

Collapse of a fish population after exposure to a synthetic estrogen

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Municipal wastewaters are a complex mixture containing estrogens and estrogen mimics that are known to affect the reproductive health of wild fishes. Male fishes downstream of some wastewater outfalls produce vitellogenin (VTG) (a protein normally synthesized by females during oocyte maturation) and early-stage eggs in their testes, and this feminization has been attributed to the presence of estrogenic substances such as natural estrogens [estrone or 17 β -estradiol (E2)], the synthetic estrogen used in birth-control pills [17 α -ethynylestradiol (EE2)], or weaker estrogen mimics such as nonylphenol in the water. Despite widespread evidence that male fishes are being feminized, it is not known whether these low-level, chronic exposures adversely impact the sustainability of wild populations. We conducted a 7-year, whole-lake experiment at the Experimental Lakes Area (ELA) in northwestern Ontario, Canada, and showed that chronic exposure of fathead minnow (*Pimephales promelas*) to low concentrations (5–6 ng·L⁻¹) of the potent 17 α -ethynylestradiol led to feminization of males through the production of vitellogenin mRNA and protein, impacts on gonadal development as evidenced by intersex in males and altered oogenesis in females, and, ultimately, a near extinction of this species from the lake. Our observations demonstrate that the concentrations of estrogens and their mimics observed in freshwaters can impact the sustainability of wild fish populations.

endocrine disruptors | fathead minnow | municipal wastewaters | population-level effects | whole-lake experiment

There is considerable evidence that fishes inhabiting waters that receive untreated municipal wastewaters or effluents from municipal wastewater treatment plants (MWTPs) are exposed to chemicals that affect reproductive endocrine function. Male fish downstream of some wastewater outfalls produce vitellogenin (VTG) mRNA and protein, associated with oocyte maturation in females, and early-stage eggs in their testes (1–3). This feminization has been linked to the presence of estrogenic substances such as the natural estrogen 17 β -estradiol (E2) and the synthetic estrogen 17 α -ethynylestradiol (EE2) (4). Natural and synthetic estrogens are not completely broken down in current MWTP processes (5), and, as a result, are discharged into receiving waters in both treated and untreated wastewaters and found in the aquatic environment at low parts per trillion concentrations (typically <5 ng·L⁻¹; refs. 6–8). These effluents contain mixtures of individual estrogens and their mimics that differ in their ability to elicit estrogenic responses (e.g., ref. 9). For this reason, total estrogenicity (expressed as E2 equivalents) of an effluent or water sample is determined either by summing concentrations of individual compounds after adjusting these concentrations by the compound's estrogenic potency (relative to E2; ref. 10) or with bioassays (e.g., refs. 10–12). Using these approaches, total E2 equivalents of up to 147 and 17 ng·L⁻¹ have been measured in final effluents and surface waters, respectively (10–12). Within this group of substances, the estrogen used in birth-control pills, EE2, is one of the more potent estrogens

present (9) and has been linked to the feminization of male fishes in rivers receiving municipal wastewater (4, 6).

Despite growing documentation on the feminization of male fishes in waterways receiving municipal effluents, a critical question in the field of endocrine disruption research is whether these low-level, chronic exposures adversely impact wild populations (13). Although laboratory studies have shown decreased reproductive success of fish exposed to <1–5 ng·L⁻¹ of EE2 (14, 15), it is unknown whether this response would be observed in wild populations and whether it would result in a subsequent decline in abundances. To assess the ecological risk posed by this class of compounds, we must understand population-level effects of estrogens and their mimics on aquatic organisms.

