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Effect-based and chemical analytical methods to monitor estrogens under the European Water Framework Directive

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Detailed description of the authors contributions can be found in the Supplementary Information (SI, Table S1).

46 **Abstract**

47 The European Decision EU 2015/495 included three steroidal estrogens, estrone, 17 β -estradiol and 17 α -
48 ethinyl estradiol, in the “watch-list” of the Water Framework Directive (WFD). As consequence, these
49 substances have to be chemically monitored at the level of their environmental quality standards, which
50 can be challenging. This project aimed to identify reliable effect-based methods (EBMs) for screening of
51 endocrine disrupting compounds, to harmonise monitoring and data interpretation methods, and to
52 contribute to the current WFD review process. Water and wastewater samples were collected across
53 Europe and analysed using chemical analyses and EBMs. The results showed that 17 β -estradiol
54 equivalents were comparable among methods, while results can vary between methods based on the
55 relative potencies for individual substances. Further, derived 17 β -estradiol equivalents were highly
56 correlated with LC-MS/MS analyses. This study shows that the inclusion of effect-based screening
57 methods into monitoring programmes for estrogens in surface waterbodies would be a valuable
58 complement to chemical analysis.

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64 *Keywords*

65 Science-policy interface Estrogen screening Endocrine disruption

66 Surface and waste water assessment Emerging pollutants EU watch-list

67 Steroid analyses *In vitro* bioassays Integrated effects of mixtures

68

69 1 State of the Art

70 Over the past two decades, numerous scientific studies have demonstrated that endocrine disrupting
71 chemicals (EDCs) elicit adverse effects on sensitive aquatic species, such as fish [1-7]. Steroidal
72 estrogens, like the natural hormones estrone (E1) and 17 β -estradiol (E2), as well as the synthetic
73 hormone 17 α -ethinyl estradiol (EE2), are of particular environmental concern [8-11]. Due to their steady
74 release via waste water effluents into surface waters [12, 13] and their high biological activity, even very
75 low concentrations of E2 and EE2 have been shown to cause reproductive toxicity with negative effects
76 at the population level [14-16]. As a consequence, E1, E2, and EE2 were included in a European Union
77 (EU) Water Framework Directive (WFD) “watch-list” [17-20]. The WFD watch-list mechanism aims to
78 collect high-quality monitoring data on concentrations of emerging pollutants and potentially hazardous
79 substances, whose currently available monitoring information shows either quantitative or qualitative
80 deficiencies [21]. To collect more high-quality data, listed substances have to be monitored at
81 representative EU sampling sites for a period of at least 12 and up to 48 months. The watch-list
82 mechanism is expected to support future substance prioritisation processes, enable the implementation
83 of measures, and facilitate environmental risk assessment across the EU.

84 Chemical monitoring of estrogens for the watch-list mechanism is challenging, because the **European**
85 Commission set maximum acceptable method detection limits (MDLs) at EQS levels of 400 pg/L for E1
86 and E2, and 35 pg/L for EE2 [18, 22]. Most routine analytical methods used by the Member States
87 cannot meet these requirements, especially for EE2, **based on** [23, 24]. Hence, the quality assessment of
88 water bodies based on current methods is a challenge for the detection/quantification limits that are too
89 high to detect if EQS are being exceeded or not. Effect-based methods are able to detect estrogenic
90 substances at **sub-ng or even pg levels** and have the potential to be used as a complementary screening
91 tool [12, 25-27]. In addition, they do not require *a priori* knowledge of the substances to be monitored,
92 as they are able to determine the biological response caused by complex mixtures of unknown
93 compounds. Thus, effect-based methods may be suitable to serve as a valuable link between chemical
94 analytical and ecological quality assessments, since the effects can **rarely** be linked to individual
95 compounds.

96 As described in **an** EU technical report, which was elaborated in the context of the Chemical Monitoring
97 and Emerging Pollutants (CMEP) expert group under the Common Implementation Strategy (CIS) of
98 the WFD, effect-based tools can be categorised into three main groups: Bioassays (*in vitro*, *in vivo*),
99 biomarkers, and ecological methods [28]. With regard to steroidal estrogens and other EDCs, *in vitro*
100 reporter gene assays have been used predominantly to determine the total estrogen receptor (ER)
101 mediated estrogenicity of an environmental sample [29]. Among the most commonly applied assays are
102 *in vitro* methods such as estrogen receptor transactivation assays (ER-TAs), which use various cell types
103 including yeast, human and other mammalian cell lines that were transfected with a human estrogen
104 receptor coupled to a reporter gene [30]. Activation of the ER leads to the expression of the reporter

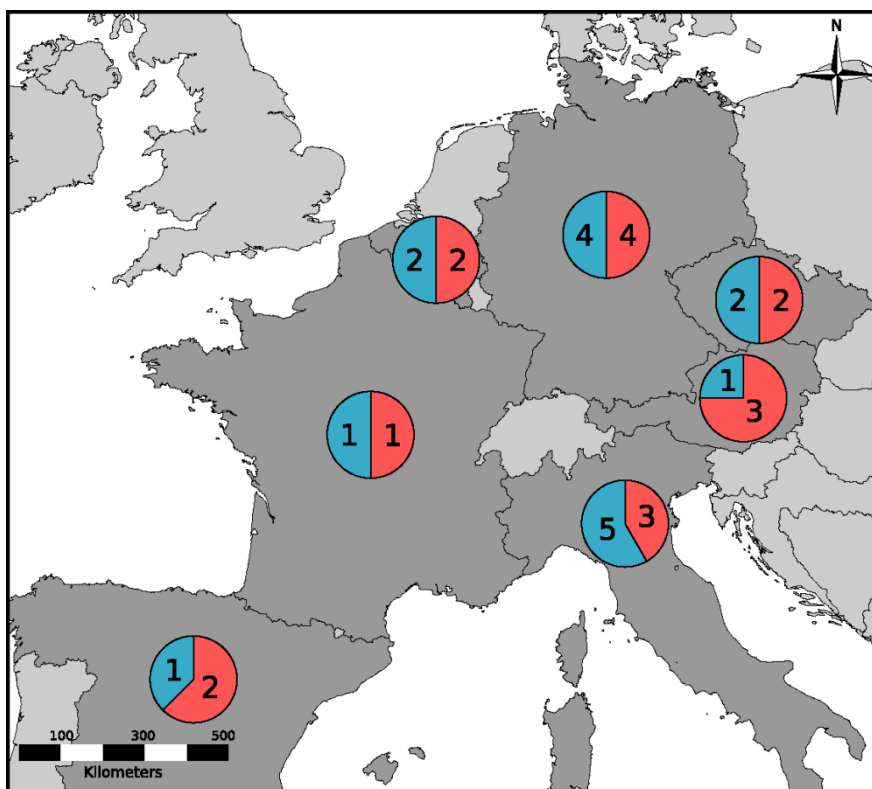
105 gene product, usually an enzyme that modifies another chemical, causing a quantifiable response. The
106 resulting estrogenic potential of a sample is expressed as an E2 equivalent concentration (EEQ),
107 indicating the estrogenic activity of the sample or sample dilution in terms of equivalency to the
108 estrogenic activity of the corresponding E2 reference concentration [31].

109 Although ER-TAs are highly advantageous methods for the detection of ER activation and
110 quantification of very low estrogen concentrations in surface waters [23], these methods are not included
111 within current WFD monitoring programmes [20]. One reason for this is the lack of data that
112 demonstrate their applicability as a monitoring and screening tool in combination with chemical
113 analytical methods (see e.g. [14]). Such information would greatly increase their regulatory acceptance.
114 As a response to this need, an EU-wide project involving 24 research organisations and environmental
115 agencies from 12 countries was carried out to evaluate the usefulness of specific *in vitro* methods for
116 identifying the presence of the watch-list substances, E1, E2, and EE2, in surface and waste waters. The
117 project aimed to compare the chemical and effect-based data resulting from the analysis of 16 surface
118 and 17 waste water treatment plant effluent samples. Analyses were conducted in seven participating
119 laboratories using different LC/MS- (three laboratories) and effect-based methods (five laboratories).
120 The objectives of the study were (i) the demonstration of reliable effect-based screening methods for the
121 monitoring of estrogenic EDCs in waste water and surface water, (ii) the harmonisation of data
122 interpretation methods, and (iii) providing recommendations for the implementation of cost-effective
123 and reliable effect-based methods in WFD monitoring programmes.

124 **2 The Project**

125 **2.1 Sampling**

126 A total number of 16 surface water (SW) and 17 waste water (WW) samples were collected according to
127 a protocol developed by the participants (SI, Part A). Selected sampling sites were located in seven
128 European countries in Central and Southern Europe (Figure 1): Austria (1 SW/ 3 WW), Belgium (2/2),
129 Czech Republic (2/2), France (1/1), Germany (4/4), Italy (5/3), and Spain (1/2). Sample collection was
130 carried out from September to November 2015 by ten participating institutions. The samples were taken
131 based on prior knowledge on their contamination with estrogens and represented a gradient of
132 contamination from high to moderate.



133
 134 **Figure 1:** Samples taken in various European States (dark grey). The circles indicate the number of surface water (blue) and
 135 waste water samples (red) taken in each country.

136 **2.2 Sample preparation**

137 The sample preparation included the filtering of a part of the SW (see SI, Part A) and all WW samples
 138 over glass fibre filters (Millipore, type 4, retention 2.7 µm, circle size 4.7 cm). Since a filtration step can
 139 have an impact on the composition of a sample and its estrogenic activity [32], the filtration step was
 140 investigated during a feasibility study prior to the main study presented here. The results of the pre study
 141 did neither show a significant reduction in estrogenicity in the control nor in tested environmental
 142 samples (data not shown). Subsequently, all samples were enriched by means of solid-phase extraction
 143 (SPE; 11 L sample to 11 mL extract) and extracts were passed over silica gel (SiOH) columns (methods
 144 focusing on E1, E2 and EE2). While for surface water each extract was split into eleven 1 mL aliquots
 145 that were each passed over a single SiOH column, for waste water a single column was inadvertently
 146 used to treat the whole extract (11 mL). For LC-MS/MS analysis this means that matrix was less
 147 efficiently removed from WW extracts (relative to SW extracts) and higher matrix loads would have
 148 impeded low LOQs in WW LC-MS/MS analysis. For bioassay analysis this means that, should
 149 additional ER-agonists (i.e. other than E1, E2 and EE2) have been present in the extracts, a reduced
 150 clean-up efficiency would have reduced ER-agonist removal which in turn would have caused enhanced
 151 effects in bioassays. Full details of sample preparation are provided in SI, Part A.

152 **2.3 Chemical and effect-based analyses**

153 Participating laboratories received spiked reference samples, blanks and encoded water extracts. The
154 chemical analyses were conducted in three different labs, which applied an LC-MS/MS with negative
155 electron spray ionisation (detailed information in SI, Part D Table S2). The effect-based methods were
156 conducted in five different labs: **Estrogen Receptor Chemical Activated LUciferase gene eXpression**
157 **(ER-CALUX)** at Biodetection Systems (BDS), **luciferase-transfected human breast cancer cell line**
158 **(MELN) gene-reporter assay** at INERIS [33], ER-GeneBLAzer assay at the Helmholtz Centre for
159 Environmental Research (UFZ) [34], the **stably transfected human estrogen receptor-alpha**
160 **transcriptional activation Assay using hER α -HeLa-9903 cells** (HeLa-9903 assay) at RECETOX [35],
161 and **planar Yeast Estrogen Screen** (pYES) at the German Federal Institute of Hydrology (BfG) [36, 37].
162 The pYES is a method, which combines a chromatographic separation of the sample by thin layer
163 chromatography (TLC) with a subsequent performance of the YES on the planar surface of the TLC-
164 plate [38-40]. Like the common assays which are performed in micro-well-plates, this approach allows
165 the quantification of the overall estrogenic activity present in the sample by means of E2-equivalence
166 concentrations. Furthermore, like methods based on LC/MS, it also allows the estimation of
167 concentrations of individual estrogenic compounds, e.g. E1, E2 and EE2, due to the chromatographic
168 separation of the sample. For this purpose the respective standard compounds are used for a calibration
169 on the same TLC plate – in the present study E1, E2, EE2, and **estriol** (E3) were applied in a mixture at
170 three different levels. Due to the limited separation power of the thin layer chromatography compared to
171 HPLC and GC in particular, a co-migration of estrogenic compounds cannot be excluded. Therefore,
172 under the assumption of effect addition, the estimated individual concentrations represent the possible
173 maximal concentration of the respective compound. This approach can be used to identify and quantify
174 substance groups causing ER-activation.

175 **2.4 Blanks and positive controls**

176 Ultrapure water (11 L) was used as extraction blank. An extraction blank was included with each
177 extraction run of 10 samples, subjected to clean-up and distributed the same as the sample extracts.
178 Further, each analysis using effect-based methods included a negative control. To avoid solvent effects
179 on cell viability, its concentrations did not exceed a **defined** value (see SI, Part D Table S3). As positive
180 controls for ensuring the validity and enabling a comparison of the methods, surface water samples
181 (11 L each) from the Netherlands were spiked with E2 and EE2 at two concentrations by the central lab
182 (BDS). The “low spike” (600 pg/L) represented a concentration slightly above the proposed EQS for E2
183 (400 pg/L). The “high spike” (6000 pg/L) represented a concentration that is quantifiable with high
184 certainty by both effect-based and chemical methods.

185 **2.5 Data evaluation – effect-based methods**

186 Raw data and information on relative enrichment factors (REF) of the extracts were collected from
187 participating laboratories. The REF expresses the combination of: 1) sample enrichment using SPE and
188 2) extract dilution steps in each of the applied effect-based methods. Estrogenic activity of the extracts
189 was expressed as E2-equivalence concentration (pg EEQ/L water) (described in detail in SI, Part B).
190 Briefly, dose-response curves of the reference compound, E2, and the dilution series of the water
191 extracts and blanks were fitted using a five-parametric non-linear regression with normalised data. The
192 concentration of the positive control (E2) needed to induce 10 % effect of the maximum E2-induction
193 (PC₁₀), was calculated. Subsequently, the relative REF of the sample, that stimulates the assay at PC₁₀
194 level was determined by interpolation. The PC₁₀ reference concentration was divided by the
195 corresponding sample dilution (REF) to obtain the EEQ of the sample. EEQs derived by the PC₁₀
196 method are presented in the results section.

197 **2.6 Data evaluation – chemical analysis**

198 Internal standard calibration and interpolation using a linear regression model were performed to
199 determine concentrations (pg/L) of the individual steroidal estrogens in sample extracts. Identification of
200 selected analytes was performed based on two to three Multiple Reaction Monitoring (MRM) transitions
201 between the precursor ion and two or three most abundant product ions, depending on the laboratory
202 where analyses were done. The first transition was used for quantification purposes whereas the second
203 and third transitions were used to confirm the presence of the target compound in the sample. Quantified
204 analytes were identified by comparing the retention time (RT) of the corresponding standard and the
205 ratio between two ion transitions recorded ($\pm 20\%$) in the standard and water samples.

206 **2.7 Calculation of sample-dependent LOD and LOQ**

207 The Limits of quantification (LOQ) for effect-based methods the LOQs were calculated as 3-fold the
208 standard deviation (SD) of the averaged response of the negative control on each assay plate. The effect
209 level of 3-fold the SD was interpolated from the E2 reference curve and divided by the REF of the
210 sample to derive the LOQ. The actual reporting for effect-based methods occurred at the 10% effect
211 level which was always above LOQ (typically at 2-5 % effect levels).

212 In case of the chemical analysis the limits of detection (LOD) were determined for each compound in
213 each sample based on the signal intensity of the internal standards or the analyte peak by a signal-to-
214 noise (S/N) ratio of 3:1 and LOQ by a S/N ratio of 10:1.

