1	Endocrine disruption in aquatic systems: Up-scaling research to
2	address ecological consequences
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14	ABSTRACT
15	Endocrine disrupting chemicals (EDCs) can alter biological function in organisms at
16	environmentally relevant concentrations and are a significant threat to aquatic biodiversity,
17	but there is little understanding of exposure consequences for populations, communities and
18	ecosystems. The pervasive nature of EDCs within aquatic environments and their multiple
19	sub-lethal effects make assessments of their impact especially important but also highly
20	challenging. Herein, we review the data on EDC effects in aquatic systems focusing on
21	studies assessing populations and ecosystems, and including how biotic and abiotic processes
22	may affect, and be affected by, responses to EDCs. Recent research indicates a significant
23	influence of behavioural responses (e.g. enhancing feeding rates), transgenerational effects
24	and trophic cascades in the ecological consequences of EDC exposure. In addition,

25	interactions between EDCs and other chemical, physical and biological factors generate									
26	uncertainty in our understanding of the ecological effects of EDCs within aquatic ecosystems.									
27	We illustrate how effect thresholds for EDCs generated from individual-based experimental									
28	bioassa	ays of the types commonly applied using chemical test guidelines (e.g. Organisation for								
29	Econor	mic Co-operation and Development [OECD]) may not necessarily reflect the hazards								
30	associa	ated with endocrine disruption. We argue that improved risk assessment for EDCs in								
31	aquatic	e ecosystems urgently requires more ecologically oriented research as well as field-								
32	based a	assessments at population-, community- and food-web levels.								
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34	Key wo	ords: aquatic pollution, ecotoxicology, endocrine disrupting chemicals, food webs,								
35	popula	tions.								
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57 I. INTRODUCTION

VI.

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Future directions

Endocrine disrupting chemicals (EDCs) remain an active topic in contemporary 58 ecotoxicology due to their proven environmental impacts (Zhou, Cai & Zhu, 2010; Wang & 59 Zhou, 2013) and postulated health effects (Kabir, Rahman & Rahman, 2015). Over the past 60 decade published work on EDCs has provided a strong mechanistic understanding of 61 exposure effects (Colborn, vom Saal & Soto, 1993; Tyler, Jobling & Sumpter, 1998; Kloas et 62 al., 2009; Söffker & Tyler, 2012; Orton & Tyler, 2012; Tijani, Fatoba & Petrik, 2013). Far 63 less consideration, however, has been given to processes and interactions controlling the 64 effects of EDCs at broader ecological scales, including inter- and intra-specific interactions 65 within populations and food webs (Segner, 2011; Brodin et al., 2014; Schoenfuss et al., 66 67 2015). Understanding the effects of EDCs on processes operating at these broader scales is essential, but also challenging, because their effects can be pervasive and they are generally 68 sub-lethal in nature. Although EDCs can induce deleterious effects in a wide range of 69 70 organisms across different trophic levels (Brander, 2013), there is insufficient knowledge for environmental regulators to assess the impacts and risks posed by EDC pollution to 71

populations, communities and ecosystems (e.g. Mills & Chichester, 2005; Hallgren *et al.*,
2012).

Herein, we evaluate critically the known and potential effects of EDCs on natural ecological systems. We highlight a need for EDC research to incorporate processes and effects at broader spatial and temporal scales, illustrating how such studies have helped to advance our understanding of EDC impacts beyond common approaches to EDC testing. We also suggest an integrated research strategy for EDCs that develops previous designs from other pollutants to generate more environmentally relevant data. Finally, we consider further research needs to understand better the effects of EDCs on natural systems.

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82 II. THE BENEFITS OF UP-SCALING EDC RESEARCH

The requirement for information on population effects of EDC exposure to inform ecological 83 risk assessments has led to the extrapolation of individual-based experimental bioassays (e.g. 84 Jobling et al., 2002b; Miller & Ankley, 2004; Gutjahr-Gobell et al., 2006; Lange et al., 2008; 85 Brander et al., 2016). Such extrapolations assume, however, that the effects of EDC exposure 86 within individual-based bioassays generally show simple, direct and invariant relationships 87 with impacts on populations and communities, even if safety factors are used to account for 88 uncertainties associated with these extrapolations. Assessments involving wild populations, 89 however, demonstrate discontinuities between the results of individual- and population-level 90 91 assessments (Jobling et al., 2002a; Brown et al., 2005; Lange et al., 2011; Hamilton et al., 2014). Fundamental differences in the ecological processes represented within micro-, meso-92 and macroscale assessments (Fig. 1) are potentially responsible for this disparity. 93 94 Specifically, these differences include the nature of the EDC exposure regime, possible compounding environmental influences (e.g. multiple stressors), and the fact that multiple 95 effect mechanisms may operate through trophic interactions across food webs at the 96

macroscale (Hamilton et al., 2016a). There are several potential inconsistencies in findings 97 about endocrine disruption from different biological, spatial and temporal scales. For 98 99 example, cause-effect relationships reflect the methods used and scales at which studies are completed and this creates a challenge in determining mechanistic relationships and emergent 100 effects at broader spatio-temporal extents. As an example, feminisation at the individual level 101 102 would suggest significant potential population effects, but studies at broader spatial scales 103 have indicated that population-level effects depend on mating-system dynamics (White et al., 104 2017). On the one hand, the low cost of sperm production relative to eggs means that males 105 are able to fertilise multiple females, thus the feminisation of males may have little effect on population dynamics (White et al., 2017). On the other hand, mating systems may prevent 106 male promiscuity, meaning that feminisation and minor alterations in the sex ratio result in 107 negative effects on populations (White et al., 2017). Currently, little consideration is 108 generally given to natural complexity in ecological and toxicological processes within 109 experimental research designs (see Barton, 2003). Models developed for up-scaling from 110 individual-based assessments to population scales are therefore inherently weak, and may 111 even be flawed, as they provide limited appreciations of wider controls on higher levels of 112 biological organisation. Factors such as density-dependence, adaptation, trophic interactions, 113 likelihood of population exposure (habitat preferences), as well as species-specific life-114 history traits of organisms, are all likely to have a significant impact on endocrine disruption, 115 116 yet none of these characteristics are considered in common experimental assessments used to investigate the ecological impacts of EDC exposure. 117 Research that considers processes over longer periods of time (e.g. entire life cycles) and at 118

120 limitations associated with most current experimental ecotoxicology bioassays (Geiszinger *et*

higher levels of biological organisation (e.g. populations and food webs) overcomes several

121 *al.*, 2009). The complexity associated with analysis of mesocosm and field scenarios,

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however, has restricted the uptake of these research designs. Furthermore, many field studies 122 are characterised by correlation and weak inference in comparison to well-established 123 124 mechanistic knowledge developed under more-controlled experimental conditions. A combination of experimental and field-based studies across a range of ecological scales is 125 thus required for an improved understanding of population- and food-web-level responses to 126 EDC exposure. This approach has, however, had relatively little uptake (Patiño & Carr, 2015) 127 128 and studies assessing the effects of EDCs at community and food-web scales remain scarce 129 (Boxall et al., 2012). Contemporary studies have consequently called for a greater focus on 130 broader scale ecological and toxicological processes (Brodin et al., 2014; Kidd et al., 2014).

