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- ¹ Improvements in estimating bioaccumulation metrics in
- ² the light of toxicokinetics models and Bayesian inference
- ³ Aude Ratier¹, 2 \cdot Christelle Lopes¹ \cdot
- ⁴ Sandrine Charles^{1,*}

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Université de Lyon, Université Lyon 1, CNRS UMR5558, Laboratoire de Biométrie et Biologie Evolutive, 69100 Villeurbanne, France.

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Institut National de l'Environnement Industriel et des Risques (INERIS), Parc ALATA BP2, 60550 Verneuil en Halatte, France.

^{*}Corresponding author: E-mail: sandrine.charles@univ-lyon1.fr

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22 abstract

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The surveillance of chemical substances in the scope of Environmental Risk 23 Assessment (ERA) is classically performed through bio-assays from which data 24 are collected and then modelled. Some statistical analysis base on the fitting 25 of toxicokinetic (TK) models to assess the bioaccumulative capacity of chem-26 ical substances via the estimation of bioaccumulation metrics as required by 27 regulatory documents. Given that bio-assays are particularly expensive and 28 time consuming, it is of crucial importance to deeply benefit from all informa-29 tion contained in the data. By revisiting the calculation of bioaccumulation 30 metrics under a Bayesian framework, this paper presents improvements in the 31 classification of the bioaccumulative capacity of chemical substances. A meta-32 analysis of a data-rich TK database was performed, considering the uncer-33 tainties around the bioaccumulation metrics. The subsequent results appeared 34 sufficiently statistically robust to propose the replacement of the single me-35 dian estimate to decide of the class to which assign a chemical substance. The 36 main recommendation is to use the 75th percentile of the uncertainty interval 37 of the bioaccumulation metrics, which revealed a better criterion to classify a 38 chemical substance, and in the same way as the conventional method in 90%39 of cases. 40

Keywords: Environmental Risk Assessment, classification of chemical sub stances, REACH regulation, accumulation-depuration test.

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43 1 Introduction

Chemical substances, present in the environment as a result of human ac-44 tivities, are of extreme concern due to their persistence, to their capacity in 45 accumulating within non-target living organisms and to their potential toxi-46 city on the different levels of biological organization all along trophic chains 47 (Cousins et al., 2019; Popek, 2018). In particular, this reveals crucial to bring 48 reliable and precise information on the bioaccumulative capacity of the dif-49 ferent chemical substances, on what depend the concentrations internalized 50 by organisms almost exclusively. This stage then conditions the way in which 51 relevant links can be established between the exposure concentrations and the 52 likely damages on life-history traits (Arnot and Gobas, 2006; Chojnacka and 53 Mikulewicz, 2014; Armitage et al., 2021). 54

In Europe, chemical substances are governed by the REACH regulation, 55 adopted by the European Union to improve the protection of human health 56 and the environment from the risks that can be raised by chemical substances 57 (European Commission, 2006). In principle, REACH applies to all chemical 58 substances, also stipulating the need to reduce animal testing. To comply with 59 the regulation, manufacturers must identify and manage the risks linked to 60 the chemical substances in demonstrating that they can be safely used. For 61 example, in order to evaluate the bioaccumulative capacity of chemical sub-62 stances, produced or imported above 100 tonnes per year, the REACH regula-63 tion requires the calculation of bioaccumulation metrics. According to Ratier 64 et al. (2021b), we use the generic expression "bioaccumulation metrics" to de-65 note either bio-concentration factors (BCF) used when exposure is via water, 66 biota-sediment accumulation factors (BSAF) when exposure is via sediment 67 or biomagnification factors (BMF) when exposure is via food. For substances 68 produced or imported between 10 and 100 tonnes per year, bioaccumulation 69

