1	Rise of toxic cyanobacterial blooms is promoted by agricultural intensification in the basin of
2	a large subtropical river of South America
3	
4	Running head - 45 characters including spaces: Agro-intensification promotes Cyanobacteria
5	
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21	
22	Summary
23	Toxic cyanobacterial blooms are globally increasing with negative effects on aquatic ecosystems,
24	water use and human health. Blooms' main driving forces are eutrophication, dams construction,
25	urban waste, replacement of natural vegetation with croplands, and climate change and variability.

27 basins. Here, we performed a historical analysis of cyanobacterial abundance in a large and

The relative effects of each driver have not still been properly addressed, particularly in large river

important ecosystem of South America (Uruguay river, c.a. 1,900 km long, 365,000 km² basin). We 28 29 evaluated the interannual relationships between cyanobacterial abundance and land use change, 30 river flow, urban sewage, temperature and precipitation from 1963 to the present. Our results 31 indicated an exponential increase in cyanobacterial abundance during the last two decades, 32 congruent with an increase in phosphorus concentration. A sharp shift in the increase rate of 33 cyanobacterial abundance after the year 2000 was identified, resulting in abundance levels above 34 public health alert since 2010. Path analyses showed a strong positive correlation between 35 cyanobacteria and cropland area at the entire catchment level, while precipitation, temperature and 36 water flow effects were negligible. Present results help to identify high nutrient input agricultural 37 practices and nutrient enrichment as the main factors driving toxic bloom formation. These 38 practices are already exerting severe effects on both aquatic ecosystems and human health and 39 projections suggest these trends will be intensified in the future. To avoid further water degradation 40 and health risk for future generations, a large scale (transboundary) change in agricultural 41 management towards agroecological practices will be required.

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43 Key words. Cyanobacterial blooms, land use, crops, temperature, precipitation, health risk.

44 INTRODUCTION

45

The occurrence of massive toxic blooms of cyanobacteria is an environmental and public health 46 47 problem worldwide. Most blooms produce metabolites that are toxic to animals and humans, which are collectively referred to as cyanotoxins (Chorus & Welker, 2021; Preece et al., 2017; Svirčev et 48 49 al., 2019). These cyanotoxins can lead to sickness and cause deaths of aquatic animals, including 50 negative impacts on human health and other activities (Azevedo et al., 2002; Giannuzzi et al., 2011; 51 Vidal et al., 2017; Zhang et al., 2015). Over the past few decades, an upsurge in the frequency, 52 geographic distribution and severity of such blooms has recurred in many areas of the planet 53 (Preece et al., 2017) and forecast models indicate that both bloom frequency and intensity will 54 continue to increase (Fang et al., 2022). Therefore, disentangling the main driving causes to define 55 management actions is of fundamental importance (Sukenik & Kaplan, 2021).

56

57 Cyanobacterial bloom forcings

58 Understanding cyanobacterial toxic bloom dynamics is a complex issue and there are several 59 driving forces linked to their occurrence and toxicity. However, there is a 50-year consensus in the 60 scientific community about the fundamental role that the input of nutrients to water bodies (due to 61 cultural eutrophication) plays in driving algal biomass increment and cyanobacterial dominance 62 (Huisman et al., 2018; Paerl et al., 2018). The identification of nutrient enrichment as a relevant 63 driver can be traced back to early empirical studies (e.g. Vollenweider & Kerekes, 1982; Trimbee & 64 Prepas, 1987), including manipulative experiments at the ecosystem-scale (Schindler, 1977) and has 65 most recent support in cross-system field and laboratory studies of large data sets at different latitudes (Downing et al., 2001; Kosten et al., 2012). These studies demonstrate that bloom forming 66 cyanobacteria benefit over other phytoplankton groups under high nutrients and low turbulence 67 68 conditions (e.g. Kim et al., 2017; Scheffer et al., 1993).

69

70 Cultural eutrophication is originated by multiple human activities including agriculture, cattle rising

and poorly-treated sewage water from human settlements (Fang et al., 2022), which were traced back by paleolimnological studies (Bueno et al., 2021). Among the multiple human-related driving forces, agriculture is a prominent and persistent cause of diffuse nutrient loads and the world largest source of both phosphorus and nitrogen pollution to water bodies (Bennett et al., 2001; Chakraborty et al., 2017; MacDonald et al., 2011; Withers et al., 2014). A strong linkage has been identified between the industrialized cutting-edge production of crops and the emergence of toxic cyanobacteria outbreaks (Chakraborty et al., 2017; Massey et al., 2020).

78

79 There are other drivers that also promote cyanobacterial blooms and might exert synergistic effects 80 with eutrophication (Meerhoff et al., 2022). Most cyanobacterial organisms have relatively low 81 specific growth rates and thus toxic populations are not able to proliferate under low to moderate flushing (Bakker & Hilt, 2016; Kruk et al., 2021). Therefore, in eutrophic rivers, dam construction 82 83 favors blooms by increasing water residence time (Massey et al., 2020; Paerl & Huisman, 2009). 84 High temperatures also accelerate cyanobacterial metabolism, promoting their growth, the 85 formation of blooms under stratified waters (Paerl & Huisman, 2008) and the stimulation of toxin 86 production (Martínez de la Escalera et al., 2017; Massey et al., 2020). Changes in precipitation modify water residence time and geographical distribution of cyanobacterial blooms (Massey et al., 87 88 2020; Kruk et al., 2021). Moreover, the replacement of natural land cover with crops and the 89 modification of trophic food webs accelerates the effects of eutrophication and climate change (Alcántara et al., 2022; Meerhoff et al., 2022). Thus, understanding the importance of individual 90 91 drivers and potential synergies among them are both relevant to provide management actions to 92 stakeholders at different levels (Fang et al., 2022; Alcántara et al., 2022; Kosten et al., 2012;).

