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3	Removal of metronidazole from aqueous media by C. vulgaris
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5	Sufia Hena ^a , Leo Gutierrez ^b and Jean-Philippe Croué ^{a, c, *}
6	
7	^a Curtin Water Quality Research Centre, Department of Chemistry, Curtin University, Australia
8	^b Facultad del Mar y Medio Ambiente, Universidad del Pacifico, Ecuador
9 10	^c Institut de Chimie des Milieux et des Matériaux IC2MP UMR 7285 CNRS, Université de Poitiers, France
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21 22	* Corresponding author Email : jean-philippe.croue@curtin.edu.au (Jean-Philippe Croué)
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25 Highlights

27	1.	Metronidazole is efficiently removed during algal treatment.

- 28 2. Metronidazole is removed via sorption onto the biomass.
- 29 3. Metronidazole showed toxicity to *C. vulgaris* at stationary phase of life cycle.
- 30 4. The removal efficiency of *C. vulgaris* decreased with increase of antibiotic.

31 Abstract

This current study investigated the removal of metronidazole from aqueous media by C. 32 vulgaris. Two different initial sizes of inoculum (0.05 and 0.1mg/L) were tested for a wide 33 concentration range of metronidazole (1-50µM). The effect of metronidazole concentrations on 34 biomass production was studied for 20 days. The exopolymeric substances (EPS) were 35 quantified and correlated with the removal of antibiotics from aqueous media. Specifically, MDZ 36 37 stimulated the production of EPS in C. vulgaris, which played the major role in the adsorption of 38 this antibiotic. Also, metronidazole significantly influenced the zeta potential of C. vulgaris in the test cultures, indicating a change in surface characteristics. This decrease in surface negative 39 40 charge caused auto-flocculation phenomena at a stationary phase. Chronic and acute toxicity experiments showed that metronidazole was harmful to C. vulgaris at stationary phase. Results 41 from this study would advance our knowledge on the treatment of metronidazole-contaminated 42 43 waters with C. vulgaris as a green technology-oriented process.

44

45 Keywords: Adsorption; C. vulgaris; EC50; metronidazole

47 **1.** Introduction

The excessive use of pharmaceuticals and their improper disposal pose risks to the environment 48 and public health [1]. Especially, antibiotics (i.e., widely used for preventing bacterial diseases in 49 humans and livestock) are continuously introduced in natural aquatic environments [2, 3]. 50 Despite their low concentrations, antibiotics potentially affect the quality of drinking water, 51 human health, and ecosystems [4]. The residual antibiotics occurring in aquatic environments 52 53 would lead to the emergence of resistance among targeted and non-targeted strains of 54 microorganisms [5-7]. Also, surface waters contaminated with antibiotics expand the variety of resistant microorganisms and the transfer of genetic material between multidrug-resistant 55 56 bacteria [8, 9].

Metronidazole is a first-generation derivative of 5-nitroimidazole drugs developed for 57 therapeutic uses [10, 11]. Metronidazole is an effective prescription medication used for treating 58 59 several infectious diseases caused by anaerobic bacteria and protozoa in humans and livestock [12, 13], showing minimal side effects and a broad spectrum of applications [14]. In general, 60 antibiotics have a relatively short half-life in open environments (e.g., water bodies or soils) [2]; 61 however, metronidazole is highly persistent in nature due to its high solubility [15] and low 62 biodegradability [16]. Also, the removal of metronidazole from water and wastewater during the 63 treatment train is not efficient [17]. Deblonde et al., [18], reported a mere 38.9% removal of 64 metronidazole from a wastewater treatment plant. Nevertheless, because of its high toxicity and 65 mutagenic effect on targeted and non-targeted microorganisms in the environment [12], it is 66 highly necessary to develop efficient protocols and (bio) technologies to remove this antibiotic 67 68 from aqueous media.

Several methods have been reported for the removal of metronidazole from aqueous media. 69 Briefly, chemical processes such as reduction [19], Fenton and photo-Fenton process [20], UV-70 photolysis [21], and physical treatments (e.g., adsorption) [22, 23] have been mainly tested. 71 However, high energy and chemical consumptions, as well as the generation of secondary waste 72 are the main constraints for their successful (economic) implementation [24, 25]. On the other 73 hand, Rivera-Utrilla et al., [22] had reported that microorganisms in wastewater treatment plants 74 75 cannot efficiently degrade metronidazole. Activated sludge process is the most common 76 biological treatment method to remove most of the drugs from aqueous media; however, this process fails to efficiently remove most of the antibiotics [26]. In general, antibiotics are not 77 78 readily biodegradable [27]. Antibiotics are mainly removed by sorption onto activated sludges in wastewater treatment plants [4, 28]. However, antibiotics desorb from the activated sludge to the 79 aqueous media during its reuse in aeration tanks to maintain a fixed concentration of activated 80 81 microorganisms [29]. According to Grandclement et al., [30] wastewater treatment plants using activated sludge are not specifically designed to remove antibiotics. 82 83 Simultaneously, the activated sludge in wastewater treatment plants promotes the occurrence of

antibiotic-resistant bacteria due to the exposure of bacteria to antibiotics present in wastewater [31]. Therefore, the efficient treatment of antibiotics present in wastewater to ensure biosafety for humans and their environment is a difficult task [32].