The fathead minnow (*Pimephales promelas*) is a common species in North America, and its range extends from the southern United States to northern Canada (16). It is an important food source for numerous game fish species, such as lake trout (*Salvelinus namaycush*), walleye (*Sander vitreus*), and northern pike (*Esox lucius*). In the lakes used for this study, fathead minnow have a lifespan of \approx 4 years, but few individuals live past 2 years of age (17). Asynchronous spawning starts in early summer and extends for a period of \approx 2 months; multiple females will typically spawn in the nest of a single male, who will care for developing eggs until hatching. Sexual maturity occurs during the second year of life, so most fish spawn for only one season. Therefore, 2 consecutive years of reproductive failure cause catastrophic declines in abundance (18, 19). Fathead minnow have been widely adopted by the scientific community for toxicity testing because of their widespread distribution, ease of laboratory culture, well characterized biology and life history, and the large family (Cyprinidae) of fishes that they represent (20–22). It is a freshwater equivalent of the “miner's canary.”

We describe the results of a 7-year, whole-lake study at the Experimental Lakes Area (ELA) in northwestern Ontario, Canada, to assess the subcellular-level through population-level effects of the potent synthetic estrogen EE2 on fathead minnow. The concentrations of EE2 achieved in the experimental lake, Lake 260, during the 3 years of additions were within the range of those observed in untreated and treated municipal wastewaters (5, 6) and below the total E2 equivalents (with EE2 having

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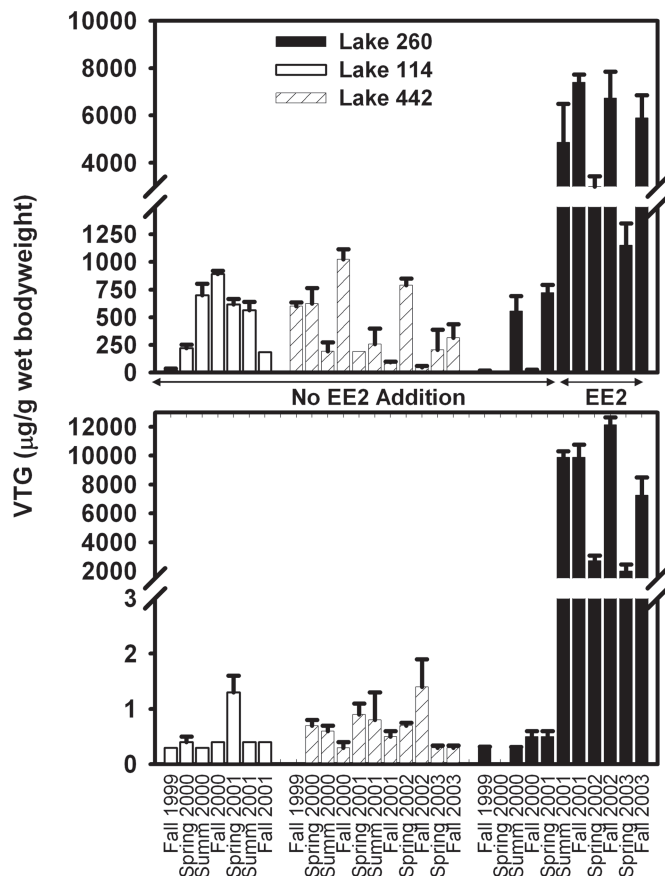
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Abbreviations: CPUE, catch-per-unit-effort; E2, 17 β -estradiol; EE2, 17 α -ethynylestradiol; VTG, vitellogenin.

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an E2 equivalent of 0.19–1.9; ref. 23) measured in effluents and receiving waters (10–12).

Results and Discussion

Measures of VTG mRNA and protein are used to assess exposure of fishes to estrogens and estrogen mimics, and elevated concentrations in males are indicative of the presence of these substances in effluents or surface waters (1, 3). We used these endpoints to assess the subcellular responses of fathead minnow from Lake 260 to EE2 additions. Seven weeks after the first estrogen additions to Lake 260 began in 2001, VTG concentrations in male and female fathead minnow were elevated when compared with preaddition or reference lake data (Fig. 1). Whole-body homogenates of males from Lake 260 had concentrations of VTG that were three orders of magnitude greater than reference samples, and this response was sustained in each of the 3 years of EE2 additions. Female fathead minnow also produced more VTG after exposure to EE2 when compared with reference samples (reference lake fish had 2.5% of the VTG concentrations observed in treated fish from Lake 260; Fig. 1), although their response was less dramatic than that observed for male fish. Liver VTG mRNA was measured in samples collected from the EE2-amended and reference lakes concurrently with those used for VTG protein analyses. Results from these mRNA analyses showed patterns similar to the VTG protein expression. Over the 3 years of dosing, male fathead minnow from Lake 260 had mean \pm SD normalized liver VTG mRNA values ranging from 0.422 ± 0.685 to 1.22 ± 0.181 . Liver VTG mRNA values