93

94 Watersheds in Southern South America

South America is one of the regions most affected by global agriculture, where large-scale cash
crops have replaced natural vegetation (Doughty, 2010; Graesser et al., 2018; Perez et al., 2021;
Tucci & Clarke, 1998). Large proportions of land are dedicated to agricultural production with

annual crops, such as soybean, maize, wheat and rice (Giller et al., 2021). These large-scale cash
crop areas have transformed land use dynamics with important implications for food security,
biodiversity conservation, and revenues (Bueno et al., 2021; Graesser et al., 2018; Tucci, 2001;
Tucci & Clarke, 1998).

102

103 Within South America, the Uruguay river is a major and vital ecosystem (c.a. 1,900 km long) with a transboundary catchment area of 365,000 km² shared by three countries (Brazil, Argentina and 104 105 Uruguay). This river supports key ecosystem services, drinking water for human consumption, recreation and tourism, navigation, as well as biological production at different trophic levels 106 including fisheries (de Vasconcelos et al., 2014; Tucci & Clarke, 1998). It integrates the Río de la 107 108 Plata basin, the second largest in South America (Guerrero et al., 1997; Milliman et al., 2008) and represents an important region for agriculture and livestock production in the world (de 109 110 Vasconcelos et al., 2014; Tucci, 2001; Tucci & Clarke, 1998). During the last decades, natural 111 grasslands supporting extensive livestock production have been replaced with large scale croplands, causing degradation of ecosystems services (de Vasconcelos et al., 2014; Modernel et al., 2016). 112 113 High-resolution palaeoceanographic studies in the Río de la Plata demonstrated the effect of 114 continental human activities on the aquatic ecosystems (Pérez et al., 2021a; Pérez et al., 2021b) attributed to land-use intensification and dam construction leading to significant soil erosion and 115 116 transport/deposition into the inner shelf, enhanced by the El Niño Southern Oscillation (ENSO) 117 events (Bonachea et al., 2010; Pérez et al., 2021a; Pérez et al., 2021b). The Uruguay river holds 10 large dams (> 10 km^2) and there are more than 90 further dams either planned or under construction 118 (http://globaldamwatch.org) (Pérez et al., 2021a), which cause strong modifications to its 119 120 hydrological cycle (Depetris & Pasquini, 2007a, b; Milliman et al., 2008; Tucci & Clarke, 1998).

121

Toxic cyanobacterial blooms are frequently registered in the Uruguay River (Bordet et al., 2017;
Debastiani Júnior et al., 2016; O'Farrell & Izaguirre, 2014) and exported to the Río de la Plata
Estuary (Nagy et al., 2002; Kruk et al., 2017; Martínez de la Escalera et al., 2017; Segura et al.,

125 2017). These blooms negatively impact health, including human intoxications (Giannuzzi et al., 126 2011; Vidal et al., 2017), local economies, tourism, and related recreational activities (Kruk, 127 Martínez de la Escalera et al., 2021). Long term records of toxic cyanobacterial blooms and the 128 analysis of their forcings are not common in large watersheds worldwide (Doubek et al., 2015; Salk 129 et al., 2022), and particularly in the Southern Hemisphere, where these type of analysis are scarce 130 (Silvarrey Barruffa et al., 2021). The objective of this study was to disentangle the contribution of 131 different environmental drivers in determining cyanobacterial abundance. With this aim, we analyzed temporal trends which showed an increase in the frequency and severity of cyanobacterial 132 133 blooms over a 60-year period in a large subtropical river basin (Uruguay river basin). Also, causal 134 path models were used to evaluate the relative importance of each driving forcing (land use change, 135 urban sewage, river flow and climate variability (temperature, precipitation) in explaining 136 cyanobacterial abundance at the entire-basin level.

137

138 METHODS

139

140 Study area

The Uruguay river basin lies between 28°10'S and 37°08'S (Di Persia & Neiff, 1982) and it is shared by Brazil (percentage of surface area: 50.1%), Uruguay (32.5%) and Argentina (17.4%) (Figure 1). The river origin is located in Brazil at 200 meters above mean sea level and its main tributary is the Negro River (500 km long). Together with the Paraná River are the main tributaries of the Río de la Plata Estuary.

