# Development of an Ecotoxicological Test with a Soil Microalgae Species

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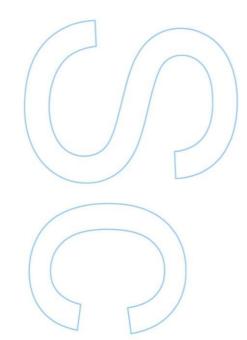
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O Presidente do Júri,



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# Abstract

Microalgae form an important component of soil microflora, accounting for up to 27% of the total soil microbial community. This group of soil organisms has important soil functions as for example: i) to contribute for soil formation, being one of the first colonizers of bedrock and, for improving soil structure; ii) to promote organic carbon sequestration in soils and, iii) to maintain soils fertility. Despite, the recognized importance of microalgae to soil functions and soil quality, this group of organisms is not considered in the battery of species used to assess the risks of new or existing contaminants to soil biota. Taking this into account, a microalga species was isolated from biological soil crusts, collected in a Portuguese soil in the Center region of the country, and was genetically identified as *Micractinium inermum*, a Chlorophyta species with a worldwide distribution. This species was then used to pursue the main objective of this work, *the development of an ecotoxicological test procedure with a soil microalga, using soil as a test substrate*.

To meet this main objective, the following specific tasks were accomplished: 1) considering that soil is a semi-aquatic habitat, the adequacy of the standard protocol for growth inhibition tests with freshwater microalgae and cyanobacteria (OECD 201) was tested for *M. inermum*. Copper (II), used as Copper (II) Sulfate Anhydrous (CuSO<sub>4</sub>) and an herbicide, a soluble concentrate formulation containing 28.85% of glyphosate (GLY), the Roundup Max® were selected as reference substances, given the availability of toxicity data for these two compounds. With this task it was also intended to compare the sensitivity of the soil species with that of freshwater species commonly used in ecotoxicological tests. 2) a more ecologically relevant test methodology, using the artificial OECD soil, was developed.

Three independent tests were performed for each reference substances both following the existing standard protocol and the one developed in this study, after obtaining growth curves for the species, both in aqueous  $BG_{11}$  culture medium and in OECD artificial soil. Growth curves were useful for establishing the exposure durations (5 and 13 days, in aqueous and soil media, respectively) and the validity criteria of the tests (an increase of 14 and 100 times, in cell's density and in chlorophyll content of the soils, in the CTLs, respectively).

The EC<sub>50</sub> values recorded and the corresponding 95% confidence intervals obtained for both contaminants on the aqueous tests and on soil tests proved the repeatability of the test procedures as they fit in the same range of values. The results recorded, following the OECD 201 standard protocol, showed that *M. inermum* was more sensitive than freshwater green microalgae, reinforcing the importance of including tests with species from the soil, to prevent an underestimation of the risks to the overall soil biota. Additionally, *M. inermum* displayed a similar or a lower sensitivity to the reference substances tested when exposed to them in the artificial soil. These results, highlight the role of soil components in the bioavailability of contaminants, and subsequently the importance of this test procedure, for assessing the risks of contaminants, in more realistic exposure scenarios, to prevent overestimation of risks.

KeyWords: soil microalga, *Micractinium inermum*, reference substances, new ecotoxicological test procedure

## Resumo

As microalgas formam uma componente importante da microflora do solo, representando até 27% da comunidade microbiana total do solo. Este grupo de organismos do solo possui funções importantes no solo, como por exemplo: (i) contribuir para a formação do solo, sendo um dos primeiros colonizadores do leito de rocha, melhorando a sua estrutura; ii) promover o sequestro de carbono orgânico nos solos e iii) manter a fertilidade do solo. Apesar da importância do papel das microalgas nas funções do solo e na qualidade do solo, este grupo de organismos não é considerado no conjunto de espécies usadas para avaliar os riscos de contaminantes novos ou existentes para a biota do solo. Levando isso em consideração, espécies de microalgas foram isoladas de crostas biológicas do solo, recolhidas num solo português na região Centro do país, e foram geneticamente identificadas como *Micractinium inermum*, uma espécie Chlorophyta com distribuição mundial. Esta espécie foi então utilizada para acalçar o objetivo principal deste trabalho, ou seja, o *desenvolvimento de um procedimento de teste ecotoxicológico com uma microalga de solo, usando solo como substrato de teste.* 

Para atender a este objetivo principal, foram realizadas as seguintes tarefas específicas: 1) considerando que o solo é um habitat semi-aquático, a adaptação do protocolo padrão para testes de inibição de crescimento com microalgas de água doce e cianobactérias (OECD 201) foi testada para *M. inermum.* Cobre (II), utilizado como sulfato de cobre (II) anidro (CuSO<sub>4</sub>) e um herbicida, formulação concentrada solúvel contendo 28,85% de glifosato (GLY), o Roundup Max®, foram selecionados como substâncias de referência, dada a disponibilidade de dados de toxicidade para estes dois compostos. Com esta tarefa também se pretendeu comparar a sensibilidade das espécies do solo com as espécies de água doce comumente usadas em testes ecotoxicológicos. 2) Como segunda tarefa, foi desenvolvida uma metodologia de teste ecologicamente mais relevante, usando o solo artificial da OCDE.

Três testes independentes foram realizados para cada substância de referência, tanto seguindo o protocolo padrão existente como o desenvolvido neste estudo, após obtenção de curvas de crescimento para a espécie, tanto em meio aquoso BG<sub>11</sub> como em solo artificial da OCDE. Curvas de crescimento foram úteis para estabelecer as durações de exposição (5 e 13 dias, em meio aquoso e solo, respetivamente) e os critérios de validação

dos testes (aumento de 14 e 100 vezes, na densidade celular e no teor de clorofila dos solos, nos CTLs, respetivamente).

Os valores de  $EC_{50}$  registados e os correspondentes intervalos de confiança a 95% obtidos para ambos os contaminantes nos testes aquosos e nos testes de solo, provaram a reprodutibilidade dos procedimentos de teste, uma vez que se encaixam na mesma gama de valores. Os resultados registados, seguindo o protocolo padrão da OECD 201, mostraram que *M. inermum* foi mais sensível que as microalgas verdes de água doce, reforçando a importância da inclusão de testes com espécies do solo, para evitar uma subestimação dos riscos à biota total do solo. Além disso, *M. inermum* apresentou uma sensibilidade semelhante ou menor para as substâncias de referência testadas quando expostas a elas no solo artificial. Isto demonstrou o papel dos componentes do solo na biodisponibilidade de contaminantes e, subsequentemente, a importância deste procedimento de teste, para avaliar os riscos de contaminantes, em cenários de exposição mais realistas, para evitar a sobrestimação dos riscos.

Palavras-chave: microalga de solo, *Micractinium inermum*, substâncias de referência, novo procedimento de teste ecotoxicológico

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# 1. Introduction

## 1.1. Terrestrial microalgae

Algae are a polyphyletic group of organisms found in a wide variety of habitats, usually in damp places or in aquatic environments (Hoffmann, 1989; Shimmel et al. 1985) and play an important role in the equilibrium of aquatic ecosystems, being key players in primary productivity and biogeochemical cycles (Fayez et al. 2007, Wang et al. 2018). Although most green algae typically occur in aquatic environments, many species also live in terrestrial habitats (also considered as semi-aquatic habitats), representing a major component of the microbial flora, occurring on exposed aero-terrestrial interface zones (Holzinger et al. 2013, Rindi, 2011), where their ecological importance is remarkable. Mostly microscopic, the green algae benefit from their small size which gives them the ability to grow in diversified habitats (Lewis et al. 2004a), playing an important role in environmental carbon sequestration and being responsible for 50% of the total photosynthesis on the Earth (Renuka et al. 2018). Light is the major factor affecting algal growth, supplying the energy to support its metabolism (Nam et al. 2015a) and consequently its productivity (Simionato et al. 2013).

The term algae include all photosynthetic eukaryotes included in several Phylum of the Plantae kingdom, as for example Chlorophyta and Charophyta Phyla to which belong the green algae (Ruggiero et al. 2015)<sup>12</sup>. The colonization of terrestrial habitats by descendants of Streptophyta algae started about 470-450 million years ago and was one of the most important occurrences in the evolution of life on Earth (Becker et al. 2009).

The first compilation of a species list, including soil algae, was presented by Graebner (1895), describing communities of Northern Germany (Metting, 1981). Green algae compared to other algal groups have proven to be particularly prepared for colonizing terrestrial habitats (Gaysina et al. 2013) and based on molecular and morphological evidence, terrestrial algae derived from many independent terrestrial colonization events by different lineages of aquatic algae, both marine, and freshwater, evolving actively even in environments considered too hostile to support their existence (Lam et al. 2010). Therefore,

<sup>&</sup>lt;sup>1</sup> <u>http://algaevision.myspecies.info/node/3510</u>, Last accessed on: 07/12/18

<sup>&</sup>lt;sup>2</sup> <u>http://www.algaebase.org/browse/taxonomy/?id=142046</u>, Last accessed on: 7/12/18

there is a wide variety of algae taxa recognized as terrestrial algae (Hoffmann, 1989; Lund, 1962; Shields, 1962) such as, *Chlamydomonas* (Pröschold et al. 2007; Pentecost 2002), *Chlorococcum infusion* (Bold, 1930; Chae et al. 2016), *Chlorella* (Luo et al. 2010; Bock et al. 2011), *Scenedesmus* sp. (Lewis et al. 2004b; Bahar et al. 2013), *Klebsormidium* sp. (Škaloud, 2014; Ryšánek et al. 2016), *Desmococcus* sp. (Paul, 2006; Broady, 1993) *Geminella terricola* renamed as *Interfilum* sp. (Petersen, 1932; Graham et al. 2000; Mikhailyuk et al. 2008) and recently found in soil, the species of the genera *Micractinium* (Luo et al. 2010).

Green algae occur in almost every habitat, including the most extreme ones (Starks et al. 1981) such as: i) extreme arid hot deserts (Kitzing et al. 2015; Fletcher et al. 2008; Lewis et al. 2004b); ii) polar regions (Ryšánek et al. 2016; Davydov, 2016; Holm-Hansen, 1964; Broady, 1996; Broady, 1993), where often are the only primary producers (Hoffmann, 1989); iii) mountain areas and rocks (Johansen et al. 2007; Bischoff, 1963); iv) tropical regions (Lam, 2010) and v) airborne (Sharma et al. 2007). In extreme habitats the occurrence of desiccation conditions, low nutrients availability, extreme temperatures, and UV radiation are common features, therefore prevent the loss of water and keep cellular homeostasis under stress is vital (Lin et al. 2014; Holzinger et al. 2016). In these conditions of great abiotic stress, terrestrial algae developed morphological and physiological adaptations and occupied sheltered microhabitats in which conditions are less severe (Hoffmann, 1989). Terrestrial algae also have the capacity to disperse in the air, resulting in different dispersion patterns and indicating that algae are likely to have a global distribution (Sharma et al. 2007).

Microalgae are an important component of the soil environment and perform several important functions. In nature, most microalgae are found in association with other aerobic or anaerobic microorganisms. In this symbiosis, the molecular oxygen from algal photosynthesis is used as an electron acceptor by bacteria to degrade organic matter (Subashchandrabose et al. 2011). These organisms contribute to soil formation as primary colonizers (Davey et al. 1993) and generate energy and matter fluxes, having a significant impact in soil carbon pool by contributing to the organic content of soils (Subashchandrabose et al. 2013). The existing information about the mechanisms used by terrestrial algae to prevent cellular damage is still scarce but it is known that terrestrial algae are able to survive in a wide temperature range and have developed several mechanisms to prevent and/or counteract the damage caused by UV radiation which is known to cause

inhibition and several types of damage in their photosynthetic apparatus, with the production of mycosporine-like amino acids (MAAs) and reactive oxygen species (ROS) (Rindi, 2011; Lin et al 2014). The production of mucilage and accumulation of osmotic active compounds, the maintenance of homeostasis during dehydration and regulation of water status are the most common anti-desiccation strategies (Holzinger, 2013; Holzinger, 2016). Proline, SOD (superoxide dismutase) and carotenoids may be some of the molecular components involved in the tolerance to desiccation in green algae (Lin et al., 2014) and also in ROS elimination. Other strategies to avoid stress are the ability to aggregate cells to form biofilms of several layers and the decrease of dispersion by aggregation with secondary colonizers (lichens, mosses) and soil particles. The outer layers are fully exposed to the environment and susceptible to damage and at the same time they effectively protect the cells underneath by their water-holding and by their filtering capacity of the excess radiation on the surface. Without the radiation "umbrella" offered by the pigmented species in the outer layer, the species in the inner layer would experiment high mortality (Sharma et al. 2007; Holzinger et al. 2016; Belnap, 2003).

Due to their high capacity for morphological and physiological adaptations to different environments, as previously mentioned, algae often act as pioneer microorganisms in terrestrial habitats (Lin et al. 2013), playing an important role for the establishment of other trophic levels (Nam et al. 2015b). Soil algae excrete growth-promoting substances and biologically active compounds such as enzymes and ions that can affect other components of soil communities, including plants (Hastings et al. 2014). These photosynthetic active microorganisms form an intimate association with the soil particles that exist within or just above the upper millimeters of the soil (Bérard et al. 2004) and with the existing communities, contributing to the formation and integrity of soils (Renuka et al. 2018). In natural conditions, filamentous soil algae and cyanobacteria like *Klebsormidium* sp., *Lynbya* sp., *Nostoc* sp., form multiple layer structures, because of their sticky mucilaginous sheaths, that glue soil particles together forming biological soil crusts (BSC) (Schulz et al. 2016).

Biological soil crusts are a great help in resisting desertification, boosting desert soil formation and changing nutrient circulation (Wu et al. 2013). Algae and cyanobacteria, two main components of BSC, play a critical role in the soil succession and contribute with organic matter towards soil formation, favoring conditions for further colonization by plants, lichens, mosses and vascular plants (Zhao et al. 2009). Generally, cyanobacteria colonize the soil surface during the early stages and green algae, lichen and moss crusts establish

over time (Zhang et al. 2017a). Thus, the degree of algae distribution in the soil profile varies and their role at particular stages of soil development and in different soils can be different (Rahmonov et al. 2007). Knowledge on the diversity and distribution of terrestrial microalgae in BSC still falls behind of that for marine and freshwater environments. The terrestrial environment is still one of the less studied biological resources (Arguelles et al. 2017).