215 When comparing LOQs of effect-based methods with those of chemical analyses the various key
216 differences between the two approaches need to be taken into account (for further background see SI,
217 Part C).

2.8 Comparison of chemical and biological analysis

The EEQ_{bio} is the ratio of the effect concentration of the reference compound estradiol $EC_{50}(E2)$ (pg/L) and the sample $EC_{50}(\text{sample})$ (Equation 1) and was derived in this study using the PC_{10} approach (see above). The EEQ_{chem} was calculated from the sum of the relative effect potencies REP_i times the detected concentration of estrogenic chemical i , c_i [41]. The REP , in turn, is the ratio of the effect concentration of the reference compound estradiol $EC_{50}(E2)$ and the chemical i 's $EC_{50}(i)$ (Equation 2).

$$EEQ_{bio} = \frac{EC_{50}(E2)}{EC_{50}(\text{sample})} \quad (1)$$

$$EEQ_{chem} = \sum_{i=1}^n REP_i \cdot c_i = \sum_{i=1}^n \frac{EC_{50}(E2)}{EC_{50}(i)} \cdot c_i \quad (2)$$

Due to the analytical method detection limits of E2 and EE2, we evaluated the potential contribution of non-detected estrogens to the overall $EEQ_{chem, LOD/2}$ using Equation 3, where values below the LOD (“non-detects”) were included as $LOD/2$. If the analytical lab reported data as $<LOQ$, we used $LOQ/2$ in Equation 3 instead of $LOD/2$. In Equation 3, n refers to the total number of chemicals included in the analysis, m refers to the number of chemicals below LOD. C_i is the average value of three analytical measurements,

$$EEQ_{chem, LOD/2} = \sum_{i=1}^{n-m} REP_i \cdot c_i + \sum_{j=1}^m REP_j \cdot LOD_j/2 \quad (3)$$

2.9 Correlation analysis

The correlation analysis among effect-based methods (EEQ_{bio}) was performed with GraphPad Prism, using the Pearson correlation (r). [42].

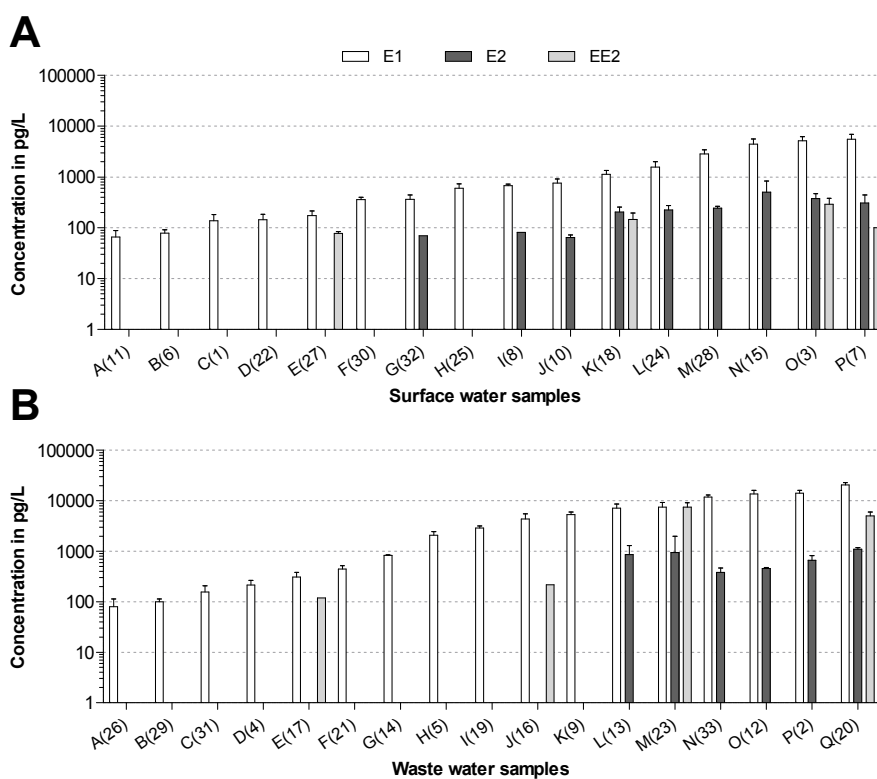
3 Results and discussion

3.1 Reference chemicals and validation

All essential criteria for method performance were fulfilled in this study (described in more detail in the SI, Part E). As shown in Table S4 (SI, Part E), the chemical analytical as well as effect-based methods showed good recovery in the spiked samples. No estrogenic activity or quantifiable concentrations of E1, E2, and EE2 were measured in the blank samples (i.e. procedure-, extraction- and solvent blanks). As the derived effect concentrations in the effect-based methods and chemically measured EE2 concentrations matched with the nominal concentrations of the spiked samples, the observed effects can be ascribed to the samples themselves.

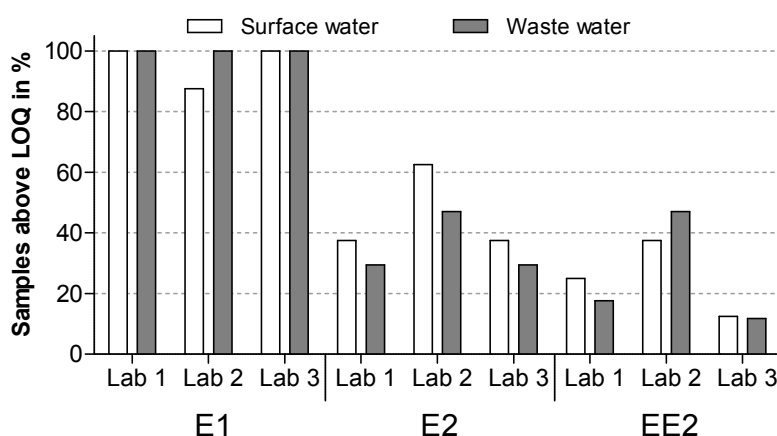
245 **3.2 Results of chemical analysis**

246 Measured concentrations of the three estrogens E1, E2 and EE2 differed widely between sampling sites
 247 as well as between surface and waste water samples. Differences among SW samples can be explained
 248 by **varying river characteristics, e.g. flow (dilution factor), or temperature, as** well as differences in
 249 estrogenicity of treated WW, that are released into the SW. The results of the analyses, which are
 250 summarised in Figure 2, show a 3.2 to 3.6 times higher mean concentration for E1 and E2 in WW
 251 (Figure 2B) compared to SW (Figure 2A). Due to **the highly contaminated** WW sample M(23), possibly
 252 influenced by an industrial discharge of EE2, the mean concentration of EE2 across all WW samples
 253 was approximately 20 times higher compared to SW (Figure 2). Estrone (E1) was quantified in all
 254 samples. For E1 maximum concentrations of 5.6 ng/L (sample P(7)) and 20.5 ng/L (sample Q(20)) in
 255 SW and WW were measured, respectively. E2 was the second most frequently quantified estrogen and
 256 measured above LOQ in nine of 16 SW and six of 17 WW samples. Measured concentrations ranged
 257 from 0.4 ng/L (sample N(33)) to 1.1 ng/L (sample Q(20)) in WW, and from 0.06 ng/L (sample J(10)) to
 258 0.5 ng/L (sample N(15)) in SW. The synthetic EE2 was least frequently quantified and measured above
 259 LOQ in four of 16 SW and four of 17 WW samples with a maximum concentration of 0.3 ng/L in SW
 260 sample O(3) and 7.5 ng/L in WW sample M(23). These concentration ranges and patterns are in
 261 accordance with recent review studies [43, 44].



262
 263 **Figure 2: Chemical analytically measured concentrations for SW (A) and WW extracts (B) above LOQ for E1, E2 and**
 264 **EE2.** The bars show the mean concentration of all three applied methods for each analyte showing results > LOQ, the standard
 265 deviation is shown when two or three methods reported results. The sample-dependent LOQs are listed in the supplementary
 266 information together with the measurement data of analytical methods (SI, Part F, Table S6 and S7).

267 Our results underline the analytical difficulties that have recently been highlighted for E2 and EE2 by
 268 several studies and workshops [16, 45], stressing the challenges that emerge for routine methods used in
 269 national monitoring programmes. Despite the use of quite advanced chemical analytical techniques
 270 (status 2015), the detection and quantification of E2 and EE2 in SW and WW samples was problematic
 271 in some cases. While it was possible to quantify E1 in almost all samples, the percentage of
 272 quantifications was significantly reduced for E2 and even more for EE2 (Figure 3). This was partially
 273 due to the fact that insufficient silica gel was used to reduce the matrix effects in WW. WW is
 274 considered as worst-case regarding matrix effects [46, 47].



275
 276 **Figure 3: Mean percentage of quantified (>LOQ) samples for each substance in SW and WW.** The sample-dependent
 277 LOQs are listed in the supplementary information together with the measurement data of the analytical methods (SI Part F,
 278 Table S7).

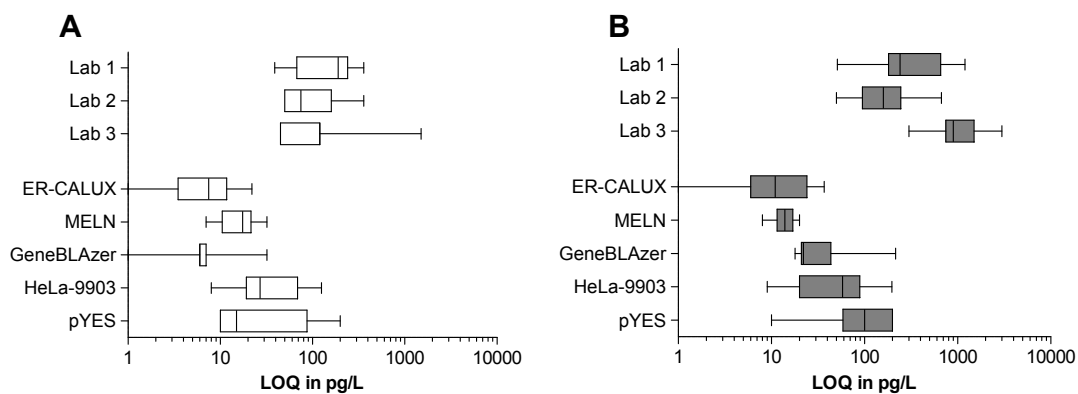
279 However, the quantification of substances itself is not the only challenge faced by those routinely
 280 applying analytical methods for watch-list monitoring. According to the EU Commission Decision
 281 2015/495, which established the first watch-list, the indicative methods applied by Member States have
 282 to meet the minimum requirement for method detection limits (MDL) equal to the proposed EQSs of E1
 283 at 3.6 ng/L, E2 at 0.4 ng/L and EE2 at 0.035 ng/L [18]. To take into consideration the matrix effects of
 284 different waters, LODs and LOQs had to be calculated for each sample (SI Part F, Table S7). The three
 285 techniques used in the current study were able to meet MDL requirements for E1 in all SW and WW
 286 samples. Also for E2, in 96 % of surface water samples and 94 % of waste water samples detection was
 287 possible at the level of the proposed EQS. In the case of EE2, the minimum criteria were not met, since
 288 only 56 % and 16 % of SW and WW samples, respectively, could be monitored at the EQS level. These
 289 findings are in accordance with a recent report from 2015, which showed that the lowest LOQ found in
 290 literature at that time was sufficient for compliance monitoring of E1 and E2 in inland surface waters,
 291 while the criteria were not met for EE2 by several Member States [24]. It has to be pointed out that, in
 292 this project, the silica clean-up step for the sample extracts differed between WW and SW samples (see
 293 methods section) favouring the presence of polar compounds in extracts of WW samples. This
 294 difference likely reduced the sensitivity of the analytical method for the target compounds in WW
 295 samples. Furthermore, sample extraction was performed at pH 3 possibly increasing concentrations of

296 humic acids and thus lowering sensitivity of LC/MS-based methods applied. Under ideal conditions, we
297 estimate that analytical methods can achieve LODs and LOQs of a factor 2 to 3 lower in WW samples.
298 It has to be recognised that the LODs of chemical analytical methods used exclusively for steroidal
299 estrogens already significantly decreased from 2013 (LOD E2 and EE2 of 100 pg/L) to 2015 (E2: 60
300 pg/L, EE2: 85 pg/L) and will certainly decrease further [16, 23].

301 Nevertheless, if steroidal estrogens were to be included in the EU priority list for monitoring, very strict
302 minimum performance criteria would apply. As stated in the Commission Directive 2009/90/EC, an
303 analytical method used for monitoring of priority substances needs a LOQ equal or below a value of
304 30 % of the EQS [48]. These requirements can presently be met only for E1, but not for E2 or EE2 in all
305 SW. Regarding the quantification of E2, and EE2, existent routine analytical techniques still lag behind
306 the requirements. This result is supported by two recent reviews on the performance of current analytical
307 methods that have shown that 35 % of reviewed methods complied with the EQS for E2, while only one
308 method complied with the EQS for EE2 [49, 50]. In order to not only detect but also quantify at such
309 low concentrations as required for regulatory monitoring application, a further decrease of LOQs is
310 necessary, which is difficult to achieve for routinely used non-tailored analytical methods in the short-
311 term.

312 **3.3 Quantification limits of chemical-analytical and *in vitro* effect-based methods**

313 The LOQs for all methods applied in this study are summarised in Figure 4. Since E2 is used as the
314 reference compound for all effect-based methods, the LOQ of E2 is shown for the chemical-analytical
315 methods as an example. When comparing LOQs across the different methods it has to be taken into
316 account that LOQs were derived along different approaches (see method section and SI, Part C for
317 further details). The effect-based *in vitro* methods were generally able to quantify effects at one to two
318 orders of magnitude lower concentrations than the analytical methods used. For effect-based methods,
319 LOQs ranged between 0.002 ng/L and 0.2 ng/L for SW as well as WW, while for chemical-analytical
320 methods LOQs for E2 were 0.04 ng/L to 1.5 ng/L in SW and 0.05 ng/L to 3 ng/L in WW. This increase
321 in LOQs for chemical-analytical methods in WW samples (Figure 4B) compared to surface water
322 (Figure 4A) can be ascribed to the higher complexity of the waste water matrix [46, 47] as well as the
323 less efficient clean-up used for WW samples.



