

# **Carbon nanotubes alter agrosystem multifunctionality**

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25 CNT exposure also led to a significant decrease of  $H_2O$  flux in the system. As a result, the 26 multifunctionality of the agrosystem was significantly decreased at 0.1 mg.kg $^{-1}$ . Structural equation 27 modeling suggested that CNT impact on bacteria population in general, and on bacteria implied in 28 denitrification and  $CO<sub>2</sub>$  emission were the main factors influencing the multifunctionality index.

#### 29 **1. Introduction**

30 Nanotechnology feeds the hopes of several research domains due to its many potential applications. 31 Indeed, nanotechnology could be an important lever to the convergence of knowledge and 32 technology for the benefit of the society and could represent an opportunity for progress in the 33 twenty-first century<sup>1</sup>. To develop nanotechnology, the production and use of nanomaterials have 34 increased over the years<sup>2</sup>. Among nanomaterials, particular attention from industries is focused on 35 carbon nanotubes (CNTs) because of their remarkable physical, electronic and chemical properties  $^3$ , 36 and mainly because they can combine several of them. Consequently, CNTs are now found in 37 products such as electronics, tires or sport equipment  $4$ , and subsequently end up into the 38 environment during their journey from production to recycling  $5$ . In addition, they tend to 39 accumulate in sewage sludge used as fertilizer in agriculture <sup>5</sup> and modelling predicts that 40 concentration of carbonaceous nanomaterials in biosolids could reach up to 10 mg.kg<sup>-1</sup> while in 41 antural soil it could be up to 0.01 mg.kg<sup>-1 6</sup>. Furthermore, CNTs might be used as nano-fertilizer as 42 their application on plants (for biofuel applications) and crops leads to increased yield according to 43 several studies  $^{7,8}$ . Thus, their concentration in ecosystems, in particular in agrosystems, will possibly 44 increase in the future and their effects on this type of ecosystems need to be further examined in 45 order to ensure a sustainable use of CNTs.

According to the literature, CNTs can be taken up by plants<sup>7,9,10</sup> and have contrasted impacts on 47 plants <sup>7</sup>. They can act on plants from the germination stage with an activation of this process  $^{11-14}$ . 48 Furthermore, upon exposure, plant growth was reported to either increase, decrease, or even 49 unmodified (based on root systems, biomass production, and number of flowers and fruits)<sup>8,15</sup>. When 50 toxicity was detected, it was mostly associated with an overproduction of reactive oxygen species 51 (ROS), antioxidant enzymatic activities, necrotic lesions of cells and leaf tissues as well as root and 52 leaf morphological changes, decreased chlorophyll, and programmed cell death  $16-19$ . These divergent 53 results might be related with the influence of confounding factors which have been understudied so 54 far such as plant species  $^{20}$  or soil type  $^{21}$ . Additionally, CNTs can also influence gas exchanges of 55 plants with their environment. For example, CNT exposure at 250 mg. $L^{-1}$  increased carbon (C) assimilation rate of *Zea mays* and also decreased stomatal conductance and transpiration rate 57 possibly as a result of H<sub>2</sub>O use modulation  $^{22}$ . The same results were observed on *Orthosiphon stamineus* with a decrease in transpiration rate, stomatal conductance and net photosynthesis following a dose-dependent relationship with the increase of CNT concentration (0, 700, 1400 and  $\,$  2100 mg.L<sup>-1</sup>)<sup>23</sup>. Plants were also water-stressed as demonstrated by the positive correlation between 61 water use efficiency and CNT concentration  $^{23}$ . At lower concentrations (25 mg.L<sup>-1</sup> and 54 mg.L<sup>-1</sup>), no effects of CNT exposure were observed on *Arabidopsis thaliana* on ambient photosynthetic rate, stomatal conductance, transpiration, intrinsic water use efficiency and instantaneous water use 64 efficiency , It is also worth mentioning that these reported studies were performed in hydroponics which is not a realistic exposure scenario for agrosystems; usually not taking into account interactions with other biotic and abiotic components of the environment.

 In parallel, interactions of CNTs with micro-organisms have also been investigated. Microscopy observations have evidenced that CNTs can interact with bacterial cell membranes; they can disrupt 69 integrity of the membrane structure by oxidative stress and/or physical damage  $25-28$ . In addition, most studies have reported a decrease in soil microbial activity following exposure to CNTs and a loss 71 of viability  $28-32$ . Furthermore, CNT and other carbon nanomaterial application can also cause changes 72 in microbial community structure  $29,32-34$ . Finally, microbial functions such as organic matter 73 decomposition, nitrogen (N) cycle and  $CO<sub>2</sub>$  fixation could be modified  $35-38$ .

