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## Effects of water accommodated fractions of crude oils and diesel on a suite of biomarkers in Atlantic cod (*Gadus morhua*)

T.F. Holth<sup>a,\*</sup>, D.P. Eidsvoll<sup>a</sup>, E. Farnen<sup>b</sup>, M.B. Sanders<sup>c</sup>, C. Martínez-Gómez<sup>d</sup>, H. Budzinski<sup>e</sup>,  
T. Burgeot<sup>f</sup>, L. Guilhermino<sup>g</sup>, K. Hylland<sup>a</sup>

<sup>a</sup> Department of Biosciences, University of Oslo, PO Box 1066 Blindern, N-0316 Oslo, Norway

<sup>b</sup> Norwegian Institute for Water Research (NIVA), Gaustadaléen 21, N-0349 Oslo, Norway

<sup>c</sup> CEFAS Weymouth Laboratory, The Nothe, Barrack Road, Weymouth, Dorset DT4 8UB, England

<sup>d</sup> Instituto Español de Oceanografía, San Pedro del Pinatar, 30740 Varadero 1, Spain

<sup>e</sup> ISM, Université Bordeaux 1, 351 crs de la Libération, 33405 Talence, France

<sup>f</sup> IFREMER, Unit of research in Biogeochemistry and Ecotoxicology, Rue de l'Île d'Yeu, BP 21105, 44311 Nantes, France

<sup>g</sup> CIIMAR - Interdisciplinary Centre of Marine and Environmental Research, Laboratory of Ecotoxicology and Ecology & ICBAS-Institute of Biomedical Sciences of Abel Salazar, Department of Population Studies, Laboratory of Ecotoxicology, University of Porto, Rua dos Bragas 289, 4050-123 Porto, Portugal

\*: Corresponding author : T.F. Holth, tel.: +4722855600 ; fax: +4722854726 ; email address : [t.f.holth@ibv.uio.no](mailto:t.f.holth@ibv.uio.no)

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### Abstract:

The aim of this study was to characterise concentration- and time-dependent responses in juvenile Atlantic cod (*Gadus morhua*) following exposure for one and three weeks to the water-soluble fraction (WAF) of three weathered oils: Arabian Light crude oil (ALC), North Sea crude oil (NSC) and ship-diesel. The sum of polycyclic aromatic hydrocarbons (PAH) in water was highest after one week of exposure and within environmentally relevant concentrations. PAH metabolites in bile confirmed exposure to and uptake of PAHs. Hepatic cytochrome P450 1A (CYP1A) gene expression (here and later the phrase gene expression is used synonymously to gene transcription, although it is acknowledged that gene expression is also regulated by, e.g., translation and protein stability) increased dramatically following exposure to all three oil types (fold-change up to 165), and there was a time lag between gene and protein expression. Hepatic CYP1A protein concentration and ethoxyresorufin-O-deethylase (EROD) activity were however more variable among individuals and treatments than gene expression. Hepatic and gill EROD increased in fish exposed to WAF from the two crude oils, but not in fish exposed to WAF from diesel. Exposure to diesel appeared to induce oxidative stress to a larger extent than exposure to crude oils. Other biomarkers (glutathione S-transferases, acetylcholine esterase, vitellogenin) did not appear to respond to the exposure and hence did not discriminate among oils. Biomarker responses in cod after exposure to weathered crude oils and diesel suggested the CYP1A system and oxidative stress markers to have the highest potential for discriminating among different oil types and to monitor the environmental consequences of spills.

**Keywords** : Atlantic cod ; Biomarkers ; Weathering ; crude oil ; ship-diesel ; water accommodated fraction ; time-course

## 1 Introduction

A range of oils with different characteristics is transported in European waters, all of which may be released to the environment as spills. Since the properties of the oils differ, it is important to understand how they affect marine organisms. A fingerprint of different sublethal responses in marine organisms for individual oils can be used to monitor the immediate effects of a spill and identify the oil causing the effects. Discharges of crude and refined oils to sea have the potential for long-term impacts on populations of marine organisms (e.g. Jewett et al., 2002; Monson et al., 2011). Polycyclic aromatic hydrocarbons (PAHs), metals, alkylphenols and naphthenic acids are some of the groups of compounds present in most oil types known to induce adverse effects on marine organisms and populations (e.g. Kavanagh et al., 2012; Meier et al., 2007; Myers et al., 2003; Vieira et al., 2009). Although the effects of accidental oil spills on marine ecosystems have been assessed for decades, the number of established biological effect methods (biomarkers) and knowledge on their sensitivity and time course development to different oil types is still limited (Martínez-Gómez et al., 2010). Following some of the largest accidental oil tanker spills in European waters (“Haven”, “Braer”, “Sea Empress”, “Erika” and “Prestige”), the use of biomarkers included assessment of PAH metabolite formation, genotoxicity, biomarker gene expression (here and later the phrase gene expression is used synonymously to gene transcription, although it is acknowledged that gene expression is also regulated by, e.g., translation and protein stability), lysosomal membrane integrity, cytochrome P450 1A (CYP1A) protein quantity and

1 ethoxyresorufin-O-deethylase (EROD) activity, antioxidant enzyme activities as well  
2 as histopathology in a variety of fish species (Amat et al., 2006; Budzinski et al.,  
3 2004; Claireaux et al., 2004; Harvey et al., 1999; Kirby et al., 1999; Marigómez et al.,  
4 2006; Martínez-Gómez et al., 2009; Pietrapiana et al., 2002; Raingeard et al., 2009;  
5 Stagg et al., 2000; Viarengo et al., 2007). These studies have shown that biomarkers  
6 are valuable tools to quantify the severity of pollution impacts on organisms and  
7 populations, but also that there are challenges in interpreting biomarker responses. For  
8 example, the potential differences in biomarker responses to oils from different  
9 geographical regions are not well known, although such knowledge could assist in  
10 selecting the optimal biomarkers to be used in post-spill monitoring (Martínez-Gómez  
11 et al., 2010).

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The intensity and specificity of biological responses may differ among oil types due to differences in oil composition. Even within a defined geographical region such as the North Sea, oils and their water-accommodated fractions (WAFs) may differ substantially in their composition of most toxic substances due to natural causes (e.g. PAHs, Utvik, 1999). Weathering processes may further modify oil composition and toxicity following discharge into the environment. For example, photoenhanced toxicity of crude and bunker oil has been observed for Pacific herring (*Clupea pallasii*) development (Barron et al., 2003; Incardona et al., 2012). The speed and efficacy of such degradation processes may depend on oil type, but also on intensity of irradiation, mechanical breakdown and dispersion (waves, wind, gravel), ambient temperature and rate of biotransformation by microorganisms (Sauer et al., 1998). Weathering does however not always decrease oil toxicity; weathered oils generally have altered ratios of higher to lower molecular weight PAHs since the lighter PAHs are more liable to evaporate or be used as energy source by microorganisms (Neff et

1 al., 2000; Wang and Fingas, 1995). Therefore, portions of an oil spill may remain  
2 bioactive for prolonged periods of time. Other common confounding factors forcing a  
3 complex interpretation of effects may be biological and ecological differences among  
4 sentinel species; differences in genetic expression, enzymatic activities and levels of  
5 other biomarkers in distinct tissues; seasonal differences; poorly known time-scale of  
6 biomarker responses and/or adaptation and recovery processes of the selected species;  
7 background/natural contamination as well as climatic conditions (e.g. Claireaux et al.,  
8 2004; Lee and Anderson, 2005; Stagg et al., 2000; Viarengo et al., 2007).

19 Some PAH compounds are known to affect well-characterised  
20 biotransformation systems in fish, such as CYP1A. Dioxin-like PAHs may activate  
21 the aryl hydrocarbon receptor (AhR) pathway, causing an increase in CYP1A mRNA  
22 and protein (see e.g. Goksøyr and Förlin, 1992). Furthermore, ligands of the AhR may  
23 also induce CYP1A enzymatic activity, generally measured as EROD activity (Whyte  
24 et al., 2000). In addition to indicating the presence of dioxin-like substances,  
25 induction of CYP1A activity may increase bioactivation of some PAHs causing  
26 embryotoxicity and genotoxicity (Myers et al., 2003; Hodson et al., 2007). Aneugenic  
27 and/or clastogenic DNA damage may also result in the formation of micronuclei  
28 (Jaylet et al., 1986). Quantification of micronuclei has been successfully used for  
29 decades as a biomarker for exposure to genotoxic compounds in both marine and  
30 freshwater species (Bolognesi and Hayashi, 2011).

48 Several known (and unknown) substances in oil WAFs may affect the  
49 endocrine system of fish (Thomas et al., 2004b; Thomas et al., 2009). Some PAHs,  
50 alkylphenols or their metabolites have been shown to have estrogenic activity  
51 (Martínez-Gómez et al., 2013), to induce vitellogenin protein synthesis in male fish  
52 (Sherry et al., 1999), and to delay gonadal development (Holth et al., 2010; Meier et  
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al., 2007). PAHs and redox-active trace metals are also known to cause oxidative stress in tissues, which may lead to macromolecule inactivation and breakdown (Almeida et al., 2012; Van Der Oost et al., 2003). Oxidative stress may be monitored using several different techniques, of which the most common is monitoring of lipid peroxidation levels (thiobarbituric acid reactive substances; TBARS) (Hermes-Lima et al., 1995), and measuring the activities of antioxidant enzymes, e.g. glutathione *S*-transferases (GST; Van Der Oost et al., 2003). Furthermore, WAFs from oil may contain compounds that inhibit acetylcholine esterase (AChE) (Holth and Tollefsen, 2012), an enzyme associated with the hydrolysis of acetylcholine in the synaptic cleft. Reduced AChE activity may lead to acetylcholine accumulation, continuous nerve stimulation, ataxia and death, and this activity has been extensively used as a biomarker for organophosphorous insecticide exposure in both marine fish and invertebrates (Costa, 2006; Fulton and Key, 2001), but has also been shown to be affected by other environmental contaminants such as some PAHs and metals (Roche et al., 2002; Vieira et al., 2009).

In the current study, we aimed to assess how WAFs from three weathered oil types would affect a suite of biomarkers in Atlantic cod (*Gadus morhua*). Juvenile Atlantic cod were exposed for one and three weeks to WAFs from three commonly transported oil types in Europe: Arabian Light crude oil, North Sea crude oil and ship-diesel. Tissues were sampled and analysed for PAH metabolites in bile, CYP1A gene expression and protein quantity in liver, EROD activity in liver and gills, lipid peroxidation levels (TBARS) and GST activity in gills, AChE activity in muscle, vitellogenin concentration in plasma and the frequency of micronuclei in erythrocytes.

