

## Effects-Based Monitoring of Bioactive Contaminants Associated with Municipal Wastewater Treatment Plant Effluent Discharge to the South Platte River

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## Introduction

- Over the last several years, contaminants of emerging concern (CECs), including pharmaceuticals, personal care products, and pesticides, as well as in vitro bioactivities have been detected in the South Platte River near Denver, Colorado, USA at concentrations and bioactivities among the highest in the nation (Bradley et al., 2017; Conley at al., 2017).
- However, given the lack of available biological effects data for most of the compounds detected, potential ecological risks associated with these contaminants are unclear.
- The present study builds upon several years of preliminary chemical monitoring research on the South Platte River as a basis for further investigation of estrogen receptor (ER), glucocorticoid receptor (GR), and peroxisome proliferator activated receptor gamma (PPARγ)-bioactive contaminants and potential responses elicited in caged fish.
- To assess the spatial and temporal distribution of CECs, surface water samples were collected and analyzed approximately bi-monthly throughout 2018 and 2019 at 11 sites along the South Platte River with varying proximities to a wastewater treatment plant (WWTP) and surrounding tributaries. Water samples were analyzed for a suite of pharmaceuticals, pesticides, steroid hormones, and wastewater indicators as well as screened for *in vitro* biological activities.
- To evaluate the potential bioavailability and significance of these biologically-active contaminants, adult fathead minnows (*Pimephales promelas*) were caged at six sites along the South Platte River at various proximities to a WWTP as well as in surrounding tributaries.

### **Objectives:**

- Assess the spatial and temporal distribution of CECs and ER, GR, and PPARy bioactivities associated with a historically bioactive WWTP effluent along the South Platte River.
- Utilize a caged fish system to determine the potential *in vivo* bioavailability of bioactive contaminants.

## Methods

### Water Samples

- Grab or 5 d composite water samples collected approximately every other month from May 2018 to October 2019 during the Spring to Fall seasons
- Solid phase extraction for in vitro bioassays and steroid measurements
- Analyzed for a suite of pharmaceuticals, pesticides, wastewater indicators, steroid hormones, and *in vitro* biological activities (e.g., ER, GR, and PPARy activities)

### Analytical Chemistry

- Instrumental determination of 252 analytes was conducted on the surface water samples (131 pharmaceuticals and personal care products, 69 pesticides, and 52 wastewater indicators) by the Region 8 EPA Laboratory (Denver, CO).
- Steroidal estrogen concentrations determined by the GLTED EPA Laboratory (Duluth, MN)

### In vitro Bioassays

- Multiplexed *in vitro* Trans-FACTORIAL<sup>TM</sup> bioassay used to evaluate human nuclear receptors (Attagene, RTP, NC)
- Estrogen receptor agonist assay (T47D-Kbluc; Wilson et al. 2004)
- Glucocorticoid receptor agonist assay (Indigo Biosciences, State College, PA)
- Peroxisome proliferator activated receptor-gamma assay (Indigo Biosciences)