 From this brief overview of the literature, we can see that if the impact of CNTs in single-organism 75 exposure condition has been studied so far, very little has been done at the ecosystem scale  $39,40$ . However, having a functional overview of contaminant impacts at this scale is of tremendous importance as plant/soil micro-organism interactions drive multiple processes and functions 78 supporting ecosystem services such as food production . In particular, aboveground-belowground

 interactions drive biogeochemical cycles (C, N, P) through several processes such as photosynthesis, mineralisation, nitrogen fixation, nitrification, denitrification, among others. One way to tackle this 81 gap of knowledge is the use of multifunctionality index as developed in ecology  $42-44$ . This score 82 provides an easily interpretable measure of the ability of different entities composing an ecosystem (bacteria and plants, for example) to sustain multiple functions simultaneously by gathering the 84 selected functions in an index .

85 The objective of this study was thus to investigate the impacts of CNTs at both the organism (crop plant) or community (micro-organisms) scales and at the microcosm scale by implementing a 87 multifunctionality score. More specifically, after exposure to CNTs their toxicity was assessed on (1) the abundance of microbial communities and in particular those related to the N cycle by quantifying marker genes by qPCR, (2) the organic matter decomposition process using tea bags, (3) the plant growth (height, biomass, foliar surface area, dry matter content), (4) the plant physiology (photosynthetic pigment and secondary metabolite concentrations by spectrophotometry and plant 92 mineral nutrition) and (5) the greenhouse gas exchanges of the system ( $CO<sub>2</sub>$ , H<sub>2</sub>O, CH<sub>4</sub>). From these different variables, the impact of CNTs on ecosystem functioning was determined by an integrative approach based on multifunctionality scores and, structural equation modeling was used to better understand the relationships among the different drivers and multifunctionality.

#### **2. Material and methods**

#### **2.1. CNT synthesis and characterization**

 CNTs were synthesized at the Interuniversity Center of Materials Research and Engineering (CIRIMAT, Toulouse University). More information is available in Supporting Information.

 CNTs were characterized using transmission electron microscopy (JEOL 1400F, operated at 120 kV,), Raman spectroscopy (Labram HR 800 Yvon Jobin equipped with a He laser at 633 nm) and BET

 method (on freeze-dried samples). They were oxidized multiwalled CNTs (MWCNTs) with length 104 varying from 1 to 20  $\mu$ m, an average diameter of 22.5  $\pm$  5.0 nm (Figure S1) and a specific surface area 105 of 60 m<sup>2</sup>.g<sup>-1</sup>. CNTs were suspended in deionised water at 100 mg.L<sup>-1</sup> and homogenized for 15 min in an ultrasonic bath before use.

#### **2.2. Soil characterization and contamination**

 The experiment was performed with a sandy soil (LUFA-Speyer 2.1) sieved to 2 mm with 86.0% of sand, 12.6% of loam and 1.4% of clay. It contained 0.58 wt. % of organic carbon, 0.04 wt. % of 111 nitrogen, with a pH of  $5.0 \pm 0.1$  (H<sub>2</sub>O extraction according to NF ISO 10390:2004), a water retention 112 capacity of 30g/100g and a cation exchange capacity of  $2.9 \pm 0.2$  meq.100g<sup>-1</sup>. More details on soil characteristics are available in supporting information (Table S1). Soils were contaminated by the addition of the required volume of CNT suspension to reach a final concentration of 0.1 or 10 mg.kg<sup>-1</sup> dry soil while maintaining the water holding capacity at 25%; the first concentration being close to 116 environmentally predicted concentrations <sup>6</sup> and the second one representing a worst-case scenario (use of nano-fertilizer, for instance). They were then mixed manually for 10 min every day for a week 118 to ensure that CNTs were homogeneously distributed in the soil.

#### **2.3. Microcosm set-up**

121 Three hundred grams of soil were distributed in each microcosm ( $L = 45$  cm,  $D = 10$  cm). A tea-bag 122 (Clipton green tea sencha, 1.13  $\pm$  0.32 g) was introduced after the addition of 50 g of soil in each microcosm to study organic matter decomposition. Prior to introduction, tea bags were infused in water at 90°C to remove tea molecules which could have induced bias in microbial community behaviour, dried at 70°C during 48 h and weighted. One seed of *Zea mays* L*.* subsp. mays was added, 126 plants were watered *ad libitum*. For each condition (CNT concentration of 0, 0.1 and 10 mg.kg<sup>-1</sup>),  microcosms were replicated five times with a total of 15 microcosms. The experiment lasted for 6 weeks in an environmental chamber with controlled parameters (10 h/14 h day/night cycle, 129 24°/18 °C and a hygrometry rate of 85%).

