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In vitro effects-based method and water quality screening model for use in pre- and post-distribution treated waters

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Abstract

Recent urban public water supply contamination events emphasize the importance of screening treated drinking water quality after distribution. *In vitro* bioassays, when run concurrently with analytical chemistry methods, are effective tools to evaluating the efficacy of water treatment processes and water quality. We tested 49 water samples representing the Chicago Department of Water Management service areas for estrogen, (anti)androgen, glucocorticoid receptor-activating contaminants and cytotoxicity. We present a tiered screening approach suitable to samples with anticipated low-level activity and initially tested all extracts for statistically identifiable endocrine activity; performing a secondary dilution-response analysis to determine sample EC_{50} and

Competing Interests

The Authors have no competing interests to declare.

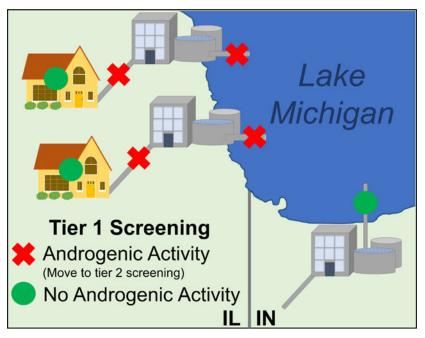
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biological equivalency values (BioEq). Estrogenic activity was detected in untreated Lake Michigan intake water samples using mammalian (5/49; median: 0.21 ng E2Eq/L) and yeast cell (5/49; 1.78 ng E2Eq/L) bioassays. A highly sensitive (anti)androgenic activity bioassay was applied for the first time to water quality screening and androgenic activity was detected in untreated intake and treated pre-distribution samples (4/49; 0.93 ng DHTEq/L). No activity was identified above method detection limits in the yeast androgenic, mammalian anti-androgenic, and both glucocorticoid bioassays. Known estrogen receptor agonists were detected using HPLC/MS-MS (estrone: 0.72-1.4 ng/L; 17α -estradiol: 1.3-1.5 ng/L; 17β -estradiol: 1.4 ng/L; equol: 8.8 ng/L), however occurrence did not correlate with estrogenic bioassay results. Many studies have applied bioassays to water quality monitoring using only relatively small samples sets often collected from surface and/or wastewater effluent. However, to realistically adapt these tools to treated water quality monitoring, water quality managers must have the capacity to screen potentially hundreds of samples in short timeframes. Therefore, we provided a tiered screening model that increased sample screening speed, without sacrificing statistical stringency, and detected estrogenic and androgenic activity only in pre-distribution Chicago area samples.

Graphical Abstract



Keywords

Effects-based method; estrogen; androgen; tapwater

1. Introduction

Despite great strides in physical and chemical drinking water treatment over the last century (Cutler and Miller, 2005; Schoenen, 2002), the growing global human population has depleted many freshwater sources and increased reliance on effective water treatment

processes (UNWAPP, 2017). In vitro effects-based methods (bioassays), when run concurrently with analytical chemistry methods, have gained popularity over the last decade as effective tools to evaluate the efficacy of water treatment processes (Conley et al., 2017b; Escher et al., 2014; Jia et al., 2015; Jia et al., 2016; Medlock Kakaley et al., 2020; Plewa and Wagner, 2015; Shi et al., 2018; Snyder and Leusch, 2018; Zhen et al., 2018). Much of the development and application of bioassays to water quality has focused on surface (Blackwell et al., 2019; Blackwell et al., 2017; Conley et al., 2017a; Jeong et al., 2012), waste (Dong et al., 2017; Dong et al., 2019; Leusch et al., 2018) and recycled water (Escher et al., 2014; Jia et al., 2015). In these water sample types, the detected biological activity can be relatively high. However, effects-based methods can also make exceptional tools for screening water samples that contain very low-level concentrations of individual contaminants (e.g. treated drinking water). This versatility results from bioassays' inherent ability to detect cumulative biological activity from the sum of all present contaminants. Typically, biological activity in an environmental sample, reported in biological equivalency values (BioEq), is calculated by fitting a sigmoidal concentration-response curve to cell (or organism) responses from treatments of serial dilutions of a sample and determining an EC₅₀ value and sample relative potency (compared to method reference compound). However, biological activity detected in treated drinking water can fall below an EC_{50} value, despite sample enrichment. Therefore alternative approaches to the traditional sigmoidal concentration-response curves to determining sample activity are necessary (Escher et al., 2018b).

In a preceding pilot study we sought to fill data gaps in chemical occurrences in point-of-use drinking water (tapwater, TW) using both bioassays and analytical chemical methods (Bradley et al., 2018). TW samples collected from locations across the United States suggested human exposure to mixtures of trace level organic and inorganic compounds that are not required to be monitored in TW. Herein we seek to identify potential sources, movement and transformation of biologically-active contaminants in treated drinking water during distribution. In a multi-agency collaborative study of pre- and post-distribution treated TW representing the Chicago Department of Water Management service areas, an extensive targeted-chemical toolbox (540 organic and 35 inorganic analytes) was employed and no detected chemical concentrations exceeded U.S. Environmental Protection Agency (USEPA) Maximum Contaminant Levels in any treated TW sample (Bradley et al., 2020). However, multiple exceedances of health-based advisories (e.g., maximum contaminant level goals; MCLG) in untreated and pre-distribution water samples, together with the recognized order of magnitude analytical underestimation in the TW exposure (350,000+ commercial compounds (Bradley et al., 2017) and unquantifiable transformation products (Dobson, 2004; Vasquez et al., 2014)), raised concerns for potential biological effects and therefore supported a non-targeted effects-based screening.

