

1 Local nutrient regimes determine site-specific environmental triggers of cyanobacterial and  
2 microcystin variability in urban lakes

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4 Som Cit Sinang<sup>1,2</sup>, Elke S. Reichwaldt<sup>1</sup> and Anas Ghadouani<sup>1\*</sup>

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6 <sup>1</sup>Aquatic Ecology and Ecosystem Studies, School of Civil, Environmental and Mining

7 Engineering, The University of Western Australia, 35 Stirling Highway, M015, Crawley WA

8 6009, Western Australia, Australia

9 <sup>2</sup>Present address: Faculty of Science and Mathematics, Sultan Idris Education University,

10 35900 Tanjong Malim, Perak, Malaysia

11

12 \*Corresponding author:

13 Ph: +61-8-6488-2687

14 Fax: +61-8-6488-1015

15 Email: [anas.ghadouani@uwa.edu.au](mailto:anas.ghadouani@uwa.edu.au)

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26 **Abstract**

27 Toxic cyanobacterial blooms in urban lakes present serious health hazards to humans and  
28 animals and require effective management strategies. In the management of toxic  
29 cyanobacteria blooms, understanding the roles of environmental factors is crucial. To date, a  
30 range of environmental factors have been proposed as potential triggers for the  
31 spatiotemporal variability of cyanobacterial biomass and microcystins in freshwater systems.  
32 However, the environmental triggers of cyanobacteria and microcystin variability remain a  
33 subject of debate due to contrasting findings. This issue has raised the question, if the  
34 relevance of environmental triggers may depend on site-specific combinations of  
35 environmental factors. In this study, we investigated the site-specificity of environmental  
36 triggers for cyanobacterial bloom and microcystin dynamics. Our study suggests that  
37 cyanobacterial dominance and cyanobacterial microcystin content variability were  
38 significantly correlated to phosphorus and iron concentrations. However, correlations  
39 between phosphorus and iron with cyanobacterial biomass and microcystin variability were  
40 not consistent between lakes, thus suggesting a site specificity of these environmental factors.  
41 The discrepancies in the correlations could be explained by differences in local nutrient  
42 concentration and the cyanobacterial community in the systems. The findings of this study  
43 suggest that identification of significant environmental factors under site-specific conditions  
44 might be an important strategy to enhance positive outcomes in cyanobacterial bloom control  
45 measures.

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47 **Keywords:** Cyanobacterial variability; Microcystin variability; Environmental triggers;  
48 Nutrients; Site-specific; Bloom management.

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## 52 **1 Introduction**

53 Urban lakes often serve as recreational spaces for communities and habitats for wildlife (Yan  
54 et al., 2012; Liu, 2014) . To date, urban lakes continue to deteriorate due to increased  
55 anthropogenic activities and often face water quality problems including toxic cyanobacteria  
56 blooms (Maria Pineda-Mendoza et al., 2012; Reichwaldt and Ghadouani, 2012; Lei et al.,  
57 2014; Sun et al., 2014; Zhang et al., 2014). This issue has received great attention from **water**  
58 **authorities world-wide** as it presents health hazards to humans and animals who either  
59 directly or indirectly received services provided by urban lakes (O'Bannon et al., 2014;  
60 Rastogi et al., 2014; Waajen et al., 2014). **The management of toxic cyanobacterial blooms is**  
61 **often challenging due to the variability in cyanobacteria biomass and microcystins (Rolland**  
62 **et al., 2013; Carey et al., 2014).**

63

64 Cyanobacterial biomass and the amount of microcystins being produced during toxic  
65 cyanobacterial blooms can vary significantly on a spatial basis (Reichwaldt et al., 2013;  
66 Sinang et al., 2013; Thi Thuy et al., 2014; Waajen et al., 2014). Past studies have found large  
67 variations in the percentages of potentially toxic cyanobacteria and microcystin concentration  
68 between spatially isolated populations (Sitoki et al., 2012; Li et al., 2014). Furthermore, it  
69 was reported that the variability of cyanobacterial biomass in lakes only explained a small  
70 fraction of the variability in microcystin concentration (Sinang et al., 2013; Eva and Lindsay,  
71 2014). These findings highlight the importance to fully understand the roles of environmental  
72 factors controlling cyanobacteria and microcystin variability.

73

74 It has been suggested that cyanobacterial biomass and microcystin variability largely depends  
75 upon physical, chemical and biological properties of the water bodies (Engström-Öst et al.,

76 2013; Lehman et al., 2013; Paerl and Otten, 2013; Ruiz et al., 2013). A range of  
77 environmental factors, including nitrogen and phosphorus (Schindler, 2012; Srivastava et al.,  
78 2012; Chaffin and Bridgeman, 2014; Van de Waal et al., 2014), TN:TP ratio (Smith, 1983;  
79 Wang et al., 2010b; Van de Waal et al., 2014), temperature (Davis et al., 2009; Rolland et al.,  
80 2013), salinity (Tonk et al., 2007), and iron (Ame and Wunderlin, 2005; Nagai et al., 2007;  
81 Wang et al., 2010a) have been shown to have pronounced effects on either cyanobacterial  
82 dominance, microcystin production, or both. Nevertheless, the results between studies differ,  
83 and there is no clear understanding of the roles of these environmental factors as the triggers  
84 of cyanobacterial dominance and microcystin production. It therefore remains an important  
85 challenge for bloom management to fully understand the mechanisms behind toxic  
86 cyanobacterial bloom development. For instance, regardless of many studies suggesting the  
87 important role of phosphorus, reduction of internal and external phosphorus concentration is  
88 not always successful in preventing the occurrence of toxic cyanobacterial blooms in water  
89 bodies (Lewis and Wurtsbaugh, 2008; Amano et al., 2010; Koreiviene et al., 2014).

90

91 By taking into account the contrasting findings of earlier studies, including inconsistent  
92 outcomes of nutrient reduction strategies, we suggest that the environmental triggers of  
93 cyanobacterial and microcystin variability may vary between water bodies. Therefore, the  
94 main objective of this study was to investigate the site-specificity of environmental triggers  
95 for cyanobacterial biomass and microcystin variability in a local urban lake system. More  
96 specifically, the objectives were to (1) determine the variability of cyanobacterial biomass  
97 and microcystin concentration in a set of local urban lakes, (2) identify the site-specific  
98 relationship between environmental factors and cyanobacterial biomass or microcystin  
99 dynamics.

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## 102 **2 Material and methods**

### 103 **2.1 Study lakes**

104 This study was carried out in Jackadder Lake (31°54'30"S, 115°47'36"E), Bibra Lake  
105 (32°5'25"S, 115°49'16"E) and Yangebup Lake (32°6'56"S, 115°49'33"E) located on the  
106 Swan Coastal Plain, Western Australia (Fig. 1). Sampling was carried out between January  
107 and March 2010. These lakes are shallow with mean depth of 2.1 m, 1.1 m, and 2.5 m for  
108 Jackadder Lake, Bibra Lake and Yangebup Lake, respectively. Jackadder Lake and  
109 Yangebup Lake are permanent lakes while Bibra Lake is subjected to seasonal drying due to  
110 progressive decline in groundwater levels over the Jandakot Mound. Jackadder Lake has an  
111 area of 7.18 ha, is surrounded by 6.6 ha of parkland and is draining a 152 ha catchment area,  
112 (Arnold, 1990; Woodward, 2008). Water levels in Jackadder Lake are maintained by the  
113 input of surface runoff via 10 drain inlets (Rajah 1991, as cited in Kemp, 2009). Jackadder  
114 Lake receives water from the Herdsman Lake catchment area and Osborne Park main drain  
115 during dry summers (Department of Planning, 2010). Bibra Lake has a size of 135 ha with an  
116 open water area of approximately 100 ha (Strategen, 2009) and is located within a 250 ha  
117 catchment are. This lake is surrounded by urban areas and a golf course and serves as habitat  
118 for many species of water birds (Kemp, 2009). Water enters Bibra Lake via direct rainfall  
119 recharge onto the lake surface or from surface runoff from the surrounding catchment  
120 (Strategen, 2009). Yangebup Lake has a total area of 90.5 ha with an open water area of  
121 approximately 68 ha, and is surrounded by residential, agriculture and industrial areas.  
122 Yangebup Lake is a groundwater through-flow wetland that accepts groundwater from the  
123 east and discharges groundwater to the west (Dunlop, 2008). Yangebup Lake receives urban  
124 runoff from three stormwater drains and additionally serves as a compensation basin for the  
125 South Jandakot Drainage system with an approximate area of 200 km<sup>2</sup>. This includes

126 receiving water from neighbouring Thomson Lake when it reaches its maximum water level.  
127 Once Yangebup Lake reaches its maximum allowable water level, water is pumped into  
128 nearby Cockburn Sound (Environmental Protection Authority, 1989). The hydrology of  
129 Jackadder, Bibra and Yangebup lakes is mainly affected by the strong seasonal rainfall  
130 pattern due to the Mediterranean climate. The region's mean annual rainfall is reported as  
131 771.5mm and monthly mean rainfall is 35.1 , 156.3 , 433.3 , and 144.2 mm during summer,  
132 autumn, winter and spring, respectively (Bureau of Metereology, 2014). In response, the  
133 maximum water levels in all lakes occur in September and October, and the minimum water  
134 levels occur in March and April at the end of summer months (Davis et al., 1993). The  
135 region's mean maximum annual temperature is 24.5 °C and monthly maximum temperature  
136 are 30.9°C, 25.4°C, 18.0°C, and 22.6°C during summer, autumn, winter and spring,  
137 respectively (Bureau of Metereology, 2014). Prolonged stable thermal stratification is usually  
138 prevented in these lakes during summer due to continuous or intermittent wind mixing that  
139 creates a homogeneous environment throughout the water column (Davis et al., 1993; Arnold  
140 and Oldham, 1997).

