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Luke R. Iwanowicz

U.S. Geological Survey, liwanowicz@usgs.gov

Vicki S. Blazer

U.S. Geological Survey

Christopher P. Guy

U.S. Fish and Wildlife Service, chris_guy@fws.gov

Alfred E. Pinkney

U.S. Fish and Wildlife Service, fred_pinkney@fws.gov

John E. Mullican

Inland Fisheries Management Division

See next page for additional authors

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Authors

Luke R. Iwanowicz, Vicki S. Blazer, Christopher P. Guy, Alfred E. Pinkney, John E. Mullican, and David A. Alvarez

REPRODUCTIVE HEALTH OF BASS IN THE POTOMAC, USA, DRAINAGE:
PART 1. EXPLORING THE EFFECTS OF PROXIMITY TO WASTEWATER TREATMENT
PLANT DISCHARGELUKE R. IWANOWICZ,*† VICKI S. BLAZER,† CHRISTOPHER P. GUY,‡ ALFRED E. PINKNEY,‡ JOHN E. MULLICAN,§
and DAVID A. ALVAREZ||

†National Fish Health Research Laboratory, U.S. Geological Survey, 11649 Leetown Road, Kearneysville, West Virginia 25430

‡U.S. Fish and Wildlife Service, Chesapeake Bay Field Office, 177 Admiral Cochrane Drive, Annapolis, Maryland 21401

§Maryland Department of Natural Resources Fisheries Service, Inland Fisheries Management Division, 10932 Putman Road,
Thurmont, Maryland 21788, USA

||U.S. Geological Survey, Columbia Environmental Research Center, 4200 New Haven Road, Columbia, Missouri 65201

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Abstract—Intersex (specifically, testicular oocytes) has been observed in male smallmouth bass (SMB; *Micropterus dolomieu*) and other centrarchids in the South Branch of the Potomac River, USA, and forks of the Shenandoah River, USA, during the past five years. This condition often is associated with exposure to estrogenic endocrine-disrupting chemicals in some fish species, but such chemicals and their sources have yet to be identified in the Potomac. In an attempt to better understand the plausible causes of this condition, we investigated the reproductive health of bass sampled up- and downstream of wastewater treatment plant (WWTP) effluent point sources on the Potomac River in Maryland, USA. Smallmouth bass were sampled from the Conococheague Creek and the Monocacy River, and largemouth bass (LMB; *Micropterus salmoides*) were collected near the Blue Plains WWTP on the mainstem of the Potomac River. Chemical analyses of compounds captured in passive samplers at these locations also were conducted. A high prevalence of intersex (82–100%) was identified in male SMB at all sites regardless of collection area. A lower prevalence of intersex (23%) was identified in male LMB collected at the Blue Plains site. When up- and downstream fish were compared, significant differences were noted only in fish from the Conococheague. Differences included condition factor, gonadosomatic index, plasma vitellogenin concentration, and estrogen to testosterone ratio. In general, chemicals associated with wastewater effluent, storm-water runoff, and agriculture were more prevalent at the downstream sampling sites. An exception was atrazine and its associated metabolites, which were present in greater concentrations at the upstream sites. It appears that proximity to effluent from WWTPs may influence the reproductive health of bass in the Potomac watershed, but inputs from other sources likely contribute to the widespread, high incidence of testicular oocytes.

Keywords—Smallmouth bass Endocrine disruption Potomac Intersex

INTRODUCTION

During 2003, a high prevalence of intersex (33–80%; specifically, testicular oocytes [TOs]), was identified in male smallmouth bass (SMB; *Micropterus dolomieu*) collected from numerous sites within the South Branch of the Potomac River (WV, USA). Subsequent surveys of SMB in a regionally distinct section of the Potomac River drainage (the Shenandoah River, VA, USA) revealed an even higher prevalence of TOs (80–100%) than found in the South Branch [1]. The observation of TOs in gonochoristic fish is not an unprecedented finding in wild fish, but it is uncommon and, therefore, is frequently used as a biomarker of exposure to estrogenic endocrine-disrupting chemicals (EDCs) [2–4]. Thus, the apparent widespread prevalence of this condition in the Potomac River drainage suggests the presence of biologically relevant point and nonpoint sources of these chemical compounds. Interestingly, given that other centrarchids, such as largemouth bass (LMB; *Micropterus salmoides*), at these sites have a lower prevalence of TOs compared with SMB (unpublished data from our laboratory), SMB may be particularly sensitive to exposure to putative EDCs. Currently, however, the causes for the condition and presence of other adverse responses are unknown.

The Potomac River basin is located in four U.S. mid-Atlantic states (MD, PA, VA, and WV) and the District of Columbia, and it is the second largest tributary of the Chesapeake Bay. The basin encompasses more than 23,600 km² and receives effluent from multiple industrial sources and urban combined sewer overflows as well as from nonpoint sources [5]. Wastewater treatment plants (WWTPs) are included as contributors of point-source effluent into this system. Though not previously examined in the Potomac River basin, WWTPs are recognized as a general source of EDCs. Examples include both natural and synthetic substances, such as polychlorinated biphenyls, phthalates, pesticides, heavy metals, alkylphenols, polycyclic aromatic hydrocarbons, 17 β -estradiol, 17 α -ethinylestradiol, and bisphenol A [6–8] (<http://coastalscience.noaa.gov/documents/techmemo149.pdf>). Treatment of wastewater before discharge differs between WWTPs, and not all of the common strategies effectively remove EDCs. Consequently, different WWTPs vary in their ability to remove particular classes of EDCs [9,10].

Wastewater treatment plants are only one potential source of chemicals that may adversely affect the general and reproductive health of fish. Runoff from agricultural land has been shown to contain hormones and pharmaceutical chemicals [11,12]. Additionally, pesticides and herbicides applied on agricultural, public, and residential lands are transported to aquatic ecosystems during rain events [13,14]. Leachates from landfills [15], polychlorinated biphenyls [16,17], polycyclic aro-

* To whom correspondence may be addressed
(liwanowicz@usgs.gov).

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Table 1. Water-quality parameters at collection sites within the Potomac (USA) drainage measured at the time of fish collections

Site	Water-quality parameter				
	Temperature (°C)	Conductivity (mS/cm)	Dissolved oxygen (mg/L)	% Dissolved oxygen saturation	pH
Conococheague, upstream	21.1	0.511	7.1	77.4	6.5
Conococheague, downstream	21.7	0.441	9.0	101.4	7.3
Monocacy, upstream	23.1	0.365	10.9	124.2	7.0
Monocacy, downstream	21.8	0.487	10.5	117.2	7.0
Blue Plains	25.8	0.615	6.8	100.0	6.8

matic hydrocarbons [18,19], and other compounds from industrial effluents, atmospheric deposition, spills, and storm-water runoff also contribute endocrine-modulating chemicals to the aquatic environment.

Intersex may be used as biomarker of reproductive health, but other bioindicators of reproductive health in fish populations are frequently evaluated as well. Examples include morphological indicators, such as changes in gonadosomatic index (GSI) and secondary sex characteristics [20,21]. Histologic observations within the gonads such as atresia, foci of pigmented cells, and gamete stage also are useful [22]. Lastly, physiological measures of plasma vitellogenin (VTG) [23] and sex steroid hormone concentrations are common endpoints of endocrine disruption [24,25].