## 2 Materials & Methods

### 2.1 Chemicals and weathering of oils

The oils, Arabian Light crude oil (ALC), North Sea crude oil (NSC) and marine gas oil (ship-diesel), were selected based on their differences in geographic origin and composition of main groups of hydrocarbons (figure 1). Details of oil composition were presented in Radović et al. (2012), in which the diesel used in this work was diesel B. Artificial weathering was performed according to the method described by Smith et al. (2005): the oils were mixed with seawater (1:1 v v<sup>-1</sup>) and stirred using a magnetic stirrer in a fume hood for 24 hours at room temperature. Following a resting phase, water and oil was separated. The weathered oils were mixed with clean gravel (4-5 mm diameter) at concentrations of 2 and 6 grams oil kg<sup>-1</sup> gravel (dichloromethane was used as solvent). Treated gravel was air dried in a fume hood for 24 hours at room temperature, frozen at -20°C and stored until use. Gravel used for the control group was treated similarly, including solvent addition, evaporation and freezing. Unless otherwise mentioned, all chemicals were purchased from Sigma Aldrich (St. Louis, Oregon, USA).

### 2.2 Fish and experimental setup

Juvenile Atlantic cod were exposed in 4 replicates to two doses of WAFs from three different oils in addition to a seawater control. Cod were purchased from Profunda AS (Barstadvik, Norway). Cod length and weight were  $17.8 \pm 0.2$  cm and  $47.6 \pm 1.3$  g (average  $\pm$  SE). Following transportation, cod juveniles were acclimated to system water for at least two weeks before introduction into the exposure system. Cod juveniles were fed with 0.5 grams of boiled commercial Faeroese shrimps per fish per day throughout the study, except for three consecutive days before the samplings to

1 promote concentration of bile. System water intake was at 34 meters depth from the  
2 outer Oslofjord, with an average temperature of 10.2°C and salinity of 32.0‰. A total  
3 of 56 fish were distributed into 28 tanks (i.e. two fish per tank) giving one control  
4 group and six exposed groups, each of 4 replicate tanks (figure 2). Tank volumes were  
5 20 L and the nominal flow rate was 50 mL min<sup>-1</sup> tank<sup>-1</sup>, equivalent to 3.6 water  
6 volume exchanges per day and a fish loading rate of 1.3 g L<sup>-1</sup> day<sup>-1</sup>. Oxygen  
7 concentration was monitored throughout the exposure period (>5 mg L<sup>-1</sup>). Tanks were  
8 covered with lids and separated by blinds in the interspaces. Weathered oil WAFs  
9 were produced by allowing seawater to percolate through columns containing oil-  
10 covered gravel (Carls et al., 1999). Clean and soiled gravel (650 g) were introduced  
11 into their respective columns (d=75 mm, h=350 mm) and flow was applied 24 hours  
12 before the experiment started to reduce eventual initial elevated concentrations of  
13 volatile organic compounds (VOC) or oil droplets. All tubings were made of Teflon™  
14 and tanks were of a whole glass construction with no silicone seams. The experiment  
15 was run continuously for 3 weeks. There was no mortality during the experiment  
16 period.

### 2.3 Tissue sampling

23 One cod was sampled from each tank one and three weeks after the start of exposures.  
24 In addition, eight fish were sampled prior to the start of the exposure to indicate the  
25 background levels of the biomarkers being measured (pre-exposure control samples).  
26 All fish were anaesthetised by brief immersion in MS-222 (Sigma-Aldrich AS; 0.5 g  
27 L<sup>-1</sup>) and sacrificed by a blow to the head. Length and weight were recorded and blood  
28 was collected from the caudal vein using a heparinised syringe. A drop of blood was  
29 immediately prepared for micronucleus assessment. The remainder of the blood  
30 sample was centrifuged (2,000 x g for 5 minutes) to separate blood cells and plasma,  
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1 which were frozen in separate tubes in liquid nitrogen. Two gill arches were excised:  
2 one gill arch was immediately assessed for EROD activity, the other was snap frozen  
3 in liquid nitrogen. The cod were dissected, the gall bladder excised and frozen at -  
4 20°C. The liver was weighed, split into replicates and frozen on liquid nitrogen. A  
5 muscle sample was excised from the area below the first dorsal fin and immediately  
6 snap-frozen on liquid nitrogen. The gonads were not sufficiently developed for visual  
7 sex determination.  
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#### 10 **2.4 Monitoring of PAHs and VOCs in water and PAH metabolites in cod bile**

11 Water samples (100 mL) were collected in amber glass flasks from each tank one and  
12 three weeks after the start of exposures, frozen at -20°C and stored until analysis  
13 (approx. 1 year). Before use, the flasks were washed 3 times in acetone and  
14 autoclaved. Teflon caps were used.  
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17 Water samples were analysed for PAHs according to a method adapted from the one  
18 described in De Perre et al. (2014). Briefly, SPME (solid phase micro-extraction)  
19 analyses of 9 mL water samples (in 10 mL vials) were performed with commercially  
20 available PDMS (polydimethylsiloxane 100 µm) fibers from Supelco (Bellefonte,  
21 USA). After the immersion of the fiber in the water sample (60 minutes; 30°C; 250  
22 rpm), it was immediately desorbed into the GC-MS injection port (5 min; 270°C).  
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25 Water samples were analysed for VOC by headspace SPME analyses. Ten mL of  
26 water was put in 20 mL vials and extraction performed in the headspace part of the  
27 vial with commercially available PDMS (30 min; 50°C; 250 rpm). The PDMS fiber  
28 was immediately desorbed into the GC-MS injection port (10 min; 220°C; pulsed  
29 splitless). The GC flow was 1.3 mL min<sup>-1</sup> (helium 6.0 constant flow) and the  
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1 temperature program was as follows: start at 20°C (10 min); increase to 200°C (5  
2 min) at 10 °C min<sup>-1</sup> and finally to 300°C (9 min) at 25°C min<sup>-1</sup>.  
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4 Analyses of PAHs and VOCs in water were performed in automated mode using a  
5 Gerstel MPS2XL (RIC, Belgium) and the GC was an Agilent 7890A model (Agilent  
6 Technologies, France) equipped with a 5975C mass selective detector, operated with  
7 an energy of ionization of 70 eV (electronic impact). The column used was an HP-  
8 5MS-UI ((5%-phenyl)-methylpolysiloxane; 30 m × 0.25 mm i.d.; 0.25 µm film;  
9 Agilent Technologies, France).  
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11 For the determination of PAHs and VOCs, the mass spectrometer was operated in the  
12 Selected Ion Monitoring (SIM) acquisition mode and the molecular ions were sought  
13 (PAHs: table S1; VOCs: table S2). Quantification was performed using internal  
14 perdeuterated standards (for PAHs at least one per aromatic class) added prior to  
15 SPME extraction in the sample directly by gravimetry (few µL of ethanolic solution;  
16 table S1 and S2 for PAHs and VOCs, respectively). The response factors of the  
17 different compounds were measured by injecting a solution of unlabelled standards  
18 mixed with deuterated compounds used as surrogate standards (determined at  
19 different levels of concentrations).  
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21 The procedural limits of detection (LODs) and quantification (LOQs) were  
22 determined by analysis of spiked milli-Q (MQ) and natural mineral water at various  
23 concentrations, for a signal to noise ratio of three for LODs and ten for LOQs. Limits  
24 of detection (LODs) were comprised between 0.1 – 0.5 ng L<sup>-1</sup> for PAHs and 0.5 – 2  
25 ng L<sup>-1</sup> for VOCs. Limits of quantification (LOQs) were comprised between 0.3 - 1.5  
26 ng L<sup>-1</sup> for PAHs and 1.5 – 6 ng L<sup>-1</sup> for VOCs. Analytical variability was determined  
27 on spiked MQ and mineral water samples (n = 5) and was found to be below 10% for  
28 all PAHs and below 20% for all VOCs.  
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1 Bile samples were analysed for PAH metabolites, according to a method adapted from  
2 Mazéas and Budzinski (2005), Wessel et al. (2013) and Dévier et al. (2013). Briefly,  
3 metabolites were extracted after deconjugation by solid phase extraction (SPE).  
4 Purified extracts were analysed by UPLC/MSMS (Wessel et al. 2013, Le Dû-Lacoste,  
5 2008). The reported sums of hydroxylated non-alkylated PAHs included 1+2 OH-  
6 naphthalenes ( $\Sigma$ -OH-naphthalenes), 1+2+3+4+9 OH-phenanthrenes ( $\Sigma$ -OH-  
7 phenanthrenes) and 3+9 OH-benzo[a]pyrenes ( $\Sigma$ -OH-BaP). Quantification was  
8 performed using internal standards (deuterated hydroxy-pyrene, 1-OHPyrd9) added  
9 prior to SPE extraction and the response factors were measured by injecting a solution  
10 of unlabelled standards mixed with deuterated compounds used as surrogate standards  
11 (table S3). LODs and LOQs were 0.3 and 1 ng g<sup>-1</sup>, respectively. Analytical variability  
12 was determined on spiked MQ, mineral water and bile samples (n = 5) and was below  
13 20% for all the compounds. For details on quality assurance and control procedures  
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## 34 **2.5 Hepatic RNA isolation, cDNA synthesis and RT-qPCR**

35 Livers were homogenized at 5,500 rpm for 3 x 10 seconds in tubes containing  
36 zirconium beads and 96% ethanol using a Precellys homogenizer (Bertin  
37 Technologies, Paris, France). Following centrifugation for 2 minutes at 10,000 x g,  
38 the resulting pellet was homogenized in 400  $\mu$ L lysis buffer (E.Z.N.A. Total RNA Kit  
39 I, Omega Bio-Tek, Norcross, GA). Following another centrifugation (2 min; 10,000 x  
40 g), nucleic acids were precipitated from supernatant by addition of ethanol (1:1 v v<sup>-1</sup>).  
41 RNA was purified on filter columns using the E.Z.N.A. Total RNA Kit I following  
42 the manufacturers instructions. RNA quantity and quality were determined using a  
43 nanodrop spectrophotometer (Nanodrop Products, Wilmington, DE) and a 2100  
44 bioanalyzer instrument (Agilent Technologies Inc., Santa Clara, CA), respectively.  
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1 Synthesis of cDNA (RNA input 1  $\mu\text{g}$ ) was performed using the qScript cDNA  
2 supermix (Quanta BioSciences, Gaithersburg, MD). A standard curve (200, 40, 8, 1.6  
3 ng total RNA) was included giving a 91% reaction efficiency. Primer sequences for  
4 CYP1A used for RT-qPCR (PerfeCta SYBR Green Fastmix with low ROX, Quanta  
5 BioSciences) were CCAACTTACCTCTGCTGGAAGC (forward) and  
6 GGTGAACGGCAGGAAGGA (reverse) (Rees et al., 2003). Annealing temperature  
7 was 60°C.  
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## 10 **2.6 Hepatic CYP1A protein quantity and EROD activity**