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142 These lakes were selected due to differences reported on physicochemical properties, levels  
143 of cyanobacterial biomass and microcystin concentration. Based on an earlier study, these  
144 lakes represent systems with low, medium and high cyanobacterial biomass and microcystin  
145 concentration. Mean cyanobacterial biomass was reported as 28, 108, and 80  $\mu\text{g chl-}a\text{ L}^{-1}$  in  
146 Jackadder, Bibra and Yangebup Lake, respectively. Mean cellular microcystin concentrations  
147 ( $\text{mg g}^{-1}$  cyanobacterial dry mass) was 4.8  $\text{mg g}^{-1}$  in Jackadder Lake, 35  $\text{mg g}^{-1}$  in Bibra Lake  
148 and 1.7  $\text{mg g}^{-1}$  in Yangebup Lake (Sinang et al., 2013).

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## 151 **2.2 Sampling and analyses**

152 The lakes were sampled **twice a month** between January and March 2010. Three samples  
153 were collected from the same three points at each lake on every sampling occasion. Bibra  
154 Lake dried up in late February; therefore no samples were taken from this lake in March. On-  
155 site measurements and samples were taken from shore sites at a water depth of 0.6 to 1 m.  
156 Temperature (Temp), pH and Salinity (Sal) were measured on-site with a WP-81 probe (TPS  
157 Pty Ltd) at a depth of 0.6 m. Grab water samples for cyanobacteria, microcystin and total  
158 phosphorus quantification were taken from approximately 0.15 m below the surface to avoid  
159 surface scum. Although there was a slight difference in the depth from which the samples for  
160 the physicochemical and water samples were taken, this is not expected to influence the  
161 interpretation of the results, as an earlier study in these lakes (Arnold and Oldham 1997)  
162 indicated that the water bodies at these shallow shore sites are well mixed with respect to  
163 physicochemical variable as shown in **Fig. 2**. Water samples were stored immediately in glass  
164 bottles in the dark on ice. Parameters analysed from these samples were total phosphorus  
165 (TP), total dissolved phosphorus (TDP), total iron (TFe), total dissolved iron (TDFe), total  
166 nitrogen (TN), total dissolved nitrogen (TDN), ammonium (NH<sub>4</sub><sup>+</sup>), cyanobacterial biomass,  
167 total phytoplankton biomass, intracellular and extracellular microcystin fractions. Samples  
168 for dissolved nutrient analyses were pre-filtered with a 0.45µm syringe filter (Acrodisc, HT  
169 Tuffryn) before freezing at -20°C.

170

### 171 **2.2.1 Nutrients and phytoplankton biomass**

172 TP and TDP concentrations were analyzed using the ascorbic acid method, while TFe and  
173 TDFe concentrations were analyzed with the Phenanthroline method, according to standard  
174 methods (APHA, 1998). TN, TDN, and NH<sub>4</sub><sup>+</sup> were analyzed at the South Coast Nutrients  
175 Analysis Laboratory, Albany, Western Australia with the standard colorimetric methods on a

176 segmented flow auto-analyser (Alpkem, Wilsonville, OR, USA). Cyanobacterial and total  
177 phytoplankton chlorophyll-*a* were measured with a top-bench version of a Fluoroprobe (bbe  
178 Moldaenke, Germany). The FluoroProbe measures chl-*a* fluorescence and differentiates four  
179 groups of phytoplankton (chlorophytes, cryptophytes, diatoms, and cyanobacteria) by their  
180 specific fluorescence emission spectrum (Beutler et al., 2002). The fluorescence is used to  
181 calculate total biomass of each phytoplankton group that is expressed as chl-*a* concentration  
182 equivalents ( $\mu\text{g chl-}a \text{ L}^{-1}$ ) (Beutler et al., 2002; Ghadouani and Smith, 2005). FluoroProbe  
183 chl-*a* measurements were validated against chl-*a* data of samples extracted according to  
184 standard methods (APHA, 1998) (linear regression analysis:  $R^2 = 0.94$ ,  $N = 32$ ,  $P < 0.05$ ). In  
185 our study, chl-*a* fluorescence as measured by FluoroProbe was used as a proxy for  
186 cyanobacterial biomass (Geis et al., 2000; Eisentraeger et al., 2003).

187

188 For quantification of cyanobacterial biomass and to separate the intracellular from the  
189 dissolved microcystin fraction, water samples were filtered through pre-combusted and pre-  
190 weighed 47 mm GF/C filter papers. Filter papers containing particulate organic matter were  
191 dried for 24 hours at  $60^\circ\text{C}$  and re-weighed to obtain total dry weight (Harada et al., 1999).  
192 These filter papers were then moistened with Milli-Q water and kept frozen (at  $-20^\circ\text{C}$ ) until  
193 intracellular microcystin extraction. As we were interested in the microcystin concentration  
194 per unit cyanobacterial dry mass, cyanobacterial dry mass was calculated from the total dry  
195 mass (from the filters) by adjusting it to the percentage of cyanobacteria measured with the  
196 FluoroProbe. Cyanobacterial dry mass was only used for microcystin quantification.

197

198 Water samples collected for cyanobacterial identification and enumeration were preserved  
199 with acidic Lugol's iodine solution (5 g  $\text{I}_2$ +10 g KI, 20 ml distilled water and 50 ml of 10%  
200 acetic acid) and cyanobacteria were identified to the genus level using phytoplankton



201 taxonomic guideline (Komarek and Hauer, 2011). The relative abundance of each  
202 cyanobacterial genera (cells or colonies ml<sup>-1</sup>) was determined from 10-50 ml of sample using  
203 an inverse microscope (Utermöhl, 1958) and converted into biovolume per ml (µm<sup>3</sup> ml<sup>-1</sup>) by  
204 multiplying the mean cell or colony biovolume (µm<sup>3</sup>) with the total cells or colonies per ml  
205 (cells or colonies ml<sup>-1</sup>). Mean cell or colony biovolume for each cyanobacterial genus was  
206 calculated by finding the geometric figure that best approximated the shape of each genera,  
207 and by measuring the dimension of 20 individual cells or colonies (Hillebrand et al., 1999). A  
208 minimum of 200 cells or colonies of the most abundant cyanobacteria were counted for each  
209 sample. Different cyanobacterial species within each genus can vary in size by several orders  
210 of magnitude. However, as we measured the mean biovolume of each cyanobacterial genus,  
211 differences in sizes between species are evened out as a larger mean is expected, if larger  
212 species are more abundant and vice versa. The calculated mean biovolume of each  
213 cyanobacterial genus was used to compute the dominant cyanobacteria genera in the studied  
214 lakes.

215

### 216 **2.2.2 Microcystin extraction and quantification**

217 Filters were freeze-thawed twice to break the cells prior to methanol extraction (Lawton et  
218 al., 1994). Filters were placed into centrifuge tubes and 5 ml of 75% methanol-water (v/v)  
219 was added. Filters were sonicated on ice for 25 min, followed by gentle shaking for another  
220 25 min. The extracts were then centrifuged at 3273g (Beckman and Coulter, Allegra X-12  
221 Series) for 10 min at room temperature. Extracts were carefully transferred into conical  
222 flasks, and two more extractions were done per filter. All three extracts were pooled and  
223 diluted with Milli-Q to 20% methanol (v/v).

224

225 Intracellular microcystin extracts and the pre-filtered water containing dissolved  
226 (extracellular) microcystin were subjected to solid-phase extraction (SPE) (Waters Oasis  
227 HLB) for clean-up and concentration with a loading speed of  $< 10 \text{ ml min}^{-1}$ . SPE cartridges  
228 were then rinsed with 10 ml of 10, 20 and 30% methanol-water (v/v), before microcystin was  
229 eluted with 100% methanol + 0.1% trifluoroacetic acid (TFA) and evaporated with nitrogen  
230 gas at  $40^\circ\text{C}$ . Finally, samples were re-dissolved in 30% acetonitrile and analysed with high-  
231 performance liquid chromatography (HPLC) by using the Alliance 2695 (Waters, Australia)  
232 with a PDA detector (1.2 nm resolution) and an Atlantis T3  $3\mu\text{m}$  column (4.6 x 150mm i.d).  
233 Mobile phases used were acetonitrile + 0.05% v/v TFA and Milli-Q water + 0.05% TFA.  
234 Microcystin peaks were separated using a linear gradient as described in Lawton et al., (1994)  
235 but with a maximum acetonitrile concentration of 100% and a run time of 37 min. Column  
236 temperature was maintained at  $37.5 \pm 2.5^\circ\text{C}$ . The limit of detection per microcystin peak was  
237 1.12 ng. Microcystin variants were identified based upon their typical absorption spectrum  
238 detected by PDA detector at 238 nm (Meriluoto and Codd, 2005). Commercially available  
239 microcystin-LR standard (Sapphire Bioscience, Australia; purity  $\geq 95\%$ ) was used to  
240 quantify microcystin concentrations. Throughout this manuscript we refer to the total  
241 concentration of microcystin variants per sample as microcystin concentration.

