L s^{-1} , all under a 12 h/12 h light/dark cycle. Before TBZ contamination, stream biofilms were allowed to adapt for one week to the indoor channels by recirculating tap water (previously dechlorinated using an active carbon filter). After TBZ contamination, the water was not replaced during the 22-day experiment but N and P were added to prevent nutrient limitation in the biofilms (weekly added quantities: $225 \mu\text{g N L}^{-1}$ and $65 \mu\text{g P L}^{-1}$). For plankton, LP and P-site communities were exposed to the three different treatments in triplicate. Bottles containing plankton communities (18 in total) were incubated at 20 °C under a 12 h/12 h light/dark cycle and without nutrient addition. Air circulation was supplied to prevent anoxia in the plankton communities.

Solutions of TBZ (C₁₆H₂₂CIN₃O, Sigma Aldrich; 10 mg L^{-1} in deionized water) were added to the biofilm and plankton communities, and biological community and water samples were taken on days 0, 3, 7, 10, 14, 17, 22 post addition. For biofilms, samples were scraped from glass tiles and suspended in the water of the corresponding channel for subsequent biological analyses. Plankton samples were taken with a pipette directly from bottles after stirring. Assays to determine substrate-induced respiration, bacterial production, chlorophyll concentration and photosynthetic activity were processed immediately after sampling, while samples taken to determine bacterial density were fixed in 2% formaldehyde until analysis. Water samples to determine nutrient concentrations were taken into glass flasks (100 mL) and analyzed immediately. Further water samples taken to analyze TBZ concentration (100 mL) and TBZ metabolites (500 mL) were stored frozen ($-20 \text{ }^\circ\text{C}$) until analysis.

2.3. Chemical analyses

Water temperature and pH were measured at each sampling time using portable meters (WTW 3110, Germany). Filtered (0.45 μm) water samples were analyzed to determine the concentration of inorganic nutrients [soluble reactive phosphorus (SRP), dissolved inorganic nitrogen (DIN = ammonia + nitrite + nitrate) and dissolved organic carbon (DOC)] following French standard operating procedures.

TBZ quantification was determined by UPLC (ACQUITY, Waters, Manchester, UK) using a Hypersil Gold PFP stationary phase (100 m \times 2.1 mm \times 1.9 μm) with detection at 220 nm. The mobile phase consists of 50% acetonitrile/50% H₂O at 0.5 mL/min. Analysis was performed at 30 °C with an injection volume of 6 μL . TBZ degradation metabolites were determined by LC/Q-TOF (Waters/Micromass, Manchester, UK). For metabolite analyses, 25 μL were injected onto a reverse-phase column (X terra, Waters, C18, 3.5 μm , 100 mm \times 2.1 mm) at a flow rate of 0.3 mL min⁻¹ with a mobile phase composed of acetonitrile (solvent A) and acidified water (solvent B: formic acid 0.3% v/v; pH 2.6). The gradient was 5% A from 0–5 min, 5–95% A (linear) from 5–15 min, and 95% A from 15–25 min. The desolvation chamber and ion source block temperatures were set at 350 °C and 100 °C, respectively. The voltages for the probe and ion source components were set to 2.1 kV for the stainless steel capillary, 35 V for the sample cone, and 2 V for the extractor cone. A collision energy gradient (15–35 eV) was used for preliminary fingerprints, and specific energies of 15–45 eV were used to unambiguously assign the MS/MS fragments.

2.4. Biological analyses

2.4.1. Bacterial density

Bacterial densities were quantified by DAPI staining for biofilms (Porter and Feig, 1980) and by flow cytometry for plankton communities (Tadonléké et al., 2005). For biofilms, suspensions corresponding to 1 cm² of scraped biofilm were sonicated for 2 min to facilitate disaggregation, diluted 1:2, and stained for 5 min with 4,6-diamidino-2-phenylindole (DAPI, Sigma-Aldrich) at a final concentration of 2 $\mu\text{g mL}^{-1}$. Suspensions were then filtered through 0.2 μm irgalan black-stained polycarbonate membranes (Nuclepore, Whatman, England) and mounted on microscope slides. Fifteen fields were counted per filter, accounting for a total of 400–800 cells, using a fluorescence microscope (Nikon Labophot-2, Japan) at 1000 \times magnification. Bacterial densities were calculated as cells per unit of biofilm surface area in cm².

For biofilms, additional bacterial suspensions (1 cm² of biofilm surface area) were diluted 1:2 and stained with propidium iodide (PI) and SYTO 9™ (BacLight) for 15 min at room temperature in the dark. Samples were subsequently filtered through 0.2 μm irgalan black-stained polycarbonate membranes and mounted on microscope slides. Counting of dead (red-stained) and live (green-stained) bacterial cells was performed on 20 random microscope fields using an epifluorescence microscope (Nikon Labophot-2, Japan) at 1000 \times magnification (Freese et al., 2006). The ratio between live and dead bacterial cells (L/D bacterial ratio) was calculated.

For plankton, total bacterial abundance was determined using a flow cytometer (FacsCalibur, Becton-Dickinson) equipped with a laser emitting at 488 nm. Samples were stained with SYBR Green II (Invitrogen), and an internal standard (yellow–green 0.92 μm fluorescent latex beads) was included. Samples were run at low speed (10–12 $\mu\text{L min}^{-1}$), and bacterial abundances were calculated as cells mL⁻¹.

2.4.2. Bacterial activity

Bacterial activity in biofilms was estimated using substrate-induced respiration assays (Campbell et al., 2003) as modified by Tlili et al. (2011b). Bacterial activity in plankton was estimated by means of bacterial production using ³H-leucine incorporation assays (Tadonl  k   et al., 2005). For biofilms, deep-well microtiter plates were filled with biofilm suspensions (500  L) of the corresponding treatment, and 30  L of glucose (6.2 mg C per well) was added. Controls without glucose were also included to determine basal respiration. Immediately after inoculation, the detection microplate (containing cresol red dye) was sealed to the deep-well plate with a silicone seal. The assembled MicroResp system was incubated in the dark at room temperature for 15 h. After incubation, detection microplate absorbance was measured at 590 nm (Biotek Synergy HT spectrophotometer), amounts of released CO₂ were calculated, and the results were expressed in  g CO₂ mgAFDW⁻¹ h⁻¹.

For plankton, bacterial production was estimated from ³H-leucine (Leu) incorporation by bacteria after prior determination of saturating concentration (Tadonl  k   et al., 2005 and Tadonl  k   et al., 2009). The samples were counted in a scintillation counter (2100-TR, Packard Instruments). The results were expressed in pmol [³H]leucine L⁻¹ h⁻¹ incorporated.

