



- 1 Increase of bacterial community induced-tolerance to Cr in response to soil
- 2 properties and Cr level in the soil
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10 ABSTRACT

| 11 | Chromium (Cr) pollution in soils is a global concern that should be assessed. Pollution |
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| 12 | Induced Community Tolerance (PICT) methodology is a highly sensitive tool that can |
| 13 | directly indicate metal toxicity in the microbial community. Ten soils with a wide range |
| 14 | of properties were spiked with 31.25, 62.5, 125, 250, 500, 1000 and 2000 mg $\text{Cr}\cdot\text{kg}^{-1}$, in |
| 15 | addition to the control. Bacterial growth (using [³ H]-leucine incorporation technique) was |
| 16 | used to determine PICT, that is, whether bacterial communities developed tolerance in |
| 17 | response to Cr additions to different soil types. Some bacterial communities did not grow |
| 18 | normally at 1000 or 2000 mg $Cr \cdot kg^{-1}$, probably due to high Cr toxicity, while others did. |
| 19 | Regarding below 500 mg Cr·kg ⁻¹ , bacterial communities showed two responses |
| 20 | depending on soil type: 7 of the 10 studied soils showed increased tolerance to Cr, while |
| 21 | for the remaining 3 soils did not develop tolerance to Cr. Furthermore, the Cr level from |
| 22 | which bacterial communities developed tolerance was dependent on the soil, i.e. Cr was |
| 23 | more toxic in some of studied soils. The Cr effect on microbial communities was mainly |
| 24 | determined by Dissolved Organic Carbon (DOC) and the fraction of Cr extracted with |
| 25 | distilled water (H ₂ O-Cr) ($R^2 = 95.6$ %). Their effect on Cr in the soil might lead to an |
| 26 | increase in toxicity (selection phase of PICT). |

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- 30 Keywords:

31 PICT (Pollution-Induced Community Tolerance), bacterial growth, Cr (chromium),

- 32 dissolved organic carbon, metal bioavailability, risk assessment
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35 **1. Introduction**

Chromium (Cr) is a highly toxic non-essential metal for microorganisms and 36 plants, that may naturally occur at high concentrations from parent materials, e.g. 37 serpentine rocks (Adriano, 2001; Cervantes et al., 2001). The world average content of 38 Cr in soils is 60 mg kg^{-1} , but in soils developed from mafic and volcanic rocks can reach 39 up to 10000 mg·kg⁻¹ (Gonnelli and Renella, 2013). Cr contents up to 2879 and 3865 40 41 mg·kg⁻¹ were reported for serpentine soils in Galicia (NW Spain) and Albania, respectively, (Covelo et al., 2007; Shallari et al., 1998). Anthropogenic activities, e.g. 42 metallurgical industry, also lead to Cr accumulation in soils (Kabata-Pendias, 2011). Up 43 to 195, 88 and 6228 mg·kg⁻¹ Cr were found in urban, agricultural and industrial soils, 44 respectively (Srinivasa Gowd et al., 2010; Wei and Yang, 2010). Speciation and 45 adsorption on soil solid surfaces are the main processes controlling Cr toxicity in soils 46 47 (Adriano, 2001; Shahid et al., 2017). Despite the various Cr oxidation states, Cr(III) and 48 Cr(VI) are the most stable and common forms in soils. Cr(VI) is considered the most toxic 49 form of Cr, while Cr (III) is less mobile, less toxic and presents mostly precipitated 50 (Kabata-Pendias, 2011). The adsorption of Cr on soil solid surfaces depends on several 51 factors, e.g. soil pH, clay content, organic matter or Fe hydroxides (Bolan and Thiagarajan, 2001; Bradl, 2004; Dias-Ferreira et al., 2015; Gonnelli and Renella, 2013; 52 Kabata-Pendias, 2011). 53

In the assessment of metal pollution, the toxic metal effect on soil microorganisms should be considered, because of their key role in maintaining soil ecosystem functions (Nannipieri et al., 2003). Lower microbial diversity, enzymatic activity, C mineralization and microbial biomass were found in Cr-polluted soil in comparison to unpolluted soil (Dotaniya et al., 2017; He et al., 2016; Pradhan et al., 2019). The potential nitrification and microbial abundance were inhibited with the increase of Cr level in the soil (Zhang