Here, we present the results of a study designed to investigate the reproductive health of bass inhabiting waters near three WWTPs. The primary goals of this research were to further evaluate the extent of reproductive abnormalities in bass within the Potomac drainage and to investigate potential causes of the high prevalence of TOs and other adverse effects using a suite of biological endpoints. A companion study [26] quantified the presence of polar and nonpolar waterborne chemical contaminants and their *in vitro* estrogenicity to define the differences in chemical fingerprints between the sample locations.

MATERIALS AND METHODS

Sampling sites

A list of the 66 major WWTPs, defined as those with capacity of more than 0.5 million gallons per day (mgd; 1 mgd = 3.78 million liters per day), in the Maryland (USA) and Washington, DC, portions of the Potomac watershed was generated and reviewed. Site selection was governed by the principle criteria that bass were present, that they could be collected without adversely affecting the local populations, and that collection could occur immediately downstream and at least 15 km upstream of a WWTP. Sites along Conococheague Creek and the Monocacy River were selected because they conformed best to these criteria. Water temperatures ranged from 21.1 to 23.1°C at the four tributary sites and was 25.8°C at the lower mainstem site. All water-quality parameters (Table 1) were similar among sites and within acceptable ranges for bass growth, survival, and health [27].

Conococheague Creek (Fig. 1A) originates north of Chambersburg (PA, USA) and flows through Fairview (MD, USA; site of a U.S. Geological Survey [USGS] gauging station, 01614500) and into the Potomac River at Williamsport in Washington County (MD, USA). Land use within the 911-km² watershed is agricultural (61%), forested (34%), and urban (5%). The estimated minimum baseflow for 7 d over a 10-year period (7Q-10) for Conococheague Creek, using data from the nearest

USGS gauging station, is 40.1 mgd [28] (www.mde.state.md.us/assets/document/ConocoTech_Memo%20Final.pdf). The discharge flow of the WWTP (Fig. 1B), which uses a modified Ludzack–Ettinger treatment process, is 4.1 mgd, or approximately 10% of the minimum baseflow at the downstream site. Agricultural runoff is the primary anthropogenic input at the upstream site. A privately owned structure, Kemps Mill Dam, is located immediately upstream of the WWTP outfall. This structure is considered to be a major impediment to upstream fish migration.

The Monocacy River (Fig. 1A) is approximately 93 km in length and forms near the Maryland and Pennsylvania (USA) border. From there, it flows south between Frederick and Carroll counties (MD, USA), through the City of Frederick (MD, USA), and into the Potomac River. Approximately 60% of the Monocacy watershed is agricultural, 33% forested, and 7% urban. Two WWTPs are suspected to influence the selected downstream Monocacy site (Fig. 1B). The larger is the Frederick WWTP plant, which uses the anaerobic–anoxic–aerobic treatment process and has an average daily flow capacity of 8.0 mgd. The Fort Detrick (MD, USA) WWTP has a discharge flow of 1.0 mgd [29]. Fish collections for this site occurred from the outflow of the Frederick WWTP to approximately 1.5 km downstream. Although a 7Q-10 is not available for the Monocacy River at this site, a USGS gauging station (01643000) is located just upstream [30] (<http://nwis.waterdata.usgs.gov/md/nwis/annual>). A conservative estimated contribution of a 1.4% volume of stream flow from these two WWTPs was determined by dividing their combined maximum discharge rates by the annual flow rate for 2005. Average annual flow was calculated using historical flow rates measured at this gauging station. In drought years, such as 2002, the contribution to stream flow by these WWTPs may approximate 4% volume. The upper Monocacy River site is influenced primarily by agriculture, similar to the upper Conococheague Creek site.

A site at the Blue Plains WWTP in Washington, DC (Fig. 1A), also was selected for sampling, because it is the largest plant in the Potomac River watershed and the world's largest advanced (tertiary) WWTP. It serves the District of Columbia as well as Montgomery and Prince George counties in Maryland and Fairfax and Loudon counties in Virginia, and it has a design capacity to treat 370 mgd [31] (<http://www.dc.wasa.com/about/facilities.cfm>). The percentage effluent during baseflow conditions could not be estimated, because the area is in the tidal region of the Potomac River and no USGS stream gauges were located nearby.

Fish collections

Bass were collected by boat or barge electrofishing between September 6 and 14, 2005. This time marks the seasonal onset