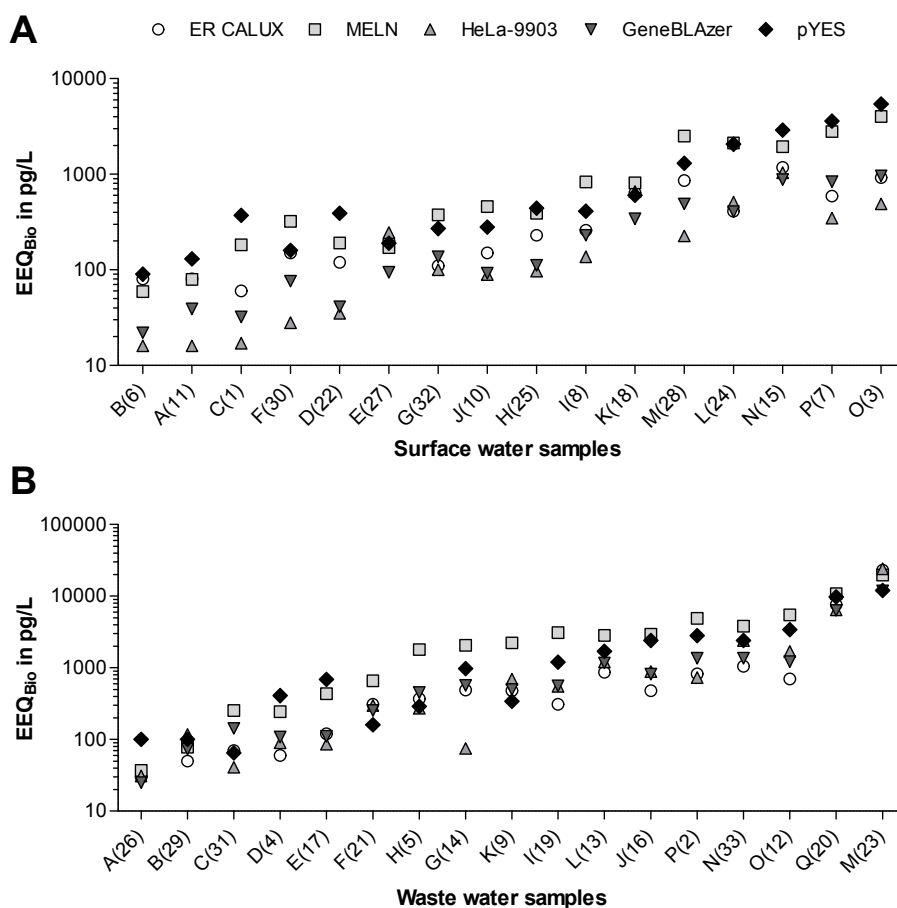
324

325 **Figure 4: Sample-dependent LOQs in surface water (A) and waste water (B) extracts.** For the chemical analytical method
 326 the LOQ of E2 is shown as an example and for the effect-based methods the LOQ of the integrated effects is represented. Plots
 327 indicate the distribution of data, thereby the bottom and the top of the box are the first and third quartiles, while the line inside
 328 the box is the median. The whiskers show the minimum and maximum of all data.

329 3.4 Measured estrogenic effects

330 As a result of these low effect-based quantification limits, estrogenic activities were detected in all tested
 331 samples. As expected, highest EEQs were measured in WW samples (Figure 5A and B). In SW, EEQ_{bio}
 332 ranged from 0.16 ng/L measured with HeLa-9903 in sample B(6) to up to 5.4 ng/L measured with pYES
 333 in sample O(3). In WW, the lowest EEQ_{bio} of 0.03 ng/L was measured in sample A(26) with ER-
 334 GeneBLAzer, while the highest EEQ_{bio} of 24 ng/L was measured in sample M(23) with HeLa-9903.
 335 Further, it is evident that EEQ_{bio} for SW samples determined with the MELN, as well as the pYES, were
 336 higher (> 50 %) than the EEQ_{bio} measured with the other effect-based methods. A possible reason for
 337 this pattern, which was less pronounced in WW, could be a higher sensitivity of the MELN and pYES
 338 towards E1 (see SI Part F, Table S8), combined with a larger proportion of E1 in surface water.
 339 Additionally, alterations in the method's performance occur due to differences between the test systems,
 340 which was already mentioned in previous studies [23, 44, 51] and is further discussed for this project in
 341 an associated publication [52].

342



343

344 **Figure 5: Measured E2-equivalents for all SW (A) and WW (B) extracts.** The symbols show the EEQs for each bioassay,
 345 which were calculated according to the method described in section 2.5. The sample-dependent LOQs are mentioned in the
 346 supplementary information, together with the measurement data of effect-based methods (SI Part F, Table S8 and S9).

347 3.5 Comparison of chemical analysis and *in vitro* effect-based methods

348 We cannot a priori expect consistency between EEQ_{chem} calculated from E1, E2, and EE2 concentrations
 349 and EEQ_{bio} . Although the extraction and clean-up method focused on E1, E2, and EE2, other natural
 350 estrogens and xenoestrogens (both agonists and antagonists) might still be present in the extracts and
 351 contribute to the mixture effects detected by effect-based methods. Thus, there can be situations where
 352 EEQ_{chem} is lower than EEQ_{bio} because: 1) agonists other than E1, E2, and EE2 were present in the
 353 sample but not quantified by LC-MS/MS analyses or 2) some target compounds were present but below
 354 LOQ or LOD, thus they were not included in EEQ_{chem} but still contributed to EEQ_{bio} . Alternatively,
 355 EEQ_{chem} can be higher than EEQ_{bio} when antagonists suppress the response of the assay.

356 For ER-CALUX, the comparison of EEQ_{bio} with EEQ_{chem} (Figure 6A) indicated an underestimation of
 357 EEQ_{bio} by EEQ_{chem} at low concentrations of steroidal estrogens. When E1 concentrations are low,
 358 typically E2 and EE2 concentrations are below LOQ (Figure 2). However, as stated above, also below
 359 their LOD/LOQ, these chemicals may be present and contribute to the biological mixture effect (i.e.
 360 EEQ_{bio}). We therefore also calculated the $EEQ_{chem,LOD/2}$ that uses the LOD/2 or LOQ/2 for those E2 and
 361 EE2 concentrations below the LOD or LOQ. The increase in EEQ_{chem} , due to the inclusion of LOQ/2

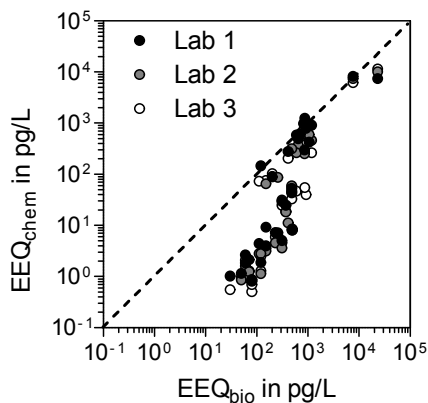
362 and LOD/2 data (SI, Part F, Table S10-14), shifts the $EEQ_{chem} - EEQ_{bio}$ data cluster towards the one-to-
363 one line (Figure 6B). In fact, there is now a slight overestimation of the biological effect in the range
364 where EEQ concentrations are low (up to ca.100 pg/L). The fact that the agreement between EEQ_{chem}
365 and EEQ_{bio} has become much better (going from Figure 6A to 6B) is a good indication that E2 and EE2
366 are indeed present and were captured by effect-based methods.

367 The situation for MELN is markedly different from that of ER-CALUX. For MELN the direct
368 comparison between EEQ_{chem} and EEQ_{bio} is already very good (Figure 6C). In fact, EEQ_{chem} tends to be
369 above EEQ_{bio} already before adding the additional EEQ_{chem} component using LOD/2 or LOQ/2 for E2
370 and EE2. The inclusion of LOD/2 or LOQ/2 in the EEQ_{chem} calculation caused a notable overestimation
371 of EEQ_{chem} for almost all samples (>90 % of data above the 1 to 1 line in Figure 6C). The other three
372 bioassays show results that are intermediate between ER-CALUX and MELN, with a general trend
373 towards a slight underestimation of EEQ_{chem} for samples with low EEQ_{bio} and an overestimation after
374 adding LOD/2 or LOQ/2 (see Figure S1).

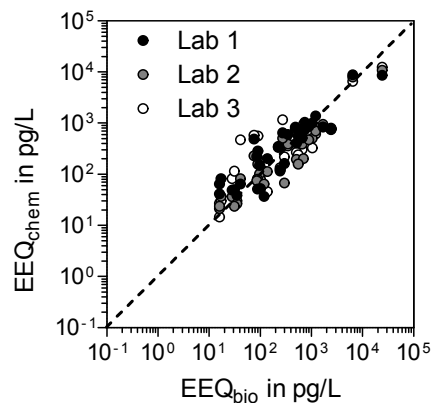
375 The marked differences between ER-CALUX and MELN are not unexpected. MELN has the highest
376 relative E1 effect potency of all tested bioassays (0.29 compared to 0.01 for ER-CALUX; Table S5).
377 Thus, EEQ_{chem} results for MELN are strongly based on E1 concentrations – a compound that was always
378 measured (except for a few samples by Lab 2, Figure 3). Consequently, for MELN the relative
379 contribution of E2 and EE2 at LOD/2 or LOQ/2 on top of measured E1 concentrations is relatively small
380 though still noticeable for samples with low EEQ concentrations (compare Figure 6C to 6D).

ER-CALUX

A EEQ_{chem}

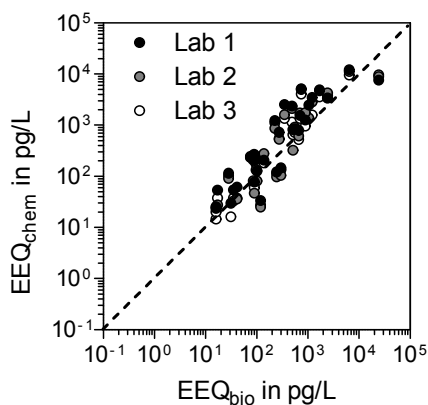


B $EEQ_{chem + LOQ/2 \text{ or } LOD/2}$

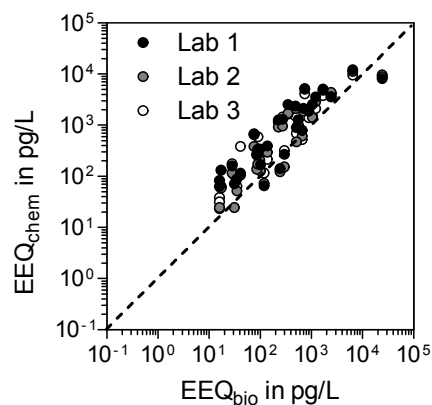


MELN

C EEQ_{chem}



D $EEQ_{chem + LOQ/2 \text{ or } LOD/2}$



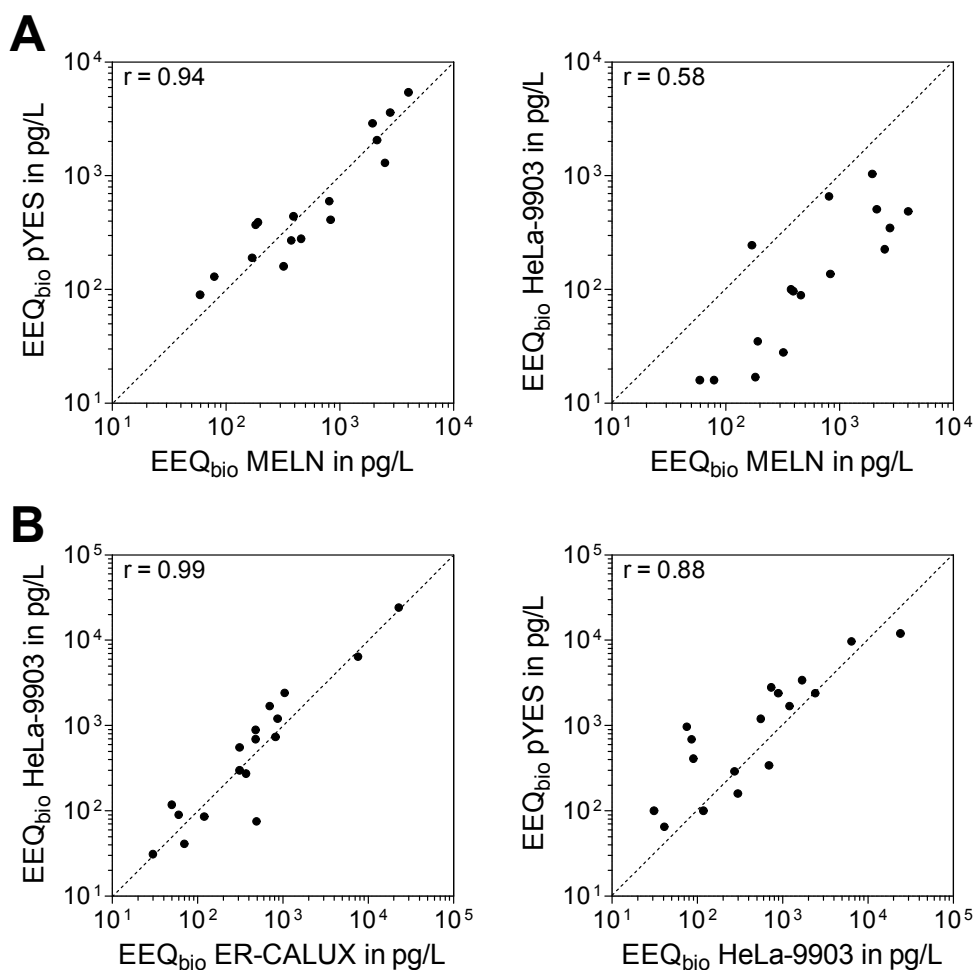
381

382 **Figure 6: Comparison of EEQ_{chem} with EEQ_{bio} .** Exemplary graphs are shown for the ER-CALUX (A, B) and MELN assay
383 (C, D) (further figures in the SI, Part G). Graphs on the left show the EEQ_{chem} derived from values $>LOQ$, while the graphs on
384 the right show the $EEQ_{chem + LOD/2 \text{ or } LOQ/2}$ calculated by including $LOD/2$ or $LOQ/2$. The dashed line indicates perfect agreement
385 of EEQ_{chem} with EEQ_{bio} .

386 3.6 Comparison of effect-based methods

387 To compare the five effect-based methods amongst each other, a correlation analysis was conducted by
388 plotting the EEQs of one method against the EEQs of all other methods for SW samples and WW
389 samples, respectively (Figure 7).

390



391

392 **Figure 7: Exemplary graphs of correlation analysis of effect-based methods for SW (A) and WW (B) showing the**
 393 **strongest and weakest correlations.** The correlation analysis was based on the method described in section 2.9. The dashed
 394 line indicates perfect agreement of the compared effect-based methods. All correlations were significant with a p value <0.0001
 395 except for MELN and HeLa-9903 (top right panel) which had a p value \approx 0.01. Further graphs are shown in SI, Part H, Figures
 396 S2 and S3.

397 The results of this analysis are summarised in **Table 1 and Table 2** and show a strong correlation and
 398 thus good comparability of pYES, MELN and ER-CALUX. For SW samples, the strongest correlations
 399 were seen for pYES/MELN ($r^{\circ}=0.94$) and pYES/ER-GeneBLAzer ($r^{\circ}=0.94$), while the weakest
 400 correlation was determined for MELN/HeLa-9903 ($r^{\circ}=0.58$). For WW samples, test results correlated
 401 strongly among all methods (Table), and the strongest correlation ($r^{\circ}=0.99$) was observed for ER-
 402 CALUX/HeLa-9903. It is known that effect-based methods differ in their REPs for individual ER-
 403 agonists [53-55] which can explain that results obtained by the HeLa-9903 assay correlated less strongly
 404 with other test results . Based on these differences effect-based methods can be split into two groups:
 405 pYES and MELN with high E1 REP and ER-CALUX, HeLa-9903 and ER-GeneBLAzer with lower E1
 406 REP.

407 **Table 1: Pearson correlation coefficients of all bioassays for SW.** The values were calculated according to the method
 408 mentioned in section 2.9. All correlations were significant with a p value <0.0001 (***) and a p value ≈ 0.01 (*).