146

The Uruguay river hydrographic basin climate is located in a humid subtropical (*sensu* Köppen) and is set upon sedimentary and volcanic soils. It comprises mainly three dominant biomes from headwaters to the Río de la Plata: i) upper basin: mixed agro-eco-region, ii) middle basin: Campos biome and iii) lower basin: South American Pampas biome (Argentina, Uruguay) (Saurral et al., 2008; Viglizzo & Frank, 2006). The upper river is steep (43 cm km⁻¹), comprises 750 km, has a

maximum average flow of 9.387 m³sec⁻¹ and exhibits large annual variations in water level. The Itá 152 Hydroelectric Power Plant is the largest reservoir in this area (lake area: 141 km², basin area: 153 45,800 km²) and it became operative in 2000. Additionally, eight dams are now built in the river 154 155 and other 89 are planned for the next few years (http://globaldamwatch.org, Mulligan et al., 2020) (Figure 1). The middle zone of the river is 800 km long, has a 9 cm km⁻¹ slope, and does not contain 156 large reservoirs. The lower river is 350 km with a slope 3 cm km⁻¹ and its hydrology is strongly 157 influenced by the Salto Grande reservoir, constructed for hydroelectricity production (Concordia, 158 Salto, 31°16'29"S 57°56'18"W). Salto Grande became first became operative in 1979 with a 159 catchment area of 224,000 km² and a lake surface area of 783 km². Finally, at Nueva Palmira, the 160 Uruguay river flows into the Río de la Plata Estuary, and then into the Atlantic Ocean (Figure 1). 161

162

163 Cyanobacterial abundance and limnological variables

Cyanobacterial information including abundance and blooms presence was compiled for the lower 164 165 Uruguay river from different data sources, attaining a total number of 1.712 cases from 1963 to 2019. From the total number of cases, 1,424 were retrieved from sampling sites located either 166 upstream or in the Salto Grande (SG) dam and 288 from sites downstream this dam. The largest part 167 168 of the data set was obtained from publicly accessible reports of the Uruguayan Government Water Institution (i.e., Obras Sanitarias del Estado, OSE, 2009), for the period between 1963 and 2008. 169 This information included cyanobacterial cells abundance (cells mL⁻¹) in surface water, dominant 170 171 cyanobacterial species and the presence of blooms in six sites next to three water treatment plants: 1) upstream SG reservoir: Bella Unión, 2) in the SG reservoir: Belén and Constitución, and 3) 172 downstream the reservoir in Salto, Fray Bentos, Paysandú and Bassotti. From 2008 until 2019 we 173 174 gathered information from other institutional sources, including the Argentina-Uruguay Administrative Commission (CARU, 2019, https://www.caru.org.uy/web/), SG Joint Technical 175 176 Commission (CTM, Argentina-Uruguay, https://www.saltogrande.org/) and the National Environmental Observatory (MVOTMA-OAN, 177 of Uruguav 2019; https://www.ambiente.gub.uy/oan/). These data sources included cvanobacterial cells abundance, 178

179 cyanobacterial species, and fecal coliforms abundance (as fecal indicator bacteria, FIB) in 16 sites 180 from Bella Union (Artigas) to the SG reservoir main channel next to the dam. Total phosphorus 181 concentration in water (mg L⁻¹) was retrieved for the same sites for the period 2008 to 2019, 182 including information from CTM (N = 245), CARU (N= 25) and our own monitoring program 183 (Kruk et al., 2015; N = 72).

184

In all cases, surface water samples were taken with bottles and fixed with Lugol solution. 185 Cyanobacterial and other phytoplankton species were counted with comparable methods with 186 Sedgwick Rafter or Utermöhl counting chambers using a light microscope and estimating the cell 187 abundance (cell mL⁻¹) using the settling technique (Utermöhl, 1958). Utermöhl chambers were used 188 189 for counting samples with low phytoplankton abundances, while Sedgwick chambers were used for 190 high abundance samples including those dominated by cyanobacteria and with presence of blooms. The abundance of cyanobacteria was also used to determine alert levels following (Chorus & 191 Welker, 2021; level 1: 2.000 - 100.000 cell mL⁻¹, level 2: higher than 100.000 cell mL⁻¹, vigilance 192 stage: cell abundance higher than 200 cell mL^{-1}). 193

194

The effects of human settlements were estimated based on the abundance of fecal indicator bacteria (FIB). Samples for FIB detection and quantification were taken from surface water at the same area as cyanobacteria but with a higher frequency from 1987 to 2020, attaining a total number of 2,127 cases. For the 1987-2005 period, the information was retrieved from the binational Procon-CARU project (<u>https://www.caru.org.uy/web/2020/05/programa-procon-caru-1987-2005-recopilacion-y-</u> validacion-de-datos/). For the 2014-2020 period, the information source was the Uruguay National Environmental Observatory (OAN) (<u>https://www.ambiente.gub.uy/oan/</u>).

202

203 Climatological and hydrological variables

Temperature and precipitation data in the Uruguay river watershed from 1963 to 2020 were obtained from the historical climate database produced by the Climatic Research Unit (CRU) of the 206 University of East Anglia and provided by the World bank climate knowledge portal, corresponding 207 to number 252 watershed (https://climateknowledgeportal.worldbank.org/watershed/252/climate-208 data-historical). The information included annual average values of maximum, minimum and 209 average daily temperatures (°C) and the average annual precipitation (mm). Historical data of 210 Uruguay river flow rates were retrieved from the Argentinian National System of Information 211 (https://snih.hidricosargentina.gob.ar) and corresponded to station 3802 in Paso de los Libres. Monthly average water flow $(m^3 \text{ sec}^{-1})$ was selected from 1963 to 2020 and expressed as the 212 213 maximum value and the accumulated flow per year.