2.4.3. Chlorophyll-a concentration and photosynthetic activity

“In vivo” measurements of chlorophyll-a fluorescence in biofilms and plankton communities were performed using the Phyto-PAM fluorometer (Heinz Walz, Germany). In dark-adapted communities (ca. 30 min), Phyto-PAM excites chlorophyll fluorescence at different measuring lights (470, 520, 645 and 665 nm), and the reading at 665 nm (F₀) is proportional to chlorophyll-a concentration (Rysgaard et al., 2001). Photosynthetic efficiency measurements were based on the maximal quantum yield (YII) of algae which reflects number of functional photosystem II (Schreiber et al., 2002). A single saturation pulse was applied to calculate maximal quantum yield, which was later computed considering F₀ (steady-state fluorescence) and F_m (maximum fluorescence after saturation pulse) parameters.

2.5. Statistical analyses

Three-way ANOVA tests were applied on biological descriptors and on water chemical characteristics to test differences between control and TBZ-contaminated treatments in LP and P-site communities over time. Biological descriptors were classified as functional (substrate-induced respiration, bacterial production, L/D cell ratio, photosynthetic efficiency) and structural (bacterial density and chlorophyll-a concentration). The sources of variation were calculated for Time (T), Treatment (control, 2 and 20  g L⁻¹) and Site (clean vs. polluted) and their corresponding interactions. Multiple comparison tests (Tukey test) were also applied on the three-way ANOVA to determine differences between treatments. Prior to all ANOVA analyses, biological and chemical variables were log-transformed and resultant probability values adjusted by the Dunn-Sidak correction. SPSS Statistics 17.0 was used for analyses.

The percentage of difference (D) of biological descriptors in TBZ treatments (D_{treatment}) relative to controls (D_{control}) was calculated using the following equation:

$$D(\%) = \frac{D_{\text{treatment}} - D_{\text{control}}}{D_{\text{control}}} \times 100.$$

3. Results

3.1. Physical and chemical characteristics of the water

Water temperature and pH conditions were similar between biofilm (Temperature = 20.4–20.6 and pH = 8.5–8.6) and plankton (Temperature = 20.5–20.9 and pH = 8.1–8.3) TBZ experiments, whereas water nutrient concentrations (DOC, DIN and SRP) varied between samples from LP and P sites (biofilms and plankton) and between TBZ treatments (only biofilms) (Fig. 1, Table 1).

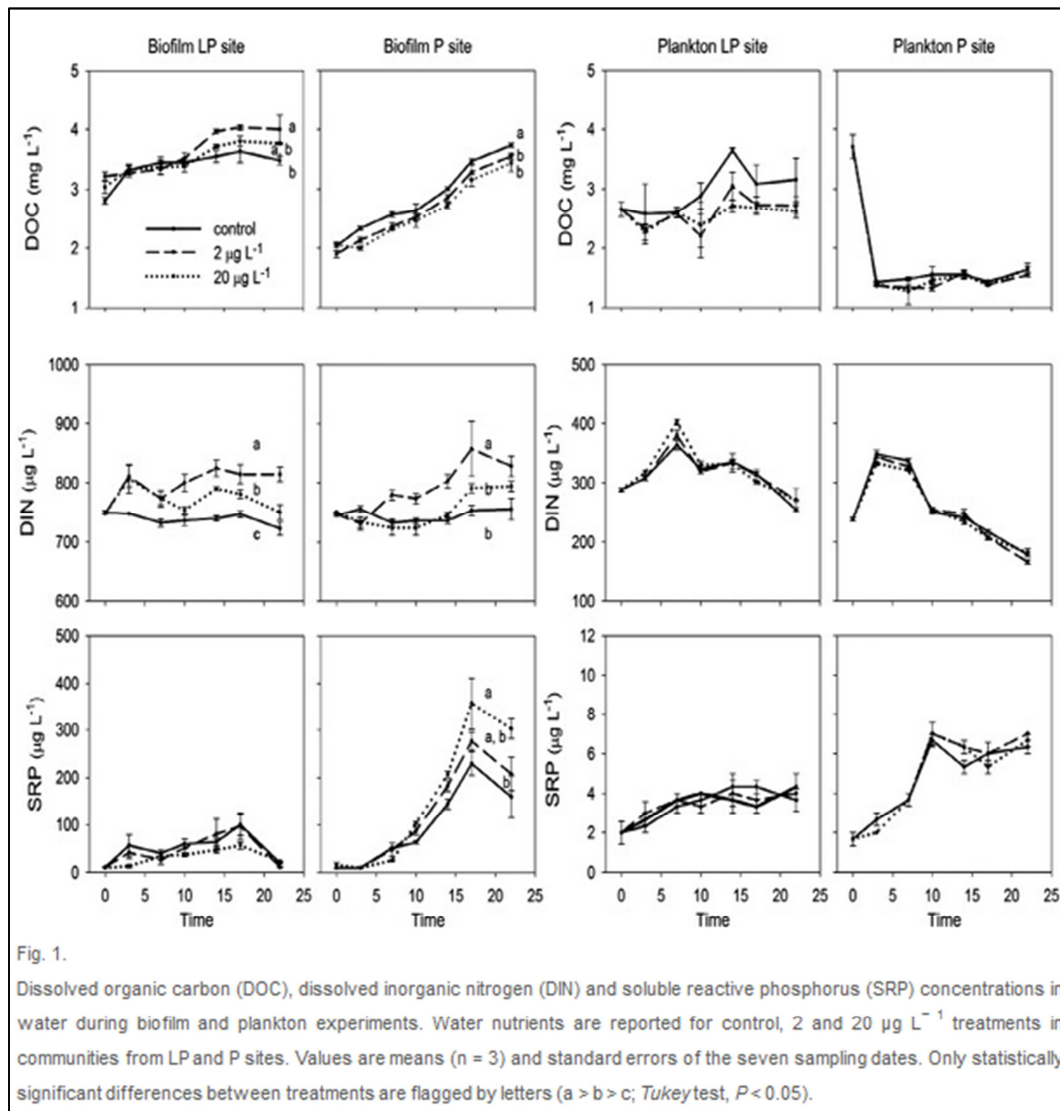


Fig. 1.

Dissolved organic carbon (DOC), dissolved inorganic nitrogen (DIN) and soluble reactive phosphorus (SRP) concentrations in water during biofilm and plankton experiments. Water nutrients are reported for control, 2 and 20 µg L⁻¹ treatments in communities from LP and P sites. Values are means (n = 3) and standard errors of the seven sampling dates. Only statistically significant differences between treatments are flagged by letters (a > b > c; Tukey test, P < 0.05).