The study of terrestrial algae is still a challenge since ecological parameters such as population density, biomass, and productivity are often difficult to evaluate (Hoffman, 1989; Kabirov et al. 2009). The current classification of green microalgae is characterized by the compromise between the concept of conventional and modern phylogenetic species (Gustavs et al. 2011). Modern classification systems are polyphasic incorporating not only morphological and ultrastructural features but also molecular genetics in species identification (Wojciechowski et al. 2013). As a result, the molecular phylogeny of green algae expanded rapidly as molecular approaches were applied in their taxonomic identification (Lewis et al. 2004a). Even though the majority of the recently described terrestrial green algae have been isolated from various extreme habitats, since are more attractive for current biodiversity research, terrestrial microalgae from temperate regions in Europe have been described using the polyphasic approach (Barcyte et al. 2017).

## 1.2. Terrestrial Ecotoxicology

Terrestrial ecosystems are known to provide a complex range of essential ecosystem services regulated by soils (Chagnon et al. 2015). The soil is one of the fundamental components for supporting life on Earth and the processes that occur within soil drives the ecosystem and global functions (Jeffery et al. 2010). Over the years soil has had multiple definitions, according to the Soil Science Society of America, from a soil scientist's perspective, soil is the layer of generally loose mineral and organic material that is affected by physical, chemical and/or biological processes at or near the planetary surface and usually holds liquids, gases, and biota. The soil ecosystem also performs numerous functions and services, ranging from providing food to filtering the water being also used as a platform for human activities, a potential source of vital products, such as antibiotics and the habitat of a great biodiversity of species (Es van, H., 2017; Jeffery et al. 2010). Thus, the contamination of this resource has become a global environmental, economic and

societal problem (Nam et al. 2015b). The degradation of soils due to various anthropogenic stress factors is alarming (Filser et al. 2008). Depending on the habitat and mobility, terrestrial organisms may be more or less exposed to chemicals present in the topsoil layer (van Gestel, 2012).

The study of the toxic effects that chemical substances have on soil populations and communities, as well as the analysis of their multiple interactions in ecosystems, are the focus of the multidisciplinary field of ecotoxicology and has gained a fundamental role in the regulation of existing and new chemical substances and in the assessment and management of environmental risks (Expósito et al. 2017). The degree of contamination is frequently evaluated by comparing the measured concentrations in the bulk soil with specific threshold values. Toxicity assays appeared as a complementary and a valuable tool for evaluating the hazard of chemical substances on the viability, growth, and reproduction of cells and organisms (Arouja et al 2011), as only organisms reflect the available fraction of contaminants in soils. In this context the development of more rapid and sensitive assays, become particularly important for evaluating the potential impact of contaminants on soil biota, aiming to establish threshold levels of toxicants to protect natural communities (Sabatini et al. 2009). With this purpose several standard protocols were developed for testing chemicals with different soil species, assessing both acute and chronic effects, namely with: i) soil invertebrates, using oligochaetes [guidelines OECD 207, OECD 222, ISO 11268 (OECD, 1984; OECD, 2016; ISO, 2012)], collembolans [guidelines OECD 232, and ISO, 11267 (OECD, 2016; ISO, 2014)] and enchytraeids [OECD 220 and ISO 16387 (OECD, 2016; ISO 2014)]; ii) terrestrial plants [OECD 227, OECD 208, ISO 22030, ISO 11269 (OECD, 2006; ISO 2005; ISO 2012)] and, iii) microorganisms and microbial processes [ guidelines OECD 216; OECD 217; ISO 16072, ISO 17155; ISO 14238 (OECD, 2000; OECD, 2000; ISO, 2002; ISO 2012; ISO 2012)]. Despite all the available protocols, Van Gestel (2012), concluded that risk assessment processes would benefit from an extension of the available battery of toxicity tests for soils and by paying more attention to exposure, bioavailability, and toxicokinetics of contaminants, to better understand the longterm consequences of chemical exposure at an individual, population and community level. Further, it has been reinforced that apart from measuring the relevant parameters and meeting the environmental requirements, effective toxicity tests should be quick, simple and reproducible.

Algal toxicity tests are extensively applied to assess the effects of hazardous substances, especially of those intentionally applied in the environment, like pesticides, which may attain aquatic compartments, because of their sensitivity and ecological relevance in the food webs. The endpoint currently assessed in these tests is the inhibition of growth of an algal population, exposed to aqueous solutions of contaminants or to aqueous environmental matrices, under continuous light conditions for 72h, following the standard protocols OECD 201 or ISO 8692 (OECD, 2011; ISO 2012). Algae growth inhibition tests are also used to test soil elutriates or soil lixiviates, to assess the ability of this environmental matrix to retain contaminants preventing their mobility to the water resources (Antunes et al. 2010). However, until now, soil algae species, have never been proposed as possible test species on available guidelines to increase the representativeness of soil biota, in evaluations targeting the risks to this compartment. Further, more ecological relevant test methodologies, for soil algae have never been proposed and are needed given the importance of this group of soil organisms.

Soil algae are distributed in nearly all terrestrial environments, both on and beneath the soil surface (Nam et al. 2015b; Metting, 1981). Since it is impossible to cover species of all relevant groups of organisms for each environmental compartment, at least appropriate indicator species have to be identified for risk assessment purposes (Filser et al. 2008) as well as multiple relevant endpoints (Nam et al. 2016) to be measured on these bioindicators, that could reflect the mechanisms of toxicity of contaminants, as well as the vulnerability of the test species. Terrestrial habitats are poorly represented in toxicity testing and therefore, microalgae species isolated from soil would be another important representative group of the terrestrial ecosystems and can be potentially used (Dominguez-Morueco et al. 2014).

The sensitivity of microalgae to toxic substances is species-dependent (Expósito et al. 2017) and may vary by orders of magnitude (Levy et al. 2007) so there is a great degree of uncertainty in extrapolations between species of algae species. Thus the best approach is to incorporate a high number of species, representative of all major taxonomic groups when the aim is to protect the overall ecosystem (Lewis, 1995).

The terrestrial algae are a relatively neglected group among soil microbiota and there is limited information regarding the assessment of the impacts of toxic substances on this biological group (Bérard et al. 2004; Nydahl et al. 2015; Chae et al. 2016; Subaschandrabose et al. 2015). An interesting study was conducted by Nam et al. (2016), who suggested the paper-disc soil method as an easy-to-use approach. In this method, a

flat-bottom 24-well microplate was filled in with LUFA 2.2 soil that was contaminated with different concentrations of copper. A paper disc was placed in each well and pressed to adhere to the soil surface. The green microalgae Chlorococcum infusionum in the exponential phase was inoculated on each paper disc and incubated for 6 days. The results revealed that the copper inhibited the growth zone of C. infusionum, compared to the control, showing an EC<sub>50</sub> value of 148 mg/kg. However, it was argued that the paper disc might act as a boundary layer between the contaminated soil and the soil algae, decreasing the bioavailability of the toxic through the binding of particles to the filter paper. This means that in direct contact with contaminated soil, soil microalgae tend to become more exposed to the toxicants, so a new approach should be conducted applying the soil algae directly to the contaminated soil in a given exposure time period, assessing the effects on the growth rate of algae through the chlorophyll content analysis. In this context, reference contaminants are used to assess the repeatability and reliability of test organism responses and of test procedures over a specific period of time, and further, they are used in inter-laboratory testing to judge comparability of results. Reference contaminants should also be used when new organisms or protocols are introduced in order to establish an expected dose-response relationship for a range of concentrations (Blaise et al. 2005, Orr et al. 1990). Like copper, other reference contaminants can be used, such as 3,5-dichlorophenol and potassium dichromate suggested in the OECD 201 guideline for testing chemicals with freshwater algae and cyanobacteria (OECD, 2006). Reference contaminants should be selected within the most widely used and concerning contaminants, for which a large set of data is already available

Several metals are essential micronutrients for all living organisms and play essential roles in fundamental biochemical processes that sustain life on Earth. Metals are persistent elements that cannot be destroyed or degraded, as they exist as natural constituents of the Earth's crust (Huertas et al. 2014; Chen et al. 2016). However, with the increase of metal emissions from anthropogenic activities, related with their extraction and with their intentional use (e.g. copper is a widely used fungicide in agriculture) a large proportion of these emissions are accumulated in soils and sediments due to their high sorptive capacity (Arouja et al. 2011; Machado et al. 2015), where they attain toxic levels to biota, in several areas. For this reason, metals are one of the groups of contaminants more deeply assessed in the literature (Debelius et al 2009).

Microalgae exhibit high metal binding capacities due to the presence of polysaccharides, proteins or lipids on the surface of their cell walls which contain functional groups that can act as binding sites for metals (Priya et al. 2014; Yu et al. 1999). In the last few years, some studies have been carried out with different species of microalgae isolated from soil and different contaminants. Bahar et al. (2012) and Subramaniyam et al. (2016), for example, evaluated the toxicity of arsenite and iron in microalgae isolated from uncontaminated soils, determining the EC<sub>50</sub> values for both contaminants. Subashchandrabose et al. (2015), evaluated the toxicity of cadmium using the microalgae, Chlorococcum sp., isolated from a contaminated soil concluding that this species was less sensitive, with an EC<sub>50</sub> of 2.85 mg/L, when compared with the same species isolated from an uncontaminated site, with an EC<sub>50</sub> value of 0.41 mg/L (Krishnamurti et al. 2004), indicating the ability of this species in developing tolerance to toxicants. The unique properties of green microalgae, such as high sensitivity and repeatability along with the need to increase the available taxa for use in toxicity bioassays, especially in terrestrial habitats, make them a suitable choice for the development of standardized bioassays of toxicity (Dominguez-Morueco et al. 2014). Copper (mainly in the form of copper sulfate), is a widely used fungicide in agriculture, contaminating not only soils but also the aquatic environment, posing a direct risk of toxic effects on the ecosystems (Kungolos et al. 2008). It is an essential micronutrient as it acts as a cofactor for a number of enzymes (Chen et al. 2016), but it can also be toxic at concentrations as low as 1 µg/L (Levy et al. 2007). At high concentrations Cu becomes toxic, inhibiting the growth of algae due to adverse effects on the same cellular processes for which it has an essential role, as enzymes activity and photosynthetic electron transport (Leal et al. 2016). In plants, at the cellular level, copper plays an essential role in signaling transcription and protein trafficking machinery and for a healthy plant growth, it needs to be acquired from the soil and transported throughout the plant. In higher concentrations, copper becomes toxic to the plant affecting the photosynthetic machinery, being the photosynthetic reaction center II (PSII) more sensitive to copper toxicity than the PSI, by inhibiting the oxygen formation. The tolerance to toxic levels might be related to potential mechanisms at the cellular level involved in detoxification (Yruela et al. 2005). In green algae, the process is the same, copper inhibits photosynthesis including mainly light reactions thus damaging the photosynthetic organisms (Chen et al. 2016). A study by Afkar et al. (2010), indicated that copper toxicity at high concentrations may be due to the oxidative potential of copper that causes reduction of chlorophyll,

decreases oxygen evolution rates and causes depletion of ATP by inhibition of enzymes involved in the cellular metabolism. A similar study indicated that for several metals, such as copper, the induction of ROS was observed for two different microalgae, to neutralize the oxidative damage induced by copper stress (Hamed et al. 2017). Copper is also one of the most studied metals in what regards their phytotoxic effects. Wang et al. (2018), for example, evaluated the photosynthetic and biochemical responses of *Closterium ehrenbergii* to copper as CuSO<sub>4</sub> and CuCL<sub>2</sub>, determining the EC<sub>50</sub> values of 0.202 mg/L and 0.245 mg/L, respectively. The data obtained showed a considerable decrease in pigment levels and photosynthetic efficiency, while inducing the generation of ROS in cells with increased exposure time, indicating a decrease in algal growth rate and cells density. Taking this and many other studies into account, data is available for comparing the sensitivity between species, and between species from different compartments.

The common practice to use copper in multiple anthropogenic activities such as mining, agriculture, several industries is affecting its concentrations in soils, which are attaining concerning levels. At the European Union levels, there is no common agreement on the copper threshold for the definition of risk but some legislation and several studies propose that a copper threshold value of 100 mg/kg should be assumed (Ballabio et al. 2018). In Portugal, the same level depends on soil pH, but the same threshold was proposed, for soils with pH between 5.5-7.0, by the law by decree 276/2009 (MAOTDR, 2009). However, Canadian and USEPA guideline values are lower, and Caetano et al. (2015) by using a Portuguese natural soil, for the derivation of a soil screening value for copper, proposed a limit of 31.8 mg/kg. Thus, in a study conducted by Panagos et al. (2018), using the Land Use Cover Area survey (LUCAS Topsoil), containing soil samples from 27 countries of European Union, collected between 2009-2012, showed that Cyprus is estimated to have the highest mean copper concentration in topsoil (53.41 mg/kg) and, in other regions such, Malta, west Greece, east Bulgaria, Catalonia and Algarve, the mean copper concentration is between 30-50 mg/kg. The data also showed that in Portugal, there was one site with a copper concentration higher than 100 mg/kg, being France the country with more sites surpassing this threshold. Although the application rate is not known, the model used by Panagos et al. (2018), resulted in a prediction of an average consumption of copper of 8.1 kg/ha for permanent crops, which is higher than the permit value (a maximum of 6.4 kg of Cu/ha). While countries such as Australia and the United States still have no

restrictions, this has led to some European governments to set limits to copper usage. The Netherlands, for example, has banned copper use (Mackie et al. 2012).

Herbicides account for about 40% of the pesticide amounts used worldwide (Sihtme et al. 2013) and their widespread application led to its accumulation in soils affecting aquatic (Ermis et al. 2009) and terrestrial biota, from microorganisms to plants and animals (Hackenberger et al. 2018). The effects of pesticides in soils can range from acute and chronic toxicity in organisms to many sub-lethal and indirect effects on behavior, functional roles, predator-prey relationships, and food web dynamics (Chagnon et al. 2015).