214

215 Land Use

Annual land use and land cover maps from 2000 to 2019 for the entire Uruguay river basin were obtained from an assembly of classifications generated by MapBiomas initiatives (Table 1). The MapBiomas project is an international multi-institutional initiative whose goal is to generate annual land use and land cover maps based on the use of satellite images provided by Landsat 4, 5, 7 and 8 and automatic classification processes of images. These included MapBiomas Pampa (Baeza et al. 2022) and Atlantic Forest Tri-national initiatives (Souza et al., 2020).

222

Land use and land cover were grouped into six classes natural forest, forest plantation, grasslands and wetlands, farming, non-vegetated areas (that include bare soil and urban infrastructure) and water bodies and rivers (Souza et al., 2020). Farming includes annual crops (i.e. maize, soybean, wheat, etc.) and sown pastures. Information for a longer period (from 1989 and 2019) was only available for the Brazilian portion of the basin. Collection 6 of Map-Biomass Brazil was used to separate the composition of the class "annual crops and pasture" into different land use covers, such as: pastures, soybeans, other summer crops and a "mosaic of pastures and annual crops".

230

231 Data analysis

232 Cyanobacterial abundance was log_{10} transformed for most analyses and graphical representations. 233 Linear models were performed to evaluate the change in cyanobacterial abundance and total 234 phosphorus yearly average with time and were compared using the differences in the Akaike 235 information criteria (ΔAIC). The fit of the segmented models followed a well developed technique 236 based on a linearization of the problem that allow to estimate slopes and intercepts before and after 237 a breakpoint, as well as a confidence interval for the breakpoint (Muggeo, 2003) which is implemented in the {segmented} package (Muggeo, 2008) in the R software. The relationships 238 239 between total phosphorus and cyanobacterial abundance, and between total phosphorus and land 240 use areas were explored with correlations and linear models. Due to the lack of sufficient available 241 data points and to the heterogeneity in the methodological approaches applied to estimate nitrogen 242 in water, this nutrient was not included in the analysis.

243

244 To evaluate the effect of the unequal sample size and the increasing sampling effort with time on 245 the estimation of cvanobacterial abundance (Hallegraeff et al., 2021) we conducted different strategies (Supplementary material 1); i) aggregating the information as yearly average both 246 247 including and excluding the periods with less data (between 1995 and 2003), ii) aggregating the information every 5-years and performing unweighted and weighted regressions (weights 248 proportional to the number of data points per period), and iii) using bootstrapping type of re-249 250 sampling to produce 250 smaller samples of 15 cases and then producing 250 temporal linear 251 models with the yearly average in each period. In iii) the number of significant models with positive 252 slopes was registered. The models with similar response variable were compared using AIC 253 (Supplementary material 2).

254

Temporal trends in temperature (average, maximum, minimum), precipitation average, river flow (accumulated and maximum), and log_{10} cyanobacterial abundance were explored for each data set. The presence of white noise, was evaluated in the residuals of the segmented regression on yearly clustered information, using Ljung-Box and turning points tests. The results did not allowed to discard white noise and therefore specific time series analyses were not necesary. Spearman nonparametric rank correlation and the variance inflation factor (VIF) were used to evaluate multicollinearity among variables, including climatic, hydrological and land use.

262

263 Causal path models were constructed to evaluate the relative contribution as well as the interactions 264 of the different potential driving factors of the annual average of \log_{10} (cyanobacterial abundance+1) (logCya) using {"Lavaan"} package. This model framework is specially designed to 265 266 evaluate competing models representing hypothesis based on theoretical arguments (Petraitis et al., 1996). We *a priori* introduced eight path models, including the effect of climate (precipitation: PP, 267 maximum temperature: Tmax), hydrology (accumulated river flow: Oacum), and land use (area of 268 269 land with annual crops and pastures and natural area) (Figure 2). Covariates were standardized to 0-1 prior to the analysis. To evaluate the existence of multicollinearity among the explaining 270 271 variables, we inspected the VIF. No significant multicollinearity was retrieved as VIF values were 272 in all cases lower than four. Competing models were compared based on the AIC and paths coefficients were estimated including their standard error (SE), z statistics and the associated p 273 274 value. The Fisher's C statistic was calculated to evaluate whether the data follow the causal hypothesis specified in the dyacyclic aggregated paths. For the selected model, the root mean square 275 error of approximation (RMSEA), the explained variance (R^2) and the comparative fit index (*cfi*) 276 277 were estimated. Models were performed for the 1985-2019 period using land use covariates from the Brazilian catchment area. 278

279

280 **RESULTS**

281 Temporal Changes in cyanobacterial abundance and composition

Total abundance of cyanobacterial cells in the Salto Grande reservoir and downstream increased by 4 to 5 orders of magnitude from 1963-1970 (average 33 cell L⁻¹, maximum = 53 cell L⁻¹) to 2010-2020 (average 21281 cell L⁻¹, maximum = 1.69 x 10^6 cell L⁻¹; Figure 3) and phytoplankton communities shifted from either diatom- or chlorophyte-dominated to cyanobacteria-dominated. The dominant cyanobacterial genera shifted from the first decades when *Merismopedia*, *Lyngbya*, *Planktothrix* and other genera from the Oscillatoriales order dominated, to *Dolichospermum* and *Microcystis* after the 2000s. In recent years, blooms were in most cases dominated by members of the *Microcystis aeruginosa* complex.