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metrics are not mandatory but still required to classify chemical substances 70 as persistent, bioaccumulative and toxic (abbreviated PBT) or very persistent 71 and very bioaccumulative (abbreviated vPvB). Most of European countries 72 classify substances as bioaccumulative (abbreviated "B") if the bioaccumula-73 tion metric is in [2000; 5000], or very bioaccumulative (abbreviated "vB") if it 74 is > 5000 (European Commission, 2006). Other regulations around the world 75 (Saito et al., 2011; Government of Canada, 1999; Agency, 1979) classify as "B" 76 a chemical substance with a bioaccumulation metric ranging in [1000; 5000]. 77 Chemical substances with a bioaccumulation metric in [1000; 2000] are always 78 classified as low bioaccumulative (abbreviated as " ℓB "). These classifications 79 are summarized in Wassenaar et al. (2020) and by Hartmann et al. (2014). 80 Bioaccumulation metrics are calculated from toxicokinetic (TK) parame-81

ter estimates by fitting a TK model (usually a one-compartment model) to 82 experimental data collected during bioaccumulation tests (e.g., OECD (2008, 83 2012)). Bioaccumulation tests provide internal concentration measurements 84 from two-phase experiments: a first phase (the "accumulation" phase) during 85 which organisms are exposed via one or several uptake route(s) (water, pore 86 water, sediment and/or food) to a given chemical substance, kept constant 87 over time; a second phase (the "depuration" phase) during which organisms 88 are transferred into a clean medium where elimination processes take place. 89 The TK model is expected to account for all uptake routes and elimination 90 processes (including excretion, dilution by growth and/or metabolization) to 91 better describe the overall kinetics in terms of internal concentrations. From 92 TK parameter estimates (namely, uptake and elimination rates), two types of 93 bioaccumulation metrics can be calculated: the "kinetic" bioaccumulation met-94 ric, as the ratio between uptake and elimination rates; and the "steady-state" 95 bioaccumulation metric as the ratio of the internal concentration at steady 96 state and the constant exposure concentration; the steady state is defined as 97

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⁹⁸ the plateau concentration that is expected at the end of the accumulation ⁹⁹ phase. In theory, the "steady-state" bioaccumulation metric should be calcu-¹⁰⁰ lated only if the internal concentrations measured at the last three time points ¹⁰¹ of the accumulation phase are not significantly different.

In addition to the experimental procedure of bioaccumulation tests, the 102 OECD guidelines also explain how to obtain bioaccumulation metrics depend-103 ing on the exposure routes that have been considered within the experiments. 104 If a one-compartment TK model is often sufficient. So, in case the goodness 105 of fit is poor this may be an indication that first order kinetics does not ap-106 ply suggesting that a more complex model should be employed. For example, 107 one of the most common complexities to account for is fish growth during 108 the bioaccumulation test (OECD, 2012). Surprisingly, in such a case, guide-109 lines only recommend to seek advice from bio-statistician and/or pharmaco-110 kineticist experts. In addition, very few tools exist to easily perform TK anal-111 yses. To name but a few, there are the Excel macro by Gobas et al. (2020), 112 the "bcmfR" package (OECD, 2012) and the free open-source $MOSAIC_{bioacc}$ 113 web service (http://umr5558-shiny.univ-lyon1.fr/mosaic-bioacc/) that 114 has been recently updated (Ratier et al., 2021b; Charles et al., 2021a). Only 115 MOSAIC_{bioacc} is entirely generic whatever the species-compound combination 116 of interest, allowing to account for different exposure routes and several elim-117 ination processes simultaneously, automatically adapting the fitted TK model 118 according to the input experimental data. It provides all possible bioaccumu-119 lation metrics accordingly, namely BCF, BSAF and/or BMF. 120

Recently, the European Food Safety Authority (EFSA) strongly advocated the need to associate uncertainties with model parameter estimates (EFSA Scientific Committee, 2018) in general, emphasizing this need for toxicity indicators in particular (Ockleford et al., 2018). This requirement is complementary to the previous regulation with regard to the bioaccumulation metrics. Indeed,