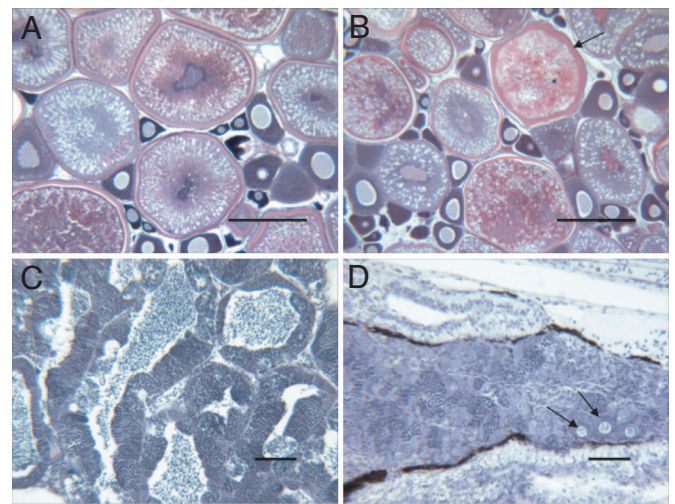


Fig. 2. Gonadal tissue sections from fathead minnow sampled in early May 2003. (A) Ovary from reference Lake 442 with small, dark, primary-stage oocytes situated between cortical alveolar-stage oocytes and large vitellogenic oocytes. (B) Ovary from EE2-amended Lake 260 demonstrating an atretic follicle (arrow). (C) Testis from reference Lake 442. (D) Testis from EE2-amended Lake 260 demonstrating intersex; arrows indicate primary-stage oocytes among the remnants of testicular tissue. (Scale bars: A, B, 300 μm ; C, D, 80 μm .)

for males from the two reference lakes were on average <0.1 – 1.6% of those for the EE2-exposed males. In addition, males from Lake 260 had VTG mRNA values (0.904 ± 0.437) that were more than an order of magnitude higher than female fish (0.045 ± 0.048) collected from the reference lakes during the same time periods. Females collected from Lake 260 in the spring and mid-summer of each year of EE2 additions had gene expression that ranged from 0.834 ± 0.879 to 0.378 ± 0.053 , and this response was 82–152% of the values in females from the reference lakes. These among-lake differences were highest in the fall; for example, females collected in 2002 from Lake 260 exhibited levels of VTG mRNA expression (0.849 ± 0.793) that were more than an order of magnitude higher than those of reference females (0.045 ± 0.048). This trend was also observed in the accompanying protein data, demonstrating that EE2 amendments prolonged the expression of VTG mRNA and protein in males and females well past the breeding season.

in females beyond the normal breeding season, impacts on gonadal development, as evidenced by intersex in males and altered oogenesis in females, and a near extinction of this species from the lake. Likely because of its short life-cycle, the fathead minnow was the first to show population collapse in this experiment. This response implies that short-lived fish species may generally be at greatest risk from exposure to estrogens and their mimics, but chronic exposure of longer-lived species to these substances may result in the loss of these populations as well.

Materials and Methods

The experimental system, Lake 260, is located in an undisturbed watershed at the Experimental Lakes Area, has a surface area of 34 ha and a maximum depth of 14 m, and contains naturally reproducing populations of lake trout, white sucker (*Catostomus commersoni*), pearl dace, and fathead minnow, as well as several other small-bodied fish species. We studied Lake 260 and two nearby reference systems, Lakes 114 and 442, for 2 years before EE2 additions, 3 years during the amendments to the experimental lake, and 2 years after the additions were discontinued. The physical, chemical, and biological characteristics of Lake 260 and the reference lakes are detailed in refs. 24 and 28.