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132 III. ADVANCES IN BROAD-SCALE EDC RESEARCH

Here, we assess critically recent findings derived from EDC research focusing on processes
operating at broad spatial and temporal scales and highlight the limitations associated with
using experimental bioassays conducted without due consideration of natural system
dynamics. This builds upon previous conceptual reviews of the role of theoretical ecology in
enhancing ecotoxicological studies (e.g. Relyea & Hoverman, 2006).

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139 (1) Biotic interactions and trophic transfer of EDCs through food webs

The effects derived from EDC exposure within natural systems are variable and influenced by biological processes including competitive interactions and predation. Only a few examples exist regarding how biotic factors affect the severity of endocrine disruption, but a suite of processes appear to provide an important influence on the risk associated with EDC exposure within ecosystems. The behaviour of organisms in response to EDC exposure, in particular, can result in important ecological effects and in some cases, behavioural changes enhance adverse effects of EDC exposure (Melvin & Wilson, 2013). As well as providing the

potential to exacerbate an effect at higher levels of biological organisation, interactions 147 among individuals can also buffer the observed effects of EDC exposure. An example of this 148 149 is density-dependent compensatory effects in zebrafish Danio rerio (Hamilton) populations that have been shown to alleviate negative individual reproductive effects of octylphenol 150 exposure (Hazlerigg et al., 2014). Effects such as those detailed above are rarely considered 151 or captured in laboratory-based studies and the consequences of these alterations could 152 153 exacerbate the effects of EDCs at higher levels of biological organisation and within natural systems. 154

155 Biotic and abiotic processes can influence the trophic transfer of EDCs within aquatic ecosystems. Alkylphenols, pyrethroids, polychlorinated biphenyls (PCBs), polybrominated 156 diphenyl ethers (PBDEs) and diclofenac have been shown to partition, accumulate and 157 magnify within components of aquatic food webs (see Table 1) and exhibit different entry 158 and transfer pathways within the environment (Burreau et al., 1997, 2006; Correa-Reyes et 159 al., 2007; Corcellas, Eljarrat & Barcelo, 2015; Muggelberg et al., 2017). Many EDCs are 160 hydrophobic in nature and readily partition out of the water column through adsorption to 161 both suspended and benthic sediments (Petrović et al., 2001). Consequently, a significant 162 proportion of the total pollutant load entering aquatic food webs is likely to be through 163 benthic taxa interacting with sediments (e.g. sediment ingestors) (Brooks, Gaskell & Maltby, 164 2009; Wu et al., 2009). Dietary transfers, however, are not the main route of uptake for many 165 EDCs, and for selected compounds (e.g. carbamazepine and diphenhydramine) direct 166 adsorption from the water column is a major route for their bioaccumulation (Du et al., 2014, 167 2015, 2016). This transfer of EDCs directly from the water column into aquatic organisms 168 can occur either through passive adsorption, whereby the skin and respiratory surfaces enable 169 diffusion or via assimilation of EDCs adhering to suspended organic matter (Zhou et al., 170 2007). In natural systems, it is likely that most EDCs enter organisms by multiple uptake 171

pathways. Thus, EDC exposure within natural systems may be intermittent, as in dietaryintake, or possibly continuous *via* the water column.

174 Upon entry into organisms the transfer of EDCs within aquatic food webs is affected by a series of biological controls, including the organism's physiology, and *via* biotic interactions. 175 The biological traits of organisms, including functional feeding guilds, influence the 176 177 bioaccumulation, biomagnification and transfer of EDCs (Muñoz et al., 2009; Damásio et al., 178 2011). Bioaccumulation can vary across trophic levels (Ruhí et al., 2015), but even within the 179 same trophic level individual biological traits, including size, can influence EDC uptake 180 (Sidney et al., 2016). Many organisms exhibit an ability effectively to eliminate selected EDCs from tissues, thereby mitigating their accumulation via diet or water and subsequent 181 transfer (Norman et al., 2007; Al-Ansari et al., 2013). These assessments demonstrate the 182 importance of biological interactions in the trophic transfer of EDCs within natural systems 183 and indicate why responses may deviate from those expected from experimental, laboratory-184 based exposure assessments on individual organisms. Further research is, however, required 185 to understand better the influence of biological traits on the bioaccumulation and ecological 186 risk of EDCs. 187

Interactions between the direct effects of endocrine disruption and the subsequent transfer of 188 EDCs through ecosystems may also occur, supporting that alterations in individual-level 189 effects may have consequences for wider biological systems (Brooks et al., 2009). A specific 190 191 illustration of this is provided by Brodin et al. (2013, 2014) where an increased feeding rate of perch (Perca fluviatilis L.) in a behavioural response to oxazepam exposure resulted in 192 enhanced consumption of its damselfly prey (Coenagrion hastulatum Charpentier), and in 193 turn an increase in the transfer and bioaccumulation of oxazepam. These examples illustrate 194 that ecological risks for some EDCs that affect ecosystem processes (e.g. feeding behaviour 195

and bioaccumulation potential) may be greater than commonly appreciated within aquaticecosystems.

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199 (2) Adaptation to EDC exposure

Individuals, populations and food webs have varying levels of resilience to environmental 200 stressors (Harrison, 1979), but in most cases organisms in aquatic ecosystems are able to 201 202 persist at low levels of stress, even in multi-stressor environments (Vinebrooke et al., 2004). There is little field-based information, however, on the ecological and evolutionary resilience 203 204 of individuals and populations to endocrine disruption, although the presence of adaptation is widely displayed within experimental assessments (see Wu, Siu & Shin, 2005). Many 205 existing studies do not assess adaptations directly, instead indicating the reduction in effect 206 207 size over the duration of exposure, which occurs more rapidly for individuals in comparison to populations and communities (Wu et al., 2005). Several field studies have identified 208 populations of aquatic organisms resistant to certain EDCs. For example, Weston et al. 209 (2013) indicated that point mutations at the pyrethroid target site (voltage-gated Na⁺ channel) 210 in Hyalella azteca (Saussure) populations meant that resistant individuals did not experience 211 212 the neurotoxic effects observed in non-resistant populations, instead exhibiting oxidative stress only at considerably higher pyrethroid concentrations. Varying levels of resistance 213 were found across several populations. Adaptation has also been observed within fish 214 215 assemblages (Hamilton et al., 2016b). Both the Atlantic tomcod (Microgadus tomcod Walbaum) and the Atlantic killifish (Fundulus heteroclitus L.) can adapt to polycyclic 216 aromatic hydrocarbon (PAH) and PCB exposure in natural systems (Clark et al., 2010; 217 Wirgin et al., 2011), but through different mechanisms. In M. tomcod a six-base deletion in 218 the aryl hydrocarbon receptor 2 (AHR2) restricted inducible gene expression and was 219 responsible for the observed resistance to EDC exposure (Wirgin et al., 2011). In 220

comparison, resistance in *F. heteroclitus* individuals was generated by single nucleotide
polymorphisms in the regulatory regions of the cytochrome P4501A gene (Clark *et al.*, 2010;
Reid *et al.*, 2016).