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for the authorisation dossiers of active chemical substances, the current regu-126 latory document only ask for a single mean or a median value as bioaccumu-127 lation metrics (European Commission, 2013). Such a practice is still common 128 today, probably because of a lack of efficient and easy tools to handle com-129 puter resources, specifically designed to automatically provide uncertainties 130 on any model output. However, recent recommendations have clearly been es-131 tablished when using toxicokinetic-toxicodynamic (TKTD) models (Baudrot 132 and Charles, 2019), while Charles et al. (2021b) highlighted how critical it 133 is to take uncertainty into account when assessing the toxicity of a chemical 134 substance to a range of non-target terrestrial plants thus revisiting the species 135 sensitivity distribution (SSD) approach. 136

Today, when bioaccumulation metrics need to be estimated, MOSAIC_{bioacc} 137 is one of the only tools in support of a facilitated in-depth quantification of 138 the uncertainties; on the contrary, Gobas et al. (2020) tool only gives a stan-139 dard deviation. While available on-line since May 2020, the number of users 140 of MOSAIC_{bioacc} is continuously growing all over the world, whether they 141 are from academia, regulatory bodies or industry (380 recordings these last 142 6 months). A publicly available database accompanies the MOSAIC_{bioacc} ser-143 vices (http://umr5558-shiny.univ-lyon1.fr/mosaic-bioacc/data/database/ 144 index_readme.html), with more than 200 accumulation-depuration data sets 145 collected within published scientific papers (Ratier and Charles, 2021). All 146 data sets are automatically analyzed with $MOSAIC_{bioacc}$, full analysis reports 147 being made available via the database directly. This database is dynamically 148 supplemented as new data sets are retrieved from the recent scientific litera-149 ture, or directly deposited by researchers upon request. 150

The aim of the present paper is to propose improvements in the estimation of bioaccumulation metrics in association with the quantification of their uncertainty with the perspective to reinforce the statistical foundations leading

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to the classification of chemical substances as (low, medium or very) bioac-154 cumulative or not. To this end, we first present the last updates brought to 155 MOSAIC_{bioacc}, especially an innovative prediction device that can be specif-156 ically used in designing new experiments in full respect of the 3R principles 157 (Replacement, Reduction and Refinement) ensuring animal welfare and quality 158 of science (Prescott and Lidster, 2017). Then, the added value of accounting for 159 uncertainty of bioaccumulation metrics is underlined through a meta-analysis 160 of the TK database associated with MOSAIC_{bioacc}. Finally, we demonstrate 161 how influential may be the consideration of uncertainty when classifying chem-162 ical substances according to the current regulatory intervals into which the 163 bioaccumulation metric estimates fall. We conclude with a revisited workflow 164 to improve environmental risk assessment if it would be adopted by regulatory 165 bodies. 166

¹⁶⁷ 2 Calculations and predictions of bioaccumulative capacity

This section gives a brief overview of the different features of MOSAIC_{bioacc}. A focus is first made on the calculation of bioaccumulation metrics. Then the new prediction tool is introduced, to close with illustrative case studies.

171 2.1 Calculations of bioaccumulation metrics

Recent updates have been brought to the MOSAIC_{bioacc} web application to increase the speed of calculations and improve titse user-friendliness. Above all, a new R-package is today available on the official CRAN web site (https: //CRAN.R-project.org/package=rbioacc) that allows to similarly perform all MOSAIC_{bioacc} calculations and graphs directly in the R software with ready-to-use dedicated functions (Ratier et al., 2021a). The new version of MOSAIC_{bioacc} has entirely been rewritten to reduce the length of the source

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code and take advantage of this new package. Above all, MOSAIC_{bioacc} is now 179 based on a tabbed presentation that clarifies and facilitates browsing from one 180 step to the next. A special tab gives all bioaccumulation metrics, appropriately 181 calculated according to the input observed data that the user has uploaded. 182 By default, the kinetics BCF, BSAF or BMF values are delivered, displayed 183 via their entire posterior probability distribution then summarized with their 184 median (that is the 50% quantile) and their 95% uncertainty interval (bounded 185 by the 2.5 and 97.5% quantiles). In addition, users can ask for the steady-state 186 corresponding bioaccumulation metrics if they consider it relevant according 187 to the duration of the accumulation phase, and the fact of having actually 188 reached the plateau. 189