	MELN	ER-GeneBLAzer	HeLa-9903	pYES
ER-CALUX	0.81 ***	0.91 ***	0.86 ***	0.76 ***
MELN		0.93 ***	0.58 *	0.94 ***
ER-GeneBLAzer			0.77 ***	0.94 ***
HeLa-9903				0.61 *

409
 410 **Table 2: Pearson correlation coefficients of all bioassays for WW.** The values were calculated according to the method
 411 mentioned in section 2.9. All correlations were significant with a p value <0.0001 (***)

	MELN	ER-GeneBLAzer	HeLA-9903	pYES
ER-CALUX	0.94 ***	0.98 ***	0.99 ***	0.89 ***
MELN		0.98 ***	0.94 ***	0.97 ***
ER-GeneBLAzer			0.97 ***	0.96 ***
HeLa-9903				0.88 ***

412 **4 Conclusions and trends**

413 By including E1, E2, and EE2 in the watch-list of the WFD, the European Commission recognised the
 414 need to assess environmental occurrence and impact of these endocrine disrupting substances. However,
 415 the current WFD monitoring approach, which is based on chemical analytical measurements and
 416 compliance with specific EQSs, has been shown to be limited with regard to the ability to detect these
 417 substances at required concentrations [18, 51]. As demonstrated in this study, chemical analytical
 418 methods (status 2015) were unable to quantify the steroidal estrogens E2 and EE2 at EQS concentrations
 419 in all samples although E1 was measured effectively. Using effect-based methods, EEQ concentrations
 420 could be determined in all samples. As these EEQ concentrations are the responses to mixtures of
 421 known as well as unknown substances, effect-based methods have the potential to be highly valuable
 422 tools complementing routine monitoring and water quality assessment for estrogenic compounds. Effect-
 423 based methods are of particular regulatory interest as tools to screen and prioritise samples for further
 424 analysis by chemical analytical methods. Furthermore, DIN/EN/ISO standards to determine the
 425 estrogenic potential of water samples – covering human cell lines (e.g. ER-CALUX) and yeast based
 426 assays – will be available in early 2018 under ISO/DIS19040. The availability of such standards will
 427 facilitate the integration of effect-based methods into regulatory schemes.

428 Our study showed that EEQ results obtained from all effect-based methods applied were comparable –
 429 especially at higher concentrations found in WW – but results can vary between methods based on the

430 relative effect potencies for individual substances. This has to be considered for the interpretation of data
431 and determination of threshold values. As stated above: 1) *in vitro* effect-based methods cannot deliver
432 single substance based measurements, but are suitable to assess overall estrogenicity in water samples
433 and 2) results of these methods need to be confirmed by advanced chemical analysis. Along these lines,
434 the inclusion of effect-based methods into monitoring programmes as a screening tool (detailed
435 description in Kase et al., [52]) for estrogenic substances in surface water bodies would be a valuable
436 complement to chemical analysis currently foreseen by the Directive 2013/39/EU and WFD [28, 56, 57].

437

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461

462 **Conflict of interests**

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465

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633

Supplementary Information

Effect-based and chemical analytical methods to monitor estrogens under the European Water Framework Directive

Sarah Könemann, Robert Kase, Eszter Simon, Kees Swart, Sebastian Buchinger, Michael Schlüsener, Henner Hollert, Beate I. Escher, Inge Werner, Selim Ait-Aïssa, Etienne Vermeirssen, Valeria Dulio, Sara Valsecchi, Stefano Polesello, Peter Behnisch, Barbora Javurkova, Olivier Perceval, Carolina Di Paolo, Daniel Olbrich, Eliska Sychrova, Rita Schlichting, Lomig Leborgne, Manfred Clara, Christoph Scheffknecht, Yves Marneffe, Carole Chalon, Petr Tušil, Premysl Soldán, Brigitte von Danwitz, Julia Schwaiger, Maria Isabel San Martín Becares, Francesca Bersani, Klara Hilscherova, Georg Reifferscheid, Thomas Ternes, Mario Carere

Table S1: Author contributions calculated according to Clement Prabhakar, T. [1].

	Ideas	Work			Writing	Stewardship	Ideas	Work			Writing	Stewardship	Sum	Total contribution
		Sampling	Labwork	Data eval.				Sampling	Labwork	Data eval.				
weighting	0.1	0.2	0.2	0.2	0.3	0.2	0.1	0.2	0.2	0.2	0.3	0.2	1.0	
Sarah Könemann				15.0	30.0		0.0	0.0	0.0	2.3	9.0	0.0	11.3	11.3
Robert Kase	44.4			15.0	13.0	10.5	2.2	0.0	0.0	2.3	3.9	2.1	10.5	10.5
Mario Carere	44.4				13.0	10.5	2.2	0.0	0.0	0.0	3.9	2.1	8.2	8.2
Eszter Simon				25.0	10.0		0.0	0.0	0.0	3.8	3.0	0.0	6.8	6.8
Kees Swart			39.1				0.0	0.0	5.9	0.0	0.0	0.0	5.9	5.9
Sebastian Buchinger				20.0	5.0	5.3	0.0	0.0	0.0	3.0	1.5	1.1	5.6	5.6
Michael Schlüsener			17.4	10.0			0.0	0.0	2.6	1.5	0.0	0.0	4.1	4.1
Henner Hollert					5.0	10.5	0.0	0.0	0.0	0.0	1.5	2.1	3.6	3.6
Beate Escher					8.0	5.6	0.0	0.0	0.0	0.0	2.4	1.1	3.5	3.5
Inge Werner						15.3	0.0	0.0	0.0	0.0	0.0	3.1	3.1	3.1
Selim Ait-Aissa			8.7	10.0			0.0	0.0	1.3	1.5	0.0	0.0	2.8	2.8
Etienne Vermeirssen					5.0	5.3	0.0	0.0	0.0	0.0	1.5	1.1	2.6	2.6
Valeria Dulio						10.5	0.0	0.0	0.0	0.0	0.0	2.1	2.1	2.1
Sara Valsecchi		8.3			2.5		0.0	1.2	0.0	0.0	0.8	0.0	2.0	2.0
Stefano Polesello		8.3			2.5		0.0	1.2	0.0	0.0	0.8	0.0	2.0	2.0
Peter Behnisch				5.0		5.3	0.0	0.0	0.0	0.8	0.0	1.1	1.8	1.8
Olivier Perceval	11.1					5.3	0.6	0.0	0.0	0.0	0.0	1.1	1.6	1.6
Daniel Olbrich			8.7				0.0	0.0	1.3	0.0	0.0	0.0	1.3	1.3
Eliska Sychrova			8.7				0.0	0.0	1.3	0.0	0.0	0.0	1.3	1.3
Rita Schlichting			8.7				0.0	0.0	1.3	0.0	0.0	0.0	1.3	1.3
Isabel San Martin Becares			8.7				0.0	0.0	1.3	0.0	0.0	0.0	1.3	1.3
Lomig Leborgne		8.3					0.0	1.2	0.0	0.0	0.0	0.0	1.2	1.2
Manfred Clara		8.3					0.0	1.2	0.0	0.0	0.0	0.0	1.2	1.2
Christoph Scheffknecht		8.3					0.0	1.2	0.0	0.0	0.0	0.0	1.2	1.2
Yves Marneffe		8.3					0.0	1.2	0.0	0.0	0.0	0.0	1.2	1.2
Carole Chalon		8.3					0.0	1.2	0.0	0.0	0.0	0.0	1.2	1.2
Petr Tusil		8.3					0.0	1.2	0.0	0.0	0.0	0.0	1.2	1.2
Premysl Soldan		8.3					0.0	1.2	0.0	0.0	0.0	0.0	1.2	1.2
Brigitte von Danwitz		8.3					0.0	1.2	0.0	0.0	0.0	0.0	1.2	1.2
Julia Schwaiger		8.3					0.0	1.2	0.0	0.0	0.0	0.0	1.2	1.2
Francesca Bersani		8.3					0.0	1.2	0.0	0.0	0.0	0.0	1.2	1.2
Klara Hilscherova						5.3	0.0	0.0	0.0	0.0	0.0	1.1	1.1	1.1
Thomas Ternes						5.3	0.0	0.0	0.0	0.0	0.0	1.1	1.1	1.1
Georg Reifferscheid						5.3	0.0	0.0	0.0	0.0	0.0	1.1	1.1	1.1
Barbora Javurkova					3.0		0.0	0.0	0.0	0.0	0.9	0.0	0.9	0.9
Carolina di Paolo					3.0		0.0	0.0	0.0	0.0	0.9	0.0	0.9	0.9
Sum	99.9	99.6	100.0	100.0	100.0	100.0	5.0	14.9	15.0	15.0	30.0	20.0	99.9	100.0

Calculation of the minimal total contribution (MTC): $MTC = \frac{0.3 \cdot 100}{n}$ with n = 36

1 Part A: Sampling and sample preparation

2 Given that chemicals from a broad range of compound classes can interact with the estrogen receptor – both in an
3 agonistic and in an antagonistic way – there is a wide scope of possible sources of contamination of the samples. To
4 allow for a robust analysis of the samples, any contamination needs to be avoided.

5 Samples are taken by means of a scoop. The sample container on the scoop was cleaned before use by rinsing its in
6 and outside **three times** with acetone. Rinsing is performed by spraying acetone from a PTFE wash bottle onto the
7 scoop. Subsequently, the container was rinsed using ambient water (i.e. river water or effluent that will be sampled)
8 by dipping it in ambient water and filling and emptying the container **five times**. After this preparation, bottles were
9 filled one after the other. One person – **using clean nitrile gloves** – removed the lid from the first bottle and places
10 this lid on a clean surface. The other person used the scoop to collect a sample and poured the sample into the
11 bottle. Sample scooping and addition of sample to the bottle was repeated until the bottle was 2/3 full (i.e. 3.75 L of
12 sample). Next, the lid was removed from bottle 2 and used to seal bottle 1; now bottle 2 was be filled. This process
13 was continued until all bottles were filled. The last bottle in the sampling series received the lid that came off the
14 first bottle. In order to prevent degeneration of substances, 3.75 mL of 3 M H₂SO₄ were added to adjust the pH to
15 approximately 3. As soon as possible, the samples were frozen and sent to the central lab on dry ice, where they
16 were stored at -20 °C until extraction.

17 The further preparation of samples generally involved the extraction of samples according to the SPE procedure to a
18 1000-fold concentration. For the solid-phase extraction (SPE) 11 L of each sample were thawed overnight at room
19 temperature. Then the samples were homogenised and divided into 1 L aliquots. By addition of 3 M H₂SO₄ the pH
20 was adjusted to 3. Approximately 40 % of the surface water samples **(A(11), B(6), D(22), E(27), I(8), M(28), O(3))** and
21 all waste water samples were filtered over glass fiber filters (Millipore, type 4, retention 2.7 µm, circle size 4.7 cm)
22 before extraction. A 1 L procedure blank with ultrapure water adjusted to pH 3 was processed in parallel. In case a
23 sample needed to be filtered, the procedure blank was filtered as well. Prior to the extraction, the cartridges
24 (Phenomenex Strata C₁₈-E, 55 µm, 70 Å, 500 mg/6 mL) were conditioned with 6 mL of n-hexane, 2 mL of acetone,
25 6 mL of methanol and finally 10 mL of ultrapure water adjusted to pH 3. Subsequently, each cartridge was loaded
26 with 1 L of sample using a continuous flow setup with 5 mL/min. To determine the exact volume of the sample that
27 was extracted, the glass bottles were weighed before and after the extraction process. After 30 min of drying with
28 full vacuum, the cartridges were eluted with 4 mL acetone. Pertinent extracts were combined and the collection
29 tubes were rinsed with 1 mL acetone. Then the extracts were evaporated to a final volume of 200 µL acetone.
30 Procedural blank extracts were subjected to the same protocol.

31 Subsequently, all samples underwent a silica clean-up. For the clean-up step, all extracts were reconstituted in 1 mL
32 n-hexane / acetone (65:35, v/v) and applied to a silica column (CHROMABOND SiOH, Macherey-Nagel, 6 mL, bed size
33 1000 mg), which was previously activated (3 h, 85 °C) and conditioned with 10 mL n-hexane / acetone (65:35, v/v).
34 For surface water extracts one column was used for one mL of sample, whereas for waste water extracts one column
35 was used for all 11 mL of the sample. After sample application the column was rinsed with 2 mL of n-hexane /
36 acetone (65:35, v/v) and eluted with 4 mL of the same mixture. The extracts were then dried with N₂ at 45 °C and
37 reconstituted in 11 mL ethanol resulting in a 1000-fold concentration of the initial water sample (enrichment factor
38 1000).

39 The final 1 mL of all extracted samples including the blanks and the spiked water reference samples were then sent
40 to all project partners to be analysed by LC MS/MS as well as effect-based techniques.

41

42 Part B: Data evaluation and EEQ calculation

43 Data collection

44 The bioassays of this study are based on the same principle (i.e. binding of a ligand to an estrogen receptor with a
45 subsequent activation of a reporter gene), however they vary in the measured endpoint (i.e. luciferase activity by
46 bioluminescence or the photometric measurements after the induction of the β -galactosidase enzyme) and in
47 certain experimental parameters (e.g. cell types and density, % and type of organic solvent in the exposure medium,
48 cell culture and exposure mediums, and extracts dilution applied in the assay).

49 Each participating laboratory received the same number of E1, E2 and EE2 stocks, ethanolic extracts of extraction
50 blanks, positive controls and encoded surface and wastewaters (1 mL) sent by the extraction lab together with the
51 list of weights of the extracts. The participating labs were asked to analyze all blanks, references, controls and
52 sample extracts in dilution series following their own protocol and report any further handling the extracts
53 underwent during the analysis (e.g. solvent exchange, dilution etc).

54 All raw data together with the relative enrichment factors (REFs - the multiplication of the enrichment factor of the
55 sample extraction and the dilution factor of the extract in each of the bioassays) and the test concentrations
56 (dilutions) of the samples were provided by the participating laboratories and evaluated centrally.

57 Data evaluation

58 Bioassay data were normalised based on the measured response for the negative control (i.e. negative control) in
59 the respective bioassay (0%) and the highest induction measured for the reference substance (100%). The PC₁₀
60 approach was used as EEQ derivation procedure for data evaluation, [2]

61 First the measured bioassay responses were fitted using the five-parameter logistic regressions (Equation 1). This
62 model - unlike the four-parameter model - is suitable to handle asymmetric curves, that are often observed when
63 analysing environmental samples [2].

$$64 \quad y = \frac{a - d}{\left(1 + \left(\frac{x}{c}\right)^b\right)^f} + d \quad (1)$$

65 $y =$ the calculated effect measure (e.g. corrected absorbance) at concentration x

66 $x =$ the compound concentration which activates the test system to effect measure y

67 $a =$ minimum response (mean value fixed to the measurement of the solvent blank; bottom curve point)

68 $d =$ maximum response (mean value of y with the maximal activation of the test; curve plateau)

69 $c =$ the curve point of inflection, in case of a symmetric concentration-response curve C equals the mean effect concentration
70 at which the estrogenic effect reaches half of its maximum (EC50 - 50 % effect concentration)

71 $b =$ Slope refers to the steepness of the curve

72 $f =$ asymmetry factor, which reflects the asymmetry of the curve. In case of a symmetric curve f is equal to one.

73

74 The REF of the sample at a certain effect level (e.g. 10 %-effect) of the reference compound E2 - was calculated by
75 inserting the associated effect measure into the inverse of the function describing the concentration response
76 relationship of the sample. The respective REF of the sample equals the PC_x of the reference compound. The
77 appropriate effect level of the reference compound was defined as 10 % for each bioassay. This effect level in the
78 sample extracts was above the assays sample-dependent LOQs and we assumed that this level can be reached and
79 quantified by low contaminated surface water samples as well. Further, the 10 % threshold is statistically different
80 from the control. The estrogenic activity of the sample was determined by dividing the PC₁₀ by the derived REF
81 (REF₁₀) and expressed as pg EEQ/L water.