290

291 After 2000, blooms became highly frequent along with an exponential increase in both maximum and average cell abundance (Figure 3). Surface scums (i.e., water discoloration events) became 292 293 frequent from 2000 to 2007, particularly in the lower Uruguay river (Figure 1). Further, the risk level alerts also increased with time, from no risk to vigilance (cell abundance higher than 200 cell 294 mL^{-1}) in the period between 1963 to 2005, then from vigilance to at least one level 1 alert (2.000 -295 100,000 cell mL⁻¹) per year, and since 2006 at least one level 2 alert (100,000 cell mL⁻¹) per year 296 was observed. During the last decade (2010-2020), 155 and 46 alerts of level 1 and 2, were 297 298 respectively reported. In addition, the temporal persistence of blooms throughout the year increased 299 from isolated occasions in summer to continuous presence, including even the cold seasons (winter and autumn). Between 1960 and 2000 no winter or autumn blooms occurred; between 2000 and 300 301 2010, 24 cases of level 1 alert and one case level 2 alter were observed in winter. In the last decade, 302 65 level 1 and 16 level 2 alerts developed during winter and autumn.

303

304 Temporal dynamics of yearly average cyanobacterial abundance, presented two different periods 305 and was well characterized by a segmented log-linear model: I) a first period (1963-2000) where average cell abundance remained constant with time (slope [95%CI]= -0.02[-0.05 to 0.01]), a 306 breakpoint after year 2000 [CI95%=1995 to 2005], and II) a second period characterized by an 307 308 exponential increase (slope [95%CI]= 0.21[0.13 to 0.30]) (Figure 3). The results remained similar either by including or excluding years with less data (between 1995 and 2003) or aggregating the 309 310 information into 5-year periods. Linear weighted regressions also provided similar trends and coefficients and thus we decided to keep the simpler approach. In addition, when data were 311 aggregated in 5-year bins and the lineal model was performed 250 times, the fitted temporal models 312

indicated a positive trend of cyanobacterial abundance increase in 100% of the cases, beingsignificant in 93% of the cases (Supplementary material 2).

315

316 Climate and hydrological changes observed in the Uruguay river basin

Average, maximum and minimum annual temperature at the watershed level were significantly correlated (minimum vs mean temperature: r=0.93; maximum vs mean temperature: r=0.84; minimum vs maximum temperature: r=0.58; N=60, all p <0.01) (Figure 4A). Minimum annual temperatures presented significant positive temporal linear trends since the beginning of the time series (T_{min} =-22.11 + 0.018 year; N= 60, R²=0.32, p <0.01). A simple linear trend was chosen with respect to the more complex segmented model as the difference in AIC was low (Δ AIC<2).

323

Precipitation (PP) in the watershed showed an apparent linear temporal trend in the period 1963-2020 but this trend was not significant (p >0.05; $R^2=0.05$) and no breakpoints were detected (Figure 4B). Neither the accumulated flow nor the annual maxima of the Uruguay river flow presented temporal trends (p >0.05; $R^2=0.004$ and p >0.05; $R^2=0.025$ respectively) (Figure 4C).

328

329 Correlation analyses between climatic variables for the period 1963-2020 indicated that average annual precipitation and minimum temperature had a weak positive linear correlation (r=0.52, 330 p<0.01). Precipitation presented a strong correlation with water flow, either accumulated (r=0.83, p 331 <0.01) or maximum flow (r=0.73, p <0.01). Minimum temperature exhibited a weak but significant 332 correlation with yearly precipitation and water flow (r_{PP}=0.52; r_{Omax}=0.42; r_{Oacum}=0.4; all p values 333 334 <0.01) while maximum yearly temperature displayed a negative weak correlation with accumulated 335 water flow (r_{Oacum}=-0.35). Maximum and accumulated water flow were strongly correlated 336 (Supplementary material 3).

337

338 Changes in indicators of human and cattle effects

From 1987 to 2020 the collected FIB exhibited a variability of six orders of magnitude, ranging from undetectable to 2 x 10^6 UFC $100mL^{-1}$, a median value of 160 and a mean of 5,875 UFC $100mL^{-1}$ (Figure 5). A negative temporal trend in FIB concentration was described by a log-linear model (slope[95%CI]= -0.017 [-0.02 to -0.013]) with a low but significant explained variance (R²=0.04, p<0.001, N=2119).

