1	Supporting information
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3 4	Method development for effect-directed analysis of endocrine disrupting compounds in human amniotic fluid
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26	
27	15 pages containing 7 figures and 2 tables

### 28 S1. – Chemicals

29  $\alpha$ -Minimal Essential Medium (MEM) with GlutaMax (32561-029), Dulbecco's Modified 30 Eagle Medium/Nutrient Mixture F-12 medium (DMEM/F12) with GlutaMax (31331-028), 31 DMEM/F12 medium without phenol red (11880-028), Leibovitz's L-15 medium (L-15, 11415-32 049), L-15 medium without phenol red (21083-027), Hank's Balanced Salt Solution (HBSS, 33 14175-053), penicillin/streptomycin (P/S, 15140-122), trypsin (25300-054) and foetal calf 34 serum (FCS) were purchased from Gibco (Eggenstein, Germany); G418 (ant-gn-1, ant-gn-5, 35 10832-42-2) was obtained from InvivoGen (Toulouse, France); Steady-Glo® Luciferase Assay 36 system (E2550) was purchased from Promega (Leiden, The Netherlands). Methanol (MeOH) 37 and dichloromethane (DCM) were purchased from Thermo Fisher Scientific (Loughborough, 38 UK); acetone was obtained from BDH Laboratory (Poole, Dorset, UK); dimethyl sulfoxide 39 (DMSO, 99.9+%) was obtained from Alfa Aestar (MA, USA). All solvents used were of HPLC 40 grade. Ammonium hydroxide solution (ACS reagent, 28.0-30.0% NH3 basis) was purchased 41 from Sigma Aldrich (Poole, Dorset, UK); LC-MS grade water (LiChrosolv®) was obtained 42 from Merck KGaA (Darmstadt, Germany); chloroform (ACS reagent, 99.8+%) was purchased 43 from Acros Organics (Geel, Belgium). Two hydroxylated metabolites of brominated flame 44 retardants, namely 6-hydroxy-2,2',4,4'-tetrabromodiphenyl ether (6-OH-BDE-47) and 3-45 hydroxy-2,2',4,4',5,5'-tetrabromodiphenyl ether (3-OH-BDE-153) were generously provided 46 by Dr. Lillemor Asplund (Stockholm University). All other reference compounds were 47 purchased from Sigma Aldrich (Poole, Dorset, UK) at the highest commercially available 48 purity. 4-methyl-2,4-bis(4-hydroxyl- phenyl) pent-1-ene (MBP), a potent metabolite of 49 bisphenol A, was synthesized and its identity and purity was confirmed spectrometrically 50 (NMR, IR and HR-MS) as described below.

#### S1.1 – Synthesis of 4,4'-(4-methylpent-1-ene-2,4-diyl)diphenol (MBP)

52 Materials and Methods: NMR spectra were recorded on a Bruker Avance 500 (125.78 MHz 53 for <sup>13</sup>C) using the residual solvent as internal standard (<sup>1</sup>H:  $\delta$  2.05 ppm, <sup>13</sup>C{<sup>1</sup>H}:  $\delta$  206.26 ppm 54 for acetone-d<sub>6</sub>. Chemical shifts ( $\delta$ ) are given in ppm and coupling constants (*J*) are given in 55 Hertz (Hz). Melting points were recorded on a Büchi M-565 melting point apparatus. 56 Electrospray Ionization (ESI) high resolution mass spectroscopy was carried out using a Bruker 57 micrOTOF-Q instrument in negative ion mode. Infrared spectra were recorded neat using a 58 Shimadzu FTIR-8400s spectrophotometer and wavelengths are reported in cm<sup>-1</sup>. Flash 59 chromatography was performed on Silicycle Silia-P Flash Silica Gel (particle size 40-63 μm, 60 pore diameter 60 Å) using the indicated eluent. Thin layer chromatographic (TLC) analysis 61 was performed with Merck F254 silica gel-60 and visualized by UV light.