We added EE2 (Schering AG, Berlin, Germany) three times weekly to Lake 260 starting in May 2001, after the lake had stratified, and additions were continuous during the open-water seasons of 2001–2003 for a period of 20–21 weeks. For each EE2 addition, the required mass [calculated by using its predetermined half-life in the water column (K.A.K., unpublished data), the volume of the epilimnion, and the mean concentration of EE2 measured the previous week] was dissolved in 30 ml of DIG-grade methanol (Caledon Laboratories, Georgetown, ON, Canada) and added to the propeller wash of the boat as it was driven in transects across the lake. Mean \pm SD epilimnetic concentrations of EE2 were quantified weekly by using replicate samples collected at each of five sites around the lake. EE2 was extracted onto preconditioned C-18 columns (Supelco, Oakville, ON, Canada), eluted with 100% DIG methanol, and quantified along with the internal standard testosterone by using standard RIA techniques and GC-MS confirmation for 10% of the samples. Additional details on extraction methods and EE2 concentrations have been described in refs. 24 and 28. Seasonal mean \pm SD EE2 concentrations in Lake 260 were 6.1 ± 2.8 ($n = 189$), 5.0 ± 1.8 ($n = 169$), and 4.8 ± 1.0 ($n = 170$) $\text{ng}\cdot\text{L}^{-1}$ in 2001–2003, respectively.

For VTG mRNA and protein analyses, fathead minnow were collected by using live-minnow traps in the spring and fall of 1999–2003 and in the summer of 2000 and 2001 from Lake 260 and two reference lakes ($n = 4$ –10 individuals per sampling date for each analysis). Each fish was handled according to Canadian

Animal Care protocols (approved by the Freshwater Institute Animal Care Committee) and individually euthanized in pH-buffered tricaine methanesulfonate ($250 \text{ mg}\cdot\text{L}^{-1}$). For VTG analyses, whole fish were placed in sterile plastic bags, immediately frozen on dry ice, and kept at -90°C until analyzed. Whole-body homogenates were used to quantify the VTG by using the indirect competitive ELISA methods detailed previously (24). Reagent VTG from fathead minnow was obtained from the Core Biomarker Facility (University of Florida, Gainesville, FL). The primary antibody used for the analysis was a mouse anti-carp VTG monoclonal antibody (ND-2D3; Biosense Laboratories, Bergen, Norway) raised against VTG from common carp (*Cyprinus carpio*). Fish captured for VTG mRNA analysis were euthanized and dissected immediately after capture, and their livers were placed in RNAlater (Ambion, Austin, TX) for transport to the laboratory. Liver VTG mRNA was measured by using quantitative real-time PCR and normalized to 18S ribosomal gene expression, an invariant ribosomal gene, by using methods detailed previously (29).

Impacts on fathead minnow gonadal development in Lake 260 were best examined in the spring of each year when gonads were mature and not affected by the asynchronous spawning season. Fish were collected from Lake 260 and the two reference lakes each spring from 1999 to 2003 as described above; whole fish were euthanized, immersed in Bouin's solution, and processed by using standard histological techniques as described previously (30).

Index trap netting (adults and young-of-the-year) and minnow traps (adults only) were used to examine changes in the fathead minnow population size and structure in the fall during each of the 7 years in both Lake 260 and reference Lake 442. Each September, at least three trap nets were set in each lake and fished from 7 to 33 consecutive days; nets were emptied every 2–3 days. Lake 260 and Lake 442 CPUE data (not shown) were also derived from fall minnow trapping (30 traps set per lake for 10–11 consecutive days and emptied daily) and compared with CPUE data from fall trap netting (shown in Fig. 3).

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