Water nutrient concentration comparisons between LP and P sites revealed that DOC concentration was greater in LP-site communities whereas SRP concentration was greater in P-site communities during the 22-day experiment (Fig. 1). Conversely, water DIN concentration was similar between LP and P-site biofilms but not LP and P-site plankton (average DIN in LP 313.4 ± 7.3 µg L⁻¹ and DIN in P 259 ± 13 µg L⁻¹, Table 1). Water nutrient concentrations (especially SRP) tended to increase during experiments, except in P-site plankton communities where decreases of DOC (– 62% between Time 0 and Time 3) and DIN (– 26% between Time 3 and Time 22) concentrations were observed (Fig. 1).

Table 1.

Results of a three-way ANOVA test applied on biological descriptors of biofilm (B) and plankton (P) communities, as well as on water chemical characteristics, during TBZ experiments. Probability values (previously adjusted by the Dunn-Sidak correction) are given for each of the sources of variation tested [Time (T), Site (S), Treatment (TBZ)] and their interactions. Statistically significant *P* values are highlighted in bold.

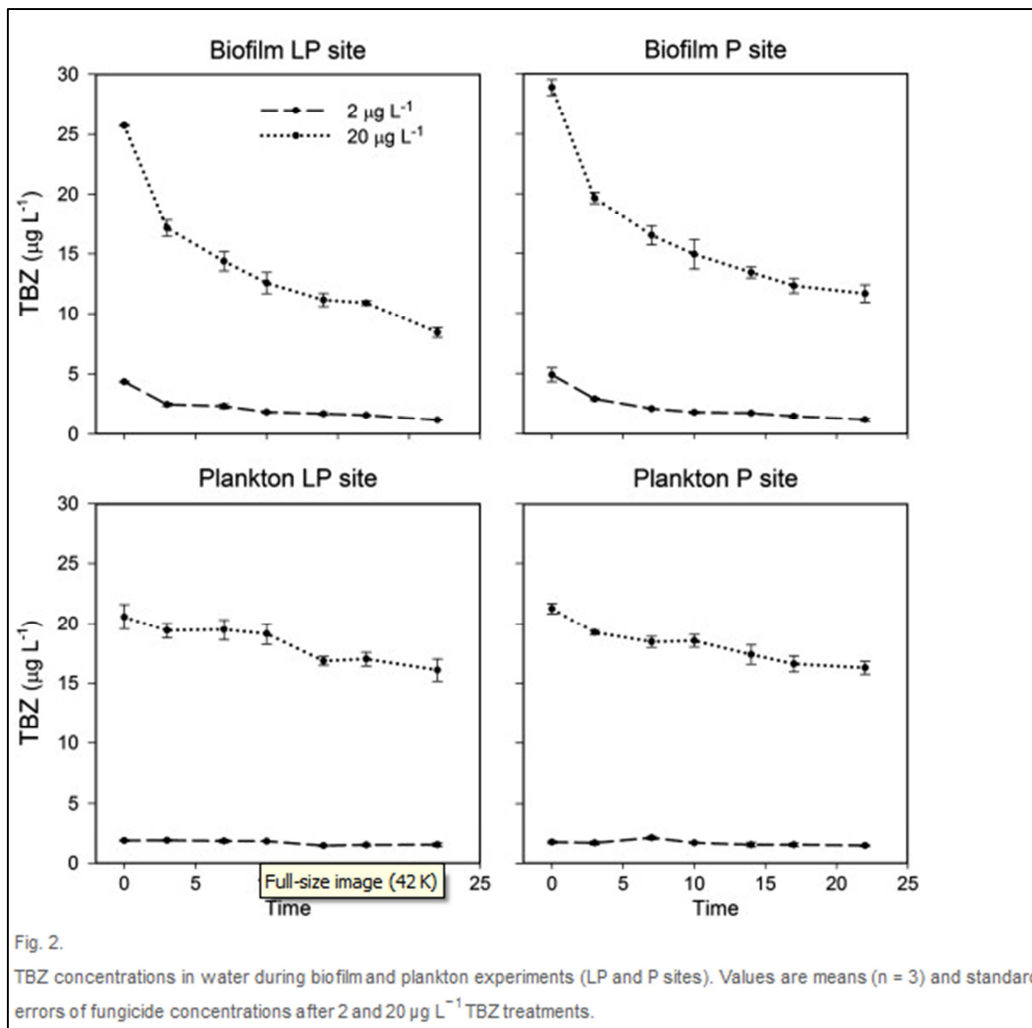
Biological descriptors									
	Bacterial densities		Live/Dead	Bacterial activity		Chlorophyll- <i>a</i> concentration		Photosynthetic efficiency	
	B	P		B	P	B	P	B	P
T	< 0.001	< 0.05	< 0.001	< 0.001	< 0.001	< 0.001	< 0.001	< 0.001	< 0.001
S	0.645	0.315	< 0.001	< 0.001	0.190	< 0.001	< 0.001	0.300	< 0.001
TBZ	0.959	< 0.001	1	0.827	0.984	0.980	0.952	< 0.001	0.793
T × S	< 0.05	< 0.001	< 0.001	0.270	< 0.001	< 0.001	< 0.001	< 0.001	< 0.001
T × TBZ	< 0.05	0.226	1	0.226	0.920	1	1	< 0.001	0.746
S × TBZ	< 0.05	0.559	< 0.05	< 0.001	0.706	0.964	0.992	< 0.001	< 0.01
T × S × TBZ	< 0.05	0.878	1	0.201	0.448	1	0.816	< 0.001	0.979

Water chemistry						
	DOC		DIN		SRP	
	B	P	B	P	B	P
T	< 0.001	< 0.001	0.096	< 0.001	< 0.001	< 0.001
S	< 0.001	< 0.001	0.466	< 0.001	< 0.001	< 0.001
TBZ	< 0.05	0.267	< 0.05	0.991	0.337	0.988
T × S	< 0.001	< 0.001	< 0.05	< 0.001	< 0.001	< 0.001
T × TBZ	0.340	1	< 0.05	< 0.05	0.457	1
S × TBZ	< 0.001	0.538	0.674	< 0.05	< 0.001	0.868
T × S × TBZ	0.767	1	0.875	< 0.05	0.184	0.975

Water DOC, DIN and SRP concentrations varied between control and TBZ treatments in biofilm experiments but not in plankton experiments (Table 1). In biofilms, TBZ treatments (at 2 and 20 $\mu\text{g L}^{-1}$) had greater DOC, DIN and SRP concentrations in comparison to controls (except for SRP in LP, Tukey test, Fig. 1). Chronologically, the differences in nutrient concentrations between control and TBZ treatments for biofilms were observed first for DIN (Time 3) and later for DOC and SRP (Time 14) concentrations.

