

Supporting Information to the Article:

Comparison of polycyclic aromatic hydrocarbon uptake pathways and risk assessment of vegetables from waste-water irrigated areas in northern China

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Material and methods

Sample extraction and cleanup

Soil samples were extracted by accelerated solvent extraction (ASE) (Dionex ASE 300). Ten grams of soil samples blended with 15g anhydrous sodium sulfate were extracted in 34 mL stainless steel vessels. Extractions were performed with 1:1 mixture of dichloromethane and acetone (pesticide grade, Fisher Company, USA) at 120 °C and 1500 psi for a 5 min heat-up followed by a 10 min static extraction. The vessels were then rinsed with 17 mL of the same solvent and the extracts were purged from the sample cell using pressurized nitrogen at 1500 psi.

The ASE extracts were concentrated using a vacuum rotary evaporator (R-201, Shanghai Shenshen) at 38°C before being transferred to a silica gel/alumina column for sample cleanup. A glass column (20 mm outside diameter, 450 mm length) was first filled with 120 mm of silica gel (pre-rinsed with n-hexane), then packed with 60 mm of alumina (pre-rinsed in n-hexane). 20 mm of anhydrous sodium sulfate was added to the top of alumina. During the process, n-hexane was added to ensure the column remained soaked. Columns were pre-eluted with 15 mL of hexane, followed by introduction of the extract and an elution of 70 mL of methylene chloride/hexane (v/v=3:7). All elutions were carried out at 2 mL/min. The eluates collected from the silica column were concentrated to 1.0 mL with a rotary evaporator and nitrogen blowdown (SHE812, Beijing Shuaien). The samples were sealed in vials and stored at -18 °C before analysis. The vegetable samples were subjected to similar extraction, involving circa 0.2 g dried shoot samples and circa 0.4 g root samples were used. Solvent blanks were analyzed within each batch.

Sample analysis

Soil and vegetable extracts were analyzed by an Agilent 7890 GC equipped with an Agilent 5975 mass spectrometer under the selected ion monitoring mode (SIM). Total 16 PAHs including naphthalene (Nap), acenaphthylene (Any), acenaphthene (Ace), fluorene (Fle), phenanthrene (Phe), anthracene (Ant), pyrene (Pyr), fluoranthene (Flu), benz[a]anthracene (BaA), chrysene (CHR), benzo[b]fluoranthene (BbF), benzo[k]fluoranthene (BkF), benzo[a]pyrene (BaP), indeno[1,2,3-cd]pyrene (IncdP), benzo[ghi]perylene (BghiP) and dibenzo[a,h]anthracene (DahA) were determined in all samples. A 30 mm×0.25 mm i.d.×0.25 mm film thickness HP 5MS capillary column (Agilent Technology) was used with helium as the carrier gas at a constant flow rate of 1 mL / min. Splitless injection of 1 µL of the sample was conducted with a 7683 autosampler (Agilent Technology). The GC oven temperature was programmed from 50 °C (2 min) to 200 °C (2 min) at 20 °C /min, then to 240 °C (2 min) at 5

°C min before reaching 290 °C at 3 °C min and held for 15 min. The injector and detector temperatures were 280 °C and 300 °C, respectively.

PAHs in the samples were identified by the retention time and the abundance of quantification ions/confirmation ions with respect to authentic PAH standards. Automated library searching was performed using the National Institute of Standards and Technology (NIST) Mass Spectral Database. Quantization was performed using the five-point calibration curve for individual components. Detection limits were 1.5-3.6 $\mu\text{g kg}^{-1}$ dry weight for PAHs. Results were expressed on a dry weight basis.

Table S1. Abraham Solvent Descriptors, Experimental and Predicted Log Dimensionless Octanol-Water (Log Kow) and Air-Water Partition Coefficients (log Kaw) for the EPA-16 PAHs