Resistance, and/or adaptation has significant implications for the potential broad-scale effects 224 of endocrine disruption in aquatic systems. A recent example in *H. azteca*, showed that 225 populations pre-exposed to the pyrethroid pesticide Permethrin were able to persist under 226 higher environmental concentrations (>210 ng l^{-1}) than those populations which were not 227 pre-exposed (Muggelberg et al., 2017). This adaptation meant that resistant individuals 228 229 provided a source of dietary exposure for fathead minnows (*Pimphales promelas* Rafinesque) under conditions within which non-resistant individuals cannot survive. Within natural 230 systems, adaptation of individuals or populations leads to an enhanced risk of 231 bioaccumulation with increasing concentrations of EDCs. Adaptation to endocrine disruption 232 indicates that organisms may be able to persist at environmentally relevant concentrations of 233 EDCs, yet it also suggests potential for increased flux of EDCs through food webs. Changes 234 in the bioaccumulation and transfer of EDCs potentially lead to increases in the body burden 235 of higher trophic-level organisms, increasing the likelihood of adverse effects across the 236 237 aquatic food web.

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239 (3) Long-term, life-cycle and transgenerational EDC effects

There have been relatively few assessments of EDCs for long exposure durations, over full organism life cycles and/or over multiple generations, even though many organisms will be exposed for prolonged periods of time. Chronic exposure studies that have been undertaken have provided several significant advances. Firstly, in most cases they have shown that the effects are greater than for short-term exposures (Keiter *et al.*, 2012; Tassou & Schulz, 2013). Secondly, different health effects have been identified for longer-term exposures in

comparison to short-term exposures. For example, for 17α -ethinyloestradiol (EE2) exposure, 246 effects reported on mating behaviour, growth and survival in D. rerio individuals differed 247 248 between exposure periods of 0-21 and 0-75 days post-fertilisation (Segner et al., 2003). Thirdly, unanticipated effects have been identified following chronic exposures to EDCs. 249 250 Exposure of rainbow trout (Oncorhynchus mykiss Walbaum) eggs to an environmental oestrogen, bisphenol A (BPA), over a range of concentrations including 300 and 3000 ng l⁻¹ 251 resulted in lower energy levels in larvae to first feeding, reductions in specific growth and 252 restricted food conversion ratios (Birceanu, Servos & Vijayan, 2015). Finally, chronic 253 exposure studies have helped to highlight life-stage-specific susceptibilities to the effects of 254 EDCs. Schäfers et al. (2007) illustrated that the chronic effects on sexual differentiation in D. 255 *rerio* resulting from lifelong exposure to 10 ng l⁻¹ of EE2 were more pervasive than the 256 reversible effects induced by exposure extending over the period of gonadal differentiation 257 258 only.

It must be emphasised that not all EDC effects are necessarily permanent; some are transient 259 in nature and the organism may recover after the exposure is removed. Examples include the 260 261 reported partial recovery from the effects of EE2 (5 ng \vdash^1) on gonad differentiation in *D*. rerio after a five-month depuration period post-EE2 exposure (Nash et al., 2004). Complete 262 recovery of biological function was observed in a full-life-cycle analysis of D. rerio after 263 exposure to EE2 (3 ng l⁻¹) (Fenske *et al.*, 2005). Here exposure to EE2 from the fertilised egg 264 stage for 118 days post-fertilisation inhibited gonad differentiation in males, but a 58-day 265 post-exposure period of depuration resulted in resumption and subsequent completion of 266 testicular differentiation. Reproduction in D. rerio has also been shown to recover completely 267 after exposure to zearalenone; exposure to 1000 ng l⁻¹ zearalenone for 140 days induced a 268 female shift in the population sex ratio, but a subsequent period of depuration for 42 days 269 resulted in recovery of relative fecundity (Schwartz et al., 2013). The ability to recover will, 270

in part, depend on EDC exposure concentration and the consequent nature and severity of 271 effect(s). In other studies on D. rerio, e.g. Schäfers et al. (2007) and Baumann et al. (2014), 272 individuals did not show full recovery following EE2 exposure at 9.3 ng l^{-1} or trenbolone (an 273 androgen used as a growth promotor for cattle in the USA) exposure at 30 ng l⁻¹. The length 274 of both exposure and period for depuration thus appear to be important in weighing up the 275 potential for biological impacts of EDCs in natural systems. The fact that EDCs can act 276 277 through multiple pathways means that it is especially difficult to identify chronic and lifestage-specific effects (Sohoni & Sumpter, 1998). Pinpointing these effects is further hindered 278 279 by the fact that effect mechanisms for many EDCs are not well defined. As an example, phthalate esters [e.g. di-n-butyl phthalate and di(2-ethylhexyl)phthalate] have been identified 280 as both oestrogen receptor agonists and androgen receptor antagonists (Takeuchi et al., 281 2005). Furthermore, exposure to these compounds maintains a range of individual-level 282 effects, including alterations in cellular proliferation, biosynthesis and apoptosis, as well as 283 several immune responses (Milla, Depiereux & Kestemon, 2011; Mankidy et al., 2013). 284 Thus, when considering the spatio-temporal dynamics of EDC pollution within aquatic 285 systems it is important to assess all the effects that may manifest. In natural systems, 286 exposure to EDCs in periodic urban run-off inputs may result in different effects compared 287 with continuous emissions from wastewater treatment works (WwTWs). 288 Transgenerational studies on the effects of EDCs further highlight the importance of 289 290 considering temporal scale in effect analyses. There is a mounting consensus that EDC exposure effects can span multiple generations, and may induce different impacts in offspring 291 compared with the parental generation (Skinner, Manikkam & Guerrero-Bosagna, 2011; 292 293 Bhandari, vom Saal & Tillitt, 2015). Some of the adverse effects observed in subsequent generations have been shown not to be induced through the direct modulation of DNA 294 sequences, but rather through permanent alterations in the epigenome promoting 295

transgenerational phenotypes (Skinner et al., 2011; Head, 2014). This mechanism can 296 promote transmission of potentially susceptible phenotypes to the offspring of affected 297 298 organisms, and may enhance the adverse impacts of EDCs within subsequent generations (Sowers et al., 2009). Exposure during early life or at particularly susceptible life stages can 299 also have effects that span the lifetime of the affected organism and potentially lead to 300 301 adverse effects in subsequent generations (Head, 2014). These changes can be through 302 somatic and gametic effect pathways (Faulk & Dolinoy, 2011). Consequently, epigenomic 303 changes resulting from EDC exposure may lead to transgenerational effects, and possibly 304 different population-level impacts within natural systems because of cumulative adverse effects in multiple generations (Bernal & Jirtle, 2010). 305