3.2. TBZ concentrations

TBZ concentration analyses in water (Fig. 2) confirmed the exposure of biofilm and plankton communities to low and high fungicide doses. TBZ concentrations at the beginning of experiments (Time 0) were $4.6 \pm 0.7 \mu\text{g L}^{-1}$ (low dose) and $27.3 \pm 1.9 \mu\text{g L}^{-1}$ (high dose) for biofilms, and $1.8 \pm 0.2 \mu\text{g L}^{-1}$ (low dose) and $21 \pm 1.3 \mu\text{g L}^{-1}$ (high dose) for plankton. TBZ concentrations decreased over the three-week experiment, revealing an attenuation of fungicide concentration in the water phase. However, this decrease was greater in biofilms (range 60–75% between Time 0 and Time 22) than in plankton communities (range 15–18%). Secondary metabolites of TBZ, such as hydroxyl-tebuconazole and desbutyl-tebuconazole, were detected in water at the end of biofilm experiments (Time 22, data not shown).



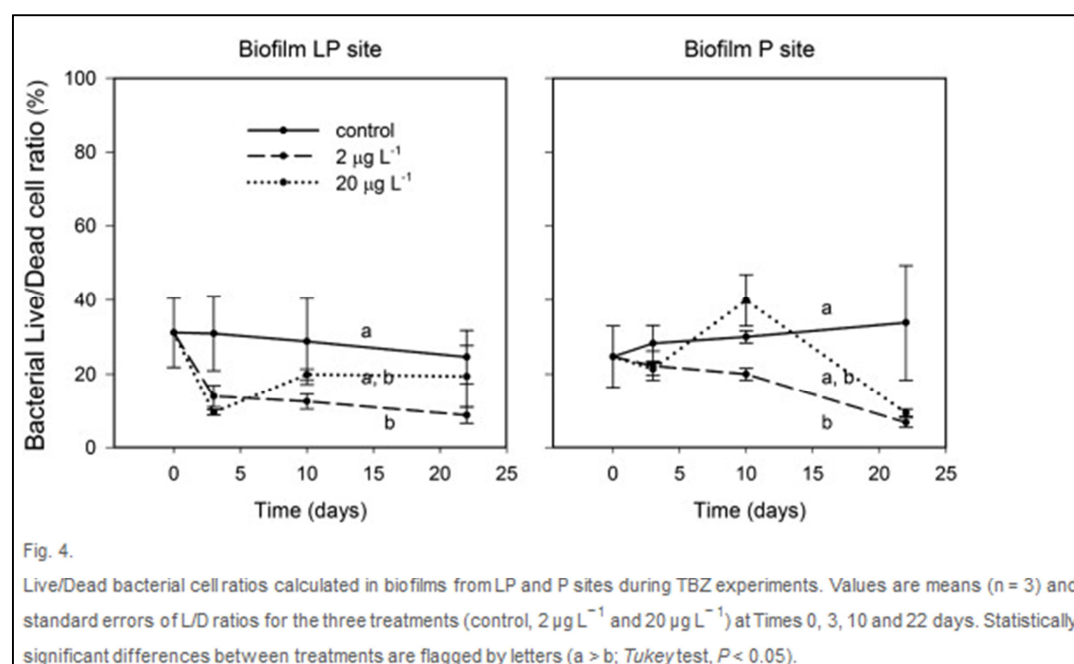
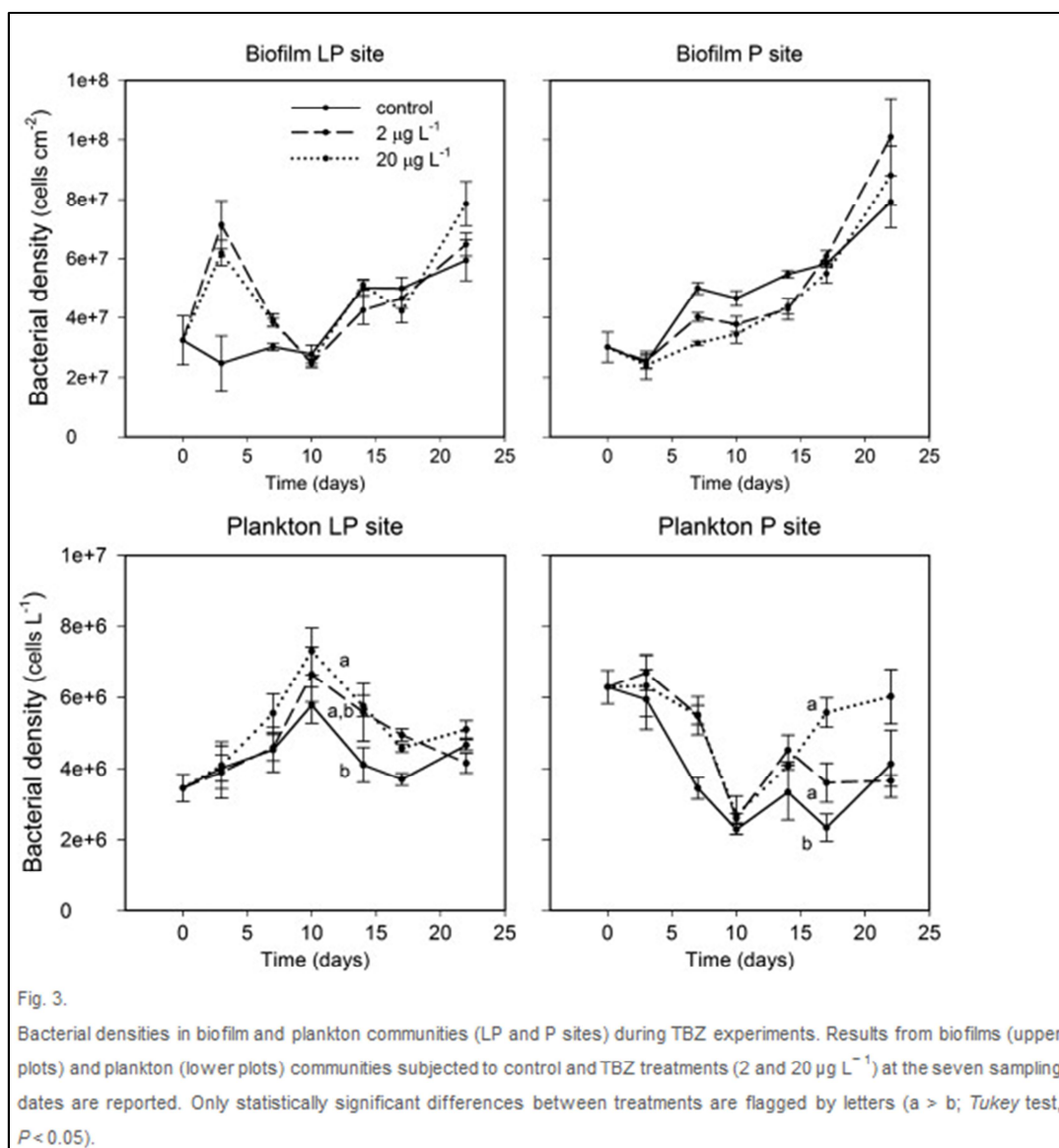
3.3. Bacterial communities