62 4-Methyl-2,4-bis-(p-hydroxyphenyl)pent-1-ene (MBP): Bisphenol A (2.00 g, 8.76 mmol) was 63 dissolved in concentrated sulfuric acid (6.86 mL) and the orange solution was poured over ice 64 water (116 mL) under vigorous stirring. The resulting solids were collected by filtration. The 65 filtrate was washed with MTBE (3 x 20 mL) and the previously obtained solids were dissolved 66 in the combined organic layers. Traces of sulfuric acid in the organic layer were neutralized 67 with saturated aqueous NaHCO<sub>3</sub>, after which the organic layer was dried over Na<sub>2</sub>SO<sub>4</sub>, filtered 68 and concentrated in vacuo. The residue was purified by flash silica gel chromatography 69 (gradient, CH<sub>2</sub>Cl<sub>2</sub>:EtOAc 97.5:2.5  $\rightarrow$  90:10) to give MBP (235 mg, 20 %) as a white solid.



#### 70 Figure S1.1 - Proton NMR spectrum of synthesized MBP

EJA284 Pure

71 72 <sup>1</sup>H NMR: (500 MHz, Acetone-d<sub>6</sub>)  $\delta$  8.16 (br s, 2H), 7.19 (ddd, 4H, J = 11.0 Hz, 9.0 Hz, 2.5 Hz), 6.74 (ddd, 4H, 4H, 5.1 Hz), 6.74 (ddd, 5.1 Hz), 6.74 (ddd, 5.1 Hz), 6.74 (dddd, 5.1 Hz), 6.74 (dddd, 5.1 Hz), 6.74 (ddddd, 5.1 Hz), 6.74 (ddddd 73 *J* = 11.0 Hz, 9.0 Hz, 2.0 Hz), 5.05 (d, 1H, *J* = 2.5 Hz), 4.68 (s, 1H), 2.78 (s, 2H), 1.17 (s, 6H);



74 Figure S1.2 - <sup>13</sup>C NMR spectrum of synthesized MBP

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 <sup>75</sup>
 <sup>13</sup>C NMR: (125 MHz, Acetone-d<sub>6</sub>) δ 157.5, 155.9, 147.4, 141.4, 135.7, 128.4 (2C), 127.6 (2C), 115.7 (2C), 115.3 (2C), 115.0, 50.1, 38.6, 29.4 (2C);



78 Figure S1.3 - Infra Red (IR) spectrum of synthesized MBP

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80 IR (neat) vmax (cm<sup>-1</sup>): 3265 (s), 1610 (m), 1512 (s), 1437 (m), 1366 (m), 1221 (s), 1175 (s), 895 (m), 829 (s), 555 (s);

# 81 Figure S1.4. - HRMS (ESI) spectrum of synthesized MBP

			Mass	Spectru	um Sma	artForn	nula R	eport			
Analysis Info Analysis Name Method Sample Name Comment		D:\Data\HRMS2015\hrms1215\Elwin_1_01_13914.d caffeine_neg_elwin2.m Elwin EJJA284 Attempt 2						Acquisition Date Operator Instrument / Ser#	12/14/2015 8:49:12 AM Marek Smoluch micrOTOF-Q 32		
Acquisition F Source Type Focus Scan Begin Scan End	Parameter	ESI Not active 50 m/z 3000 m/z	lon F Set Set	Polarity Capillary End Plate Offse Collision Cell R	t	Negative 4500 V -500 V 200.0 Vpp		Set Nebulizer Set Dry Heater Set Dry Gas Set Divert Valve	1.6 B 210 ° 8.0 M Sour	iar 'C min ce	
Intens. x10 <sup>5</sup> 3 2 1	133.0661		267.1	403				433 2041	-M:	567,2773	18-11
100 Meas. m/z 133.0661	150 Formula C 18 H 18 O 2	200 m/z  err  [ppm : 133.0659 1.5	250 ] err [mDa] 9 -0.3	300 rdb N-Rui 10.0 c	3 e e <sup></sup> Conf k even	50 mSigma 120.9	400 # Sigma 1	450	500	550	'n

HRMS (ESI): m/z calculated for C<sub>18</sub>H<sub>19</sub>O<sub>2</sub> [M-H]<sup>-</sup> 267.1391, found 267.1403, C<sub>18</sub>H<sub>18</sub>O<sub>2</sub> [M-2H]<sup>2-</sup> 133.0659, found 133.0661; melting point 127.3-129.5 °C.