242

### 243 **2.3 Data processing and statistical analyses**

244 Surface water temperatures were between  $19.9$  and  $28.7^\circ\text{C}$  during the study period. However,  
245 the onsite measurements of surface water temperatures were dependent on the time of  
246 sampling and varied by up to  $3.9^\circ\text{C}$  over the course of a day. Therefore, maximum air  
247 temperature on each sampling day recorded by weather stations located nearest to the studied  
248 lakes was used as a substitute for surface water temperature in all analysis. Autocorrelation  
249 Function (ACF) and Partial Autocorrelation Function (PACF) were calculated (SPSS 17.0) in

250 order to verify if autocorrelation exists between the data points. The analyses revealed that  
251 autocorrelation coefficients for all parameters were within the upper and lower confidence  
252 limits, thus suggest independency between data from each sampling date. In this study,  
253 cellular (intracellular) microcystin concentration was expressed as  $\mu\text{g}$  (microcystin-LR mass  
254 equivalents) per g cyanobacterial dry mass to illustrate cyanobacterial microcystin content.  
255 Extracellular microcystin was expressed as the fraction of extracellular microcystin  
256 concentration per total microcystin concentration to allow the quantification of the proportion  
257 of microcystin released into the water column in comparison to the total microcystin being  
258 produced. For all variables, no significant differences (ANOVA,  $P > 0.05$ ) were detected  
259 between three samples collected from three different points in each lake per sampling date.  
260 Therefore, average values of all variables per sampling date were calculated from the three  
261 samples. Between lakes variability of physicochemical factors, cyanobacterial biomass and  
262 microcystin were analysed with one-way ANOVA (SPSS 17.0) with post hoc test (Least  
263 Significance Difference; LSD) as all assumptions for an ANOVA were met (homogeneity of  
264 variances, normality). Bivariate correlation analysis was carried out to identify the  
265 environmental variables which significantly correlate with cyanobacterial fraction, cellular  
266 microcystin concentration and extracellular microcystin fraction (SPSS 17.0). Site-specificity  
267 analysis was performed with a General Linear Model (SPSS 17.0) to identify if the  
268 correlation between environmental variables and cyanobacterial biomass and microcystin  
269 concentration was different between lakes. The site-specificity was determined by the  
270 significant interaction between lake and environmental variable in predicting the variability  
271 of cyanobacteria biomass or microcystin concentration. Redundancy analysis (RDA) was  
272 used to identify the best combination of explanatory variables to explain the variability of  
273 cyanobacterial biomass and microcystin concentration (R version 2.15.1) for each lake.  
274 Canonical ordination (999 permutations) was computed with standardised explanatory and

275 response variables. All data was log transformed to meet the assumption of normality. RDA  
276 analysis on Bibra Lake was conducted without the inclusion of pH and temperature due to an  
277 inadequate number of data points (residual d.f < 0). In all analyses, results were considered  
278 significant at  $P < 0.05$ .

279

## 280 **3 Results**

### 281 **3.1 Physical and chemical characteristics of studied lakes**

282 The three lakes were significantly different in most physicochemical factors except for pH,  
283 Temp and TDFe (Table 1). Salinity, phosphorus, nitrogen and iron, either as total or  
284 dissolved forms (except TDFe), were significantly different between all lakes (one-way  
285 ANOVA). The posthoc tests (LSD) indicates that Jackadder and Yangebup Lake did not  
286 differ in TP, TDP, and  $\text{NH}_4^+$ , however, both lakes were different to Bibra Lake. Furthermore,  
287 all lakes were different in salinity, TN, TDN, and TFe. Jackadder and Yangebup Lakes can  
288 be classified as eutrophic, while Bibra Lake can be classified as hypereutrophic, based on the  
289 mean TP concentrations (Carlson, 1977). Nitrogen limited condition in a lake is usually  
290 defined when TN:TP weight ratio is less than 10 (Graham et al., 2004). As our result showed  
291 that TN:TP ratios below 10 were rare, the studied lakes were not associated with persistent  
292 nitrogen limitation.

293

### 294 **3.2 Variability of cyanobacterial biomass and microcystin concentration**

295 Cyanobacterial communities in all lakes contained potentially toxin-producing cyanobacteria  
296 including *Microcystis* spp., *Planktothrix* spp., *Anabaenopsis* spp., *Anabaena* spp and  
297 *Nodularia* spp. (Fig. 3) with *Microcystis* spp. being the most abundant cyanobacterial genera  
298 in all lakes. Mean total cyanobacterial biomass was  $5.41 \mu\text{g L}^{-1}$ ,  $29.60 \mu\text{g L}^{-1}$ ,  $15.14 \mu\text{g L}^{-1}$  in  
299 Jackadder, Bibra and Yangebup Lake, respectively (Fig. 4A). Cyanobacterial biomass varied

300 within an order of magnitude on a temporal basis in Bibra and Jackadder Lake (Jackadder: 1 -  
301 12  $\mu\text{g L}^{-1}$ , Bibra: 5 - 83  $\mu\text{g L}^{-1}$ , Yangebup: 8 - 32  $\mu\text{g L}^{-1}$ ) (Fig. 4A). Although cyanobacterial  
302 biomass was significantly higher in Bibra Lake compared to the other two lakes ( $F_{(2,45)}= 7.62$ ,  
303  $P < 0.05$ ), the cyanobacterial fraction (the ratio of cyanobacterial chlorophyll-*a* to total  
304 phytoplankton chlorophyll-*a*) in this lake was significantly lower than in Jackadder and  
305 Yangebup Lake ( $F_{(2,45)}= 3.59$ ,  $P < 0.05$ ) (Fig. 4B). Cyanobacterial fraction ranged between  
306 0.05 to 0.71 in Jackadder Lake, 0.16 to 0.68 in Yangebup Lake, and 0.11 to 0.51 in Bibra  
307 Lake. The post hoc tests indicated that Jackadder and Yangebup Lakes did not differ in  
308 cyanobacterial biomass and cyanobacterial fraction, but both lakes were different to Bibra  
309 Lake.

310

311 Cellular microcystin concentration ( $\text{mg g}^{-1}$  cyanobacterial dry mass) varied over three orders  
312 of magnitude in Jackadder Lake, and two orders of magnitude in both Bibra Lake and  
313 Yangebup Lake (Fig. 4C) throughout the sampling events. Mean cellular microcystin  
314 concentrations were 0.407  $\text{mg g}^{-1}$  in Jackadder Lake, 0.233  $\text{mg g}^{-1}$  in Bibra Lake, and 0.150  
315  $\text{mg g}^{-1}$  in Yangebup Lake. Cellular microcystin concentration was not significantly different  
316 between lakes ( $F_{(2,45)}= 2.07$ ,  $P > 0.05$ ). Mean extracellular microcystin fraction was 0.18 in  
317 Jackadder Lake, 0.04 in Bibra Lake, and 0.26 in Yangebup Lake (Fig. 4D). The post hoc tests  
318 indicated that Bibra Lake was the only lake that had a significantly different extracellular  
319 microcystin fraction when compared to other lakes ( $F_{(2,45)}= 6.49$ ,  $P < 0.05$ ).

320

### 321 **3.3 Relationship between environmental factors and cyanobacterial fraction or** 322 **microcystin concentration**

323 Most environmental factors were weakly, but significantly correlated with cyanobacterial  
324 fraction and microcystin concentrations when data from all lakes were combined (Table 2).

325 The correlations presented in Table 2 suggested that, in general, cyanobacterial dominance in  
326 the phytoplankton community was favored at relatively lower nutrient concentrations as it  
327 was negatively correlated to TP, TDP, TFe, and TDFe. In contrast, cyanobacterial dominance  
328 was positively correlated with TN:TP ratio. Cellular microcystin concentration was positively  
329 correlated with phosphorus and iron, but not nitrogen. TDFe showed the strongest positive  
330 correlation with cellular microcystin concentration, followed by TP, TFe, and TDP. Cellular  
331 microcystin was also negatively correlated with TN:TP ratio. In contrast to cellular  
332 microcystin, extracellular microcystin fraction was negatively correlated with salinity, TP,  
333 TDP, TN, TDN, and positively correlated with TN:TP ratio.

334

### 335 **3.4 Site-specific relationship between environmental factors and cyanobacterial fraction** 336 **or microcystin concentration**

337 **Most of the significant** correlations between environmental factors and cyanobacterial  
338 fraction, cellular microcystin concentration or extracellular microcystin fraction were  
339 different between lakes (Table 2). The correlations between cyanobacterial fraction and TP,  
340 TDP, TFe and TN:TP ratios were significantly different between lakes, while the correlation  
341 with TDFe was consistent between lakes. In terms of cellular microcystin concentration, the  
342 correlations with TP, TDP, and TFe were significantly different between lakes, while the  
343 correlations with TDFe and TN:TP ratio were consistent between lakes. The correlations  
344 between extracellular microcystin fraction and salinity, TDP, and TDN were significantly  
345 different between lakes, while the correlation with TP, TN and TN:TP ratio was consistent  
346 between lakes.