11 The hepatic CYP1A protein content was analysed by a semi-quantitative enzyme-  
12 linked immunosorbent assay (ELISA) as described in Holth et al. (2008), with the  
13 following modifications. Livers were homogenised in potassium phosphate buffer (5  
14 mL  $\text{g}^{-1}$  tissue; 0.1M pH 7.8; 0.15M KCl; 1 mM DTT; 5% glycerol) using glassware  
15 and a teflon pistil with tight fit (Potter-Elvehjem). Homogenates were centrifuged at  
16 12,000 x  $g$  for 20 minutes at 4°C to remove cell debris. The supernatant was  
17 centrifuged at 50,000 x  $g$  for 2 hours at 4°C. The microsomal pellets were  
18 resuspended in potassium phosphate buffer (0.1M pH 7.8; 0.15 M KCl; 1 mM EDTA;  
19 20% glycerol) and frozen at -80°C before quantification of CYP1A protein (LOQ 4.5  
20 mOD  $\mu\text{g}^{-1}$  protein). The EROD activity was measured in S9 fraction of liver  
21 homogenates as described in Burgeot et al. (1996) using the method of Burke and  
22 Mayer (1974), modified for plate readers by Galgani and Bocquene (1991).  
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## 51 **2.7 EROD activity in gills**

52 The assay was performed as described in Jönsson et al. (2002), with the following  
53 modifications. One gill arch was excised and placed in 0.5 mL ice-cold HEPES-  
54 cortland buffer pH 7.7 (0.13M NaCl, 6.0 mM HEPES, 5.6 mM glucose, 5.1mM KCl,  
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1 2.4 mM NaH<sub>2</sub>PO<sub>4</sub>, 1.6 mM CaCl<sub>2</sub>, 0.9 mM MgSO<sub>4</sub>) in a 24-well microtiter plate. The  
2 assay was started by replacing HEPES-cortland buffer with 0.5 mL room tempered  
3 EROD-buffer (HEPES-cortland buffer containing 1 μM dicumarol and 20 μM  
4 resorufin ether). After 2 minutes, the EROD-buffer was replaced with 0.5 mL fresh  
5 room tempered EROD-buffer. The samples were incubated in the dark for 30 minutes  
6 at room temperature. Two hundred microlitres of samples, blanks (EROD-buffer) and  
7 a resorufin standard dilution series ranging from 6 – 200 nM were transferred in  
8 duplicates to a 96-well plate and fluorescence (535/585 nm) determined (Bio-Tek  
9 FL800x; LOQ 1.7 fmol min<sup>-1</sup> filament<sup>-1</sup>). The number of incubated gill filaments was  
10 counted (approximately 30 filaments per fish) and results were reported as fmol  
11 resorufin minute<sup>-1</sup> filament<sup>-1</sup> (Jönsson et al., 2003).  
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## 27 **2.8 Glutathion S-transferase (GST) activity in gills**

28 Gill filaments were added 4 mL potassium phosphate buffer (0.1M, pH 6.5) and  
29 homogenized using glassware and a teflon pistil with tight fit (Potter-Elvehjem).  
30 Homogenized samples were centrifuged (9,000 x g, 30 minutes, 4°C) and supernatant  
31 used to determine broad spectred GST activity as described by Habig and Jakoby  
32 (1981), adapted to microtiter plates by Frasco and Guilhermino (2002). Three  
33 replicates (100 μL) of blanks and samples were added to microtiter plates and  
34 reactions started by addition of 200 μL potassium phosphate buffer containing 10 mM  
35 glutathione and 10 mM 1-chloro-2,4-dinitrobenzene. Optical density was monitored  
36 (340 nm) every 20 seconds for 5 minutes using a microtiter plate spectrophotometer  
37 (Bio-Tek PowerWave 340) at 25°C. The enzymatic activity was calculated from the  
38 slope of the absorbance curve and was expressed in μmol min<sup>-1</sup> mg<sup>-1</sup> protein (LOQ  
39 6.2 μmol min<sup>-1</sup> mg<sup>-1</sup> protein).  
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## 2.9 Acetylcholinesterase (AChE) activity in muscle

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3 Approximately 1 g of muscle was added 4 mL potassium phosphate buffer (0.1M, pH  
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5 7.2) and homogenized using glassware and a teflon pistil with tight fit (Potter-  
6  
7 Elvehjem). Homogenized samples were centrifuged at 5,000 x g for 5 minutes at 4°C  
8  
9 to remove cell debris. The supernatant was diluted 10 times in potassium phosphate  
10  
11 buffer and frozen at -80°C until analysis. The determination of AChE activity was  
12  
13 performed according to the method described by Ellman et al. (1961), adjusted for use  
14  
15 in microtiter plates (Guilhermino et al., 1996). Four replicates (50 µL) of blanks and  
16  
17 samples were added to microtiter plates and reactions started by addition of 250 µL  
18  
19 potassium phosphate buffer containing 0.5 mM acetylthiocholine iodide, 0.33 mM  
20  
21 5,5'-dithiobis-(2-nitrobenzoic acid) and 0.6 mM sodium bicarbonate. Optical density  
22  
23 was monitored (414 nm) every 2 minutes for 10 minutes using a microtiter plate  
24  
25 spectrophotometer (Bio-Tek PowerWave 340) at 25°C. Protein concentration in  
26  
27 muscle homogenates was determined using the method of Bradford et al. (1976) as  
28  
29 indicated in Frasco and Guilhermino (2002). The enzymatic activity was calculated  
30  
31 using the extinction coefficient of 5-thio-2-nitrobenzoic acid (TNB) ( $\epsilon = 1.36 \times 10^4$   
32  
33 mL mmol<sup>-1</sup> cm<sup>-1</sup>) from the slope of the absorbance curve and was expressed as nmol  
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35 thiocholine min<sup>-1</sup> mg<sup>-1</sup> protein (LOQ 0.8 nmol min<sup>-1</sup> mg<sup>-1</sup> protein).  
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## 2.10 Lipid hydroperoxide (LPO) concentration in gills

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48 Gill samples (filaments) were homogenized (1:10 g wt v<sup>-1</sup>) in potassium phosphate  
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50 buffer (0.1 M, pH 7.4) and endogenous LPO determined through measurements of  
51  
52 thiobarbituric acid reactive species (TBARS), according to Ohkawa et al. (1979) and  
53  
54 Bird and Draper (1984). Artificial lipid oxidation was prevented by addition of 0.2  
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56 mM butylhydroxytoluene (Torres et al., 2002). Gill homogenates (0.2 mL) were  
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1 added 1 mL 12% trichloroacetic acid, 1 mL 0.73% thiobarbituric acid and 0.8 mL 60  
2 mM Tris-HCl (pH 7.4) containing 0.1 mM DTPA. Following incubation (60 minutes,  
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4  
5 100°C) the solution was centrifuged at 12,000 x g for 5 minutes and optical density  
6  
7 determined (535 nm) using a microtiter plate spectrophotometer (Bio-Tek  
8  
9 PowerWave 340). LPO concentrations were expressed as nmol TBARS g<sup>-1</sup> wt (LOQ:  
10  
11 202 nmol TBARS g<sup>-1</sup> wt).  
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### 15 **2.11 Vitellogenin protein quantity in plasma**

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17 Plasma samples were analysed for the presence of vitellogenin (VTG), the female  
18  
19 specific egg yolk protein, using a validated homologous ELISA method (Scott et al.,  
20  
21 2002, 2006a, 2006b). The limit of detection was 93 ng mL<sup>-1</sup>, estimated from 2 x  
22  
23 standard deviation of 15 blank measurements.  
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### 28 **2.12 Micronucleus in erythrocytes**

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30 After three weeks of exposure, a blood smear was made from one drop of heparinized  
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32 blood on a microscope slide. Samples were air dried at room temperature and stored  
33  
34 in the dark at 4°C until analysis. The samples were fixed (3:1 v v<sup>-1</sup> methanol:acetic  
35  
36 acid) for 5 minutes, air dried and incubated for 40 minutes in 3% giemsa stain. The  
37  
38 samples were washed twice (tap water), air-dried and a cover slide was mounted.  
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40 Quantification of micronuclei in 5,000 cells per sample was assessed using a  
41  
42 microscope (Olympus BX41TS, Tokyo, Japan) at 400x magnification, following the  
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44 scoring criteria as summarised in Baršienė et al. (2012).  
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### 52 **2.13 Statistical analyses**

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54 Statistical analyses were performed using JMP® version 8.0.2 (SAS Institute Inc.,  
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56 Cary, NC, USA) and Microsoft Excel 2008 for Mac (Microsoft Corp., Redmond,  
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1 WA). One-way ANOVA was performed on data from cod sampled one and three  
2 weeks after the start of exposures for analysis of effects of treatments. Furthermore, a  
3 one-way ANOVA was performed on data from each treatment group (including pre-  
4 exposure control) for analysis of within group temporal development. Samples for  
5 both types of analyses were treated as independent, assuming negligible effects of fish  
6 density for the latter analysis (sampled fish were terminated on each occasion). The  
7 general model assumptions (homoscedasticity and variance to mean relationships)  
8 were evaluated using Levene's test ( $p < 0.05$ ) and residual plots, respectively. If  
9 necessary, the data was  $\log^{10}$ -transformed and re-evaluated. Following significant  
10 ANOVAs a Dunnett's post-hoc test was performed (Sokal and Rohlf, 1994). When  
11 parametric tests were not applicable, non-parametric Kruskal-Wallis was used  
12 followed by a non-parametric post-hoc test (Conover, 1999). Applied transforms can  
13 be seen in table S4. If any value within a group was reported as below LOQ, the data  
14 set was not statistically tested. Furthermore,  $p$  values were adjusted for multiple  
15 comparisons before non-parametric post-hoc tests using the formula  $p_{adj} = p / k \times (k -$   
16  $1)$ , where  $k$  = the number of groups for comparison. Fold-change values for each  
17 biomarker (except vitellogenin and micronucleus) were calculated as ratios between  
18 values for individuals exposed to WAF from oil or diesel and the mean of the  
19 respective control group. For statistical testing of micronucleus data, samples were  
20 assigned a categorical variable (MN present or MN absent). Data were compiled in a  
21 contingency table and a Pearson chi-square test applied (Sokal and Rohlf, 1994).  
22 Unless otherwise mentioned, quantification limits were estimated from 3 x standard  
23 deviations of several blank measurements.  
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### 3 Results