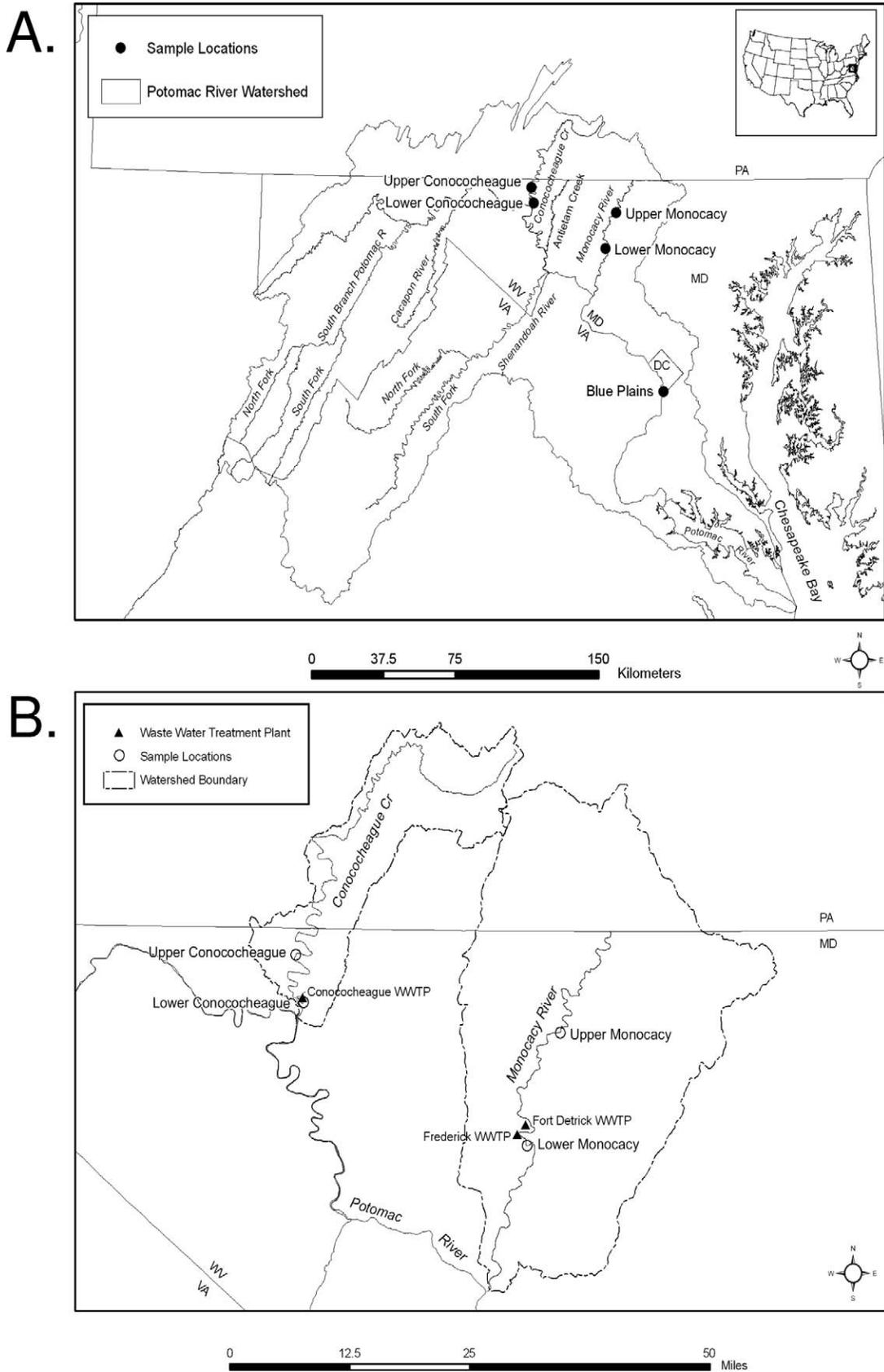


Fig. 1. (A) Collection site locations (USA) for smallmouth bass (*Micropterus dolomieu*) and largemouth bass (*Micropterus salmoides*) within the Potomac drainage. (B) Sample locations in special reference to wastewater treatment plants (WWTPs).

of reproductive recrudescence. Collections would have less impact on the population at this time than during the spring, when nest-building and spawning occurs, and the data could be compared to those of other field studies that assessed reproductive health of bass and were conducted in the fall [32–36]. Thirty mature bass (total length, >200 mm) were collected at each site in an attempt to obtain 10 males and 10 females. Smallmouth bass were collected at the sites on Conococheague Creek and the Monocacy River. Because no SMB were caught on the mainstem site of the Potomac near the Blue Plains plant, LMB were collected at this site (Fig. 1). Largemouth bass were only captured downstream of the Blue Plains site. Fish collected at the downstream WWTPs and Blue Plains were collected within 1 km of the discharge. Upstream and downstream sites on the same river were separated by at least 15 km.

Fish were killed with a lethal dose of tricaine methane sulfonate (FiquelTM; Argent), weighed on a portable scale to the nearest 0.1 g, and measured for total length to the nearest millimeter. Gonads and livers were weighed on an analytical balance with a calibrated tolerance of 0.01 g. Blood was drawn from the caudal vein with a heparinized syringe, transferred to vacutainers containing 62 U of sodium heparin (Fisher Scientific), and stored on wet ice. Blood was centrifuged for 10 min at 1,000 *g* and 4°C to hasten plasma separation within 3 h of collection. Plasma was removed, aliquoted into cryovials, and stored at –80°C until assayed for VTG and reproductive hormones.

A necropsy-based assessment similar to that described by Schmitt et al. [37] accompanied the sample collection. Grossly visible lesions and abnormalities were recorded. Gonads were removed and weighed for GSI determination as follows:

$$[\text{gonad wt}/(\text{body wt} - \text{gonad wt})] \cdot 100$$

Gonads were fixed in Z-fixTM (Anatech LTD) for histological evaluation. Otoliths were removed from the SMB and used for age determination [38].

Histopathological and biochemical procedures

Fixed gonads were dehydrated in alcohol, infiltrated with paraffin, sectioned at a thickness of 6 μm , and stained with hematoxylin and eosin [39]. Gonad sections were examined microscopically and staged, and any abnormalities were ranked from 0 to 4 (absent to severe) as described by Blazer [22]. Individual oocyte were staged as follows: Stage 1, immature (nucleolar); stage 2, early vitellogenic (cortical alveolar); stage 3, midvitellogenic (yolk droplet); stage 4, mature (yolk hydrates); and stage 5, postovulatory follicles. The stage for an ovary was based on the most prevalent oocyte stage present. At least five sections along the ovary were examined. The percentage of atretic eggs was determined by counting 200 oocytes and calculating the percentage of degenerating or necrotic oocytes. At least five pieces along the length of the testes were sectioned, and TO prevalence and severity were scored as described by Blazer et al. [1]. Male gonad stage was scored as follows: Stage 1, predominantly spermatogonia or spermatocytes; stage 2, approximately equal portions of spermatocytes, spermatids, and spermatozoa; stage 3, primarily spermatozoa; and stage 4, postspawn.

Plasma VTG concentrations were measured using a direct enzyme-linked immunosorbent assay (ELISA) with monoclonal antibody 3G2 and were carried out at the Center for Human and Environmental Toxicology, University of Florida (Gainesville, FL, USA), as described by Denslow et al. [40]. Con-

centrations of the unknowns were determined from the standard curves and using the Softmax Pro TM Program (Molecular Devices). The limit of detection was 0.001 mg/ml. Intra- and interassay variability were less than 10%.

Plasma hormone concentrations (17 β -estradiol and testosterone) were measured using radioimmunoassay according to the method described by Sower and Schreck [41]. Briefly, plasma samples were extracted twice in a 10-fold excess of diethyl ether, blown to dryness, and solubilized in 200 μl of room-temperature Dulbecco's phosphate-buffered saline (Sigma-Aldrich) containing 0.1% Knox gelatin (PG). One hundred microliters of antiestradiol antiserum (anti-17 β -estradiol Ab-244 anti-estradiol-6-BSA; Gordon Niswender) diluted 1:65,000 in PG buffer or antitestosterone antiserum (R156/7; Coralie Munroe) diluted 1:30,000 in PG buffer were added to each sample tube, vortexed, and incubated at room temperature for 30 min. The same volume of PG buffer was added to tubes designated to determine nonspecific background and total counts per minute (CPM). Following incubation, 100 μl of tritiated 17 β -estradiol or testosterone (5,000 CPM in PG buffer) were added to all tubes, vortexed, and incubated at room temperature for 60 or 30 min, respectively. Samples were immediately cooled in an ice bath for 30 min, and 500 μl of ice-cold charcoal-dextran solution (0.63% alkaline charcoal and 0.4% dextran in PG buffer) were added. Samples were vortexed, incubated on ice for 15 min, and centrifuged at 2,200 *g* for 20 min at 4°C. The supernatant was then decanted into scintillation vials containing 4 ml of OptiPhase HiSafe 2 scintillation fluid (PerkinElmer) and mixed by inversion. Sample CPM was measured using a Tri Carb liquid scintillation counter (PerkinElmer), and mean sample CPM was determined over an 8-min integration time. All samples were run in duplicate, and plasma steroid values were interpolated from a standard curve using curve-fitting algorithms in Prism for Windows[®] (Ver 4.03; GraphPad Software). Standard curves consisted of nine dilutions of 17 β -estradiol (E0950-000; Steraloids) or testosterone (A6950-000; Steraloids). Sample values were rejected and reevaluated if the coefficient of variation between duplicate tubes exceeded 10%.