**Table S1.** Concentration of the referencecompounds in the spike mix used during methoddevelopment of extraction and fractionation ofamniotic fluid.

Compound	Concentration in the spike mix (µg/mL)
4-OP	0.788
PP	0.394
MnBP	0.708
BPA	0.570
BPS	0.679
MBP	0.336
TCS	0.423
OH-PCB-61	0.729
PFOS	0.263
60H-BDE-47	0.378
30H-BDE-157	0.741
daidzein	0.705
genistein	0.188
enterolactone	0.308
FICZ	0.345
E2	0.693
E1	0.292
E4	0.797

#### 91 S2 - Extraction procedures

92 Solid phase extraction (SPE): MCX columns (Oasis, 6 cc, 150 mg/3 mL, 30 µm particle 93 size, Waters Corp., Milford, MA) and HLB columns (Oasis, 6 cc, 150 mg, 30 µm particle size, 94 Waters Corp., Milford, MA) were conditioned with 3 mL of methanol and 3 mL of deionised 95 water. Matrix (5ml of LC-MS grade water or AF) was spiked with the reference mix (100  $\mu$ L), 96 pH was adjusted to 2 with 98% formic acid (for MCX extraction only), and then samples were 97 loaded on the cartridges at a rate of 1 mL/min. The MCX and HLB cartridges were rinsed with 98 3 mL of 2% formic acid (in deionized water) and with 5% methanol respectively, and then air 99 dried for 1 min. The analytes were eluted from the MCX cartridges with 2 x 2 mL of 100% 100 methanol and 2 x 2 ml of 5% of ammonium hydroxide, and from the HLB cartridges with 4 x 101 2ml of 100% methanol. Deionised water (0.2 mL) was added to the extracts and the extracts 102 put in a water bath (max. temperature 40°C) where the solvent was evaporated under a gentle 103 stream of nitrogen to a final volume of 0.2 mL. The samples were then reconstituted with 0.3 104 mL of 10% methanol in deionised water, transferred to glass vials for analysis and analysed on 105 LC-ESI-MS/MS for chemical recoveries.

106 Dispersive liquid/liquid extraction (DLLE): 5 ml of LC-MS grade water or amniotic 107 fluid was spiked with the reference mix (100  $\mu$ L) and pH adjusted to ~4.5 with sodium 108 acetate/acetic acid buffer (0.1 M). DLLE was performed by addition of DCM (0.5 mL) or 109 chloroform (0.5 mL) only, or by addition of a mixture of acetone (1ml) (dispersive solvent) 110 and DCM (0.5 mL) or chloroform (0.5 mL) (extraction solvent). Samples were then vortexed 111 2 x for 15 seconds and centrifuged at 4081 x g for 15 min at 25°C. The supernatant was 112 removed, 50 µL of deionised water was added, the organic layer evaporated under gentle 113 stream of nitrogen and the residue reconstituted with 10% MeOH to a final volume of 0.5 mL. 114 Samples were transferred to glass vials and analysed on LC-ESI-MS/MS for chemical 115 recoveries.

116 Table S2. Parameters used during LC-ESI-MS/MS Multiple Reaction Monitoring (MRM) detection of the collision cell parameters and monitored precursor/product ions. Limit of detection (LOD)

1	1	7	analytes	inc	ludin	g
		'	unurytes	me	ruum	5

118 for each analyte was calculated as described in the method section.