#### 3.1 PAH and VOC in water

After one week, there were significantly higher concentrations of most PAHs in water in all 6 g oil kg<sup>-1</sup> gravel treatments compared to the control (table 1). WAFs of 2 g oil kg<sup>-1</sup> gravel were in most cases not significantly different from control (table 1). There were no significant differences for VOC in water among treatments (table 1). WAF of 6 g diesel kg<sup>-1</sup> gravel contained the highest concentrations of ΣPAH and alkylated naphthalenes: approximately twice the concentrations produced by 6 g NSC kg<sup>-1</sup> gravel and two to three times the concentrations produced by 6 g ALC kg<sup>-1</sup> gravel (table 1). The concentrations of alkylated phenanthrenes were similar between the respective doses of all WAFs, but concentrations of dibenzothiophene and its alkylated derivatives were 6 to 50 times higher in the WAF of the ALC treatments than in the WAFs of the NSC and diesel treatments (table 1).

Three weeks after the start of exposures, ΣPAH concentrations in all treatments were close to background and only ΣPAH concentrations in WAF of 6 g NSC kg<sup>-1</sup> gravel were significantly higher than the control (table 1). VOC concentrations in WAF of 6 g diesel kg<sup>-1</sup> gravel were significantly higher than the control (table 1).

Detailed data on PAHs on VOCs in water can be found in tables S5 and S6, respectively.

#### 3.2 PAH metabolites in bile

Exposure to crude oil or ship-diesel resulted in significantly increased concentrations of PAH bile metabolites for the high concentrations of all three oils after one week ( $p = 0.01$ , Dunnetts test for sum OH-PAH; table 2), but not for the low concentration or after three weeks exposure. Bile metabolite concentrations decreased



1 significantly for all treatments from 1 to 3 weeks (GLM;  $p = 0.0003$  for sum OH-  
2 PAH; table 2). Ratios between low molecular weight ( $\leq 4$ -ring OH-PAH) and high  
3 molecular weight (5-ring OH-PAH) were higher in fish exposed to diesel than in fish  
4 exposed to crude oils ( $p = 0.005$ , Kruskal Wallis for ratio data, table 3).  
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### 10 **3.3 Hepatic CYP1A gene expression and protein**

11 CYP1A gene expression was significantly increased in cod from all groups after  
12 one week of exposure when compared to the control group (table 4; figure 3A) and to  
13 the pre-exposure control group (table 4). Three weeks after the start of exposures, cod  
14 from all treatment groups except 2 g diesel  $\text{kg}^{-1}$  gravel still had increased hepatic  
15 CYP1A gene expression compared to the control group (table 5; figure 3B) and the  
16 pre-exposure control group (table 5).  
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28 CYP1A protein in liver was significantly increased in cod exposed for three  
29 weeks to WAF from 6 g ALC and 6 g diesel  $\text{kg}^{-1}$  gravel compared to fish from the  
30 control group (table 5; figure 3B), but there were no significant differences compared  
31 to the pre-exposure control group (table 5). There were no significant differences in  
32 hepatic CYP1A protein after one week of exposure (table 4; figure 3A).  
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### 41 **3.4 EROD activity in liver and gills**

42 Hepatic EROD activity was significantly increased after one week in cod  
43 exposed to WAF from 2 g NSC  $\text{kg}^{-1}$  gravel and 6 g ALC  $\text{kg}^{-1}$  gravel compared to the  
44 control group (table 4; figure 3A). Three weeks after the start of exposures, hepatic  
45 EROD activity was still significantly increased in cod exposed to WAF from 6 g ALC  
46  $\text{kg}^{-1}$  gravel, and also in cod exposed to WAF from 6 g NSC  $\text{kg}^{-1}$  gravel (table 5; figure  
47 3B). Compared to cod from the pre-exposure control group, hepatic EROD activity  
48 was increased in cod exposed to WAF from 2 and 6 g NSC  $\text{kg}^{-1}$  gravel after one and  
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1 three weeks, respectively, as well as in cod exposed for three weeks to WAF from 6 g  
2 ALC kg<sup>-1</sup> gravel (table 4 and 5).  
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4  
5 Gill EROD activity was significantly induced after three weeks in cod exposed  
6 to WAF from 2 and 6 g NSC kg<sup>-1</sup> gravel as well as 2 and 6 g ALC kg<sup>-1</sup> gravel  
7 compared to cod from the control group, but activity related to dose only in the ALC  
8 treatment (table 5; figure 3B). Cod exposed to ALC and NSC WAF had significantly  
9 induced activities compared to cod from the pre-exposure control group (table 5).  
10 After one week of exposure, gill EROD activity in cod from the control group, ALC  
11 and NSC groups was significantly increased compared to cod from the pre-exposure  
12 control group (table 4), and a dose related trend was apparent in all treatments (F=2.5,  
13 p<0.06).  
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### 27 **3.5 Glutathione S-transferase (GST) activity in gills**

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29 There were no significant differences in GST activity in gills in cod among  
30 groups after one and three weeks of exposure (table 4 and 5; figure 4). Compared to  
31 cod from the pre-exposure control group, there was a significant reduction in gill GST  
32 activity after three weeks in cod exposed to WAF from 6 g diesel kg<sup>-1</sup> gravel (table 5).  
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### 41 **3.6 Lipid peroxidation (TBARS) in gills**

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43 After three weeks the level of gill TBARS was significantly increased in cod  
44 exposed to WAF from 2 g diesel kg<sup>-1</sup> gravel (table 5; figure 4B). One week after the  
45 start of exposure there were no significant differences among groups (table 4; figure  
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### 3.7 Acetylcholinesterase (AChE) activity in muscle

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3 Compared to the pre-exposure control group, AChE activity in muscle was  
4 significantly reduced after one week in cod from the control group and in cod exposed  
5 to WAF from ALC oil and WAF from 6 g NSC kg<sup>-1</sup> gravel (table 4). Similarly, after  
6 three weeks the AChE activity in muscle was significantly reduced in cod from the  
7 control group and in cod exposed to WAF from ALC oil (table 5). There were no  
8 significant differences in muscle AChE activity among exposure groups within the  
9 different timepoints.  
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### 3.8 Vitellogenin in plasma

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22 Due to a large fraction of cod with plasma vitellogenin concentrations below the  
23 limit of detection (LOD; 93 ng vtg mL<sup>-1</sup>), no statistical analysis was performed (table  
24 S7). In cod from the pre-exposure control group, four out of eight values were below  
25 LOD and the remaining four were close to LOD (<107 ng vtg mL<sup>-1</sup>). More fish had  
26 vitellogenin concentrations in plasma above LOD after three weeks of exposure than  
27 after one week of exposure. Most of the fish exposed to the highest oil concentrations  
28 for three weeks had vitellogenin concentrations in plasma above LOD (table S7).  
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### 3.9 Micronucleus in erythrocytes

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43 There were no significant differences between treatments in the frequency of  
44 fish with micronuclei present/absent in erythrocytes (table S8).  
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## 4 Discussion

52  
53 Atlantic cod was exposed to water-accommodated fractions (WAFs) from three  
54 weathered oils (ALC, NSC, ship-diesel) in a continuous flow-through system for three  
55 weeks. The selected concentrations of oil simulated a light contamination: a  
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1 discolouring of the gravel was only observed when applying 6 g oil kg<sup>-1</sup> gravel. The  
2 concentrations of ΣPAH in WAFs one week after the start of the experiment were in  
3  
4 the environmentally relevant ng L<sup>-1</sup> range (12.5 – 184.9): depending on weather  
5  
6 conditions and the distance from an oil spill site, hydrocarbon concentrations in  
7  
8 seawater in areas impacted by oil spills may range from ng L<sup>-1</sup> concentrations to short  
9  
10 periods of concentrations in the mg L<sup>-1</sup> range (Elordui-Zapatarietxe et al., 2010;  
11  
12 González et al., 2006; Reddy and Quinn, 2001; Stagg et al., 2000). ΣPAH  
13  
14 concentrations in seawater receiving oil production wastewater have been estimated to  
15  
16 be in the lower (30-50) ng L<sup>-1</sup> range (Harman et al., 2009), a similar concentration  
17  
18 range as in the current experiment. For comparison, PAH reference concentrations for  
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20 OSPAR region IV were suggested to be in the pg L<sup>-1</sup> range (OSPAR Commision,  
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22 2004).  
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28  
29 There was no relationship between water concentrations of PAHs after one week  
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31 and the general characterisation of oil hydrocarbons in the original oils. The oil  
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33 containing the largest fraction of mono- and polyaromatic compounds (Arabian Light;  
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35 39.6%) did not result in the highest concentrations of ΣPAHs or alkylated  
36  
37 naphthalenes/phenanthrenes in water. There are many possible explanations for this  
38  
39 observation, predominantly associated with properties of the exposure system and the  
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41 WAF produced. We have no measurement of what came out of the columns the first  
42  
43 week, during which there could have been a depletion of some of the components for  
44  
45 some oils. Arabian Light contained more asphaltthenes than any of the other oils.  
46  
47 Asphaltthenes could have produced a more viscous matrix, thereby reducing the  
48  
49 leaching rate of other components of the oil. The rate of release of the smaller PAHs  
50  
51 may be larger from lower doses than from higher doses (Nahrgang et al., 2010). Such  
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53 effects may cause water PAH concentrations to differ among oil types, and not  
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1 necessarily in correlation to parent oil hydrocarbon composition. The concentrations  
2 of dibenzothiophenes and its derivatives were higher in the ALC treatments than in  
3 the NSC and diesel treatments, in correspondence with the higher sulphur content  
4 (including PAH heterocycles) of this oil type (Radović et al., 2012). The temporal  
5 reduction in PAH concentrations in water between one and three weeks was an  
6 expected result of the columns used for generating exposure (Carls et al., 1999), and  
7 further augments the environmental relevance of the study. By using biological  
8 biomarkers, variations in water contaminant concentration are integrated over time  
9 (see e.g. Aas et al., 2000a), reducing the complexity of impact assessment.  
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22 The increased concentrations of PAH metabolites in bile after one week of  
23 exposure indicated that these PAHs were bioavailable and taken up, metabolised and  
24 excreted to bile by the fish in all exposed groups. The range of concentrations of 1-  
25 OH-pyrene in bile was lower but close to concentrations observed in soles caught  
26 following the “Erika” oil spill (Budzinski et al., 2004). During the experiment, fish  
27 were fed twice a week thus bile was regularly evacuated to the intestines. After three  
28 weeks of exposure, most PAH metabolite concentrations in bile were reduced to  
29 within the range of PAH metabolite concentrations in control group fish and  
30 corresponded well with the observed temporal reductions in water PAH  
31 concentrations. PAH metabolite concentrations in bile may therefore indicate short  
32 term exposure, but depending on feeding status of the fish metabolite concentrations  
33 also have the potential to remain elevated for weeks following exposure to PAHs or  
34 dispersed oil (Aas et al., 2000a, 2000b).  
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53 The dose-related increase in hepatic CYP1A mRNA in all treatments at both  
54 time points indicated that the WAFs from ALC, NSC and diesel contained AhR  
55 agonists that activated CYP1A gene transcription (Goksøyr and Förlin, 1992). The  
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1 associated temporal reductions of PAH concentrations in water demonstrated that  
2 hepatic CYP1A gene expression in Atlantic cod, in addition to being a sensitive  
3 biomarker, could be expected to last for weeks following an oil exposure incident.  
4 Whether this was a result of a continued DNA transcription, high mRNA stability,  
5 low rate of mRNA degradation or other species-specific factors is not known, but a  
6 rapid CYP1A mRNA turnover (days) has been indicated in other fish species  
7 following intraperitoneal injections of CYP1A inducers (Durieux et al., 2012;  
8 Kloepper-Sams and Stegeman, 1989; Nahrgang et al., 2009). An extended period  
9 (four weeks) of CYP1A mRNA induction was however also observed in polar cod  
10 (*Boreogadus saida*) following exposure to WAF from a NSC oil (Nahrgang et al.,  
11 2010).