60 et al., 2022). Bacterial diversity was negatively correlated with total and available Cr, while microbial community structure was altered (Zhang et al., 2021). However, 61 sometimes differentiating if the microbial response is due to Cr toxicity or to soil 62 properties variation is a difficult task (Liu et al., 2019), in addition to the complex 63 biogeochemical behaviour of Cr in soils (Ao et al., 2022). Therefore, a microbial indicator 64 specifically related to Cr toxicity that reduces interference of other soil properties is 65 66 needed to assess the Cr toxicity, such as the Pollution Induced Community Tolerance (PICT) methodology. PICT is a sensitive tool that can be used as a direct indicator of 67 metal toxicity in the microbial community (Blanck, 2002). PICT methodology is based 68 on the selective pressure that the metal exerts on a microbial community, which favoured 69 70 the proliferation of more tolerant species over the more sensitive ones. Thus, the microbial community that was exposed to the pollutant should show higher tolerance than that of 71 72 the unexposed reference microbial community (Blanck, 2002; Tlili et al., 2016). PICT 73 methodology has been successfully applied to assess Cr pollution in soils or sediments 74 (Gong et al., 2002; Ipsilantis and Coyne, 2007; Ogilvie and Grant, 2008; Santás-Miguel 75 et al., 2021; Shi et al., 2002a, 2002b; Van Beelen et al., 2004). The microbial community 76 tolerance should be quantified in a short-term assay by a sensitive endpoint, such as bacterial growth measured using [³H]-leucine incorporation (Berg et al., 2012; Boivin et 77 al., 2006; Lekfeldt et al., 2014). Despite the high sensitivity and specificity, the PICT 78 79 methodology might present some difficulties, mainly due to the influence of soil 80 properties (Blanck, 2002; Lekfeldt et al., 2014). Shi et al. (2002b) found similar values of PICT to Cr and Pb both at low and high Cr (263 $g \cdot kg^{-1}$) and Pb (10000 mg $\cdot kg^{-1}$) levels, 81 respectively, suggesting that different soils affected Cr and Pb bioavailability. Similarly, 82 Shi et al. (2002a) did not found bacterial community tolerance to Cr (or Pb), regardless 83 84 of exposure history to Cr (or Pb), suggesting that several factors (organic matter, pH,





| 85 | redox potential) might influence metal availability. Boivin et al. (2006), Fernández- |
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| 86 | Calviño et al. (2012) and Fernández-Calviño and Bååth (2016) also reported different |
| 87 | tolerance values to heavy metals in soils with similar values of metals but different soil |
| 88 | properties. Soil properties may affect PICT development due to effects on metals |
| 89 | speciation, adsorption and bioavailability (Bradl, 2004; Shahid et al., 2017). |
| 90 | We hypothesize that soil pollution with Cr induces the development of bacterial |
| 91 | community tolerance to Cr, but the magnitude of the increases depends on soil |
| 92 | physicochemical characteristics. Therefore, we aim to determine the induced bacterial |
| 93 | community tolerance to Cr in response to the addition of different Cr levels to 10 soils |
| 94 | with variable properties. We also aim to assess the importance of soil properties on the |
| 95 | increase of bacterial community tolerance to Cr. |
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98 2. Materials and Methods

99 2.1 Soil samples

Soil samples were the same used previously in Campillo-Cora et al. (2021a, 2020) to 100 101 study Cr adsorption and fractionation in soils with different properties, mainly in terms of organic matter and pH. In brief, ten remote forest locations in Galicia (NW Spain) were 102 103 selected to avoid heavy metal pollution. Locations were also selected to obtain soil 104 samples with a range of different physicochemical properties (Macías-Vázquez and Calvo 105 de Anta, 2009). Superficial soil samples (0-20 cm) were taken using an Edelman probe and, once in the laboratory, were air-dried, homogenized, sieved (2 mm mesh) and stored 106 107 until analysis.

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109 2.2 Soil properties

110 A detailed description of the chemical analysis is given in Campillo-Cora et al. (2020) 111 and in Supplementary Information. The properties of the 10 soils can be found in Tables 112 S1 and S2. In brief, soil samples presented a wide range of textures (19-71 % Sand, 13-113 67 % Silt, 14-32 % Clay). A wide range of soil pH_w and pH_K was found: 4.0-7.5 and 3.0-114 6.9, respectively. Similarly, OM oscillated between 10-29 %. A range from 2 to 29 115 cmol_c·kg⁻¹ was obtained for eCEC. A large range was obtained for DOC: 0.14 to 0.70 116 g·kg⁻¹. Chromium total content varied from 7 up to 394 mg·kg⁻¹.

