CONffIDENCE: Contaminants in food and feed: Inexpensive detection for control of exposure

Multiplex Screening of Persistent Organic Pollutants in Fish Using Spectrally Encoded Microspheres

Anastasia Meimaridou^a, Kamila Kalachova^b, Weilin L. Shelver^c, Milan Franek^d, Jana Pulkrabova^b, Willem Haasnoot ^a, Michel W.F. Nielen ^{a,e}



WP1a NCE

Figure 1: The 3 different microbead sets coated with the protein conjugates Figure 1: the 3 different microbead sets coated with the protein conjugates of BaP-BSA, PCB7-OVA and PBDE47-BSA are combined to capture the mixture of a monoclonal antibody against BaP (MabBaP) and the two polyclonal antibodies against the PCB77 (PAbPCB77) and PBDE47 (PabPBDE47). The binding of the mixture of antibodies to the mixture of the three different coated microbeads was inhibited by the presence of the respective free POPs in the fish extract. Detection and quantification of the immune complexes were obtained via combined in a frame dum antibidie noise to many a public incomplexes in the fish a combination of secondary antibodies against mouse or rabbit immunoglobulin (IgG) conjugated to the fluorescent protein R-phycoerythrin (PE).



3-plex FCIA to fish samples & Aroclors

Target POPs	Matrix	Goodness of the 4P R ² *	Curve steepness (ming ⁻¹) ⁶	IC ₅₀ ppb in the 3plex FCIA ^c	IC _{so} ppb in ELISAs
PCB77 (A)	Buffer	0.9968	-0.8	20±2	2-15 34
	Fish extract	0.9927	-0.6	55±5	Not measured
BDE47 (B)	Buffer	0.9992	-0.7	2±0.1	0.135 30
.,	Fish extract	0.9902	-0.7	2±0.4	Not measured
BaP (C)	Buffer	0.9857	-1.3	0.4±0.1	0.3 32
(0)	Fish extract	0.9435	-1	4±0.5	Not measured



Each point represents the average of six replicates \pm SD.

Table 2: Contaminated fish samples with their fat contents, levels of the target POP's (BaP, PCB77 and BDE47), as measured with GC-MS, and the percentages of inhibition of the maximum responses as was measured in the 3-plex FCIA with (+) or without (-) the clean-up

Fishes	Target POPs measured	Fat content %	µgkg-1 as measured in GC-MS	Clean- up	% of inhibition of maximum response in 3plexFCIA
Smoked trout	BaP	10	0.06	•	5 0 5
Smoked trout	BaP	11	1	-	80±2
Smoked trout	BaP	14	5	-	80±5
Smoked trout	BaP	13	14.7	-	80±3
Trout	PCBs/BDEs	2	n.d.	+	0±0.1
Chub	BDE47	1.5	0.43	+	45±2
Chub	BDE47	2	4.93	+	56±5
Chub	BDE47	2	9	+	50±4
Chub	PCB77	1.5	1.95	-	22±2

RIKILT

WAGENINGENUR



Figure 4: Relative inhibition (B/B0) of the maximum MFIs (B0) caused by the addition of 1000 ppb of Aroclor 1232, 1242 and 1248 respectively, to buffer (µg L-1) (A) and fish (µg kg-1) (B), applying a concentration step of 2.5 (2.5 g of fish mL-1 of extract), as measured in the 3-plex FCIA. The different Aroclors spiked fish samples were extracted with (+) or without (-) elean-up using the simplified extraction procedure described in this paper. paper.

USDA

RIKILT - Institute of Food Safety, Wageningen UR, P.O. Box 230, 6700 AE Wageningen, the Netherlands Fax: +31 317 417717. E-n
^b Department of Food Chemistry and Analysis, Institute of Chemical Technology Prague, Technická 3, 166 28 Prague 6, Czech Repul
^c USDA-ARS Biosciences Research Laboratory, P.O. Box 5674, Fargo, ND 58105, USA
^d Department of Biotechnology, Veterinary Research Institute, Hudeova 70, 621 32 Brano, Czech Republic
^e Laboratory of Organic Chemistry, Wageningen University, Dreijenplein 8, 6703 HB Wageningen, the Netherlands

The research leading to these results has received funding from the European Community's Seventh Framework Programme (FP7/2007-2013) under grant agreement n 211326: CONffIDENCE project



f	
ure	
eu ich	





Figure 3: Dose-res se curves obtained with the 3- plex FCIA in buffer Figure 3: Dose-response curves obtained with the 5-plex FCIA in buffer (ϕ) and fish-extract ($\mathbf{0}$) for the three main POPs representatives analysed: this current study: (A) PCB77 (fish extract (-) clean-up), (B) BDE47 (fish extract (+) clean-up) and (C) BaP (fish extract (-) clean-up), The relative binding (B/B0) was calculated by dividing the MFI response (B) of each concentration by the MFI obtained in a solution without the analyte (B0). analysed in Solid lines show curves fitted with the four-parameters (4P) model. Each point represents the average of six replicates \pm SD.