347

348 **The differences in the correlations between environmental factors and cyanobacterial**  
349 **fraction, cellular microcystin concentration or extracellular microcystin fraction between**

350 lakes are shown in Fig. 5, Fig. 6, Fig. 7 and Table 3. In Bibra Lake, no significant correlation  
351 between cyanobacterial fraction and TP (Fig. 5A) or TN:TP (Fig. 5E) were found.  
352 Additionally, the correlations between cyanobacterial fraction and TDP (Fig. 5B) and TFE  
353 (Fig. 5C) were positive in Yangebup and Bibra lakes. Meanwhile, the correlation between  
354 cyanobacterial fraction and TDFe was negative in all lakes (Fig. 5D). In terms of cellular  
355 microcystin concentration, its correlation with TP was weak in Bibra Lake (Fig. 6A). The  
356 correlation between cellular microcystin concentration and TDP (Fig. 6B) was not significant  
357 in Yangebup Lake. The correlation between cellular microcystin concentration and TFe (Fig.  
358 6C) was negative only in Bibra Lake. Meanwhile, the correlations between cellular  
359 microcystin concentration and TDFe (Fig. 6D) and TN:TP (Fig. 6E) were positive and  
360 negative in all lakes, respectively. In terms of extracellular microcystin fraction, the  
361 correlation with salinity (Fig. 7A) was not significant in Bibra Lake. In addition, its correlation  
362 with TDP (Fig. 7C) was significant only in Yangebup Lake. The correlation between  
363 extracellular microcystin fraction and TDN (Fig. 7E) was positive in Jackadder Lake. The  
364 correlations between extracellular microcystin fraction and TP (Fig. 7B) and TN:TP (Fig. 7F)  
365 were negative and positive in all lakes, respectively.

366

### 367 **3.5 Multivariate analysis of environmental factors and the variability of cyanobacterial** 368 **fraction and microcystin concentration**

369 RDA analyses performed with forward selection by permutation ( $n_{perm} = 999$ ) showed  
370 significant relationships ( $P < 0.05$ ) between the measured environmental factors and the  
371 combined variability of cyanobacterial fraction, cellular microcystin concentration and  
372 extracellular microcystin fraction for each lake. The canonical ordination showed that 75%  
373 (Jackadder Lake;  $R^2_{adj.} = 0.75$ ;  $F=5.726$ ), 80% (Bibra Lake;  $R^2_{adj.} = 0.80$ ;  $F=5.888$ ) and 75%  
374 (Yangebup Lake;  $R^2_{adj.} = 0.75$ ;  $F=5.804$ ) of the combined variability of cyanobacterial

375 fraction, cellular microcystin concentration and extracellular microcystin fraction can be  
376 explained by the measured environmental factors (Fig. 8A - C). The environmental factors  
377 that were closely correlated to the cyanobacterial fraction, cellular microcystin concentration  
378 and extracellular microcystin fraction were not always the same between lakes. TDP was  
379 only correlated to either cyanobacterial fraction or cellular microcystin concentration in Bibra  
380 and Jackadder Lakes (Fig. 8A, 8B) but not in Yangebup Lake (Fig. 8C). Additionally, TFe  
381 was positively correlated to cyanobacterial fraction only in Bibra Lake (Fig. 8B) but not in  
382 the other two lakes (Fig. 8A, 7C). In comparison to the other factors, TDFe was always  
383 consistently correlated to cyanobacterial fraction and cellular microcystin concentration  
384 variability.

385

#### 386 **4 Discussion**

387 The correlation between the environmental factors and cyanobacterial and microcystin  
388 variability were different between lakes. This is a strong indication that the relevance of  
389 environmental triggers of cyanobacterial fraction, cellular microcystin concentration, and  
390 extracellular microcystin fraction may depend on site-specific combinations of environmental  
391 factors. Our results suggest that the site-specificity of environmental triggers may be related  
392 to spatial heterogeneity of the respective environmental factor, as each factor could present at  
393 different concentration regimes in each lake. Graham et al., (2004) and Dolman et al., (2012)  
394 have suggested that the correlations between the environmental factors and cyanobacterial  
395 biomass and microcystin concentration could change when the concentrations of the  
396 respective environmental factors increased from low to high in systems. Our results support  
397 these previous findings as the between lakes consistencies in the correlations between  
398 cyanobacterial fraction and cellular microcystin variability with TP, TFe and TDFe were  
399 closely related to the levels of TP, TFe and TDFe concentrations in each lake. Mean TP and



400 TFe concentrations in Bibra Lake were one order of magnitude higher than in Jackadder and  
401 Yangebup Lakes, while mean TDFe concentrations in all lakes ranged within the same order  
402 of magnitude (Table 1). This could explain why the correlations between cyanobacterial  
403 fraction and cellular microcystin variability with TP and TFe were significantly different  
404 across lakes, while TDFe was not (Table 2). Further, the correlation between cyanobacterial  
405 fraction and TP was only significant in Yangebup and Jackadder Lake, which both had lower  
406 TP concentrations than Bibra Lake, in which no significant correlation was found.  
407 Meanwhile, the correlation between cellular microcystin concentration and TFe was negative  
408 only in Bibra Lake, where TFe was present at significantly higher concentrations compared to  
409 the other two lakes. This indicates that the effect of environmental factors on cyanobacterial  
410 and microcystin variability may depend on site-specific factors such as concentration  
411 regimes, even in non-nutrient limited lakes. Therefore, a generalization by only using  
412 concentrations of nutrients might not be sufficient for future management of lakes.

413

414 The site-specificity of the environmental triggers of cyanobacterial and microcystin  
415 variability may also be a consequence of the variation of cyanobacterial communities  
416 between the systems. TFe was negatively correlated to cyanobacterial fraction in Jackadder  
417 and Yangebup Lake, while in Bibra Lake, a positive correlation (even not may not  
418 significant) was observed between the two (Fig. 8A, B). The cyanobacterial community in  
419 Jackadder Lake was composed of only one nitrogen-fixing cyanobacterial genera (Fig. 3). In  
420 contrast, multiple nitrogen-fixing cyanobacterial genera were present in Bibra Lake.  
421 Nitrogen-fixing cyanobacteria are known to utilize more iron in comparison to non nitrogen-  
422 fixers (Wilhelm, 1995). Therefore, the site-specific correlation between TFe and  
423 cyanobacterial fraction may be explained through a greater iron requirement of the

424 cyanobacterial community in Bibra Lake, in comparison to the cyanobacterial community in  
425 Jackadder Lake.

426

427 Due to the potentially site-specific environmental triggers of cyanobacterial and microcystin  
428 variability, the results presented in this study are important for the management of these lakes  
429 or lakes with similar physical, chemical and biological characteristics. In this study, the  
430 variability of cyanobacterial fraction was negatively correlated with TP, TDP, TFe, TDFe;  
431 positively correlated with TN:TP ratio. These correlations illustrate that in our study,  
432 cyanobacteria dominated under lower phosphorus availability (Amano et al., 2010). Although  
433 cyanobacteria as a group can dominate under a wide range of conditions, high phosphorus  
434 concentrations have been shown to potentially limit the ability of cyanobacteria to become  
435 dominant in the phytoplankton community (Chorus and Bartram, 1999; Reynolds et al.,  
436 2006). One reason for that is the higher growth rate of other phytoplankton groups compared  
437 to cyanobacteria, and, as such, their ability to utilize nutrients faster under high nutrient  
438 conditions. This can explain the negative correlation between cyanobacterial fraction and  
439 phosphorus concentration found in our study, and, maybe as a consequence of this, a positive  
440 correlation with TN:TP. In terms of iron, low availability was correlated to high  
441 cyanobacterial fraction in these lakes. This result indicated that cyanobacteria pose a  
442 competitive advantage to dominate the phytoplankton community under low iron availability.  
443 Cyanobacteria are capable to alter their cellular iron requirements, and increase the ability to  
444 utilize iron at a low concentration, through the present of siderophores (Boyer et al., 1987;  
445 Lee et al., 2011). As reported in the Nagai et al., (2007), cyanobacteria including *Microcystis*  
446 spp. and *Planktothrix* spp. can produce siderophores and become a superior competitor under  
447 iron limited conditions. These results indicate that phosphorus and iron reduction in water

448 bodies might not be a sufficient remedial strategy against the occurrence of toxic  
449 cyanobacterial bloom.

450

451 In contrast to cyanobacterial fraction, the variability of **cellular** microcystin concentration was  
452 positively correlated to TP, TDP, TFe, TDFe and negatively correlated with TN:TP and  
453  $\text{NH}_4^+$ . **High availability of phosphorus relative to other nutrients** is required for energy and  
454 material supply in microcystin biosynthesis as microcystin production in cyanobacterial cells  
455 is an energy intensive process (Vezie et al., 2002). This is further supported through the  
456 observed negative correlation between cellular microcystin and TN:TP ratio, as low  
457 microcystin production is expected under conditions where **phosphorus is present at lower**  
458 **concentrations** in relation to other nutrients. In addition, the positive correlation between iron  
459 and cellular microcystin concentration is in agreement with earlier studies which suggested  
460 that iron plays an essential role in many metabolic pathways including microcystin  
461 biosynthesis in cyanobacteria (Jiang et al., 2008; Wang et al., 2010a).