82 Calculation of 17β-estradiol equivalents (EEQs) for test samples

83 First, the effect measure y_{PCx} at the x%-effect of the reference compound was calculated by the following equation
84 (Equation 2):

$$85 \quad y^{PCx} = \left(\frac{d-a}{100} \cdot e \right) + a \quad (2)$$

86

87 y_{PCx} = the effect measure at the x%-effect level of the reference compound

88 e = the x%-effect level of the reference (e.g. 10 for the estimation of the 10%-effect level)

89 d = the mean value of y with the maximal activation of the test (curve plateau derived from the concentration response
90 relationship of the reference by the curve fitting)

91 a = the mean value of y without estrogenic effects (bottom curve point derived from the concentration response relationship
92 of the reference by the curve fitting)

93

94 Then, y_{PCx} was inserted into the inverse of the function describing the concentration-response relationship of the
95 sample for calculating the REF of the sample that induce the bioassay to the same extend as the PC_x of the reference
96 compound (Equation 3).

$$97 \quad REF = \left(\left(\frac{a-d}{y_{PCx}-d} \right)^{\left(\frac{1}{f}\right)} - 1 \right)^{\left(\frac{1}{b}\right)} \cdot C \quad (3)$$

98

99 REF = the REF of the sample that induce the same effect as the PC_x of the reference compound

100 y_{PCx} = the effect measure at the x%-effect level of the reference compound

101 d = the mean value of y with the maximal activation of the test (curve plateau derived from the concentration response
102 relationship of the sample by the curve fitting)

103 a = the mean value of y without estrogenic effects (bottom curve point derived from the concentration response relationship
104 of the sample by the curve fitting)

105 C = the curve point of inflection derived from the concentration response relationship of the sample by the curve fitting

106 b = Slope refers to the steepness of the curve and is proportional to the slope of the function at C (derived from the
107 concentration response relationship of the sample by the curve fitting)

108 f = assymetry factor, which reflects the assymetry of the curve and derived from the concentration-response relationship of
109 the sample by the curve fitting. In case of a symmetric curve f is equal to one.

110

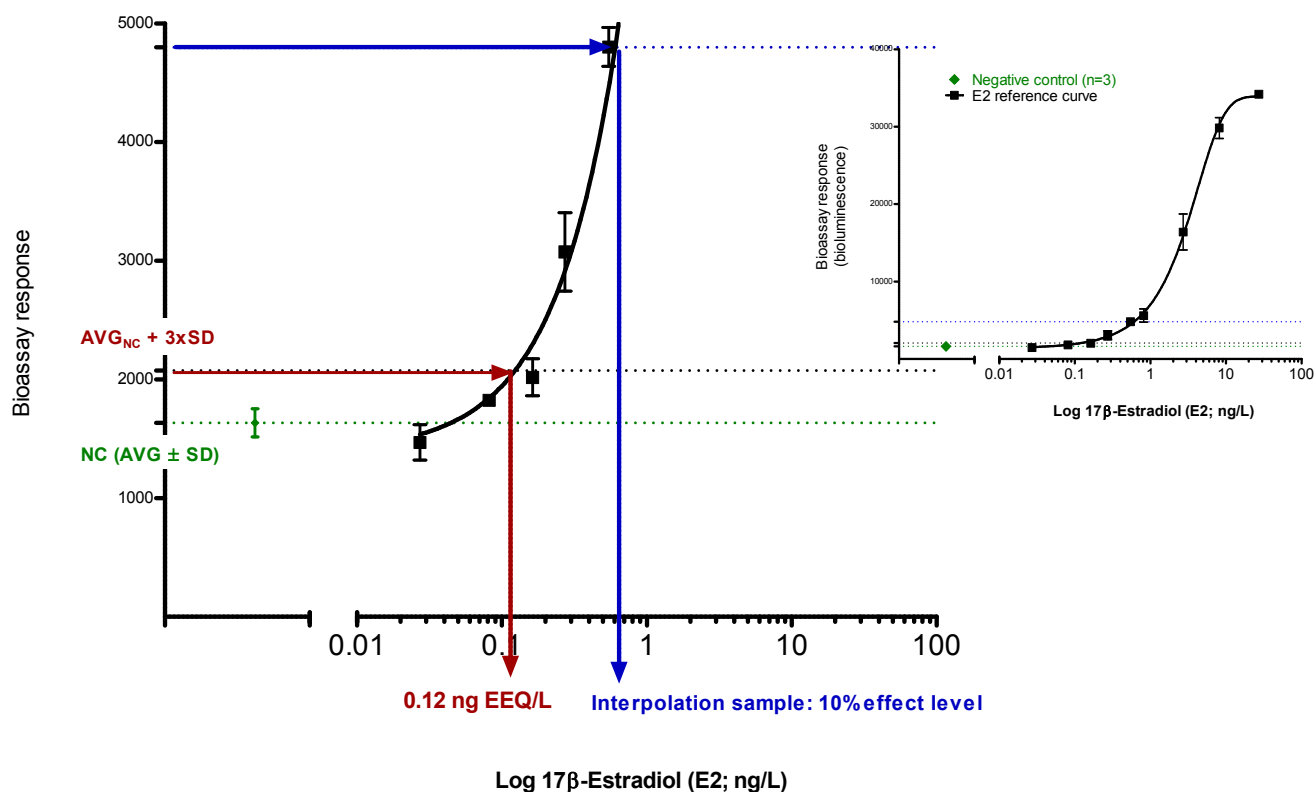
111 The estrogenic potential of the sample as was estimated by the quotient of PC_x of the reference compound, E2 and
112 the calculated REF_s of the sample.

113

114 Part C: LOQ determination for bioassays and LC-MS/MS analyses

115 LOQs for bioassays

116 The sample-specific LOQs in the bioassays were determined based on the variability of the response of the negative
117 controls tested along the sample(s) and the highest sample-specific REF (relative enrichment factor) tested. The
118 average response of the negative control (SC) replicates plus three times the standard deviation (SD) was
119 interpolated from the E2 dose-response curve. This resulting E2-equivalent concentration (ng EEQ/L) was then
120 divided by the highest tested REF of the sample to obtain a sample-specific LOQ. The figure below illustrates the
121 procedure using an example from ER-CALUX.



122

123 **Figure S1. Graphical representation of the sample-specific LOQ determination in the ER α -CALUX bioassay.** The small figure shows the full E2
124 dose-response curve, the large figure shows the lower part of the curve for better clarity. The response measured for the negative control was
125 1634 ± 118 RLU (relative light units; bioluminescence). Based on this measurement the LOQ threshold was determined: $AVG_{\text{negative control}} + 3 \times SD$
126 $= 2075$ RLU. This value was then interpolated from the E2 curve and an EEQ of 0.12 ng/L was obtained. Dividing this EEQ by the highest REF of
127 the sample (in this particular case: 23.3) leads to a sample-specific LOQ of 0.005 ng EEQ/L (5.3 pg EEQ/L).

128 It is important to note two aspects: 1) sample EEQ concentrations were derived using the 10 % effect level of the E2
129 curve (PC₁₀); 2) sample curve fitting was only performed when “at least two effect measurements above 10 % were
130 obtained” (see Section E of the SI).

131 Considering the first point: a “PC₁₀ reporting level LOQ” for the above sample can be interpolated from the E2 curve
132 along the blue arrows. The 10 % effect level equates to 4720 RLU and intercepts the E2 curve at 0.70 ng/L. For the
133 sample above, with an REF of 23.3, the sample-specific PC₁₀ reporting level LOQ is 29 pg/L.

134 Considering the second point: for a robust fitting of the PC₁₀, a higher sample EEQ concentration than 29 pg/L is
135 required and the effective LOQ is thus higher than the PC₁₀ derived LOQ.

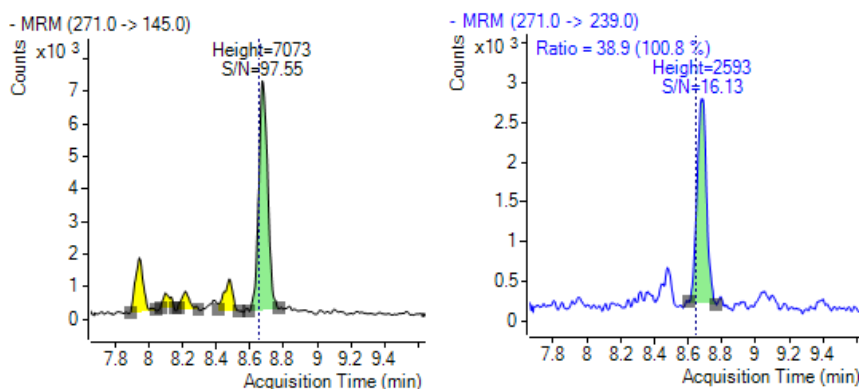
136

137 LOQs for LC-MS/MS

138 The figure below illustrates the procedure followed by Lab 2 for two examples. In the first example, the analyte was
139 detected above a S/N of 10, in the second example the analyte was not detected and standard addition was used
140 to determine the LOQ. Both the quantifier and the qualifier should have a S/N ≥ 10 .

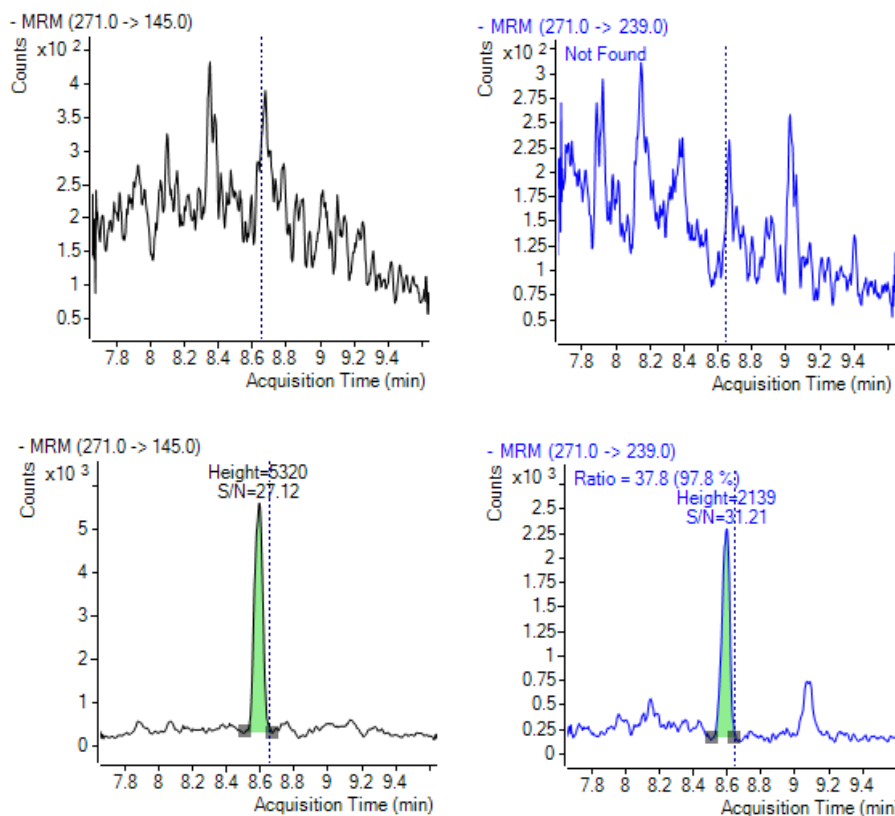
141 Example 1

142 The first panel shows the E2 peak in a wastewater sample. With a concentration of 0.55 ng/L and an S/N of 97 (graph
143 on the left; quantifier) and S/N 16 (graph on the right; qualifier) the data were normalized to a S/N of 10 as follows:
144 $0.55 \text{ ng/L} / (16/10) = 0.34 \text{ ng/L}$. This theoretical concentration should have a S/N of 10:1 and is defined as LOQ for this
145 analyte in this sample.



146
147 Example 2

148 The top panel shows the absence of a peak (quantifier on the left and qualifier o the right). The bottom panel shows
149 the same sample with the addition of 0.5 ng/L with a S/N of 27 and S/N 31. After normalization to a S/N of 10 the
150 LOQ was calculated 0.19 ng/L (i.e. $0.5 \text{ ng/L} / (27/10) = 0.19 \text{ ng/L}$).



151

152

153

154

Table S2: Detailed information on analytical methods used for the analysis of steroidal estrogens.

	Laboratory 1	Laboratory 2	Laboratory 3
Sample preparation	Extracts were evaporated to dryness under gentle N ₂ stream and then reconstituted to 0.2 ml reconstituting solution (NH ₄ OH 0.1%: AcN, 9:1, % v/v)	No additional clean-up was performed on the samples, except for sample 20 and sample 23. For these samples the silica SPE clean-up was repeated as carried out by BDS (but this time using 1 SPE column per extract) Generally, 500 µL extract were taken, IS added, evaporated till dryness, and reconstituted in 50 µL MeOH and 50 µL Water; concentration factor 1:5000	Internal Standards were added. Extracts were evaporated to dryness under gentle N ₂ stream and then reconstituted in 0.5 ml methanol followed by 0.5 ml water.
System	LC-MS/MS	LC-MS/MS	LC-MS/MS
Pumps	Binary Solvent Manager, Model UPB, Waters (Milford, MA, USA).	Binary Pump: Agilent G7120A Quaternary Pump for post-column addition: G1311C	Agilent G1312B binary LC pump, a G1310B isocratic LC pump
Autosampler	Sample Manager, Model UPA, Waters (Milford, MA, USA).	Multisampler: Agilent G7167B	Agilent G1367E
Ionization method	ESI Negative	ESI Negative	ESI Negative
Detector	QTRAP 5500 MS	Agilent 6495 QQQ	Sciex 6500 Qtrap One Quantifier MRM transition Two Qualifier MRM transition
Flow rate	400 µl/min	500 µL/min	300 µL/min
Injection volume	30 µl	20µL	10, 30, 100 µL
Column	Hypersil GOLD, 1.9 µm, 50 x 2.1 mm, Thermo Scientific	ACQUITY UPLC BEH Shield RP18 Column, 130Å, 1.7 µm, 2.1 mm X 100 mm, [Waters 186002854] at 40°C	Poroshell C18-EC column (3 x 50 mm, 2.7 µm, Agilent) at 25 °C
Gradient	A: NH ₄ OH 0.1 % B: Acetonitrile 0 to 0.5 min 10% B, from 1 to 5 min 40% B, from 5 to 6 min 90% B, from 6.5 to 12 min 10% B	A: 5mM NH ₃ in water B: Methanol, postcolumn addition of 0.05ml/min 600µM NH ₄ F 0 to 0.19 min, 10% B; from 0.19 to 0.2 min, 10 to 30% B; from 0.2 to 9.0 min, 30 to 74% B; from 9.0 to 10.3 min 74 to 100% B; from 10.3 to 14 min, 100% B; equilibration time 3min before analysis	A: Water B: Acetonitrile 0 to 0.5 min 10% B; from 0.5 to 1.0 min 10 to 45% B; from 1.0 to 9.0 min 45 to 60% B; from 9.0 to 9.1 min 60 to 98% B; from 9.1 to 12 min 98% B; from 12.0 to 12.1 min 98% to 10% B, from 12.1 to 15.0 min 10% B
Internal Standards	E1 13C3, E2 d4, EE2 d4	Estrone-2,4,16,16-d4 (D-3650, cdn isotopes), Estradiol-2,4,16,16-d4 (DLM-2487-0, CIL), 17-α-EE2-2,4,16,16-D4 (DLM-4691, CIL)	Estrone 2, 4, 16, 16 - d4 (E2-d4), 17β-estradiol 2,4,16,16-d4, and 17α-ethinyl estradiol 2, 4, 16, 16-d4 (EE2-d4) were obtained from Toronto Research Chemicals (North York, ON, Canada)

Table S3: Detailed information on effect-based methods used for determination of EEQs.