Data analysis

Before comparative statistical analyses, all data were tested for normality using the Shapiro–Wilks *W* test and for homogeneity of variance with Levene's test. Given that all data did not conform to the assumptions of parametric statistics, the conservative, nonparametric Kruskal–Wallis test was selected. Data from the Blue Plains site were not included in statistical analyses, because LMB rather than SMB were collected. Differences in the prevalence of male bass with intersex and fish with measurable VTG were analyzed using Fisher's exact test. All statistical analyses were performed using SyStat[®] 11 for Windows (SyStat Software). Differences were considered to be significant when $p < 0.05$.

RESULTS

Morphometric and fish health indicators

At both Conococheague sites, an equal number of male and female SMB were collected. Only seven males were obtained at the Monocacy lower site, despite sampling 30 fish. At the upstream site, one male fish was misidentified as a female in the field; hence, unequal numbers were sampled. At the mainstem site, only seven LMB females were collected (Tables 2 and 3).

Table 2. Morphometric results for female bass from collection sites within the Potomac (USA) drainage^a

Site	<i>n</i>	Length (mm)	Weight (g)	Age	Condition factor
Conococheague, upstream	10	321 ± 17.3 A	484.3 ± 87.0 A	3.6 ± 0.5 A	1.35 ± 0.03 A
Conococheague, downstream	10	261 ± 8.8 B	233.2 ± 27.0 B	3.1 ± 0.2 A	1.27 ± 0.03 A
Monocacy, upstream	9	271 ± 18.8	284.2 ± 61.8	2.6 ± 0.4	1.26 ± 0.03
Monocacy, downstream	13	269 ± 14.2	280.0 ± 51.0	2.7 ± 0.4	1.27 ± 0.04
Blue Plains	7	370 ± 21.1	831.4 ± 180.5	ND	1.48 ± 0.01

^a Data are presented as the mean ± standard error. Values followed by different uppercase letters indicate that the upstream and downstream sites are significantly different (Kruskal–Wallis test, $p < 0.05$). ND = no data available.

Female SMB collected at the upstream Conococheague site were larger ($p = 0.008$), but not statistically older, than those collected downstream. Female bass collected at the upstream and downstream sites on the Monocacy were similar in size, age, and condition factor (Table 2). The LMB collected at the Blue Plains (mainstem of the Potomac River) site were larger than the SMB collected from the Conococheague or Monocacy rivers; unfortunately, their age was not determined.

Male SMB collected upstream on the Conococheague were heavier, with a significantly ($p = 0.009$) higher median condition factor and approximately the same age as those collected downstream. On the Monocacy, the males downstream were similar in size (length and weight), had a similar condition factor, but were significantly older than those collected at the upstream site (Table 3).

Reproductive indicators

All fish collected were sexually mature. The median GSI of female SMB collected at the upstream Conococheague site was approximately double and significantly ($p = 0.005$) higher than that of females from the downstream site (Table 4). Vitellogenin ranged from not detected (ND) to 3.378 mg/ml upstream and from ND to 0.802 mg/ml downstream, and it was detected in 8 of 10 females (80%) at each site. The median VTG concentration in downstream females was significantly ($p = 0.003$) lower than that of fish collected upstream (Table 4). This coincided with 90% of the female gonads collected at the upstream site being at stage 2, whereas only 60% of the downstream gonads were at stage 2 (Fig. 2A). The remaining 10 and 40% of ovaries, respectively, contained oocytes at stage 1 (Fig. 2B). Atretic follicles (Fig. 2C) were present in females from both collection sites. The average percentage atretic follicles per ovary was 8.6% (standard deviation [SD] = 4.9%) and 4.2% (SD = 4.1%) for females collected at the up- and downstream sites, respectively.

On the Monocacy, the median GSI of the upstream and downstream female SMB was similar. Only 56% of the female SMB collected upstream and 77% of those collected downstream had measurable VTG, and no significant difference was found in concentrations between sites. Those collected down-

stream, however, had only 32% (or 21% if using the mean of only those with measurable levels) of the amount of those collected upstream. Ovaries with stage 2 oocytes were observed in 77 and 85% of the females collected at the upstream and downstream sites, respectively. The average percentage atretic follicles per ovary was 3.8% (SD = 4.2%) and 6.0% (SD = 5.4%) for females collected at the up- and downstream sites, respectively.

At the mainstem site, the mean GSI of the female LMB was the lowest observed (i.e., 0.43), and though 100% had measurable VTG, the mean concentration was lower than that measured in SMB at the two upstream sites (Table 4). All female gonads collected at this site were stage 1.

Male SMB collected at the upstream site on the Conococheague had approximately triple the GSI of those collected downstream. Sixty percent of the upstream male bass had measurable VTG (range, ND to 0.337 mg/ml), whereas 90% of the male downstream SMB had circulating VTG (range, ND to 0.306 mg/ml). All testes (100%) were at stage 2 at the upstream site. Downstream, however, only 60% male gonads were at this stage, and the remaining 40% were stage 1. No significant difference was found in the mean concentration of plasma VTG, but the mean concentration in the upstream bass was more than double that of those collected downstream (Table 5). No statistical difference ($p = 0.303$) was found in the number of males with detectable VTG. A high percentage of SMB with TOs (Fig. 2D and Table 2) was observed at both upstream and downstream sites on the Conococheague (100 and 90%, respectively).

On the Monocacy, the upstream males had a median GSI similar to that of the downstream males (Table 5). Male gonads from the upper Monocacy were predominantly stage 1 (55%), with the remaining 45% at stage 2, whereas testes from the lower site were mostly stage 2 (75%), with 25% at stage 1. Vitellogenin was observed in 45% of male SMB collected upstream (range, ND to 0.081 mg/ml) and in 33% of those collected downstream (range, ND to 0.278 mg/ml), and 80 and 100%, respectively, had TOs. Eighty-five percent of the LMB collected at the Blue Plains site had measurable VTG, and the mean VTG concentration at this site was the highest measured.

Table 3. Morphometric results of male bass from collection sites within the Potomac (USA) drainage^a

Site	<i>n</i>	Length (cm)	Weight (g)	Age	Condition factor
Conococheague, upstream	10	292 ± 12.7 A	342.7 ± 40.2 A	2.9 ± 0.3 A	1.33 ± 0.03 A
Conococheague, downstream	10	255 ± 10.4 A	207.7 ± 24.1 B	2.6 ± 0.4 A	1.21 ± 0.03 B
Monocacy, upstream	11	245 ± 11.3	200.1 ± 32.0	1.7 ± 0.2	1.27 ± 0.02
Monocacy, downstream	7	264 ± 10.6	247.2 ± 28.2	2.6 ± 0.4	1.22 ± 0.03
Blue Plains	13	353 ± 10.1	743.4 ± 87.3	ND	1.61 ± 0.05

^a Data are presented as the mean ± standard error. Values followed by different uppercase letters indicate that the upstream and downstream sites are significantly different (Kruskal–Wallis test, $p < 0.05$). ND = no data available.