Bioassays that detect steroid hormone-active compounds have previously compensated for gaps in targeted analyte coverage (Bradley et al., 2017; Conley et al., 2017a; Medlock Kakaley et al., 2020). Therefore, we applied a suite of bioassays focused on steroid hormone signaling pathways to Chicago TW extracts to screen pre- and post-distribution water filtration plant (WFP)-treated water. Included in the suite was a highly sensitive method for detecting (anti)androgenic activity (Hartig et al., 2007) never before applied to water quality screening. Further, we provide a tiered screening model for sample analysis applicable to

routine water quality monitoring that expedites drinking water sample extract evaluation while maintaining statistical stringency.

2. Methods

2.1 Sample Collection

A total of 49 samples of raw-water intake from Lake Michigan (WFP-UT), WFP-treated/ pre-distribution water (WFP-TW), and post-distribution tapwater (TW) from private residences (cold water samples without pre-cleaning, screen removal, or flushing; exact locations of private residences have been anonymized) were taken from Chicago, Illinois and East Chicago, Indiana between July and December 2017 using procedures that were previously described (Supporting Information) (Romanok et al., 2018). Sampling sites (38 total) were selected based on community volunteers and represented a broad geographical coverage of the City of Chicago Department of Water Management water filtration plants' (WFP) service areas (Figure 1; Chicago North, CN: 16 sites; Chicago South, CS: 18 sites; East Chicago, EC: 4 sites). Only pre-distribution samples were screened from the City of East Chicago Utilities Department service area and were included for comparison to Chicago, IL pre-distribution samples. Field blanks were collected as part of the quality assurance and quality control protocols at three sites (CS-WFP-TW, EC-WFP-UT1 and EC-WFP-TW1). Sample aliquots from each sampling site were allocated for multiple analysis types *e.g.*, analytical chemistry, endocrine bioassays and cytotoxicity bioassay.

2.2 Analytical Chemistry

Water samples were analyzed by U.S. Geological Survey (USGS) using 14 organic (540 unique analytes) and 7 inorganic (37 analytes) analytical methods (Supporting Information) and a full report of detected analytes were reported previously (Bradley et al., 2020). Of the 550 analytes, several known estrogen, androgen, and glucocorticoid receptor agonists were quantified using HPLC/MS-MS (Yost et al., 2013; Yost et al., 2014), as described previously (Bradley et al., 2020; Romanok et al., 2018), and are reported and discussed herein.

2.3 Endocrine Bioassays

A 4 L sample from each site was extracted into 400 μL methanol using the solid phase extraction methods described previously (Supporting Information) (Romanok et al., 2018). Two 100 μL extract aliquots were shipped overnight on ice to both U.S. Environmental Protection Agency, Research Triangle Park, NC (mammalian bioassay) and USGS Leetown Science Center, Kearneysville, WV (yeast bioassay) for *in vitro* screening of steroid hormone activity.

2.3.1 Mammalian Bioassays—The T47D-KBluc cell line (American Type Cell Culture, ATCC, Manassas, VA; #CRL-2865) (Wilson et al., 2004; Wilson et al., 2002) has been applied to chemical and environmental sample testing for estrogenic activity previously (Conley et al., 2017a; Conley et al., 2017b). Cell culture maintenance and sample screening were conducted as previously described (Wilson et al., 2004), with exceptions (Supporting Information) (Medlock Kakaley et al., 2020). Briefly, treatments included methanol vehicle control, serially diluted water samples, 17-beta estradiol (E2; CAS#: 50-28-2; Sigma)

standard curve or 1 μ M ICI 182,780 (antagonist control; Tocris Bioscience) in competition with E2 in Dulbecco's Modified Eagle's Medium (DMEM) with 10% dextran-coated charcoal treated fetal bovine serum (DCC-FBS).

The CV1 cell line (ATCC; CCL-70), which is naturally devoid of glucocorticoid and androgen receptor (AR) was cultured and transduced with the human glucocorticoid receptor (GR) to test for glucocorticoid activity in water samples as previously described (Hartig et al., 2002; Medlock Kakaley et al., 2018). Approximately 5×10^6 cells were transduced with the human GR (Ad/GR4) and MMTV-luciferase (Ad/Luc7), with a multiplicity of infection (MOI) of 1 and 50, respectively. Cells were plated at a rate of 22,000-33,000 cells/well, and exposed to cell media with either the methanol vehicle control, serially-diluted water extract samples, dexamethasone standard curve (Dex; CAS#: 50-02-2; Sigma) or 300 nM mifepristone (antagonist control; CAS: 84371-65-3; Sigma) in competition with Dex for 24 hrs. CV1-cells were frozen at -80° C with lysis buffer and thawed at room temperature before analyzing with luminometer.

Using a similar CV1 transduction system, water extract samples were also tested for (anti)androgenic activity using a CV1-chAR bioassay (Hartig et al., 2007). Culturing, transduction and exposure methods are identical to the CV1-hGR bioassay with the following exceptions. Cells were transduced with chimpanzee androgen receptor (chAR) and the MMTV-luc gene with MOIs of 1 and 50, respectively. In the AR agonist assay cells were exposed to media with either methanol vehicle control, 4,5α-Dihydrotestosterone (DHT; CAS#: 521-18-6; Sigma) standard curve (0, 3, 10, 30, 100, 300, 1000, 3000 pM), hydroxyflutamide (OHF; antagonist control; CAS#: 52806-53-8; Sigma) in competition with DHT, or serially diluted extract samples. In the AR antagonist assay, standards and samples were diluted with a concentration of AR agonist that produces approximately 80% of the maximal response (100 pM DHT) in RMPI. Cells were exposed to media with either methanol vehicle control (0, 1, 3, 10, 30, 100, 300, 1000, nM), or serially diluted water extract samples. Antagonist activity was quantified as OHF equivalents (OHFEq).