Of note is the fact that contemporary assessments of EDCs in the laboratory are confined to a 306 307 restricted range of short-lived species suitable for experiments; for fish, notably D. rerio, P. promelas and medaka (Oryzias latipes Temminck & Schlegel). Whilst these taxa are 308 convenient as study models, they may not necessarily allow the accurate prediction of effects 309 within populations of longer-lived organisms which may accumulate greater levels of EDCs 310 over longer periods of time and have slower generational turnover, and thus a lower ability to 311 adapt in response to toxicological impacts. Further efforts to understand long-term exposure 312 effects across a wider range of taxa are urgently required. 313

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315 (4) Interactive mixtures of EDCs

Wastewater effluents and other pollutant sources are often composed of highly complex
mixtures, and interactions between EDCs and of EDCs with other chemicals could alter their
biological effects (Keiter *et al.*, 2012; Schoenfuss *et al.*, 2015). The potential for additive
effects of EDCs and other chemicals is significant. Most experiments on EDCs, however,
have assessed only the effects of individual chemicals, with a small number of exceptions

(e.g. Thorpe et al., 2003; Brian et al., 2007). A range of adverse, sub-lethal impacts may 321 occur that are not always predictable from assessments of individual components 322 (Kortenkamp, 2007; Viñas, Jeng & Watson, 2012) or via simple additive-effect modelling 323 (Silva, Rajapakse & Kortenkamp, 2002). Compounds with dissimilar modes of action may 324 induce novel effects, operating through multiple mechanisms (Viñas et al., 2012). Sárria et al. 325 (2011) demonstrated that exposure to EE2 and tributyltin (TBT) caused alterations in the 326 327 behavioural responses of juvenile black-striped pipefish (Syngnathus abaster Risso). TBT depressed the burst-swimming response known to result from EE2 exposure, whilst EE2 328 329 influenced the alterations in the time spent in secluded areas generated by high concentrations of TBT. Consequently, when mixtures of EDCs combine with processes such as competition 330 and predation, a range of complex and often unpredictable effects can result. 331 There are also reports of a non-monotonic dose–response relationship resulting from 332 exposure to EDC and their mixtures (Vandenberg et al., 2012). Non-monotonic dose-333 response relationships are not unique to EDCs, but they have been reported more frequently 334 for EDCs than for other toxicants (Vandenberg, 2014), in part reflecting the use of more 335 sensitive endpoints or the wider range of concentrations tested (vom Saal et al., 2010; 336 Vandenberg et al., 2013; Vanderberg, 2014). Controversially, it has been proposed that 337 hormesis, where marked beneficial low-dose effects are observed, may be responsible for the 338 non-monotonic dose-response relationships (Calabrese, 2005). This conclusion has been 339 340 disputed, with some arguing that the impacts of oestrogenic EDCs always remain negative irrespective of concentration (Weltje, vom Saal & Oehlmann, 2005). Many examples exist of 341 non-monotonic dose-response relationships for EDCs with markedly different 342 physicochemical properties. Pyrethroid pesticides, for example, generally exhibit greater 343 negative effects at lower concentrations (Brander et al., 2016), and BPA shows a non-344

monotonic transcriptional-effect response (Villeneuve et al., 2012). There appears to be a 345 346 wide range of effects that exhibit non-monotonic relationships with several EDCs. 347 The identification of non-linear, non-monotonic, and in some cases hormetic, relationships across many studies has led some authors to suggest that effects observed at high EDC 348 349 concentrations may not represent those at environmentally relevant concentrations or for 350 mixtures of EDCs (Beausoleil et al., 2013; Vandenberg, 2014). Thus, the lowest observed 351 effect levels (LOELs) recorded within experimental bioassays may not accurately extrapolate 352 to the lowest concentrations present within natural systems (Vandenberg *et al.*, 2012). It has 353 been suggested that alternative relationships (U- or inverted U-shaped) may better reflect effects associated with environmental EDC exposure (Vandenberg et al., 2014; Vandenberg 354 & Bowler, 2014; Zoeller & Vandenberg, 2015). This challenges the concentration-specific 355 understanding of endocrine disruption within natural systems and poses a significant 356 challenge for risk assessment if true (Futran Fuhrman, Tal & Arnon, 2015). 357

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359 (5) EDCs within the context of multiple stressors

Accounting for environmental variation is crucial in determining the effects of EDC exposure 360 within natural systems, as multiple covariant environmental variables influence observed 361 effects within natural systems (Daughton, 2004; Damásio et al., 2011). Previous assessments 362 have used the statistical and environmental control provided by experimental bioassays to 363 eliminate confounding relationships between influential variables present within natural 364 environments. However, interactions between multiple stressors ultimately dictate the relative 365 severity of EDC exposure and subsequent ecological risk within ecosystems (Hooper et al., 366 2013). Recent research has demonstrated the importance of assessments incorporating and 367 accounting for exogenous environmental characteristics, such as water temperature, 368 physicochemical conditions and biotic interactions. These abiotic and biotic stressors may 369

interact with one another as well as with EDCs to affect the outcome in exposed organisms. 370 A modelling study by An et al. (2009) assessing wild roach (Rutilus rutilus L.) populations 371 372 demonstrates the potential for interactive effects of multiple stressors. Here, the feminisation of individuals generated by endocrine disruption appeared to have negligible effects on 373 population extinction risk, yet the combination of exposure and selective fishing practices 374 375 resulted in significant increases in local population extinction rates. The feminising effect of 376 oestrogenic EDCs in isolation does not always result in significant population effects (see 377 Hamilton *et al.*, 2016*a*) and in some cases the population-level threats from masculinisation 378 are greater than from feminisation. The relative threat of both feminisation and masculinisation, however, is dependent on the optimal sex ratio of individual populations 379 (White et al., 2017). Fish species exhibiting a non-linear mating function (non-linear 380 response of reproductive capacity to changing sex ratio) did not exhibit reduced reproductive 381 output when few males were present, however, the overall reproductive output of the 382 population was significantly reduced by declines in the relative abundance of females (White 383 et al., 2017). 384