Adsorption constants determined from Freundlich and Langmuir models (batch experiments) are presented in Table S3, obtained from Campillo-Cora et al. (2020). The different Cr fractions from extractions using distilled water, CaCl₂ and DTPA are shown in Table S4, obtained from Campillo-Cora et al. (2021a)

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123 2.3 Experimental design and bacterial community tolerance to Cr determination

Sieved soil samples were rewetted until reaching 60 - 80% of water holding capacity 124 125 (Meisner et al., 2013). To rewet, soil samples were spiked with seven Cr solutions (made 126 from K₂Cr₂O₇) and one of distilled water, to obtain the following final Cr levels in soils: 2000, 1000, 500, 250, 125, 62.5, 31.25 and 0 mg Cr·kg⁻¹ soil. Each Cr solution was added 127 separately and in triplicate, finally obtaining 240 microcosms (10 soils x 8 [Cr] x 3 128 129 replicates). Once soil samples were spiked with Cr, microcosms were incubated in the dark at 22 °C for two months, to ensure the reactivation of bacterial communities (Meisner 130 131 et al., 2013).

132 After the incubation period, bacterial community tolerance to Cr was estimated 133 through the PICT methodology (Blanck, 2002). The homogenization-centrifugation 134 technique was performed to extract soil bacterial communities (Bååth, 1992). The 135 bacterial community tolerance to Cr was determined as previously for Cu (Fernández-136 Calviño et al., 2011), with modifications based on suggestions by Lekfeldt et al. (2014). 137 For this purpose, each microcosm was distributed in three 50 mL centrifuge tubes and 138 MES buffer was added in a ratio 1:10 soil/buffer (20 Mm pH 6; 4-139 Morpholineethanesulfonic acid, CAS no: 4432-31-9) (Lekfeldt et al., 2014). The 140 suspensions soil/MES were mixed using a multi-vortex at maximum intensity for 3 min. 141 This step was followed by low-speed centrifugation to remove most of the fungal biomass (1000 x g, 10 min) (Bååth, 1994; Bååth et al., 2001; Rousk and Bååth, 2011). Soil 142 143 supernatants, i.e. bacterial suspensions, were filtered through glass wool and 1.5 mL aliquots were transferred into 2 mL micro-centrifugation tubes. A volume of 0.15 mL of 144 145 different Cr concentrations (made from K₂Cr₂O₇) was added to micro-centrifugation tubes, obtaining nine Cr concentrations (3.3 x 10⁻⁴ to 10⁻⁸ M) plus a blank (0.15 mL of 146 distilled water). Then, the ³H-leucine incorporation method was used to estimate bacterial 147





- growth (Bååth et al., 2001). A volume of 0.2 μ L [³H]Leu (37 MBq mL⁻¹ and 5.74 TBq mmol⁻¹. Amersham) with non-labelled Leu (19.8 μ L) was added to each tube, resulting in 300 nM Leu in the bacterial suspensions. Bacterial suspensions were incubated for 8 h at 22°C. Bacterial growth was stopped with 75 μ L of 100% trichloroacetic acid. The washing procedure and subsequent radioactivity measurement were carried out according to Bååth et al. (2001). Radioactivity was measured by liquid scintillation counting using a Tri-Carb 2810 TR (PerkinElmer, USA)
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- 156 2.4 Data analysis
- 157 2.4.1 Estimation of bacterial community tolerance to Cr (log IC₅₀)

158 A dose-response curve was obtained for each soil microcosm. To compare the dose-159 response curves, i.e. inhibition curves, with each other, bacterial growth was expressed 160 as relative bacterial growth. For each inhibition curve, generally, the four lowest added 161 metal concentrations to bacterial suspensions not showed bacterial growth inhibition 162 (Figure 1). Thus, relative bacterial growth was calculated by dividing all bacterial growth 163 data by the average of results from the four lowest added metal concentrations (including 164 blank), obtaining comparable dose-response curves. From each dose-response curve, log 165 IC₅₀ was determined as a tolerance index, i.e. Cr concentration resulting in 50% inhibition 166 of bacterial community growth. Higher log IC₅₀ values mean higher bacterial community 167 tolerance to Cr, and lower log IC_{50} values mean lower bacterial community tolerance to 168 Cr. Log IC₅₀ was calculated using the following logistic model (Fernández-Calviño et al., 2011): 169

170 $Y = c/(l + e^{b(X - a)})$ (equation 1)