	ER α CALUX	MELN	ER-GeneBLazer	HeLa-9903	pYES
Cell-line	Human U2OS osteosarcoma cells stably transfected with 3xHRE-TATA-Luc and pSG5-neo-hER α constructs using calcium phosphate co-precipitation method	Human MCF-7 breast cancer cells stably transfected with an ERE- β Globine-luciferase construct. This cell line expresses endogenously hER α and hER β while only hER α is functional and activates the reporter gene	GeneBLazer [®] ER alpha -UAS-bla GripTite™ cells (genetically engineered from HEK293) using β -lactamase reporter gene under control of a UAS response element	Human HeLa-9903 cervical tumor cells stably transfected with two constructs: (i) the hER α expression construct (encoding the full-length human receptor), and (ii) a firefly luciferase reporter construct bearing five tandem repeats of a vitellogenin Estrogen-Responsive Element (ERE) driven by a mouse metallothionein promoter TATA element.	<i>Saccharomyces cerevisiae</i> BJ3505 transfected with the plasmids YEPE10 and YRPEG3. YEPE10 contains a CUP1::hER fusion encoding the human estrogen receptor alpha cloned from the MCF-7 human cell lineage under the control of the metallothionein promoter CUP1. YRPEG3 contains the fusion gene 2ERE-CyC1::lacZ. This fusion gene express β -galactosidase under the control of the CyC1 promoter from <i>S. cerevisiae</i> fused to two copies of the vitellogenin A2-gene from <i>Xenopus laevis</i> .
Endpoint for estrogenicity	Luciferase activity	Luciferase activity	β -Lactamase via fluorescence (FRET) reagent with combined fluorescence cytotoxicity measure	Luciferase activity	β -Galactosidase activity
Assay medium	Phenol red free D-MEM/F 12) containing 5% DCC-FCS	Phenol red free DMEM containing 3% DCC-FBS (v/v)	Phenol red free DMEM (Gibco 10569-010) containing 2% DCC-FBS (Gibco 12676-011), 1% Penicillin-Streptomycin (Gibco 15140-122), 1mM Sodium Pyruvate (Gibco 11360-070), 0.1 mM NEAA (Gibco 11140-050)	Phenol red free DMEM - F12 (Sigma Aldrich, USA) + fetal calf serum (10% v/v) for maintenance. For experiments: Medium DMEM - F12 (Sigma Aldrich, USA) + dialyzed DCC-FBS (10% v/v)	28 mg/ml Yeast nitrogen base w/o amino acids, 130 mg/ml Glucose, 150 μ g/ml L-Lysine-HCL, 100 μ g/ml L-Histidine-HCL, 25 μ g/ml CuSO ₄ •7H ₂ O, supplemented with Ampicillin and Streptomycin
Assay format	96-well microtiter plates	96-well microtiter plates	384-well microtiter plates	96-well microtiter plates	High performance thinlayer plate (silica)
Cells per well seeded	1 x 10 ⁴	0.5-1 x 10 ⁴	2 x 10 ⁴	1 x 10 ⁴	n.a.
Dosing of cells	After 24 h seeding, the growth medium is replaced by the assay medium containing the reference compound or the sample extract in respective dilutions (100 μ l/well).	After 24h seeding in medium (100 μ l/well) and another 50 μ l/well of assay medium containing the test chemical or sample (3x concentrated) is added to the cells. Only internal wells on the assay plate	8 μ l/well of dosing media containing the test chemical or sample	Cells are seeded for 24h in medium (100 μ l/well) and another 100 μ l/well of dosing media containing the test chemical or sample is added for cells exposure. Total volume of media is therefore 200 μ l/well.	Overnight-culture of yeast cells is adjusted to 1000 FNU. 5 ml of the cell suspension is applied evenly on a TLC-plate (10x20 cm) after chromatographic separation of the extract. Applied volumes of the

	are used.				extracts vary between 5 µl and 100 µl depending on the expected contamination level.
Incubation period for exposure	22-24h	16-24h	16h	24h	3h
Number of replicate wells	3	3 (manual method) or 4 (automated method)	2 replicates + minimum 2 independent repeats of the assay	3	n.a. Assay is repeated three times independently
Negative controls	0.1% DMSO in the assay medium in triplicate pro assay plate (negative control)	0.1-0.5% DMSO in the assay medium (depending on the expected low activities) and assay medium without DMSO	32 replicates of assay medium per assay plate (No negative controls ^a)	0.1% (v/v) MeOH on every plate in triplicates and medium in triplicates	5 µl of Ethanol for negative control, additionally 100 µl of the extracted field blank sample applied on a separate lane on the TLC-plate
Concentration range tested of the reference compound, 17β-estradiol (E2)	0.03-27 ng/L The full concentration range was tested on each assay plate	0.03 – 27 ng/L The full concentration range was tested in each experimental series and a fixed concentration of E2 (10 nM) E2 on each assay plate (in in sextuplicate)	0.3 – 545 ng/L The full concentration range (6 - 12 concentrations in duplicate) was tested on each plate	0.001 - 10 ng/L The full concentration range (at least 5 concentrations in triplicate) was tested on each plate	1 pg/L to 10 pg/L E2 and EE2, 10 pg/L to 100 pg/L E1, 100 pg/L to 1000 pg/L E3
Cell harvesting at the end of the exposure and response detection	Exposure medium was removed and cells were lysed with 30 µL Triton-lysis buffer to open up the cell membrane. Then the luciferin substrate mix was added to the cells and the luciferase activity was measured at 0.1 min/well	Cells were first washed with PBS buffer (optional) and 50 µL of medium containing 30 mM of D-luciferin substrate was added. Cells were not lysed. After 5 min plates were read on a luminometer, 1s per well.	8µl per well and incubated for 2h at room temperature Fluorescence was measured immediately after adding the substrate buffer (time 0h for correction of autofluorescence) and then after 2h-incubation using the same gain for both measurements	50 µL of cell lysis solution enriched by luciferin solution was added to each well. Plates were read with a luminometer after 10 min of shaking (150 rotations/min) at room temperature	After exposure the TLC-plate is developed by the application of the following buffer: 10 mg/ml Na ₂ HPO ₄ •2 H ₂ O, 0.75 mg/ml KCl 0.25 mg/ml MgSO ₄ •7 H ₂ O, 1 mg/ml SDS, 0.5 mg/ml MUG, pH=7.0 After application the plate is incubated for 15 min at 37°C. The fluorescence-signal of the developed Methyl-umbelliferon is detected at 254 nm.

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Abbreviations: **DCC-FCS**: Dextran-charcoal treated Fetal Calf Serum; **DCC-FBS**: Dextran-charcoal treated Fetal Bovine Serum; **D-MEM**: Dulbecco's modified Eagle's medium; **MUG**: 4-Methylumbelliferyl β-D-galactopyranoside; **SDS**: Sodium dodecyl sulfate

^aThe extracts were dried and resuspended in assay medium. DMSO stock solution of reference compound is diluted 5x10⁵-times in the assay medium. Therefore, the amount of solvent in the positive control is negligible and negative control was not tested.

^b Phosphate buffered saline

Part E: General and specific validity criteria of the bioanalytical and chemical analysis methods

- No estrogenic activity or quantifiable concentration of E1, E2 and EE2 were measured in the blank samples (i.e. procedure-, extraction- and solvent blanks)
- Derived effect concentrations in the bioassays and chemically measured EE2 concentrations matched with the nominal concentrations of the spiked samples
- Participating laboratories reported their raw data in a standardized format and all data were then analysed centrally in a harmonized way.

Table S4: Measured chemical concentrations and 17 β -estradiol (E2) equivalent concentrations (EEQ) of the positive control samples with high or low concentrations of E2 and EE2 are compared with the nominal chemical concentrations and calculated nominal EEQ concentrations.

Spiked water	Nominal spike conc.	Chemically determined spike conc. *	Measured conc. as % of nominal spike conc.	EEQ determined in the <i>in vitro</i> assays **	EEQ _{nominal} ***	EEQ as % of EEQ _{nominal}
	pg/L	pg/L		pg/L	pg/L	
E2 _{low}	600	617	103%	772	678	114%
E2 _{high}	6000	5651	94%	4914	5717	86%
EE2 _{low}	600	770	128%	914	975	94%
EE2 _{high}	6000	6341	106%	6134	7548	81%
AVG \pm SD			108 \pm 15%			94 \pm 14%

*Averaged spiked concentration (pg/L) of the three measured values provided by the analytical labs (JRC, BfG, OZ)

**Averaged EEQ (pg/L) obtained in the 5 *in vitro* bioassay

***EEQ_{nominal} (pg/L) was calculated by multiplying the chemically measured and averaged E2 or EE2 concentrations by their assay-specific relative potency (REP). In all four spiked water samples E1 concentration was measured. Similarly to E2 and EE2, the chemically measured E1 concentrations were also translated into EEQ concentrations and added to the pertinent EEQ_{nominal} concentration.

Effect-based methods

- Each participating laboratory used the same batch of certified E2 standard as reference compound in their bioassay and the same batch of E1 and EE2 standard solutions to determine their assay-specific relative potencies (REPs). REPs are summarised in Table S4.
- Sample dilutions with observed cytotoxicity were not taken for data evaluation.
- Additionally to the assessment criteria of the respective laboratory (e.g. acceptable CV of replicate measurements), the reported data fulfilled sufficient criteria elements that allowed for a robust data evaluation. The evaluation was performed collectively by a group of three experts. Bioassay data should have included a full reference dose-response curve (below 10% and above 90%) and had sufficient values at the lower effect levels (e.g. <10% effect) for both reference and samples. These latter criteria were especially important for the PC₁₀ EEQ derivation method. In the case of incomplete curves, at least two effect measurements above 10% were obtained that allowed curve fitting.

Table S5: Assay-specific mass-based 17 β -estradiol relative potency factors (REPs). REP shows the relative potency of the compound in the certain bioassay compared to the reference compound (E2 in this case). REPs were determined by dividing the 50% effect concentration (EC₅₀; g/L) of the reference compound by the EC₅₀ of the test compound (g/L).

REP (g _{E2} /g _i)	E1	E2	EE2
ER α -CALUX	0.01	1.0	1.2
MELN	0.29	1.0	0.79
ER-GeneBLazer	0.08	1.0	1.67
HeLa-9903	0.02	1.0	1.18
pYES	0.11	1.0	1.0

LC-MS/MS

- Only peaks with a signal-to-noise (S/N) ratio > 10:1 were evaluated and quantified.
- The acceptable retention time deviation relative to the pertinent internal standard was <0.2 min within the same sequence and the uncertainty of the data qualifier <30%.

Table S6: Results of the three chemical analytical methods. Measured concentrations are shown for surface and waste water in pg/L.

		E1 in pg/L			E2 in pg/L			EE2 in pg/L		
		Lab 1	Lab 2	Lab 3	Lab 1	Lab 2	Lab 3	Lab 1	Lab 2	Lab 3
Spikes and Blanks	Blank	<LOD	<LOD	<LOD	<LOD	<LOD	<LOD	<LOD	<LOD	<LOD
	Blank-Spike	<LOD	<LOD	<LOD	<LOD	<LOD	<LOD	<LOD	<LOD	<LOD
	E2 600	1020	760	800	741	571	540	<LOD	<LOD	<LOD
	E2 6000	773	531	640	6022	5531	5400	<LOD	<LOD	<LOD
	EE2 600	703	524	560	<LOD	<LOD	<LOD	685	874	750
	EE2 6000	204	108	120	<LOD	<LOD	<LOD	6170	6654	6200
Surface water samples	A(11)	82	<LOQ	51	<LOQ	<LOD	<LOD	<LOD	<LOD	<LOD
	B(6)	88	<LOQ	70	<LOQ	<LOD	<LOD	<LOD	<LOD	<LOD
	C(1)	186	96	130	<LOQ	<LOQ	<LOQ	<LOD	<LOD	<LOD
	D(22)	189	114	130	<LOQ	<LOD	<LOQ	<LOD	<LOD	<LOD
	E(27)	221	136	160	<LOQ	<LOQ	<LOQ	73	75	85
	F(30)	400	318	360	<LOQ	<LOD	<LOQ	<LOD	<LOD	<LOD
	G(32)	444	282	360	<LOQ	<LOQ	70	<LOD	<LOD	<LOD
	H(25)	732	461	600	<LOQ	<LOQ	<LOQ	<LOD	<LOQ	<LOD
	I(8)	717	691	630	<LOQ	81	<LOD	<LOD	<LOD	<LOD
	J(10)	933	702	640	<LOQ	59	70	<LOD	<LOD	<LOD
	K(18)	1348	1106	960	261	171	180	186	161	90
	L(24)	1971	1124	1600	259	<LOQ	190	<LOD	<LOQ	<LOD
	M(28)	3353	2212	3000	267	230	240	<LOD	<LOQ	<LOD
	N(15)	5626	3307	4400	862	431	220	<LOD	<LOQ	<LOD
	O(3)	6186	5220	4000	311	441	<LOD	354	230	<LOD
	P(7)	7136	4846	4700	403	215	<LOD	100	<LOQ	<LOD
Waste water samples	A(26)	103	<LOQ	56	<LOQ	<LOD	<LOD	<LOD	<LOD	<LOD
	B(29)	116	87	96	<LOQ	<LOQ	<LOD	<LOD	<LOQ	<LOD
	C(31)	215	125	130	<LOD	<LOQ	<LOD	<LOD	<LOD	<LOD
	D(4)	268	162	210	<LOQ	<LOQ	<LOD	<LOD	<LOQ	<LOD
	E(17)	390	284	250	<LOQ	<LOD	<LOD	119	<LOQ	<LOD
	F(21)	509	364	460	<LOQ	<LOD	<LOD	<LOD	<LOD	<LOD
	G(14)	852	816	810	<LOQ	<LOD	<LOD	<LOD	<LOQ	<LOD
	H(5)	2502	1848	1900	<LOQ	<LOQ	<LOD	<LOD	<LOQ	<LOD
	I(19)	2955	3150	2500	<LOQ	<LOQ	<LOD	<LOD	<LOQ	<LOD
	J(16)	4351	5490	3300	<LOQ	<LOQ	<LOD	<LOD	218	<LOD
	K(9)	5350	6020	4700	<LOQ	<LOQ	<LOD	<LOD	<LOQ	<LOD
	L(13)	8022	7971	5500	1177	551	<LOD	<LOD	<LOQ	<LOD
	M(23)	9485	5940	7200	<LOQ	1670	220	6124	7022	9400
	N(33)	10606	13080	12000	323	478	350	<LOD	<LOQ	<LOD
	O(12)	14746	15300	11000	461	471	430	<LOD	<LOQ	<LOD
	P(2)	14482	16000	12000	839	540	620	<LOD	<LOQ	<LOD
Q(20)	22648	21000	18000	1031	1074	1200	5864	5160	4000	

Table S7: Limits of quantification (LOQs) of the three chemical analytical methods. Determined concentrations are shown for surface and waste water in pg/L.