The use of microalgae as green technology is a viable option to remove antibiotics from water. Moreover, microalgal cultures mitigate atmospheric CO₂ while the biomass can be used for biofuel feedstocks. Briefly, microalgae are known for efficiently removing xenobiotic compounds from water, including a range of pharmaceuticals. Garcia-Rodríguez et al., [33] studied the role of microalga *Spyrogira* sp. in photolysis degradation and the contribution of light

92 in the total removal of six antibiotics from the sulfonamides and tetracyclines families. Du et al., [34] investigated the response of green algae Chlorella pyrenoidosa towards the removal of 93 ceftazidime antibiotic. Guo and Chen [2] showed the removal of cephalosporins from 94 wastewater using Chlorella pyrenoidosa. Conversely, Magalhães et al., [16] unsuccessfully 95 attempted to remove metronidazole from water using Microcystis protocystis (cyanobacteria). 96 From these previous studies, chlorophytes have clearly shown more resistant to stress conditions 97 98 during water treatment. Chlorella sp. is a chlorophyte which has been extensively used for the removal of antibiotics such as 7-amino cephalosporanic acid [32], cephalosporins [2], 99 norfloxacin [35], levofloxacin [36], and enrofloxacin [37]. However, to the best of our 100 101 knowledge, metronidazole removal using C. vulgaris has not yet been reported.

This current study has investigated the metronidazole removal capacity of C. vulgaris and the 102 dominant mechanisms involved in the removal. The impact of metronidazole at different 103 104 concentrations on the growth rate of C. vulgaris was examined. Additionally, the tolerance and chronic toxicity (EC50) of C. vulgaris to this antibiotic as a function of its stage of life cycle was 105 106 investigated. Finally, the exo-polymeric substances (EPS) production by C. vulgaris during exposure to metronidazole, its influence on the surface properties of C. vulgaris, and its role 107 during the removal of this antibiotic were comprehensively studied. Results from this study will 108 advance our knowledge of the treatment of metronidazole-contaminated waters with C. vulgaris. 109 110 The protocols followed in the current investigation can be extended to the treatment of different antibiotics-contaminated aquifers using green technology-oriented processes. Simultaneously, a 111 dual-approach can be used in the system to treat wastewater and to produce valuable biomass as 112 feedstock for biofuel, through anaerobic digestion. This latter process would aim to completely 113 degrade any antibiotic residue [61]. 114

2. Materials and Methods

116 2.1. Chemical reagents

Metronidazole (HPLC grade; >99% purity) was obtained from Sigma-Aldrich. All the chemicals,
solvents, and reagents used were of analytical grade. A stock solution of 1mM metronidazole
(MDZ) was prepared with 30% methanol and 70% MilliQ water (v/v) and stored in a dark
airtight glass vial at 4°C for no longer than 30 days for further experiments.

121

122 2.2. Algae Culturing and batch experiments

123 *Chlorella vulgaris* (*C. vulgaris*) was obtained from CSIRO Marine Research Hobart, Tasmania. 124 *C. vulgaris* was re-cultured in 2 L conical flasks with a working volume of 1L in BG11 media 125 (Table 1), at $25\pm4^{\circ}$ C. The cultures were exposed to fluorescent lamps to provide the irradiance of 126 µmol photon /m²/s until the stationary phase was attained.

127 Two sets of batch experiments were conducted. The freshly prepared culture media were 128 inoculated with two different sizes of inoculum of *C. vulgaris* for obtaining 0.05 g/L and 0.5 g 129 L^{-1} biomass (i.e., dry biomass vs optical density calibration curve) in 3 L of Erlenmeyer flasks 130 with 2 L of working volume. The cultures were spiked with increasing concentration of MDZ (1-131 50 μ M) except for the control cultures of each batch. All experiments were conducted in 132 triplicate.

133

134 2.3. Determination of biomass and chlorophyll-a

The determination of the biomass of the test and control cultures was conducted every day by measuring the optical density at 680 nm as described in [38]. A batch study was conducted to evaluate the specific growth rate using the following formula; 138 $\mu = (\text{In } N_2 - \text{In } N_1) / (t_2 - t_1)$

139 Where, N_1 and N_2 are biomass concentrations (mg L⁻¹) at day t_1 and t_2 , respectively, during the 140 exponential phase of growth. All experiments were conducted in triplicate.

The extraction of chlorophyll-a and its quantification was conducted as reported by Xiong et al., [39]. A 10 mL of *C. vulgaris* was centrifuged at 4500 rpm for 10 min. The supernatant was discarded and the pellet was collected. The pellet was re-suspended in 10 mL of methanol:water (9:1; v/v) and incubated in 20 mL centrifuge tubes for 10 min. The incubated culture was centrifuged again at 4500 rpm for 10 min. The absorbance of the supernatant was measured at 665 nm and 652 nm and the chlorophyll-a concentration of the extract was calculated using the following equation:

148 Chlorophyll-a ($\mu g L^{-1}$) = 16.29 × A₆₆₅ - 8.54 × A₆₅₂

Where, A₆₆₅ and A₆₅₂ are the absorbance of the supernatant at wavelength 665 nm and 652 nm,
respectively.