#### **2.4. Bacterial abundance, functional genes and litter decomposition**

 The bacterial abundance (16S rRNA) and the abundance of selected genes indicators for N cycling 133 (nitrification and denitrification) were analysed by qPCR<sup>46</sup>. To summarize, after DNA isolation from contaminated soil using the extraction DNeasy PowerSoil kit ©QIAGEN, the total abundance of soil bacteria was measured by targeting the universal 16S rRNA gene. The following genes related to N cycle were quantified: ammonia monooxygenase A gene (*amoA*) of ammonia oxidizing archaea (AOA) and ammonia oxidizing bacteria (AOB), nitrite reductase (*nirK*) and nitrous oxide reductase (*nosZ*) 138 genes. The same experimental procedure as described in Vijayaraj *et al.* <sup>47</sup> was used. Results were expressed relative to the initial gene abundances at the beginning of the experiment to account for initial variability among microcosms (Figure S2; absolute results after exposure are also available in Figure S3).

 To study litter decomposition (C cycle), tea bags were removed and weighted after the 6 weeks of exposure which permitted to calculate a decomposition average rate as follows:

#### D M D

 where Mi and Mf represent the initial and final mass (mg dry weight) of the tea bag and the duration of exposure is expressed in days.

**2.5. Plant morphology and metabolism**

 Upon exposure, indicators of the C cycle and biomass production were gathered through the determination of the effective quantum yield of maize plants (Diving-Pam underwater fluorometer, Walz, United Kingdom), the height, the fresh and dry biomasses and the leaf surface area (using Image J software). Part of the leaf biomass was oven-dried (60°C until constant weight) to determine elemental concentrations by ICP-AES (see section 2.7). The remaining biomass was used fresh for methanol-based extraction to quantify photosynthetic pigments (chlorophyll a and b) and secondary 154 metabolites *i.e.* total phenolic compounds, flavonoids and tannins as described in Leroy et al. <sup>48</sup> More details on the phytotoxicity markers are available in SI.

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#### 157 **2.6. Greenhouse gas exchange measurements and analysis**

158 When plants were sufficiently high (*i.e.* sixth week), H<sub>2</sub>O, CO<sub>2</sub> and CH<sub>4</sub> exchanges were recorded 159 every second for 3 minutes. To do so, an airtight flux chamber (V = 0.06 m<sup>3</sup>) was connected to a trace 160 gas analyser (LI-7810, ©LI-COR inc, United States).

161 Net ecosystem exchange of  $CO<sub>2</sub>$  (NEE) was measured using a transparent chamber and a LED light 162 while ecosystem  $CO<sub>2</sub>$  respiration (RECO) was measured using a darkened chamber. Gross primary 163 productivity (GPP: CO<sub>2</sub> uptake as a result of photosynthesis), was calculated as the difference 164 between NEE and RECO.

165 The slope of gas concentration was calculated and used to determine CO<sub>2</sub> flux rates ( $\mu$ mol.m<sup>-2</sup>.h<sup>-1</sup>). 166 using R software and the "flux"  $49$  and "gasfluxes"  $50$  packages. For NEE, calculations used an 167 exponential nonlinear function to calculate the change in the concentration in the chamber over time <sup>51</sup>. RECO was calculated using linear regression of gas concentrations in the chamber over time. The 169 same was done with H<sub>2</sub>O and CH<sub>4</sub> to obtain fluxes in presence of light and in the dark. Positive values 170 for the  $CO<sub>2</sub>$  and CH<sub>4</sub> fluxes indicate C release from the system while negative values indicate C uptake 171 by the mesocosm.

#### **2.7. Sample preparation and chemical analysis by ICP-AES**

174 Plant samples were digested using 1 ml of a mix of HNO<sub>3</sub> and HCl (3:1 v/v) and 1 ml of H<sub>2</sub>O<sub>2</sub> in a close 175 vessel in a microwave system (MARS 2, ©CEM corporation, USA: ramp time: 20-25 min; holding 176 time: 10 min; temperature: 190 °C). Digested samples were then diluted in 5% HNO<sub>3</sub> before analysis on an ICP-AES (ARCOS FHX22, ©AMETEK Spectral, United States) along with control samples (blanks with only chemicals and standard reference material NIST SRM 1570a: Trace Elements in Spinach Leaves).