	E	L	V ^a	A	B	S	log K _{ow} (-) 25°C	log K _{aw} (-) 25°C
Nap	1.340 ^b	5.161 ^d	1.085	0.00 ^b	0.20 ^b	0.92 ^b	3.33 ^g	-1.73 ^g
Any	1.750 ^c	6.175 ^c	1.216	0.00 ^c	0.20 ^f	1.14 ^c	3.84 ^l	-2.20 ^m
Ace	1.604 ^d	6.469 ^c	1.259	0.00 ^d	0.20 ^d	1.04 ^d	3.94 ^g	-2.63 ^j
Fle	1.588 ^d	6.922 ^c	1.357	0.00 ^d	0.20 ^d	1.03 ^d	4.68 ^g	-2.57 ^g
Phe	2.055 ^b	7.632 ^c	1.454	0.00 ^b	0.29 ^b	1.29 ^b	4.57 ^g	-2.92 ^g
Ant	2.290 ^b	7.568 ^c	1.454	0.00 ^b	0.28 ^b	1.34 ^b	4.63 ^h	-3.34 ^m
Flu	2.377 ^e	8.827 ^c	1.585	0.00 ^e	0.20 ^e	1.55 ^e	5.23 ^g	-3.35 ^g
Pyr	2.808 ^d	8.833 ^c	1.585	0.00 ^d	0.29 ^d	1.71 ^d	4.79 ⁱ	-4.38 ^g
BaA	2.992 ^b	10.291 ^c	1.823	0.00 ^b	0.35 ^b	1.70 ^b	5.91 ^h	-4.68 ^j
Chr	3.027 ^b	10.334 ^c	1.823	0.00 ^b	0.36 ^b	1.73 ^b	5.91 ^g	-4.56 ^j
BbF	3.194 ^c	11.609 ^c	1.954	0.00 ^c	0.44 ^f	1.82 ^c	5.81 ^g	-4.68 ^g
BkF	3.194 ^c	11.607 ^c	1.954	0.00 ^c	0.44 ^f	1.91 ^c	6.11 ^g	-4.75 ^g
BaP	3.625 ^e	11.715 ^c	1.954	0.00 ^e	0.44 ^e	1.98 ^e	6.13 ^g	-5.58 ^g
IcdP	3.610 ^c	12.690 ^c	2.084	0.00 ^f	0.44 ^f	1.90 ^f	6.64 ^l	-4.92 ^k
DahA	4.000 ^c	12.960 ^c	2.192	0.00 ^c	0.46 ^f	2.04 ^c	7.06 ^l	-5.59 ^m
BghiP	4.073 ^e	13.264 ^c	2.084	0.00 ^e	0.46 ^e	1.90 ^e	6.22 ^g	-5.26 ^g

a) Abraham and McGowan, 1987; b) Torres-Lapasio et al., 2004; c) Abraham, 1993; d) Abraham et al., 1994a; e) Abraham et al., 1994b; f) assumed from homologues; g) De Maagd et al., 1998; h) Yalkowsky and Valvani, 1979; i) Paschke et al., 1999; j) Bamford et al., 1999; k) Tenhulscher et al., 1992; l) Predicted using the Abraham descriptors and the relevant equation in (Goss, 2005); m) Predicted using the Abraham descriptors and the relevant equation in (Mintz et al., 2007)

Table S2 BCFs of vegetable shoot and root

type	PAH	Radish	Cauliflower	Chinese cabbage	Rape	Leek	Leaf lettuce
shoot	Nap	2.43	3.14	3.75	2.94	2.2	4.7
	Any	0.54	0.91	0.81	0.32	0.8	0.5
	Ace	1.11	2.72	1.55	1.24	0.95	1.4
	Fle	2.27	3.69	4.6	2.47	1.62	2.8
	Phe	2.6	3.23	3.87	3.89	2.23	4.1
	Ant	1.51	1.79	2.09	1.17	0.93	1.5
	Flu	0.76	0.96	2.16	0.67	0.62	0.9
	Pyr	0.63	0.64	1.56	2.19	0.74	1.0
	BaA	0.43	0.62	0.47	0.33	0.36	2.0
	Chr	0.46	0.72	0.73	0.51	0.82	1.7

	BbF	0	0.06	0	0	0	0.0
	BkF	0	0.19	0	0	0	0.0
	BaP	0.2	0.19	0.61	0.55	0.15	0.4
	IcdP	0.14	0.11	0.23	0.49	0	0.6
	DahA	0.06	0.18	0.85	0.68	0	1.1
	BghiP	0.82	0.39	0.63	1.08	0.22	0.9
root	Nap	1.21	1.73	2.34	1.09	1.39	4.2
	Any	1.27	1.61	2.02	1.68	1.85	6.6
	Ace	3.66	4.75	4.22	3.84	2.28	5.5
	Fle	3.74	5.63	4.86	3.71	4.75	5.6
	Phe	1.64	2.14	1.18	2.12	2.3	3.1
	Ant	1.88	1.41	1.78	1.89	0.37	4.1
	Flu	0.66	0.98	2.06	1.26	0.35	0.7
	Pyr	0.81	1.13	2.73	2	0.47	0.9
	BaA	0.75	0.72	0.91	1.11	0.46	2.1
	Chr	1.58	2.04	2.1	3.3	2.38	4.0
	BbF	0.23	0	0.25	0.18	0.16	0.1
	BkF	0.12	0	0.09	0.11	0.06	0.3
	BaP	0.53	0.32	0.45	0.41	0.29	0.4
	IcdP	0.2	0.25	0.15	0.15	0.2	0.8
	DahA	0.11	0	0.02	0.1	0.16	0.0
	BghiP	1.05	0.8	0.94	0.92	0.21	0.3

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