Table 4. Gonadosomatic indices (GSI), plasma vitellogenin (VTG), and estrogen to testosterone ratio (E:T) of female bass from collection sites within the Potomac (USA) drainage^a

Site	GSI	% Samples with VTG	Plasma VTG (mg/ml) ^b	E:T
Conococheague, upstream	1.26 ± 0.35 A	80 A	1.247 ± 0.485 A	0.98 ± 0.14 A
Conococheague, downstream	0.63 ± 0.06 B	80 A	0.119 ± 0.105 B	0.91 ± 0.05 A
Monocacy, upstream	0.94 ± 0.11 A	45	1.885 ± 1.014	0.89 ± 0.17
Monocacy, downstream	0.82 ± 0.09	77	0.395 ± 0.118	0.95 ± 0.07
Blue Plains	0.43 ± 0.05	100	0.465 ± 0.251	0.76 ± 0.09

^a Data are presented as the mean ± standard error. Values followed by different uppercase letters indicate the upstream and downstream sites are significantly different (Kruskal–Wallis or Fisher's exact test, $p < 0.05$).

^b Mean of only those fish with a measurable amount of VTG.

At Blue Plains, in only 23% was the presence of TO noted (Table 5).

Median plasma estradiol concentrations of the SMB were significantly lower in the male fish at both sites when compared to the female fish. On the Conococheague, the downstream males had approximately double the median plasma estradiol of those upstream, whereas on the Monocacy, upstream and downstream levels were similar. Females on the Conococheague were similar upstream and downstream, whereas on the Monocacy, bass from the downstream site had higher concentrations compared to those in the upstream females. Median male and female estradiol concentrations were similar in the LMB from Blue Plains (Fig. 3).

Median plasma testosterone concentrations of LMB were similar between males and females. On the Conococheague, the upstream males had a lower mean concentration compared with those downstream, as did the females. On the Monocacy, the median concentrations of the upstream and downstream males were similar. The upstream females had a lower concentration than the downstream females (Fig. 4). These differences, however, were not statistically different.

The median estradiol to testosterone ratios (E:T) of the female SMB were all similar and approximately one; however, the median E:T of the female LMB was lower, at 0.73. Male SMB on both rivers had a higher median E:T downstream than upstream. The SMB males at the downstream sites had an E:T similar to that of the male LMB (Table 5).

DISCUSSION

Abnormal reproductive physiology resulting from unavoidable exposure to wastewater effluent has been documented in wild fish populations for more than a decade [2,6,42–44]. Although WWTP effluent contains a milieu of biologically active substances, those with estrogenic potential have garnered the most attention from the scientific community [45,46]. Numerous plasma-associated physiological measures (i.e., VTG and steroid hormones) have been used as indicators of exposure to estrogenic chemicals. The presence of intersex (principally TOs in males), however, continues to rival these measures, because it appears to be a less transient indicator of estrogenic exposure. Experimental evidence purports that intersex may be induced early in development, during critical

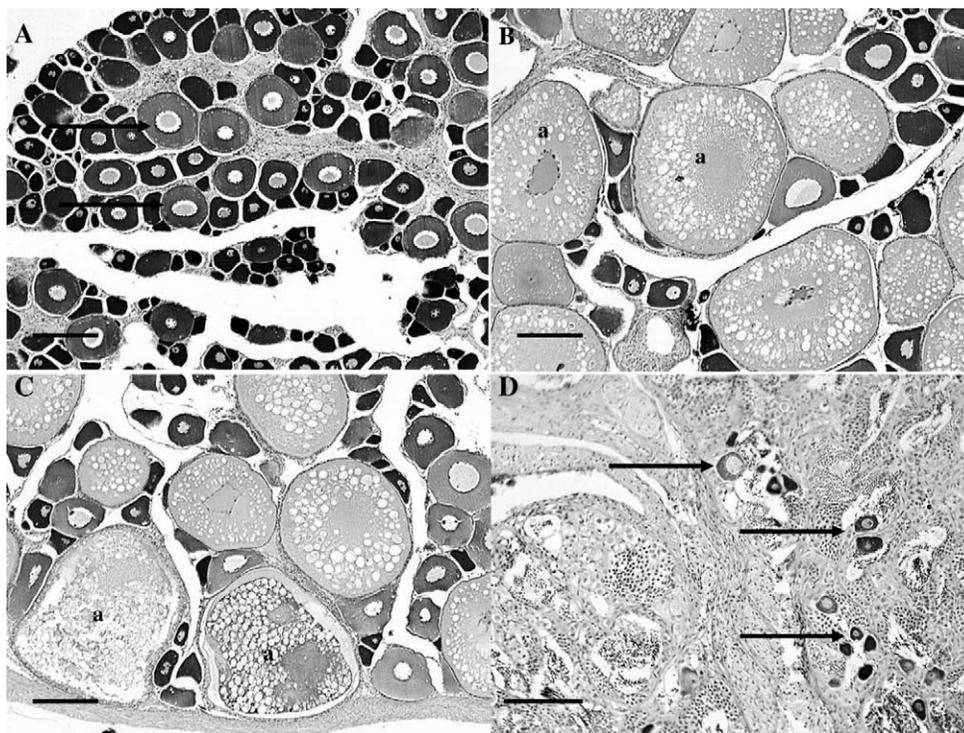


Fig. 2. Microscopic observations in bass gonads. (A) Stage 1 ovary with oocytes that have only progressed to the perinucleolar stage (arrows). (B) Stage 2 ovary containing oocytes that have progressed to the cortical alveolar stage (a). (C) Atretic oocytes (a) within a stage 2 ovary. (D) Oocytes (arrows) within the testis of a bass. Hematoxylin-and-eosin stain; bar = 200 μm (A–C) and 100 μm (D).

Table 5. Gonadosomatic index (GSI), testicular oocytes (TO), plasma vitellogenin (VTG), and estrogen to testosterone ratio (E:T) ratio of male bass from collection sites within the Potomac (USA) drainage^a

Site	GSI	TO		VTG		E:T
		Prevalence (%)	Severity index	% Samples with VTG	Plasma concentration (mg/ml) ^b	
Conococheague, upstream	0.39 ± 0.06 A	100 A	2.1 ± 0.3 A	60 A	0.117 ± 0.051 A	0.37 ± 0.10 A
Conococheague, downstream	0.13 ± 0.06 B	90 A	1.8 ± 0.4 A	90 A	0.050 ± 0.036 B	0.58 ± 0.08 B
Monocacy, upstream	0.30 ± 0.04	82	1.2 ± 0.3	45	0.059 ± 0.010 A	0.46 ± 0.13
Monocacy, downstream	0.30 ± 0.15	100	1.9 ± 0.3	33	0.143 ± 0.085 A	0.51 ± 0.15
Blue Plains	0.06 ± 0.003	23	0.2 ± 0.2	85	0.577 ± 0.165	0.51 ± 0.07

^a Data are presented as the mean ± standard error. Values followed by different uppercase letters indicate the upstream and downstream sites are significantly different (Kruskal–Wallis or Fisher's exact test, $p < 0.05$).