All cells for T47D-KBluc and CV1-hGR/chAR screening assays were plated in 96-well luminometer plates. Each standard, control, or sample was run in quadruplicate, and each sample screen was at least duplicated *i.e.* different cell passage number. After 24 hr *in vitro* exposure, cells were visually scored for cytotoxicity (scale of 1–5) and any wells with cells exhibiting cytotoxic effects were excluded from subsequent analysis (Bhatia and Yetter, 2008; Conley et al., 2017a). Cells were washed, lysed and immediately following injection of luciferase reaction buffer and *Firefly* D-luciferin substrate, luminescence was quantified by measuring luminescence every 0.2 sec for 5 sec using a FLUOstar luminometer (BMG Labtech, Cary, NC) (Medlock Kakaley et al., 2020). The method detection limits (MDL) for mammalian bioassays were determined using the equation described below and were as follows; 0.0044 ng E2Eq/L, 0.3 ng DHTEq/L, 67.10 ng OHFEq/L, and 1.06 ng DexEq/L for the T47KBluc, CV1-chAR, CV1-chAR in antagonist mode, and CV1-hGR bioassays, respectively.

2.3.2 Yeast Bioassays—To add depth to our *in vitro* analysis of endocrine activity in the water extract samples (Blackwell et al., 2019; Könemann et al., 2018; Medlock Kakaley et al., 2020), the bioluminescent yeast estrogen screen (BLYES; 490 BioTech, Knoxville, TN) also was used to assess estrogenicity (estrogen receptor-alpha only) in the water extract samples as previously described (Conley et al., 2017a; Sanseverino et al., 2005), with modifications (Ciparis et al., 2012). Extracts were also screened for androgenicity and glucocorticoid activity using yeast strains DSY-1555 and MCY-105, respectively, as previously described (Kassotis et al., 2016).

Complete method details are provided in the Supporting Information. Briefly, each assay plate included either a 17 β -estradiol, DHT, or hydrocortisone standard curve, vehicle controls, and sample extracts. All treatments were added as 10 µL volumes to wells of solid white 96-well plates and methanol was allowed to evaporate to dryness prior to the addition of media and yeast reporter. Final sample dilutions were 1:20. The reference hormone/ environmental contaminant-induced chemiluminescent signal was measured using a SpectraMax M4 microplate reader (Molecular Devices, San Jose, CA) in luminescent mode (100 ms integration time). Endocrine activity within each sample, relative to respective standard curves, was determined using a four-parameter logistic regression (SoftMax Pro 6.6.6, Molecular Devices) and corrected for sample enrichment. Method detection limits for the BLYES, DSY-1555, and DSY-105 bioassays were 0.10 ng/L 17 β -estradiol, 1.3 ng/L dihydroxytestosterone, 1.1 ng/L hydrocortisone, respectively.

2.4 Cytotoxicity

The Chinese hamster ovarian (CHO) cell cytotoxicity assay measures the reduction in cell density (surrogate for viability) as a function of the concentration of the test (Supporting Information) (Plewa et al., 2002; Plewa et al., 2010; Wagner and Plewa, 2017). Water sample aliquots (20 L) were extracted into DMSO as described previously (Daiber et al., 2016). DMSO concentrate was diluted in F12+ 5% fetal bovine serum cell culture medium (F12-FBS). In a 96-well flat-bottomed microplate, a blank control (no cells), negative control (F12-FBS plus cells) and experimental samples were analyzed concurrently. Initially, all negative and experimental wells contained 3×10^3 CHO cells. A covered microplate was rocked at 37° C for 10 min, ensuring even cell distribution, and incubated for 72 hr at 37° C under 5% CO₂. Cells fixed in methanol were stained with crystal violet and washed with DMSO: methanol (3:1 v/v)l. Fixed cells were incubated at room temperature for 10 min and subsequently analyzed at 595 nm with a SpectraMaxTM microplate reader.

Detailed statistical methods for CHO cytotoxicity assay were conducted as described previously and are provided in the Supporting Information (Box et al., 1978; K and M, 2008). Experimental treatments were normalized to percent of negative control treatment (100% survival) and LC_{50} values were converted into mean cytotoxicity index (CTI) values (CTI = $10^3 \times LC_{50}^{-1}$) (Box et al., 1978; K and M, 2008). One-way analysis of variance (ANOVA) tests were conducted to determine the lowest concentration factor that induced cytotoxicity compared to negative control (*P* 0.05).

2.5 Screening Model and Data Analyses

Steroid hormone activity is typically low, and often undetectable, in treated point-of-use TW, therefore we used a tiered screening process (Figure 2). In tier one, the presence of estrogenic/(anti)androgenic/glucocorticoid activity in each sample extract was determined. All 49 water sample extracts were screened at 10x and 5x final assay sample enrichment factors. Mean fold change above concurrent control of treatment replicates were normalized to percent maximal assay activity (based on assay-specific hormone standard saturation levels). Sample extracts with significant within plate increases/decreases in percent maximal activity of respective agonist/antagonist reference compound were identified with the General Linear Model Procedure using the Least Square Means Statement and Dunnett's multiple comparison procedure.

In tier two, only active samples from tier one were screened using a series of 2-fold serial dilutions (1:1000, 1:2000, 1:4000, 1:8000, etc). Biological equivalency values (BioEq; reported in ng Reference/L) were calculated (Conley et al., 2017a; Medlock Kakaley et al., 2020) and graphically represented for each positive sample using the equation,

 $BioEq = \frac{(Reference EC_{50})}{(Sample EC_{50})(Sample EF)}$

where the sample EC_{50} was a unitless bioassay sample dilution and was extrapolated using mean observed transcriptional activation values. The enrichment factor (EF) = 10,000 (prior to assay dilution) due to 4,000-fold concentration of sample during extraction (4 L water sample) and dried extract resuspension 0.4 mL methanol. Reference EC_{50} values are provided in Table S1.

The minimum detectable concentration (MDC) for hormone activity was determined as described previously (Conley et al., 2017b), and the method detection limit (MDL), where

$$MDL = \frac{MDC}{Final \ Sample \ EF}$$

and Final Sample EF = 10 (10,000x Enrichment Factor and 1,000x assay dilution), was used in the final step of the tiered screening process. Samples were identified as a positive hit for activity depending on whether the BioEq value was above or below the respective assay MDL.