Glyphosate (GLY) under the trade name Roundup® was introduced in the market by Monsanto Company during the 1970's (Pérez et al. 2011) and is the major herbicide used worldwide, being also used increasingly because approximately 80% of commercially grown genetically modified plants are tolerant to GLY (Nicolas et al. 2016). It is a systemic and nonselective herbicide that inhibits the activity of ESPS (5-enolpyruvyl shikimate-3-phosphate) synthase enzyme belonging to the shikimic acid metabolic pathway in plants resulting in the inhibition of the synthesis of some aminoacids and with subsequent growth inhibition (Sáenz et al. 1997). Apart from plants, the shikimate pathway is also present in fungi and bacteria, however, this does not necessarily mean that GLY operates as in plants.

The toxic effects of GLY have been studied in different aquatic organisms, by studying endpoints as growth, the content of pigments in tissues and photosynthesis parameters (Romero et al. 2011). Since algae is a sensitive taxonomic group to herbicides (Eugenio et al. 2018), the exposure of several freshwater algae species to glyphosate, for either 72h or 96h, provided a wide range of species-specific  $EC_{50}$  values, ranging from 3.5 mg/L to 55.9 mg/L (Annett et al. 2014), representative of the great variability in species sensitivity. Soil algae may be also endangered by herbicides since they are found on the surface of the soil and have a high potential for sorption and contact with herbicides (Wegener et al. 1985). Algae are known to be comparatively sensitive to many chemicals and the inclusion of these organisms in the test batteries has been shown to improve the capacity to predict the most sensitive ecosystem responses (Vendrell et al. 2009, Ermis et al. 2009). Tsui et al. (2003), concluded that algae, which have metabolic pathway similar to higher plants were, therefore, more susceptible to the herbicidal effect of IPA (isopropylamine) salt of GLY than non-photosynthetic organisms.

A study conducted by Nicolas et al. (2016), evaluated the toxicity of a commercial formulation containing GLY as an active ingredient on a soil filamentous fungus, *Aspergillus* 

*nidulans.* The data indicated that GLY was toxic to this species at low doses, far below the recommended agricultural application rates. However, the authors also noted that the residues presently found in soils are much lower than the tested concentrations. (Nicolas et al. 2016). In another study, Zabaloy et al. (2012), evaluated the potential effects of GLY treatments on microbial community structure and function in soils from two sites in Argentina, with the application of two different doses of GLY (0.15 and 150 mg/kg) incubated for 7 days. The results indicated that a single exposure of soils to GLY caused only minor changes to microbial community structure and function. However, microbial respiration in uncontaminated soils increased in response to GLY applications, reflecting a possible stress response of most sensitive species. Chronically exposed soils did not show this response, most likely due to the gradual elimination of most GLY sensitive species.

Another issue related with the use of GLY is the toxicity of aminomethlphosphonic acid (AMPA), one of the glyphosate's main metabolites, which has been classified as persistent in soils, raising concern regarding the widespread use of this herbicide in agriculture and forestry. A study conducted by Dominguéz et al. (2016), evaluated the effect of AMPA on mortality and reproduction of the earthworm species *Eisenia andrei* using the standard available protocol. The results showed that AMPA had no significant effects on mortality except at the highest concentration (2500  $\mu$ g/kg) tested in the chronic assay. The data, however, concluded that juveniles from parents exposed to contaminated soils had a reduced growth, something that can also limit their beneficial roles in key ecosystem functions.

Since the use of this herbicide is increasing its accumulation on soils, it is important to address its impacts on sensible soil communities, such as of microalgae, which have important soil functions. Effect-related research, which has addressed pesticides and herbicides in a rather constant proportion of published papers for more than 20 years does not reflect the actual number of active ingredients applied in the United States and Europe (Köhler et al. 2013). In European soils the information on the effect of herbicides is fragmented and outdated and more studies with non-target species are urgently needed to support the authorization of these compounds based on ecologically relevant information, to better protect ecosystems.

# 2. Aim of the thesis

Currently, soil microalgae are not taken into account in the battery of standard ecotoxicological assays, for assessing the risks of new and existing contaminants to soil biota. Thus, it is of utmost importance to develop and validate a new test procedure that can potentially be standardized in the future by competent organizations, and this was the main goal of this thesis. To accomplish this main objective, the following specific objectives were pursued:

- To perform the genetic identification of one terrestrial microalgae species isolated from soil biological crusts (SBC);
- To perceive if the standard guideline OECD 201 (OECD, 2006) for testing the effects of chemical substances in freshwater algae and cyanobacteria can be used for testing terrestrial microalgae and cyanobacteria and to depict the adaptations required;
- To perceive if the soil microalgae species is more sensitive than the freshwater species, thus supporting the relevance of performing ecotoxicological tests with soils species;
- To develop a new test procedure for exposing the soil microalgae to chemical substances, copper (Cu) in the form of Copper (II) Sulfate Anhydrous (CuSO<sub>4</sub>) and the herbicide glyphosate (GLY), in particular of its commercial formulation RoundUp ULTRA Max<sup>®</sup>, which contains 360 g/L of GLY or 28.85% (p/p) in the artificial OECD soil, to improve the ecological relevance and thus increase the realism of these tests for predicting the risks to soil communities.

# 3. Material and Methods

### 3.1. Microalgae isolation and culture conditions

Biological soil crust samples were collected from the surface of the soil (geographical coordinates 40°26'37.15"N/ 8°26'33.03"W). The samples were placed in a 50.0 mL falcon tube and taken to the laboratory. In the laboratory, using the binocular magnifier (Motic), some of the greener areas of the collected BSC samples were selected and were grown in sterilized Erlenmeyer's with 20.0 mL of a liquid medium, appropriate for the growth of photosynthetic microorganisms, the BG<sub>11</sub> medium (Rippka et al. 1992), whose composition can be found in Annex A. The working cultures were placed in an orbital shaker at 100 rpm, at 24 ± 2 °C, photoperiod of 16h<sup>L</sup>:8h<sup>D</sup> hours and a light intensity of 65  $\mu$ mol m<sup>-2</sup>s<sup>-1</sup> provided by 30W cool-white fluorescent lamps. Isolation and purification were performed by serial dilutions of liquid cultures and by applying the streak plating method using agar BG<sub>11</sub> medium (with 1.5% agar) (Temraleeva et al. 2016), previously autoclaved at 121 °C, 1 bar and for 30 min, to isolate different microalgae species. The plates were incubated at  $24 \pm 2$ °C, for 2 to 3 weeks and the morphology of the colonies was inspected under optical microscopy. From the different isolates, one green microalgae species was selected for being used as test species in soil ecotoxicological tests. Thus, therein in this section, only the work performed with this species will be described.

Liquid cultures of the isolated microalgae were prepared in 50.0 mL sterile Erlenmeyer flasks containing BG<sub>11</sub> medium with air-permeable stoppers covered with aluminum foil, by inoculating a single colony removed from the plates with a loop, and by dipping and shaking the loop in the medium. Cultures were incubated in an orbital shaker under the same conditions described above for 15 days (figure 1). All the procedures were carried out under aseptic conditions in a laminar flow chamber previously sterilized with ultraviolet light for 15 minutes.

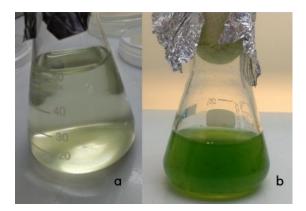


Figure 1 - Inoculum of the green microalgae from the BSC, day 1 (a) and after 15-days of growth (b).

### 3.2. Genetic Identification of the BSC selected microalgae

### 3.2.1. DNA extraction

The Maxwell® 16 System (Promega Corporation, USA) was used to extract the genomic DNA, following the protocols provided by the manufacturer and the extracted DNA was stored at -20°C until PCR amplification.

### 3.2.2. PCR amplification

To identify the isolated species genomic DNA was subjected to PCR amplification using a set of primers designed for the partial amplification of the 18S and ITS regions (Internal Transcribed Spacer) of rDNA from algae as shown in figure 2. The two pairs of primers used for PCR amplification of the rRNA gene are described in table 1. All reactions were carried out using a Bio-Rad MJ Mini<sup>™</sup> Thermal Cycler. For each PCR reaction, the final volume was 20 µL, containing 10 µg of DNA template, 0.4 µM of each forward and reverse primers and 10 µL of NZYTaq 2x Green Master Mix (NZYtech, Lisbon, Portugal). The volume was completed with ultrapure water. The PCR thermal cycling conditions were as follows: initialization at 95°C for 5 min, followed by 30 cycles of denaturation at 94°C for 30s, annealing at 52°C for 30s, elongation at 72°C for 2 min and a final extension at 72°C for 7

min, followed by hold step at 4°C. The PCR products were checked on 0.8% (p/v) agarose gel electrophoresis following the protocol described by Maniatis et al. (1982) using 1x TAE buffer [Tris-acetate (0.04M); EDTA (0.001M)] (Sambrook and Russell, 2001) with 0.75  $\mu$ L of Green Safe (NZYtech, Lisbon, Portugal) to allow the visualization of DNA bands under UV light. NZYDNA Ladder VIII (NZYtech, Lisbon, Portugal) was used as molecular size markers.

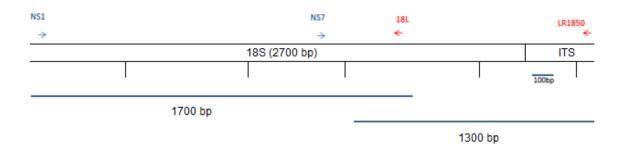


Figure 2 - Representative scheme of the 18S rDNA, the hybridization sites of the primer pairs used for PCR amplification and the expected size of the amplified fragments.

Table 1 – Sequences of the specific primers for the amplification of the 18S rRNA gene and the corresponding size of the amplified fragments.							
Primers	Sequence (5' to 3')	Size (bp)	Reference				
NS7m (Forward)	5'-GGCAATAACAAGTCTGT-3'	1300	Mikhailyouk et al. 2008				
LR1850 (Reverse)	5'-CCTCACGGTACTTGTTC-3'						
NS1 (Forward)	5'-GTAGTCATATGCTTGTCTC-3'						
18L (Reverse)	5'-CACCTACGGAAACCTTGTTACGACTT-3'						

#### 3.2.3. DNA purification and sequencing

The DNA fragments amplified (please see section 2.2.2.) were separated in an electrophorese agarose gel (0.8%), observed using an UV-transilluminator (Molecular Imager® Gel Doc<sup>™</sup> XR System, BioRad) and extracted from the gel by excision with a razor blade. The purification of the PCR products was carried out with NZYGelPure Kit (NZYtech, Lisbon, Portugal), following the manufacturer's instructions. Then, samples were re-run in

an agarose gel to confirm the integrity. Finally, DNA concentration was quantified by measuring the absorbance at 260 nm, and DNA purity was calculated using the relation between the absorbance at 260 nm and 280 nm (OD<sub>260</sub>/OD<sub>280</sub>) using a microplate reader (Thermo Scientific<sup>™</sup> Multiskan<sup>™</sup> GO). The DNA concentration was calculated using the following Beer-Lambert law equation [1] (Swinehart et al. 1962).

$$Abs = \mathcal{E} * l * c \quad [1]$$

Where,

*E* - absorptivity (50 ng.cm/µL)

- I path length of the beam of the light through the material sample
- c concentration of DNA

The DNA fragments were sent to STAB Vida (Lisbon) for sequencing and the DNA sequences obtained were aligned using CLUSTAL program (<u>https://www.ebi.ac.uk/Tools/msa/clustalo/</u>), analyzed by using the Basic Local Alignment Search Tool (BLAST) (<u>http://www.ncbi.nlm.nih.gov/BLAST/</u>), performing a comparative search against the database of existing sequences.

# 3.3. Ecotoxicological tests in aqueous medium following available standard protocol for microalgae

### 3.3.1. BSC microalga growth curves in a liquid medium

For the purpose of determining the BSC microalga growth curves in liquid medium, two representative culture media were tested for comparison: the BG<sub>11</sub> medium and the MBL medium whose detailed composition can be found in Annex A. An aliquot of 1 mL of the working culture was spread in a plate with agar BG<sub>11</sub> medium (with 1.5% agar). The plate was incubated for 2 weeks at  $24 \pm 2$  °C, photoperiod of  $16h^{L}$ :8h<sup>D</sup> hours and a light intensity of 65 µmol m<sup>-2</sup>s<sup>-1</sup> provided by 30W cool-white fluorescent lamps. A colony from the plate was then inoculated in both 25 mL of MBL and 25 mL of BG11 media in an orbital shaker at 100

rpm, at 20  $\pm$  2 °C, under continuous illumination with a light intensity of 65 µmol m<sup>-2</sup>s<sup>-1</sup> provided by 30W cool-white fluorescent lamps for 5 days. The cultures in both media were incubated in the same conditions as described above, for more 15 days, and were performed in triplicate. In order to analyze if it was necessary to adjust the exposure duration previewed in the standard protocol available for freshwater microalgae testing (OECD, 2006) the growth curve of the microalga isolated from the BSC was determined. Aliquots of 1.5 mL were collected from each culture replicate, over 10-days, in sterile conditions using a laminar flow chamber. The aliquots were collected for microtubes, after a good homogenization of each Erlenmeyer and for each one, the optical density at 440 nm was measured and the number of cells per mL was counted using and Neubauer chamber and an optical microscope (MOTIC BA310). The procedure described was performed three times each one with three replicates to evaluate the repeatability of the growth curve. The growth parameter followed (cell's density) is expressed as a mean  $\pm$  standard deviation.

#### 3.3.2. Reference substances test solutions

In order to assess the sensitivity of the isolated microalga from BSC, test solutions of both reference substances selected were prepared. The chosen reference substances were copper (Cu) in the form of Copper (II) Sulfate Anhydrous (CuSO<sub>4</sub>) and the herbicide glyphosate (GLY), in particular of its commercial formulation RoundUp ULTRA Max<sup>®</sup>, which contains 360 g/L of GLY or 28.85% (p/p). The copper concentrations used were [0.30; 0.24; 0.14; 0.08; 0.05; 0.03] mgCu/L diluted in BG<sub>11</sub> medium and the GLY concentrations were [7.32; 5.86; 4.69; 3.75; 3.0; 0.75] mgGLY/L. The range of concentrations for both copper and GLY were obtained with range finding tests.