Compound	Q1	Q3a	Q3b	DT	EP	DP (Q/C)	CE (Q/C)	CXP (Q/C)	LOD (M)
4-OP	205.18	106.0	189.0	25	-10	-55	-28	-15	4.4 x 10 <sup>-10</sup>
PP	179.13	91.9	136.0	25	-10	-55	-32	-13	6.0 x 10 <sup>-11</sup>
MnBP	221.05	121.1	70.9	25	-12	-80	-20	-13	2.0. x 10 <sup>-10</sup>
BPA	227.20	133.1	212.0	25	-11	-90/-90	-40/-26	-13/-13	3.7 x 10 <sup>-09</sup>
BPS	249.10	107.9	156.0	25	-10	-70	-38	-17	3.7 x 10 <sup>-11</sup>
MBP	267.20	133.1	117.2	25	-11	-100/-150	-25/-50	-13/-15	9.4 x 10 <sup>-12</sup>
TCS	287.09	34.9	241.1	25	-10	-40	-48	-7	5.4 x 10 <sup>-11</sup>
OH-PCB-61	306.95	270.8	34.9	25	-10	-55/-55	-30/-82	-19/-17	9.9 x 10 <sup>-11</sup>
PFOS	498.95	79.9	98.9	25	-10	-170	-84	-17	9.9 x 10 <sup>-12</sup>
60H-BDE-47	500.75	78.8	80.8	25	-10	-75	-40	-11	2.7 x 10 <sup>-11</sup>
30H-BDE-157	658.55	80.8	576.6	25	-10	-105/-105	-100/-38	-13/-41	5.8 x10 <sup>-12</sup>
daidzein	253.14	132.0	223.0	25	-10	-90/-90	-54/-36	-7/-13	1.7 x 10 <sup>-11</sup>
genistein	269.10	133.1	63.1	25	-10	-135/-135	-46/-50	-11/-11	2.9 x 10 <sup>-11</sup>
enterolactone	297.16	253.0	107.0	25	-10	-110/-150	-28/-50	-17/-15	2.7 x 10 <sup>-11</sup>
FICZ	283.30	254.2	255.3	25	-10	-90	-40	-13	2.1 x 10 <sup>-11</sup>
E2	271.15	183.0	145.0	25	-10	-80/-50	-56/-56	-11/-17	4.1 x 10 <sup>-11</sup>
E1	269.12	145.0	142.9	25	-10	-105	-52	-23	1.6 x 10 <sup>-11</sup>
E4	303.12	241.0	273.0	25	-14	-90	-28	-15	3.8 x 10 <sup>-11</sup>

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Q1, precursor ion (m/z); Q3a, quantification product ion (m/z); Q3b, confirmation product ion (m/z); DT, dwell time (ms); DP, declustering potential (V); EP, entrance potential (V); CE, collision energy (V); CXP, collision cell exit potential (V)

### 122 S3 – Cell culture

123 The estrogen receptor-mediated luciferase reporter gene (ER-Luc) human breast 124 carcinoma (VM7Luc4E2) cells<sup>1</sup>, the androgen/glucocorticoid receptor-mediated luciferase 125 reporter gene (AR-Luc) human breast carcinoma (MDA-kb2) cells<sup>2</sup> and the aryl hydrocarbon 126 receptor enhanced green fluorescent protein reporter gene (DR-GFP) mouse hepatoma (H1G1.1c3) cells<sup>3</sup> were routinely maintained in 75 cm<sup>2</sup> canted neck tissue culture flasks 127 128 (Greiner Bio-One Ltd.). ER-Luc and DR-GFP cells were maintained in DMEM/F12 and  $\alpha$ -129 MEM medium respectively, supplemented with 10% FCS and 1% P/S (culture media) and 130 incubated in a humidified incubator at 37 °C and 5% CO<sub>2</sub>. DR-GFP cells were additionally 131 supplemented with 1% G418. AR-Luc cells were maintained in Leibowitz-15 (L-15) medium 132 supplemented with 10% FCS (culture media) and incubated in a humidified incubator at 37 °C 133 without additional CO<sub>2</sub>. Cells were subcultured when 70-80% confluent.