12 The effects on CYP1A protein were however not as clear. Hepatic CYP1A  
13 protein was induced after three weeks in some of the treatments (ALC and diesel), but  
14 with large variability among individuals (table 5; figure 3B). Such a lag between  
15 CYP1A mRNA transcription and protein expression was expected and has also been  
16 observed in other fish species (e.g. Durieux et al., 2012; Nahrgang et al., 2009). The  
17 large variability within groups in CYP1A protein expression could be due to  
18 individual differences in response times and sensitivity to potential agonists or  
19 inhibitors. Binding affinity of PAHs to the teleost AhR may predict the potency for  
20 CYP1A protein induction (Billiard et al., 2002), thus for single compound exposures a  
21 good correlation between CYP1A gene and protein expression would be expected. No  
22 compounds are yet known to inhibit the translation of CYP1A mRNA to protein,  
23 although fluoranthene may inhibit EROD activity (Willett et al., 2001). Translation  
24 and/or degradation of mRNA is regulated by miRNAs, which may be altered by  
25 xenobiotics (Jenny et al., 2012; Tsuchiya et al., 2006).

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There was an increase in gill and/or hepatic EROD activity in fish exposed to WAF from ALC and NSC oils, also corresponding to an increase in hepatic CYP1A protein in cod exposed to WAF from ALC for 3 weeks. The trends were less consistent in the same biomarkers in cod exposed to WAF from NSC oil. WAF from diesel did not induce gill or hepatic EROD activity, but did clearly induce CYP1A gene and protein expression after 3 weeks of exposure. The EROD activity was in a similar (liver) and lower (gills) range than EROD activity in cod exposed to 1 ppm North Sea crude oils (Abrahamson et al. 2008; Aas et al., 2000a). The lack of response relationships to PAH concentration may indicate concentration dependent effects of antagonists on EROD activity or differences in toxicokinetics (Curtis et al., 2011; Sturve et al., 2006; Viarengo et al., 1997; Willett et al., 2001). The dose-related effect on hepatic CYP1A gene expression in all treatments, as well as the induction of CYP1A protein and lack of EROD induction in liver in cod exposed to diesel WAF, suggests the former to be the more probable scenario. The ratios between low ( $\leq 4$ -rings) and high ( $> 4$ -rings) molecular weight PAH metabolites in bile were higher in the cod exposed to the diesel than to the NSC and ALC WAFs. Some low molecular weight PAHs and metals have been shown to inhibit EROD activity (Pathiratne and Hemachandra, 2010; Willett et al., 2001; Whyte et al., 2000). It is also possible that endocrine disruptors in oil produce crosstalk between the estrogen and aryl hydrocarbon receptors that could inhibit *in vivo* EROD activity in liver (Kirby et al., 2007). Only low vitellogenin concentrations in a few individuals were observed in the current study indicating a low potential for estrogen receptor activation. The higher ratio of saturates to aromatic compounds in the diesel and NSC oils may have modulated (inhibited) EROD activity in cod exposed to these oil WAFs.

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The activity of the phase II glutathione S-transferase (GST) isozymes in gills was not affected by the oil WAF treatments, indicating that the WAFs were poor GST inducers in Atlantic cod or that the GST enzymatic conjugating capacity was already sufficient (tables 4 and 5; figure 4). In fish, other phase II enzymes such as glucuronosyltransferases and sulfotransferases may also contribute significantly to, or dominate the conjugating capacity but substrate selectivity may vary (Celander et al., 1993; Van Der Oost et al., 2003; Willett et al., 2000). Although some GSTs may be induced through AhR activation (George, 1994), it has been shown that GSTs are not as easily inducible as the phase I system and that exposure to PCDDs and PAHs may cause both induction and inhibition of GST activities (for summary see Van Der Oost et al., 2003). A lack of GST induction or insufficient GST conjugating capacity could increase the residence time for electrophiles in the liver; a risk factor for chemical carcinogenesis in cod compared to other species with an easily inducible GST system such as starry flounder (*Platichthys stellatus*) or sea bass (*Dicentrarchus labrax*) (Kerambrun et al., 2012; Varanasi et al., 1987).

WAF from diesel appeared to have a larger potential than oil WAFs for producing oxidative damage in gills (TBARS). Although differences between crude oil and diesel exposure were moderate, the highest fold-change values of TBARS in gills were observed in cod exposed to WAF from diesel after both one and three weeks of exposure (figure 4). A low potential of North Sea crude oil for producing oxidative damage in liver was previously observed in Atlantic cod following exposure to 0.5 ppm North Sea oil for 15 days (Sturve et al., 2006). The increase over time in TBARS concentration in cod exposed to the lower diesel WAF concentration may indicate that the higher diesel WAF exposure activated lipid metabolism and thus regeneration mechanisms more efficiently than the lower diesel WAF exposure. An



1 oxidation of polyunsaturated fatty acids may induce regulatory feedback mechanisms  
2 on the expression of lipid metabolism genes (Sampath and Ntambi, 2005), thus  
3 increasing their rate of anabolism. Effects of a mixture of PAHs on the expression of  
4 key genes related to lipid metabolism (e.g. upregulation of *fasn* and *cebpa*) and  
5 oxidative stress (e.g. in iron metabolism) have previously been demonstrated in  
6 zebrafish (*Danio rerio*) and Atlantic cod (Holth et al., 2008, 2010). Multivariate  
7 analyses indicated that TBARS and gill EROD were the two biomarkers  
8 discriminating best between the oil types after one week of exposure (Holth et al.,  
9 2012).

21 Lipid peroxidation, generally thought to be a consequence of oxidative stress  
22 (Kappus, 1987), may also be linked to perturbations such as decreased lysosomal  
23 membrane stability, carcinogenesis and micronuclei formation (Brinkmann et al.,  
24 2013; Köhler et al., 2002). The lack of differences in frequency of fish with  
25 micronuclei present/absent among treatments may indicate a low potential, especially  
26 of crude oil WAFs, to produce chromosomal aberrations on the implemented  
27 timescale in cod erythrocytes (table S8). The micronuclei frequencies in individuals  
28 exposed to diesel WAF for three weeks were clearly higher than the suggested  
29 micronucleus background assessment criteria (Davies et al., 2012). There might also  
30 be both species and tissue differences in susceptibility for micronuclei formation, and  
31 although peripheral blood cells are the most commonly used cell type they may be  
32 less responsive than e.g. cephalic kidney cells or hepatocytes (Baršienė et al., 2006;  
33 Pietrapiana et al., 2002; Rybakovas et al., 2009; Viarengo et al., 2007). This may be a  
34 result of a relatively long erythrocyte turnover rate, estimated to extend 100 days in  
35 some fish species (Soldatov, 2005). Increasing the interval from exposure to sampling  
36 may therefore also improve sensitivity for detecting damage in erythrocytes, but at the  
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1 same time complicate the exposure scenario, especially in wild collected fish.  
2 Erythrocyte micronuclei formation appear to be a relevant biomarker following long-  
3 term exposure, especially to diesel type oil spills, but assessing other tissues may  
4 improve sensitivity and reduce the latency period also for other oil types.  
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10 Acetylcholine esterase activity (AChE) in cod muscle was not affected by  
11 exposure to the oil or diesel WAFs. WAFs from other oils and wastewater from oil  
12 production facilities may however contain compounds capable of inhibiting fish  
13 AChE both *in vitro* and *in vivo* (Casini et al., 2006; Holth and Tollefsen, 2012; Vieira  
14 et al., 2008). In the current experiment, a reduction of AChE activity in muscle was  
15 observed in cod from some of the treatments (including the control group), when  
16 compared to cod from the pre-exposure control group. Such a reduction in AChE  
17 activity could be a response to a reduced swimming activity or confinement stress (as  
18 suggested in e.g. Payne et al., 1996), and demonstrates the value of including pre-  
19 exposure controls. The range of enzymatic activities was similar to or even slightly  
20 higher than AChE activity measured in muscle from cod collected in the Baltic Sea  
21 (Schnell et al., 2008).  
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39 There was a low potency for vitellogenin protein induction of the NSC, ALC and  
40 diesel oil WAFs. After one week, vitellogenin concentrations in plasma were below  
41 the assay detection limit (93 ng mL<sup>-1</sup>) in most of the cod. After three weeks most of  
42 the cod exposed to WAF from 6 g kg<sup>-1</sup> gravel of NSC, ALC or diesel had plasma  
43 vitellogenin concentrations above 93 ng mL<sup>-1</sup> and up to 670 ng mL<sup>-1</sup>, even though  
44 PAH concentrations were reduced to background levels during the time-course of the  
45 experiment. Four weeks of exposure to 0.5% produced water from Ekofisk did not  
46 induce vitellogenin synthesis in juvenile cod (Sundt et al., 2012), but juvenile cod  
47 exposed to 1% produced water from Oseberg C for 78 days still had induced  
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1 vitellogenin concentrations in plasma following two months of recovery in natural  
2 seawater (Meier et al., 2010). The *in vitro* estrogen receptor agonist potency has been  
3 shown to vary significantly between oil fields (Thomas et al., 2004a, 2004b).  
4 Considering the huge potential for vitellogenin synthesis in juvenile cod (in the mg  
5 mL<sup>-1</sup> plasma range) (Meier et al., 2010), the concentrations measured in the current  
6 study were considered close to background. Possibly, a longer exposure period would  
7 have produced higher vitellogenin concentrations and more differentiated results  
8 between oil types.  
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19 Some of the responses changed significantly in the control group over the  
20 duration of the experiment. It is unlikely that this was due to solvent in the gravel,  
21 which would have evaporated, but reflect changes from the fish adapted to the tanks.  
22 In our view, tank controls are the most appropriate for comparing with exposures, but  
23 such changes over time suggest a need for further investigation of the time required  
24 for adaptation and the extent to which the exposure in itself affects responses.  
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## 34 35 36 **5 Conclusions**