¹⁹⁰ 2.2 Prediction of bioaccumulation matrics to optimize experiments

A new prediction tab have been added to $MOSAIC_{bioacc}$ that allows interac-191 tive simulations of a TK model under a constant or a time-variable exposure 192 profile. The main aim of this prediction tool is to assist experimenters in opti-193 mizing the design of new experiments, based on previous TK analyses. Indeed, 194 when studying a new chemical substance and/or a new species, when study-195 ing a new chemical substance and/or a new species, some information needed 196 to be known in advance in full respect of scientific ethic in terms of experi-197 ment on living organisms and chemical use and recycling. For example, the 198 exposure concentration, the duration of the accumulation and the depuration 199 phases as well as the number of time points at which internal concentrations 200 should be primarily measured need to be defined in advance. Then, based on 201 previous TK analyses for given close species/compound combinations, TK pa-202 rameter estimates can serve to simulate what could be expected when planing 203 additional time points and/or extending the accumulation phase for example. 204

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Above all, benefiting of the Bayesian framework, the uncertainty around parameter estimates can be propagated towards predictions and any function of the parameters (Baudrot and Charles, 2019), in a way that is particularly useful when environmentally realistic exposure scenarios have to be run in numbers.

210 2.3 Illustrations with case studies

We provide a collection of case studies as supplementary information (SI, see the .pdf file) to illustrate various situations where the prediction tool can be helpful:

Case study 1 Plan an experiment for an already studied species exposed to 214 a different but chemically similar compound (i.e., with a mode of action215 expected to be close), without accounting for the parameter uncertainty; 216 Case study 2 Compare several species exposed to a same chemical substance 217 accounting for the uncertainty around parameter estimates coming from 218 a previous TK analysis conducted on a species phylogenetically (or tax-219 onomically) close to the new set of interest. In such a case, the user will 220 need to enter the required input information but also a tabular file with 221 the joint posterior distribution of the parameters, either coming from a 222 previous $MOSAIC_{bioacc}$ TK analysis or a home-made TK implementation. 223 Case study 3 A prediction for a same species/compound combination but 224 for different exposure scenarios for which the user may have observed data 225 to which simulations can be compared as a validation step of the exploited 226 TK model. 227

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²²⁸ 3 Matter of uncertainty in estimating bioaccumulation metrics

Regarding accumulation-depuration data, one output of great interest is the 229 bioaccumulative capacity of a chemical substance which is assessed through 230 the appropriate bioaccumulation metrics according to the exposure route(s) 231 (either the BCF, the BSAF and/or the BMF). So, benefiting of the probability 232 distributions of these metrics (coming from the propagation of the parameter 233 uncertainty) is crucial to catch their precision. This latter may indeed influence 234 the classification of the substance as bioaccumulative or not, and if bioaccu-235 mulative, influence the choice between "B" or "vB" categories. In order to 236 illustrate this critical issue for ERA, we present below a meta-analysis of the 237 TK database associated to MOSAIC_{bioacc} (Ratier and Charles, 2021). 238

239 3.1 The accumulation-depuration TK database

The TK database currently contains 211 accumulation-depuration data sets 240 collected from a literature review and corresponding to a total of 56 studies. 241 The 211 data sets encompass 52 genus, 124 chemical substances, three different 242 exposure routes (water being the main one, sediment and food), 34 data sets 243 with also biotransformation data (that is metabolization data). Figure 1 shows 244 several tree maps performed with the treemap R-package (Tennekes, 2017) 245 allowing to visualise the proportion of each chemical category (Figure 1-a) and 246 each genus (Figure 1-b) among the 211 data sets. Pesticides and hydrocarbons 247 are the most represented chemical substances within the TK database, what 248 can be explained by the predominance of data sets on freshwater invertebrates 249 and fish. Gammarus and Daphnia are the most represented genus, probably 250 due to less ethic exigence with them, while fish studies are rarer. Gammarus 251 and Daphnia are also the genus for which the most biotransformation data are 252 available. 253