Fish per cage $6M, 6F$ Exposure duration $5$ daysSamples collectedMucus (metab) Plasma (steroids) Muscle (QPCR) Adipose (QPCR) Liver (QPCR & metab) Gonad (QPCR & metab) $\circ$ Nuscle (QPCR) Adipose (QPCR) Liver (QPCR & metab) Gonad (QPCR & metab)Water samples $5$ d composite at each site $Gene Expression$ Plasma (steroids) Muscle (QPCR) Liver (QPCR & metab) Gonad (QPCR & metab)Water samples $5$ d composite at each site $GR-related:$ $\cdot vtg = VitellogeninData analysisGene Expression Data: (Analyzed using Statistica)\cdot Analyzed using a one-way ANOVA, followed by Duncan's orDunn's post-hoc testPORPA\gamma-related:\cdot acoxl = Acyl-coenzyme A oxidase 1 palmitoyl\cdot lpl = Lipoprotein lipase\cdot fabp3 = Fatty acid binding protein 3\cdot fabp1bl = Fatty acid binding protein 1b1\cdot pparg = peroxisome proliferator activated receptor$	Cages per site	2	<b>Caged Fish Endpoints</b>		
Tish per cagetow, orExposure duration5 daysSamples collectedMucus (metab)Plasma (steroids)Plasma (steroids)Muscle (QPCR)Adipose (QPCR)Liver (QPCR & metab)Gonad (QPCR & metab)Gonad (QPCR & metab)Gonad (QPCR & metab)Water samples5 d composite at each siteData analysisGRe-related:Gene Expression Data: (Analyzed using Statistica)•Analyzed using a one-way ANOVA, followed by Duncan's or Dunn's post-hoc test•Differences considered significant at $p < 0.05$ In vitro Bioassay Data: (Analyzed using GraphPad Prism):•Analyzed using a non-linear regression (log-agonist vs. response)	Fish species	Pimephales promelas			
Exposure duration5 daysSamples collectedMucus (metab) Plasma (steroids) Muscle (QPCR) Adipose (QPCR) Liver (QPCR & metab) Gonad (QPCR & metab)• Ouantitative polymerase chain reaction (QPCR): • Power SYBR Green RNA-to-Ct 1-step kit • Taqman RNA-to-Ct 1-step kitWater samples5 d composite at each site• $Taqman RNA-to-Ct 1-step kit$ <b>Data analysis</b> Gene Expression Data: (Analyzed using Statistica) • Analyzed using a one-way ANOVA, followed by Duncan's or Dunn's post-hoc test• $m^2cl = Glucocorticoid receptor• sgkl = Serum/glucocorticoid regulated kinase 1• nfkbiaa = Nuclear factor of kappa light polypeptideenhancer in B-cells inhibitor, alpha aPPARY-related:• acoxl = Acyl-coenzyme A oxidase 1 palmitoyl• lpl = Lipoprotein lipase• fabp3 = Fatty acid binding protein 3• fabp1bl = Fatty acid binding protein 1bl• pparg = peroxisome proliferator activated receptor$	Fish per cage	6M, 6F			