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39 Juvenile Atlantic cod were exposed for one and three weeks to two environmentally  
40 realistic concentrations of water-accommodated fractions from Arabian Light crude  
41 oil, North Sea crude oil and ship-diesel. PAH metabolites measured in cod bile  
42 confirmed that oil components were bioavailable and metabolised by the fish. There  
43 were clear differences in responses in the CYP1A pathway between oil types:  
44 exposure to WAF from ALC oil clearly induced EROD activity in gills and liver.  
45 Responses in EROD activity of exposure to WAF from NSC were less clear, and  
46 there was no induction of EROD activity after exposure to WAF from diesel. This  
47 could be due to interactions from other components in oil and may lead to  
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1 underestimation of the potential for EROD induction. The gene expression of CYP1A  
2 in liver was a sensitive biomarker for all tested oil types, and the expression was  
3 maintained for three weeks although concentrations of PAHs in water and their  
4 metabolites in bile were reduced to background. A time lag between CYP1A mRNA  
5 and protein expression was observed in some of the treatments (ALC, diesel), but the  
6 extent of CYP1A protein expression was variable between treatments and individuals.  
7 In contrast, WAF from diesel appeared to have higher potential for inducing lipid  
8 peroxidation (TBARS) in gills and micronuclei in erythrocytes than crude oil WAFs.  
9 GST activity, vitellogenin synthesis and AChE activity did not appear to be  
10 modulated by any of the treatments.  
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## 26 **6 Acknowledgements**

27 This work was performed at the Biological Station Drøbak (University of Oslo) in  
28 Drøbak, Norway. Assistance from students at the University of Oslo during fish  
29 sampling, and technical contribution of Joana Almeida, Luís Luís and Dr. Luís Vieira  
30 (AChE/GST determinations) were greatly appreciated. This work was financially  
31 supported by: the RCN – Research Council of Norway, Norway; Defra – Department  
32 for Environment and Rural Affairs, UK; the French National Agency of Research  
33 ANR; MICINN – Ministry of Science and Innovation, Spain; and the Portuguese  
34 Foundation for the Science and Technology of Portugal (FCT: ERA-  
35 AMPERA/0001/2007), through the project European concerted action to foster  
36 prevention and best response to Accidental marine Pollution – AMPERA  
37 (ERACCT2005- 016165) within the framework of the EU ERA-Net initiative (6<sup>th</sup>  
38 Framework Program).  
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Table 1

Treatment	Dose (g kg <sup>-1</sup> gravel)	Weeks of exposure	ΣVOC (ng L <sup>-1</sup> )x'	ΣPAH (ng L <sup>-1</sup> )	ΣC1N (ng L <sup>-1</sup> )	ΣC2N (ng L <sup>-1</sup> )	ΣC3N (ng L <sup>-1</sup> )
Control	0	1	82 ± 10	5.0 ± 1.6	13.4 ± 3.7	23.3 ± 5.1	21.0 ± 5.5
		3	71 ± 10	4.9 ± 1.4	11.7 ± 2.5	23.8 ± 2.4	19.9 ± 2.3
Arabian Light crude	2	1	320 ± 228	12.5 ± 3.1	49.6 ± 4.1	146.0 ± 44.5	327.1 ± 174.6
		3	177 ± 29	6.9 ± 1.9	18.2 ± 4.9	25.9 ± 2.5	23.1 ± 3.9
Arabian Light crude	6	1	609 ± 446	25.5* ± 5.2	206.8* ± 84.9	669.5* ± 232.2	1160.0* ± 197.4
		3	122 ± 14	8.3 ± 0.2	17.1 ± 3.0	26.1 ± 3.1	26.3 ± 3.8
North Sea crude	2	1	527 ± 432	18.4* ± 1.2	42.2 ± 9.4	96.4 ± 11.7	170.2 ± 7.9
		3	109 ± 39	7.1 ± 1.4	9.9 ± 2.1	18.6 ± 3.7	15.0 ± 0.5
North Sea crude	6	1	463 ± 236	97.2* ± 16.3	288.5* ± 112.1	1036.3* ± 286.0	1204.7* ± 206.9
		3	165 ± 35	11.7* ± 1.6	8.4 ± 1.4	18.7 ± 2.5	23.8 ± 2.0
Diesel	2	1	383 ± 200	19.0* ± 4.0	56.7 ± 18.5	154.1 ± 48.3	219.5 ± 70.0
		3	154 ± 43	6.3 ± 0.6	10.3 ± 3.2	18.2 ± 2.2	20.2 ± 2.3
Diesel	6	1	1076 ± 419	184.9* ± 84.6	588.7 ± 315.9	1986.2* ± 1135.7	2166.3* ± 1324.6
		3	232* ± 21	10.1 ± 1.4	11.9 ± 1.9	23.4 ± 0.7	29.9 ± 1.7

continues..

Table 1 continued

Treatment	Dose (g kg <sup>-1</sup> gravel)	Weeks of exposure	ΣC4N <sup>a</sup> (ng L <sup>-1</sup> )	ΣC1P <sup>a</sup> (ng L <sup>-1</sup> )	ΣC2P <sup>a</sup> (ng L <sup>-1</sup> )	ΣC3P <sup>a</sup> (ng L <sup>-1</sup> )	DBT <sup>a</sup> (ng L <sup>-1</sup> )	ΣC1DBT <sup>a</sup> (ng L <sup>-1</sup> )
Control	0	1	18.7, 33.2, nq	1.5, 1.8, nq	4.7, 6.9, nq	3.4, 4.5, nq	0.8, nq, nq	1.2, nq, nq
		3	nq	2.3, 3.8, nq	nq	nq	nq	nq
Arabian Light crude	2	1	346.4 ± 186.3	22.7 ± 11.8	39.4 ± 12.2	17.9 ± 2.2	11.9 ± 5.8	74.8 ± 40.0
		3	nq	2.1, 3.0, nq	5.1, 6.8, nq	nq	0.4, 0.3, nq	0.7, 1.4, nq
Arabian Light crude	6	1	657.7 ± 40.4	52.5 ± 7.8	54.6 ± 13.2	18.8 ± 2.4	55.5 ± 13.5	189.6 ± 24.9
		3	14.2, nq, nq	2.6, nq, nq	5.1, 6.2, nq	nq	0.3 ± 0.0	1.4, nq, nq
North Sea crude	2	1	179.5 ± 20.3	22.3 ± 6.5	52.9 ± 2.2	24.2 ± 2.2	0.9 ± 0.1	5.1 ± 0.6
		3	nq	1.7, 2.2, nq	nq	nq	0.2, nq, nq	nq
North Sea crude	6	1	542.6 ± 58.6	112.7 ± 19.6	107.6 ± 19.1	37.3 ± 8.5	8.5 ± 1.0	23.3 ± 2.5
		3	17, 23, nq	2.0, 2.4, nq	7.5 ± 1.0	10.6 ± 0.8	0.3, 0.5, nq	1.0, nq, nq
Diesel	2	1	227.0 ± 83.1	15.3 ± 5.0	41.5 ± 12.0	19.9 ± 6.3	0.9, 0.9, nq	2.6, 3.1, nq
		3	20.2, nq, nq	nq	7.8, nq, nq	6.9, nq, nq	nq	nq
Diesel	6	1	1043.0 ± 535.8	91, 219, nq	102.6 ± 29.5	33.1 ± 4.7	2.0 ± 0.7	4.1 ± 1.5
		3	30.3 ± 1.2	3.8, nq, nq	9.3 ± 1.3	9.5 ± 0.4	0.5, nq, nq	nq

nq = not quantifiable/below limit of quantification

VOC: volatile organic compounds; C1N: C1-naphthalenes; C2N: C2-naphthalenes; C3N: C3-naphthalenes; C4N: C4-naphthalenes; C1P: C1-phenanthrenes; C2P: C2-phenanthrenes; C3P: C3-phenanthrenes; DBT: dibenzothiophene; C1DBT: C1-dibenzothiophenes

Sum VOC: benzene, toluene, ethylbenzene, xylene, isopropylbenzene, n-propylbenzene, 1,3,5-trimethylbenzene, terbutylbenzene, 1,2,4-trimethylbenzene, secbutylbenzene, 4-isopropyltoluene, butylbenzene and naphthalene.