462

463 These results illustrate that reducing phosphorus and iron concentrations in water bodies  
464 **could** potentially reduce the overall toxicity of cyanobacterial bloom, even though it might  
465 not completely prevent the occurrence of cyanobacterial bloom. In terms of  $\text{NH}_4^+$ , our results  
466 suggest that reducing  $\text{NH}_4^+$  concentrations may be associated with higher microcystin  
467 concentration. This is possible as toxic cyanobacterial genotypes are known to be favored  
468 under low inorganic nitrogen conditions (Ame et al., 2003).

469

470 Environmental conditions influencing the release of microcystin into the environment,  
471 besides cells lyses, are-not well understood (Rohrlack and Hyenstrand, 2007; Barrington et  
472 al., 2013). Our results showed that correlations exist between extracellular microcystin

473 fraction and nutrients, however, the correlations could be direct or indirect ones. If they are  
474 direct, our results suggest that regardless of the potentially low microcystin production,  
475 cyanobacteria may release microcystins at lower nitrogen and phosphorus concentrations. Th  
476 would support by the hypothesis that microcystin is involved in nutrient competition in the  
477 phytoplankton community (Huisman and Hulot, 2005).

478

## 479 **5 Conclusions**

480 The current approach to water body restoration and the prevention of toxic cyanobacterial  
481 blooms relies on reducing nutrient loading into water bodies and limiting the availability of  
482 nutrients in the water column. This approach might not always be successful in preventing  
483 the occurrence of cyanobacterial blooms, due to the roles of physicochemical factors on  
484 cyanobacteria and microcystin variability being dependent on the site-specific combination of  
485 environmental factors. Thus, it is important to take into account the effect of spatial  
486 heterogeneity in the management of toxic cyanobacterial blooms. Site-specific studies may be  
487 required to determine the factors causing cyanobacterial dominance and microcystin  
488 production in different systems with different characteristics such as the hydrology, land use  
489 and water chemistry.

490

491 In our study, the dominance of cyanobacteria in the phytoplankton community is correlated to  
492 lower phosphorus and iron concentrations in the systems. In contrast, cyanobacteria required  
493 higher phosphorus and iron concentrations in the water column to produce a high amount of  
494 microcystin. Therefore, reducing phosphorus and iron concentration in the water column  
495 might not be a sufficient remedial strategy against the occurrence of toxic cyanobacterial  
496 bloom, if these nutrients are still available in sufficient amount to support the growth of

497 highly competitive cyanobacteria. However, reducing phosphorus and iron could reduce the  
498 amount of microcystin being produced within cyanobacterial cells.

499

## 500 **6 Acknowledgements**

501 This project was funded by the Australian Research Council's Linkage Project funding  
502 scheme (LP0776571) and the Water Corporation of Western Australia. We wish to thank  
503 Professor Pierre Legendre, Laura Firth and Kevin Murray for their valuable statistical advice,  
504 and Liah Coggins for her help in the editing of the manuscript. During the study, Sinang, S.C.  
505 was supported by a scholarship from Universiti Pendidikan Sultan Idris (UPSI) and Malaysia  
506 Government.

507

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711

712

Table 1. Physical and chemical properties of the three lakes throughout the sample period (Jan – March 2010).

Factors	Jackadder Lake (N =18)		Bibra Lake (N =12)		Yangebup Lake (N =18)		ANOVA
	Mean ± SD	Range	Mean ± SD	Range	Mean ± SD	Range	
pH	8.7 ± 0.3	8.1 – 9.0	8.9 ± 0.2	8.5 – 9.2	8.9 ± 0.4	7.5 – 9.3	F <sub>(2,45)</sub> = 2.16
Temp	33.0 ± 4.9	27.4 - 42.7	35.7 ± 4.7	30.8 - 43.0	34.7 ± 4.1	30.8 - 43.0	F <sub>(2,45)</sub> = 1.31
Sal (ppk)	0.4 ± 0.04	0.3 – 0.4	2.9 ± 1.0	1.7 – 4.1	0.9 ± 0.1	0.8- 1.1	F <sub>(2,45)</sub> = 99.08 *
TP (µg L <sup>-1</sup> )	44.0 ± 28.0	20.0 – 131.6	598.1 ± 362.0	214.7 – 1145.9	64.8 ± 44.2	24.0 – 168.0	F <sub>(2,45)</sub> = 40.28 *
TDP (µg L <sup>-1</sup> )	17.6 ± 4.8	12.0 – 26.7	67.9 ± 51.3	16.0 – 180.0	23.2 ± 7.6	13.3 – 40.7	F <sub>(2,45)</sub> = 15.27 *
TFe (µg L <sup>-1</sup> )	123.3 ± 66.2	63.6 – 261.8	192.1 ± 43.4	138.2 – 289.3	81.5 ± 24.1	48.4 – 122.9	F <sub>(2,45)</sub> = 18.91 *
TDFe (µg L <sup>-1</sup> )	69.2 ± 66.3	20.0 – 200.0	89.1 ± 30.4	38.6 – 154.1	52.9 ± 28.9	11.2 – 92.6	F <sub>(2,45)</sub> = 2.15
NH <sub>4</sub> (µg L <sup>-1</sup> )	100.8 ± 54.9	30.0 – 180.0	191.5 ± 33.8	150.0 – 250.3	86.3 ± 45.6	30.0 – 160.0	F <sub>(2,45)</sub> = 20.04 *
TN (mg L <sup>-1</sup> )	1.3 ± 0.4	0.7 – 2.2	11.7 ± 5.2	4.9 – 17.3	3.5 ± 0.8	1.9 – 5.2	F <sub>(2,45)</sub> = 59.38 *
TDN(mg L <sup>-1</sup> )	0.8 ± 0.2	0.4 – 1.1	8.7 ± 3.0	4.9 – 14.0	2.4 ± 0.3	1.9 – 2.8	F <sub>(2,45)</sub> = 104.98 *
TN:TP	35.6 ± 14.9	11.1 – 76.1	23.1 ± 10.0	10.3 – 41.1	68.6 ± 29.9	25.0 – 124.1	F <sub>(2,45)</sub> = 19.51 *

N = number of samples

SD = standard deviation

\* =  $P < 0.05$

1 Table 2. Correlation coefficients (R) between the environmental factors and cyanobacterial  
 2 fraction, cellular microcystin concentration and extracellular microcystin fraction analyzed  
 3 from combined data from all lakes using bivariate correlation analysis.

4

Factor N = 48	Cyanobacterial Fraction (%)	Cellular microcystin concentration ( $\mu\text{g g}^{-1}$ )	Extracellular microcystin fraction (%)
pH	-0.108	0.227	-0.297
Temp	0.018	-0.246	0.078
Sal (ppk)	-0.250	0.067	<b>-0.374*</b>
TP ( $\mu\text{g L}^{-1}$ )	<b>-0.337*</b>	<b>0.399*</b>	<b>-0.392</b>
TDP ( $\mu\text{g L}^{-1}$ )	<b>-0.357*</b>	<b>0.296*</b>	<b>-0.427*</b>
TFe ( $\mu\text{g L}^{-1}$ )	<b>-0.570*</b>	<b>0.343*</b>	-0.037
TDFe ( $\mu\text{g L}^{-1}$ )	<b>-0.777</b>	<b>0.590</b>	-0.064
NH <sub>4</sub> ( $\mu\text{g L}^{-1}$ )	0.105	-0.267	-0.114
TN ( $\text{mg L}^{-1}$ )	-0.236	0.085	<b>-0.375</b>
TDN ( $\text{mg L}^{-1}$ )	-0.265	0.095	<b>-0.400*</b>
TN:TP	<b>0.423*</b>	<b>-0.446</b>	<b>0.386</b>

5 Significant ( $P < 0.05$ ) factors are highlighted in bold;

6 \* Indicates site-specific correlation of the respective environmental variables. Site-specific  
 7 correlation was determined through a significant interaction between lake and  
 8 environmental variable when analyzed in General Linear Model ( $P < 0.05$ );

9 N = number of samples.

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13 Table 3: Correlation coefficients (R) between the environmental factors and cyanobacterial fraction, cellular microcystin concentration and  
 14 extracellular microcystin fraction analyzed for each lake using bivariate correlation analysis.