#### **2.8. Statistical analyses**

 For statistical analyses, 1-way ANOVAs were performed on the toxicity endpoints to test the effects of the addition of CNTs. The normality, independence and homoscedasticity of the residues were checked using Shapiro, Durbin-Watson and Breusch-Pagan tests, respectively. Data were transformed to log or square root if one of the three conditions of validity was not met. A non- parametric Kruskal-Wallis test was performed in the case of failed transformations. All statistical analyses were performed on RStudio software (version 3.4.1). Principal component analyses (PCA) were performed on the elemental concentrations in plants to assess the impact of CNTs on plant nutrition.

 Different functions with corresponding indicators were selected to be representative of the two main 191 types of functions provided by agroecosystems: ecological functions and nutritional functions<sup>52</sup>. Ecological functions of agroecosystems include fluxes of energy and nutrients such as primary production, decomposition, and biological fixation by plants. These functions are basic ecosystem processes but are key to defining the global functioning of the system. As an extension of these basic functions, nutritional functions have to be measured alongside their ecological counterparts as

 agroecosystems' primary goal is food production. Nutritional functions encompass a broad suite of functions such as crop production, nutrient quality or resistance to environmental stress. In our case, we gathered the functions and variables as follows: N cycle including both nitrification (with *amoA*- AOA and *amoA*-AOB gene abundances as indicators) and denitrification (*nirK, nosZ*), C cycle including litter decomposition, ecosystem respiration, methane emission, and gross primary production 201 (related to bacterial abundance, quantum yield, chlorophyll, crop biomass), H<sub>2</sub>O cycle (plant dry matter content), plant nutrition (Z-score based on all elements quantified by ICP-AES), and plant 203 resistance (total phenolic compounds and tannins). Using the packages "dplyr" <sup>53</sup> and "tidyverse" <sup>54</sup> 204 as well as the code library "mulitidiv" , the data were standardized and implemented in a unique Z- score, an index representing the multifunctionality of the ecosystem (Allan et al., 2015, Delgado- Baquerizo et al., 2016). A negative Z-score indicates a low multifunctionality and a positive Z-score a high multifunctionality of the considered agrosystem.

208 We then combined CNTs, the multifunctionality of the agrosystem and its drivers to determine which function was influenced most by CNTs and then drove a change in the multifunctionality of the agrosystem. Specifically, we constructed a structural equation model (SEM) using the package 211 "Iavaan" <sup>56</sup>, which draws from a hypothesis-driven *a priori* model based on agrosystem theory and accounting for our expectation that individual agrosystem functions may respond either positively or negatively to CNTs, and therefore differently influence agrosystem multifunctionality. We considered all available variables to investigate how individual indicators of each function related to agrosystem multifunctionality (that is the full model). Then, the full model was simplified by step-wise exclusion of variables with non-significant weights and/or significant collinearity, as estimated by AIC and chi-217 squared statistics, until a minimal adequate model showing specific linkages remained . We diagnosed model fits using chi-squared statistics (*P* > 0.05), root-mean-square error of approximation index (RMSEA ≤0.1), standardized root-mean-square residual index (SRMR ≤0.1) and comparative fit 220 indices (CFI  $\geq$  0.95), and included variables and paths in the final model based on chi-squared statistics (*P* < 0.05) and AIC values of the model.

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235 other conditions ( $p_{\text{litter decomposition}}$  < 0.01) with on average 9.1 mg.j<sup>-1</sup> vs. 7.8 mg.j<sup>-1</sup> (Figure 1B).

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#### 238 **3.2. Crop plant response to CNT exposure**

239 At the end of exposure, the proportion of dry weight was higher for plants exposed to 10 mg.kg<sup>-1</sup> of 240 CNT with on average +17.4% compared to control plants ( $p_{DWFW}$  < 0.01, Figure 2A). No significant 241 difference was found on the other growth parameters tested: height, fresh weight (FW) and leaf area 242 with on average 54.23 cm, 3.54 g and 126.34 cm<sup>2</sup>, respectively whatever the condition ( $p_{height} = 0.7$ , 243 *pFW* = 0.6, *pleaf* area = 0.5; Figure 2 B-C-D).

244 At the metabolic level, no difference on the quantum yield was detected between maize grown on 245 control soils and those exposed to CNTs (*pquantum yield* = 0.3, Figure 3A). However, the concentration of 246 total chlorophyll was on average 34% and 9% lower in maize exposed to 0.1 and 10 of mg.kg<sup>-1</sup> CNT 247 than in maize in control condition (*pChlorophyll* < 0.01, Figure 3B). Furthermore, phenolic compounds

248 decreased by 15% when exposed to 0.1 mg.kg<sup>-1</sup> of CNT compared to control condition ( $p_{\textit{phenolic compounds}}$ 249  $\leq$  0.01, Figure 3C). Similarly, maize exposed to the two CNT concentrations (0.1 and 10 mg.kg<sup>-1</sup>) 250 decreased by 18% their tannin content in leaves compared to control soils (*ptannin* = 0.01, Figure 3D).