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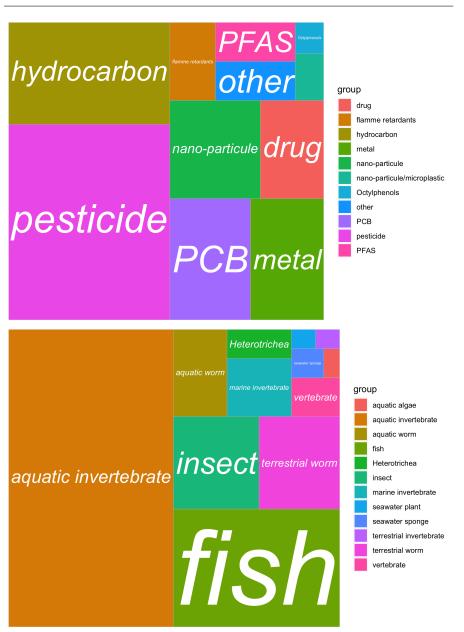


Fig. 1 Tree maps of chemical categories (upper panel) and genus categories (lower panel) available in the TK database available at http://lbbe-shiny.univ-lyon1.fr/mosaic-bioacc/data/database/TK_database.html.

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The 211 data sets of the TK database were fully analysed with MOSAIC_{bioacc}. Corresponding reports are available from the database itself, as well as references from which data have been extracted. From these analyses, we conducted a meta-analysis of the bioaccumulation metrics to specifically illustrate how much accounting for uncertainty matters when characterizing the bioaccumulation capacities of chemical substances.

²⁶⁰ 3.2 Meta-analysis of the TK database

Even if not required within regulatory documents, associate the uncertainty 261 to a bioaccumulation metric is of crucial importance (Wassenaar et al., 2020). 262 Among the 211 data sets, a total of 137 corresponds to an exposure via wa-263 ter for which the MOSAIC_{bioacc} analysis then provides a BCF probability 264 distribution. Based on the median and the 95% uncertainty interval of these 265 BCF estimates (Figure 2), aquatic invertebrates have the highest values, when 266 predominantly exposed to pesticides or metals, among which the genus Gam-267 marus has the most BCF estimates greater than 5000, classifying the corre-268 sponding chemical substances as "vB". The SI (see the .html file) provides an 269 additional figure with all the 211 estimated bioaccumulation metrics, without 270 distinguishing BCF from BSAF and BMF estimates. Note that from this Fig-271 ure, we would have concluded to similar trends. On the other hand, based on 272 the median of the BCF estimates as required by the regulation, only 22.7%273 of the BCF values classify chemical substances as "B", that is with BCF me-274 dians higher than 1000. Nevertheless, as a matter of fact, this classification 275 does not account for the precision of the BCF estimates. Hence, this raises the 276 question of using another criterion to decide whether a chemical substance is 277 bioaccumulative or not, and at which bioaccumulative capacity. 278