171 where Y is the measured level of Leu incorporation, c is the bacterial growth rate without added Cr, b is a slope parameter indicating the inhibition rate, X is the logarithm of Cr 172 173 added, and a is log IC₅₀. 174 To detect whether bacterial community tolerance increase from different studied soils occurs, $\Delta \log IC_{50}$ was determined as the difference between log IC₅₀ value from each 175 Cr level in soil (2000, 1000, 500, 250, 125, 62.5 or 31.25 mg Cr · kg⁻¹) and the control soil 176 177 (0 mg Cr·kg⁻¹). A difference of 0.3 was taken as a reference value to determine if bacterial 178 community tolerance increased since it represents twice the Cr concentration in terms of added Cr to bacterial suspensions. If $\Delta \log IC_{50}$ is higher than 0.3, we will consider an 179 increase in bacterial community tolerance to Cr (Fernández-Calviño and Bååth, 2016, 180 2013). 181

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183 2.4.2 Estimation of bacterial community tolerance increase to Cr (multiple linear
184 regression analyses)

185 A multiple regression analysis, using the backward elimination method, was performed 186 to obtain an equation that allows estimating the increase in bacterial community tolerance 187 to Cr ($\Delta \log IC_{50}$) from soil properties. As the inhibition curves for some soils did not fit the logistic model (equation 1) for the highest Cr concentrations (1000 and 2000 mg·kg-188 ¹), $\Delta \log IC_{50}$ from 500 mg·kg⁻¹ was used for estimations. Once the equation was 189 estimated, determining factors were verified: linearity, error independency, residues 190 homoscedasticity, residuals normality, autocorrelation, collinearity and presence of 191 outliers. All statistics were performed using IBM SPSS Statistics 25 software (IBM, 192 193 USA).





195 **3. Results and discussion**

| 196 | 3.1 Bacterial community tolerance to Cr in Cr-polluted soils with different properties |
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| 197 | Figure 1 shows bacterial growth inhibition curves obtained for each microcosm. |
| 198 | Generally, a sigmoid dose-response behaviour is observed in the inhibition curves, |
| 199 | indicating that when the added Cr concentration to bacterial suspension was low, relative |
| 200 | bacterial growth was close to 1, while decreased when the Cr concentration increased. |
| 201 | Most of the bacterial growth data fitted the logistic model, obtaining $R^2 \ge 0.87$, (Table |
| 202 | S5). However, some data from 1000 and 2000 mg Cr·kg ⁻¹ did not fit the logistic model, |
| 203 | i.e. bacterial populations were not able to normally grow probably due to high Cr toxicity. |
| 204 | In the case of 2000 mg \cdot kg ⁻¹ , bacterial populations only grew normally in 4 of the 10 |
| 205 | studied soils, while at 1000 mg \cdot kg ⁻¹ they grew normally in 7 soils. These differences in |
| 206 | bacterial growth for the same Cr levels may indicate the influence of soil properties on |
| 207 | Cr availability, as was previously suggested by Van Beelen et al. (2004). They found |
| 208 | tolerant communities to Cr(III) in polluted soils with high Cr levels (2894 mg \cdot kg ⁻¹) but |
| 209 | also reported that microbial communities from soils polluted with 3935 mg $\text{Cr}{\cdot}\text{kg}^{\text{-1}}$ did |
| 210 | not show tolerance to Cr(III), suggesting the influence of soil properties on metal toxicity. |
| 211 | Therefore, in order to determine which properties influence Cr toxicity, the data of 1000 |
| 212 | and 2000 mg $Cr \cdot kg^{-1}$ were not considered in the following analysis. |

The log IC₅₀ values determined from inhibition curves using the logistic model (equation 1) are presented in Table 1. Bacterial community tolerance to Cr (log IC₅₀) greatly varied between soils, even in the reference soils with no added Cr, log IC₅₀ oscillated from -6.40 (S8) up to -3.88 (S6) (log units). The variation of bacterial community tolerance to Cr in the reference soils may be an indicator that the development of PICT is dependent on soil type. In addition, this bacterial community tolerance to Cr fluctuation in reference soils, together with the natural Cr content in soils (7 – 394 mg·kg⁻