Studies assessing temperature and EDC exposure indicate that stressor-EDC interactions may 385 386 take multiple forms, with EDC exposure in some cases driving alterations in the effects of temperature increases (Jenssen, 2006), while in other cases temperature determines the 387 severity of ecological effects derived from EDC exposure (Moe et al., 2013). The importance 388 389 of interactions between two stressors has been relatively well demonstrated by contemporary research, yet these studies are still not representative of the true complexity present within 390 natural systems. More recent research has attempted to encapsulate a greater number of 391 392 stressors. For example, Brown et al. (2015) showed that a combination of EDC exposure, temperature increases and inbreeding led to a significantly skewed sex ratio in D. rerio 393 populations. Increases in temperature (28-33 °C), clotrimazole exposure (2000 and 10000 ng 394

1⁻¹) and inbreeding together had an additive effect, with a marked increase in the male-skew 395 of populations relative to the effects generated by individual stressors. The results of 396 397 multiple-stressor studies have indicated additive and synergistic interactions between stressors and endocrine disruption, but this depends on the level of biological organisation 398 included (Fischer, Pomati & Eggen, 2013; Sulmon et al., 2015). Consequently, such 399 processes are significant in altering the observed effects of EDC exposure whilst also 400 401 demonstrating the need for analyses to encapsulate the effects of ecological processes on sub-402 lethal EDC impacts.

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404 (6) Effects of population genetics on responses to EDC exposure

Interactions between the wider spatial connectivity of aquatic environments (e.g. isolated and 405 406 connected populations) and chemical contamination can have marked effects on the genetic diversity present within populations (Bickham et al., 2000). Genetics, specifically genetic 407 diversity, can play an important role in determining the effects of EDC exposure, with 408 reductions in genetic diversity derived from inbreeding potentially increasing the adverse 409 ecological effects of EDC exposure (Bickley et al., 2013). Söffker, Stevens & Tyler (2012) 410 reported that despite a generally similar response of genetically divergent D. rerio 411 populations to EE2 exposure, differences in their breeding biology and response sensitivity 412 were apparent. Inbreeding within laboratory fish stocks is a major issue for experimental 413 414 assessments of EDCs, especially when intending to inform further research in systems involving outbred individuals (Brown et al., 2009). Although perhaps of limited value for 415 building understanding of the effects of EDCs in outbred populations, experimental bioassays 416 using inbred individuals may be useful for indicating the increased susceptibility of isolated 417 natural populations to EDC exposure. In the event of habitat reconnection, whereby inbred 418 and outbred populations interact, adverse impacts on fertility within inbred populations can 419

420 facilitate a reduction in reproductive output of inbred individuals (Bickley *et al.*, 2013).

421 Assessments analysing interactions between genetic diversity and endocrine disruption within

natural populations however remain scarce, and future research is required to test several
hypotheses relating to genetic diversity and endocrine disruption across the wider aquatic
environment.

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426 (7) Trophic cascades and other indirect effects of EDCs

427 Direct effects of endocrine disruption may cause alterations in processes and interactions 428 within aquatic ecosystems, in turn generating indirect effects across wider levels of biological organisation (Relyea & Hoverman, 2006; Schulz et al., 2015). Such secondary effects may 429 result from changes in competition and predation interactions within food webs, and 430 subsequent release from biotic stressors (Knight et al., 2005). Similar trophic cascades have 431 been identified to result from other anthropogenic contaminants, such as petroleum 432 hydrocarbons and heavy metals (Fleeger, Carman & Nisbet, 2003). Very few studies, 433 however, have assessed these phenomena for EDCs. These indirect processes could alter the 434 perceived impacts of EDC exposure within natural populations, as well as affect the transfer 435 of EDCs within food webs. Indirect effects may occur through several mechanisms. Knapp et 436 al. (2005) demonstrated that changes in nutrient fluxes resulting from invertebrate mortality 437 in response to deltamethrin exposure (2000 ng l^{-1}) increased microbial community biomass. 438 439 A more commonly observed indirect mechanism is provided by the adverse effects of EDC exposure within predator assemblages and a subsequent top-down cascade through the food 440 web. Alterations in the structure of invertebrate communities have been recorded in response 441 to failed recruitment of secondary-consumer fish species when an entire Canadian lake was 442 dosed with EE2 (5–6 ng l⁻¹) over a period of three summers (Kidd *et al.*, 2014). A similar 443 example exists in a differently structured ecosystem, with endocrine disruption in R. rutilus 444

populations resulting in a reduction in predation of phytoplankton and increased copepod
abundance (Hallgren *et al.*, 2014). The indirect effects of endocrine disruption and their
influence over multiple trophic levels further indicates the potential for the observed effects
of EDC exposure within natural systems to deviate from those predicted from experimental
laboratory bioassays.

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451 IV. LIMITATIONS OF EDC IMPACT ASSESSMENTS

The results of assessments of the impacts of EDCs at broad spatial and temporal scales depart 452 453 significantly from predictions from laboratory-based experimental studies. These results highlight: (1) the limitations of using individual-based bioassays to predict the effects of 454 EDCs at population- and food-web scales, (also see Forbes et al., 2010; Hommen et al., 455 2010), and (2) the need for research at a range of spatial and temporal scales to advance 456 knowledge of broad-scale ecological effects and risk assessment. The restricted scope of 457 common experimental assessments has been highlighted previously (Matthiessen, 2008; 458 Lecomte et al., 2013), with calls for additional data to inform existing protocols and 459 enhanced higher-tier tests to replace unsuitable testing methods (Taenzler et al., 2007). 460 Although frameworks such as the OECD guidelines promote an increase in the complexity of 461 assessments (Gourmelon & Ahtiainen, 2007), the methodologies used in these assessments 462 inherently simplify the large range of controls on the effects of EDCs present within natural 463 systems. Population-level interactions, including density-dependent relationships such as 464 intra-specific competition, provide inherent controls on the effects of EDC exposure within 465 the environment, yet these controls remain absent from ecological impact and risk 466 assessments (Mills & Chichester, 2005). The low ecological complexity inherent in these 467 protocols therefore appears to provide a major constraint on the accuracy and wider 468 applicability of such tests. 469

Models developed from standard, individual-based bioassay protocols currently provide 470 limited value for the investigation of the effects of EDCs within natural systems. As 471 472 identified by Hazlerigg et al. (2014), isolation of the effects of chemical-mediation from other sub-lethal effects may underlie the underestimation of population-level impacts in 473 model scenarios. Although population-level models are suggested as a method for generating 474 environmentally relevant predictions across natural systems (Forbes, Calow & Sibly, 2008; 475 476 Forbes et al., 2010, 2011), extrapolating from overly simplified experimental data must be done with caution. Furthermore, the availability of limited data at higher levels of biological 477 478 organisation (e.g. populations) restricts the validation of model simulations (Rose et al., 1999; Forbes et al., 2008; Raimondo et al., 2009). The application of these models to the 479 prediction of EDC effects across aquatic environments thus remains prone to inaccuracies 480 (Munns et al., 2008). 481