Samples collectedMucus (metab) Plasma (steroids) Muscle (QPCR) Adipose (QPCR) Liver (QPCR & metab) Gonad (QPCR & metab)Power SYBR Green RNA-to-Ct 1-step kitWater samples5 d composite at each site $Taqman RNA-to-Ct 1-step kit$ <b>Data analysis</b> Gene Expression Data: (Analyzed using Statistica) •Analyzed using a one-way ANOVA, followed by Duncan's or Dunn's post-hoc test $GR-related:$ • $nr3cl =$ Glucocorticoid receptor • $sgkl =$ Serum/glucocorticoid regulated kinase 1 • $nfkbiaa =$ Nuclear factor of kappa light polypeptide enhancer in B-cells inhibitor, alpha a <b>PPARy-related</b> : • $acoxl =$ Acyl-coenzyme A oxidase 1 palmitoyl • $lpl =$ Lipoprotein lipase • $fabp3 =$ Fatty acid binding protein 3 • $fabp1bl =$ Fatty acid binding protein 1b1 • $pparg =$ peroxisome proliferator activated receptor	Exposure duration	5 days	• • • • • • • • • • • • • • • • • • • •		
<ul> <li><i>nr3c1</i> = Glucocorticoid receptor</li> <li><i>sgk1</i> = Serum/glucocorticoid regulated kinase 1</li> <li><i>nfkbiaa</i> = Nuclear factor of kappa light polypeptide enhancer in B-cells inhibitor, alpha a</li> <li><i>PPARγ-related:</i></li> <li><i>acox1</i> = Acyl-coenzyme A oxidase 1 palmitoyl</li> <li><i>lpl</i> = Lipoprotein lipase</li> <li><i>fabp3</i> = Fatty acid binding protein 3</li> <li><i>fabp1b1</i> = Fatty acid binding protein 1b1</li> <li><i>pparg</i> = peroxisome proliferator activated receptor</li> </ul>	Samples collected	Plasma (steroids) Muscle (QPCR) Adipose (QPCR) Liver (QPCR & metab)	<ul> <li>Power SYBR Green RNA-to-Ct 1-step kit</li> <li>Taqman RNA-to-Ct 1-step kit</li> </ul> Gene Targets: <u>ER-related:</u>		
<ul> <li>Data analysis</li> <li>Gene Expression Data: (Analyzed using Statistica)</li> <li>Analyzed using a one-way ANOVA, followed by Duncan's or Dunn's post-hoc test</li> <li>Differences considered significant at p &lt; 0.05</li> <li>In vitro Bioassay Data: (Analyzed using GraphPad Prism):</li> <li>Analyzed using a non-linear regression (log-agonist vs. response)</li> <li>PARγ-related:</li> <li>acox1 = Acyl-coenzyme A oxidase 1 palmitoyl</li> <li>lpl = Lipoprotein lipase</li> <li>fabp1b1 = Fatty acid binding protein 3</li> <li>fabp1b1 = Fatty acid binding protein 1b1</li> <li>pparg = peroxisome proliferator activated receptor</li> </ul>	Water samples	5 d composite at each site	• $nr3cl = G$ lucocorticoid receptor		
• Significant response threshold = Mean + $3(SD)$ of the	Gene Expression Data •Analyzed using a one-wa Dunn's post-hoc test •Differences considered s <i>In vitro</i> Bioassay Data: •Analyzed using a non-lir response)	: (Analyzed using Statistica) ay ANOVA, followed by Duncan's or ignificant at $p < 0.05$ a (Analyzed using GraphPad Prism): hear regression (log-agonist vs.	<ul> <li><u>PPARγ-related</u>:</li> <li><i>acox1</i> = Acyl-coenzyme A oxidase 1 palmitoyl</li> <li><i>lpl</i> = Lipoprotein lipase</li> <li><i>fabp3</i> = Fatty acid binding protein 3</li> </ul>		