151

152 2.4. *Toxicity of MDZ for* C. vulgaris

The toxicity of MDZ was calculated as the Effective Concentration (EC50) of the stressor 153 (MDZ) at which 50% of the growth rate of C. vulgaris was inhibited. The inhibition in growth 154 was calculated as a function of chlorophyll-a content. To determine the chronic toxicity of MDZ, 155 a batch experiment was conducted at different concentrations of MDZ (65-90 µM) with an initial 156 biomass concentration of 0.1 g L^{-1} . The microalgae were exposed to antibiotic for 240 hours (10 157 days). However, for acute toxicity experiments, a 0.1 g L⁻¹ culture at stationary phase (20 days 158 old culture) was spiked with 1-50 µM antibiotics for 96 hours. The growth rate was calculated in 159 160 terms of chlorophyll-a ($\mu g L^{-1}$).

161 2.5. Extraction and analysis of Exo-polymeric substances (EPS)

Samples of 50 mL of C. vulgaris cultures at late exponential phase (i.e., which started 162 approximately from day 10, Fig 1) were kept in the fridge at 4°C to settle the cells. The 163 supernatants were decanted and used for EPS analysis, while the settled cells were re-suspended 164 in 50 mL of MilliQ water in a beaker. The culture was stirred for 2 hours at 1000 rpm using a 165 magnetic stirrer to extract the EPS from the cell wall of C. vulgaris. Then, the cells were 166 167 separated by centrifugation at 1000 rpm for 10 minutes. The second supernatants were collected 168 for analysis of bound EPS. The pellets of the centrifuged biomass were dried at 60°C. The EPS were extracted at every alternate day starting from day 10 until day 20. 169

170 30 mL of all supernatants were used for measuring the dry weight of EPS. Every supernatant of 171 the tests and control cultures of age 10 days and beyond were used for EPS determination. 30 mL 172 of supernatants were dried in an oven at 60°C until a stable weight was achieved to calculate the 173 ratio of the dry weight of EPS to biomass.

174

175 2.6. Zeta potential

The Zeta (ζ) potentials of *C. vulgaris* from the test (low and high inoculum spiked with 5 μ M MDZ) and control cultures were analyzed on every alternate day from day 0 till day 20 using a Zetasizer (Malvern). 10mL of cultures were collected from the Erlenmeyer flasks. The biomass was allowed to settle down. The supernatants were discarded and biomass was re-suspended in MilliQ water. The suspended *C. vulgaris* was pipetted in a folded cuvette. The zeta potential was measured 5 times and the average and standard deviation were calculated.

182

183 2.7. Analysis of MDZ

184 Samples of 5 mL from test bioreactors were collected and filtered by a syringe filter (0.45 µm) and analyzed using an HPLC equipped with UV/VIS detector (Agilent 1100 series, USA) at 320 185 nm. A 250 mm*4.6 mm C18 5-µm reverse phase HPLC column (AlltimaTM, GRACE) was used 186 and the mobile phase consisted of 0.1% phosphoric acid solution and methanol (80:20 v/v) with 187 a 1 mL min⁻¹ flow rate. To analyze the removal mechanism and removal efficiency of MDZ via 188 C. vulgaris, flocculated biomasses of microalgae were collected from the bottom of a bioreactor 189 190 (i.e., a random selection was conducted from a 10 µM MDZ high biomass bioreactor on day 20) 191 in triplicate. The biomasses were washed thrice with 5 mL MilliQ to remove the unbounded molecules of MDZ from biomass using filter paper (GF/C; 1.2 µm) and the filtrates were 192 193 collected in HPLC vial for analysis after every wash. The absence of MDZ peak after third wash approved the adequacy of three washes of collected biomasses for further steps of analysis. The 194 washed biomasses were collected on glass Petri dishes separately and vacuum dried at 60°C in 195 196 temp controlled oven for 24 hours till they achieved constant weight. 0.5 mg of dried biomass was suspended into 20 mL of ethyl acetate followed by sonication for 15 minutes and 197 centrifuging at 12,000 rpm for 10 minutes [16]. The biomass was separated from ethyl acetate 198 using a syringe filter (GF/C; 1.2 µm). The filtrate was vacuum dried at 40°C, followed by 199 diluting the remaining residue in 20 mL fresh culture media. The final diluted residue (i.e., 200 expecting desorbed MDZ in solution) was filtered with a 1.2 µm GF/C syringe filter and 201 collected in HPLC vial for analysis. The results were used to elucidate the MDZ removal 202 efficiency of *C. vulgaris* and the mechanism involved during the removal process. 203

204

205 **3. Results and discussion:**

206 3.1. *Effects of MDZ on the growth of* C. vulgaris:

Two sizes of inoculum (0.05 and 0.5 g L^{-1}) were selected to study the effect of MDZ on C. 207 *vulgaris* concentration in solution. These results were compared with those of control culture. As 208 depicted in Fig 1a-b (Refer to Table S1a-b for standard deviations), the growth of the cultures 209 can be divided into three phases. The early exponential phase was calculated for 10 days from 210 the beginning of the culture due to the highest growth rate during this period for all tests and 211 212 control culture regardless of inoculum size. Interestingly, the specific growth rates of C. vulgaris during the early exponential phase were almost similar for cultures with the same inoculum size 213 but different concentrations of MDZ (Table 2). Clearly, there was no impact of MDZ on C. 214 215 vulgaris during this early exponential phase. Remarkably, there was no removal of MDZ from 216 any of the test bioreactors during the early exponential phase (Figure 2a-b, Table S2a-b). During the late exponential phase (following the first 10 days of growth), the biomass increased with 217 218 similar specific growth rate for the same inoculum size. These results indicate that MDZ does not 219 impact the specific growth rate of the culture. However, MDZ hampered the duration of the late exponential growth phase, where MDZ decreased the surface charge of the cells causing them to 220 flocculate and settling down (Fig 3). 221