482

483 V. THE NEED FOR MULTI-TIER INTEGRATED RESEARCH FOR STUDIES ON 484 EDCS

Low environmental concentrations of EDCs, coupled with their high propensity for sub-lethal 485 impacts, means that assessments at broader scales are essential for understanding the true 486 implications of EDC exposure. Nonetheless, the complex mechanisms through which 487 endocrine disruption can occur requires a detailed causal understanding which is difficult to 488 derive from large-scale studies (e.g. mesocosm or field assessment) (Schindler, 1998; Forbes 489 et al., 2010). The requirement for a multi-tiered research strategy may apply to all chemicals, 490 but is arguably most relevant to EDCs due to their wide range of sub-lethal effects that 491 operate at different ecological scales, together with their potential for multiple biotic and 492 abiotic interactions within and among spatial and temporal scales. The need to develop a 493

cohesive, broad-scale biomonitoring strategy is frequently identified in reviews of 494 495 ecotoxicological risk assessments (Besse, Geffard & Coquery, 2012; Gavrilescu et al., 2015). 496 Knowledge acquired at multiple spatial and temporal scales provides a suitable framework to mitigate previous limitations and to increase our understanding of EDC effects over wider 497 ecological scales. Similar integrated research has proved effective when assessing the 498 499 complex effects of stressors within a range of ecosystems, including multiple stressors in 500 freshwater systems (Altshuler et al., 2011) and heavy metals in coastal areas (Vlahogianni et 501 al., 2007). In the case of endocrine disruption, such a focus will enable an increase in 502 mechanistic knowledge at broad scales and the development of environmentally relevant experimental bioassays (Fig. 2). The product of this framework is environmentally relevant 503 knowledge at a range of scales, enabling the provision of suitable information (and 504 505 uncertainties) to practitioners and managers, potentially facilitating a reduction in adverse EDC effects across aquatic environments. 506

As in other research fields (see Culp et al., 2000), experiments on individuals can initially be 507 used to understand the direct impacts of stressors at the organism level, and these can then be 508 translated to research designs operating at broader scales. The multi-tiered research strategy 509 that we propose here, unlike other more-specific ecosystem-based strategies, is applicable to 510 a wide range of ecosystems and a suite of EDCs. Furthermore, it surpasses previous 511 methodological designs which focus more on the identification of ecological risk (using 512 513 experimental bioassays) and subsequent biomonitoring programs (e.g. Maruya et al., 2013), rather than providing a framework for understanding the risks within all levels of biological 514 organisation across ecosystems. Microcosm assessments within this research strategy allow 515 for an assessment of EDC exposure on reproductive morphology, physiology and behaviour, 516 in turn allowing for mechanistic knowledge at the organism and sub-organism scales. 517 Similarities and discrepancies identified between individual- and population-level 518

assessments can in turn indicate the population-level processes and controls (e.g. density 519 dependence and habitat-mediated exposure) influencing the effects of EDCs within 520 521 populations of aquatic organisms. Significant effects identified at the population level can be used to pinpoint areas of research suitable for further individual-based studies. In terms of 522 food-web assessments, the initial direct effects identified within individual-based assessments 523 524 can indicate the potential for indirect effects and trophic cascades, allowing for the derivation 525 of a suitable research design to identify these processes within natural systems. Furthermore, 526 the high replicability and mechanistic understanding developed within individual-based 527 studies provides a valuable tool for broad-scale assessment, enabling causal relationships to be derived for processes observed within aquatic food webs. The combination of individual-, 528 population- and food-web-level analyses can therefore enable improved realism of 529 investigations, and facilitate up-scaling of results to suitable levels for utilisation by 530 practitioners. 531

532

533 VI. FUTURE DIRECTIONS

534 (1) Spatial variation in EDC concentrations across aquatic environments

Contemporary research focuses on up-scaling EDC exposure to populations and food webs 535 within aquatic environments. The spatial coverage of these assessments, however, is 536 restricted when using individual systems to exemplify the wider conditions present across the 537 landscape. An example of this is the focus on WwTWs and their downstream impacts across 538 aquatic systems. A focus on wild populations and the effects of regulated effluent discharges 539 (containing EDCs) has made significant contributions to establishing the effects of effluent 540 discharges on aquatic organisms across aquatic environments. However, a focus on WwTWs 541 discharges has also led to limitations in our understanding of the spatial variation in EDC 542 occurrence and their impacts within and between different types of aquatic systems. Up-543

scaling research strategies to landscape scales to understand these spatial variations is much 544 needed to extend our knowledge of the effects of EDCs within natural systems. This will 545 546 enable improved impact and risk assessment, with practitioners able to assess more accurately the degree to which potential concerns vary across the aquatic environment. 547 Water-quality data regarding WwTWs discharges are available in many countries, 548 549 consequently high-risk WwTWs can be targeted for regulation and remediation. A range of 550 techniques are available to achieve this objective, including spatial and statistical modelling. 551 Modelling at extremely broad scales has identified variations in emission of steroidal 552 oestrogens between catchments, highlighting spatial variation in effects (Zhang et al., 2014). Furthermore, a significant role of mixing zones in determining the distribution of EDCs has 553 been identified at high resolutions (~500 m) (Pagsuyoin, Lung & Colosi, 2012). Assessments 554 investigating intra-catchment variation, along aquatic continuums and among systems, 555 however, are scarce. Understanding how EDC concentrations and subsequent exposure varies 556 557 at this scale is extremely important for River Basin Management strategies currently employed by water managers. 558

559

560 (2) EDC transfers across food webs

A detailed understanding of the transfer of EDCs across entire aquatic food webs is not yet 561 available, with studies predominantly focusing on bioaccumulation and biomagnification of 562 EDCs within upper trophic levels (Berglund, Nyström & Larsson, 2005). Assessments aiming 563 to evaluate entire food webs are generally restricted to a small range of organisms 564 representing several trophic levels. Controls on food-web organisation, such as environmental 565 conditions, may significantly influence EDC bioaccumulation, biomagnification and effects, 566 whilst a range of other biological factors also provide important regulatory impacts. The 567 extent to which these factors enhance (or mitigate) the transfer of toxicants through food 568

webs, however, remains relatively unknown. Moreover, although existing studies document 569 relatively variable relationships between biological controls and bioaccumulation of different 570 571 EDCs across aquatic food webs, explanations for such variability are absent. Future work is required to detail the specific pathways of accumulation and magnification throughout the 572 lower trophic levels to understand the routes of dietary EDC exposure and biomagnification 573 574 within higher trophic-level organisms. The first stage will be identifying the role of biotic and 575 EDC-specific processes in controlling trophic transfers. Comprehensive biological-trait 576 databases for aquatic organisms, such as Tachet *et al.* (2010), provide a valuable resource for 577 such work.