**U.S. Environmental Protection Agency** Office of Research and Development

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## Results

**Bioactivity Surveillance** 

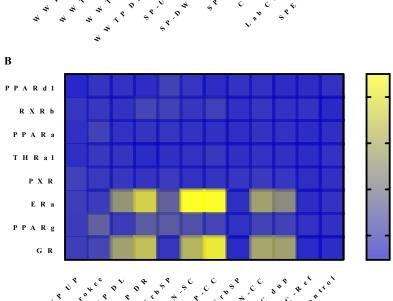


Figure 2. Results of the Attagene Trans-FACTORIAL<sup>TM</sup> assay, evaluating human nuclear receptor bioactivities in composite surface water extracts during caged fish exposures in A) August 2018; B) August 2019. Results are shown only for the endpoints that had a significant response and expressed as fold-change relative to background response. Yellow = higher activity; Blue=lower activity.

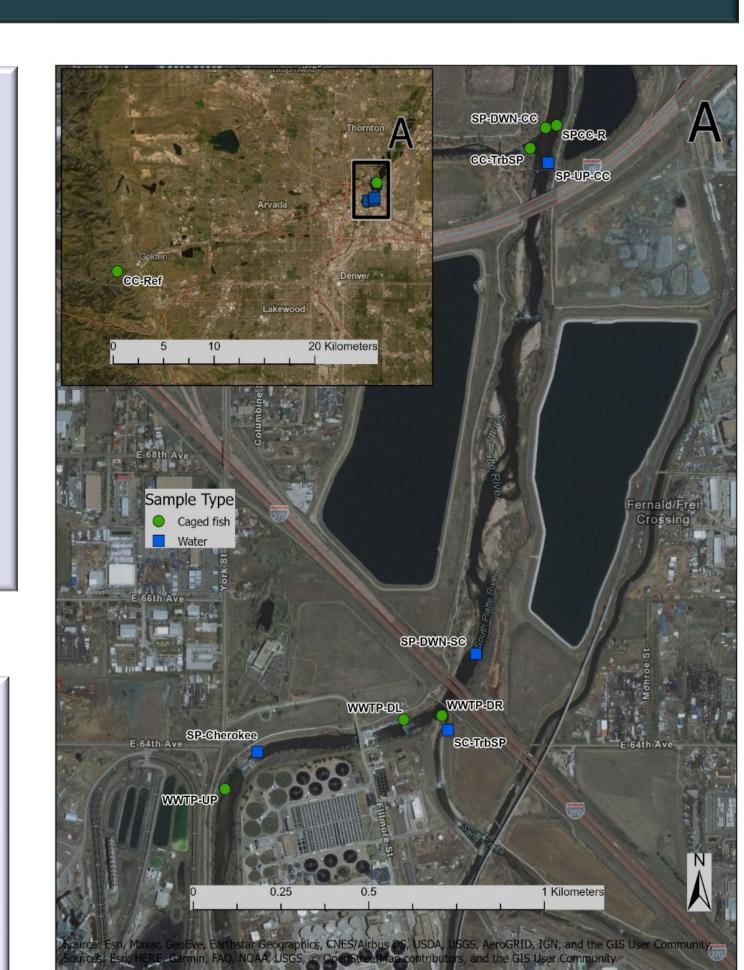
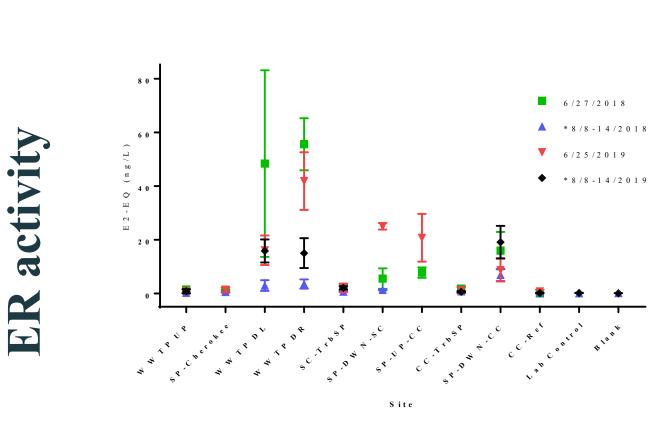


Figure 1. Map of field sites along the South Platte River and surrounding tributaries in Denver, CO.

	Site Name	Site Abbreviation
1	Clear Creek Reference	CC-Ref
2	WWTP Upstream	WWTP-UP
3	S. Platte Cherokee	SP-Cherokee
4	WWTP Downstream Left	WWTP-DL
5	WWTP Downstream Right	WWTP-DR
6	Sand Creek Tributary of S. Platte	SC-TrbSP
7	S. Platte Downstream of Sand Creek	SP-DWN-SC
8	S. Platte Upstream of Clear Creek	SP-UP-CC
9	Clear Creek Tributary of S. Platte (Caged fish 2019 only)	CC-TrbSP
10	S. Platte Downstream of Clear Creek	SP-DWN-CC
11	S. Platte Confluence with Clear Creek- Right Bank (Caged fish 2018 only)	SPCC-R



**Figure 3**. Estrogenic activity reported as estimated 17β-estradiol equivalents (E2-EQ) of grab and composite surface water sample extracts. Data represented by mean  $\pm$  SD of sample replicates within assays. \*Caged fish sample dates.