#### 3.3.3. Ecotoxicological test in BG11 medium with the BSC microalga

The ecotoxicological tests with the reference substances followed the standard protocol OECD No. 201 for the freshwater algae and cyanobacteria, growth inhibition test (OECD, 2006).

For this purpose, 400 µL of the BSC microalga from the working culture was inoculated in a 250 mL Erlenmeyer with 150 mL of BG<sub>11</sub> medium and incubated in a shaker at 100 rpm,  $20 \pm 2$  °C and under continuous illumination with a light intensity of 65 µmol m<sup>-2</sup>s<sup>-1</sup>, provided from 30W cool-white fluorescent lamps for 4 days. The tests were conducted in 24-well microplates and each one was filled with 900 µL either of the test solution (at different concentrations) or of the BG<sub>11</sub> medium in the controls (CTLs) and 100 µL of the BSC microalga inoculum (figure 3). The number of cells in this inoculum was previously counted, by using a Neubauer chamber, and it was diluted to obtain a cell's density of 10<sup>5</sup> cells/mL. The initial cells density in each well of the microplates was 10<sup>4</sup> cells/mL. The tests were conducted under the same conditions above described for the working cultures and the inoculum preparation. Growth inhibition tests were conducted for 5 days and four replicates (in four different wells of the microplates) were prepared for each concentration of the reference substances tested. The 24-well microplates were placed on an orbital shaker during the test period, at 130 rpm, to minimize sedimentation, and its place in the orbital shaker was changed every day to ensure equal light conditions to all the plates. At the end of the test, the cell's density in each well was counted as previously described and algae growth rate was calculated by using the following equation [2] (OECD, 2006). The procedure described was performed three times each one with four replicates to evaluate the repeatability of the ecotoxicological test. The growth parameter followed (cell's density) is expressed as a mean ± standard deviation.

$$\mu_{i-j} = \frac{\ln X_j - \ln X_i}{t_j - t_i} (day^{-1}) \ [2]$$

Where,

 $\mu_{i-j}$  – average specific growth rate from time i to j;

X<sub>i</sub> – biomass at a specific time i;

 $X_j$  – biomass at a specific time j;



Figure 3 - Microplates incubated in the orbital shaker at 130 rpm (a) and example of a 24-well after a 5-days of incubation period (b). Each line of wells has the four replicates for the CTL and for two of the tested concentrations. The wells of the last column and of the last line were filled with water to reduce evaporation.

### 3.4. Ecotoxicological tests in soil with the BSC microalga

### 3.4.1. Standard artificial soil

The standard artificial soil, called OECD soil was used as a test substrate for developing a more ecological relevant methodology for the ecotoxicological test with the BSC microalga. This soil is composed by a mixture of 70% fine quartz sand, 20% kaolin clay and 10% of finely ground sphagnum peat. The pH of the mixture was adjusted to  $6.0 \pm 0.5$  by adding calcium carbonate (OECD, 1984). The soil pH was measured with a pH probe (Jenway 2510) in a 1:5 (w/v) soil: water suspension. The maximum water holding capacity (WHC<sub>max</sub>) was calculated following an adaptation of the standard protocol (ISO, 2008) and was measured as follows: samples of the soil were placed in flasks (3 replicates), which bottom was replaced by a filter paper and then immersed for 3 h in water. Then, the flasks were placed over absorbent paper for free draining for 2 h and the saturated soil was weighed. The soil was dried for 24 h, at 105 °C and weighed again and the average WHC<sub>max</sub> was calculated.

#### 3.4.2. BSC microalga growth curves in soil

To determine the optimum exposure time for the soil ecotoxicological test with the BSC microalga, growth curves for the species were obtained. The inoculation of the microalga in the soil was performed on sterile, disposable plastic Petri plates of 90x15 mm. To each Petri plate was added 12g of OECD soil which was moistened with the volume of BG<sub>11</sub> necessary to adjust the WHC<sub>max</sub> to 100% (approx. 2.5 mL).

The BSC microalga inoculum was prepared as described in section 2.1, the number of cells was counted and adjusted by dilution to 10<sup>7</sup> cell/mL. To each plate 1.0 mL of inoculum was spiked on the soil with a micropipette, ensuring that the inoculum was homogeneously distributed on the soil surface. All the procedure was carried out in aseptic conditions (figure 4). The initial inoculum concentration of 10<sup>7</sup> cell/mL was chosen based on previous tests.

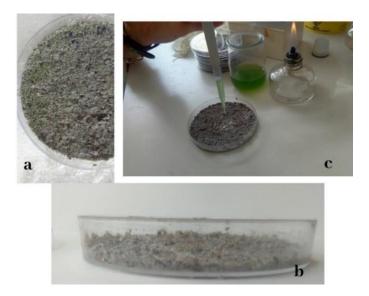


Figure 4 - Petri plate with an OECD soil layer. Top view (a) and side view (b). Soil inoculation with the BSC microalga (c).

The Petri plates were incubated at  $24 \pm 2$  °C under photoperiod  $16h^{L}:8h^{D}$ , light intensity of 125 µmol m<sup>-2</sup>s<sup>-1</sup> provided from 30W cool-white fluorescent lamps for 13 days (figure 5). Three replicates (3 Petri plates) were prepared as described and left to rest for 2h and then for each one, the soils were thoroughly mixed and placed in a 50.0 mL falcon tube for

immediately chlorophyll *a* quantification, in order to determine the initial content provided by the inoculum added to the soil. The remaining replicates were collected after 4, 6, 8, 11 and 13 days and the same procedure was performed. Each Petri plate was daily rehydrated with 1.0 mL of the BG<sub>11</sub> medium by using a sterilized Pasteur pipette, during the 9 days of growth. The procedure described was performed three times each one with three replicates to evaluate the repeatability of the growth curve. The growth parameter followed (chlorophyll *a* content) is expressed as a mean  $\pm$  standard deviation.

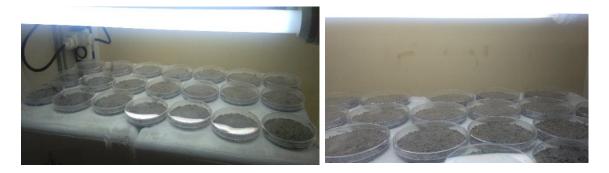


Figure 5 – Petri plate plates with OECD soil incubated for obtaining BSC microalga growth curves.

### 3.4.3. Pigment content

The quantification of the chlorophyll *a* content of the OECD soil, inoculated with the BSC microalga, was based on Nam et al. (2015). Soil samples were thoroughly mixed and placed in a 50.0 mL falcon tubes. Afterward, 5.0 mL of methanol 100% (v/v) were added to each sample ensuring that the entire soil was drowned. The tubes were vortexed for 10 seconds. The chlorophyll *a* extraction was made in the dark to prevent its degradation. To simplify the process, the tubes were covered with aluminum foil and placed in an orbital shaker at  $\pm$  230 rpm for at least 2 h. After, the tubes were centrifuged (Centrifuge 5804R, Eppendorf) at 11,000 rpm for 10 min. The supernatant was transferred to a 1.5 mL microtubes and placed in a centrifuge for 5 minutes at 13,000 rpm. The supernatant was measured at 663 nm in a

spectrophotometer. If necessary, the extracts were diluted with 100% (v/v) methanol. A blank test was performed using BG<sub>11</sub> medium instead of algal suspension.

The chlorophyll *a* content was calculated using the following equation [3] (Meeks and Castenholz, 1971):

$$C\left(\frac{\mu g}{mL}\right) = OD663nm * 12,7 * DF$$

Where,

C – concentration (µg/mL)
 OD – optical density at 663 nm
 DF – dilution factor

### 3.4.4. Preparation of test solutions

For the ecotoxicological test on the soil the two test solutions of both reference substances selected were prepared according to the section 2.3.1. However, different concentrations for both substances were used. The copper concentrations used were [0.51; 0.25; 0.13; 0.06; 0.03] mg Cu/g<sub>soil</sub> diluted in BG<sub>11</sub> medium and the GLY concentrations were [5.86; 4.69; 3.75; 3; 0.75] mg GLY/g<sub>soil</sub>. The range of concentrations for both Cu and GLY were obtained with range finding tests.

### 3.4.5. Ecotoxicological test in the OECD soil and with the BSC microalga

The ecotoxicological test in the soil was performed on sterile, disposable plastic Petri plates of 90x15 mm. To each Petri plate was added 12g of OECD soil which was moistened with the amount in volume of BG<sub>11</sub> necessary to adjust the WHC to 100% (approx. 2.5 mL). Then 1.0 mL of the test substance (for the different concentrations) were added homogenously through the soil in the proper concentrations and left to stabilize for 24 hours. The BSC

microalga inoculum was prepared as described in section 2.1, the number of cells was counted and adjusted by dilution to  $10^7$  cell/mL. To each plate 1.0 mL of inoculum was spiked on the soil with a micropipette, ensuring that the inoculum was homogeneously distributed on the soil surface. All the procedure was carried out in aseptic conditions, as shown in figure 4. The Petri plates (treatments and control) were incubated at  $24 \pm 2$  °C under photoperiod  $16h^{L}:8h^{D}$ , light intensity of 125 µmol m<sup>-2</sup>s<sup>-1</sup> provided from 30W cool-white fluorescent lamps for 13 days. The initial sample was left to rest for 2h and then thoroughly mixed and placed in a 50.0 mL falcon tube for chlorophyll *a* quantification. After the 13-days of exposure (determined based on growth curves), the same process for chlorophyll *a* content (please see section 2.4.2.) was made for all the soil samples. The procedure described was performed three times each one with three replicates to evaluate the repeatability of the ecotoxicological test.

#### 3.5. Statistical analysis

The average growth rate and the average percentage of inhibition towards the CTL for each concentration of the reference substances tested were calculated and are presented with the corresponding standard deviation. For each test, the data was analyzed for the homogeneity of variances and normality of data by using the Levene's test and by the Shapiro-Wilk test, respectively. One-Way ANOVA was used to test for significant differences in microalga growth rate between the concentrations of each reference substance tested, followed by a Dunnet test to look for differences from the CTL. A significance level of 0.05 was used for all tests. When ANOVA assumptions were not met, the Kruskal-Wallis non-parametric test was used to support the One-Way ANOVA test. The EC<sub>50</sub> was calculated using non-linear interpolation methods of regression analysis with the statistical program STATISTICA version 13.3. The growth curves of the microalga, both in liquid and soil medium were obtained by fitting a non-linear sigmoidal 4PL, X is log(concentration) model to data, using the software GraphPad Prism version 6.

# 4. Results

## 4.1. Identification of the BSC selected microalga

### 4.1.1. Morphological characterization of the microalga

The morphological characterization of the microalga species isolated from BSC collected in a Portuguese soil was performed under an optic microscopic observation as shown in figure 6. This unicellular green alga is characterized by spherical to ovoid cells, arranged in colonies when the concentration in the medium increases.

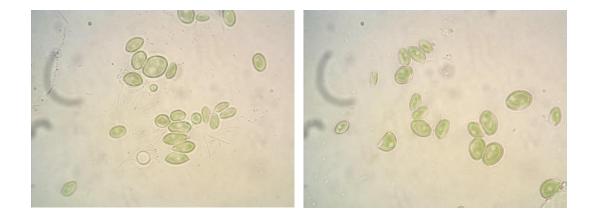


Figure 6 – General aspect of the morphology of the cells of the soil microalga, at the end of the isolation process with a 100X amplification under optical microscopy.

### 4.1.2. Genetic identification of the BSC selected microalga

Genomic DNA of the BSC selected microalga was amplified by PCR using primers for 18S rRNA and ITS1 (Internal Transcribed Spacer 1). The PCR products, with molecular sizes in agreement with the expected,1300 bp and 1700 bp, respectively (figure 7), were sequenced.

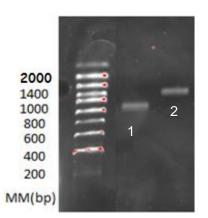


Figure 7 - Electrophoretic separation of the DNA fragments, amplified by PCR using the pair of primers (1) NS7m/LR1850 and (2) NS1/18L.

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The BLASTn analyses of the sequences obtained showed a 98% identity with the genera *Micractinium*, in particular with the species *Micractinium inermum*. The corresponding BLAST (figure 8) and sequences are provided in Annex B.