2.6 Statistical Analysis

Statistical analyses were performed using SAS statistical software (Cary, NC USA) and graphs were generated using GraphPad Prism version 7.02 for Windows (GraphPad Software, LaJolla CA, USA).

3. Results

We efficiently screened raw WFP intake from Lake Michigan, WFP-treated pre-distribution water and point-of-use tapwater from locations near and around Chicago, IL (Table 1 and

Figure 1), for multiple types of biological activity using a tiered testing model applicable to water quality monitoring (Figure 2). Overall, no treated point-of-use TW sample produced significant responses in endocrine activity (estrogenic, androgenic, anti-androgenic or glucocorticoid) or cytotoxicity. Estrogenic activity and androgenic activity were detected in raw WFP intake samples above method detection limits (Figure 3, 4 and Tables S2, S3, and S4).

Estrogenic activity was detected in untreated WFP intake waters using both the T47D-KBluc and BLYES bioassays (Figure 3A and B). All five extract samples that produced responses above method detection limits for estrogenicity in the T47D-KBluc bioassay (MDL: 0.0044 ng estradiol equivalents($E2Eq_{T47D-KBluc}/L$) also produced positive responses in the BLYES bioassay (MDL: 0.10 ng $E2Eq_{BLYES}/L$). Estrogenic activity ranged from 0.04 to 0.52 ng $E2Eq_{T47D-KBluc}/L$ (Mean: 0.21 ± 0.14 SD) and 0.19 to 1.93 ng $E2Eq_{BLYES}/L$ (Mean: 1.24 ± 0.84). A linear regression analysis was conducted to compare the magnitude of estrogenic activity (ng E2Eq/L) in corresponding samples with positive detections. There was a distinct positive relationship between the results of the two bioassays (Figure 4; R² = 0.82), although ng $E2Eq_{T47D-KBluc}/L$ were consistently an order of magnitude lower than the corresponding ng $E2Eq_{BLYES}/L$.

In Bradley *et al.* (Bradley et al., 2020), many known estrogen receptor agonists were targeted environmental analytes and four were detected above established laboratory method reporting limits (Tables S5 and S6). Estrone (MRL: 0.5 ng/L) was detected in East (EC-WFP-TW1, Sept.) and South Chicago WFP-treated pre-distribution water (CS-WFP-TW, Nov.), one East Chicago WFP intake (EC-WFP-UT2, Sept.), and one South Chicago point-of-use samples extract (CS-TW-15). Both 17α-estradiol (MRL: 0.5 ng/L) and 17β-estradiol (MRL: 0.5 ng/L) were detected in one South Chicago pre-distribution water sample (CS-WFP-TW1, Nov.), and Equol (MRL: 0.5 ng/L), a non-steroidal phytoestrogen, was detected in a South Chicago point-of-use samples extract (CS-TW-14) (Table S6).

We used the inaugural application of an androgenic activity bioassay to illustrate a model for water quality monitoring with endocrine bioassays (Figure 2). Androgenic activity (significantly higher than vehicle control) was detected in two untreated intake and two treated pre-distribution water samples from North and South Chicago using the CV1-chAR bioassay (Figure 5). All four samples with quantifiable androgenic activity contained levels above the bioassay MDL (0.30 ng dihydroxytestosterone equivalents(DHTEq)_{CV1-chAR}/L). Activity ranged from 0.77 to 1.17 ng DHTEq_{CV1-chAR}/L (Mean: 0.94 \pm 0.16), yet no sample produced androgenic activity above the yeast assay MDL for androgen activity (1.3 ng DHTEq_{DSY-1555}/L). Further, of the targeted androgen receptor agonist analytes assessed, none were reported in the water extract samples above method detection limits by Bradley *et al.* (Table S5 and S6) (Bradley et al., 2020).

The CV1-chAR bioassay can also be used in an antagonist mode, to evaluate the presence of compounds that might interfere with, or reduce, androgen receptor activity. No sample exhibited significant antagonist activity compared to the 100% maximal activity of 100 pM dihydrotestosterone positive control. The MDL for the CV1-chAR antagonist bioassay, 67.10 ng OHFEq/L, was calculated using the previously reported methods (Conley et al.,

2017b), with the exception of using the lower 95% prediction interval around the 100% maximal 100 pM DHT response.

The CV1 cell line transduced with the human glucocorticoid receptor (CV1-hGR) can detect samples with glucocorticoid activity, but no samples contained significant activity compared to vehicle control (MDL: 1.06 ng $DexEq_{CV1-hGR}/L$). Similarly, the chemiluminescent yeast bioassay for glucocorticoid activity (MDL: 1.1 ng $DexEq_{MCY-105}/L$) did not produce any positive detections for glucocorticoid activity. Of the targeted glucocorticoid receptor agonist analytes assessed, none were reported in the water extract samples above method detection limits (Table S5 and S6) (Bradley et al., 2020).

Notably, the only sample with a significant increase in cytotoxicity from the CHO bioassay also produced measurable levels of estradiol equivalents in both estrogenic activity bioassays (EC-WF-UT1-Sept). Cytotoxicity (using cytopathogenic effect) was not apparent in either of the estrogenic bioassays (no reduction in cell viability, no reduction in bioluminescent signal) which may have resulted from variations in contaminant mixtures produced by the respective extraction methods.