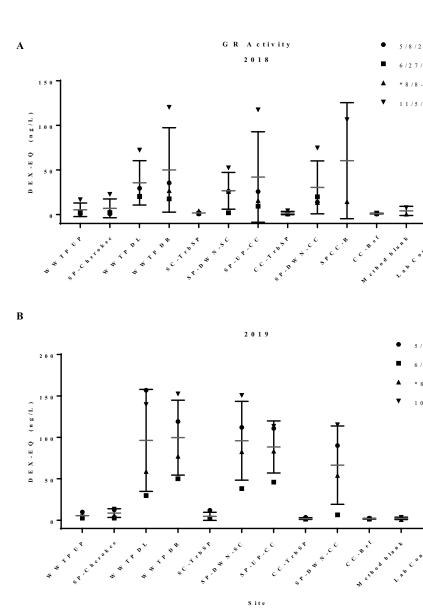


Figure 5. GR-activity of grab and composite surface water extracts reported as a dexamethasone equivalents (DEX-EQ). A) 2018 samples; B) 2019 samples. Data represented by mean  $\pm$  SD of sample replicates within assays. \*Caged fish sample dates.



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No significant in vitro response.

# Acknowledgements/References

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- 51: 4781-4791. Wilson, VS, Bobseine, K, Gray, L.E.J. 2004. Development and characterization of a cell line that stably expresses an estrogen-responsive luciferase reporter for the detection of estrogen
- agonist and antagonists. Toxicol Sci 81; 69-77.

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### In vitro response

• 5 / 8 / 2 0 1 8

6 / 2 7 / 2 0 1 8

▼ 11/5/2018

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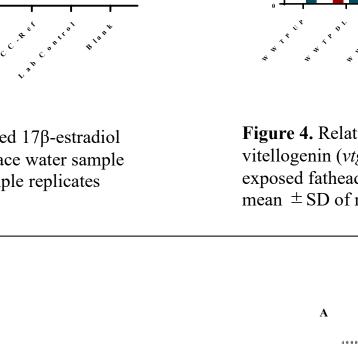
• 5/1/2019

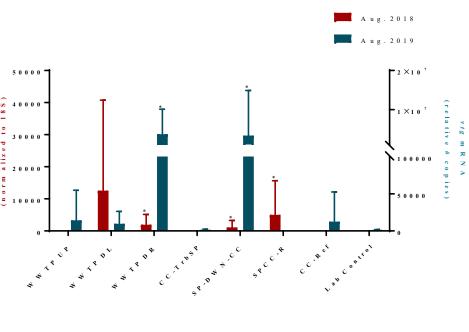
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In vivo response

Figure 4. Relative transcript abundance of vitellogenin (vtg) mRNA in male livers from fieldexposed fathead minnows. Data represented by mean  $\pm$  SD of n = 9-14 fish per treatment.

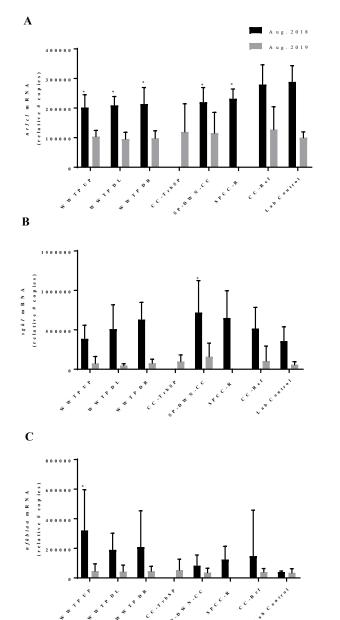
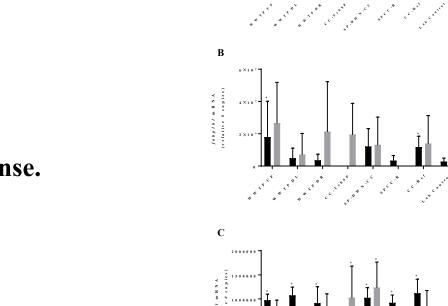


Figure 6. Relative abundance of predicted glucocorticoid receptor (GR)-related transcripts (*nr3c1*, *sgk1*, and *nfkbiaa*) in male livers from field-exposed fathead minnows. Data represented by mean  $\pm$  SD of n = 9-14 fish per treatment.