3.3.1. Bacterial densities

Bacterial densities at the beginning of the experiment (Time 0) were similar between LP-site (3.2×10^7 cells cm⁻²) and P-site (3×10^7 cells cm⁻²) biofilms but different between LP-site (3.4×10^6 cells L⁻¹) and P-site (6.3×10^6 cells L⁻¹) plankton communities. Cell densities showed significant temporal variation in biofilm and plankton microcosm experiments, but only plankton communities were affected by TBZ (Table 1).

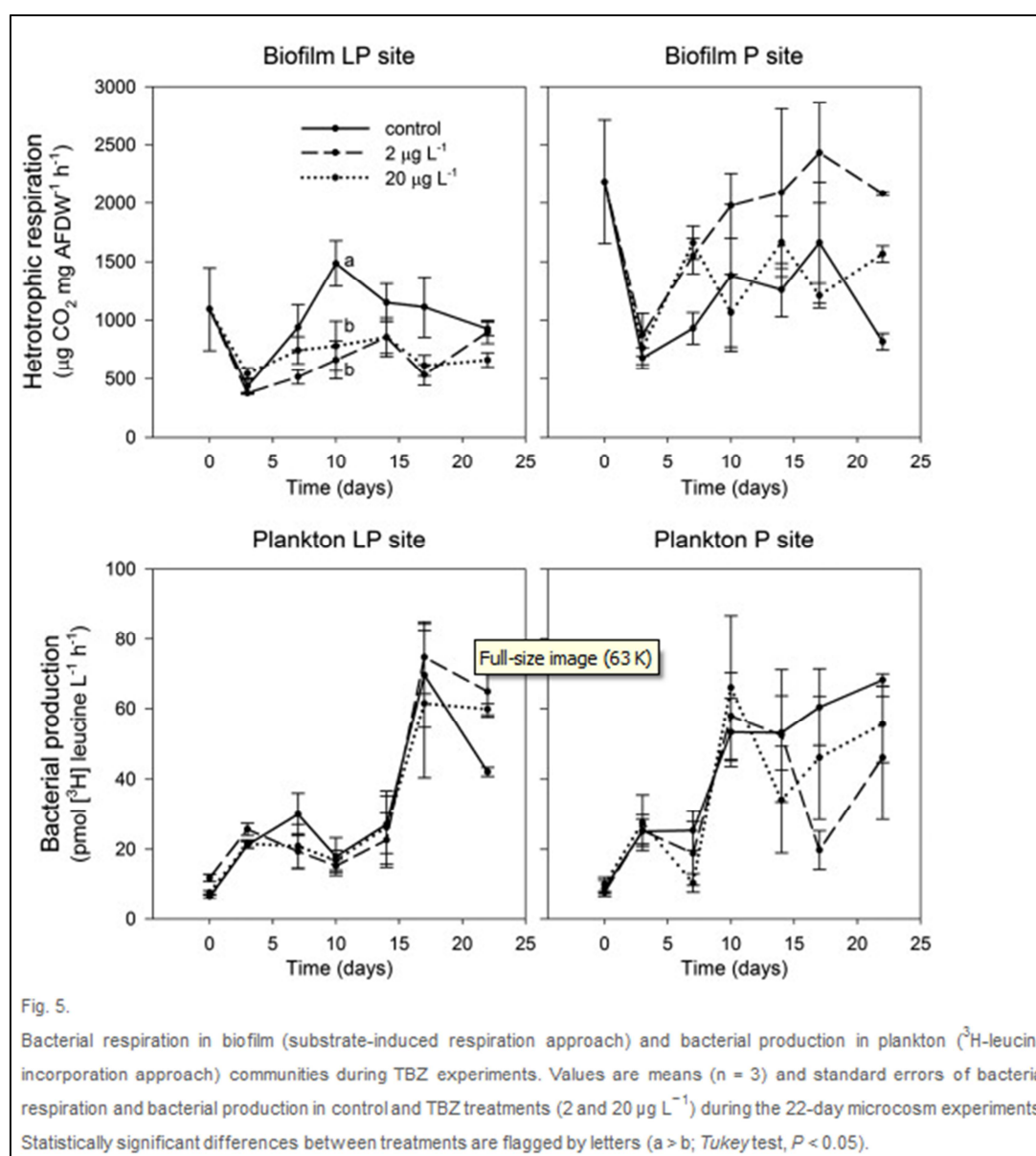
TBZ increased bacterial densities in plankton communities from the two studied sites (Fig. 3). The increases were globally stronger at the high TBZ dose (+ 21% in LP-site plankton and + 48% in P-site plankton; Fig. 3, Tukey $P < 0.05$) but appeared relatively transient over time (weak differences at Time 22). TBZ effects on biofilm bacteria were also significant but were site- and time-dependent (Table 1). While TBZ treatment resulted in increases in bacterial densities in LP-site biofilms at Time 3 and Time 6, it tended to decrease bacterial densities in P-site biofilms between Time 6 to Time 10.

L/D cell ratios (only available for biofilms) in the controls averaged 0.23 for the LP site and 0.31 for the P site between Time 0 and Time 22 (Fig. 4). The presence of TBZ led to different patterns of decrease in L/D cell ratios in communities from the LP and P sites (average - 48% and - 35%, respectively) (Site \times TBZ effect, Table 1). These effects on biofilm L/D cell ratios were fairly persistent over time and were constant between TBZ doses



3.3.2. Bacterial activity

SIR rates measured in control biofilms from the two sites remained fairly constant over the experiment ($1145 \pm 117 \text{ mgCO}_2 \text{ mgAFDW}^{-1} \text{ h}^{-1}$, average Time 0–Time 22), except for a sharp decrease at Time 3, independently of TBZ exposure (Fig. 5). TBZ decreased SIR in LP-site biofilms (range: -20 to -28%) compared to controls (Site \times TBZ effect, Table 1). This negative effect was more intense between Time 10 and Time 17 and was independent of TBZ dose (Tukey test, $P < 0.05$, Fig. 5). Conversely, TBZ tended to stimulate SIR in P-site biofilms (range: $+23$ to $+48\%$), but this effect was not statistically significant.