The duration of the late exponential phases was different in different test bioreactors. The longest 222 duration was observed for control cultures as 8 days; nevertheless, the control culture of high 223 224 inoculum attained a lower specific growth rate than control culture and low inoculum (Table 2). 225 The test cultures with low inoculum remained in late exponential phase for 5-6 days, while high 226 inoculum test cultures showed the shortest duration of 2-3 days. The longer late exponential phase for low inoculum cultures might be due to: 1) a lower concentration of microalgal cells 227 present in the media than higher inoculum bioreactors for the same amount of nutrients, and 2) 228 late auto-flocculation in low inoculum cultures. The late auto-flocculation in low inoculum test 229

230 bioreactors caused biomass to remain in suspension media for a longer time and hence increased the duration of the late exponential phase. The concentrations of MDZ start decreasing during 231 late exponential phase in all cases regardless of inoculum sizes (0.05 and 0.5 g L^{-1}) or initial 232 concentrations (1-50 µM) of MDZ as shown in Fig 2a-b. These results confirmed that MDZ 233 played a major role in auto-flocculation as further explained in next section. In all test 234 bioreactors, the stationary phase did not last for more than 24 hours, which was then followed by 235 a quick decline in biomass content. The reason for this rapid decrease in suspended biomass was 236 due to auto-flocculation; however, the removal of MDZ continued even during the decline in 237 biomass phase. 238

239

240 3.2. *Effect of MDZ on zeta potential of* C. vulgaris:

241 Fig 3 revealed the electronegative zeta potential of C. vulgaris from control culture and test 242 bioreactor spiked with 5 µM MDZ (i.e., which was randomly selected as a representative case). In both cases, the electronegative zeta potential decreased with the age of the culture. 243 Throughout the growth cycle of the microalgae, the concentration of the functional groups 244 present on the cell membrane and cell wall changes and influences the surface properties of the 245 cells [40]. However, during the active growth phase the lower intercellular interaction caused a 246 247 higher net electronegative zeta potential [41]. At late exponential growth phase, the metabolism 248 of the cells and the mobility decreased. Simultaneously, the changes on the cell surface caused 249 intercellular interactions which resulted in a lower electronegative zeta potential [41]. Xia et al., [42] reported that the decrease in the electronegative zeta potential of the C. vulgaris was also 250 due to the increase in the effective (hydrodynamic) size of the cells (i.e., cell with bound EPS) at 251

a later stage of its life cycle. Decho and Gutierrez [43] considered EPS as an extended part of thecell.

Fig 3 showed that the electronegative zeta potential of the *C. vulgaris* decreased more significantly in the test bioreactors than their respective control cultures. Among the test bioreactors, the cultures with low inoculum size showed a higher electronegative zeta potential than the high inoculum size culture at day 12 and beyond. The high inoculum culture contained more biomass, which attained a stationary phase earlier than low inoculum cultures. Since the EPS forms at late exponential or stationary phase of life cycle [43], high inoculum cultures start forming EPS earlier than the low inoculum culture (Table 3).

261 As mentioned earlier, the net negative zeta potential of the microalgal cell decreased due to change in functional groups present on the cell wall, which would be directly associated with the 262 metabolic conditions of the cells [40]. However, in the current study, the additional decrease in 263 264 zeta potential compared to their control cultures could be due to the charge dispersion interaction of MDZ with algal cells (discussed in section 3.4) as well as the increase in the effective size of 265 the cell. The effective size of the cells would be larger in test cultures compared to their 266 respective control cultures due to the formation of bound EPS onto the cells in test cultures 267 (Table 3). 268

A drop in zeta potential in the test cultures indicated a decrease in the repulsive electrostatic forces between the cells which led to auto-flocculation. The auto-flocculation takes place when van der Waals forces overcome the repulsive electrostatic forces [44]. In this study, the repulsive electrostatic forces were most likely decreased due to the adsorption of EPS around the cells and also because of its interaction with MDZ in its scaffolds.

275 3.3. Effect of MDZ on the EPS production of C. vulgaris

EPS is produced by microalgae at a later stage of life cycle [43]. Microalgae produce EPS for cell protection as a waste of metabolic processes [45]. There are two types of EPS produced by *C. vulgaris*; a) EPS soluble (free) in media, and b) loosely bound (adsorbed) EPS which assists in auto-flocculation of cells [46]. EPS is generally discussed as a three-dimensional architecture scaffold providing space for trapping foreign compounds and slowing the diffusion rate, physical stability of cells, digestion of xenobiotic substances, and communication between cells (i.e., quorum sensing) [43].