^b Mean of only those fish with a measurable amount of VTG.

stages of sexual differentiation in gonochoristic fish, though exceptions certainly exist. That same research demonstrates that intersex is not induced by similar estrogenic stimuli at later life stages in some species [47]. Interestingly, exposure to estrogenic stimuli later in life may exacerbate the severity of TOs in instances when primary exposure occurred at the critical stages of sexual development [43].

In the present study, estrogenic endocrine disruption, as evidenced by the presence of TOs, was observed at all study sites. The incidence of TOs in male SMB collected from Conococheague Creek and the Monocacy River exceeded 80%, which is higher than that generally observed in SMB from the South Branch of the Potomac but similar to that observed in SMB from the Shenandoah River [1]. When intersex is induced in male SMB from this watershed is not known, but it may occur during critical developmental stages. Male LMB collected at the Blue Plains site had a much lower prevalence of TO compared with that in SMB at the other samples sites. Considering that the Blue Plains site generally had higher levels of most contaminants (Table 6), this may indicate differences in sensitivity to estrogenic chemicals in these closely related species or differences in chemical concentrations among preferential spawning habitat for the two species. Unfortunately, a baseline occurrence of TOs has not been deter-

mined for either LMB or SMB. Currently, the best data for SMB is from out-of-basin sites with both low human population and low to moderate agricultural land use, at which 14 to 22% of male SMB were observed to have TOs [1]. From 1995 to 2004, the USGS conducted fish health assessments, including reproductive endpoints, at large river sites in the Mississippi [32], Rio Grand [33], Columbia [34], Colorado [35], and Savannah, Pee Dee, Mobile, and Apalachicola–Chattahoochee–Flint [36] river basins (all USA). In these studies, LMB and common carp (*Cyprinus carpio*) were the target species. At a total of 55 sites, LMB males were collected, and at 32 of these sites (58%), no TOs were reported, suggesting that the baseline prevalence for LMB may be zero. It is interesting that SMB appear to have a fairly high baseline prevalence of this condition. Unfortunately, relatively little information is available regarding sexual development and sex determination in this species. For this reason, it currently is not possible to rule out factors other than, or in addition to, contaminant exposure as contributing to this condition.

The induction of VTG in male fish is perhaps the most frequently applied biomarker of estrogenic exposure, particularly near WWTP discharges [48–51]. Hepatic synthesis of this glycolipoprotein is estrogen dependent. Given the appropriate stimulus, it is produced in females and males alike;

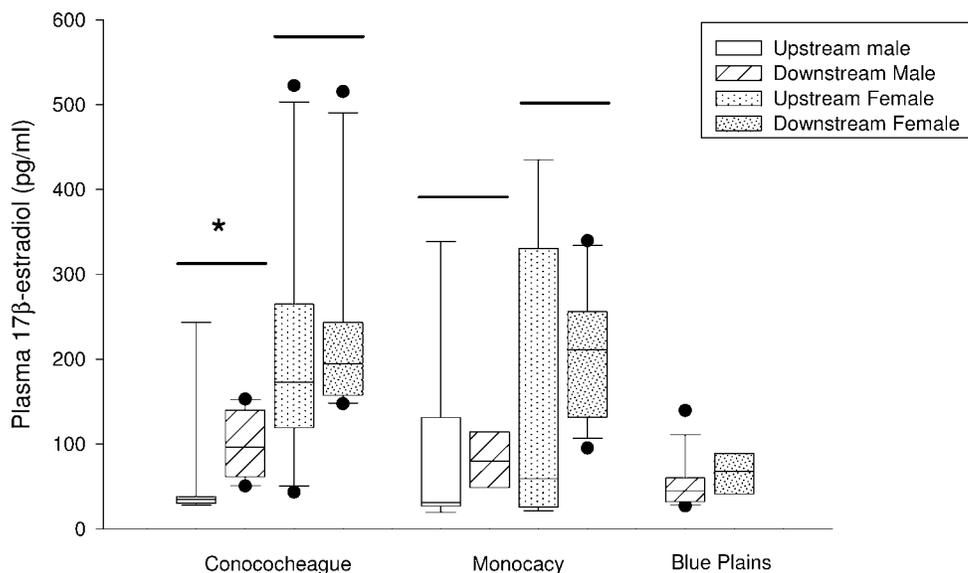


Fig. 3. Plasma 17 β -estradiol concentrations (pg/ml) of smallmouth bass (*Micropterus dolomieu*) and largemouth bass (*Micropterus salmoides*). Upstream/downstream pairs marked with an asterisk are significantly different ($p < 0.05$). Dots represent outliers, and whiskers mark the 5th and 95th percentiles.

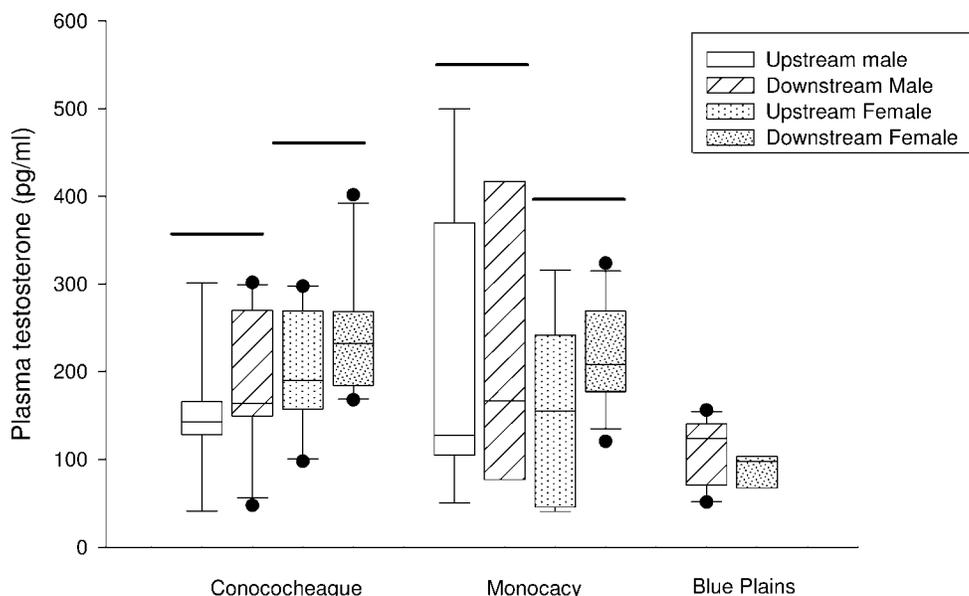


Fig. 4. Plasma testosterone concentrations (pg/ml) in smallmouth bass (*Micropterus dolomieu*) and largemouth bass (*Micropterus salmoides*) captured in the vicinity of wastewater treatment facilities. Dots represent outliers, and whiskers mark 5th and 95th percentiles.

however, under normal physiological conditions, males do not produce VTG. Vitellogenin has no known biological function in males, although antimicrobial actions have been suggested [52]. Unlike induction of TOs, the induction of VTG in male fish is rapid, and the protein is detectable in the plasma within days postexposure. Consequently, the presence of plasma VTG is more indicative than TOs of recent and/or continued exposure to estrogenic compounds. In the present study, plasma VTG in male SMB ranged from undetectable to 0.337 mg/ml.