| 220 | ¹ , Table S2), highlights the importance of selecting reference soils for PICT studies |
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| 221 | (Campillo-Cora et al., 2022a; Campillo-Cora et al., 2021b). Likewise, when Cr was added |
| 222 | to soils, bacterial community tolerance to Cr varied greatly between soils with the same |
| 223 | Cr level. A range from -6.37 (S8) to -3.56 (S6) was determined for soils polluted with the |
| 224 | lowest Cr level in soil (31.25 mg Cr·kg ⁻¹); from -6.27 (S8) to -3.79 (S7) for 62.5 mg |
| 225 | $Cr \cdot kg^{-1}$; from -6.26 (S8) to -3.65 (S7) for 125 mg $Cr \cdot kg^{-1}$; from -6.27 (S5) to -3.41 (S7) |
| 226 | for 250 mg Cr·kg ⁻¹ ; and from -6.09 (S8) to -2.87 (S3) for 500 mg·kg ⁻¹ . |
| 227 | Overall, bacterial communities showed two different responses to Cr addition to |
| 228 | the soil (Figure 2): (1) bacterial communities of S1, S2, S3, S6, S7, S8 and S10 developed |
| 229 | tolerance in response to Cr additions; while (2) bacterial communities of S4, S5 and S9 |
| 230 | did not develop tolerance following Cr addition to the soil. Based on the PICT hypothesis, |
| 231 | the bacterial community is first exposed to the metal (i.e. selection phase of PICT), and |
| 232 | if metal exerts toxicity, then the most sensitive organisms of the community will |
| 233 | disappear, while the tolerant ones will be favoured. Therefore, whether the microbial |
| 234 | community developed tolerance to Cr is a toxicity indicator. Later, the microbial |
| 235 | community tolerance is quantified through a second exposition to Cr (i.e. detection phase |
| 236 | of PICT) (Blanck, 2002; Tlili et al., 2016). Accordingly, Gong et al. (2002) and Ipsilantis |
| 237 | and Coyne (2007) reported an increase in bacterial community tolerance to Cr with |
| 238 | increasing Cr levels in soil and rhizosphere. Van Beelen et al. (2004) found that bacterial |
| 239 | community tolerance to Cr(VI) increased with increasing Cr in pore water. Ogilvie and |
| 240 | Grant (2008) determined a tendency to increase the bacterial community tolerance to Cr |
| 241 | when the Cr level increases in estuarine sediments. Our results showed that bacterial |
| 242 | community tolerance to Cr increased with increasing Cr levels in soils only in 7 of the 10 |
| 243 | soils studied (Figure 2). However, our results showed that the Cr level in soil from which |
| 244 | bacterial communities developed tolerance to Cr varied depending on the soil ($\Delta \log IC_{50}$ |





245 > 0.3). Bacterial communities from S7 and S10 showed an increased tolerance at 31.25 mg Cr·kg⁻¹, bacterial communities from S1 and S3 at 62.5 mg Cr·kg⁻¹, bacterial 246 communities from S2 and S8 at 250 mg Cr·kg⁻¹, and bacterial communities from S6 at 247 500 mg Cr·kg⁻¹. In other words, Cr was more toxic for bacterial communities depending 248 on soil type, following the sequence: S7, S10 > S1, S3 > S2, S8 > S6. In other soils, our 249 250 results show that microbial communities did not develop tolerance to Cr, even at high Cr 251 levels. For example, bacterial communities of S6 did not show tolerance to Cr even at 2000 mg·kg⁻¹ (Figure 2). Similarly, Shi et al. (2002b, 2002a) and Ipsilantis and Coyne 252 (2007) did not find tolerant microbial communities to Cr even at high Cr levels, from 447 253 up to 263000 mg Cr·kg⁻¹. Therefore, considering that Cr-pollution sometimes has no toxic 254 effect on microbial communities and that, in other cases, microbial communities are 255 affected by Cr from very low levels of Cr-pollution, including soil properties in the 256 257 assessment of Cr-pollution is highly recommended, as for other heavy metals (Campillo-258 Cora et al., 2022b).

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260 3.2 Estimation of the increase in bacterial community tolerance to Cr as a function of soil
261 properties

The bacterial community tolerance to metals may be influenced by several soil properties, 262 263 such as soil pH, clay content or organic matter content (Ogilvie and Grant, 2008; Shi et 264 al., 2002b). The effect of soil properties on bacterial community tolerance can occur in 265 soil (selection phase of PICT), or in the determination phase of PICT. The effect of the soil properties in the selection phase occurs in the soil, i.e. the first time bacterial 266 267 communities are exposed to the metal. For example, Fernández-Calviño and Bååth (2016) found that bacterial community tolerance to Cu was lower in vineyard soils with high pH 268 269 in comparison to more acid soils, as Cu toxicity was reduced. On the other hand, the effect





270 of soil properties may occur in the detection phase, i.e. confounding factors leading to 271 altered tolerance measures (Lekfeldt et al., 2014). For example, Fernández-Calviño et al. (2011) reported that the measurement of PICT to Cu was altered because of the presence 272 273 of the finer soil fraction in the bacterial suspensions when Cu concentrations were added. That is, the finer particles will bind part of the Cu added to bacterial suspensions, resulting 274 in lower available Cu, so higher Cu concentrations will be necessary to inhibit the 275 276 bacterial growth leading to apparent higher tolerance, i.e. overestimated bacterial 277 community tolerance to Cu.