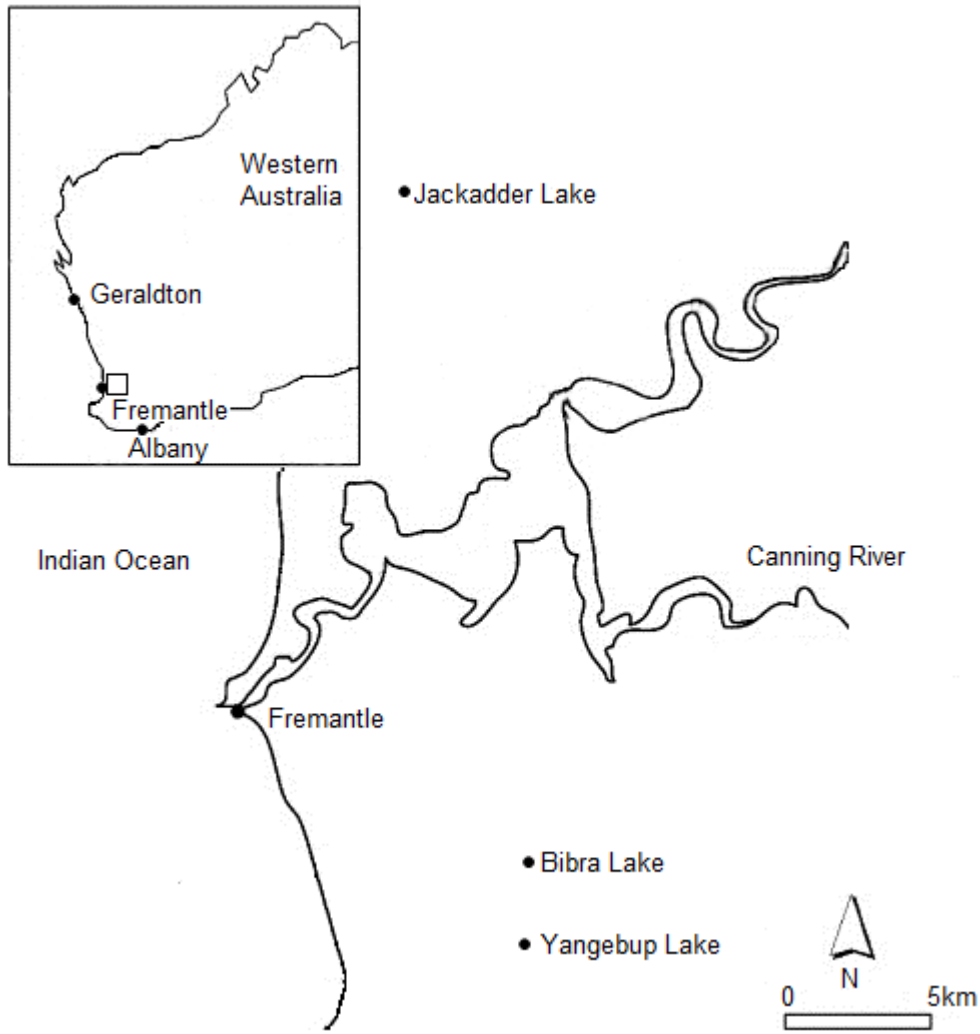
Factor N=48	Cyanobacterial fraction (%)			Cellular microcystin concentration ( $\mu\text{g g}^{-1}$ )			Extracellular microcystin fraction (%)		
	Jackadder	Bibra	Yangebup	Jackadder	Bibra	Yangebup	Jackadder	Bibra	Yangebup
pH	-0.363	<b>-0.653</b>	0.225	0.426	<b>0.762</b>	0.190	0.155	-0.714	-0.360
Temp	0.119	-0.112	0.016	-0.288	-0.185	-0.160	0.138	-0.686	0.130
Sal (ppk)	-0.423	-0.204	-0.460	0.330	0.448	<b>0.587</b>	<b>0.570</b>	-0.775	<b>-0.659</b>
TP ( $\mu\text{g L}^{-1}$ )	<b>-0.873</b>	-0.272	<b>-0.742</b>	<b>0.826</b>	0.489	<b>0.696</b>	-0.303	-0.441	-0.295
TDP( $\mu\text{g L}^{-1}$ )	-0.397	<b>-0.641</b>	0.147	<b>0.553</b>	<b>0.764</b>	0.225	-0.088	-0.498	<b>-0.587</b>
TFe ( $\mu\text{g L}^{-1}$ )	<b>-0.789</b>	0.389	-0.304	<b>0.715</b>	<b>-0.605</b>	0.230	0.380	0.499	-0.245
TDFe ( $\mu\text{g L}^{-1}$ )	<b>-0.903</b>	-0.355	-0.432	<b>0.811</b>	0.135	0.400	0.166	0.162	-0.252
NH <sub>4</sub> ( $\mu\text{g L}^{-1}$ )	0.375	0.576	<b>0.543</b>	-0.433	-0.338	<b>-0.579</b>	-0.382	0.013	<b>0.530</b>
TN ( $\text{mg L}^{-1}$ )	<b>-0.487</b>	0.035	<b>-0.628</b>	0.441	0.268	<b>0.613</b>	0.420	-0.633	-0.417
TDN ( $\text{mg L}^{-1}$ )	<b>-0.534</b>	-0.219	-0.305	<b>0.482</b>	0.533	<b>0.479</b>	0.324	<b>-0.921</b>	<b>-0.633</b>
TN:TP	<b>0.570</b>	0.299	<b>0.464</b>	<b>-0.593</b>	-0.257	-0.382	0.492	0.514	0.239

15 Significant ( $P < 0.05$ ) factors are highlighted in bold;

16 N = number of samples.

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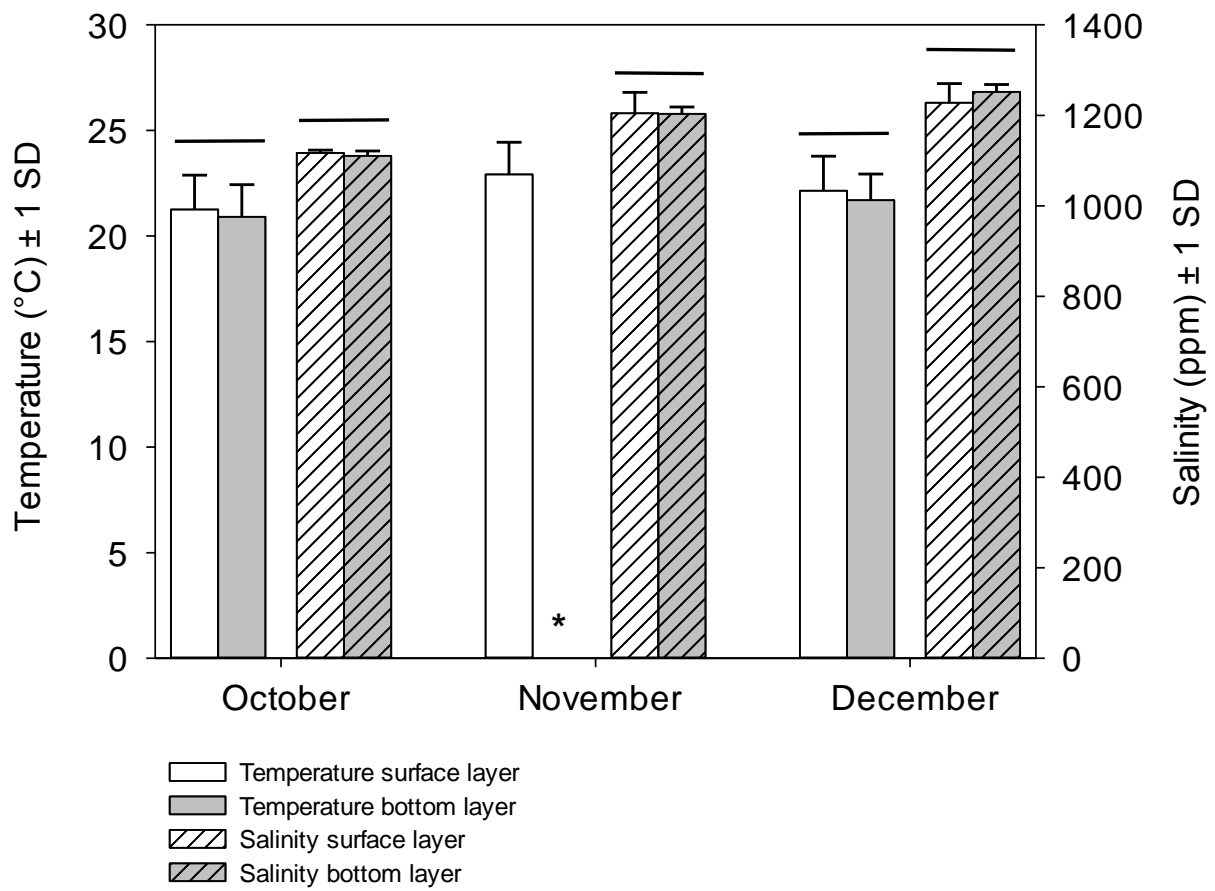
20 **Fig. 1.**The locations of three studied lakes on Swan Coastal Plain.

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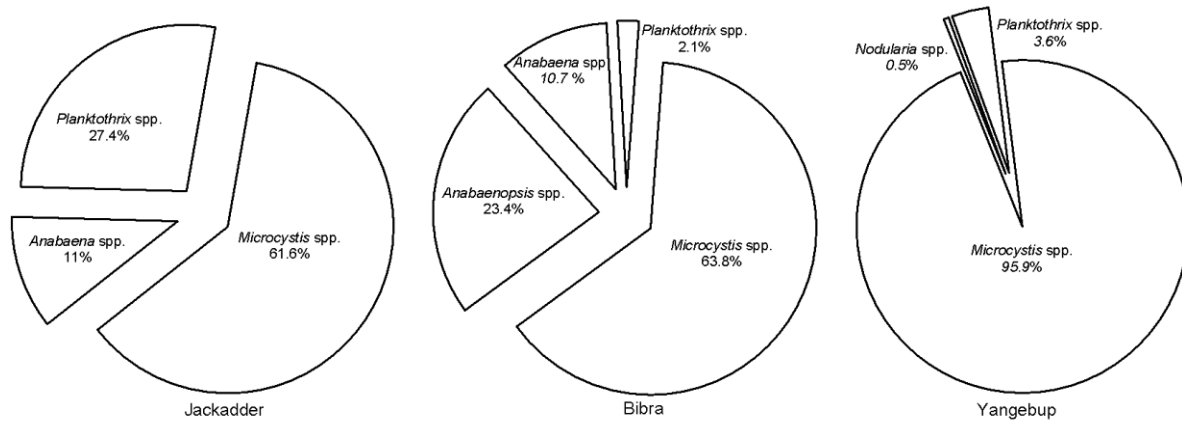
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**Fig. 2.** Temperature (°C) and salinity (ppm) in the surface and bottom layers measured at 7 sites over three months in Lake Yangebup during a previous study in 2012. \* = missing data; horizontal line indicates that no significant difference between data were detected (t-test).