 CNT exposure impacted significantly plant mineral nutrition. Indeed, the PCA on maize leaf elemental composition (Ca, Cu, Fe, Mg, Mn, Na, P, S, Zn) after exposure suggested that plants exposed to 253 0.1 mg.kg<sup>-1</sup> CNT were depleted in Fe but enriched in P, K while plants exposed to 10 mg.kg<sup>-1</sup> were enriched mainly in S; both exposure concentrations led to plant enrichment in Mg and Zn (Figure 3E, Table S2).

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#### 257 **3.3. Greenhouse gas exchange**

258 At the  $6<sup>th</sup>$  week of exposure, the NEE, RECO and CH<sub>4</sub> (in light and dark conditions) were similar 259 whatever the CNT concentration with on average 0.04 and 0.07  $mg_{CO2}$ . $m^2$ . $h^2$  and -0.004 and -0.002 260 mg<sub>CH4</sub>.m<sup>-2</sup>.h<sup>-1</sup> respectively ( $p_{NEE}$  = 0.6,  $p_{RECO}$  = 0.2,  $p_{CH4~Light}$  = 0.4,  $p_{CH4~darkness}$  = 0.9 Figure 4A-B-E-F). 261 Additionally, the GPP was also similar whatever the CNT concentration with on average -0.16 262  $mg_{\text{co2}} \cdot m^{-2} \cdot h^{-1}$  ( $p_{\text{GPP}}$  = 0.7, Figure S4). In contrast, the emission of H<sub>2</sub>O in microcosms measured with 263 light (*ie* the evapotranspiration gathering both leaf transpiration and soil evaporation) was around 264 twice lower in the presence of CNTs at the two concentrations studied than in controls with on 265 average 16.4 and 37.2 mg<sub>H2O</sub>.m<sup>-2</sup>.h<sup>-1</sup>, respectively ( $p_{H2O}$  light < 0.01, Figure 4C). This was not the case 266 without light: the emission of H<sub>2</sub>O was on average 10.5 mg<sub>H2O</sub>.m<sup>-2</sup>.h<sup>-1</sup> whatever the CNT 267 concentration  $(p_{H2O \text{ dark}} = 0.16, \text{ Figure 4D}).$ 

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#### 269 **3.4. Multifunctionality and structural equation modeling**

270 The agrosystem multifunctionality significantly decreased from 0.18  $\pm$  0.10 in control condition and

271 0.17  $\pm$  0.22 at 10 mg.kg<sup>-1</sup> CNT to -0.35  $\pm$  0.14 with the addition of 0.1 mg.kg<sup>-1</sup> (p<0.001, Figure 5A).

272 The SEM model revealed that CNT addition impacted the multifunctionality of the agrosystem mostly indirectly (path = 0.193, p > 0.05; Figure 5B). In particular, multifunctionality was altered through the 274 impact of CNTs on microbial properties which impaired the C and N cycles. Indeed, CNTs directly increased bacterial abundance (path = 0.420), which in turn positively influenced *amoA*-AOA (path = 276 0.773) and *nirK* (path = 0.705) gene abundances, as well as  $CO<sub>2</sub>$  respiration (path = 0.519) and organic matter decomposition (path = 0.349) rates. Increasing nitrification (*amoA*-AOA, path = 0.888), 278 denitrification to a lower extent ( $nirK$  gene abundance; path = 0.285) and, CO<sub>2</sub> respiration (path = 0.552) then positively and significantly influenced agrosystem multifunctionality. CNTs also directly increased the ratio between dry and fresh plant weight (path = 1.155), which in turn negatively 281 influenced methane emissions (path = -0.493) but positively influenced H<sub>2</sub>O fluxes (path = 0.682) and decomposition (path = 0.622). Additionally, CNT also directly impacted other plant parameters such as plant nutrition with a positive influence on mineral content (path = 0.516) or on plant defense with a negative influence on tannins (path = -0.621) or on water cycle overall (path = -1.327). Plant related parameters contributed to a lesser extent to the multifunctionality with either non-significant paths or paths <0.4.

**4. Discussion**

#### **4.1. Impacts of CNTs on C cycle**

291 Organic matter decomposition was 17% faster with the addition of 10 mg.kg<sup>-1</sup> of CNT compared to control microcosms. This increase was related to the increase in the relative number of universal 16S rRNA gene copies (+82%) showing an increase in bacterial density at this same condition (Figure 5B).