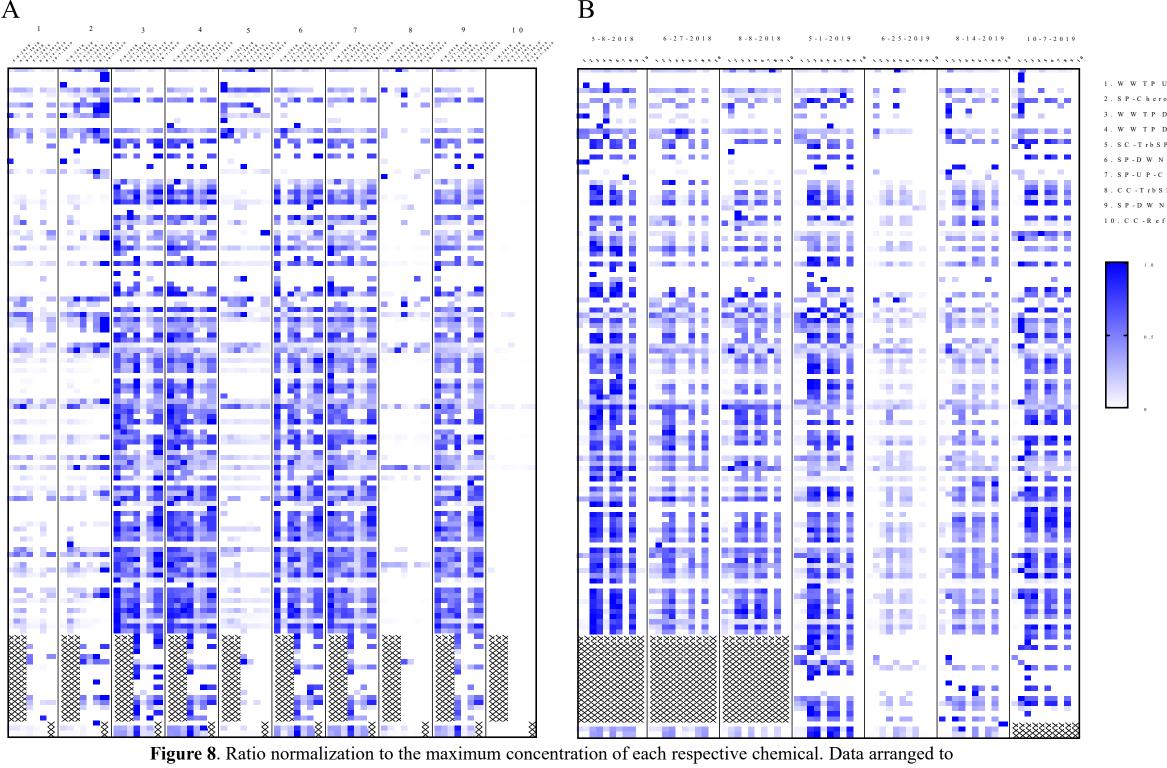


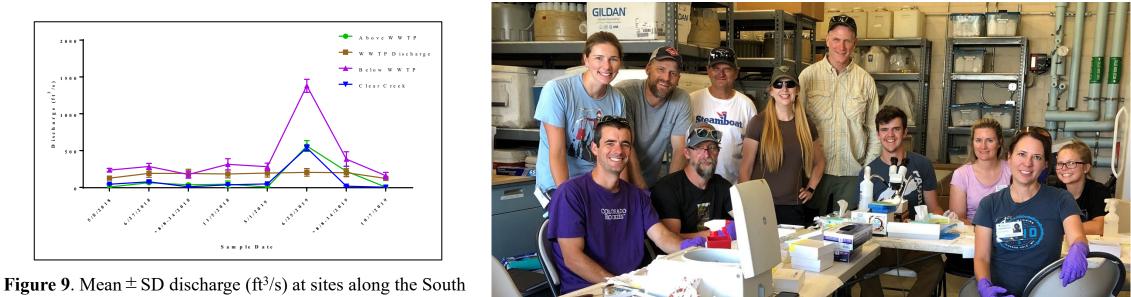
**Figure 7**. Relative abundance of predicted PPARγ-related transcripts (*pparg* 

*fabp1b1*, *acox1*, *fabp3*, and *lpl*) in male livers from field-exposed fathead minnows. Data represented by mean  $\pm$  SD of n = 9-14 fish per treatment.