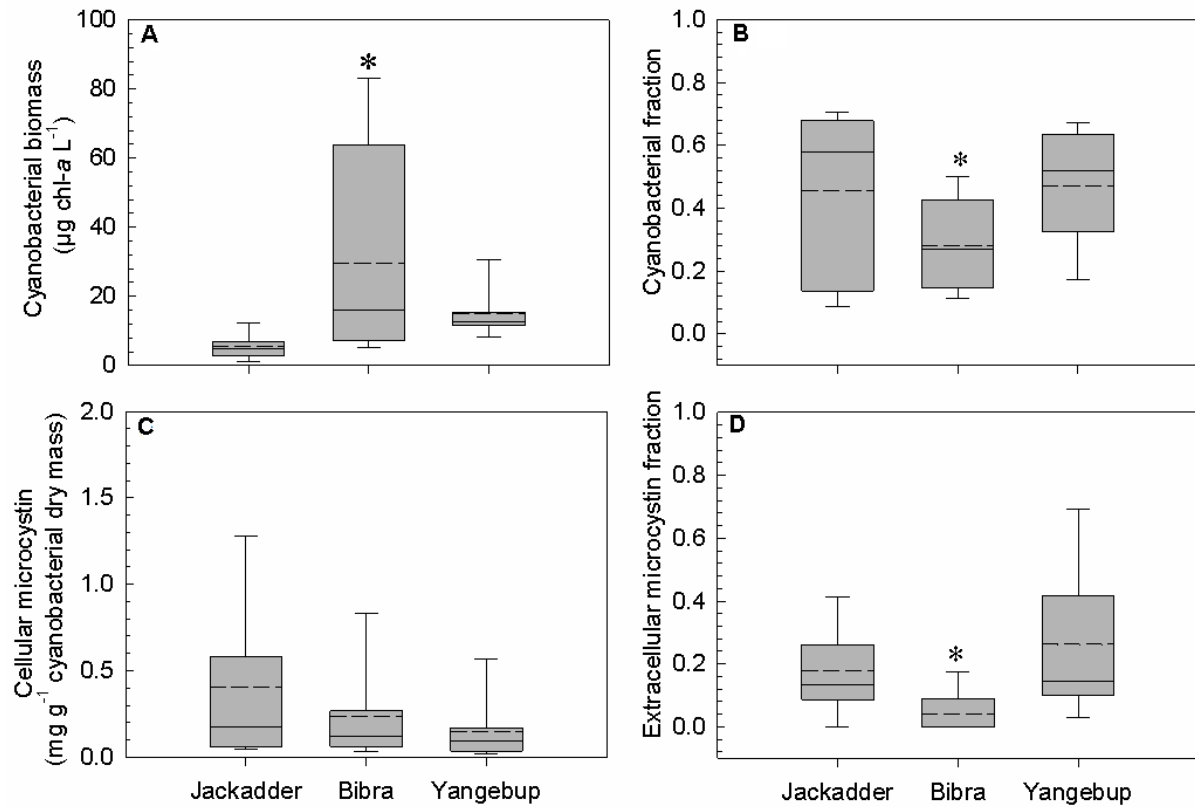




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32 **Fig. 3.** Average biomass ( $\mu\text{m}^3 \text{mL}^{-1}$ ) proportions of potentially toxic cyanobacterial genera in  
 33 Jackadder, Bibra and Yangebup lakes during the study period.

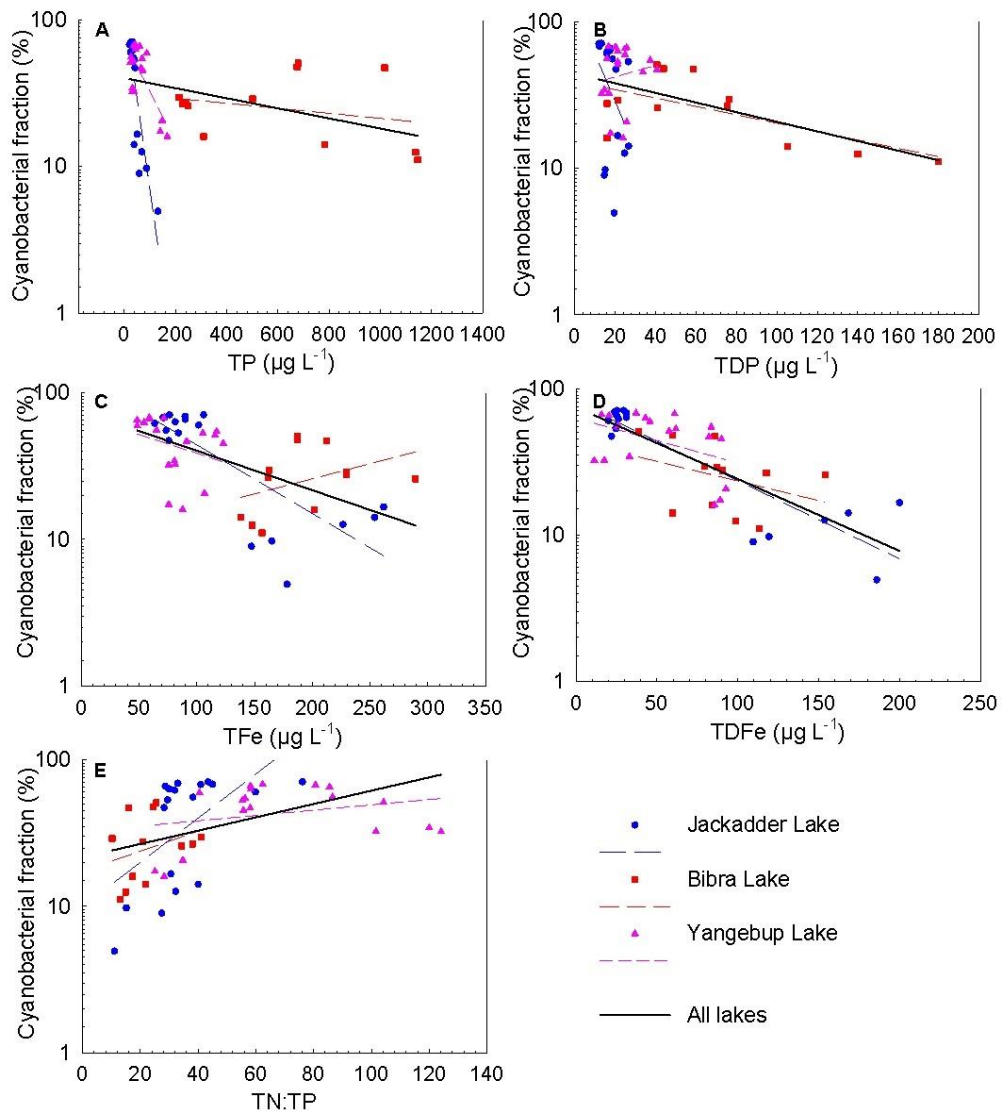
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36 **Fig. 4.** The variability of (A) cyanobacterial biomass ( $\mu\text{g chl-}a \text{ L}^{-1}$ ), (B) cyanobacterial  
 37 fraction (cyanobacterial biomass to total biomass), (C) cellular microcystin concentration ( $\text{mg}$   
 38  $\text{g}^{-1}$  cyanobacterial dry mass) and (D) extracellular microcystin fraction over time for each  
 39 lake. Boxes represent 25<sup>th</sup> to 75<sup>th</sup> percentiles; straight lines within the boxes mark the median  
 40 short dashed lines the mean; whiskers below and above the boxes indicate 10<sup>th</sup> and 90<sup>th</sup>  
 41 percentiles. Asterisks (\*) indicated lakes that are significantly ( $P < 0.05$ ) different from other  
 42 lakes.