 In parallel, plant photosynthesis did not seem to be impacted (slight trend of a decrease in the quantum yield of exposed maize, although not significant) even though there was a significant  decrease in total chlorophyll pigments for plants exposed to CNTs. Plant growth (fresh biomass production and height) was also not impacted after 6 weeks of exposure.

 In the literature, it has been reported in different articles that nanomaterials can increase organic 299 matter decomposition process  $58,59$ . A study with similar exposure duration (56 days) highlighted the same increase in 16S rRNA copies in the presence of CNTs (both single-walled CNTs and multi-walled S01 CNTs, between 0.7 and 2.6% O) at 10 and 50 g.kg $^{-1}$  <sup>60</sup>. Although, there was no significant differences 302 in 16S RNA gene copy between the control condition and the soil exposed to 0.1 mg.kg<sup>-1</sup> CNT after the 6 weeks of exposure; in this last condition, there was a significant decrease of gene copy between the beginning and the end of exposure (T-test; p=0.041) suggesting toxicity at this lower concentration. This higher toxicity of CNTs at lower concentrations has already been evidenced with 306 similar reduction of microbial biomass after exposure to carbon nanomaterials such as graphene  $32,61$ , 307 graphene oxide  $32,61,62$  and CNTs  $29,32$ . The homo and hetero-agglomeration of CNTs in soil might 308 explain this effect . Indeed, when the ionic strength of the medium and/or the CNT concentration increase, CNT tend to agglomerate and become less mobile and less prone to interactions with other environmental components <sup>63</sup>. Despite the decrease in bacteria concentration over time at 0.1 mg.kg<sup>-</sup> 311  $\,$ <sup>1</sup> of CNTs, no effect on decomposition potential was noticeable suggesting that decomposition was maybe compensated by others decomposers.

 Interestingly, in our exposure conditions the difference detected at the organisms and community levels (bacteria population and chlorophyll pigments) did not lead to any significant difference in gas 315 exchange ( $CO<sub>2</sub>$  and  $CH<sub>4</sub>$ ) at the microcosm level, suggesting some compensation mechanisms at this scale.

#### **4.2. Impacts of CNTs on N cycle**

 The quantification of genes involved in the nitrogen cycle showed an increase in *amoA*-AOA gene 320 copies up to 144% in soils exposed to 10 mg.kg<sup>-1</sup> of CNT compared to the control. A similar trend was

 noticeable on *amoA*-AOB gene copies, suggesting a dose-dependent induction of the nitrifier bacterial communities. In the literature, a study demonstrated a similar increase in *amoA*-AOA and  $amoA$ -AOB gene abundances after 120 days of 0.5 mg.L<sup>-1</sup> and 5 mg.L<sup>-1</sup> exposure to GO in constructed wetlands, while the increase was only significant for *amoA*-AOA gene abundance after exposure to  $-$  0.5 mg.L<sup>-1</sup> after 30 days <sup>36</sup>. In contrast, in another study after 180 days of wastewater exposure, the addition of carbon-based nanomaterials (single-walled CNTs, multi-walled CNTs and fullerene, 10 and μg.L-1 ) led to the decline of the relative abundance of *Nitrosomonas* which was the dominant 328 AOB and is mainly responsible for the oxidation of ammonia to nitrate <sup>38</sup>. Different parameters such as exposure media, the physico-chemical characteristics of nanomaterials or different initial 330 microbial communities could explain the discrepancies among studies  $^{64}$ . These contrasted results highlight the need for more mechanistic studies considering the influence of these confounding factors for a better environmental risk assessment. However, there was no impacts of CNT exposure on denitrifier communities (*nosZ* and *nirK*).

 Additionally, this increase in nitrifier abundance could also be an indirect effect related to the 335 increase of organic matter decomposition leading to an enhanced release of NH $_4^+$  which is the nitrifier substrate.

 These different results suggest that the population of nitrifiers could be favored under exposure to carbon nanomaterials compared to denitrifiers (*nosZ* and *nirK* genes unchanged abundance). This nitrification potential increase could lead to enhanced nitrate release and could thus result in higher plant uptake and therefore higher growth along the life cycle, in the eutrophication of water bodies as well as in biodiversity loss. If over the time course of this experiment, no significant impact was detected on plant fresh biomass, it would be interesting to assess this parameter after a full plant life-cycle exposure. This highlights also the interest of working with microcosm to be able to investigate cascading effects of a contaminant on an organism assemblage. In the literature, Mondal

345 et al.<sup>18</sup> evidenced similar results on *Brassica juncea* with an increase in DW/FW ratio after exposure 346  $\;$  to CNTs and oxidized CNTs at 2.3 and 23  $\mu$ g.L<sup>-1</sup>, respectively.