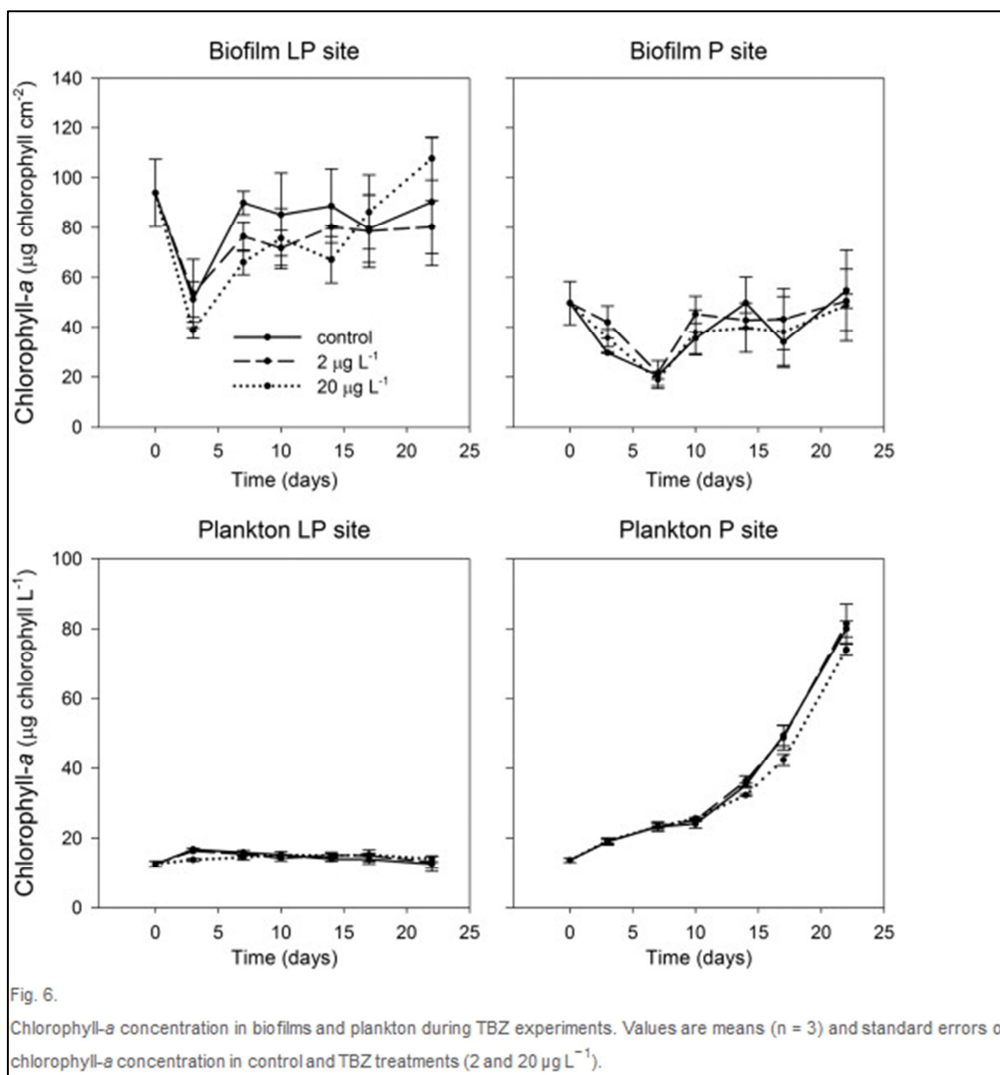


Bacterial production in plankton increased during the experiments but without an observable effect of TBZ (Table 1). Bacterial production in controls (including LP and P sites) ranged from 6 – $9 \text{ pmol } [^3\text{H}] \text{ leucine L}^{-1} \text{ h}^{-1}$ at Time 0 to 42 – $68 \text{ pmol } [^3\text{H}] \text{ leucine L}^{-1} \text{ h}^{-1}$ at Time 22. It was only at Time 22 (Fig. 5) when slight TBZ effects were observed in LP-site plankton communities (range: $+40$ to $+50\%$ compared to controls).