4. Discussion

Drinking water treatment (Murray et al., 2019) and methodologies for water quality screening (Martin et al., 2007) continue to be developed and refined. However, isolated public water supply contamination events emphasize the value of specifically assessing treated drinking water after distribution from public water filtration plants (WFP). Aging water distribution infrastructure has contributed to altered water quality post-treatment in residential taps in Flint (2015) and Detroit, Michigan (2018) (Hyde, 2015). So in 2016, we conducted a pilot study to screen residential and workplace point-of-use tapwater samples across the US to investigate potential exposures from public water supplies (Bradley et al., 2018). The current wastewater and water infrastructure study described herein was designed to limit some of the variables present in the previous study. Only one source water, Lake Michigan, was used to determine the occurrence as well as the potential changes in chemical content of water during treatment and distribution. Further, the scope of water sampling was limited to a few water filtration plants and to residential sites served by those treatment facilities.

Despite the extensive list of targeted analytes included in the current study (complete list reported elsewhere (Bradley et al., 2020)), we (Bradley et al., 2017; Conley et al., 2017a; Medlock Kakaley et al., 2020) and others (Blackwell et al., 2019; Hashmi et al., 2020; Könemann et al., 2018) have previously shown that bioassays, which produce a cumulative value for all compounds affecting a single biological endpoint, may compensate for gaps in the targeted analytical chemical coverage. Könemann *et al.* reported estrogen activity detections two orders of magnitude lower compared to concentrations of targeted estrogens in European surface water sample analyzed using both methods (Könemann et al., 2018). However, bioassays are frequently applied to water quality monitoring using only relatively small samples sets often collected from surface and/or wastewater effluent (Escher et al., 2014; Neale et al., 2015; van der Oost et al., 2017).

To realistically adapt bioanalytical tools to treated water quality monitoring (especially in the absence of robotic sample and reagent dispensing), programs must have the capacity to screen potentially hundreds of samples in short timeframes, and subsequently obtain results in real time. Considering the number of samples tested in the current study and that we expected little to no detectable biological activity in many of the samples (30 tapwater samples), a tiered screening approach was preferable over the standard dilution-response testing. Locations within the US (Denison et al., 2020) and Europe (Brack et al., 2015) are already applying bioassays to water quality monitoring and the State of California's guidance document for developing standard operating procedures for applying bioassays to water quality monitoring recommends a similar tiered approach to screening (Denison et al., 2020).

The tiered screening model we used is illustrated in Figure 2 using our inaugural application of the CV1-chAR bioassay as an example. To determine if androgenic activity was present, we screened all 49 samples using two sample dilutions and found only 4 samples with activity compared to concurrent plate vehicle control using the General Linear Model followed by Dunnet's multiple comparison procedure. A secondary *in vitro* screen was conducted on the active samples only with a series of dilutions (concentration-response) to determine how much activity was in each sample. Using the DHT standard curves run concurrently with each sample, we determined all measurable activity in androgen active samples fell above the MDL. Although no human health EBTs have been determined for the CV1-chAR bioassay, all the samples fall below the previously reported EBTs for DHTEq established by Brand *et al.* for the AR-CALUX bioassay (11 ng DHTEq/L) in drinking water (Brand et al., 2013).

The addition of the bioassay screening components to the overall study design undoubtedly increased the spectrum of chemical detection. For example, in four of the five samples that produced estrogenic activity above MDL (CN-WFP-UT-July, CN-WFP-UT-Nov, EC-WFP-UT1-Sept, and EC-WFP-UT2-Aug; Figure 3A and B), none of the targeted estrogen receptor ligands were detected above analytical method detection limits. This could result due to the presence of 1) ER ligands that were not included in targeted analysis, 2) analytes included in the targeted analysis that have not been previously identified as an ER activating compound, or 3) a mixture of known and/or unidentified targeted ER ligands present at individual concentrations below analytical detection limits. Effects-directed analysis (EDA), an emerging experimental approach, could mitigate these recurring issues in scenarios one and two (Dong et al., 2020; Dusza et al., 2019; Hashmi et al., 2020; Zwart et al., 2020). An alternative explanation for the discrepancies in chemical and biological detection exists in the creation of sample aliquots and conducting separate extraction methods. While separate extraction methods are necessary to prevent contamination that interferes with biological analysis, separate methods create inherent bias *i.e.*, extraction methods are optimized only to retain compounds that will be subsequently analyzed.

Estrogenic activity was detected in two of the four WFP intakes from Lake Michigan (T47D-KBluc), and was comparable to estrogenic activity previously detected upstream (0.14 ng E2Eq_{T47D-KBluc}/L) and downstream of wastewater treatment plants (0.16 ng E2Eq_{T47D-KBluc}/L) (Medlock Kakaley et al., 2020), and in European surface waters (0.16 ng

E2Eq_{HeLa}/L) (Könemann et al., 2018). We have previously shown differences in resulting estradiol equivalents from the two estrogenic activity bioassays (Conley et al., 2017a), which may be due in part to T47D-KBluc measuring ERa and ER β , and the BLYES bioassay measuring ERa activation only. Others assessing estrogenic activity with both human and yeast cell lines have also reported order of magnitude difference in maximum reported activity (Könemann et al., 2018). We have previously detected estrogenic activity in unfinished drinking water (pre-distribution) and treated drinking water above MDL at 0.03 (Medlock Kakaley et al., 2020) and 0.078 ng E2Eq_{T47D-KBluc}/L (Conley et al., 2017b), respectively. Estrogenic activity bioassays other than the T47D-KBluc (Shi et al., 2018), but treated water from the Chicago area WFPs did not contain measurable levels of estrogenic activity.

Previous studies that have included bioanalytical and concurrent analytical chemistry methods have directly compared detected biological activity to the environmental concentrations of estrogens by transforming each environmental concentration to biological equivalency values using bioassay-specific compound potency (Conley et al., 2017a; Jia et al., 2016; Medlock Kakaley et al., 2020; Neale et al., 2015). This type of correlation analysis clarifies whether the targeted and detected environmental compounds capture the entire measured biological activity in each sample. However, only one sample, the September 2017 EC-WFP-UT2 water extract, produced a mean bioactivity of 0.29 ng E2Eq/L (Figure 3a) and contained 0.72 ng/L estrone (Table S6). The estimated T47D-KBluc bioactivity value, based on previously reported T47D-KBluc relative potency value for estrone (1.39) (Conley et al., 2016), was 1.00 ng E2Eq/L for the September 2017 EC-WFP-UT2 sample.