When considering males with measurable VTG, mean concentrations of this protein were highest at the upstream Conococheague and downstream Monocacy sites. These concentrations were considerably lower than those measured in male LMB at the Blue Plains site (Table 5). The significance of elevated concentrations of VTG in these species is unknown, but its presence serves as a useful indication that estrogenic chemicals are present at biologically relevant concentrations. Recently, it has been shown that adverse effects are observed

Table 6. Selected concentrations of chemicals that measured above the method quantitation limit (MQL) in passive sampler extracts^a

Chemical	Conococheague		Monocacy		Blue Plains
	Upstream	Downstream	Upstream	Downstream	
Chemical estimated (pg/L)					
Total polychlorinated biphenyls	66	215	<MQL	410	2,550
Total polyaromatic hydrocarbons	4,136	4,114	3,921	16,790	21,825
Total DDTs ^b	94	144	<MQL	655	355
Hexachlorobenzene	83	58	<MQL	<MQL	59
Pentachloranisole	56	225	110	190	310
Lindane	<MQL	620	<MQL	550	<MQL
Total benzenehexachloride ^c	89	404	<MQL	93	218
Chlorpyrifos	<MQL	115	<MQL	48	475
Total chlordane ^d	43	136	35	95	620
Dieldrin	175	295	104	195	550
Heptachlor epoxide	<MQL	165	<MQL	150	405
Endosulfan + endosulfan II	630	1,470	830	1,330	6,050
Metolachlor	730	1,115	12,000	10,750	1,850
Atrazine ^e	90,000	46,850	88,000	44,000	30,900
Wastewater-related chemicals (ng/POCIS)					
Celestolide	<MDL	130	<MDL	<MDL	130
Tonalide	<MDL	120	<MDL	<MDL	515
Galaxolide	<MDL	335	<MDL	210	955
Prometon	100	115	100	<MDL	145
Tri(2-chloroethyl)phosphate	<MQL	165	<MQL	100	360

^a For more detail, see Alvarez et al. [26]. MDL = method detection limit; POCIS = polar organic chemical integrative sampler.

^b Sum of *o,p'*-dichlorodiphenyldichloroethylene (DDE), *p,p'*-DDE, *o,p'*-dichlorodiphenyldichloroethane (DDD), *o,p'*-DDT, *p,p'*-DDD, and *p,p'*-DDT.

^c Sum of α -benzenehexachloride, β -benzenehexachloride, and δ -benzenehexachloride.

^d Sum of oxychlordane, *trans*-chlordane, and *cis*-chlordane.

^e Sum of desisopropylatrazine, desethylatrazine, and atrazine.

in male fathead minnows (*Pimephales promelas*) with VTG concentrations of greater than 0.5 mg/ml, but not below this concentration [53]. Only five male LMB from the present study had concentrations exceeding this threshold, and four of these five male LMB (80%) had levels exceeding 1 mg/ml. Most striking at this site was the observation that males had VTG concentrations comparable to those of females. The detection of measurable VTG in male SMB and LMB indicates that in addition to early life exposure to estrogens (suggested by TOs), these fish also likely are exposed to estrogenic chemicals as adults. The extent and frequency of this exposure is still unknown. Additionally, the kinetics of VTG clearance in bass is unknown. This is of particular importance, because male fish do not appear to have specific mechanisms to excrete VTG [54]. Consequently, VTG remains detectable in the plasma of male fish for weeks to months postexposure [53,54]. Thus, in the present study, the presence of VTG may be the result of exposure that occurred weeks to months before our sampling. The observed variability of plasma VTG in male fish includes the possibility of recent immigration of fish from less impacted upstream areas or simply differences in the concentration of chemicals within the river based on flow dynamics. Unfortunately, published home-range studies of SMB in the Potomac drainage are not available; however, home ranges of less than 200 m have been reported for river systems in Missouri (USA) [55]. Other studies have indicated that some populations/individuals are relatively sedentary and others migratory, and that this may be affected by season [56]. Although our upstream and downstream sites were separated by a much greater distance than the reported home range for some SMB populations, without specific regional data regarding bass movement, we cannot rule out the possibility of some migration between sites. This is less likely on the Conococheague, however, because of the dam immediately upstream of the WWTP outfall.

Species of the genus *Micropterus*, like most gonochoristic fish, experience an annual reproductive cycle in which recrudescence initiates during the fall and spawning occurs in the spring. Specific dates of these events are governed largely by water temperature and photoperiod. Research conducted on Florida LMB (*Micropterus salmoides floridans*) maintained in tanks and ponds in Texas (USA) has established the baseline reproductive cycle for this (sub)species. Recrudescence is evident in October through December [57], and peak GSI occurs between late February and late April (immediate prespawn to spawn). The GSI declines rapidly postspawn in May and continues to decrease until reaching a minimum in September. A similar cycle has been documented for hatchery-raised LMB in Florida [58]. In both studies, the GSI was approximately 0.8 to 1.0 in females and 0.1 to 0.2 in males during September (water temperature, $\sim 15^{\circ}\text{C}$). In the current study, the mean GSI was 0.4 for female and 0.06 for male LMB. This is considerably lower than the September values reported above; however, this difference probably is the result of the warmer water temperature at the Blue Plains site. The onset of recrudescence at this site likely occurs later in the year, when water temperature decreases. This would explain the low GSI and sex steroid values, which probably were at baseline values during sampling. If this is the case, however, then the presence of VTG in both male and female LMB is likely the result of exposures to estrogens in the water rather than endogenous stimuli. Biological evidence of estrogenic substances in the water at this site is confirmed by results of the bioluminescence

yeast estrogen screen conducted by Alvarez et al. [26]. Additionally, concentrations of plasma VTG in male LMB normally are less than 0.05 mg/ml at any given time during the year [58]. Of particular relevance to the present study, the presence of TOs was not noted in either of the above studies.

Few studies have documented the normal changes in GSI, VTG, or reproductive hormones throughout the reproductive cycle of SMB. The GSI of naturalized lake SMB in Japan is lowest in August and begins to rise during September (water temperature, $\sim 20^{\circ}\text{C}$), to 2.0 in females and 0.8 in males [59]. In the present study, fish collected from the downstream Conococheague site had significantly lower GSI compared to fish collected from the upstream Conococheague, both for females and for males ($p = 0.005$ and $p < 0.001$, respectively). These sites experience the same photoperiod, and water temperatures are similar (Table 1). The GSIs at this site also were lower than those measured in fish from the Monocacy sites, particularly in the case of males (Table 5). Because water temperature was similar among all sites, chemical exposure from site-specific inputs is a likely explanation for the delayed onset of spermatogenic recrudescence. The GSI of females collected downstream of the WWTP on the Conococheague was significantly lower than that of females collected upstream, which is consistent with the lower plasma VTG concentrations (Table 4). Again, this observation further corroborates differences in the timing of recrudescence at the lower Conococheague site compared to that upstream. Because VTG production is related to fecundity and egg quality in individual female fish [60,61], the lower concentrations in female bass collected downstream could adversely affect egg quality and subsequent survival of fry. However, because the bass were collected in the fall, either at the onset of or early in recrudescence, further work is required to confidently document reproductive impairment and population-level effects.