578

579 (3) Validation of biomarkers for quantifying EDC effects

Biomarkers, used to identify endocrine disruption within individuals, are well established for 580 a small number of taxa, e.g. fish (Ankley et al., 2009). Methods for other taxa have received 581 less attention, and their utilisation and validation is relatively poorly developed (see Matozzo 582 et al., 2008). A recent review identified a wide range of established and novel techniques for 583 identifying endocrine disruption across environmental samples, yet there is an absence of 584 suitable data for their validation (Kudłak et al., 2015). Furthermore, the relative accuracy of 585 biomarker assessments is widely debated, with inconclusive results for some novel biomarker 586 techniques. For example, the use of vitellogenin as a biomarker of endocrine disruption in an 587 amphipod (Gammarus fossarum Fabricius) proved inconclusive as vitellogenin expression 588 was shown to vary with unexplained environmental conditions (Jubeaux et al., 2012). The 589 unknown, potentially pleiotropic, function of the vitellogenin gene within male invertebrates 590 also may limit the application of this biomarker in the assessment of endocrine disruption 591 (Jubeaux et al., 2012). Further development and validation of biomarkers specific to EDCs 592 therefore remains an important challenge (Kudłak et al., 2015). Relating the severity of 593

endocrine disruption (via biomarker assessments) to analytical quantification of 594 environmental EDC concentrations (e.g. *via* gas chromatography mass spectrometry) is 595 596 essential for advancing our understanding of endocrine disruption in natural systems. Such comparisons will allow evaluation of the robustness of biomarkers in assessing ecological 597 risk from EDCs and stimulate the refinement of in vivo methods. The currently restricted 598 599 focus a few chemicals and organisms limits the ability of practitioners to utilise biomarkers 600 for ecological risk assessment and environmental decision-making (Hutchinson et al., 2006). 601 Establishing a wider database of biomarkers for multiple species and EDCs is therefore an 602 important future goal.

603

604 (4) Applying genetics and modelling to broad-scale analysis

A significant concern surrounding EDCs is the potential for impacts on the genetic structure 605 of populations and thus on the integrity of wild populations (Coe et al., 2008). Genetic 606 assessments within natural systems, including DNA microsatellite and single nucleotide 607 polymorphism (SNP) analysis, and other sequencing methods, provide the potential to assess 608 whether EDCs affect population structure via genomic pathways (e.g. Harris et al., 2011). 609 Olmstead, Lindberg-Livingston & Degitz (2010) reported with EDC-induced sex reversal 610 identifiable from genetic polymorphisms within the western clawed frog (Xenopus tropicalis 611 Gray). As well as allowing for broad-scale analyses, these techniques enable a reduction in 612 613 the previously large number of samples required for field-based assessments to detect reproductive impacts and sex reversal at low EDC concentrations. 614 Up-scaling research into the effects of EDCs also requires improved models for populations 615 and food webs. One major constraint in currently available population models is the absence 616 of suitable parameterisation and validation data at the population level collected using field 617 assessments (Rose et al., 1999; Raimondo et al., 2009). Future models must also aim at an 618

improved representation of the biotic and abiotic controls present within natural systems 619 (Borgå et al., 2004). Complexity, nonetheless, does not always facilitate accuracy, and highly 620 621 site-specific, overly complex models may lack wider applicability (Miller et al., 2007). New model strategies, such as developed by Rose et al. (2003), provide the way forward for future 622 models, with a nested structure allowing incorporation of a range of multi-scalar data, and in 623 624 turn generating model simulations which replicate well the natural conditions found within 625 ecological systems. Such work will enable an amalgamation of laboratory and field-based 626 data, facilitating an understanding of causality and environmental relevance within future 627 research.

628

629 VII. CONCLUSIONS

(1) The ecological effects of EDCs are currently investigated by effects assessments on 630 individuals employing only a small number of different organisms under controlled 631 experimental conditions. The environmental relevance of these findings is likely to be 632 limited. Spatially and temporally up-scaling these investigations within the aquatic 633 environment is therefore vital in developing environmentally relevant knowledge and to 634 provide supporting data for practitioners to make accurate risk assessments. The hormonal, 635 sub-lethal implications of EDC exposure could lead to a range of emergent effects resulting 636 from ecological interactions. 637

(2) We have highlighted the potential benefits of applying previously derived mechanistic
knowledge at broader spatial and temporal scales to assess the ecological impacts of EDC
exposure within natural systems. A range of abiotic and biotic characteristics and processes
can alter the effects and transfer of EDCs within aquatic food webs and cause deviations of
observed effects from those identified in experimental assessments. A range of indirect
effects also occur within natural systems, thus accurate assessment of endocrine disruption

risk within aquatic ecosystems requires an appreciation of ecological processes at a range ofspatial and temporal scales.

(3) Several limitations of experimental bioassay designs are highlighted by recent research
assessing broad-scale EDC exposure. Consequently, the results of experimental bioassays
should be interpreted with caution as such investigations often poorly represent influential
controls present in natural systems. It is suggested that chemical test guidelines and models
developed using these bioassays may provide limited utility in assessing the impacts and risk
associated with EDCs.

652 (4) A complementary suite of assessments at a range of scales should be adopted within a multi-tier integrated research strategy to promote the development of environmentally 653 relevant knowledge suitable for use by practitioners. Understanding the various direct and 654 indirect impacts of EDCs, across a range of different spatial and temporal scales, should 655 allow us to determine more effectively the transfer and ecological effects of EDCs within 656 natural systems. Increasing the effectiveness of empirical and experimental research through 657 methods such as integrated frameworks is therefore an important development. 658 (5) Future research should focus on expanding field-based research across a range of different 659 aquatic environments. To achieve this objective, however, methodological and theoretical 660 advances are required to enhance their applicability to natural systems and to develop more 661 comprehensive methods of risk assessment for EDCs. 662

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1239 endocrine disrupting chemicals (EDCs): A focus on non-monotonicity. *Environmental*

Health **14**, 42.

1241 Table 1. Bioaccumulation factors (BAFs) for endocrine-disrupting chemicals (EDCs) in aquatic organisms. Chemicals are divided into

1242 organophosphates, organophosphates, pharmaceuticals, steroidal androgens and oestrogens, organobromines, pharmaceuticals,

1243 phenols and pyrethroids. Where replicates or multiple measurements were reported within studies a mean value is presented.