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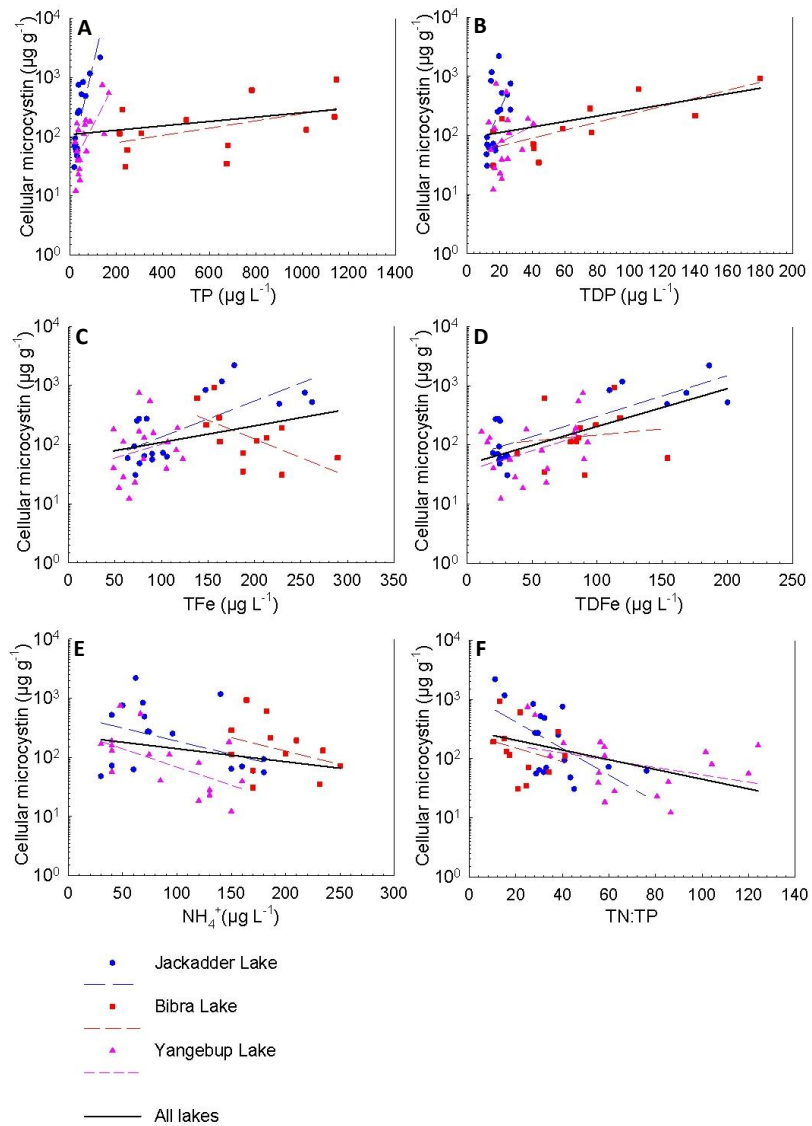


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45 **Fig. 5.** The correlations between cyanobacterial fraction and (A) TP, (B) TDP, (C) TFe, (D)

46 TDFe, (E) TN:TP in Jackadder, Bibra and Yangebup lakes during the study period.

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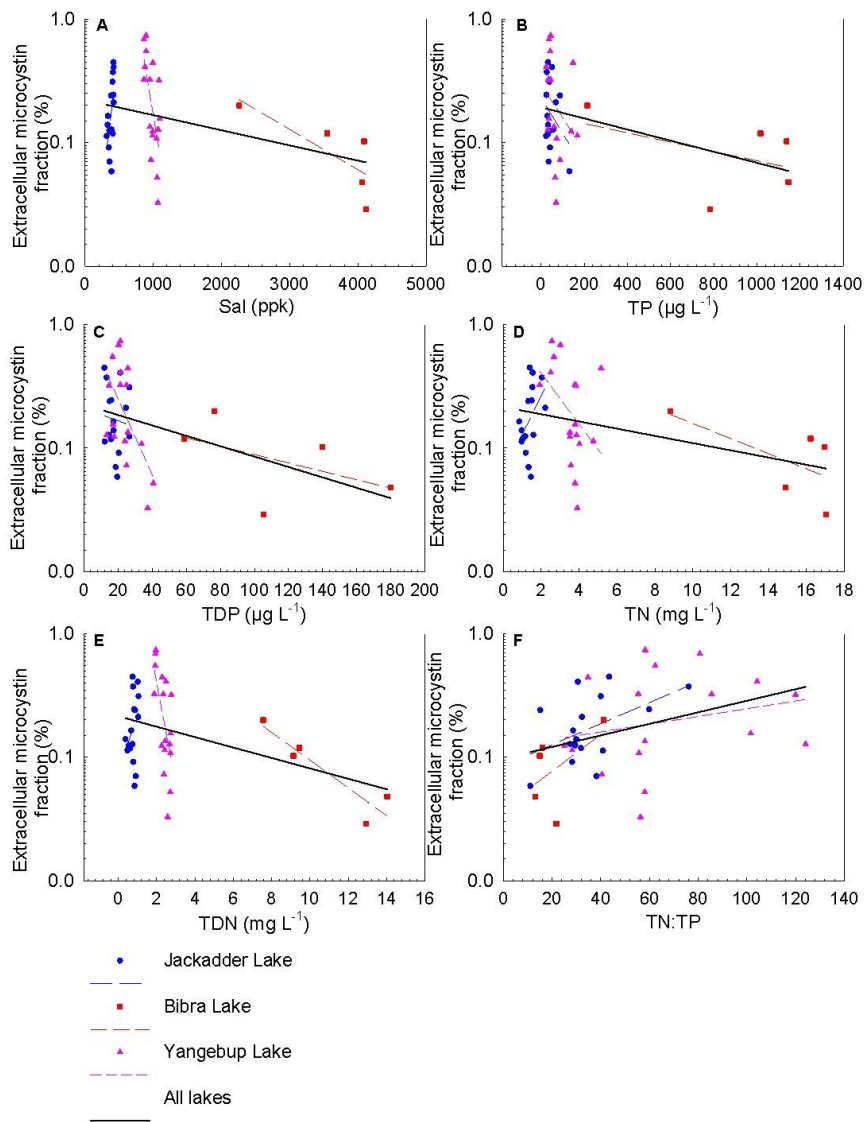
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49 **Fig. 6.** The correlations between cellular microcystin concentration and (A) TP, (B) TDP, (C)

50 TFe, (D) TDFe, (E)  $\text{NH}_4^+$ , (F) TN:TP in Jackadder, Bibra and Yangebup lakes during the

51 study period.

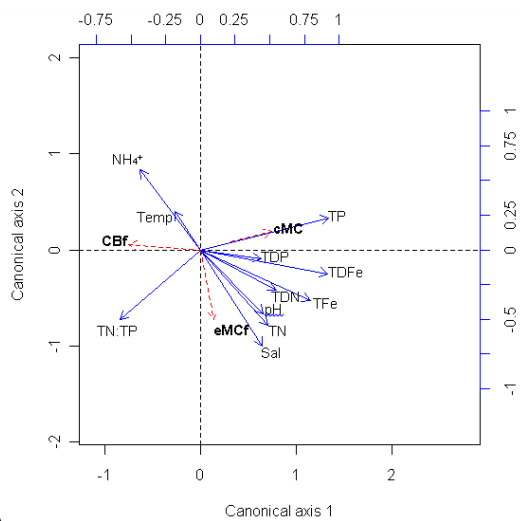
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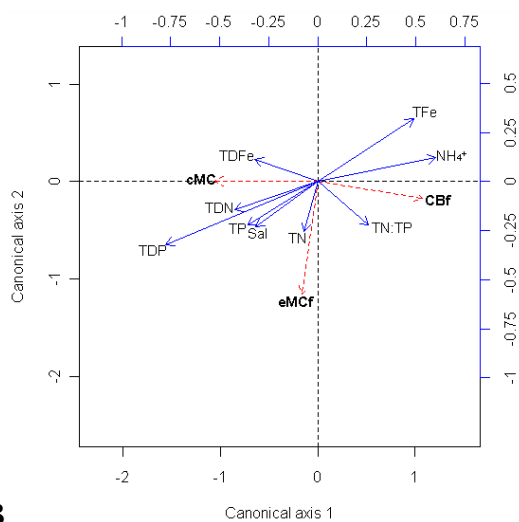
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54 **Fig. 7.** The correlations between extracellular microcystin fraction and (A) Sal, (B) TP, (C)  
 55 TDP, (D) TN, (E) TDN, (F) TN:TP in Jackadder, Bibra and Yangebup lakes during the study  
 56 period.

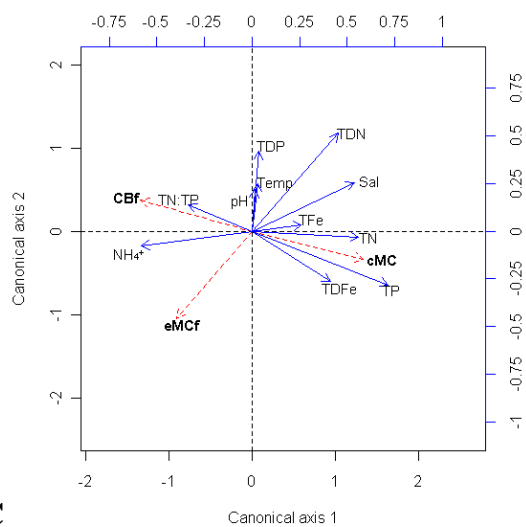
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**A**



**B**



**C**

60 **Fig. 8.** RDA biplots of environmental variables with cyanobacterial fraction (CBf), cellular  
61 microcystin (cMC) and extracellular microcystin fraction (eMCf) in (A) Jackadder Lake, (B)  
62 Bibra Lake, (C) Yangebup Lake; solid arrows = environmental variables; short dashed arrows  
63 = response variables. Canonical axis 1 and 2 represents a linear combination of the  
64 environmental variables, and axes are scaled by the square root of their eigenvalues.

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