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## 135 ER, AR and DR reporter gene assays

136 ER-Luc: Prior to experiments, ER-Luc cells were maintained in DMEM/F12 medium 137 without phenol red (assay meadia) supplemented with 10% charcoal dextran stripped FCS 138 (cdFCS) and 1% P/S for two media-changes (4 days). Cells were seeded at a density of  $2 \times 10^4$ 139 cells per well in 100 µL of assay media in clear-bottomed 96-well tissue culture dishes (Greiner 140 Bio-One Ltd.) and allowed to attach. After 24 h media was removed and 100  $\mu$ L of assay media 141 containing test chemicals or DMSO control (0.5%) was added to the wells. After 24h of 142 exposure, media was removed, cells washed with PBS (200 µL), PBS removed and lysis buffer 143 added (20  $\mu$ L). After cell lysis (15 min with shaking) the luciferase activity was determined in 144 a plate reader (LUMIstar Optima, BMG Labtech GmbH) with 100 µL of luciferase reagent 145 media injected directly to each well during measurement.

146 AR-Luc: Prior to experiments, AR-Luc cells were maintained in L-15 medium 147 supplemented with 10% cdFCS (pre-assay media) for two media-changes (4 days). Cells were 148 then seeded at a density of  $2x10^4$  cells per well in 100 µL of phenol red-free L-15 medium supplemented with 10% cdFCS (assay media) into white, clear-bottomed 96-well tissue culture 149 150 dishes and allowed to attach. After 24 h, 100 µL assay medium was added to the wells with 151 twice the intended final concentration of test chemicals or DMSO control (0.5%), to obtain the 152 desired test concentrations. After 24 h of exposure, 100 µL of media was removed and 153 SteadyGlo assay reagent (Promega UK Ltd.) (100  $\mu$ L) was added to the wells and the plate left 154 shaking for 10 min. The luciferase activity was determined as described above.

155 DR-GFP: DR-Luc cells were maintained in culture media and when confluent seeded 156 in black, clear-bottomed 96-well tissue culture dishes (DR-Luc) (Gre- iner Bio-One Ltd.) in 157 MEM Alpha media supplemented with 10% FCS, 1% P/S, but without addition of G418 (assay media). Cells were seeded at a density of  $3 \times 10^4$  cells per well in 100 µL of assay media and 158 159 left to attach. After 24 h, media was aspirated and cells exposed to test chemical or DMSO 160 control (1%) in 100  $\mu$ L of assay media. After 24 h of exposure, fluorescence was measured in 161 the intact cells (without removal of media) using a fluorescent plate reader (POLARstar 162 Galaxy, BMG Labtech GmbH) with an excitation wavelength and an emission wavelength set 163 at 485 nm and 510 nm, respectively. In order to normalize results the fluorescence gain was 164 adjusted per plate to the highest activity of the positive control. In each assay the outer wells 165 of the plate were not included in the measurements and were filled with 100  $\mu$ L assay media 166 only.

Cell viability was determined with 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl 167 168 tetrazolium bromide (MTT). Briefly, after 23 h exposure, media was removed from each well, 169 100 µl of 5 mg/ml freshly made MTT solution was added and plates were incubated at 37 °C. 170 At t= 24 h, MTT solution was removed, lysis buffer (150 uL of isopropanol) was added to the 172 with a POLARstar fluorescent plate reader.

<sup>171</sup> wells and the plate left shaking for 15 min. The absorbance at ( $\lambda ex = 590$  nm) was measured



**Figure S2.** The log*P* of reference compounds retrieved from PubChem database (www.pubchem.ncbi.nlm.nih.gov) plotted against retention times (RT) during RP-HPLC fractionation of spiked AF extract, including estetrol (E4), 17 $\beta$ -estradiol (E2) estrone (E1) and bisphenol A (BPA). Log*P* of estriol (E3), androstenedione (ASD), dihydrotestosterone (DHT) and testosterone was plotted against their predicted RTs interpolated from linear regression analysis (GraphPad Software Inc., San Diego, CA).



Figure S3. An example of a chromatogram obtained during reversed-phase fractionation of AF extract.
Sample was injected onto reverse phase Phenomenex Kinetex C18 (100mm x 2.1 mm, 5 μm pore size)
column using an Agilent 1260 system equipped with diode array detector (DAD). Absorbance was
monitored with a 253 nm wavelenght and plotted against retention time. Eluted fractions were
concentrated, resuspended in DMSO and used for bioassay analysis.



198199Figure S4. Agonistic activity of AF fractions (50 x diluted) in the ER-Luc assay. ER activity was200measured as described in the method section (n=1).

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