Imperative to the value of detected biological activity in treated tapwater is the exposure concentration at which it is predicted to adversely affect human health, *e.g.* effects-based trigger values (EBTs). Human health EBTs have been conceived using acceptable daily intakes, bioassay-specific relative potency factors, and pharmacokinetic parameters (Escher et al., 2018a). To our knowledge a T47D-KBluc specific-EBT has not been determined using either method. However, Brand *et al.* reported an EBT of 3.8 ng E2Eq/L for drinking water using a similar bioassay, ER-CALUX. Although relative potency factors of ER ligands vary between T47D-KBluc and ER-CALUX (Brand et al., 2013; Conley et al., 2016; Houtman et al., 2009; Sonneveld et al., 2005), the WFP treated water pre- and post- distribution likely do not contain levels of estrogenic compounds that would be predicted to cause adverse effects assuming standard rates of ingestion (WHO, 2000).

All estrogenic activity was detected in the intake water samples, but no environmental EBT has been developed for the T47D-KBluc. Escher *et al.* used environmental quality standards to derive EBT for several estrogen receptor agonist bioassays, which ranged from 0.1 (ER-CALUX) to 1.07 ng E2Eq/L (ISO-LYES) (Escher et al., 2018a). Given the variability in ecological EBTs, a conservative approach may be to use the predicted no effect concentration (PNEC) for long-term exposure to 17α-ethinyl estradiol (EE2; potent environmental ER agonist), 0.2 ng E2Eq/L (corrected for EE2 potency in the T47D-KBluc bioassay), presented by Caldwell *et al* (Caldwell et al., 2012; Conley et al., 2017a). Considering the Caldwell *et al.* PNEC, all samples with estrogenic activity above MDL

contain levels likely to adversely affect aquatic species. A major contributor of the estrogenic activity detected in the Lake Michigan intake water samples is likely effluent from municipal wastewater treatment plants (WWTP) from the densely populated surrounding metropolitan area. WWTPs with standard activated sludge treatment may not be sufficient to remove the all estrogenic activity (Kibambe et al., 2020), however advanced wastewater treatment technologies including ozonation and sand filtration have been shown to completely remove all estrogenic activity from WWTP influent (Gehrmann et al., 2018).

We applied the CV1-chAR bioassay, originally created for screening environmental compounds (Hartig et al., 2002; Hartig et al., 2007), to water quality screening of each water sample extract. The CV1-chAR bioassay contains the chimpanzee androgen receptor (AR) that has 99.7% overall sequence similarity to the human AR ortholog and 100% sequence similarity in the ligand- and DNA-binding domains (Choong et al., 1998). The transcriptional activation bioassay was originally designed with the non-human primate receptor due to patent limitations on the use of the human AR (Hartig et al., 2007). We previously showed that the dihydrotestosterone-mediated transcriptional activation of both AR orthologs were statistically the same (Hartig et al., 2007).

Since the Chicago area water samples were concentrated during extraction and subsequently re-diluted 1000-fold in assay media we were able to seamlessly adapt the tool to environmental sample evaluation with no cytotoxicity issues. Of the 49 screened samples (including field and lab blanks), four produced androgenic activity (median: 0.93 ng DHTEq/L) and were comparable to androgenic activity previously detected in the US (4.7 ng DHTEq/L) (Conley et al., 2017a). The androgenic activity reported by Conley *et al.* was detected in US surface waters that were historically impacted by a variety of anthropogenic sources (*e.g.* industrial, municipal, agricultural). In studies aiming to optimize effects-directed analysis protocols, a limited number of surface water sites in the Netherlands (0.2 ng DHTEq/L) (Zwart et al., 2020), and several surface water sites across Europe (2.7 ng DHTEq/L) (Tousova et al., 2017) also resulted in androgenic activity comparable to activity detected in the Chicago pre-distribution samples.

Despite the 10 androgen receptor agonists included in the HPLC/MS-MS analysis (Table S7), no androgen analytes were detected above MDLs to account for the measured *in vitro* AR activity. Similar are results presented by Tousova *et al.* assessing European surface and wastewaters. AR activity ranged 0.93–2.7 DHTEq_{MDA-KB2}/L, but AR agonists concentrations were negligible leaving the bioactivity entirely unexplained (Tousova *et al.*, 2017). In our previous assessment of US impacted surface waters, up to 96% of the *in vitro* AR activity (MDA-KB2) was accounted for by measured testosterone, androstenedione, androsterone, and dihydrotestosterone environmental concentrations (Conley et al., 2017a).

There are many existing options for screening *in vitro* (anti)androgen activity (*e.g.* MDA-KB2, GeneBLAzer, CALUX, etc.). However, the CV1-chAR bioassay had a superior Z-factor (a simple statistic to evaluate the quality of high throughput screening assays) compared to androgenic activity detection methods we have used previously (Conley et al., 2017a; Medlock Kakaley et al., 2020). On a scale of 0–1, where 0 is an unacceptable assay and 1 is an ideal assay (Zhang et al., 1999), the Z-factor for the CV1-chAR is 0.76, while a

comparable assay (which uses the same mechanism of action) we have previously used for water quality screening was 0.32 (MDA-KB2) (Conley et al., 2019).