Control culture produced more soluble (free) EPS than bound (adsorbed) EPS (Table 3). This 283 284 result explains why control culture cells did not undergo auto-flocculation. However, in test bioreactors the auto-flocculation phenomenon was common. A slight increase in bound EPS was 285 286 observed with an increase of MDZ concentration in both test cultures (low and high inoculum). The bioreactor spiked with 1 μ M MDZ produced significantly less EPS (47.7 ± 1.23 mg g⁻¹ DW 287 of C. vulgaris) than that with 5 μ M (151.6 ±3.45 mg g⁻¹ DW of C. vulgaris), which indicated 288 that MDZ stimulated the EPS production in C. vulgaris. However, concentrations higher than 5 289 μ M MDZ produced almost the same amount of EPS, which varied from 151.6-145.5 mg g⁻¹ dry 290 weight (DW) and 139.3-146.1 mg g^{-1} DW for low and high biomass bioreactors, respectively. 291 These results revealed that 5 µM MDZ was the optimum concentration of antibiotics to produce 292 maximum EPS for this specific system. 293

294

295 3.4. Removal of MDZ

MDZ was analyzed from day 0 until day 20 in test bioreactors. The removal of MDZ by *C*. *vulgaris* from aqueous media under different conditions is shown in Figure 2a-b. MDZ was not

298 removed until the early exponential growth phase. This result proved that MDZ did not adsorb 299 onto the cell surface of C. vulgaris possibly due to the low log K_{ow} (-2.3) value of MDZ. Also, the pH of the C. vulgaris culture varied between 9-10, where MDZ is expected to behave as a 300 neutral molecule [47]. Therefore, electrostatic interactions between MDZ and negatively-charged 301 C. vulgaris would not be expected. However, π -electron of the heterocyclic aromatic ring of the 302 MDZ molecule would interact with different sites of EPS (Figure 4). EPS are highly hydrated 303 304 complex polymers mainly composed of polysaccharides, proteins, and DNA [48]. The CH group 305 of the polysaccharides form CH- π bond with the π -electrons of aromatic compounds [49, 50]. Consequently, the adsorption of MDZ molecules in the scaffold of the EPS matrix would involve 306 307 van der Waals interaction between π -electron of the heterocyclic aromatic ring of the MDZ molecule and the hydrogen atom of CH group of the polysaccharides to form a CH- π bond. 308 Another possible mechanism would involve $OH-\pi$ bond with OH groups of saccharides. 309 310 However, EPS has a complex structure where the OH groups of polysaccharides involved in intermolecular interactions, might have no free OH to make bonds with MDZ [49]. Nishio [51] 311 has reported that saccharides are overall polar in nature but include some hydrophobic moieties 312 depending upon the orientation of OH groups, which have an affinity to form hydrogen bonds 313 with hydrophobic compounds [49]. The hydrophobic imidazole ring in MDZ would form $OH-\pi$ 314 bond with OH groups of hydrophobic moieties. Previous studies have reported that MDZ 315 interacts with its π electron of heterocyclic aromatic ring to form π - π dispersion bonds with other 316 aromatic compounds [22]. Protein containing aromatic amino acids would be expected to favor 317 the formation of π - π dispersion bonds for MDZ to interact with EPS. Sanvordeker et al., [52] 318 confirmed that the alkyl side chain of MDZ participates in the binding to proteins. Besides, EPS 319

320 contains a diverse range of functional groups, including amine, carboxylate, phosphate, carbonyl,321 amide, and hydroxyl which can form H-bonds with MDZ.

The removal of MDZ started after day 10 in all cases regardless of MDZ or biomass 322 concentration. Remarkably, MDZ removal and EPS formation occurred almost simultaneously. 323 These results indicated that the MDZ might be removed by EPS solely by adsorption 324 mechanism. Neither metabolites nor antimicrobial biotransformation products were observed in 325 326 HPLC chromatogram; suggesting that the EPS of C. vulgaris might not contain any metabolizing 327 enzyme for MDZ. Therefore, it is postulated that the decrease in MDZ concentration in the bioreactors would only be associated with adsorption phenomena, which eventually played a role 328 329 in increasing the concentration of EPS. As a control experiment, bound EPS was extracted from 10 µM high biomass bioreactor, sonicated for 15 minutes, followed by filtration and analysis, 330 where the presence of MDZ was clearly recorded. A bioreactor has a complex system of 331 332 suspended and flocculated biomass; thus, achieving an accurate mass balance of MDZ was not possible. However, these results showed that 0.69 mg MDZ adsorbed per 100 mg of dry weight 333 of EPS (i.e., corresponding to 59% of initial DMZ) in a 10 µM MDZ-high biomass bioreactor 334 after 20 days of incubation. The small molecular size of MDZ would favor its adsorption into the 335 scaffold of EPS, while the concentration gradient would play a major role in the diffusion of 336 MDZ into EPS [53]. 337

A comparison between the MDZ removal of low and high biomass-containing bioreactor revealed that the longer the late exponential growth phase, the lower the removal efficiency. Specifically, a longer late exponential growth phase delayed the EPS production. In the case of 5 μ M MDZ bioreactors, the low biomass reactor was in active phase until day 15, while the active or growth phase was over on day 12 in the high biomass reactor. Taking into account both biomass and EPS, it was clear that EPS initiated the adsorption; however, the higher the biomass,the better was the removal.