Chemical group	Compound	log Kow	log BCF/BAF	Approximate trophic level	Organism	Source
Organobromines	BDE-100	7.24	7.50	3	Salvelinus namaycush	Streets <i>et al.</i> (2006)
	BDE-47	6.81	7.30	3	Salvelinus namaycush	
	BDE-66	_	7.30	3	Salvelinus namaycush	
	BDE-99	7.32	6.70	3	Salvelinus namaycush	
	γ-HBCD	5.48	4.51	3	Carassius auratus	Wu et al. (2011)
	HBB	6.09	3.48	2	Cipangopaludina chinensis	
		6.09	4.47	3	Carassius auratus	
	PBDEs	6.27	0.96	2	Gammarus pulex	Tlili et al. (2012)
		6.27	0.79	2	Echinogammarus stammers	Vigano <i>et al.</i> (2009)
Organochlorines	DDE	6.51	1.65	3	Rana spp.	Albanis <i>et al.</i> (1996)
		6.51	2.40	5	Egretta garzetta	
	DDT	6.52	4.00	2	<i>Pomacea</i> spp.	Siriwong et al. (2009)
		6.52	4.40	2	Macrobranchium lanchesteri	
		6.52	6.60	2	Filopaludina mertensi	
	HCB	5.72	6.20	2	Tubifex tubifex	Egeler et al. (1997)

		5.72	2.00	2	Eisenia fetida/andrei	
	Lindane	3.80	2.20	3	Rana spp.	Albanis <i>et al.</i> (1996)
		3.80	2.35	5	Egretta garzetta	
		3.80	4.40	2	Tubifex tubifex	Egeler <i>et al.</i> (1997)
		3.80	2.50	2	Eisenia fetida/andrei	
	PCBs	6.50	7.63	3	Perca fluviatalis	Bremle <i>et al.</i> (1995)
		6.50	6.60	1	Selenastrum spp.	Stange & Swackhamer (1994)
		6.50	6.10	1	Anabaena spp.	
Organophosphates	Chlorpyrifos	4.96	5.99	2	Mytilus galloprovincalis	Serrano <i>et al.</i> (1997)
	Methidathion	2.42	5.26	2	Mytilus galloprovincalis	
	TrBT	9.49	3.37	2	Ancylus fluviatalis	Ruhi et al. (2015)
		9.49	3.61	2	Hydropsyche spp.	
		9.49	3.53	3	Phagocata vitta	
Pharmaceuticals	Carbamazepine	2.25	3.03	3	Oreochromis niloticus	Garcia <i>et al.</i> (2012)
	Diclofenac	4.01	0.92	3	Oncorhynchus mykiss	Fick <i>et al.</i> (2010)
		1.90	6.86	3	Hemiculter leucisculus	J. Liu <i>et al.</i> (2015)
	Dilitiazem	2.70	3.18	3	Oncorhynchus mykiss	Fick <i>et al.</i> (2010)
	Diphenhydramine	3.11	2.77	3	Gambusia holbrooki	Wang & Gardinali (2013)
	Erythromycin	3.16	5.67	2	Planorbidae spp.	Du <i>et al.</i> (2015)
	Gemfibrozil	4.77	4.73	3	Gambusia holbrooki	Mimeault <i>et al.</i> (2005)
	Ibuprofen	3.79	4.06	3	Oncorhynchus mykiss	Fick et al. (2010)
	Oxazepam	2.24	0.30	2	Coenagrion hastulatum	Brodin <i>et al</i> . (2014)

	Propranolol	3.48	8.29	3	Hemiculter leucisculus	J. Liu et al. (2015)
	Roxithromycin	2.75	8.87	3	Hemiculter leucisculus	
Phenols	BPA	3.40	4.97	2	Pisidium amnicum	Heinonen <i>et al.</i> (2002)
		3.40	8.48	1	Benthic algae	Yang <i>et al.</i> (2014)
	Nonylphenol	4.48	8.85	1	Isochyrysis galbana	Correa-Reyes <i>et al.</i> (2007)
		4.48	2.64	2	Lumbriculus variegatus	Mäenpää & Kukkonen (2006)
	NPEO2	4.20	3.14	1	Cladophora glomerata	Ahel <i>et al.</i> (1993); Staples <i>et al.</i> (1998)
		4.20	-0.22	3	Oncorhynchus mykiss	
Pyrethroids	Cypermethrin	5.20	5.74	2	Chironomus tentans	Muir et al. (1985)
	Deltamethrin	5.20	5.76	2	Chironomus tentans	
	Fenvalerate	5.20	4.93	2	Chironomus tentans	
	Parathion	3.83	4.62	3	Gnathopogon caerulescens	Tsuda <i>et al</i> . (1994)
	Permethrin	6.20	5.56	2	Chironomus tentans	Muir et al. (1985)
	Vamidothion	0.12	6.56	3	Gnathopogon caerulescens	Tsuda <i>et al</i> . (1994)
Steroidal Androgens	4-AD	_	5.39	2	Meretrix lusoria	S. Liu et al. (2015)
and Oestrogens	ADD	_	6.33	2	Meretrix lusoria	
	Boldenone	_	8.01	2	Meretrix lusoria	
	EE2	4.01	0.80	2	Chironomus tentans	Dussault et al. (2009)
		4.01	4.23	1	Phytoplankton	Xie <i>et al.</i> (2015)

	4.01	4.89	3	Pelteobagrus fulvidraco	
Norgestrel	3.48	6.28	2	Meretrix lusoria	S. Liu et al. (2015)
	3.48	6.14	3	Lutjanus erythopterus	
Progesterone	3.87	7.70	2	Meretrix lusoria	
Testosterone	3.32	8.29	2	Meretrix lusoria	

4-AD, 4-androstene-3,17-dione; ADD, androsta-1,4-diene-3,17-dione; BDE, Brominated Diphenyl Ether; BPA, bisphenol A; DDT, Dichlorodiphenyltrichloroethane; DDE, Dichlorodiphenyldichloroethylene; EE2, 17α-ethinyloestradiol; HBB, Hexabromobenzene; HBCD, Hexabromocylcododecane; HCB, Hexachlorobenzene; NPEO2, nonylphenol ethoxylate 2; PBDE, polybrominated diphenyl ether; PCB, polychlorinated biphenyl; TrBT, tris-(2-butoxyethyl)-phosphate; BCF, Bioconcentration factor; BAF, Bioaccumulation factor; Log Kow, octanol/water partition coefficient.

Log Kow values were taken from <u>https://pubchem.ncbi.nlm.nih.gov/</u>

	Micro	Meso	Macro
Duration Length of exposure to EDCs	Days-Weeks	Weeks–Months	Months-Years
Replicability Degree to which experiments/results can be repeated	20–40 replicates	5–20 replicates	1–3 replicates
Direct causality Level of causation that can be directly derived from results	Causation		Correlation
Taxa diversity The number of taxa that can be assessed simultaneously	1–3	3–5	5–10
Exogenous factors Natural variability encompassed by the methodology	None	Intermediate	All

- **Fig. 1.** Conceptual differences in endocrine-disrupting chemical (EDC) experimental
- 1246 framework design and expected outcomes of micro-, meso- and macroscale assessments.



1247 Fig. 2. Interrelationships and information flow between micro-, meso- and macroscale investigations for the biological impact assessment of

1248 endocrine-disrupting chemical (EDC) exposure across a range of levels of biological organisation. Solid arrows indicate transfer of knowledge.