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#### 348 **4.3. Impacts of CNTs on H2O cycle**

349 Concerning maize, the dry matter was on average 17.4% higher for plants exposed to 10 mg.kg<sup>-1</sup> of CNT compared to control plants while the fresh weights were similar, suggesting a decrease in water content. This dry matter increase has already been demonstrated to be a common response to other 352 types of stress <sup>65,66</sup>. It can be also correlated with a decrease in some secondary metabolites in maize exposed to CNTs confirming the fact that plants are suffering stress conditions. Decrease in secondary metabolites such as phenolic compounds and tannins could impair plant defense capabilities as they are implied in many stress defense mechanisms such as high light, low 356 temperatures, pathogen infection, herbivores, nutrient deficiency and exposure to metals  $67,68$ .

357 Alternatively, this decreased water content in maize may also be associated with CNT water 358 adsorption capabilities, decreasing water bioavailability for maize  $^{69,70}$ . However, the dry mater 359 content results obtained in this study do not follow the general trend observed in the literature 360 which described an increase in water content with CNT exposure  $71,72$ . However, as demonstrated by 361 Line *et al*. crop species have very contrasted response to the exposure to CNTs, which can partly be 362 explained by different plant characteristics (morphology and/or physiology)<sup>73</sup>.

363 The results related to water use obtained at the plant organism scale were confirmed at the 364 ecosystem scale with a decrease in H<sub>2</sub>O flux in light condition in the microcosms exposed to CNTs. It 365 might be related with a decreased stomatal opening in water-stressed CNT-exposed maize. Indeed, 366 several studies demonstrated a dose-dependent decrease in stomatal conductance with the 367 concentration of CNTs in *Orthosiphon stamineus*<sup>23</sup> and *Zea mays*<sup>22</sup>.

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#### **4.4. Impacts of CNTs on agrosystem multifunctionality**

 As stated by Holden et al in 2012 while writing about ecological nanotoxicology, "standard toxicity testing anchored in single-organism, dose-response characterizations does not adequately represent 372 real-world exposure and receptor scenarios and their complexities"<sup>74</sup>. A proper risk assessment should thus derive from ecology: *ie*. the study of organisms' interactions with each other and their environments and account for ecological interactions across scales from within organisms to whole ecosystems. Indeed, impact at the organism scale could propagate to the population with consequences on community and in turn on ecosystem functions. Even though this paradigm has been stated more than 10 years ago, there are only few nano-ecotoxicological studies that use this approach; probably because the upscaling is not so straightforward as it has been demonstrated also 379 for other types of contaminants . In our study, to account for different scale processes, we used two types of markers: the biomarkers directed towards organism functioning and the so-called ecomarkers (in particular gas exchange) who reflect the ecosystem functioning.

 Also in the 2010's in the field of ecology, an interesting way to approach this question has emerged with the development of multifunctionality indices that permits to gather in a same index different 384 metrics related to the different components of a considered ecosystem <sup>43</sup>. Here, it allows to conclude that the addition of CNT could lead to contrasted consequences on agrosystem multifunctionality according to CNT concentration with a decreased multifunctionality at low concentration (0.1 mg.kg-  $<sup>1</sup>$ ). At the highest concentration though, the index remained similar to the control conditions but this</sup> might be related to the fact that opposite effects were averaged, thus hiding individual effect contribution. Additionally, it is also interesting to mention that an increased multifunctionality would not necessarily indicate a benefit for the agrosystem. For instance, an increase in total phenolic 391 compounds suggests that the plant is undergoing stress  $^{67}$ . Therefore, such an increase could not be considered as an "improvement" for agrosystem functioning. More particularly, the SEM model showed that increasing multifunctionality in response to CNT was mostly driven by increasing nitrate

394 reductase and  $CO<sub>2</sub>$  respiration. suggesting that CNT addition could promote the release of 395 greenhouse gases (CO<sub>2</sub>, CH<sub>4</sub> and N<sub>2</sub>O) from the system through direct effects on the microbiome.

#### **5. Conclusion**

 In conclusion, despite some articles of the literature suggesting that CNTs could be used as fertilizer to increase crop production, our results obtained using an integrative approach in microcosms demonstrated an impairment of some of the major biogeochemical cycles with consequences both at 401 the organism and at the ecosystem level. It is also worth mentioning that unlike more "traditional" contaminants, CNT toxicity was not dose-dependent: the highest impact on the ecosystem 403 multifunctionality was observed at the lowest exposure concentration (0.1 mg.kg<sup>-1</sup>, which is environmentally relevant in agrosystems according to modeling studies available so far) rather than 405 at the highest one (10 mg.kg $^{-1}$ , simulating a worst-case scenario).