344

345 Temporal trends in land use change in the Uruguay river basin

346 The Uruguay river watershed displayed a marked transition in land use from 2000 to 2019 (Figure 1 and Figure 6). The most conspicuous patterns were a 17% increase in land use dedicated to crops 347 and pastures (23.842 km²), a 97% increase in the forestation area (10.047 km²), a decrease in 6% of 348 natural forest (3,004 km²) and a decrease in 17% of natural grasslands (23,842 km²). This last 349 350 change represents a loss of ~8% of natural grasslands replaced with intensive agricultural land use, 351 including crops and forestation. Pastures and crops, as well as forestation gains occurred along the 352 entire basin, with larger proportions in the middle and lower areas (Figure 1). In the upper basin the loss of pastures and crops were mainly detected in the littoral zones of the river (Figure 1). Data 353 354 recorded in the upper and middle basins, available for 1985 to 2020, presented the same trend and were highly correlated with the changes registered in the lower basin for the 2000-2020 period 355 (Table 2; Supplementary material 4). In addition, the proportion of land uses remained similar 356 357 throughout most of the study, therefore representing a proxy for total watershed changes (Table 2).

358

A major increase in land use dedicated to soybean production occurred along with a large decrease in other crops together with depleted pastures and the class "mosaic of pastures and annual crops". Soybeans were first introduced in 2000 but nowadays accounts for nearly 45% (4,114 km²) of the annual cropland area (Figure 7). Pastures, other crops and the "mosaic of pastures and annual crops" decreased by 34, 24 and 48%, respectively.

364

365 Importance of drivers explaining cyanobacterial abundance

366 The causal path model indicated that changes in the area dedicated to crops and pastures (CP) 367 displayed the strongest and significant positive relationship with cyanobacterial abundance (logCya) (Table 3 and Figure 8). Model 8 presented the larger difference in AIC ($\Delta AIC>35$) with the 368 remaining models and included only the path between CP and logCya as significant (Table 3, 369 Figure 8). A positive relationship between annual precipitation and accumulated flow was also 370 observed, but this relationship did not affect cvanobacterial abundance (Table 3). The selected 371 372 model Fisher's C statistic was not significant (C=1.50, df=2, p=0.47) supporting the hypothesis that data were generated following the causal relations specified in the dyacyclic aggregated path 373 (Model 8, Table 4). The RMSE of approximation was low (95% confidence interval [~0-0.36], 374 375 p=0.49) while the comparative fit index was high (*cfi*=1.00) as well as the explained variance $(R^2=0.52)$. Precipitation accounted for a high amount of variance from the accumulated water flow 376 (R²=0.79). Similar results were obtained when Tmin was included in the model instead of Tmax 377 (Supplementary material 5). 378

379

380 Relationship with total phosphorus concentration in water

381 An increase in average total annual phosphorus in water (TP) was observed with time (TPmean-5.98 + 0.003 year; R^2 =0.66, p < 0.005). Cyanobacterial cells abundance (Cya) was significantly 382 related with TP values higher than 0.2 mgL⁻¹ (N= 46, Pearson correlation: 0.46; $\log_{10}(Cya+1) =$ 383 $2.22*TP mgL^{-1} + 1.96$; df = 44, R² = 0.18, p< 0.005) (Figure 9). For TP values lower than 0.2 mgL⁻¹ 384 no significant correlation was obtained. Finally, we also obtained a positive relationship between 385 average annual TP concentration in water and area dedicated to annual crops and pastures (TP mg 386 $mL^{-1} = 1.79 \text{ *}\log_{10}(\text{Annual crops and pasture area}) - 8.93, R^2 = 0.54, p < 0.05;$ Figure 10A) and a 387 negative relationship with natural grassland area (TP mg mL⁻¹ = $-1.70*\log_{10}(Grasslands area) +$ 388 8.79, $R^2 = 0.63$, p<0.05; Figure 10B). 389

390

391 **DISCUSSION**

392 In this manuscript we evidenced the exponential increase in cyanobacterial abundance and number

393 of blooms in a relevant subtropical aquatic ecosystem of South America. A comprehensive 394 geographical long-term analysis confirmed that the increase of intensive agriculture land use was 395 the main driver. Climatic and hydrological effects on cyanobacterial abundance, otherwise, were 396 rather marginal. Moreover, results indicate that models not including the effect of land use change 397 would incorrectly point to temperature as a relevant variable, but under a causal path analysis we 398 demonstrated that its effect was negligible when compared with land use change. Anthropogenic 399 drivers of eutrophication other than land use, such as sewage or contamination by cattle were 400 discarded as the temporal trend of fecal contamination was opposite to that observed for 401 cyanobacterial abundance. Previous field studies suggest that anthropogenic factors (e.g. fertilizer, 402 gross domestic product, population) are stronger than climatic drivers (e.g. temperature, 403 rainfall)(Fang et al., 2022). Also nutrients and not temperature were identified as the key drivers for 404 cyanobacterial biomass in lakes from America (Bonilla et al., 2023).

405

406 The global trend towards the replacement of natural areas by cropping and pastures was validated 407 for the Uruguay basin. A drastic change in the type of farming and crops occurred in the new 408 millennium (Giller et al., 2021; Schlindwein et al., 2021) with a large expansion of soybean 409 production (Figure 7). These practices can trigger large phosphorous releases to water bodies, due 410 to increased phosphate fertilization and no-till practices, stratification of phosphorous content in 411 surface soil horizons and facilitation of phosphorous runoff (Carver et al., 2022). These mechanisms 412 were confirmed here with linear positive relations between total phosphorous and area of crops and 413 sown pastures, as well as between total phosphorous and cyanobacterial abundance. In future 414 analyses nitrogen inputs should also be evaluated (Paerl et al., 2016; Wurtsbaugh et al., 2019).