The equation presented in Table 2 related the increase of bacterial community tolerance to Cr ($\Delta \log IC_{50}$) with soil properties, explaining 95.6 % of the data variance (p < 0.001). Only $\Delta \log IC_{50}$ for 500 mg Cr·kg⁻¹ were used. The increase of bacterial community tolerance to Cr was estimated by using soil properties (p < 0.05): DOC and extracted Cr using distilled water (H₂O-Cr). Figure 3 shows estimated $\Delta \log IC_{50}$ versus measured $\Delta \log IC_{50}$, with a homogeneous distribution around the line 1:1 (R² = 0.95).

DOC showed a significant positive relationship with $\Delta \log IC_{50}$ (p < 0.05; Table 284 285 2), i.e. when DOC increases, the bacterial community tolerance to Cr also increases. This 286 DOC effect might be a confounding factor in the detection phase of PICT, as was previously reported for Cu (Campillo-Cora et al., 2021b; Lekfeldt et al., 2014). When 287 288 bacterial communities are extracted from soil, DOC is extracted too. Later, when Cu is 289 added to bacterial suspensions, Cu and DOC may bind together (Beesley et al., 2010), 290 reducing Cu bioavailability and altering bacterial community tolerance to Cr (overestimation). Bérard et al. (2016) reported a similar effect for microbial community 291 tolerance to Pb measurements. However, in a previous study (Campillo-Cora et al., 292 2022c), we found that when dissolved organic matter (DOM) increases on bacterial 293 294 suspensions, then bacterial community tolerance to Cr decreases, i.e. when DOM





295 increases in bacterial suspensions, Cr becomes more toxic to bacteria. Hence, the DOC 296 effect in Cr bioavailability in the detection phase should be discarded because of the positive relationship with $\Delta \log IC_{50}$ (Table 2) and attributed to an effect in the selection 297 298 phase in soil. In the soil, however, when DOC is present, Cr(VI) may be reduced to Cr(III), i.e. Cr toxicity decreases when DOC is present (Ao et al., 2022). If fact, the use 299 of organic amendments to reduce Cr toxicity in soils is very common (Abou Jaoude et 300 301 al., 2020; Mitchell et al., 2018; Yang et al., 2021). A hypothesis is that the presence of 302 DOC in soil enhanced the reduction of Cr(VI) to Cr(III) (Wittbrodt and Palmer, 1997), but during this process free radicals may also be formed (Kotaś and Stasicka, 2000), 303 increasing general toxicity for bacterial communities (Campillo-Cora et al., 2022c). In 304 305 response to increased toxicity in soil, then bacterial communities showed tolerance to Cr. Another hypothesis might be the ability of Cr(III) to coordinate various organic 306 307 compounds, leading to the inhibition of some metalloenzyme systems (Kotaś and 308 Stasicka, 2000), which might result in a more tolerant bacterial community.

309 The Cr fraction extracted with distilled water (H2O-Cr) showed a positive 310 relationship with $\Delta \log IC_{50}$ (p < 0.001, Table 2). Usually, the soluble form of heavy metals 311 represents the soil solution metal content, which is the most mobile and bioavailable form 312 (Kabata-Pendias, 2011). In the vase of Cr, probably Cr(VI) (Ao et al., 2022). Thus, H₂O-313 Cr exerts its effect in soil, during the selection phase. H2O-Cr content in soil increases as 314 added Cr level in soils increases (Campillo-Cora et al., 2021a). Whether Cr exerts 315 toxicity, the most sensitive bacterial species were removed, while the tolerant ones survived, resulting in a more tolerant community to Cr. Later, in the detection phase, 316 317 when bacterial growth is measured and Cr is added to bacterial suspensions, tolerant bacteria allow greater Cr concentrations, leading to a higher tolerant community. Van 318 319 Beelen et al. (2004) found a significant increase in microbial community tolerance to





320 Cr(VI) with Cr(VI) pore-water concentration. Similarly, Fernández-Calviño and Bååth 321 (2016) reported a positive relationship between bacterial community tolerance increase 322 (Δ log IC₅₀) to Cu versus water-soluble Cu concentrations logarithm (R² = 0.79). Kunito 323 et al. (1999) also determined a positive correlation between IC₅₀ values and soluble-324 exchangeable Cu (r = 0.76), while total Cu did not show any significant relationship (r = 325 0.013, p > 0.05).