#### **Conflicts of interest**

There are no conflicts of interest to declare.

#### **Acknowledgement**

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**Figure captions**

 **Figure 1**: A. Gene copy quantification for the universal gene 16S and for selected genes involved in N cycle: *nosZ*, *nirK*, *amoA*-AOB and *amoA*-AOA by qPCR (normalized by the initial number of gene 590 copies) and litter decomposition (B), for the 3 CNT modalities (0 = 0 mg.kg<sup>-1</sup>, 0.1 = 0.1 mg.kg<sup>-1</sup>, 10 = 10 591  $\,$  mg.kg<sup>-1</sup>) after a 6-week exposure in soil. Same lowercase letters indicate treatments that do not differ significantly (p-value > 0.05) following an ANOVA or a Kruskal-Wallis test (n = 5).

 **Figure 2**: Plant morphological parameter assessment: dry weight/ fresh weight proportion (% of dry matter) (A), height (B), fresh weight (FW, C), and leaf area (D) of maize exposed for 6 weeks in soil 596 contaminated with 3 CNT modalities (0 = 0 mg.kg<sup>-1</sup>, 0.1 = 0.1 mg.kg<sup>-1</sup>, 10 = 10 mg.kg<sup>-1</sup>). Same lowercase letters indicate treatments that do not differ significantly (p-value > 0.05) following an 598 ANOVA ( $n = 5$ ).

 **Figure 3**: Quantum yield (A), total chlorophyll (B), total phenolic compounds (C), tannin concentration (D) and principal component analysis (PCA, E) on the micro- and macro-nutrient content (Ca, Cu, Fe, Cu, Mg, Mn, Na, P, S, Zn) represented by individuals and correlation circle graphs 603 for maize leaves exposed to the 3 CNT modalities (0 = 0 mg.kg<sup>-1</sup>, 0.1 = 0.1 mg.kg<sup>-1</sup>, 10 = 10 mg.kg<sup>-1</sup>) after a 6-week exposure in soil. Cos2 corresponds to the level of variable contributions (elements) to dimensions 1 and 2. Same lowercase letters indicate treatments that do not differ significantly (p-606 value > 0.05) following an ANOVA ( $n = 5$ ).

 **Figure 4**: GreenHouse Gas exchange. Net Ecosystem Exchange (NEE, A), Ecosystem respiration (RECO, 609 B), H<sub>2</sub>O emission measured in the light (C) or in the dark (D) and CH<sub>4</sub> emission measured in the light 610 (E) or in the dark (F) for microcosms exposed to the 3 CNT modalities (0 = 0 mg.kg<sup>-1</sup>, 0.1 = 0.1 mg.kg<sup>-1</sup>,

611  $10 = 10$  mg.kg<sup>-1</sup>) after a 6-week exposure. Same lowercase letters indicate treatments that do not 612 differ significantly (p-value > 0.05) following an ANOVA or a Kruskal-Wallis (CH<sub>4</sub> in darkness) (n = 5).

**Figure 5**: A. Multifunctionality index of microcosms exposed to the 3 CNT modalities ( $0 = 0$  mg.kg<sup>-1</sup>, 614 0.1 = 0.1 mg.kg<sup>-1</sup>, 10 = 10 mg.kg<sup>-1</sup>) after a 6-week exposure. Same lowercase letters indicate treatments that do not differ significantly (p-value > 0.05) following an ANOVA (n=5). B. Structural Equation Modeling investigating the impact of carbon nanotube (CNT) addition on agrosystem multifunctionality through different functions: N cycle (nitrification represented by *amoA*-AOA gene 618 expression and denitrificiation represented by nirK gene expression, in yellow), H<sub>2</sub>O cycle (with H<sub>2</sub>O flux in the system, in blue), C cycle (including organic matter decomposition, ecosystem respiration – 620 RECO – and, CH<sub>4</sub> emission, in green) and crop resistance and nutrition (represented by tannin and mineral concentrations in plants, in red). Dashed arrows indicate a non-significant path (p values > 0.05), plain arrows indicate a significant positive correlation (p values < 0.05). The numbers adjacent to arrows are standardized path coefficients. Model fit was overall good with P = 0.436, AIC=184.0, RMSEA = 0.090, CFI = 0.988, SRMR=0.104.

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![](_page_26_Figure_1.jpeg)

**Figure 2**

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### **Figure 3**

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**Figure 5**

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