Bradley et al., 2017. Expanded target-chemical analysis reveals extensive mixed-organic contaminant exposure in U.S. streams. *Environ Sci Technol* 51: 4792-4802. Conley et al., 2017. Occurrence and in vitro bioactivity of estrogen, and glucocorticoid compounds in a nationwide screen of United States stream waters. Environ Sci Technol

Future directions:





Platte River and Clear Creek on water sample collection dates.

## **Discussion and Conclusions**

Temporal and Spatial Variation of CECs and Bioactivities

- the WWTP (Fig. 8A).
- with the highest river discharge rates (Figs. 8B & 9).
- Fig. 3) and GR (up to 156  $\pm$  28 ng/L DEX-EQ; Fig. 5)

Caged Fish Studies: In vivo Bioavailability

- a significant gene expression response.
- in male livers (Fig. 7).

### **Conclusions:**

- convergence with additional tributaries.

• Develop a receptor pull-down method to identify unknown compounds that may be responsible for GR and PPARγ activity.

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## Analytical Chemistry

3. W W T P D L 4. W W T P D R . SC-TrbSP SP-DWN-S SP-DWN-C

highlight A) spatial and B) temporal distribution of the contaminants. Numbers correspond to the sites listed in the site key on the right. Dark blue = higher relative concentrations; white = lower relative concentrations.

Sites with the highest number and concentration of detected CECs were located within the South Platte River downstream of the WWTP. Fewer and lower concentrations of CECs were detected upstream of the WWTP and in surrounding tributaries with no direct impact of

Sampling dates with the highest number and concentrations of CECs were in the spring and fall seasons (5-8-2018, 5-1-2018 and 10-7-2019). The sampling date with the overall lowest concentrations of CECs across all sites was on 6-25-2019, which also corresponded

• In vitro multi-factorial Attagene assays detected estrogen receptor (ER), glucocorticoid receptor (GR), and peroxisome proliferator activated receptor-gamma (PPARγ) bioactivities in surface water samples downstream of the WWTP (Fig. 2).

• Targeted *in vitro* transcriptional activation assays for ER, GR, and PPAR $\gamma$  corroborated bioactivities for ER (up to 55 ± 10 ng/L E2-EQ;

There was no significant PPARy activity detected for any of the sites during the bi-monthly sampling events using the Indigo PPARy kits (results not shown). Relative to PPARy activation, the Attagene assay appears to be more sensitive than the Indigo Biosciences kit.

• Significant up-regulation of male hepatic *vtg* was observed in fish exposed at sites with corresponding *in vitro* ER activity (Fig.4).

• Despite GR-activity *in vitro*, no site-related differences in GR-related gene expression were detected in male livers, suggesting observed environmental concentrations of GR-active contaminants do not induce a detectable in vivo response (Fig. 6).

Genes selected for GR-related expression were verified to be responsive to the well-known GR agonist, Dexamethasone, in a controlled laboratory study (data not shown), suggesting river concentrations associated with the 5 d caged fish exposures were insufficient to elicit

Consistent with the general lack of detectable in vitro PPARy activity, there were no significant effects on PPARy-related gene expression

Although the chemicals responsible for in vitro GR- and PPARy-mediated bioactivities are still unknown, results from the present study provide insights into the significance of these bioactivities relative to in situ fish exposures.

• Overall, both GR- and ER-related biological activities can be detected more than 1.5 km downstream of the WWTP outflow, even after

There was no clear evidence for GR- or PPARy-mediated responses in fish caged at various locations along the South Platte River with respect to proximity to the wastewater discharge. However, wastewater was clearly a major contributor of contaminant loading to the system and additional research to evaluate the possible significance of exposures to up to 150 ng DEX-EQ/L of GR-mediated activity under laboratory conditions is warranted.