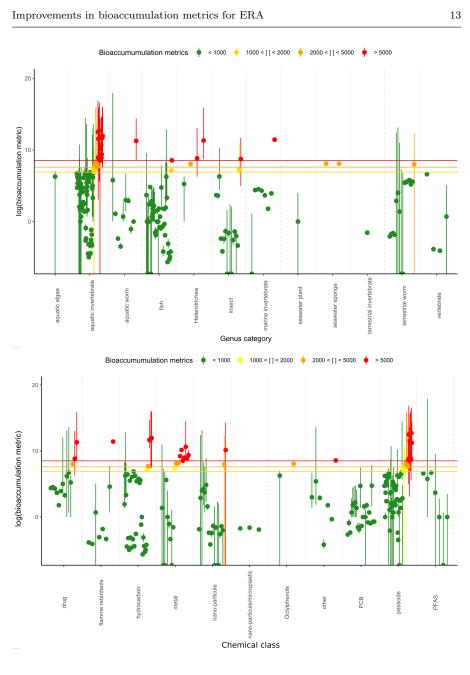


Fig. 2 Bioaccumulation metrics (in \log_{10} scale) according to genus (upper panel) and chemical (lower panel) categories. Dots represent medians of the bioaccumulation metrics, while vertical segments represent the associated 95% uncertainty intervals. Horizontal lines delineate regulatory threshold values used in the regulation (1000, 2000 and 5000, respectively) to classify chemical substances according to their bioaccumulative capacity. Bioaccumulation metrics are colored accordingly: in green when the metric is < 1000; in yellow if the metric is in [1000; 2000[; in orange if the metric is in [2000; 5000[; in red if the metric is > 5000.

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279 3.3 Towards improvements in ERA

Accounting for the precision of the bioaccumulation estimates relies on the use
of their probability distribution, that can be brought into play to define a new
decision criterion for the classification of chemical substances.
Instead of the median only, we first envisaged to consider the upper bound
of the uncertainty interval around the bioaccumulation metric estimates, that
is the 97.5% quantiles of their posterior probability distributions (referred as

Q97.5 in Table 1). This raised the question of an overestimation of the bioac-

 $_{\rm 287}$ $\,$ cumulative capacity of the chemical substances. Indeed, the number of BCF

values doubled from 22 to 45 in the "vB" category (Table 1). Such a criterion values doubled from 22 to 45 in the "vB" category (Table 1).

would classify 36.5% of the chemical substances as bioaccumulative (n = 77),

against only 22.7% (n = 48) with the usual criterion based on the median

estimate (referred as Q50 in Table 1). In terms of safety prediction, this could

²⁹² be acceptable but still based on a biased interpretation of the bioaccumulative

²⁹³ capacity.

Table 1 Number of chemical substances in each bioaccumulative capacity class according to the three decision criteria built on the 50%, the 75% or the 97.5% quantiles of the posterior probability distributions of the bioaccumulation metric estimates: Q50, Q75 and Q97.5, respectively. Abbreviations ℓB stands for low bioaccumulative, B for bioaccumulative and vB for very bioaccumulative, respectively.

Criterion	$\ell \mathbf{B} \ (BCF \in [1000; 2000[)$	B $(BCF \in [2000; 5000[)$	vB ($BCF > 5000$)
Q50	10	16	22
Q75	11	20	30
Q97.5	10	22	45

Again in the perspective of exploiting the uncertainty on bioaccumulation metric estimates to build a new classification criterion of chemical substances,

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we compared the use of the 75% quantile (referred as Q75 in Table 1) to the 296 use of the usual Q50 or the previous Q97.5, as an alternative compromise. 297 As illustrated on Figure 3, concerning the species Daphnia magna exposed to 298 phenanthrene (Wang et al., 2021), the Q50 classifies as "nB" (BCF < 1000), 299 while both the Q75 and the Q97.5 criteria classify it as "B" (BCF > 1000). 300 This example illustrates the need to consider the uncertainty to avoid "false 301 negatives", that is chemical substances as non bioaccumulative while they 302 have 75% of chance to be actually. Over the 211 chemical substances in the 303 TK database, the Q75 criterion classifies a total of 28.9% of the chemical 304 substances as "B" (n = 61). 305

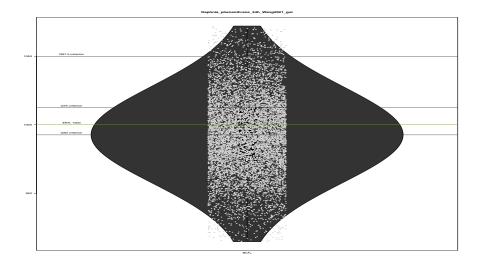


Fig. 3 Violin plot of the bioaccumulation metric posterior probability distribution for *Daphnia magna* exposed to phenanthrene (Wang et al., 2021). The green line symbolizes the threshold value at 1000 according to regulatory ERA, while grey lines stand for the 50th (Q50 criterion in table 1), 75th (Q75 criterion) and the 97.5th (Q97.5 criterion) percentiles of the distribution, respectively.