	Alignments Download - GenBank Graphics Distance tree of results						4
	Description	Max score	Total score	Query cover	E value	Ident	Accessio
	Micractinium inermum strain NLP-F014 small subunit ribosomal RNA gene, partial sequence; internal transcribed spacer 1, 5 8S ribosomal RNA gene, and internal transcribed	2215	2215	96%	0.0	98%	KF597304
	Chlorellaceae sp. MCWWS28 18S ribosomal RNA gene, partial sequence; internal transcribed spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer 2, complete	2211	2211	96%	0.0	98%	KP204570.
0	Chlorella-like algae voucher HS25 18S ribosomal RNA gene, partial sequence: internal transcribed spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer 2, com	2202	2202	96%	0.0	98%	KU641127.
	Chlorellaceae sp. MCWWW8 18S ribosomal RNA gene, partial sequence; internal transcribed spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer 2, complete	2032	2032	96%	0.0		KP204586.
							a
A	Ilignments Download - GenBank Graphics Distance tree of results						
	Description	Max score	Total score	Query cover	E value	Ident	Accessio
1	Micractinium inermum strain NLP-F014 small subunit ribosomal RNA gene, partial sequence: internal transcribed spacer 1.5.8S ribosomal RNA gene, and internal transcribed	2089	2089	99%	0.0	98%	KF597304
1	Chlorellaceae sp. MCWWS28 18S ribosomal RNA gene, partial sequence, internal transcribed spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer 2, complete	2085	2085	99%	0.0	98%	KP204570
1	Chlorella-like algae voucher HS26 18S ribosomal RNA gene, partial sequence; internal transcribed spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer 2, com	2078	2078	99%	0.0	98%	KU641127
1	Micractinium inermum small subunit ribosomal RNA gene, partial sequence, internal transcribed spacer 1, 5.8S ribosomal RNA gene, and internal transcribed spacer 2, comple	2069	2069	97%	0.0	98%	KM114868
							0
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11	Alignments 🖉 Download 🤟 GenBank Graphics Distance tree of results	Mari	Tatal	0	É	ĩ	
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	Alignments 😨 Download 🤟 GenBank. Graphics: Distance tree of results				-	Ident	Access
0	Alignments Download - GenBank Graphics Distance tree of results Description	score	score	cover 97%	value	Ident	Access AB73160
	Alignments Download CenBank Graphics Distance tree of results Description Micractinium inemum genes for SSU rRNA, ITS1, 5.8S rRNA, ITS2, LSU rRNA, culture, collection: NIES 2171	score 1847	score 1847	cover 97% 97%	value 0.0	Ident 98%	Access AB73160 KU67436
	Alignments Download CenBank Graphics Distance tree of results Description Micractinium inemum genes for SSU rRNA, ITS1, 5.8S rRNA, ITS2, LSU rRNA, culture, collection: NIES 2171 Chlorella, sp. HS-2 18S ribosomal RNA gene, eartial sequence	score 1847 1836	1847 1836	cover 97% 97% 97%	value 0.0 0.0	Ident 98% 97% 97%	Access AB73160 KU67436 KT88390
0	Alignments Description Description Micractinium: Inemum genes for SSU rRNA, ITS1, 5.8S rRNA, ITS2, LSU rRNA, culture: collection: NIES 2171 Chlorella.so, HS-2 18S ribosomal RNA gene, partial sequence Micractinium: sp. KNUA036 18S ribosomal RNA gene, partial sequence	score 1847 1836 1836	score 1847 1836 1836	cover 97% 97% 97%	value 0.0 0.0 0.0	Ident 98% 97% 97%	Access AB73160 KU67436
	Alignments Description Description Micractinium inerrum genes for SSU rRNA, ITS1, 5.85 rRNA, ITS2, LSU rRNA, culture_collection: NIES 2171 Chlorelia so, HS-2 18S ribosomal RNA gene, cartial sequence Micractinium sp, KNUA036 18S ribosomal RNA gene, cartial sequence Micractinium sp, KNUA034 18S ribosomal RNA gene, cartial sequence	score 1847 1836 1836	score 1847 1836 1836	cover 97% 97% 97%	value 0.0 0.0 0.0	Ident 98% 97% 97%	Access AB73160 KU67436 KT88390
	Alignments Description Description Micractinium inemum genes for SSU rRNA. JTS1. 5.85 rRNA. ITS2. LSU rRNA, culture_collection: NIES 2171 Chlorella.so. HS-2 15S ribosomal RNA gene, eartial sequence Micractinium sp. KNUA036 15S ribosomal RNA gene, eartial sequence Micractinium sp. KNUA036 15S ribosomal RNA gene, eartial sequence Micractinium sp. KNUA036 15S ribosomal RNA gene, eartial sequence	score 1847 1836 1836	score 1847 1836 1836 1836	cover 97% 97% 97%	value 0.0 0.0 0.0	Ident 98% 97% 97% 97%	Access AB73160 KU67436 KT88390 KM24332
	Alignments Download CanBank. Graphics Distance tree of results Description Micractinium inerrum genes for SSU rRNA. JTS1. 5.85 rRNA. ITS2. LSU rRNA. culture collection: NIES:2171 Chlorella so. HS: 2185 ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036 18S ribosomal RNA gene, oartial sequence Micractinium sp. KNUA036	score 1847 1836 1836 1836 1836	score 1847 1836 1836 1836	cover 97% 97% 97% 97%	value 0.0 0.0 0.0 0.0 0.0	Ident 98% 97% 97% 97%	Access AB73160 KU67436 KT88390 KM24332
1	Alignments       Description         Micractinium inerrum genes for SSU rRNA, ITS1, 5.85 rRNA, ITS2, LSU rRNA, culture collection: NIES 2171         Chlorelia so, HS-2 15S ribosomal RNA gene, eartial sequence         Micractinium sp, KNUA036 18S ribosomal RNA gene, eartial sequence         Micractinium sp, KNUA034 18S ribosomal RNA gene, eartial sequence         Micractinium sp, KNUA034 18S ribosomal RNA gene, eartial sequence         Micractinium sp, KNUA034 18S ribosomal RNA gene, eartial sequence         Micractinium sp, KNUA034 18S ribosomal RNA gene, eartial sequence         Micractinium sp, KNUA034 18S ribosomal RNA gene, eartial sequence         Micractinium sp, KNUA034 18S ribosomal RNA gene, eartial sequence         Micractinium inerrum small subunit ribosomal RNA gene, eartial sequence internal transcribed spacer 1, 5, 8S ribosomal RNA gene, and internal transcribed spacer 2, complete	score 1847 1836 1836 1836 1836	score 1847 1836 1836 1836 Total score	cover 97% 97% 97% 97% 97%	value 0.0 0.0 0.0 0.0 0.0	Ident 98% 97% 97% 07% 07%	Access AB7316( KU67434 KT88390 KM2433; Accessie
	Alignments       Description         Micractinium: inemum genes for SSU rRNA. ITS1, 5.85 rRNA. ITS2, LSU rRNA, cuture_collection: NIES 2171         Chlorella.co. HS-2 T85 ribosomal RNA gene, eartial sequence         Micractinium: spi, KNUA036 185 ribosomal RNA gene, eartial sequence         Micractinium: spi, KNUA036 185 ribosomal RNA gene, eartial sequence         Micractinium: spi, KNUA034 185 ribosomal RNA gene, eartial sequence         Micractinium: spi, KNUA034 185 ribosomal RNA gene, eartial sequence         Micractinium: spi, KNUA034 185 ribosomal RNA gene, eartial sequence         Micractinium: spi, KNUA034 185 ribosomal RNA gene, eartial sequence         Micractinium: spi, KNUA034 185 ribosomal RNA gene, eartial sequence         Micractinium: spi, KNUA034 185 ribosomal RNA gene, eartial sequence         Micractinium: spinal subunit ribosomal RNA gene, eartial sequence: internal transcribed seacer 1, 5.85 ribosomal RNA gene, and internal transcribed seacer 2, corrected         Micractinium: inemum genes for SSU rRNA_ITS1_S8 rRNA_ITS2_LSU rRNA_cuture_collection. NIES 2171	score 1847 1836 1836 1836 1836 1836 2002 1999	score 1847 1836 1836 1836 1836 Total score 2002	Cover 97% 97% 97% 97% 97% 0000 2000 100%	value 0.0 0.0 0.0 0.0 E value 0.0	Ident 98% 97% 97% 07% 07%	Access AB73160 KU67439 KT88390 KM2433 Access KM114866
	Alignments Description  Micractinium inemum genes for SSU rRNA. JTS1. 5.85 rRNA. ITS2. LSU rRNA. culture_collection: NIES 2171 Chlorolla.so. HS-2 155 ribosomal RNA gene, eartial sequence  Micractinium iser, KNUA036 185 ribosomal RNA gene, eartial sequence  Micractinium iser, KNUA036 185 ribosomal RNA gene, eartial sequence  Micractinium iser, KNUA034 185 ribosomal RNA gene, eartial sequence  Micractinium isermum small subunit ribosomal RNA gene, eartial sequence: internal transcribed seacer 1.5.85 ribosomal RNA gene, and internal transcribed seacer 2. complet  Micractinium isermum genes for SSU rRNA. JTS1. 5.85 rRNA. ITS2. LSU rRNA. culture_collection. NIES 2171  Micractinium isermum genes for SSU rRNA. JTS1. 5.85 rRNA. ITS2. LSU rRNA. culture_collection. NIES 2171  Micractinium isermum genes for SSU rRNA. JTS1. 5.85 rRNA. ITS2. LSU rRNA. culture_collection. NIES 2171  Micractinium isermum genes for SSU rRNA. JTS1. 5.85 rRNA. ITS2. LSU rRNA. culture_collection. NIES 2171	score 1847 1836 1836 1836 1836 1836 2002 1999 1999	score 1847 1836 1836 1836 1836 70tal score 2002 1999	Cover 97% 97% 97% 97% 97% 0000 100%	value 0.0 0.0 0.0 0.0 0.0 Value 0.0 0.0	98% 97% 97% 97% 97% 07%	Accessi KM2433 KM2433 Accessi KM114865 AB731604

Figure 8 - BLAST results for the sequences with the primers (a) NS7 LR1850+LR1850, (b) NS7 LR1850+NS7m, (c) NS1+18L and (d) NS1 18L+NS1 obtained showing a homology of 98% with *Micractinium inermum*.

### 4.2. Ecotoxicological test in liquid medium with the BSC microalga

4.2.1. Growth curves and ecotoxicological tests in aqueous medium following the available standard protocol for freshwater microalgae and cyanobacteria

In order to better characterize the growth of *Micractinium inermum*, two artificial media, the BG<sub>11</sub> medium (Rippka et al. 1992) and the Woods Hole MBL medium (Nichols, 1973) were used for growing the microalga. The results showed that *M. inermum* under a temperature of  $20 \pm 2$  °C, continuous illumination and a light intensity of 65 µmol m<sup>-2</sup>s<sup>-1</sup>, had a 3-day lag phase, reaching the exponential phase within 5-6 days and attaining the stationary phase within 7-8 days, in both media as shown in figure 9. When the stationary phase was reached, in BG<sub>11</sub> and MBL media the culture had a 45.6-fold and 28.5-fold increase in the density of cells, respectively. Based on the behavior of the microalga in both medium, the BG<sub>11</sub> medium was selected for the ecotoxicological tests with this species, as it was the medium already used for the isolation and for culturing the microalga, and it was also at this medium that the microalga displayed a greater growth.

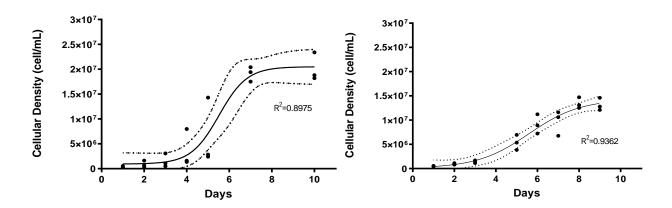


Figure 9- Growth curve of *M. inermum* cultured in two different media, BG11 (left) and MBL (right). Dots account for average values adjusted to the sigmoidal model and error bars for standard deviation. The r-squared value of the sigmoidal model adjusted to data is displayed.

These results established the time length for the ecotoxicological tests to be performed, which was set up in 5 days with an exponential growth of 14.8-fold increase. Since the growth in

the BG<sub>11</sub> medium was superior to 14 times, this level of increase was chosen as the validity criteria of the assay.

#### 4.2.2. Ecotoxicological test in BG11 medium with M. inermum

The ecotoxicological tests were carried out for 5 days and three independent tests were conducted for copper and glyphosate (figure 10 and figure 11). The  $EC_{50}$ , NOEC and LOEC values, including 95% confidence intervals based on specific growth rate (see section 3.3.3.) were calculated and are presented in table 2.

	C50, LOEC and NOEC v nd GLY exposure to <i>M. ir</i>		~	nfidence intervals based o	on specific growt	h rate after
Tests	Copper (mg/L) Glyphosate (mg		sate (mg/L)	J∕L)		
	EC <sub>50</sub>	NOEC	LOEC	EC <sub>50</sub>	NOEC	LOEC
А	0.24 [0.0;0.44]	0.03	0.05	2.94 [2.47;3.40]	< 0.75	≤ 0.75
В	0.21 [0.19;0.22]	0.03	0.05	2.72 [2.27;3.17]	< 0.75	≤ 0.75
С	0.17 [0.15; 0.20]	0.03	0.05	2.89 [2.41; 3.38]	< 0.75	≤ 0.75

All the tests were valid since there was an increase in the cell's density of the CTL of 40, 47 and 50 times, respectively, surpassing the validity criteria. The results for copper toxicity (figure 10) showed that after a 5-day exposure period there was a dose-dependent response on the growth rate of *M. inermum*. At the highest concentration of copper tested (0.3 mg Cu/L) the growth of *M. inermum* was completely inhibited, occurring total cellular death and, at the lowest concentration (0.03 mg Cu/L) the growth was stimulated. In all three independent tests, A, B and C, the results showed statistically significant differences between treatments ( $F_{A}$ = 379.19; df=43,37; p<0.01;  $F_{B}$ = 795.88; df=43,37; p<0.01 and  $F_{C}$ = 65.05; df=43,37; p<0.01). The Dunnett's multiple comparison test showed significant differences equal and above 0.05 mg/L Cu. The results of the ANOVA tests allowed the determination of NOEC and LOEC values, described in table 2.

In what regards the repeatability of the tests with *M. inermum*, it can be checked through the overlap of the confidence intervals of the estimated  $EC_{50s}$  that a good level of repeatability was obtained especially for Tests B and C for copper.

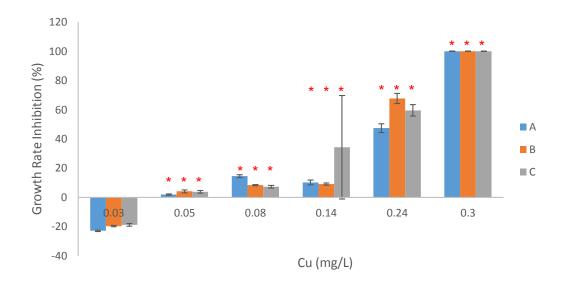


Figure 10 - Percentage of growth rate Inhibition (%) towards the control for *M. inermum*, measured by the, exposed to copper in three independent tests (A), (B) and (C). Bars account for average values and error bars for standard deviation. \* above each bar indicate statistically differences between the treatment and the control.

All the tests using glyphosate were valid since there was an increase in the growth of the microalga in the CTL of 32, 38 and 34 times, respectively, which once again is in agreement with the validity criteria. The results on glyphosate toxicity (figure 11) showed a dose-dependent increase on the % of growth inhibition towards the control after 5 days of exposure of *M. inermum* to the commercial formulation of GLY. The highest concentration promoted a complete cellular death on *M. inermum* in the soil. In all three independent tests, A, B and C, the results showed statistically significant differences between the treatments and the control ( $F_A$ = 1130.94; df=38,32; p<0.01;  $F_B$ = 2821.05; df=38,32; p<0.01 and  $F_C$ = 979.40; df=38,32; p<0.01). According to the Dunnet's test, a significant inhibition of microalga growth was recorded for all the concentrations tested, in the three ecotoxicological test, when compared to the control (p<0.05). The results of the ANOVA tests allowed the determination of NOEC and LOEC values, described in table 2. Once again, and taken into

account the  $EC_{50}$  values estimated and the overlap of the corresponding 95% confidence intervals, the three independent tests confirmed the repeatability of the procedure.