345 3.5.Effect of initial concentrations of MDZ in culture media

Wastewaters from different geographic locations contain different concentrations of 346 pharmaceuticals. Consequently, it is important to examine the MDZ removal from bioreactor as 347 a function of the initial concentration of antibiotic. Two sets of bioreactors with lower and higher 348 349 biomass concentrations were spiked with different initial concentrations (1-50 µM) of MDZ as 350 shown in Figure 5. In general, the efficiency of bioreactors decreased with an increase in the initial concentration of MDZ for a fixed period of time. The adsorption phenomena in live 351 352 biological adsorbent are complex processes, especially when the adsorbate had to stimulate the production of adsorbent. As depicted in Fig.5, only 50.80% of MDZ was removed in a low 353 biomass reactor initially containing 1 µM of MDZ. However, MDZ was completely removed in 354 355 both low as well as high biomass reactors originally containing 5 µM MDZ in 20 and 18 days of experiments, respectively. Nevertheless, concentrations higher than 5 µM of MDZ showed the 356 same trend as the higher the initial concentration of antibiotic, the lower the removal (%) under 357 constant conditions (same inoculum size and incubation period). 358

It is evident from Table 3 that the EPS production increases with increasing concentration of MDZ in culture; however, this becomes almost constant above 5 μ M MDZ. Probably, this was the reason why 1 μ M MDZ in low inoculum size test culture was not able to completely remove MDZ, whereas a 5 μ M MDZ in low as well as in high inoculum size test cultures were capable to completely remove MDZ from culture media.

364

365 *3.6. Mechanism of MDZ removal*

366 The removal of MDZ from the bioreactor was confirmed by analyzing the liquid phase using HPLC, where a decrease in concentration was recorded. The absence of new chromatographic 367 peaks during the course of the experiments and the respective analysis validated the hypothesis 368 that C. vulgaris might not participate in any biodegradation mechanism. To confirm the 369 mechanism, ethyl acetate was used to desorb the adsorbed MDZ from the flocculated biomass. 370 The sole presence of an MDZ peak from the desorbed sample confirmed that MDZ was only 371 372 removed by sorption mechanism. These results showed that 0.93 mg of MDZ was adsorbed in 1 g dry weight of C. vulgaris. However, comparing this result with those obtained for MDZ 373 removal per 1 g of dry weight of EPS for the same test bioreactor on the same day (i.e. 0.69 mg 374 375 MDZ/100 mg of dry weight of EPS in 10 µM MDZ high biomass bioreactor after 20 days of incubation; as mentioned in section 3.4) the amount recovered from dry weight of C. vulgaris 376 showed a 9.13% loss in mass balance of MDZ. This decrease in the amount of MDZ suggested 377 378 its diffusion into the microalgal cell.

379 It is a general observation that dead microalgae cells have a higher adsorption capacity than live

cells [54]. Peng et al., [55] demonstrated that live cells of freshwater algae: S. obliquus and C.

381 *pyrenoidosa* adsorbed less progesterone and norgestrel in aqueous media than their dead

- 382 biomass. Yu et al., [56] explained that the better adsorption capacity of dead biomass was due to
- 383 loss of permeability control when the cell membrane structures were destroyed.
- In this study, the adsorption capacity of live cells with EPS has been compared with live cells

without EPS. The first 10 days of culture (Figure 2 (a) and (b)) were considered as cells without

- 386 EPS while after day 10 cells started to form EPS (especially, attached or bonded) which were
- 387 considered as live cells with EPS. It is clear from Figure 2 (a) and (b) that there was no removal
- of MDZ during the initial 10 days (live cells without EPS). The removal of MDZ started with the

- 389 EPS formation (i.e. after day 10 in all test bioreactors), which proved that live cells with EPS
 390 played a major role in MDZ removal.
- 391 However, to confirm if suspended, attached, or both types of EPS were participating in removing
- 392 MDZ from water, the removal of MDZ from water was compared against the EPSs (suspended,
- attached, and total EPS) obtained from 1 μ M MDZ test bioreactors (i.e., selected as a representative case) of low and high inoculum (Figure 6). As depicted in Figure 6, it was observed that the attached EPS showed the best fit (i.e., expressed by the highest R² value) compared to the other two EPSs (R² of suspended EPS=0.49, and R² of total EPS=0.88). This high value of correlation coefficient (R² = 0.95) of the attached EPS indicated a dominant contribution of MDZ removal and good consistency between the 3D scaffold of the attached EPS
- 399 and the adsorbed antibiotic.
- 400
- 401 3.7. *EC50 of* C. vulgaris *for MDZ*

The most common toxicity assay is to calculate the effective concentration (EC50) of a stressor 402 (i.e., metronidazole) at which 50% of the growth rate of an organism (microalgae) is inhibited. 403 Photosynthesis is an essential process for the survival of microalgae, which is related to growth 404 and several other metabolic activities. Thus, any disruption in the growth of microalgae during 405 stressors exposure is related to the changes in chlorophyll (Fig. 7a-b). Antibiotics damage 406 thylakoid and chloroplast lipids as they disrupt membranes and other organelles in cells; thus, 407 destroying the photosystem II (PSII) and consequently chlorophylls and carotenoids [57]. Low 408 409 chlorophyll content, as a stress response to pharmaceuticals, has been previously observed in microalgae Tetraselmis chui, Isochrysis galbana, and C. pyrenoidosa in the presence of 410 thiamphenicol, florfenicol, and chloramphenicol [58]. 411