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Figure 4 shows another example with the genus *Enchytraeus* exposed to 306 silver nano-particles (Topuz and van Gestel, 2015), illustrating the possible 307 miss-classification of the bioaccumulative capacity of a chemical substance 308 according to the Q97.5 criterion. Indeed, instead of being classified as "B" 309 $(BCF \in [2000; 5000])$, silver nano-particles are considered as non bioaccumu-310 lative by both the Q50 (the ERA criterion) and the alternative Q75 criteria 311 (BCF < 1000). In this case, the difference in the classification comes from a 312 lack of precision of the BCF estimate which is associated with a large uncer-313 tainty range, that is the coefficient of variation (abbreviated as CV) is far over 314 0.5. This means that the BCF was delivered with a very dispersed probability 315 distribution (Figure 4. The formula for CV writes as follows: 316

317 $CV \simeq \frac{BCF_{\text{median}}}{97.5\% quantile - 2.5\% quantile}$

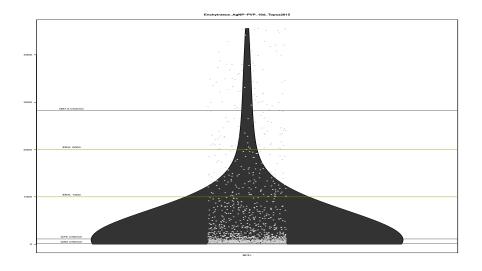


Fig. 4 Violin plot of the bioaccumulation metric posterior probability distribution for *Enchytraeus* exposed to silver nano-particles (Topuz and van Gestel, 2015). The green lines symbolize the threshold value at 1000 and 2000 according to regulatory ERA, while grey lines stand for the 50th (Q50 criterion), 75th (Q75 criterion) and the 97.5th (Q97.5 criterion) percentiles of the distribution, respectively (Table 1).

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Considering all the 211 data sets available, we performed the classification 318 of all chemical substances based on each of the Q50, Q75 and Q97.5 crite-319 ria, and compared the results in order to formulate sufficiently well-founded 320 recommendations. As illustrated on Figure 5 concerning the genus Anax ex-321 posed to chlorpyriphos (Rubach et al., 2010), the classification was the same 322 whatever the criteria. In particular, we assigned more than 90% (n = 190)323 of the chemical substances in the same class as done with the classical Q50 324 criterion (Table 1). These results support the use of the 75% quantile of the 325 posterior probability distribution of the bioaccumulation metric estimates as 326 a reasonable compromise between the classical median as currently required 327 by the regulatory ERA, and the upper bound of the uncertainty range, while 328 still accounting for the precision of the bioaccumulation metric estimates via 329 the three-quarter quantile. 330

To summarize the whole database based on our recommended Q75 crite-331 rion, we finally classified 5.2% (n = 11), 9.5% (n = 20) and 14.2% (n = 30)332 of the chemical substances as " ℓB ", "B" and "vB", respectively, confirming 333 71% of the chemical substances as non bioaccumulative. Regarding the preci-334 sion of all bioaccumulation metrics, we got a high variability among species 335 and chemical substances (see SI, see the .html file). Only considering the Q75 336 criterion, 37.4% (n = 79) of the chemical substances are associated with a 337 CV < 0.5, with the most high CV among the "nB"-classified chemical sub-338 stances (38.9%, n = 82/150). This reveals that when a chemical substances is 339 considered as bioaccumulative, what ever the class, the corresponding bioac-340 cumulation metric is precisely estimated in most of the cases. 341