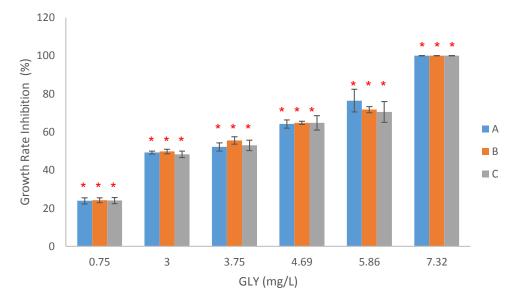


Figure 11 - Percentage of growth rate Inhibition (%) towards the control for *M. inermum* exposed to glyphosate in three independent tests (A), (B) and (C). Bars account for average values and error bars for standard deviation. \* above each bar indicate statistically differences between the treatment and the control.

## 4.3. Ecotoxicological test in artificial soil with the BSC microalga

#### 4.3.1. M. inermum growth curve in artificial OECD soil

For the soil test, pigment extraction and pigment content in soil (expressed in mg/kg) based on chlorophyll *a* quantification was the method chosen for assessing algae growth in soil, although it is recognized that the content of pigments is an indirect measure of microalga growth.

The growth of *M. inermum* on soil under a temperature of  $24 \pm 2$  °C, a photoperiod of  $16h^{L}:8h^{D}$  and a light intensity of 125 µmol m<sup>-2</sup>s<sup>-1</sup>, presented a 4-days lag-phase and an exponential growth, starting at day 6 and extending up to 13 days (figure 12). In fact, the growth test has to be repeated in a near future, for a longer period in order to determine when the stationary phase starts, however, such a long period may be too long for an

ecotoxicological test. Therefore, was decided to finish the test in the exponential growth phase as it was done for the tests in the aquatic medium.

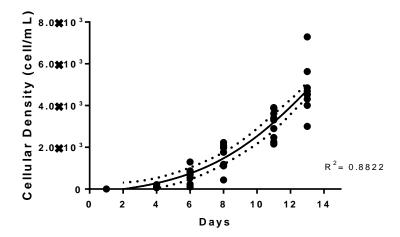


Figure 12 - Growth curve of *M. inermum* cultivated in artificial OECD soil, determined by quantification of chlorophyll a. Bars account for average values and error bars for standard deviation. The r-squared value of the sigmoidal model adjusted to data is displayed.

A 449-fold increase in pigment content on the artificial soil was recorded after the exposure period. Since the growth in the soil was superior to 100 times, this level of increase was chosen as the validity criteria of the assay. According to the results obtained, the time length of the ecotoxicological test in soil was established for 13 days.

#### 4.3.2. Ecotoxicological test with *M. inermum* in OECD artificial soil

A preliminary range-finding test with copper and glyphosate concentrations was performed for the *M. inermum*, in order to select the final range of concentrations for the ecotoxicological tests (please see chapter 2.3.6). The toxicity tests were carried out for 13 days in order to calculate the  $EC_{50}$  for growth inhibition and three independent tests were conducted for each contaminant (figure 13 and figure 14), in order to assess the repeatability of the test protocol. The  $EC_{50}$ , NOEC and LOEC values, including 95% confidence intervals based on growth rate (see section 3.3.3) were estimated or obtained and are presented in table 3.

	EC50, LOEC, and NOEC val y Cu and GLY exposure to a				n specific grow	vth rate
Tests	Copper	(mg/L)		Glyphosa	ate (mg/L)	
	EC <sub>50</sub>	NOEC	LOEC	EC <sub>50</sub>	NOEC	LOEC
А	0.19 [0.12;0.26]	< 0.03	≤ 0.03	3.43 [3.18;3.67]	< 0.75	≤ 0.75
В	0.20 [0.14;0.25]	< 0.03	≤ 0.03	3.81 [3.46;4.17]	< 0.75	≤ 0.75
С	0.17 [0.11;0.22]	< 0.03	≤ 0.03	3.37 [2.82; 3.94]	< 0.75	≤ 0.75

For copper, all the tests were valid since there was an increase in total content of chlorophyll in the CTL 107, 157 and 156 times, respectively, which is in agreement with the validity criteria. In all three independent tests, A, B and C, the results showed statistically differences between treatments ( $F_A$ = 26.44; df=20,15; p<0.01;  $F_B$ = 55.54; df=20,15; p<0.01 and  $F_c$ = 47.31; df=20,15; p<0.01). According to the Dunnet's test, statistically significant differences in terms of total chlorophyll content of the soil was recorded for all the concentrations tested, in the three ecotoxicological tests, when compared to the control (p<0.05). The results of the ANOVA tests allowed the determination of NOEC and LOEC values, described in table 3.

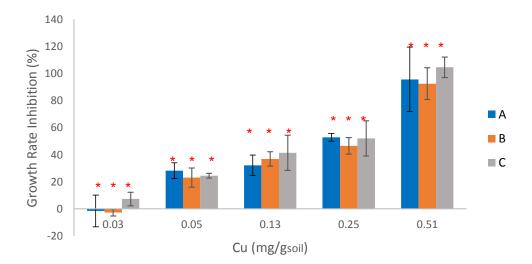


Figure 13 - Percentage of growth rate Inhibition (%) towards the control for *M. inermum* exposed to copper on soil for the three independent tests (A), (B) and (C). Bars account for average values and error bars for standard deviation. \* above each bar indicate statistically differences between the treatment and the control.

All the tests using glyphosate in soil were valid since there was an increase in total content of chlorophyll in the CTL 153, 202 and 101 times, respectively, which once again is in agreement with the validity criteria. In what regards the exposure of *M. inermum* to glyphosate in all the three independent tests, A, B and C, the results showed statistically differences between the treatments ( $F_{A}$ = 95.36; df=20,15; p<0.01;  $F_{B}$ = 33.69; df=20,15; p<0.01 and  $F_{C}$ = 62.65; df=20,15; p<0.01). According to the Dunnet's test, statistically significant differences in terms of total chlorophyll content of the soil, were recorded for all the concentrations tested, in the three ecotoxicological test, when compared to the control (p<0,05). The results of the ANOVA tests allowed the determination of NOEC and LOEC values, described in table 3.

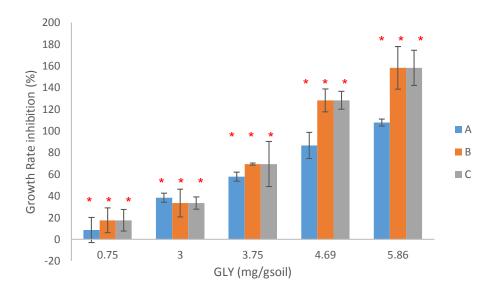


Figure 14 - Percentage of growth rate Inhibition (%) towards the control for *M. inermum* exposed to glyphosate on soil for three independent tests (A), (B) and (C). Bars account for average values and error bars for standard deviation. \* above each bar indicate statistically differences between the treatment and the control.

The results recorded, for the independent tests conducted, for the two reference substances, and considering the  $EC_{50}$  values and the corresponding 95% confidence intervals, were demonstrative of the repeatability of the test procedures.

## 5. Discussion

# 5.1. Characterization and identification of a selected microalga from the BSC

The microalga used in this study was isolated from biological crusts collected in a Portuguese soil located in Center of Portugal, sub-region of Baixo Mondego River. Genomic sequences amplified using primers for 18S/ITS were obtained and compared with those available on the NCBI GenBank database through the BLAST bioinformatic tool. *Micractinium inermum* was identified with a 98% similarity, which is considered the threshold value for differentiating two species (Kim et al. 2014), and is consistent with the morphological observations (please see section 4.1.).

The molecular data has been proven to be essential for establishing the actual phylogenetic relationships between individual green algal *taxa*. In our study we used 18S rRNA gene and the ITS1 region as molecular markers to identify the selected green alga. The 18S RNA gene is considered as the best molecular marker for studying phylogenetic relationship in eukaryotes as it is universal and is composed of highly conserved as well as variable domains (Gaysina et al. 2013, Patwardhan et al, 2014). The ITS is a non-coding region with high interspecific variability allowing differentiation of species within a genus, but low intra-specific variability preventing the separation of individuals or strains within the same species (Lin et al. 2017).

The genus *Micractinium* belongs to the Phylum Chlorophyta, Family Chlorellaceae<sup>3</sup>. *Chlorella vulgaris* is used as a bioindicator in ecotoxicological assays and belongs to the same family as *M. inermum*. *M. inermum* was first described by Fresenius (1858) and it is characterized by spherical or ovoid cells mostly arranged in colonies that propagate by auto-sporulation (Luo et al. 2006). Only two species of the *Micractinium* genus are described as having spherical cells, the *M. reisseri* and *M. inermum* (Hoshina, 2014). Recently, this genus is known to have spherical to ovoid cells with a parietal and cup-shaped chloroplast with an evident pyrenoid. It is also characterized by thin smooth cell wall capable of producing bristles (Hong et al. 2015). However, not all strains belonging to *Micractinium* are able to produce bristles. This is relevant for strains such as CCAP 211/92 which was isolated from

<sup>&</sup>lt;sup>3</sup> <u>http://www.algaebase.org/search/genus/detail/?genus\_id=43421</u>, accessed last on: 04/12/18

a soil sample collected from Mahe Island (Seychelles) (Luo et al. 2010). However, whether the presence of bristles is triggered by environmental factors needs to be proven in further studies (Pröschold et al. 2010), especially using molecular methods designed to detect differences at the DNA level (Germound et al. 2013). A study conducted by Hong et al. (2015), determined the optimal growth temperatures of the microalgae strains KNUA029, KNUA034 and KNUA032 Micractinium obtained from bloom samples in West Antarctica in liquid medium. Results through a 24-day experimental period showed that Micractinium strains from this region can tolerate lower temperatures, maintaining an optimum growth temperature of 20 °C and are able to tolerate adverse conditions. M. inermum in this work also exhibited growth to the same temperature of 20 °C, verifying that these species are mesophilic. Comparing the growth curves also in BG11 medium, obtained by Hong et al. (2015) and ours, *M. inermum* grew faster requiring half of the time to reach exponential phase, however, is important to emphasize that these species are from distinct climate regions. To the best of our knowledge, this is the first study evaluating the growth rate and the toxicity effects of contaminants on soil with *M. inermum*, since the information found in literature is mainly related to *Micractinium* sp. found in freshwater habitats (Issa et al. 2012, Varol et al. 2015). For this genus found in freshwater environments, several studies have been conducted, mainly in the biotechnology field. China et al. (2018), isolated several microalgae from freshwater reservoirs and among the isolates, *Micractinium* strains were found to grow preferably at high levels of CO<sub>2</sub>, making them preferential to produce several bioactive compounds. In line with these applications, Onay et al. (2014) and Abou-Shanab et al. (2014), both described the suitability of *Micractinium* sp. for biodiesel production. Another study by Park et al. (2017), suggested that a blended wastewater could be used to cultivate *M. inermum* and to improve microalgae biomass production. Hong et al. (2015), suggested that *Micractinium* strains from the Antarctica could be used to be a potential source of omega-3 and omega-6 fatty acids, due to its high ratio of polysaturated fatty acids.

According to the AlgaeBase, *M. inermum* does not have yet a determined habitat (Hoshina et al. 2013). This work is one more step forward to confirm the presence of this species on soil, providing more information to help comprehend the ecology of this genus. The current research still focuses mostly in other groups as bioindicators of toxicity on soil, such as plants and invertebrates, and the soil microalgae group is still to be explored. As new habitats are explored and new records are found, becomes clear that our knowledge in what regards soil microalgae still needs improvement.

# 5.2. Ecotoxicological Assays in Aqueous Medium Following Available Standard Protocol for Freshwater Microalgae and Cyanobacteria

Taking into account that soil can be considered a semi-aquatic habitat (Jeffery et al. 2010), one of the aims of this study was to assess if the guideline for testing chemicals by OECD 201(OECD, 2006) with freshwater microalgae and cyanobacteria could be used to evaluate the toxicity of contaminants with soil microalgae species. Initially was necessary to determine the growth curve for *M. inermum* in order to established the time length for the upcoming ecotoxicological tests performed on this study. The data showed that *M. inermum* needed a 5-day growth period to reach exponential phase and that the growth conditions were favorable, in both aqueous media tested, the BG<sub>11</sub> and MBL media, as demonstrated in figure 9. However, *M. inermum* clearly grew better in BG<sub>11</sub> medium and for this reason BG<sub>11</sub> medium was selected as growth media for *M. inermum* through the entire work. BG<sub>11</sub> medium and MBL media have been used in multiple studies using freshwater microalgae and cyanobacteria (Marques et al. 2012; Yeh et al. 2012; Hentati et al. 2015; Silva et al. 2016; Arguelles et al. 2017). Comparing the composition of both media, found in Annex A, the differences consist essentially in the content of nitrogen since MBL medium is considered as a nitrogen-poor medium when compared to BG<sub>11</sub>. Abou-Shanab et al. (2014), cultivated Micractinium reisseri in municipal wastewater, observing a removal of 80% on nitrogen for the tertiary effluent. Other study by Park et al. (2017), demonstrated that cultivating Micractinium inermum in a sequencing-batch cultivation with a blended wastewater medium could be a promising strategy to cultivate and grow this species. Comparing to BG<sub>11</sub>, MBL medium is also supplied with vitamins. Croft et al. 2006, explains the importance of the three most used B-complex vitamins (B12, B1, and B7) in the culture media of microalgae explaining that species with these requirements had probably lost a gene involved in the biosynthesis of that cofactor. However, China et al. (2018), showed that strains of *Micractinium* were found to produce higher amounts of B-complex vitamins, such as B1 and B6. These findings show the higher needs of this species for nitrogen in opposition to vitamins, making the BG<sub>11</sub> medium more suitable for growth. Another observation is made when comparing the time length for cellular growth with this soil microalgae species and the guideline by OECD 201 for freshwater microalgae and cyanobacteria. The OECD 201 only requires a time length for the ecotoxicological tests of 72h for continuous illumination to 96h for photoperiod, however, this study data shows that the soil microalgae, *M. inermum* requires a 5-day time length which is a slightly longer time length.