Although no glucocorticoid activity was detected in the WFP intake, pre-, or postdistribution water samples, others have detected glucocorticoid receptor activity in wastewater effluent (Chang et al., 2007; Jia et al., 2016; Medlock Kakaley et al., 2020; Schriks et al., 2010; Suzuki et al., 2015; Tousova et al., 2017) and impacted surface waters (Tousova et al., 2017). Previously, we quantified glucocorticoid activity (above MDL) ranging 6.0–43 ng DexEq_{CV1-hGR}/L in impacted surface and wastewater effluent specifically using the CV1-hGR bioassay (Conley et al., 2017a; Medlock Kakaley et al., 2020), although known glucocorticoid receptor ligands targeted in the complementary chemical analysis were not detected. In the comprehensive "budget balancing" exercise by Jia *et al.* triamcinolone acetonide, fluocinolone acetonide, clobetasol propionate, and fluticasone propionate were responsible for the majority of the detected GR activity (Jia et al., 2016). Discernibly all Chicago area samples fell below previously reported human health (drinking water) EBT of 21 (Brand et al., 2013) and 150 (Escher et al., 2015) ng DexEq/L, generated using the GR-CALUX bioassay.

5. Conclusions

We present a model that can expedite future testing of low activity samples (treated tapwater) without jeopardizing statistical stringency. Few detections of biological activity were identified in pre-distribution samples, but as we anticipated WFP treated post-distribution water samples did not produce any positive detections for endocrine activity above MDLs, and therefore likely do not contain any estrogenic, androgenic or glucocorticoid active compounds at concentrations that have the potential to cause adverse effects in humans.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Highlights

- Corresponding pre- and post-distribution treated water samples were assessed
- We applied a screening model for low level biological activity in water samples
- Inaugural use of an (anti)androgenic bioassay for water quality screening
- Estrogen and androgen activity were detected in untreated water
- No biological activity detected in tapwater above method detection limits

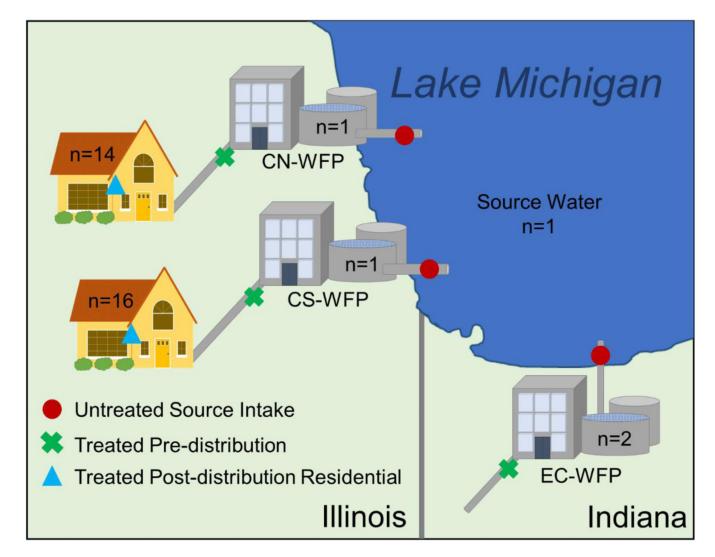


Figure 1. Water sampling scheme

Three sample types were taken from sites around the Greater Chicago metropolitan area including untreated Lake Michigan source water (red circle), water filtration plant (WFP)-treated pre-distribution water (green "x"), and post-distribution residential tapwater (blue triangle). Untreated intake and WFP-treated water was sampled at two WFPs in East Chicago, IN, one WFP in North Chicago, IL and one WFP in South Chicago. Residential post-distribution tapwater samples representing the North (n = 14) and South Chicago (n = 16) WFPs distribution areas were also sampled.

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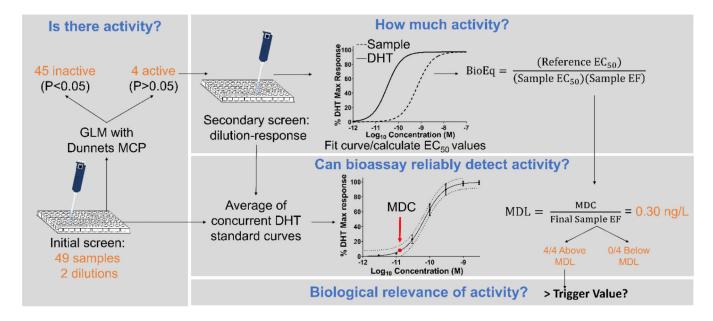


Figure 2. Tiered screening and statistical analysis model

for bioassays that measure endocrine activity in tapwater sample extracts. Here the model is presented using the CV1-chAR bioassay and Chicago area tapwater samples results. In tier one: *Is there activity*? all samples are screened with minimal dilutions and statistically compared to vehicle control (p > 0.05) using general linear model (GLM) and multiple comparison procedure (MCP). In tier two: *How much activity*? only active samples from tier one were screened again using a dilution-response to determine biological equivalency values (BioEq). Each sample BioEq is then compared to the bioassay method detection limit (MDL) and relevant trigger value.

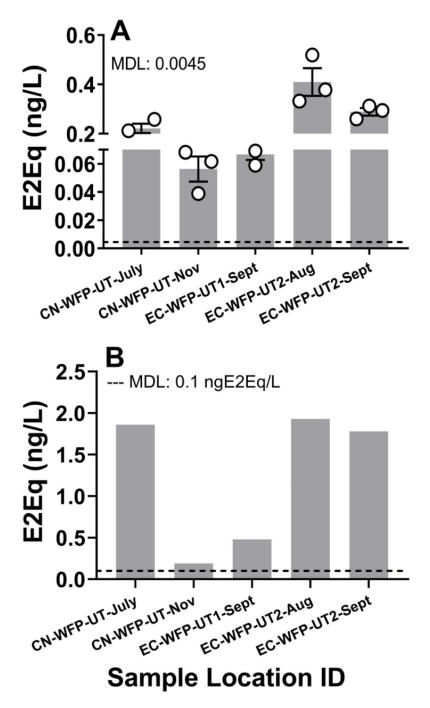


Figure 3. Estrogenic activity

was measured in extract samples from untreated Lake Michigan intakes, WFP treated predistribution waters, and post-distribution waters using the A) T47D-KBluc (biological replicate data is shown as mean ± standard deviation) and B) BLYES bioassays. Method detection limit (MDL) for T47D-KBluc was 0.0044ng E2Eq/L and 0.1 ng E2Eq/L for BLYES assay.

