

MDPI





Figure S1. Quantitative and confirmation SRMs chromatographs of target EDCs and internal and surrogate standards. A, Quantitative SRM chromatographs with compound identity noted as follows: 1: E3 ($287 \rightarrow 145$), 2:PRDN ($358 \rightarrow 327$), 3:PRNL ($359 \rightarrow 329$), 4: BPA-d₁₆($241 \rightarrow 227$), 5:BPA($227 \rightarrow 212$), 6:DES($267 \rightarrow 252$), 7:EE($295 \rightarrow 145$), 8:E2($271 \rightarrow 145$), 9:E1($269 \rightarrow 145$), 10:Eq($268 \rightarrow 143$), 11:E1-d₄($273 \rightarrow 147$), 12:Gem-d₆/50($255 \rightarrow 121$), 13: 4OP/10($205 \rightarrow 106$), 14:4NP-d₄/10($223 \rightarrow 110$), 15:4NP/10($219 \rightarrow 106$)]. B, Confirmation SRM Chromatographs. Compounds identity and selected SRM noted as follows: 1, E3 ($287 \rightarrow 171$); 2, PRDN ($327 \rightarrow 300$); 3, PRDN ($358 \rightarrow 328$); 4:PRNL ($359 \rightarrow 259$); 5:PRDL/10($359 \rightarrow 359$) 6:BPA ($227 \rightarrow 133$), 7:DES/20($267 \rightarrow 2237$), 8:DES/20($267 \rightarrow 222$), 9:EE/10($295 \rightarrow 295$), 10:EE ($295 \rightarrow 195$), 11:E2($271 \rightarrow 183$), 12:EE/10($271 \rightarrow 271$), 13:E1($269 \rightarrow 143$), 14:E1($269 \rightarrow 183$), 15:4OP/65($205 \rightarrow 205$), 15:4NP/40($219 \rightarrow 219$). For some transitions, the abundance (Y-scale response) is divided by factor of 10, 40, 50 or 65 to show response of all analytes in the chromatogram as noted after compound identity. SRM monitored noted in brackets.



Time (min)

Figure S2. The selected reaction monitoring chromatograms for BPA and internal standard BPA-d₁₆ for raw and treated wastewater in comparison with similar levels of standard. All chromatograms obtained on the same day of analysis. Chromatograms A, C, E are obtained at selected reaction monitoring transition 227→212 (quantitative SRM for BPA). Chromatograms B, C, F are obtained at selected reaction monitoring transition 241→227 (quantitative SRM for BPA-d₁₆). Sample matrix: Raw wastewater A and B; treated wastewater C and D; and 10 ng/mL BPA standard with BPA-d₁₆ E and F. All samples contain BPA-d₁₆ as an internal standard





Table S1. Linear regression equation of the best- fit curve and the R² values for all the analytes from WWTP3 for quantitative SRM.

Analyte	Y=mx+b(r ²) Solvent-	Y=mx+b(r²) standard	Y=mx+b(r²) standard	Y=mx+b(r²) standard	Y=mx+b(r²) standard
	based calibration	addition with upstream	addition with raw	addition with treated	addition with
		water WWTP3 as	wastewater from	wastewater from	downstream water
		matrix	WWTP3 as matrix	WWTP3 as matrix	WWTP3 as matrix
BPA	Y=0.0846x-0.0877(0.997)	Y=0.0193x-0.0304(0.987)	Y=0.0156x+0.1285(0.997)	Y=0.0095x+0.0508(0.997)	Y=0.0142x-0.0877(0.997)
4OP	Y=0.5314x+0.4466(0.996)	Y=0.2108x+1.5498(0.997)	Y=0.0029x-0.0231(0.995)	Y=0.2725x+1.5498(0.991)	Y=0.3684x+0.2613(0.986)
4NP	Y=0.2175x+0.4182(0.990)	Y=0.1019x+0.0082(0.997)	Y=0.0105x+0.0329(0.992)	Y=0.1445x+0.4265(0.998)	Y=0.1511x+0.1252(0.999)
E1	Y=0.0272x-0.0228(0.999)	Y=0.0201x-0.0492(0.997)	Y=0.0174x-0.0709(0.973)	Y=0.0174x-0.0709(0.973)	Y=0.0158x-0.0286(0.990)
E2	Y=0.0015x+0.002(0.999)	Y=0.0009x+0.0039(0.998)	Y=0.0017x-0.0036(0.990)	Y=0.0005x+0.0033(0.940)	Y=0.0012x-0.0104(0.963)
E3	Y=0.0045x-0.0061(0.991)	Y=0.0011x+0.0008(0.982)	Y=0.0028x+0.0924(0.979)	Y=0.0009x-0.0076(0.985)	Y=0.0009x+0.0007(0.992)
EE	Y=0.0032x+0.0034(0.997)	Y=0.0007x+0.0011(0.982)	Y=0.0025x-0.0352(0.956)	Y=0.0021x-0.0368(0.992)	Y=0.0006x+0.0006(0.993)
DES	Y=0.0231x-0.0144(0.997)	Y=0.0994x+0.0568(0.999)	Y=0.0885x-0.0591(0.968)	Y=0.0736x+0.0012(0.996)	Y=0.0865x+0.0804(0.999)
PRDN	Y=0.0077x-0.0464(0.996)	Y=0.0108x-0.1122(0.995)	Y=0.0021x+0.0389(0.985)	Y=0.0058x-0.1241(0.980)	Y=0.0062x-0.0023(0.999)
PRNL	Y=0.0163x-0.0869(0.989)	Y=0.0421x-0.1217(0.992)	Y=0.0013x+0.0138(0.989)	Y=0.0304x+0.129(0.996)	Y=0.0178x-0.0149(0.990)