Sum PAH: phenanthrene, anthracene, acenaphthylene, acenaphthene, fluorene, fluoranthene, pyrene, benzo(a)anthracene, chrysene and triphenylene

<sup>a</sup>data not statistically tested

Table 2

Treatment	Dose (g kg <sup>-1</sup> gravel)	Weeks of exposure	n	ΣOH- naphthalenes (ng g <sup>-1</sup> )	2-OH- biphenyl (ng g <sup>-1</sup> )	ΣOH- phenanthrenes (ng g <sup>-1</sup> )	3-OH- fluoranthene (ng g <sup>-1</sup> )	1-OH- pyrene (ng g <sup>-1</sup> )	1-OH- chrysene (ng g <sup>-1</sup> )	ΣOH- BaP (ng g <sup>-1</sup> )	ΣOH- PAH (ng g <sup>-1</sup> )
Pre-exposure control	0	0	3	4 ± 0.4	1.4 ± 0.4	50 ± 4	32 ± 2	90 ± 10	9 ± 1	29 ± 8	215 ± 15
Control	0	1	3	44 ± 6	8 ± 5	132 ± 38	44 ± 10	112 ± 32	15 ± 4	22 ± 5	377 ± 89
		3	2	36, 58	12, 24	115, 215	30, 58	71, 179	8, 22	21, 43	294, 599
Arabian Light crude	2	1	3	43 ± 26	24 ± 18	232 ± 61	19, nq, nq	346 ± 77	31 ± 14	36 ± 9	720 ± 159
		3	3	0.6, 0.8, nq	3.3, 9.4, nq	15 ± 11	1, nq, nq	37 ± 6	14, 21, nq	23 ± 6	92 ± 6
Arabian Light crude	6	1	2	25, 48	13, 60	107, 291	nq, nq	204, 1041	29, 32	24, 96	450, 1521
		3	3	5 ± 0.5	20 ± 8	77 ± 27	9 ± 2	79 ± 24	27 ± 11	35 ± 17	252 ± 66
North Sea crude	2	1	3	39 ± 3	10 ± 9	375 ± 168	7, nq, nq	334 ± 104	46 ± 10	37 ± 19	843 ± 284
		3	3	6 ± 0.3	40 ± 26	31 ± 7	3, nq, nq	124 ± 28	66 ± 4	76 ± 14	345 ± 17
North Sea crude	6	1	3	116 ± 48	18 ± 6	869 ± 410	27, nq, nq	784 ± 258	90 ± 20	125 ± 33	2012 ± 772
		3	3	55 ± 47	31 ± 11	611 ± 431	3, 16, nq	190 ± 71	121 ± 9	139 ± 33	1153 ± 363
Diesel	2	1	1	42	1.4	1281	42	840	23	6	2237
		3	3	5 ± 1	8 ± 3	137 ± 34	15 ± 4	445 ± 151	20 ± 6	13 ± 3	642 ± 181
Diesel	6	1	3	200 ± 142	24 ± 1	1008 ± 290	29, 81, nq	1021 ± 182	29 ± 5	12 ± 1	2332 ± 602
		3	3	9 ± 2	11 ± 3	200 ± 72	6, 14, nq	319 ± 88	14 ± 5	25 ± 7	585 ± 162

nq = not quantifiable/below limit of quantification

Table 3.

Treatment	Dose (g kg <sup>-1</sup> gravel)	n	Ratio
			<4-ring OH-PAH : 5-ring OH-PAH (median, range)
Arabian Light crude	2	6	11 (1 - 26)
Arabian Light crude	6	5	15 (4 - 18)
North Sea crude	2	6	9 (3 - 215)
North Sea crude	6	6	12 (2 - 21)
Diesel	2	4	50* (44 - 363)
Diesel	6	6	61* (21 - 240)

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Table 4

Treatment	Dose (g kg <sup>-1</sup> gravel)	n	hepatic CYP1A mRNA (normalized mean)	hepatic CYP1A protein (mOD µg <sup>-1</sup> protein)	hepatic EROD activity (pmol min <sup>-1</sup> mg <sup>-1</sup> protein)	gill EROD activity (fmol min <sup>-1</sup> filament <sup>-1</sup> )	gill GST activity (µmol min <sup>-1</sup> mg <sup>-1</sup> protein)	gill LPO level (nmol TBARS g <sup>-1</sup> tissue)	muscle AChE activity (nmol min <sup>-1</sup> mg <sup>-1</sup> protein)
Pre-exposure control	0	8	887 ± 130	48 ± 18	12.1 ± 2.5	0.3 ± 0.1	103 ± 9	127 ± 12	63 ± 3
Control	0	4	534 ± 162	56 ± 22	3.9 ± 2.2	1.9 <sup>†</sup> ± 0.4	67 ± 11	95 ± 37	47 <sup>†</sup> ± 5
Arabian Light crude	2	3	14612 <sup>**††</sup> ± 5529	53 ± 26	12.4 ± 2.1	3.2 <sup>†</sup> ± 0.9	70 ± 21	136 ± 36	44 <sup>†</sup> ± 7
Arabian Light crude	6	4	59438 <sup>**††</sup> ± 21016	79 ± 15	20.0* ± 4.1	7.3 <sup>†</sup> ± 2.2	86 ± 19	208 ± 38	45 <sup>†</sup> ± 6
North Sea crude	2	4	39881 <sup>**††</sup> ± 9729	82 ± 5	31.5 <sup>**†</sup> ± 7.2	7.8* <sup>†</sup> ± 1.1	84 ± 8	92 ± 23	58 ± 3
North Sea crude	6	4	88120 <sup>**††</sup> ± 24767	43 ± 24	19.4 ± 3.2	7.6 <sup>†</sup> ± 2.5	79 ± 16	158 ± 28	51 <sup>†</sup> ± 4
Diesel	2	4	5803 <sup>**††</sup> ± 947	49 ± 12	8.5 ± 2.8	4.0 ± 2.6	102 ± 4	122 ± 31	56 ± 6
Diesel <sup>#</sup>	6	2	6820, 33599	7, 46	1.1, 11.5	1.2, 6.1	30, 73	304, 393	41, 50

<sup>#</sup>not included in statistical tests (n<3). Individual values presented.

Table 5

Treatment	Dose (g kg <sup>-1</sup> gravel)	n	hepatic CYP1A mRNA (normalized mean)	hepatic CYP1A protein (mOD µg <sup>-1</sup> protein)	hepatic EROD activity (pmol min <sup>-1</sup> mg <sup>-1</sup> protein)	gill EROD activity (fmol min <sup>-1</sup> filament <sup>-1</sup> )	gill GST activity (µmol min <sup>-1</sup> mg <sup>-1</sup> protein)	gill LPO level (nmol TBARS g <sup>-1</sup> tissue)	muscle AChE activity (nmol min <sup>-1</sup> mg <sup>-1</sup> protein)
Control	0	4	622 ± 289	6 ± 1	9.3 ± 1.4	0.5 ± 0.1	67 ± 16	104 ± 4	50 <sup>†</sup> ± 3
Arabian Light crude	2	3	4304* <sup>†</sup> ± 1679	26 ± 11	26.4 ± 7.2	3.1* <sup>†</sup> ± 0.5	88 ± 9	153 ± 32	46 <sup>†</sup> ± 3
Arabian Light crude	6	4	23628** <sup>††</sup> ± 7090	117* ± 35	54.8** <sup>†</sup> ± 12.4	4.0* <sup>†</sup> ± 1.0	108 ± 22	200 ± 39	46 <sup>†</sup> ± 3
North Sea crude	2	4	24422** <sup>††</sup> ± 4598	62 ± 26	25.4 ± 3.9	4.5** <sup>†</sup> ± 0.6	81 ± 11	181 ± 32	57 ± 3
North Sea crude	6	4	34652** <sup>††</sup> ± 10398	71 ± 32	36.5* <sup>†</sup> ± 5.5	3.5* <sup>†</sup> ± 0.6	79 ± 8	103 ± 9	54 ± 2
Diesel	2	4	1162 ± 582	12 ± 2	11.0 ± 2.9	1.3 ± 0.4	67 ± 14	239* <sup>†</sup> ± 33	59 ± 3
Diesel	6	4	9250** <sup>††</sup> ± 2939	105* ± 35	7.7 ± 2.2	1.3 ± 0.4	61 <sup>†</sup> ± 5	134 ± 30	50 ± 7

**Table captions**

Table 1. Volatile organic compounds (VOC) and polycyclic aromatic hydrocarbon (PAH) concentrations in water ( $\text{ng L}^{-1}$ ), presented as average  $\pm$  standard error when all values were above the quantification limit ( $n=3$ ). For treatments with at least one value below the quantification limit, individual values are presented. Significant differences to the respective control groups one and three weeks after the start of exposure have been indicated ( $p<0.05$ ).

Table 2. PAH metabolite concentrations in bile ( $\text{ng g}^{-1}$ ) presented as average  $\pm$  standard error when all values were above the quantification limit ( $n=3$ ; LOQ). For treatments where  $n<3$  or at least one value below LOQ, individual values are presented (limit of detection= $0.3 \text{ ng g}^{-1}$ ; LOQ= $1 \text{ ng g}^{-1}$ ).