412 Two approaches were followed to study the growth inhibition in C. vulgaris by MDZ; chronic 413 and acute toxicity of MDZ. The chronic toxicity assay was conducted to assess its adverse effects on an organism as the result of long term exposure, while the acute toxicity is performed for 414 short term exposure. The chronic toxicity of MDZ on C. vulgaris was only observed when the 415 antibiotic is added during its growth phase, while acute toxicity was visible when the antibiotic 416 came in contact with microalgal cells at its stationary state. The results showed that the chronic 417 418 toxicity EC50 for C. vulgaris was 83.64 µM (Figure 8a), while for acute toxicity it was 13.63 419 µM (Figure 8b). The growth inhibition of C. vulgaris exposed to increasing concentrations of MDZ (i.e., as measured by the chlorophyll-a yield) is shown in Figure 7a-b for chronic and acute 420 421 toxicity, respectively.

According to Commission of the European Community (1996), pollutants with an EC50>100 mg L⁻¹ fall under the category of harmless substances, while those with EC50 from 10-100 mg L⁻¹ are included under the harmful category. Also, substances with EC50 between 1-10 mg L⁻¹, and EC50<1 mg L⁻¹ are considered as very toxic. Results from this study indicate that MDZ is harmful to *C. vulgaris* (Figures 8a-b).

Besides the efficiency of *C. vulgaris* for removing DMZ from wastewater, the biomass generated during this treatment would be a valuable feedstock for producing biofuels. Algal biomass has been commonly used as a soil conditioner, plant fertilizer, and animal feed [30]. However, the harvested biomass of *C. vulgaris* would not be recommended as a fertilizer because of its MDZ content. A scheme for biofuel production using MDZ-mediated water is proposed in Figure 9. Rawat et al., [59] have previously demonstrated the production of biogas using algal biomass through anaerobic digestion. The temperatures of these processes vary from 350-800°C [60].

- 434 Remarkably, MDZ completely decomposes at 180°C [61], which is significantly lower than the
- 435 abovementioned lowest temperature of anaerobic digestion processes.
- 436 **4.** Conclusions

In the current study, C. vulgaris was used as a novel treatment to remove metronidazole from 437 aqueous media. These results indicated that MDZ stimulated the production of EPS in C. 438 vulgaris, which played a major role in the adsorption of this antibiotic. C. vulgaris displayed an 439 440 excellent removal capacity for the target antibiotic at 5µM concentration within the range of 0.05 to $0.5g L^{-1}$ inoculum of C. vulgaris. These results showed that MDZ was only removed by 441 adsorption mechanism onto the scaffold of the EPS, which slowly diffused into the cells. The 442 removal efficiency decreased with increasing initial concentration of antibiotics. The tolerance of 443 C. vulgaris towards MDZ depended upon the stage of the life cycle, and on the exposure time to 444 445 the antibiotic. However, MDZ is marked as a harmful pollutant to C. vulgaris. Nevertheless, the recovery of biofuel from the harvested biomass through anaerobic digestion shows promising for 446 future biofuel applications. 447

- 448 **Competing interest**
- 449 The authors declare no competing interest.

450 **References:**

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Figure 1a: Growth curve of low inoculum size of *C. vulgaris* for control and test cultures at different concentrations (1-50 µM) of MDZ

- \succ The data are the mean of three replicates.
- Refer Supplementary Table 1a for the data and SD



Figure 1b: Growth curve of high inoculum size of *C. vulgaris* for control and test cultures at different concentrations (1-50 µM) of MDZ

- \blacktriangleright The data are the mean of three replicates.
- ▶ Refer Supplementary Table 1b for the data and SD



Figure 2a: Evolution of MDZ concentration during treatment using low inoculum size of *C*. *vulgaris* containing different concentration (1-50 μ M) of MDZ in aqueous media

- \blacktriangleright The data are the mean of three replicates.
- Refer Supplementary Table 2a for the data and SD



Figure 2b: Evolution of MDZ concentration during treatment using high inoculum size of *C*. *vulgaris* containing different concentration (1-50 μ M) of MDZ in aqueous media

- \succ The data are the mean of three replicates.
- Refer Supplementary Table 2a for the data and SD



Figure 3: Zeta potential of low and high inoculum size of *C. vulgaris* in control culture and culture spiked with 5 μ M MDZ on alternate days



H bond between saccharide and MDZ



 π - π dispersion bond between aromatic compound of protein (tryptophan) and MDZ



CH- π bond between saccharide and MDZ

Figure 4: Possible bonding between MDZ and components of EPS



Figure 5: Effect of initial MDZ concentration on MDZ removal



Figure 6: Correlations between the amount of attached, suspended and total EPS with MDZ removal from $1 \mu M$ test bioreactors with low and high inoculum from day 10 to day20



Figure 7a: Chlorophyll a in *C. vulgaris* during a 240-hour metronidazole chronic toxicity test. The control culture (culture without MDZ) possessed 160.22 μ g L⁻¹ after 240 hours



Figure 7b: Chlorophyll a in *C. vulgaris* during a 96-hour metronidazole acute toxicity test. The control culture (culture without MDZ) possessed 240.1 µg L⁻¹ after 96 hours



Figure 8a: Growth inhibition (%) of *C. vulgaris* by MDZ during chronic toxicity test.