		LOQ for E1 in pg/L			LOQ for E2 in pg/L			LOQ for EE2 in pg/L		
		Lab 1	Lab 2	Lab 3	Lab 1	Lab 2	Lab 3	Lab 1	Lab 2	Lab 3
Spikes and Blanks	Blank	42	50	90	240		360			300
	Blank-Spike	42	50	90	312		360			300
	E2 600		50	90		60	360			300
	E2 6000		50	90		80	360			300
	EE2 600		50	90			360		50	300
	EE2 6000		50	90			360		80	300
Surface water samples	A(11)	9	50	15	108	50	120	45	60	30
	B(6)	3	50	15	66	50	45	36	70	30
	C(1)	6	50	15	147	50	45	36	50	30
	D(22)	6	50	15	63	50	45	24	80	30
	E(27)	9	50	15	39	50	45	60	70	30
	F(30)	18	50	15	90	70	120	600	100	90
	G(32)	12	50	15	63	80	45	78	50	30
	H(25)	30	50	15	234	80	120	120	90	90
	I(8)	24	50	45	360	50	120	72	120	90
	J(10)	9	50	15	72	50	45	30	70	30
	K(18)	21	50	15	240	80	120	135	150	90
	L(24)	192	50	45	240	190	120	600	140	300
	M(28)	18	50	45	240	130	120	240	140	300
	N(15)	168	50	45	240	170	120	390	70	300
	O(3)	327	170	45	240	360	1500	60	230	900
P(7)	45	100	45	240	270	1500	72	200	300	
Waste water samples	A(26)	21	50	90	60	50	300	90	70	300
	B(29)	3	50	90	51	50	300	48	60	300
	C(31)	24	50	90	228	110	900	114	120	1500
	D(4)	12	100	90	450	160	3000	279	120	300
	E(17)	75	50	90	126	200	900	120	60	300
	F(21)	15	50	90	135	140	900	420	190	300
	G(14)	42	90	90	579	210	1500	900	290	1500
	H(5)	90	90	90	1080	670	3000	399	240	3000
	I(19)	27	50	90	480	140	900	1410	90	300
	J(16)	90	100	90	738	270	1500	2100	140	1500
	K(9)	123	50	90	1020	180	1500	720	80	3000
	L(13)	33	50	90	240	80	3000	600	110	300
	M(23)	144	1000	90	1200	290	600	810	1200	3000
	N(33)	33	70	90	240	220	900	1500	250	1500
	O(12)	306	120	90	240	290	900	1110	500	1500
P(2)	24	50	90	240	80	360	240	70	300	
Q(20)	156	90	90	240	150	900	1737	140	300	

Table S8: EEQ determined by effect-based methods using the PC₁₀ approach.

		EEQ _{Bio} in pg/L				
		ER CALUX	MELN	HeLa	ER-GeneBlazer	pYES
Surface water samples	A(11)	80	79	16	39	130
	B(6)	80	59	16	22	90
	C(1)	60	182	17	32	370
	D(22)	120	191	35	41	390
	E(27)	200	170	245	94	190
	F(30)	150	321	28	76	160
	G(32)	110	375	100	137	270
	H(25)	230	391	97	111	440
	I(8)	260	831	137	230	410
	J(10)	150	458	89	92	280
	K(18)	640	809	659	342	600
	L(24)	410	2124	509	407	2060
	M(28)	860	2498	226	487	1300
	N(15)	1180	1942	1039	876	2900
	O(3)	920	4019	487	959	5430
P(7)	590	2787	347	835	3600	
Waste water samples	A(26)	30	37	31	25	100
	B(29)	50	78	118	74	100
	C(31)	70	254	41	143	65
	D(4)	60	244	90	108	410
	E(17)	120	434	86	111	690
	F(21)	310	656	298	254	160
	G(14)	490	2066	75	567	970
	H(5)	370	1793	273	452	290
	I(19)	310	3104	553	560	1200
	J(16)	480	2936	888	813	2400
	K(9)	480	2233	692	501	340
	L(13)	870	2824	1205	1166	1700
	M(23)	22930	19716	24144	11892	12000
	N(33)	1050	3807	2407	1369	2400
	O(12)	700	5493	1695	1212	3400
P(2)	820	4928	734	1360	2800	
Q(20)	7590	10851	6442	6317	9700	

Table S9: Limits of detection (LODs) and quantification (LOQs) of effect-based methods.

	ER-CALUX		MELN		HeLa-993		ER-GeneBLazer		pYES		
	LOD in pg/L	LOQ in pg/L	LOD in pg/L	LOQ in pg/L	LOD in pg/L	LOQ in pg/L	LOD in pg/L	LOQ in pg/L	LOD in pg/L	LOQ in pg/L	
Surface water samples	A(11)	0	12	2	16	8	37	11	43	3	10
	B(6)	4	19	0	9	5	19	7	26	3	10
	C(1)	6	11	3	15	0	57	0	48	3	10
	D(22)	0	5	3	8	18	73	5	20	3	10
	E(27)	4	7	8	22	2	8	6	22	3	10
	F(30)	7	22	9	32	6	33	6	23	3	10
	G(32)	2	3	8	22	2	9	6	22	3	10
	H(25)	5	8	0	7	8	29	6	22	7	20
	I(8)	4	0	5	16		24	7	24	7	20
	J(10)	0	13	8	25	44	125	7	24	3	10
	K(18)		5	3	8		24	8	27	7	20
	L(24)	3	9	5	15		79	7	24	33	100
	M(28)	4	10	3	19		79	32	113	17	50
	N(15)		2	8	20	2	10	6	21	33	100
	O(3)	5	6	6	19		20	6	23	67	200
P(7)	1	3	7	20	5	25	6	22	33	100	
Waste water samples	A(26)	6	11	3	10	31	85	11	43	3	10
	B(29)	4	12	5	17	9	58	6	23	3	10
	C(31)	9	27	6	17	12	43	11	42	17	50
	D(4)	4	6	0	14	8	12	55	216	33	100
	E(17)	16	37	6	17	5	43	6	22	22	67
	F(21)	7	22	3	17	31	85	6	22	22	67
	G(14)	10	26	0	12	0	14	5	21	17	50
	H(5)	0	0	5	12	1	17	5	21	67	200
	I(19)	3	9	8	18	6	23	5	21	33	100
	J(16)	0	2	0	11	24	93	7	27	67	200
	K(9)	3	6	2	12	14	70	55	213	67	200
	L(13)	4	16	5	14	0	70	5	21	67	200
	M(23)	6	15	0	13	8	46	11	44	67	200
	N(33)	5	8	6	20	29	132	5	21	67	200
	O(12)	2	6	4	8	66	197	13	53	33	100
P(2)	15	36	3	17	2	9	5	18	33	100	
Q(20)	4	7	0	9	27	131	6	22	67	200	

Table S10: Sums of EEQ_{chem} based on the REPs obtained with ER-CALUX. For the left part of the table (>LOQ) only analytical data above the LOQ was included, while for the right part data below LOD or LOQ was substituted by LOD/2 and LOQ/2.

			>LOQ			<LOD = LOD/2 <LOQ = LOQ/2		
		ER-CALUX EEQ in pg/L	Sum EEQ_{chem} in pg/L			Sum EEQ_{chem} in pg/L		
			Lab 1	Lab 2	Lab 3	Lab 1	Lab 2	Lab 3
Surface water samples	N(15)	1180	918	464	264	996	506	324
	K(18)	640	498	375	298	498	375	298
	M(28)	860	301	252	270	349	336	330
	O(3)	920	798	769	40	798	769	470
	F(30)	150	4	3	4	49	35	82
	P(7)	590	594	263	47	594	383	357
	L(24)	410	279	11	206	399	190	266
	H(25)	230	7	5	6	148	99	84
	J(10)	150	9	66	76	51	80	82
	E(27)	200	90	91	104	109	116	126
	G(32)	110	4	3	74	52	53	80
	I(8)	260	7	88	6	202	112	44
	A(11)	80	1	0	1	64	21	27
	B(6)	80	1	0	1	41	23	14
Waste water samples	C(1)	60	2	1	1	83	36	30
	D(22)	120	2	1	1	38	25	30
	E(17)	120	147	3	3	210	72	213
	A(26)	30	1	0	1	49	23	111
	L(13)	870	1257	631	55	1377	697	615
	K(9)	480	54	60	47	708	198	897
	F(21)	310	5	4	5	157	65	215
	H(5)	370	25	18	19	645	497	1119
	D(4)	60	3	2	2	283	154	562
	O(12)	700	608	624	540	830	924	840
	J(16)	480	44	317	33	833	452	583
	P(2)	820	984	700	740	1032	742	800
	Q(20)	7590	8294	7476	6180	8294	7476	6180
	M(23)	22930	7444	10156	11572	8044	10156	11572
	G(14)	490	9	8	8	478	217	558
	I(19)	310	30	32	25	552	156	235
	B(29)	50	1	1	1	36	62	111
N(33)	1050	429	609	470	729	759	770	
C(31)	70	2	1	1	63	80	451	

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Table S11: Sums of EEQ_{chem} based on the REPs obtained with MELN. For the left part of the table (>LOQ) only analytical data above the LOQ was included, while for the right part data below LOD or LOQ was substituted by LOD/2 and LOQ/2.

	MELN EEQ in pg/L	>LOQ			<LOD = LOD/2 <LOQ = LOQ/2			
		Sum EEQ_{chem} in pg/L			Sum EEQ_{chem} in pg/L			
		Lab 1	Lab 2	Lab 3	Lab 1	Lab 2	Lab 3	
Surface water samples	N(15)	1942	2494	1390	1496	2545	1418	1536
	K(18)	809	799	619	530	799	619	530
	M(28)	2498	1239	871	1110	1271	927	1150
	O(3)	4019	2385	2137	1160	2385	2137	1529
	F(30)	321	116	92	104	161	117	176
	P(7)	2787	2551	1620	1363	2551	1699	1653
	L(24)	2124	831	326	654	910	476	694
	H(25)	391	212	134	174	345	209	246
	J(10)	458	271	263	256	311	272	260
	E(27)	170	122	99	114	141	124	136
	G(32)	375	129	82	174	171	128	178
	I(8)	831	208	281	183	397	297	215
	A(11)	79	24	0	15	84	23	39
	B(6)	59	26	0	20	63	25	32
Waste water samples	C(1)	182	54	28	38	132	59	64
	D(22)	191	55	33	38	89	52	64
	E(17)	434	207	82	73	270	139	262
	A(26)	37	30	0	16	72	25	106
	L(13)	2824	3503	2863	1595	3582	2906	2135
	K(9)	2233	1552	1746	1363	2156	1867	2008
	F(21)	656	148	106	133	270	154	323
	H(5)	1793	726	536	551	1318	966	1446
	D(4)	244	78	47	61	339	174	600
	O(12)	5493	4737	4908	3620	4883	5106	3818
	J(16)	2936	1262	1764	957	1907	1899	1405
	P(2)	4928	5039	5180	4100	5070	5208	4140
	Q(20)	10851	12231	11240	9580	12231	11240	9580
	M(23)	19716	7589	8940	9734	8189	8940	9734
	G(14)	2066	247	237	235	655	386	682
	I(19)	3104	857	914	725	1283	1019	915
B(29)	78	34	25	28	65	74	117	
N(33)	3807	3399	4271	3830	3596	4370	4028	
C(31)	254	62	36	38	115	107	385	

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Table S12: Sums of EEQ_{chem} based on the REPs obtained with HeLa-9903. For the left part of the table (>LOQ) only analytical data above the LOQ was included, while for the right part data below LOD or LOQ was substituted by LOD/2 and LOQ/2.

	HeLa-9903 EEQ in pg/L	>LOQ			<LOD = LOD/2 <LOQ = LOQ/2			
		Sum EEQ_{chem} in pg/L			Sum EEQ_{chem} in pg/L			
		Lab 1	Lab 2	Lab 3	Lab 1	Lab 2	Lab 3	
Surface water samples	N(15)	1039	975	497	308	1051	538	367
	K(18)	659	507	383	305	507	383	305
	M(28)	226	334	274	300	381	357	359
	O(3)	487	852	817	80	852	817	507
	F(30)	28	8	6	7	171	38	85
	P(7)	347	664	312	94	664	430	403
	L(24)	509	298	22	222	416	200	281
	H(25)	97	15	9	12	155	102	90
	J(10)	89	19	73	83	61	87	89
	E(27)	245	91	91	104	110	116	126
	G(32)	100	9	6	77	56	55	83
	I(8)	137	14	95	13	209	118	50
	A(11)	16	2	0	1	64	21	27
	B(6)	16	2	0	1	42	23	15
	C(1)	17	4	2	3	84	37	31
D(22)	35	4	2	3	40	26	31	
Waste water samples	E(17)	86	148	6	5	211	74	214
	A(26)	31	2	0	1	50	23	110
	L(13)	1205	1337	710	110	1455	775	669
	K(9)	692	107	120	94	759	258	934
	F(21)	298	10	7	9	160	68	218
	H(5)	273	50	37	38	669	514	1128
	D(4)	90	5	3	4	285	154	563
	O(12)	1695	756	777	650	974	1072	945
	J(16)	888	87	367	66	869	502	611
	P(2)	734	1129	860	860	1176	901	919
	Q(20)	6442	8403	7583	6280	8403	7583	6280
	M(23)	24144	7416	10075	11456	8016	10075	11456
	G(14)	75	17	16	16	484	222	561
	I(19)	553	59	63	50	576	186	259
	B(29)	118	2	2	2	37	62	111
N(33)	2407	535	740	590	830	887	885	
C(31)	41	4	3	3	65	81	448	

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Table S13: Sums of EEQ_{chem} based on the REPs obtained with ER-GeneBLazer. For the left part of the table (>LOQ) only analytical data above the LOQ was included, while for the right part data below LOD or LOQ was substituted by LOD/2 and LOQ/2.

	ER-GeneBLazer EEQ in pg/L	>LOQ			<LOD = LOD/2 <LOQ = LOQ/2			
		Sum EEQ _{chem} in pg/L			Sum EEQ _{chem} in pg/L			
		Lab 1	Lab 2	Lab 3	Lab 1	Lab 2	Lab 3	
Surface water samples	N(15)	876	1312	696	572	1421	754	656
	K(18)	342	679	528	407	679	528	407
	M(28)	487	535	407	480	602	524	564
	O(3)	959	1397	1243	320	1397	1243	821
	F(30)	76	32	25	29	244	65	114
	P(7)	835	1141	603	376	1141	770	710
	L(24)	407	417	90	318	584	302	402
	H(25)	111	59	37	48	209	152	133
	J(10)	92	75	115	121	119	135	130
	E(27)	94	140	136	155	159	161	177
	G(32)	137	36	23	99	89	76	107
	I(8)	230	57	136	50	257	170	95
	A(11)	39	7	0	4	73	27	32
Waste water samples	B(6)	22	7	0	6	50	30	21
	C(1)	32	15	8	10	98	47	41
	D(22)	41	15	9	10	53	40	41
	E(17)	111	230	23	20	293	106	254
	A(26)	25	8	0	4	63	30	138
	L(13)	1166	1819	1189	440	1986	1281	1024
	K(9)	501	428	482	376	1138	638	1461
	F(21)	254	41	29	37	225	105	270
	H(5)	452	200	148	152	851	683	1487
	D(4)	108	21	13	17	324	193	600
	O(12)	1212	1641	1695	1310	1950	2113	1728
	J(16)	813	348	803	264	1302	938	932
	P(2)	1360	1998	1820	1580	2064	1878	1664
	Q(20)	6317	12636	11371	9320	12636	11371	9320
	M(23)	11892	10986	13872	16494	11586	13872	16494
G(14)	567	68	65	65	608	342	732	
I(19)	560	236	252	200	869	397	434	
B(29)	74	9	7	8	48	82	141	
N(33)	1369	1171	1524	1310	1589	1733	1728	
C(31)	143	17	10	10	87	98	578	

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Table S14: Sums of EEQ_{chem} based on the REPs obtained with pYES. For the left part of the table (>LOQ) only analytical data above the LOQ was included, while for the right part data below LOD or LOQ was substituted by LOD/2 and LOQ/2.