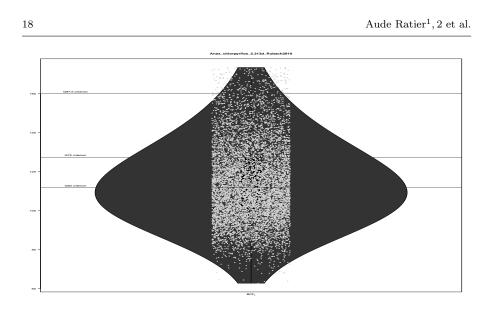


Fig. 5 Violin plot of the bioaccumulation metric posterior probability distribution for *Anax* exposed to chlorpyriphos (Rubach et al., 2010). The green lines symbolize the threshold value at 1000 and 2000 according to regulatory ERA, while grey lines stand for the 50th (Q50 criterion), 75th (Q75 criterion) and the 97.5th (Q97.5 criterion) percentiles of the distribution, respectively (Table 1).

342 4 Conclusion

Based on a meta-analysis of a TK database comprising 211 data sets, this 343 paper establishes how crucial it is to consider the uncertainty in classifying 344 the chemical substances according to their bioaccumulative capacity. Thanks 345 to the fitting of TK models under a Bayesian framework, delivering bioac-346 cumulation metrics as probability distributions, this paper gathers together 347 statistically-founded results towards the adoption a new criterion for this clas-348 sification. Indeed, it can be recommended to use the 75% quantile of the bioac-349 cumulation metric distributions in order to more precisely assign the chemical 350 substances into the four regulatory ERA categories, that is as non-, low-, 351 medium-, or very- bioaccumulative, and 90% in the same way as the current 352

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approach required today. Also associated with precise estimates of bioaccumulation metrics, this new criterion would strongly improve ERA if it would be adopted by regulatory bodies. In addition, it could easily be implemented into the MOSAIC_{bioacc} web service in support of the daily work of regulators when they need to quickly classify a set of several chemical substances.

358 Conflict of interest

³⁵⁹ The authors declare that they have no conflict of interest.

360 Consent for Publication

This manuscript has original research that has not been published previously and is not under consideration for publication elsewhere, in whole or in part.

363 Author Contributions

All authors contributed to the investigation of the TK database. Raw data collection and first analyses were performed by Aude Ratier and Sandrine Charles. The first draft of the manuscript was written by Aude Ratier and all authors commented on previous versions of the manuscript. All authors read and approved the final manuscript.

³⁶⁹ Supplementary Information and data availability

- ³⁷⁰ Supplementary information is available at https://doi.org/10.5281/zenodo.
- $_{\rm 371}$ $\,$ 5865150. All data used in this paper is downloadable from the MOSAIC $_{\rm bioacc}$
- web tool, directly from the associated TK database freely accessible at http://
- 373 lbbe-shiny.univ-lyon1.fr/mosaic-bioacc/data/database/TK_database.

374 html.

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505		egories (lower panel) available in the TK database available at
506		http://lbbe-shiny.univ-lyon1.fr/mosaic-bioacc/data/database/
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508	2	Bioaccumulation metrics (in \log_{10} scale) according to genus (up-
509		per panel) and chemical (lower panel) categories. Dots represent
510		medians of the bioaccumulation metrics, while vertical segments
511		represent the associated 95% uncertainty intervals. Horizontal
512		lines delineate regulatory threshold values used in the regula-
513		tion (1000, 2000 and 5000, respectively) to classify chemical
514		substances according to their bioaccumulative capacity. Bioac-
515		cumulation metrics are colored accordingly: in green when the
516		metric is < 1000 ; in yellow if the metric is in [1000; 2000[; in
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518		> 5000.

519	3	Violin plot of the bioaccumulation metric posterior probabil-
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521		(Wang et al., 2021). The green line symbolizes the threshold
522		value at 1000 according to regulatory ERA, while grey lines
523		stand for the 50^{th} (Q50 criterion in table 1), 75^{th} (Q75 criterion)
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529		threshold value at 1000 and 2000 according to regulatory ERA,
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543		ity distributions of the bioaccumulation metric estimates: Q50,
544		Q75 and Q97.5, respectively. Abbreviations ℓB stands for low
545		bioaccumulative, B for bioaccumulative and vB for very bioac-
546		cumulative, respectively