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327 3.3 Concluding remarks

In the present study, we aimed to improve the PICT methodology for the assessment of 328 329 soil pollution, using bacterial growth as the endpoint. Dissolved organic carbon (DOC) 330 and the fraction of Cr extracted with distilled water (H2O-Cr) were the main factors 331 controlling the Cr effect on microbial communities, determined by the increase of 332 bacterial community tolerance to Cr. The main selection pressure of Cr on the microbial 333 community presumably occurs in soil, i.e. the selection phase of PICT. In the case of 334 DOC, Cr became more toxic to bacterial communities as DOC increased in soil, leading 335 to an increase in bacterial community tolerance to Cr in response to toxicity. Secondly, 336 H₂O-Cr is related to the toxic and active form of Cr, probably Cr(VI), and the higher the 337 H₂O-Cr content in the soil, the higher the tolerance to Cr developed by bacterial 338 communities. The outcomes of this study may be helpful for normalising Cr toxicity thresholds for soil with different properties. In addition, overestimations or 339 340 underestimations of Cr toxicity based on total or bioavailable Cr content may be avoided, 341 since soil properties should be considered during risk assessment.

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557 Tables

558 Table 1

- 559 Bacterial community tolerance (expressed as $\log IC_{50}$) to different levels of Cr pollution
- 560 in the 10 studied soils (average \pm SE)

| Cr | 2000 | 1000 | 500 | 250 | 125 | 62.5 | 31.25 | 0 |
|------------------------|-------------------------|-------------------------|-------------------------|-------------------------|-------------------------|-------------------------|-------------------------|------------------------|
| (mg·kg ⁻¹) | | | | | | | | |
| Soil | Log | Log |
| | IC ₅₀ ±error | IC ₅₀ ±erro |
| S1 | -5.34±0.03 | -5.35±0.05 | -5.28±0.03 | -5.30±0.03 | -5.33±0.03 | -5.30±0.04 | -5.83±0.06 | -5.82±0.0 |
| S2 | -4.04±0.24 | -4.55 ± 0.42 | -4.61±0.21 | -4.68 ± 0.41 | -4.78±0.43 | -4.70±0.21 | -4.81±0.19 | -5.02±0.1 |
| S3 | * | * | -2.87±0.51 | -4.38±0.15 | -4.62±0.16 | -4.70±0.18 | -5.46±0.03 | -5.38±0.0 |
| S4 | -5.85±0.08 | -5.76±0.05 | -5.80±0.07 | -5.69±0.05 | -5.66±0.04 | -5.68±0.04 | -5.90 ± 0.08 | -5.66±0.0 |
| S5 | * | -4.47±0.11 | -5.80±0.19 | -6.27±0.07 | -5.86±0.10 | -5.98±0.06 | -6.02±0.10 | -6.09±0.0 |
| S6 | * | -3.47±0.06 | -3.38±0.08 | -4.48±0.13 | -4.18±0.16 | -3.97±0.12 | -3.56±0.23 | -3.88±0.1 |
| S7 | * | -3.44±0.09 | -3.35±0.07 | -3.41±0.09 | -3.65±0.11 | -3.79±0.07 | -3.85±0.05 | -4.32±0.1 |
| S8 | -3.63±0.13 | -6.03±0.06 | -6.09±0.09 | -5.90±0.09 | -6.26±0.04 | -6.27±0.03 | -6.37±0.07 | -6.40±0.1 |
| S9 | * | * | -4.32±0.27 | -4.37±0.39 | -4.70±0.23 | -4.43±0.13 | -3.82±0.05 | -4.11±0.0 |
| S10 | * | * | -4.75±0.13 | -4.64±0.09 | -4.48±0.09 | -4.69±0.09 | -4.76±0.04 | -5.16±0.0 |

561 *Unadjusted data





563 **Table 2**

- The equation for estimating bacterial community tolerance increase to Cr ($\Delta \log IC_{50(500-1)}$
- $_{0}$) was obtained by multiple regression analysis using all soil samples (n=10).