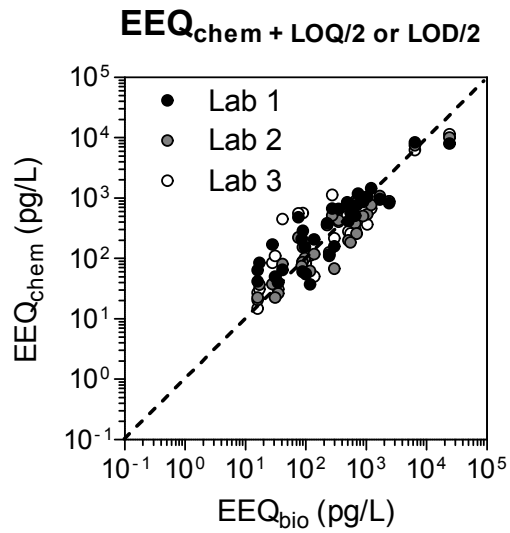
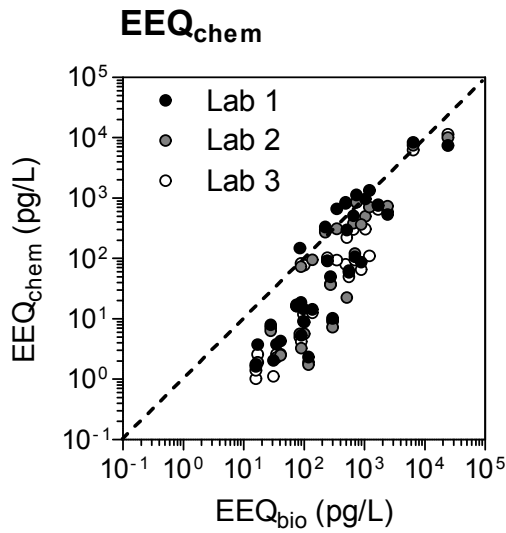
	pYES EEQ in pg/L	>LOQ			<LOD = LOD/2 <LOQ = LOQ/2			
		Sum EEQ_{chem} in pg/L			Sum EEQ_{chem} in pg/L			
		Lab 1	Lab 2	Lab 3	Lab 1	Lab 2	Lab 3	
Surface water samples	N(15)	2900	1481	795	704	1546	830	754
	K(18)	600	595	454	376	595	454	376
	M(28)	1300	636	473	570	676	543	620
	O(3)	5430	1345	1245	440	1345	1245	840
	F(30)	160	44	35	40	189	63	115
	P(7)	3600	1288	748	517	1288	848	817
	L(24)	2060	476	124	366	576	289	416
	H(25)	440	81	51	66	218	136	141
	J(10)	280	103	136	140	144	148	145
	E(27)	190	97	90	103	117	115	125
	G(32)	270	49	31	110	93	79	115
	I(8)	410	79	157	69	271	177	104
	A(11)	130	9	0	6	71	21	31
	B(6)	90	10	0	8	49	23	20
	C(1)	370	20	11	14	100	44	42
Waste water samples	D(22)	390	21	13	14	56	34	42
	E(17)	690	162	31	28	225	95	228
	A(26)	100	11	0	6	56	23	106
	L(13)	1700	2059	1428	605	2159	1483	1155
	K(9)	340	589	662	517	1219	792	1267
	F(21)	160	56	40	51	193	95	251
	H(5)	290	275	203	209	882	658	1209
	D(4)	410	29	18	23	301	158	573
	O(12)	2400	2083	2154	1640	2268	2404	1890
	J(16)	3400	479	822	363	1198	957	863
	P(2)	2800	2432	2300	1940	2472	2335	1990
	Q(20)	9700	9386	8544	7180	9386	8544	7180
	M(23)	12000	7167	9345	10412	7767	9345	10412
	G(14)	970	94	90	89	533	270	589
	I(19)	1200	325	347	275	800	462	475
B(29)	100	13	10	11	46	65	111	
N(33)	2400	1490	1917	1670	1740	2042	1920	
C(31)	65	24	14	14	81	89	414	

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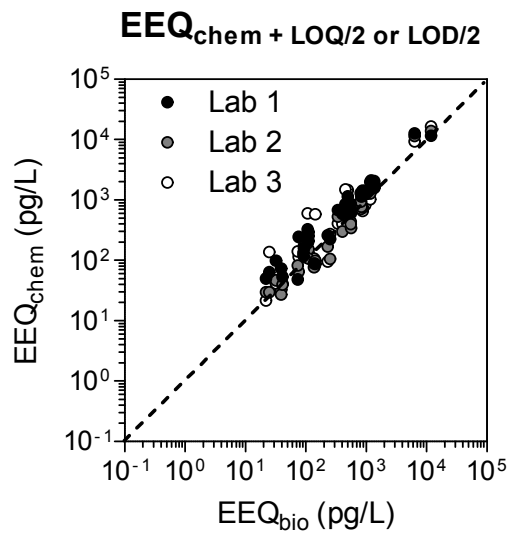
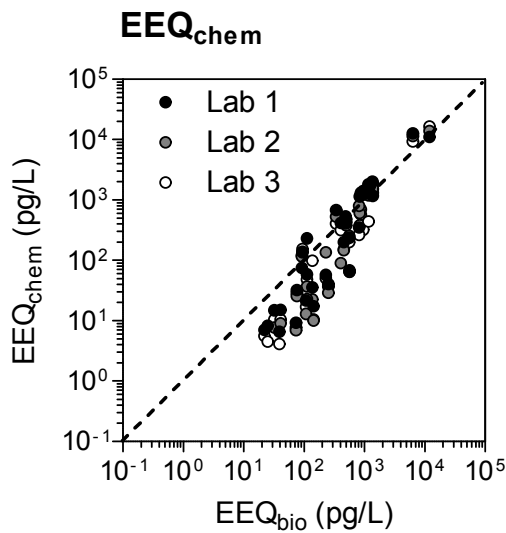
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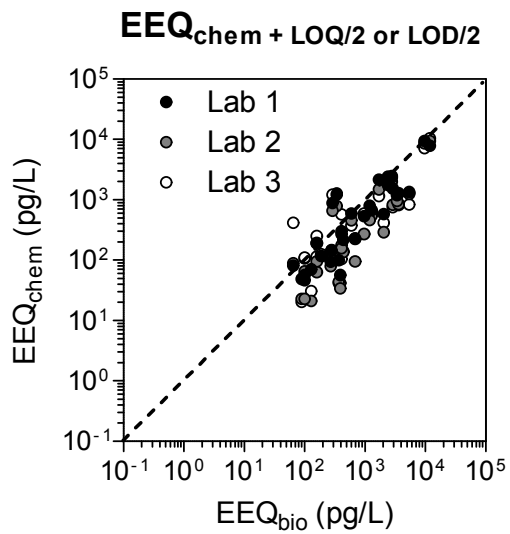
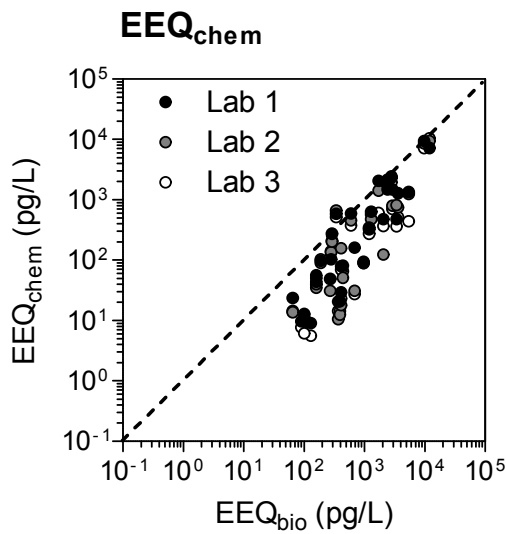
HeLa



ER-GeneBLazer



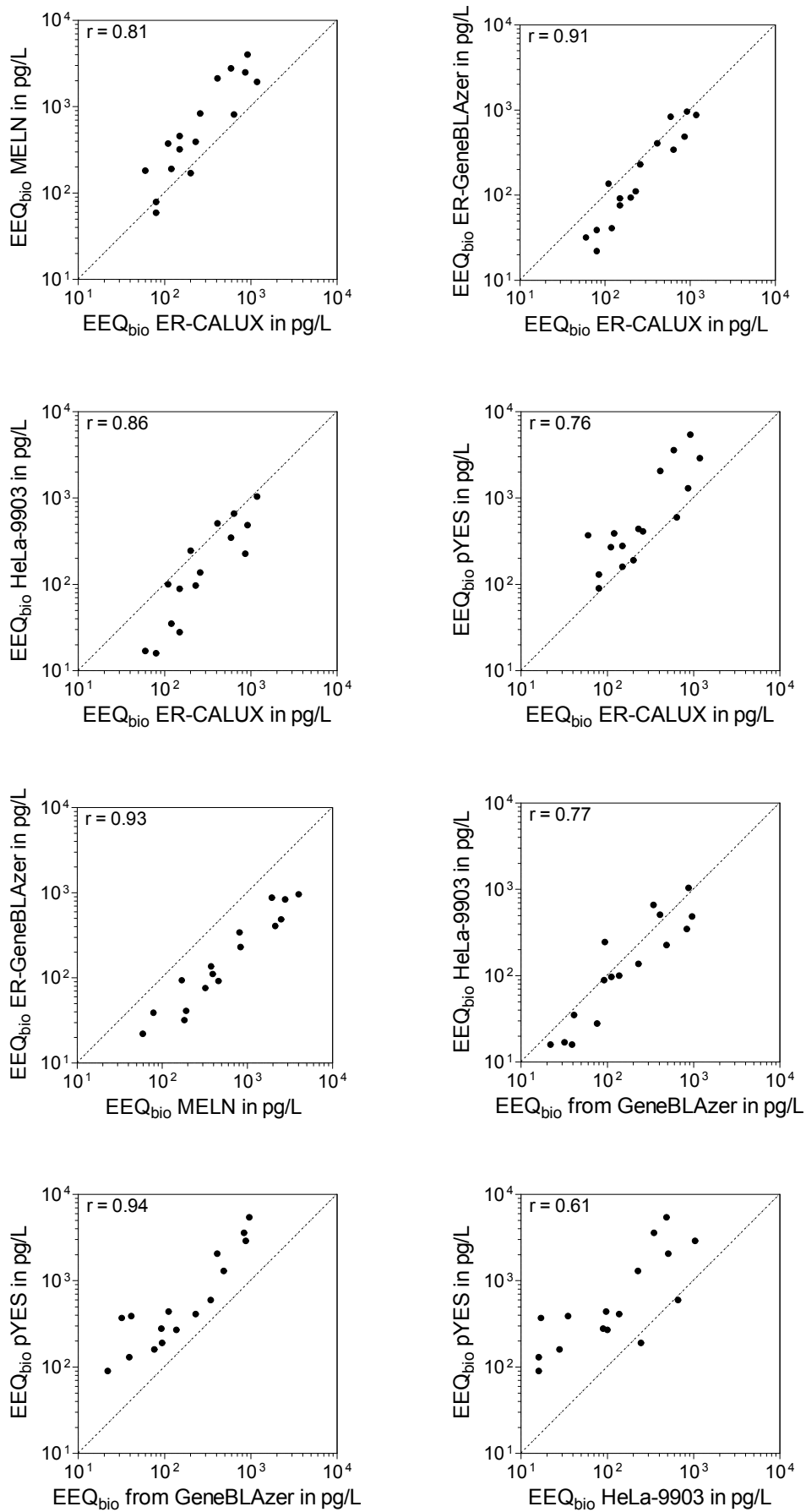
pYES



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Figure S1: Comparison of EEQ_{chem} and EEQ_{bio}. Graphs on the left show the EEQ_{chem} derived from values >LOQ, while the graphs on the right show the EEQ_{chem+LOD/2 or LOQ/2} calculated by including LOD/2 and LOQ/2. The dashed line indicates perfect agreement of EEQ_{chem} with EEQ_{bio}.



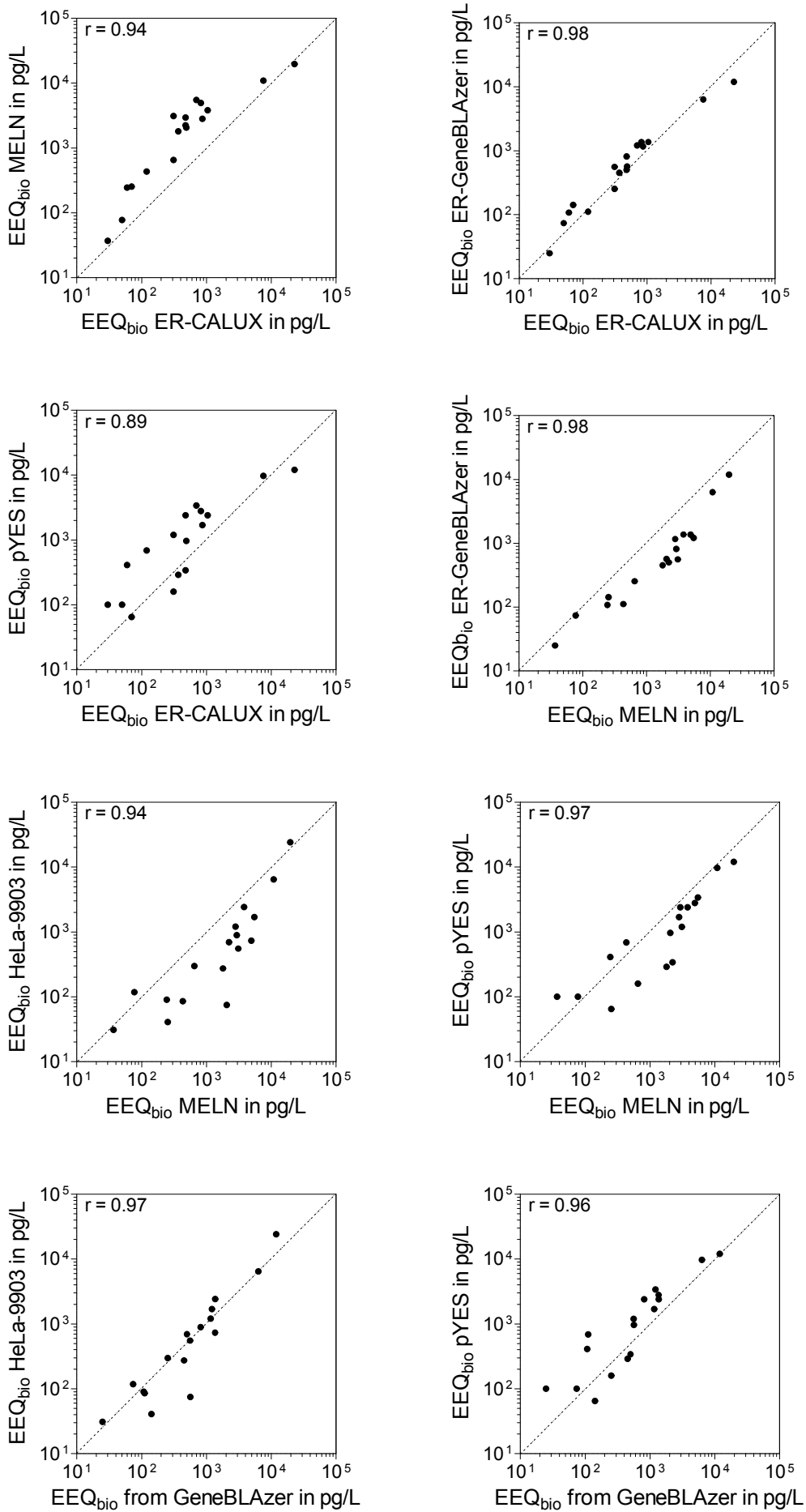
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Figure S2: Comparison of effect-based methods by correlating the measured EEQs for surface water samples. The correlation analysis was based on the method described in section 2.9 with a fixed slope of 1. The dashed line indicates perfect agreement of the compared effect-based methods.



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Figure S3: Comparison of effect-based methods by correlating the measured EEQs for waste water samples. The correlation analysis was based on the method described in section 2.9. The dashed line indicates perfect agreement of the compared effect-based methods.

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