3.4. Algal communities

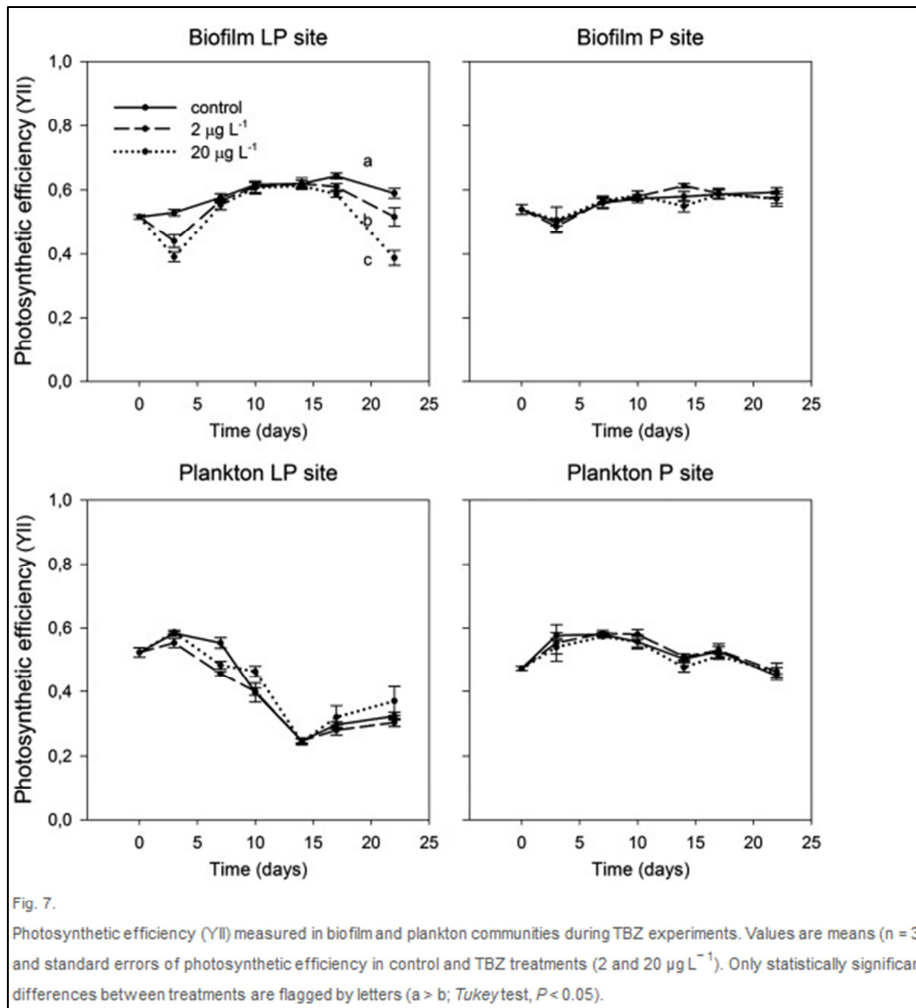
3.4.1. Chlorophyll-a concentration

Chlorophyll-a measurements in control treatments evidenced differences between biofilm and plankton LP and P-site communities (Table 1). For biofilms, average chlorophyll-a concentration in controls was greater in the LP site ($82.7 \pm 5.6 \mu\text{g chl cm}^{-2}$) than the P site ($39.2 \pm 3.9 \mu\text{g chl cm}^{-2}$), whereas for plankton communities average chlorophyll-a concentration was greater in the P site ($35 \pm 4.8 \mu\text{g chl L}^{-1}$) than the LP site ($14.5 \pm 0.5 \mu\text{g chl L}^{-1}$). Chlorophyll-a values were not affected by TBZ in either biofilms or plankton (Table 1), but showed significant temporal variations in the microcosms (Time \times Site effect; Table 1 and Fig. 6).



3.4.2. Photosynthetic efficiency

Photosynthetic efficiency (YII) values in control treatments stayed constant for biofilms (0.58 ± 0.01 LP site, 0.56 ± 0.008 P site) and plankton communities (0.42 ± 0.03 LP site, 0.52 ± 0.01 P site) over the 22 days of the experiment. TBZ only had a detectable effect on YII in LP-site biofilms (Site \times TBZ effect; Table 1, Fig. 7), which experienced a stronger decrease in YII at the higher TBZ dose treatment (-11% on average) than at the lower TBZ dose (-5.3%) (Tukey test, $P < 0.05$). These decreases were especially marked at the end of experiment between Time 17 and Time 22 (Fig. 7). Conversely, TBZ had no effect on YII in P-site biofilms or LP- and P-site plankton communities.



4. Discussion

This research has shown that biofilm and plankton microbial communities exposed to TBZ displayed different responses in some of the biological variables measured, suggesting that the effects of TBZ are modulated by lifestyle (benthic versus planktonic) and pre-exposure to pollutants (less polluted versus polluted).

Biofilm communities experienced functional alterations in presence of TBZ, as shown by a significant decrease in SIR and photosynthetic activity, especially in communities from the LP site. SIR inhibition might be partly explained by a direct TBZ effect on fungal activity in the biofilm (Table 2), as has been reported for soil microbial communities (Muñoz-Leoz et al., 2011). The observed decrease in SIR may be due to the death of fungi suggested by the increase in water DIN concentrations (Cycoń et al., 2006). Unfortunately, fungal biomass was not measured in this experiment, so we are unable to assess the proportion of dead fungi affecting SIR in biofilms. However, other studies support this view. For example, significant decreases in fungal biomass have been observed in benthic assemblages exposed for several weeks to TBZ concentrations (Bundschuh et al., 2011 and Artigas et al., 2012).

Interestingly, the reduction of SIR in biofilms coincided with increases (day 3 to 10) in bacterial cell mortality (L/D cell ratio), which also suggests a possible direct toxic effect of TBZ on bacterial community. In the same way, TBZ also impaired the photosynthetic efficiency of biofilms from the less polluted site. This effect was probably indirect (given the mode of action of TBZ; Copping and

Hewitt, 1998) and could be the result of indirect effects of pesticides on the relationships between biofilm components (Ricart et al., 2009). Previous studies have reported that the herbicide diuron affected both algal and bacterial activity in biofilms, suggesting that pesticides may cause a chain effect in biofilms (Ricart et al., 2009). The effects on algae and bacteria observed here might result from direct effects of TBZ on fungal communities and possibly effects on protozoa (Table 2). Conversely, biofilm communities from the polluted site showed more variable responses to TBZ without any significant influence on the biological parameters measured. This finding suggests that these biofilms that were pre-exposed to TBZ and to other types of fungicides (see Montuelle et al., 2010) may have developed adaptational capacities leading to an enhanced tolerance to the TBZ molecule (or “pollution-induced community tolerance”; Blanck, 2002), as already observed in biofilms for other types of pesticides (i.e. the herbicide diuron; Tlili et al., 2011a).

Table 2.
Summary of effects of TBZ in bacterial and algal communities with the hypothetical effects on fungal and protozoan communities (marked in gray) and their interactions in biofilm and plankton microbial communities from the LP and P sites. Hypothetical effects of TBZ in microbial communities were based on the mode of action of TBZ and on the literature available (marked by superscripts). Non-observed effects are marked “n. e.”.

	Bacteria (observed)	Algae (observed)	Fungi (hypothetical)	Protozoa (hypothetical)
Biofilm	LP	↓ I/D cell ratio ↓ respiration	↓ photo efficiency	↓ fungi ^{a,b} ↓ protozoa ^{c,d}
	P	↓ I/D cell ratio	n. e.	↓ fungi ^{a,b} ↓ protozoa ^{c,d}
TBZ effect	Indirect/direct(?)	Indirect	Direct	Direct
Plankton	LP	↑ bacterial density	n. e.	↓ fungi ^{a,b} ↓ protozoa ^{c,d}
	P	↑ bacterial density	n. e.	↓ fungi ^{a,b} ↓ protozoa ^{c,d}
TBZ effect	Indirect	n. e.	Direct	Direct

a. Muñoz-Leoz et al. (2011); b. Artigas et al. (2012); c. Ekelund (1999); d. Bending et al. (2007).