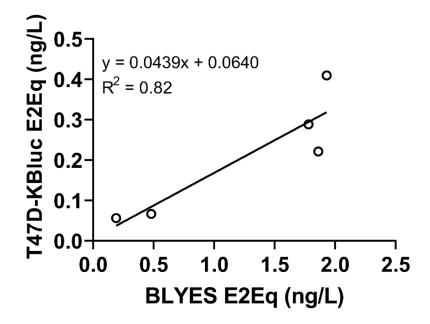


Figure 4. Estrogenic bioassay comparison

using estrogenic activity detected in water sample extracts. T47D-KBluc and BLYES bioassays are compared through linear regression analysis where y = 0.0439x + 0.0640 and $R^2 = 0.82$.

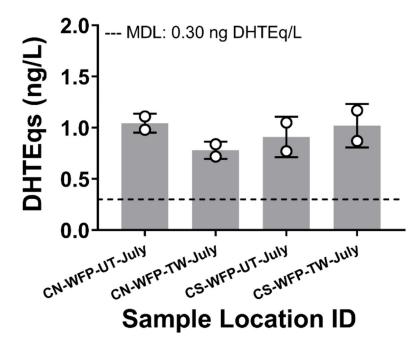


Figure 5. Androgenic activity

was measured in extract samples from untreated Lake Michigan, WFP treated predistribution, and post-distribution waters using a CV1-chAR bioassay. Biological replicate data is shown as mean \pm standard deviation and method detection limit (MDL) was 0.30 ng DHTEq/L.

Table 1.

Greater Chicago Area Sampling Site Locations, Dates, and Descriptions

Site Identifier	Sample Date	Site Location/Description
CN-WFP-UT	July 2017	Lake Michigan Intake
CN-WFP-UT	November 2017	Lake Michigan Intake
CN-WFP-TW	July 2017	Water Filtration Plant Pre-Distribution Effluent
CN-WFP-TW	November 2017	Water Filtration Plant Pre-Distribution Effluent
CN-TW-1	December 2017	Residential Treated Tapwater
CN-TW-2	November 2017	Residential Treated Tapwater
CN-TW-3	November 2017	Residential Treated Tapwater
CN-TW-4	December 2017	Residential Treated Tapwater
CN-TW-5	November 2017	Residential Treated Tapwater
CN-TW-6	November 2017	Residential Treated Tapwater
CN-TW-7	November 2017	Residential Treated Tapwater
CN-TW-8	December 2017	Residential Treated Tapwater
CN-TW-9	November 2017	Residential Treated Tapwater
CN-TW-10	November 2017	Residential Treated Tapwater
CN-TW-11	November 2017	Residential Treated Tapwater
CN-TW-12	November 2017	Residential Treated Tapwater
CN-TW-13	November 2017	Residential Treated Tapwater
CN-TW-14	December 2017	Residential Treated Tapwater
CS-WFP-UT	July 2017	Lake Michigan Intake
CS-WFP-TW	July 2017	Water Filtration Plant Pre-Distribution Effluent (Field Blan
CS-WFP-TW	July 2017	Water Filtration Plant Pre-Distribution Effluent
CS-WFP-TW	November 2017	Water Filtration Plant Pre-Distribution Effluent
CS-TW-1	December 2017	Residential Treated Tapwater
CS-TW-2	December 2017	Residential Treated Tapwater
CS-TW-3	December 2017	Residential Treated Tapwater
CS-TW-3	December 2017	Residential Treated Tapwater
CS-TW-4	December 2017	Residential Treated Tapwater
CS-TW-5	December 2017	Residential Treated Tapwater
CS-TW-6	December 2017	Residential Treated Tapwater
CS-TW-7	December 2017	Residential Treated Tapwater
CS-TW-8	December 2017	Residential Treated Tapwater
CS-TW-9	December 2017	Residential Treated Tapwater
CS-TW-10	December 2017	Residential Treated Tapwater
CS-TW-11	December 2017	Residential Treated Tapwater
CS-TW-12	November 2017	Residential Treated Tapwater
CS-TW-13	November 2017	Residential Treated Tapwater
CS-TW-14	November 2017	Residential Treated Tapwater
		······
CS-TW-15	November 2017	Residential Treated Tapwater

Site Identifier	Sample Date	Site Location/Description
EC-WFP-UT1	August 2018	Lake Michigan Intake
EC-WFP-UT1	September 2017	Lake Michigan Intake
EC-WFP-UT1	September 2017	Lake Michigan Intake (Field Blank)
EC-WFP-TW1	August 2018	Water Filtration Plant Pre-Distribution Effluent
EC-WFP-TW1	August 2018	Water Filtration Plant Pre-Distribution Effluent (Field Blank)
EC-WFP-TW1	September 2017	Water Filtration Plant Pre-Distribution Effluent
EC-WFP-UT2	August 2018	Lake Michigan Intake
EC-WFP-UT2	September 2017	Lake Michigan Intake
EC-WFP-TW2	August 2018	Water Filtration Plant Pre-Distribution Effluent
EC-WFP-TW2	September 2017	Water Filtration Plant Pre-Distribution Effluent

Each sampling event is identified with a three alpha-numeric element identifier; where the first element represents the geographical location (CN: Chicago North; CS: Chicago South; and EC: East Chicago, Indiana), the second indicates if the sample was collected pre-distribution (WFP: Water Filtration Plant) or post-distribution (TW), and the third indicates if the samples was untreated intake water (UT), or treated water (TW, or residence-specific numeric identifier). Two East Chicago WFPs were sampled and each East Chicago sample ends in a 1 or 2 indicating which plant was sampled.