The reference contaminant, copper (II), was chosen because of its well-known toxicity for different species (Moreno-Garrido et al. 2000; Debelius et al. 2009; Magdaleno et al. 2014; Chen et al. 2016), in particular for microalgae species, by affecting growth. The availability of information makes this contaminant appropriate for comparing species sensitivities, assessing the relevance of the proposal of *M. inermum* as a new test organism. The ecotoxicological test uses the inhibition of cellular division as a biomarker of the toxic mode of action of contaminants. The use of this endpoint is relevant since it is reproducible and less dependent of more demanding monitoring techniques and methodologies (Stauber et al. 2000). In the present study we evaluated the toxic effect of copper on the cells of M. inermum by measuring the growth inhibition and EC<sub>50</sub> values were estimated. The 5-day growth inhibition tests on liquid medium with *M. inermum* collected from soil showed a clearly dose-dependent response for the range of concentrations of copper tested (figure 10). The three independent tests were conducted and the validity criteria established by the OECD protocol were met with the species *M. inermum* collected from biological soil crusts. For the copper three EC<sub>50</sub> values with corresponding 95% confidence intervals (95% CI) were presented (table 2), and the repeatability of the tests was confirmed as they fit in the same range of values. The toxic effect of copper on the cellular growth was previously studied using different algal species (table 4), proving that the sensitivity to a toxic compound in microalgae is species-dependent. Different algal species appear to differ in their ability to produce certain compounds such as, intracellular phytochelatin, a metal-binding peptide, which can detoxify copper, which is reflected in the large variability in inter-species sensitivity of copper (Bossuyt et al. 2004). In multiple studies, freshwater microalgae such as C. vulgaris and P. subcapitata (presently Raphidocelis subcapitata) have been used as model species, as they are single cells that do not form aggregates in aqueous cultures (Exposito et al. 2017), they are easy to count, have a widespread geographical distribution, ecological relevance and sensitivity to toxicants (Janssen et al. 2003; Soto et al. 2011). However, sometimes, even for the same species, there is a great variability in the response to toxicants, likely due to variations among laboratories in operational procedures, growth conditions (e.g. temperature, light intensity), within other aspects. This can be observed in table 4, where, for the same species of microalgae and chosen contaminant (as Cu for example) the EC<sub>50</sub> values reported are different.

Algal cells are known to accumulate metals by physical adsorption and bioaccumulation, though, little information is available about the long-term accumulation process and effects on algae (Yan et al. 2002; Zhou et al. 2012) and, depending on the species, metals may affect algal growth by oxidative stress, respiration, nitrate uptake and loss of membrane integrity, being a strong inhibitor of photosynthesis (Machado et al. 2015; Hamed et al. 2017). However, copper is also an essential element, due to its role as cofactor of several enzymes and key role in oxygen-requiring chemical reactions (Jamers et al. 2013; Udeigwe et al. 2016). Is also involved in many physiological processes, acting as a structural element in regulatory proteins, in photosynthetic electron transport, mitochondrial respiration, oxidative stress response, cell wall metabolism and hormone signaling (Yruela et al. 2005). Thus, at lower concentrations it can cause stimulatory effects as observed in our tests with *M. inermum*.

Species	EC <sub>50</sub> (mg/L)	Reference	
	0.24 [0.0; 0.44]		
M. inermum	0.21 [0.19; 0.22]	Present work	
	0.17 [0.15; 0.20]		
Chlorella vulgaris	0.660 [0.39; 0.93]	Hadjoudja et al. 2009	
C. vulgaris	0.200	Playlock at al. 1985	
Selenastrum capricornutum*	0.400	Blaylock et al. 1985	
Pseudokirchneriella subcapitata*	0.154	Rodrigues, 2003	
C. ellipsoidea	0.489 [0.421; 0.534]	Magdaleno et al. 2014	
Scenedesmus obliquus	0.050	Yan et al. 2002	
Closterim lunula	0.200		
C. ehrenbergii	0.202	Wang et al. 2018	

Comparing the  $EC_{50}$  values from the different species on the table 4 with those obtained for *M. inermum*, this species is more sensitive to copper, and therefore when assessing environmental risks of metals to soils, based on freshwater species only, we are

likely underestimating the values. Therefore, it becomes clear the importance to ensure that these species from soil are used as bioindicators on the ecotoxicological assays, since the main purpose of these assays is to protect the soil biota.

Most of the soil herbicides are used in agriculture and enter the soil by direct application (Wegener et al. 1985). With the development of resistant crops and different formulations of herbicides it has become each time more important to assess the adverse effects of these pollutants on non-target organisms (Fayez et al. 2007). Complementing batteries of ecotoxicological tests with soil organisms, with microalgae, is of utmost importance since studies addressing the effects on this group, with important soil functions, are lower compared to studies on freshwater species (e.g. Ma et al. 2002; Lipok et al. 2010), soil invertebrates (e.g. Pereira et al. 2009; Caetano et al. 2016; Niemeyer et al. 2018) and terrestrial plants (e.g. Gavina et al. 2013).

Soil algae, can be particularly affected by chemicals intentionally applied to soils, such as pesticides in general and herbicides in particular, since they grow mainly on the soil surface. By other hand, since herbicides have low mobility and higher adsorption to soil components, they tend to stay in the upper layers of soil, ending up as being toxic to species, such as terrestrial microalgae and cyanobacteria (Wegener et al. 1985). A worldwide broad-spectrum herbicide, such as RoundUp ULTRA Max<sup>®</sup>, whose main component is GLY was tested for assessing the sensitivity of the terrestrial microalgae, *M. inermum* in liquid medium by measuring growth inhibition and,  $EC_{50}$  values were estimated. Once again, the results showed a clearly dose-dependent response at the range of concentrations of GLY tested (figure 11). The three independent tests were conducted and the validity criteria established by the OECD protocol were met with the species *M. inermum*. For GLY three  $EC_{50}$  values with a 95% confidence interval obtained were presented (table 2), and the repeatability of the test was confirmed as they fit in the same range of values. The toxic effect of GLY on the cellular growth was previously studied using different algal species (table 5), once again proving that the sensitivity to a toxic compound in microalgae is species-dependent.

Currently, the USEPA classifies the glyphosate formulations as low or non-toxic to moderately toxic for terrestrial ecosystems, based on toxicity data for the following taxonomic groups: birds, amphibians, mammals, terrestrial invertebrates and plants (Annett et al. 2014). However, there is a lack of information on the toxicity of glyphosate to soil microalgae.

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M. inermum       2         M. inermum       2         Scenedesmus obliquuos       80.         Chlorella vulgaris       55.         Pseudokirchneriella subcapitata <sup>b</sup> 55.         S. acutus       1	onfidence Interval         .94 [2.47; 3.40]         .72 [2.27; 3.17]         .89 [2.41; 3.38]         0 [47.63; 118.19]         62 [53.08; 57.56]         64.70         9.08 [8.4; 9.7]	Formulation RoundUp UltraMax®, 28.85% GLY Xnockdown 48 SL®, 48% GLY ATANOR®, 48% GLY RoundUp®, 360 g/L GLY Ron-do ®, 48%	Reference Present work Ermis et al. 2009 Romero et al. 2011 Cedergreen et al 2005	
M. inermum       2         Scenedesmus obliquuos       80.         Chlorella vulgaris       55.         Pseudokirchneriella subcapitata <sup>b</sup> 55.         S. acutus       1	.72 [2.27; 3.17] .89 [2.41; 3.38] 0 [47.63; 118.19] 62 [53.08; 57.56] 64.70	UltraMax®, 28.85% GLY Knockdown 48 SL®, 48% GLY ATANOR®, 48% GLY RoundUp®, 360 g/L GLY	Ermis et al. 2009 Romero et al. 2011 Cedergreen et al	
2       Scenedesmus obliquuos     80.       Chlorella vulgaris     55.       Pseudokirchneriella subcapitata <sup>b</sup> 55.       S. acutus     1	.89 [2.41; 3.38] 0 [47.63; 118.19] 62 [53.08; 57.56] 64.70	28.85% GLY Knockdown 48 SL®, 48% GLY ATANOR®, 48% GLY RoundUp®, 360 g/L GLY	Ermis et al. 2009 Romero et al. 2011 Cedergreen et al	
Scenedesmus obliquuos       80.         Chlorella vulgaris       55.         Pseudokirchneriella subcapitata <sup>b</sup> 55.         S. acutus       1	0 [47.63; 118.19] 62 [53.08; 57.56] 64.70	Knockdown 48 SL®, 48% GLY ATANOR®, 48% GLY RoundUp®, 360 g/L GLY	Romero et al. 2011 Cedergreen et al	
Chlorella vulgaris       55.         Pseudokirchneriella       subcapitata <sup>b</sup> S. acutus       1	62 [53.08; 57.56] 64.70	SL®, 48% GLY ATANOR®, 48% GLY RoundUp®, 360 g/L GLY	Romero et al. 2011 Cedergreen et al	
Chlorella vulgaris       55.         Pseudokirchneriella       subcapitata <sup>b</sup> S. acutus       1	62 [53.08; 57.56] 64.70	ATANOR®, 48% GLY RoundUp®, 360 g/L GLY	Romero et al. 2011 Cedergreen et al	
Pseudokirchneriella subcapitata <sup>b</sup> S. acutus	64.70	GLY RoundUp®, 360 g/L GLY	2011 Cedergreen et al	
Pseudokirchneriella subcapitata <sup>b</sup> S. acutus	64.70	RoundUp®, 360 g/L GLY	Cedergreen et a	
subcapitata <sup>b</sup> S. acutus		g/L GLY	e e	
S. acutus		3	2005	
S. acutus	0 00 [0 4: 0 7]	Ron-do ®, 48%		
S. acutus				
	9.00 [0.4, 9.7]	GLY	Saenz et al. 199	
C. saccharophila 4	0.2 [10.4; 11.2]	GLY, TC <sup>a</sup>		
	0.6 [36.7; 45.2]			
C. vulgaris 4	1.7 [37.5; 46.6]	97.5% GLY,	Vendrell et al.	
S. acutus 2	4.5 [21.9; 27.7]	TC <sup>a</sup>	2009	
S. subspicatus 2	6.0 [23.5; 28.9]			
C. vulgaris	4.69	95% GLY, TC <sup>a</sup>	Ma et al. 2002	

When we analyze table 5 it is possible to discern vast differences for the same species in what regards the EC<sub>50</sub> values obtained, which is visible for *C. vulgaris* and the different *Scenedesmus* sp. The differences in toxicity data recorded for herbicides could also be caused by differences in the chemical formulations tested in the different studies. Tsui et al. (2003), analyzed the differences in toxicity between the glyphosate and the commercial formulation, RoundUp® for *Selenastrum capricornutum*, a freshwater microalgae species, showing an EC<sub>50</sub> value of 24.7 [22.8;26.7] mg/L and 5.81 [2.36; 8.14] mg/L, respectively. Pereira et al. (2009), also studied the behavior for glyphosate as well as of its corresponding commercial formulation, determining an EC<sub>50</sub> value for *P. subcapitata* of 79 mg/L and 63 mg/L, respectively. Various types and amounts of adjuvant additives included in the formulated products have been found to improve GLY performance in different ways, playing an important role in the retention and absorption of GLY, significantly influencing the efficiency of the herbicide uptake (Travlos et al. 2017). The results on both studies clearly demonstrated that the ingredients of the commercial formulation enhanced the herbicide toxicity to microalgae, recognizing that the herbicidal activity of GLY is activated by the remaining ingredients.

The EC<sub>50</sub> estimated for the *M. inermum* in our study was must lower when compared with the values for the other test species presented in table 5, showing that this specie has a higher sensitivity to GLY in liquid medium than freshwater algae. *M. inermum* was approximately 23 times more sensitive when compared to *P. subcapitata*, to the same formulation tested. The high EC<sub>50</sub> values recorded led the authors of previous studies to conclude that glyphosate was not a hazardous herbicide for the aquatic ecosystem due to its low algae toxicity at low concentrations. Further the authors agreed that the probability of finding the concentrations tested in their study for which effects were recorded, in the environment are low (Vendrell et al. 2009). In summary the highest sensitivity of the soil microalgae tested in this work is a relevant argument for including this soil microalga species, on the battery of test species used for evaluating the risks of contaminants to soil biota.

# 5.3. Ecotoxicological Test with Soil BSC Microalgae using Artificial OECD Soil as a Test Substance

Considering that there are few standard test procedures with soil species in general and the lack of tests with microalgae soil species, in particular, it was considered important in the present work to develop a test procedure with soil microalgae, in more ecologically relevant exposure conditions i.e. by using the artificial OECD soil developed for standard ecotoxicological tests with soil organisms.

Once again was necessary to determine the growth curve for *M. inermum* in soil in order to established the time length for the upcoming ecotoxicological test. The data in this study showed that *M. inermum* needed a 13-day growth period to reach the exponential growth phase, which is longer comparing with the time length in liquid medium. With the

growth curve the validity criteria was established. The ecotoxicological test on soil uses the inhibition of growth rate as a biomarker of the toxic mode of action for the same contaminants previous tested in liquid medium. However, on soil, is not possible to use the count of cells as an endpoint. However, chlorophyll *a* content proved to be a good endpoint of algae development and its extraction proved to be reproducible, as demonstrated in the present work. In the present study we evaluated the toxic effect of copper (II) on the soil microalgae *M. inermum*, following growth inhibition, by measuring chlorophyll *a* content in the soil and the EC<sub>50</sub> values were estimated. The 13-day growth inhibition test on soil for *M. inermum* showed a clearly dose-dependent response for the range of concentrations of copper tested (figure 12). Three independent tests were conducted and the validity criteria established were met. For the copper three EC<sub>50</sub> values with the corresponding 95% confidence intervals obtained were presented (table 3), and the repeatability of the test was confirmed as they fit in the same range of values.