Previous research has demonstrated that roach (*Rutilus rutilus*) exposed to sewage treatment plant effluent had an increased incidence of atresia, decreased estradiol in females, but increased estradiol and testosterone in males [49]. In the current study, differences in steroid hormones were subtle, and significant differences were only observed in males from the Conococheague. Males downstream had significantly ($p = 0.004$) higher plasma estradiol concentrations compared with those at the upstream site on the Conococheague. Whereas concentrations of reproductive hormones are useful indicators for disruption of the reproductive cycle [24], these concentrations often are confounded by natural fluctuations and environmental factors, such as water temperature influenced by rain and runoff events [25]. Concentrations of testosterone and estradiol in male SMB have been reported previously at 1 ng/ml or less during this time of year in Japan. In female SMB, these concentrations are higher, but still less than 2 ng/ml [59]. These concentrations are considerably higher than those measured at any site during the present study, indicating that SMB in the Potomac watershed initiate recrudescence later in the year. This likely results from geographical differences. To best interpret the significance of different hormone concentrations in these fish, additional multiyear, multiseason data are required to define the normal regional reproductive cycle of SMB.

No individual chemicals were identified as the singular cause of TO and VTG induction in male bass, but a number of EDCs were measured at all sites (Table 6). Passive samplers, both polar organic chemical integrative samplers and semi-

permeable membrane devices, were deployed at the sites for approximately one month during the sampling period. More detailed information regarding the chemical data is provided by Alvarez et al. [26]. Chemical profiles differed between the sample sites and may be explained, in part, by the type of treatment used at specific plants. This also may have contributed to the biological differences observed between bass collected at the upstream and downstream sites in the Conococheague but in not the Monocacy. For instance, three wastewater-associated chemicals commonly used in fragrances (cetylolide, tonalide, and galaxolide) were identified at the downstream site on the Conococheague. Only one of these three chemicals (tonalide) was found at the downstream site on the Monocacy, and none of these three was found at either upstream site. Similarly, tri(2-chloroethyl)phosphate, used in plasticizers and flame retardants, was not found at either upstream site but was measured at both downstream sites, though at a lower concentration on the Monocacy (Table 6). In a study of three WWTPs, Thomas and Foster [62] found that the majority of acidic pharmaceuticals, caffeine, and the antimicrobial triclosan were removed during secondary treatment. The difference in the secondary treatment processes (modified Ludzack-Ettinger [MLE] vs anaerobic-anoxic-aerobic [A_2O]) at the two WWTPs in the present study may explain some of the observed differences. The Conococheague plant uses the MLE process that employs activated sludge consisting of anoxic and oxic phases for biological nutrient reduction, whereas the Frederick WWTP recently upgraded to the A_2O activated-sludge process that includes an anaerobic phase as well as the anoxic and oxic cycles [63] (<http://www.mde.state.md.us/assets/document/BRF%20Gannett%20Fleming-GMB%20presentation.pdf>). Additional pharmaceuticals may be removed during this phase. In addition, factors such as overall removal efficiencies, operational practices, the nature and concentration of the wastewater influent, and the percentage of river baseflow contributed by effluent may contribute to the differences. Estimated average annual contribution of WWTPs to flow in the Conococheague and Monocacy was approximately 1.1 and 1.4%, respectively. These are annual averages, and times exist when WWTP contribution is higher. In general, however, the percentage contribution is fairly low compared to that in other parts of the country and world where intersex has been identified. That female sex steroids were at or below the method detection limit (see Alvarez et al. [26]) and WWTP contribution to river flow is low suggest other or additional factors as being responsible, in part, for the disrupted reproductive parameters.

Hexachlorobenzene, a chemical that has been shown to alter sex hormone concentrations in crucian carp (*Carassius auratus gibelio*) [64], was present in the Conococheague but not in the Monocacy. Historically, this chemical was used through the early 1960s as a pesticide to protect the seeds of onions and sorghum, wheat, and other grains against fungus. Currently, hexachlorobenzene is not used commercially in the United States, but it is formed as a by-product during the manufacture of other solvents and some pesticides [65] (<http://www.atsdr.cdc.gov/toxprofiles/tp90.pdf>). Whereas the presence of hexachlorobenzene is a clear difference between the two rivers, its presence does not explain biological differences in SMB at the up- and downstream sites in the Conococheague. In general, chemical compounds were found at similar or higher concentrations at downstream sites compared to those at upstream sites. The most notable exception was

atrazine and its metabolites, which generally were higher at upstream sites, where agricultural input was greatest. Atrazine is a chlorotriazine herbicide used to control annual grasses and broadleaf weeds that is prominently applied where corn is grown. It is associated with intersex and reproductive anomalies in anurans [66]. The exposure of adult fish to atrazine resulted in suppressive effects on plasma androgens and induction of estrogen (both dose and time related) in goldfish [13], but no strong estrogenic effects or overt reproductive toxicity in fathead minnows [67]. To our knowledge, exposure studies during critical periods of sexual differentiation have not been conducted, and it is unknown if atrazine could induce intersex in SMB. The widespread distribution of this herbicide within the Chesapeake watershed [68], however, and particularly the concentrations measured during spring at the current sites [26], may explain the high prevalence of intersex in the Potomac watershed and should be studied in more depth.

Although natural and synthetic estrogens most commonly are associated with the induction of TOs [2–4,47] and VTG in male fish [23,40,61], no steroid hormones were detected in the passive extracts of the present study. It is important to note, however, that the detection limits for 17β -estradiol and 17α -ethinylestradiol were both 2.5 ng/L [26]. Concentrations of 17α -ethinylestradiol as low as 1 ng/L, however, have been shown to induce intersex [69]. A dose-dependent increase in VTG was induced in juvenile zebrafish (*Danio rerio*) exposed to 2 ng/L or greater of 17α -ethinylestradiol [70], and levels as low as 0.1 ng/L induced VTG synthesis in immature rainbow trout (*Oncorhynchus mykiss*) [42]. Thus, that sex steroids were not identified does not imply their absence at biologically relevant concentrations. In fact, estrogenic activity was identified in polar organic chemical integrative samples from all sites using a yeast reporter assay [26]. In short, these findings emphasize the importance of examining both biological and chemical endpoints in field studies, and the potential biological effects of chronic or intermittent exposure to complex chemical mixtures may influence in aquatic organisms.

Based on the physiological and morphological measures, potential adverse effects of WWTP effluent were observed, particularly downstream of the Conococheague WWTP. These included decreased GSI, decreased circulating VTG in female bass, and altered reproductive hormone concentrations. Vitellogenin in male bass and a high prevalence of TO were noted at both upstream and downstream sites, indicating that other sources of endocrine-modulating chemicals, such as agricultural, suburban, and urban runoff, also may influence these populations. In conclusion, though it appears that effluent from some WWTPs may affect the reproductive health of fish in the Potomac watershed, inputs from other sources likely contribute to the widespread, high prevalence of TOs.

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