415

In the middle and lower Uruguay basin, soybean expansion was at expense of the deterioration area of natural grasslands previously devoted to cattle ranching, which shows lower phosphorous exports (Goyenola et al., 2020; Modernel et al., 2016; Viglizzo et al., 2001). This resulted in increased nutrient loads and a reduced capacity of natural vegetation for filtering the contaminated water runoff from land (Goyenola et al., 2020). The gain in secondary forest in the northern basin (Rosa et
al., 2021) was not able to counteract the increment in agriculture area promoting cyanobacterial
abundance and toxic blooms.

423

424 Climatic and hydrological variables affecting cyanobacteria

425 In general terms, temperature, precipitation and river flow modulate ecosystems eutrophication 426 (Meerhoff et al., 2022; Salk et al., 2022). In this study, these variables did not present a significant 427 contribution to explain major historical changes in cyanobacterial abundance in the Uruguay river. 428 Temperature values were never limiting for cyanobacterial growth (Reynolds et al., 2002; Segura et al., 2018), and blooms occurred throughout the year including the coldest months (Kruk et al., 429 430 2017). Furthermore, it has been recently shown that different toxic cyanobacterial ecotypes exhibiting different temperature optima exist (Martínez de la Escalera et al., 2022) sustaining 431 432 potential toxicity production along the entire year in the studied system.

433

434 The obtained positive relationship between precipitation and water flow is coincident with studies 435 evaluating larger periods (1900-2020) in the lower reaches of the basin (Saurral et al., 2008; Barros et al., 2015; Díaz et al., 2020; Ungerovich et al., 2021). Also, previous studies described how water 436 flow modulates cyanobacterial blooms (Mitrovic et al., 2011; Ferrari et al., 2011; O'Farrell & 437 438 Izaguirre, 2014: Adloff et al., 2018). However, our results evidenced that in the long run (yearsdecades), land use change and its effect on eutrophication were hierarchically more important than 439 440 hydrology. Hydrological variability is expected to modulate cyanobacteria at shorter time scales 441 (days-months), promoting blooms by biomass accumulation (Bowling et al., 2013) and long-range 442 transport (Kruk et al., 2021).

443

An increase in more than four orders of magnitude in the abundance of cyanobacteria took place between 1963-2000 and 2000-2020. This type of catastrophic shifts, are commonly observed in elsewhere aquatic ecosystems (Scheffer et al., 1993) and specifically in reservoirs (Yang et al.,

2017). Phytoplankton community shifts driven by human activities, particularly nutrient 447 448 enrichment, changes rapidly from species rich communities experiencing good water quality into 449 cyanobacteria dominated communities with toxin production (Scheffer et al., 1993). These drastic 450 shifts are non-linear and reversing such a process could take far longer than it takes to arrive to degraded quality ecosystems. A clear evidence of a regime shift can be traced to the extreme of the 451 452 closure of three potabilization plants (years 2005, 2006 and 2008) (OSE, 2009), the persistence of 453 blooms throughout the year, the dramatic increase in the number of health alerts from bi-national 454 authorities (especially after 2000), and the occurrence of acute human intoxications in the river and downstream (Giannuzzi et al., 2011; Vidal et al., 2017). These bloom events reached a climax in 455 456 2019, when an unprecedented large and persistent bloom of *Microcystis* reached the Río de la Plata 457 (Aubriot et al., 2020; Kruk et al., 2021).

458

459 Future scenarios and land use changes on demand

460 Land use changes are affected mainly by economical drivers (Bueno et al., 2021; Modernel et al., 461 2016; Schlindwein et al., 2021). Brazil, Argentina and Uruguay are major producers of agricultural commodities that increased substantially in the last years to supply the demand of the world market 462 (Bueno et al., 2021; Schlindwein et al., 2021). As a result, the expansions of monocultures such as 463 soybean or eucalyptus were performed in absence of adequate conservation strategies and 464 environmental impact assessments (Bueno et al., 2021; Modernel et al., 2016; Viglizzo & Frank, 465 2006). Along with this escalation in world demand of crops, warming, increased precipitation and 466 increased river flow are expected (Díaz et al., 2013; Vera et al., 2006) with synergistic effects 467 modulating water quality, pollution and biodiversity loss (Wilkinson et al., 2022; Zabel et al., 468 469 2019). Some of the areas that currently experience restrictions on cultivation will become productive regions depending on supplementary irrigation by constructing dams. Dams modulate 470 471 river discharge and sequester sediments (Bonachea et al., 2010; Depetris & Pasquini 2007a), thus favoring eutrophication and allowing reproduction and accumulation of cyanobacteria, increasing 472 473 their toxin production and diversity (Martínez de la Escalera et al., 2017, 2022) as reported for the 474 upper reaches (Debastiani Júnior et al., 2016) and at the lower stretches (O'Farrell & Izaguirre, 475 2014) of the river. Increases in extreme precipitation events, allow rapid nutrient transport from 476 soils into rivers (Ockenden et al., 2017) and then to dams, where blooms will continue to increase in 477 frequency and magnitude. Extreme events will help transport blooms to coastal and marine 478 ecosystems along with their negative effects (Alcántara et al., 2022; Kruk et al., 2021; Preece et al., 479 2017).