| F | <i>p</i> -value | Adjusted R ² |
|--------|-----------------|--|
| 87.309 | <0.001 | 0.956 |
| | | |
| | | |
| | 011207 | 87.309 <0.001 ted using H ₂ O. Values as |

566 DOC is dissolved organic carbon $(g \cdot kg^{-1})$; H₂O-Cr is Cr extracted using H₂O. Values associated with the 567 independent variables are shown together with the standard errors (±). P-values associated with each 568 independent variable are shown below variables (in brackets)

569

570





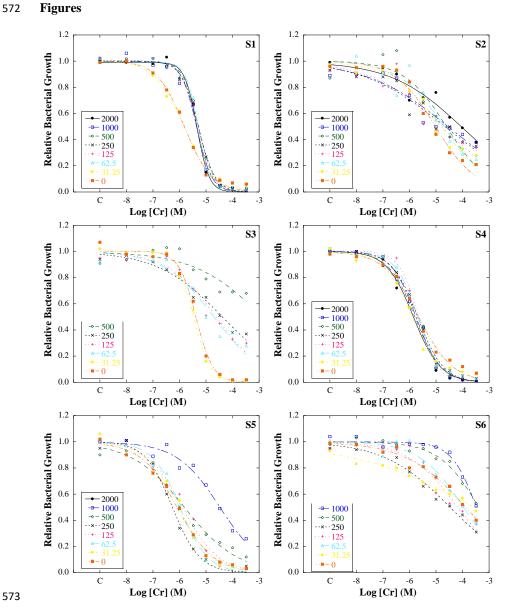
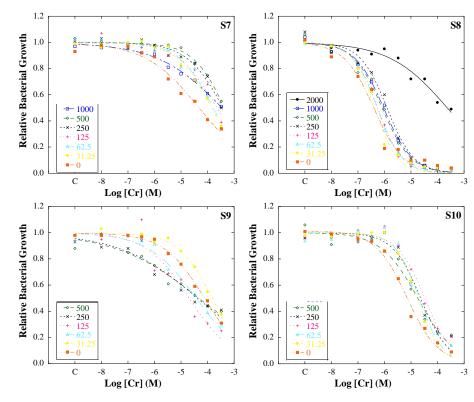


Figure 1. Bacterial growth inhibition curves for bacterial suspensions extracted from 10 soils
artificially polluted with a range of Cr concentrations: 2000, 1000, 500, 125, 62.5, 31.25 and 0
mg·kg⁻¹. Dots indicate real data measured, while the lines represent the fit of the data to the logistic
model used.







580 **Figure 1** (continued)

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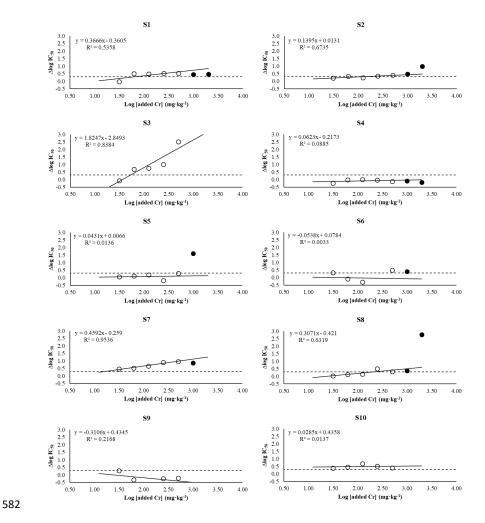
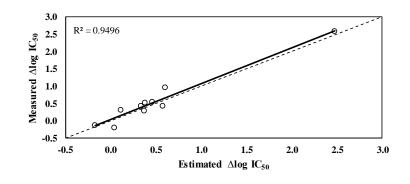
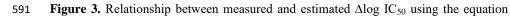


Figure 2 Bacterial community tolerance variation (expressed as $\Delta \log IC_{50}$ concerning unpolluted soil) to a range of added Cr to soil (in logarithm scale). White dots represent data from $\Delta \log IC_{50(31.25-0)}$, $\Delta \log IC_{50(62.5-0)}$, $\Delta \log IC_{50(125-0)}$, $\Delta \log IC_{50(250-0)}$ and $\Delta \log$ $IC_{50(500-0)}$. Black dots represent data from $\Delta \log IC_{50(1000-0)}$ and $\Delta \log IC_{50(2000-0)}$. Continuous lines represent linear regression fit. The discontinuous line represents the value (0.3) from which it is considered that the bacterial community has developed tolerance.









- from Table 2. The stippled line indicated a 1:1 relationship.