Comparing the EC<sub>50</sub> values recorded for *M. inermum* both in aqueous and in soil tests the results for copper were very similar. Since data for toxicity using soil microalgae species is limited, comparison on the sensitivity on soil is very challenging. A study by Kalinowska et al. 2010, compared the copper stress in *Stichococcus minor* and *Geminella terricola* isolated from Cu-polluted and unpolluted soils, reporting that *S. minor* accumulated lower amounts of copper (0,38 mM, which correspond to approximately 60 mg/L) than *G. terricola* (4,20 Mm, which correspond to approximately 670 mg/L). By comparing the toxicity of Cu to *G. terricola* and *S. minor* from unpolluted soils it was possible to conclude that *M. inermum* is much more sensitive. Both algae from this previous study were exposed to copper in aqueous medium and in our study, we exposed *M. inermum* to copper on soil, a more realistic and reliable approach.

Glyphosate was also tested on soil and the  $EC_{50}$  values were determined. The results also showed a dose-dependent response at the range of concentrations of GLY tested (figure 14). The three independent tests were conducted and the validity criteria established for soil were met. For GLY three  $EC_{50}$  values with a 95% confidence interval obtained were presented (table 3), and the repeatability of the test was confirmed as they fit in the same range of values. By comparing the  $EC_{50}$  values for GLY and for *M. inermum* on aqueous medium and on soil the results showed that on soil *M. inermum* was slightly less sensitivity. A study by Bérard et al. 2004, studied the algal community on the surface of soil, observing an apparent adaptation to the herbicide atrazine. However, after an atrazine application, only the most tolerant algae were able to grow in the soil, resulting in a more tolerant community to atrazine, despite the loss of biodiversity. Zabaloy et al. 2012, indicated that the exposure of soils to GLY caused minor changes on the microbial community structure and functions. When compared to pristine soils, soils chronically exposed to GLY didn't showed changes on microbial respiration, most likely due to the gradual elimination of GLY sensitive species and the selection of organisms acclimated for rapid mineralization of GLY. Since the information on the effects of GLY to soil microalgae is very limited, comparison is once again very difficult to make. However, reported studies point for the importance of testing the toxicity of contaminants directly with soil microalgae, and they can have impacts on the structural diversity of natural communities.

Analyzing the EC<sub>50</sub> values for both contaminants allow us, once again, to advocate the importance of ecotoxicological tests on soil, since testing in aqueous media can overestimate the risks because the role of the soil components (e.g. organic matter and clay) on the bioavailability of contaminants are not taken into account (Rieuwerts et al. 1998; Bradham et al. 2006; Takáč et al. 2009). The contaminants adsorb to the soil particles, making them less bioavailable to exert toxic effects on biota. With the limited data on soil microalgae toxicity on soil the results of this study came to reinforce that the soil microalgae group are in fact affected by the contaminants applied and found on soil and we can strongly evidence that the new guideline developed in the present work is one more step on the reliable assessment of the impact of contaminants on soils.

## 6. Conclusions

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The results in this work came to reinforce that soil microalgae should be considered in ecotoxicological evaluations as soil bioindicators, in the evaluation of the impact of different contaminants to the biota of this environmental compartment. *M. inermum* clearly showed a growth under laboratory conditions in liquid medium and artificial soil. The results also established the repeatability of the ecotoxicological tests, demonstrating the vulnerability of the soil microalga to the toxicity of both contaminants. The use of these bioassays may crucial impact on the reliability of future predictions of soil quality guidelines, as they will contribute for protecting species with relevant functions on soils protection.

However, for introducing these ecotoxicological tests in routine evaluations further research should be considered. The next steps should focus on repeating the tests in the artificial soil with other chemical compounds, to reinforce the repeatability of the procedure and the appropriateness of the validity criteria proposed. Further, testing natural contaminated soils, should also be made, in order to evaluate the potential of this ecotoxicological assay for the risk assessment of contaminated sites. Contaminated soils are challenging since when added to these soils, soil microalgae will have to compete with other species which are part of the microbial community of natural soil. Although soil sterilization may be considered for inorganic contamination, the role of sterilizing procedures in the bioavailability of contaminants in general and in the degradation of organic contaminants in particular needs to be addressed.

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# Annex A

### BG<sub>11</sub> Medium

Reference: Rippka R, Herdman H (1992) Pasteur culture collection of cyanobacteria catalogue & taxonomic handbook. 1. Catalogue of strains. Institut Pasteur, Paris.

- Use autoclave destile water.
- Add 1 mL of each stock solution (1-6) to a liter of the final solution.
- Store all stock solutions in the refrigerator.
- For solid BG<sub>11</sub>, agar should be added in a concentration of 1g to 100 mL.

able 1. Composition of sto	ock solutions	
Stock solutions	Reagent	Grams per 100 mL of distilled water
1	K <sub>2</sub> HPO <sub>4</sub>	4,00
2	MgSO <sub>4</sub> .7H <sub>2</sub> O	7,50
3	CaCl <sub>2</sub> .2H <sub>2</sub> O	3,60
	Citric Acid	0,60
4	Ferric ammonium citrate	0,60
	EDTA (dissodium salt)	0,10
5	Na <sub>2</sub> CO <sub>3</sub>	2,00
	H <sub>3</sub> BO <sub>3</sub>	0,2860
	MnCl2.4H <sub>2</sub> O	0,1810
6	ZnSO <sub>4</sub> .7H <sub>2</sub> O	0,0395
"Trace metal mix"	Na <sub>2</sub> MoO <sub>4</sub> .2H <sub>2</sub> O	0,0390
	CuSO <sub>4</sub> .5H <sub>2</sub> O	0,0079
	Co(NO <sub>3</sub> ) <sub>2</sub> .6H <sub>2</sub> 0	0,0049
7	NaNO <sub>3</sub>	a)

a. The volume of stock solution 7 is 10 mL/L;

### Woods Hole MBL Medium

Reference: Nichols, H. W. (1973) in *Handbook of Phycological Methods*, Ed. J. R. Stein, pp. 16-17. Cambridge University Press.

Table 2: Composition of stock solutions	
Stock solutions	Grams per liter distilled water
1. CaCL <sub>2</sub> .2H <sub>2</sub> O	36,76
2. MgSO <sub>4</sub> .7H <sub>2</sub> O	36,97
3. NaHCO <sub>3</sub>	12,60
4. K <sub>2</sub> HPO <sub>4</sub>	8,71
5. NaNO₃	85,01
6. NaSiO <sub>3</sub> .9H <sub>2</sub> O	28,42
7. Na <sub>2</sub> EDTA	4,36
8. FeCl <sub>3</sub> .6H <sub>2</sub> 0	3,15
9. Metal Mix <sup>1</sup>	
CuSO <sub>4</sub> .5H <sub>2</sub> O	0,01
ZnSO <sub>4</sub> .7H <sub>2</sub> O	0,022
CoCl <sub>2</sub> .6H <sub>2</sub> O	0,01
MnCl <sub>2</sub> .4H <sub>2</sub> O	0,18
Na <sub>2</sub> Mo <sub>4</sub> .2H <sub>2</sub> O	0,006
10. Vitamin stock	
Cyanocobalamin (B12)	0,0005
Thiamine HCL (B1)	0,10
Biotin	0,0005
11. Tris stock	250,0

(1) Add each constituent separately to ~750 mL of distilled water, fully dissolving between additions. Finally make up to 1 L with distilled water.

(2) Store all stock solutions in the refrigerator

To prepare MBL medium

- Add 1 mL of each stock solution (1-11) to 1 L distilled water (for species which cannot use nitrate substitute, 1 mL of NH<sub>4</sub>Cl made up to 5,4 g/L H<sub>2</sub>O).
- Adjust pH to 7.2 with HCL.
- Autoclave at 121 °C (15PSI for 15 min).

## Annex B

#### Sequences from the DNA fragments of Micractinium inermum

#### >4.3II NS7 LR1850+LR1850 1297

 ${\tt tctgtttagccttagagggatttacccctgctttgggctgcattcccaaacaacccgactcttcgaaagc}$ gtctcgtggagcctcaggatccagtcccaacggggttctcaccctctctgacgcccctttccaggggact tggggccggactgaggcagaggagcacttctatagactacaattcgccagccggaggctggagattttcaaatatqcttaaqttcaqcqqqtaqtcttqcctqaqctcaqqtcqaaaqtqtqaqacattcqqqqqtqqcaqaaccaccccgaattcctgctaggccgccagcaaagtcccctcggacaacgacaggctgtgtagcgatgcta $\verb|cctaccaagtcattgccctgcaaacggggtccatgctcaagcctctacacttcagccgacccggactcat||$ cgatgaatcgaagaggagccgggaaggccaggtccgttctatccactccaaaggagagggagagcgaggg tqtaaqccqacqctqaqqcaqacatqccctcqqccqaaqcctcqqqcqcaatttqcqttcaaaqattcqa tggttcacqgaattctgcaattcacactacqtatcqcatttcqctqcqttcttcatcqatacqqqaqcca agatatccgttgttgagagttgtctttggttagagtggggcaccgagatgcacacactagacagctttagtttaagtttaggtttggggttgggttgagttgaatataattaccgaccctacagcccagggggccccgagg ggacacccgcgggggtaaggcccgccggacctgaactgcggtcagaagaccccagacaggggaccgacgc cacctacqqaaaaccttqttacqacttctccttcctctaqqtqqqaqqqtttaatqaacttttcqqcqqct qaqaqcqqaqaccqcccccaqtcqccaatccqaacactttaccaccacacccaatcqqtaqqaqcqacqq gcqqqqqqtacaaaqqqcaqqqacqtatcaacqcaaqctqatqacttqcqcttactaqqcattcctcqtt gaaaataaaaattgcataatcattcccatcacgatgcatttaaaaatacccggggcctttcggccagggta acttcttgatggataaggaacggccgtggggccaaaa

#### >4.3II NS7 LR1850+NS7m 1217

ttggggcgcaccgcgcgctacactgatgcattcaaccgagcttagccttggccgagaggcccgggtaatctttgaaactgcatcgtgatggggatagattattgcaattattaatcttcaacgaggaatgcctagtaagcgc aagtcatcagcttgcgttgattacgtccctgccctttgtacacaccgcccgtcgctcctaccgattgggt gtgctggtgaagtgttcggattggcgactggggggggtctccgctctcagccgccgaaaagttcattaaa tcgatcgaatccactctggtaaccaaccgtcccctcgccctggtgcgagcgtcggtcccctgtctggggt  $\verb"cttctgaccgcagttcaggtccggcgggccttacccccacgggtgtcccctcggggccccctgggctgta"$ gggtcggtaattatattcaactcaacccaccccaaacctaaacttaaactaaagctgtctagtgtgtgca tctcqqtqccccactctaaccaaaqacaactctcaacaacqqatatcttqqctcccqtatcqatqaaqaa cgcagcgaaatgcgatacgtagtqtgaattgcagaattccgtgaaccatcgaatctttgaacgcaaattg cgcccgaggcttcggccgagggcatgtctgcctcagcgtcggcttacaccctcgctctccctttg gaqtqqataqaacqqacctqqccttcccqqctcctcttcqattcatcqatqaqtccqqqtcqqctqaaqt gttgtccgaggggactttgctggcggcctagcaggaattccggggtggttctgccaccccgaatgtctca ${\tt cactttcaacctgagctcaggcaagactacccgctgaacttaagcctatcaatgagccggaggaaaagaa$ actaactaggatgcccttactaacggcgaacgaaccgggcaaagctatcttgaaaaaactccagcctccgg ctggggaatgtagtctataaaattgctctctgcctcagtccggccccaattccctggaaaagggtcttcg aaaaggtgaaaaccccctttgtactgt

#### >4.31 NS1 +18L 1113

tcccccggaccaaaactttgatttctcataaggtgccggcggagtcatcgaagaaacatccgccgatccc tagtcggcatcgtttatggttgagactaggacggtatctaatcgtcttcgagccccaactttcgttctt gattaatgaaaacatccttggcaaatgctttcgcagtagttcgtctttcataaatccaagaatttcacct ctgacaatgaaatacgaatgcccccgactgtccctcttaatcattactccggtcctacagaccaacagga taggccagagtcctatcgtgttattccatgctaatgtattcagagcgtaggcctgctttgaacactctaa tttactcaaagtaacagcgtcgactccgagtccggacagtgaagcccaggagcccgtccccgacaacaa ggtgggccctgccagtgcacaccgaaacggcggaccggcaggcccgccgccagaattccact ttaactgcagcaacttaaatatacgctattggagctggaattaccgcggctgctgctggaccagagttt ttaactgcagcaacttaaatatacgctattggagctggaattaccgcggctgctgcagaactgccc tccaattgatcctcgttaaggggtttagattgtactcattccaattaccagacctgaaaaggcccagta 

#### >4.31 NS1 18L+NS1 1096

 ${\tt caggtctagtataactgctttatactgtgaaactgcgaatggctcattaaatcagttatagtttatttga}$ tggtacctactccggatacccgtagtaaatctagagctaatacgtgcgtaaatcccgacttctggaag ggacgtatttattagataaaaggccgaccqqqctctqcccqactcqcqqtqaatcatqataacttcacqa atcgcatggccttgtgccggcgatgtttcattcaaatttctgccctatcaactttcggtggtaggataga ggcctaccatggtggtaacgggtgacggaggattagggttcgattccggagagggggggcctgagaaacggc  $\verb+taccacatccaaggaaggcagcaggcgcgcaaattacccaatcctgacacagggaggtagtgacaataaa$  ${\tt taacaatactgggccttttcaggtctggtaattggaatgagtacaatctaaaccccttaacgaggatcaa}$ ttggagggcaagtctggtgccagcagccgcggtaattccagctccaatagcgtatatttaagttgctgca gttaaaaagctcgtagttggatttcgggtggggcctgccggtccgccgtttcggtgtgcactggcagggc $\verb|ccaccttgttgtcggggacgggctcctgggcttcactgtccgggactcggagtcgacgctgttactttga||$ gtaaattagagtgttcaaagcaggcctacgctctgaatacattagcatggaataacacgataggactctg gcctatcctgttggtctgtaggaccggagtaatgattaagagggacagtcgggggcattcgtatttcatt gtcagaggtgaaattcttggatttatgaaagacgaactactgcgaaagcatttgccaaggatgttttcat $\verb+taatcaagaacgaaagttgggggctcgaagacgattagataccgtcctagtctcaaccataaacgatgcc$ gactagggatcggcggatgtttcttcgatgactccgccggcaccttatgagaaatcaaagtttttgggtt ccgggggggggtatggtcgcaaggctgaaacttaaaaaaattgacgg