Conversely to biofilms, plankton communities showed a stimulation of bacterial densities in the presence of TBZ. This response was similar between plankton communities from LP and P sites and could be explained by (i) increased levels of nutrients and energy sources released from dead fungi (mostly unicellular forms in plankton; Wurzbacher et al., 2010) and utilized by bacteria, ii) direct TBZ effects on the protozoan community (Ekelund, 1999) reducing grazing pressure on the bacterial community, and iii) possible use of TBZ as an extra carbon source by the bacteria. The first two hypotheses are linked to the interaction between plankton microorganisms. For instance, grazing by protozoan communities is one of the major drivers shaping bacterial community structure in plankton (Güde, 1989). Studies using culture-dependent methods have shown direct reduction of protozoan populations by a variety of fungicides (Ekelund, 1999). Therefore, it cannot be excluded that direct negative effects of fungicides on bacterial grazers occurred in our microcosms, indirectly resulting in increased bacterial densities in the plankton communities (Table 2). The third hypothesis, concerning the microbial decay of TBZ, can be ruled out, since the carbon supplied by TBZ (0.00125 mgC L⁻¹ and 0.0125 mgC L⁻¹ in the 2 and 20 µg L⁻¹ treatments, respectively) was between two and three orders of magnitude below DOC levels in plankton microcosms (2.5–3.5 mg C L⁻¹) and was thus relatively unimportant.

The different effects of TBZ between biofilm and plankton communities could be explained by difference in the nature of microbial communities, pre-exposure to TBZ, and possibly also the specific composition of microbial communities. The pyrosequencing approach employed here on 16S and 18S rDNA genes revealed that TBZ mostly affected the eukaryotic community and that this effect was different between biofilms and plankton (Pascault et al., in prep.). These pyrosequencing results supported our argument that the TBZ-induced reduction in protozoan populations might have led to increased bacterial densities in the plankton experiments. Bending et al. (2007) also found that the effects of TBZ on fungal and protozoan community structure differed in soils with identical mineralogical composition but different microbial community composition. Apart from community composition, the presence of EPS matrix in biofilms could also have modulated the effects of TBZ compared to plankton (free-living organisms) which is left more exposed to TBZ. This hypothesis is supported by previous work performed with metal contaminants. Indeed, metal toxicity is usually found at higher concentrations in biofilms than free-living organisms, since the structure of biofilms reduces solute penetration into the cells (Guasch et al., 2010). However, the duration and frequency of contamination episodes play a major role, since toxicant diffusion within the biofilm is influenced by exposure time (Guasch et al., 2010). Given the exposure mode, it is reasonable to posit that biofilm (adapted to peak exposure scenarios, at higher TBZ concentrations in this study) and plankton (more adapted to chronic exposure, at lower TBZ concentrations in this study) microbial communities have developed different strategies to cope with contaminants in their ecosystems, but this remains a speculation that needs to be tested.

The attenuation kinetics of TBZ differed significantly between biofilm and plankton experiments, being four times greater in biofilms. Several factors can be put forward to explain the observed differences in TBZ attenuation, including adsorption processes and biotic degradation of the molecule. Previous studies indicate that TBZ is a relatively lipophilic molecule ($\log K_{ow} = 3.7$; Margni et al., 2002) that rapidly adsorbs onto organic matter (Artigas et al., 2012). Biofilms possess a matrix of extracellular polymeric substances (EPS) capable of rapidly adsorbing metal ions (Guasch et al., 2010) and organic matter (Freeman and Lock, 1995), including organic pesticides such as TBZ. Conversely, plankton cells are mostly free-living and so develop less EPS, which would mean that they have a lower pollutant adsorption capacity. Another possible explanation could be microbial decay of TBZ (Obanda and Shupe, 2009), since TBZ degradation metabolites (e.g. hydroxyl-tebuconazole and desbutyl-tebuconazole) were detected in water samples from biofilm experiments but not in samples from plankton experiments. However, the steep slopes observed in TBZ dissipation curves from biofilms suggested that an active sorption process took place between days 0 and 3 of experiment (~ 50% of total TBZ dissipated). This process may mirror that described for other organic contaminants in biofilms where two sorption phases are identified — a rapid initial process to an “equilibrium” concentration and a second phase of slower sorption (Wicke et al., 2007). Consequently, it can be argued that TBZ was mostly adsorbed by the EPS matrix rather than degraded by microorganisms, and that this potential to adsorb TBZ was greater in biofilms than in plankton communities.

The different sensitivity between communities from LP and P sites could be explained by the differences in pollution-induced community tolerances of biofilm and plankton microorganisms to TBZ. Blanck (2002) postulated that toxicants exert a selection of sensitive species which get replaced by more tolerant species in biological communities. This principle concord with the contrasting responses to TBZ observed between biofilm communities from LP and P sites, but not for the plankton. The similar sensitivity to TBZ between P-site and LP-site plankton communities might be explained by the elevated TBZ concentrations used in our experiment, as the nominal concentrations

of 2 and 20 $\mu\text{g L}^{-1}$ can be encountered during contamination peaks in rivers but not in the lakes studied (mean TBZ concentration of 0.001 $\mu\text{g L}^{-1}$ in the polluted lake). Therefore, the responses observed in our plankton experiments should be considered as extreme scenarios of TBZ contamination. Finally, increasing the TBZ dose (from 2 to 20 $\mu\text{g L}^{-1}$) did not significantly enhance TBZ effects on biofilm and plankton community descriptors. Therefore, the toxicity of TBZ for the four types of aquatic microbial communities analyzed was detectable at a relatively low concentration (2 $\mu\text{g L}^{-1}$ in this experiment). A dose-dependent response is usually expected in toxicity experiments, but many exceptions are observed, as in the case of indirect effects in complex biological communities like biofilms (bearing multiple microbial interactions) where the effects of a toxic compound do not necessarily increase linearly with pesticide dose (Ricart et al., 2009).

5. Conclusions

TBZ effects in biofilm and plankton microbial communities were moderate at the 2 and 20 $\mu\text{g L}^{-1}$ TBZ doses tested, which suggests that the actual TBZ concentrations found in the ecosystems pose relatively little ecotoxicological risk to the ecology of aquatic microbial communities analyzed. However, we found different sensitivities to the TBZ molecule between biofilms and plankton, which can be explained by the specific structure and composition of the microbial community. TBZ effects on non-targeted microalgal and bacterial communities were mostly indirect (Table 2) and seemed to be related to the multiple interactions between biofilm components (i.e. fungal and protozoan communities), though these results need to be confirmed by further investigations specifically targeting those microbial groups (Pascault et al., in prep.).

TBZ toxicity assessment in aquatic ecosystems cannot be exclusively based on single-species responses but should also cover the responses of complex microbial communities (including targeted and non-targeted organisms) to the fungicide. Our study suggests that lifestyle (benthic versus pelagic), pre-exposure to TBZ (less-polluted versus polluted sites) and exposure mode (peak exposure versus chronic exposure scenarios) of microbial communities can accentuate the variability in microorganism responses to the TBZ fungicide. To improve TBZ risk assessment in aquatic microbial communities, further research should be performed by enlarging the spatial and temporal scale of microbial responses to TBZ.

Conflict of interest

We state that this manuscript does not entail any conflict of interest.

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