Figure 8b: Growth inhibition (%) of C. vulgaris by MDZ during chronic toxicity test



Figure 9: Proposed scheme for biogas production using MDZ-containing wastewater.

Table 1: Compositions of BG-11 media in 1L MilliQ

Chemicals	Weight (mg)
K ₂ HPO ₄	75
KH ₂ PO ₄	175
MgSO4.7H ₂ O	75
NaNO ₃	250
CaCl ₂ .2H ₂ O	25
NaCl	25
Na ₂ EDTA.2H ₂ O	50
FeSO ₄ 7H ₂ O in 0.1mL H ₂ SO ₄	4.9
H ₃ BO ₃	11.42
ZnSO ₄ .7H ₂ O	1.412
MnCl ₂ . 4H ₂ O	0.232
CuSO ₄ .5H ₂ O	0.252
Co(NO ₃) ₂ .6H ₂ O	0.08
Na ₂ MoO ₄ .2H ₂ O	0.192

Table 2: Specific growth rate of *C. vulgaris* in control and test culture with different concentration of MDZ

Concentration of	Low inoculum siz	ze	High inoculum size			
MDZ						
	Specific growth rate of early exponential phase (d ⁻¹)	Specific growth rate of late exponential phase (d ⁻¹)	Specific growth rate of early exponential phase (d ⁻¹)	Specific growth rate of late exponential phase (d ⁻¹)		
Control (without MDZ)	0.279	0.055	0.095	0.036		
1µM	0.271	0.065	0.094	0.034		
5μΜ	0.273	0.060	0.096	0.029		
10 µM	0.272	0.058	0.097	0.019		
25 µM	0.271	0.046	0.093	0.035		
50 µM	0.271	0.053	0.091	0.028		

Dava	Days Control culture		Low inoculum bioreactors									
Days			liture 1µM		5 µM		10 µM		25 µM		50 µM	
	EPS (S)*	EPS (A)*	EPS (S)	EPS (A)	EPS (S)	EPS (A)	EPS (S)	EPS (A)	EPS (S)	EPS (A)	EPS (S)	EPS (A)
10	2.31±0.11	1.45±0.09	1.62±0.33	3.01±0.08	1.71±0.08	5.54±0.16	1.76±0.63	5.12±0.53	1.97±0.16	5.62±0.11	1.88±0.09	6.32±1.66
12	7.40±0.26	7.92±0.22	2.89±0.78	8.84±0.65	2.19±0.22	15.33±1.11	2.11±0.08	15.54±0.38	2.21±0.36	15.78±0.76	2.08±0.23	16.34±1.89
14	20.52±0.46	14.91±0.28	6.79±0.54	18.77±0.54	4.23±0.23	37.87±1.29	4.19±0.43	37.1±0.97	4.56±0.69	38.2±1.06	4.21±0.25	38.33±1.27
16	34.82±0.25	19.28±0.78	11.19±1.09	26.54±0.43	7.88±0.18	76.45±2.66	7.12±0.65	77.15±1.86	7.19±0.46	75.29±1.64	6.89±0.61	73.32±2.76
18	42.95±1.67	30.65±0.64	21.32±1.68	38.41±0.79	20.65±	113.62±3.54	19.9±1.06	111.3±1.69	14.06±0.87	121.23±1.39	18.88±0.28	110.12±3.18
20	48.76±1.86	37.50±2.09	39.98±1.87	47.72±1.23	33.67±1.26	146.1±2.34	31.04±0.38	142.32±2.21	24.76±1.11	148.99±2.43	30.03±0.53	139.31±5.16
						High inoculum	bioreactors	H				
10	4.27±0.22	2.11±0.13	3.12±0.08	9.32±0.65	3.25±0.11	12.25±1.07	3.29±0.38	12.11±1.09	3.41±0.09	12.43±1.66	3.52±0.33	13.11±0.87
12	9.16±0.31	8.99±0.76	4.89±0.18	20.36±0.23	4.22±0.18	26.48±1.76	4.15±0.79	25.54±1.24	4.55±0.97	26.98±1.45	4.23±0.27	27.23±1.56
14	29.12±1.55	21.34±1.56	8.77±0.54	49.42±1.43	6.78±0.16	55.15±1.18	7.12±0.68	57.75±1.76	7.67±0.76	56.67±2.12	7.45±0.87	58.43±0.98
16	37.45±1.42	34.77±1.82	13.23±0.21	59.23±1.54	11.32±0.64	107.92±2.45	12.34±0.45	78.92±2.06	12.65±1.14	76.78±2.32	10.54±1.09	76.67±1.67
18	48.72±2.64	46.34±2.44	23.45±0.67	86.43±2.69	14.43±0.98	121.4±2.27	13.98±0.88	119.12±2.46	21.33±1.12	110.43±2.34	13.78±0.95	118.34±2.65
20	54.66±1.98	51.86±1.87	39.18±2.11	110.9±2.45	26.76±1.17	151.6±3.45	25.08±1.11	148.78±2.65	32.65±1.45	140.01±4.11	24.44±1.22	145.55±3.56

Table 3: EPS production in control culture and test bioreactors

(S)*—Suspended EPS; (A)*--Attached EPS; EPS were measured in mg g⁻¹ dry weight of *C. vulgaris*

Graphical Abstract:



Schematic diagram of MDZ removal in Chlorella vulgaris culture