480

481 The first and more important step to limit the increase in cyanobacterial blooms is to reduce the 482 external anthropogenic loading of nutrients at the entire basin level as well as limiting dam construction. Otherwise, local actions will have little chance of being effective (Chorus & Welker, 483 484 2021). Major changes in agricultural practices will be required to mitigate phosphorus losses and cyanobacterial blooms under a climate change scenario (Ockenden et al., 2017) specially at the 485 486 catchment-to-coast transboundary continuum (Paerl et al., 2018; Wurtsbaugh et al., 2019). Large 487 river basins include international territories over several countries and their management requires coordinated regional actions in terms of river water use and land use policies, which are still far 488 from being implemented in the current settings (Saguier et al., 2021). As current results suggests, 489 490 the increase of surface area dedicated to high nutrient input agricultural practices and the intensification of these practices in the basin will intensify the present situation of toxic 491 492 cyanobacterial blooms in this subtropical river.

493

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Table 1. Uruguay river (UR) basin and sub-basins areas, countries, MapBiomas (MB) collections and temporal period of the information used (MapBiomas Pampa: Baeza et al. 2022, Atlantic Forest Tri-national initiatives: Souza et al., 2020). Refer to Figure 1 for visual location. Land uses and land cover were grouped into six classes: natural forest, forest plantation, grasslands and wetlands, farming (annual crops and sown pastures), non-vegetated areas (that include bare soil and urban infrastructure) and water bodies and rivers.

Basin	Area (km ²)	Country	Information origin	Temporal period	Type of information
Total UR basin	329.003	Brazil, Argentina and Uruguay	MB Brazil Collection 5, MB Atlantic Forest Collection 1 and MB Pampa Collection 1		Six classes
Upper and middle	167.130	Brazil	MB Brazil Collection 5 and 6	1985-2019	Six classes and composition of "annual crops and pasture"
Lower	161.875	Argentina and Uruguay	MB Atlantic Forest Collection 1 and MB Pampa Collection 1	2000-2019	Six classes

Table 2. Correlation between annual values of land use area from 2000 to 2019 between the lower area (LOW) and the combination of upper and middle areas (UM) of the Uruguay river basin. Land uses included were: crops and sown pastures (CP), natural forests (Nat), grasslands and wetlands (Gra), and forest plantation (For). The symbol indicates the significance level of Spearman correlation (* p <0.05, ** p <0.01).

	CP Low	CP UM	Gras Low	Gras UM	For Low	For UM	Nat Low
CP Low							
CP UM	0.73**						
Gra Low	-0.84**	-0.91**					
Gra UM	-0.72**	-0.98**	0.96**				
For Low	0.56*	0.86**	-0.91**	-0.94**			
For UM	0.67**	0.94**	-0.94**	-0.98**	0.96**		
Nat Low	-0.45*	-0.58**	0.53*	0.59**	-0.61**	-0.59**	
Nat UM	-0.54*	-0.77**	0.49*	0.68**	-0.40	-0.67**	0.52*

Table 3. List of the causal path models constructed to explain cyanobacterial abundance (logCya) sorted according to Akaike information criteria (AIC). Significant paths and their coefficients (p<0.05) are shown for each model. We recall that temperature was significant only when land uses were not included as a possible path in the model. ns: path included but with non-significant coefficients, "-": path not included in the model. CP: crops and pastures area, Tmax: maximum annual temperature, PP: precipitation, Qacum: accumulated water flow, Nat: natural class.

		paths					
Model	ΔΑΙC	CP→ logCya	Tmax→ logCya	PP→ Qacum	Qacum→ logCya	PP→ logCya	Nat→ logCya
8: all except Nat	0	0.61	ns	0.89	ns	ns	-
1: all variables	2	0.55	ns	0.89	ns	ns	ns
5: CP and Tmax	41	0.61	ns	-	-	-	-
6: only CP	41	0.70	-	-	-	-	-
4: CP and Nat	43	0.72	-	-	-	-	ns
2: all except effect of PP in Qacum	47	0.55	ns	-	ns	ns	ns
7: only Tmax	52	-	0.46	-	-	-	-
3: climatic	55	-	0.46	ns	ns	ns	-

Table 4. Coefficients estimated in the path analysis of the selected model to explain cyanobacterial abundance with climatic, land use and hydrologic variables. Their standard error (se), the z statistics (z) and the associated p values are included. The significant variable was crops and pastures area (CP in bold) and the relationship between precipitation (PP) and accumulated flow (Qacum). The remaining variables showed non-significant relationships (p>0.05). Tmax: maximum annual temperature, PP: precipitation, Qacum: accumulated water flow, Nat: natural forests class.

Variable	Coefficient	se	Z	р
PP	-0.14	0.30	-0.47	0.64
Tmax	0.23	0.16	1.39	0.17
Qacum	0.13	0.30	0.42	0.67
СР	0.55	0.22	2.32	0.02
Nat	-0.06	0.22	-0.27	0.79
PP~Qacum	0.89	0.03	9.55	< 0.001