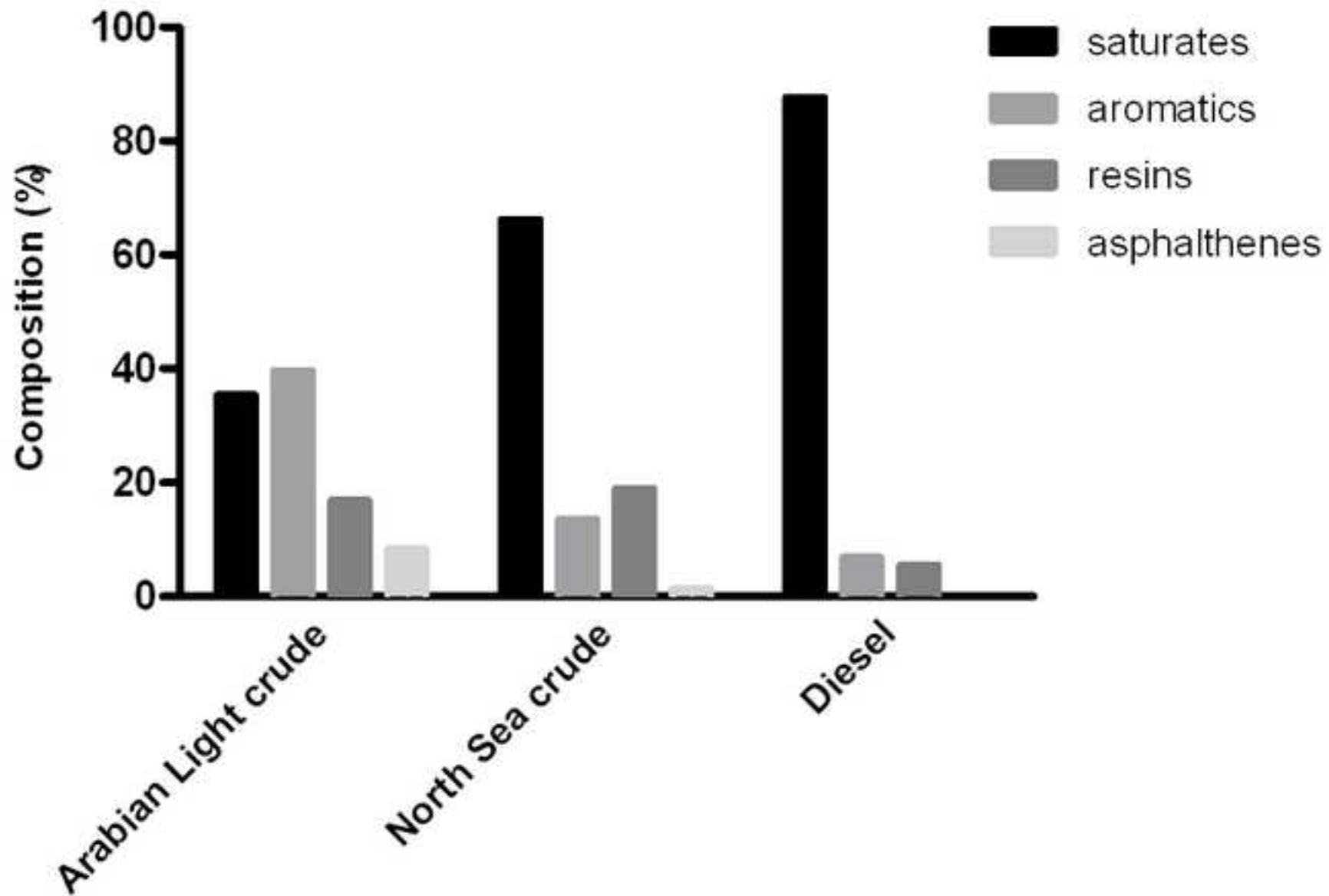
Table 3. Relationship (ratio) between  $\leq 4$ -ring OH-PAH and 5-ring OH-PAH in cod bile presented as median and min-max (non-parametric statistics applied). Significant differences between groups have been indicated (\*).

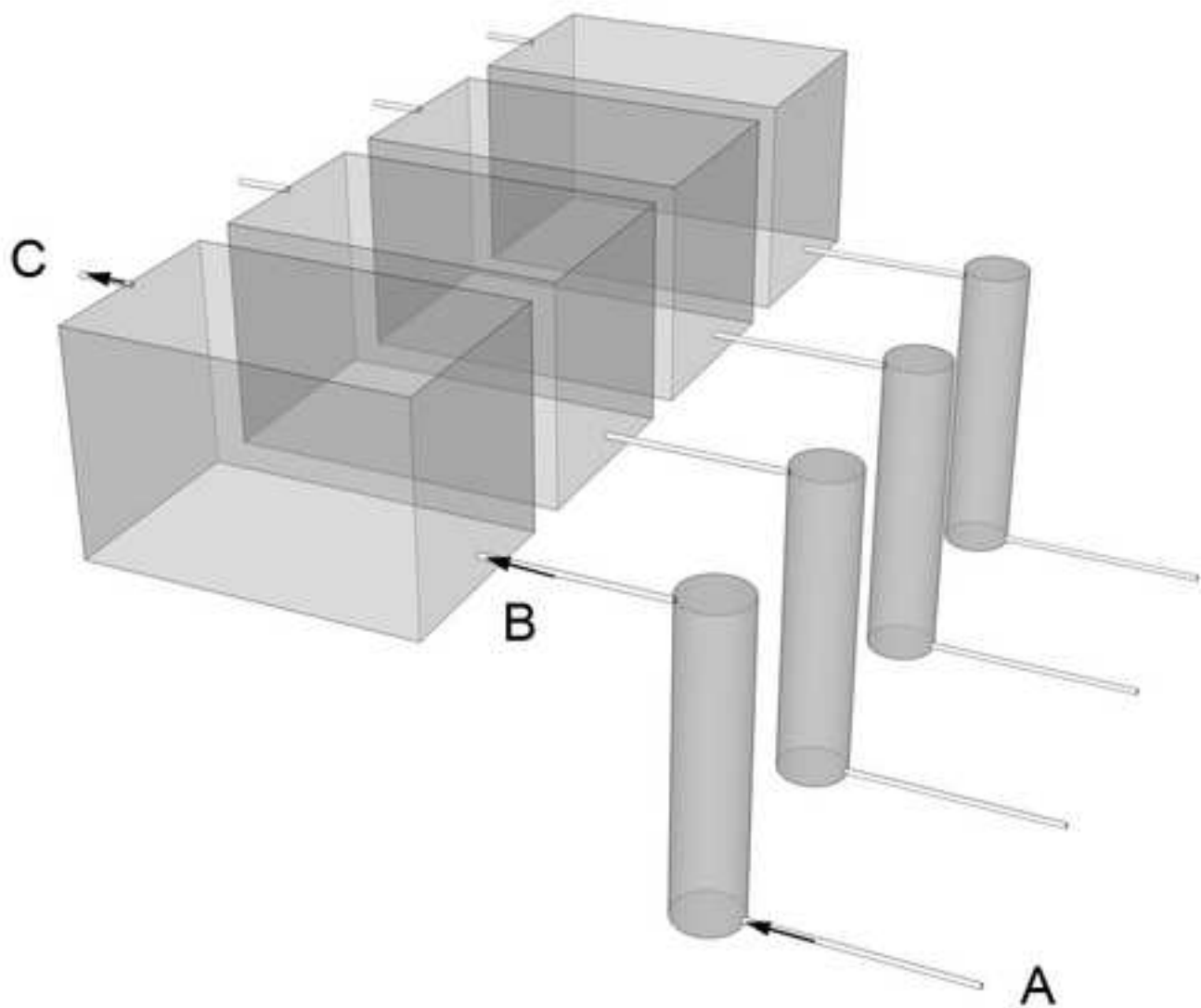
Table 4. Biomarkers in Atlantic cod from the pre-exposure control group and one week after start of exposure (mean $\pm$ SE). Significant differences to the control group (\*) and the time zero group (†) have been indicated (one symbol:  $p<0.05$ ; two symbols:  $p<0.001$ ).

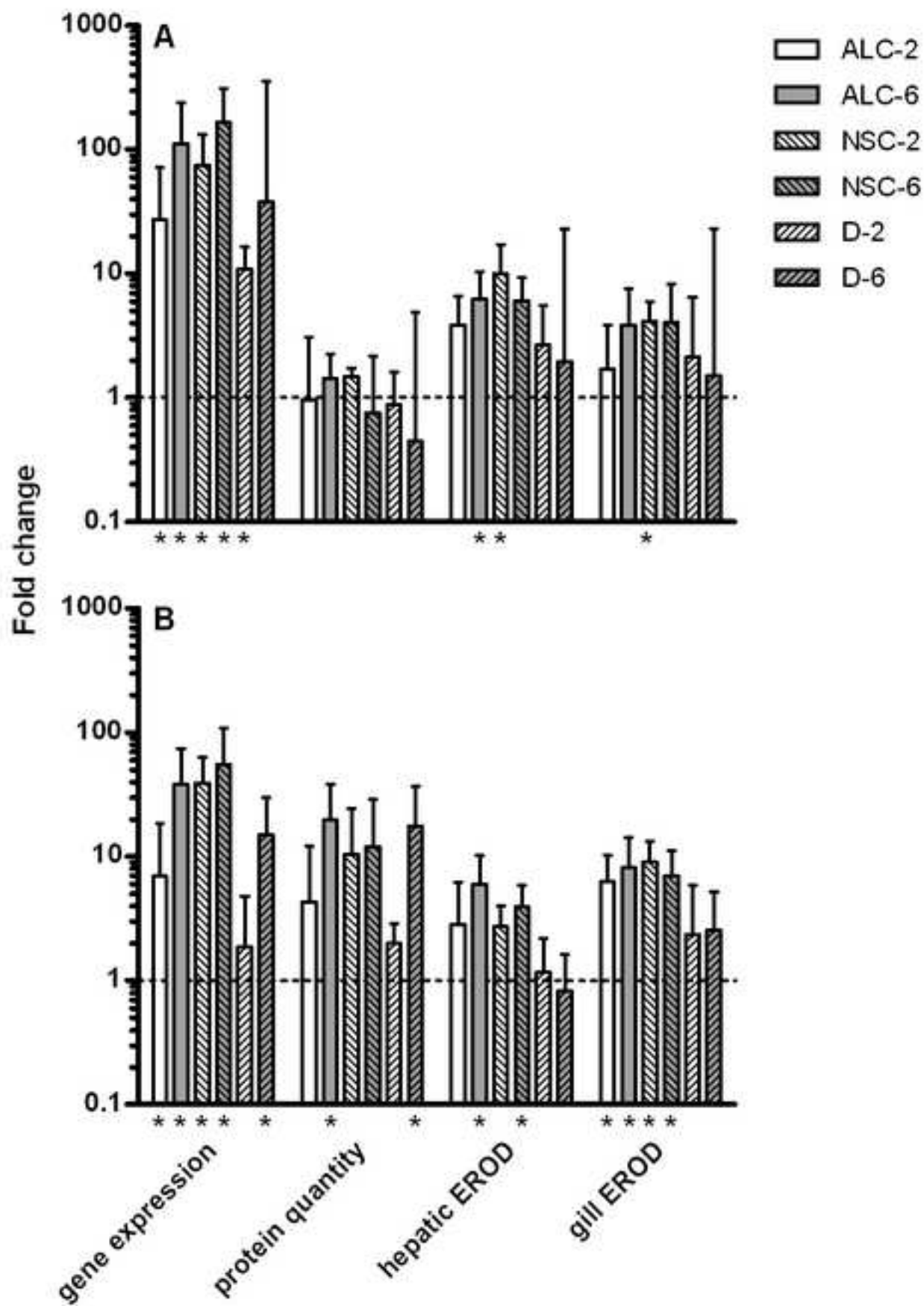
Table 5. Biomarkers in Atlantic cod three weeks after start of exposure (mean±SE). Significant differences to the control group (\*) and the time zero group (†) have been indicated (one symbol:  $p < 0.05$ ; two symbols:  $p < 0.001$ ).

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Figure 1







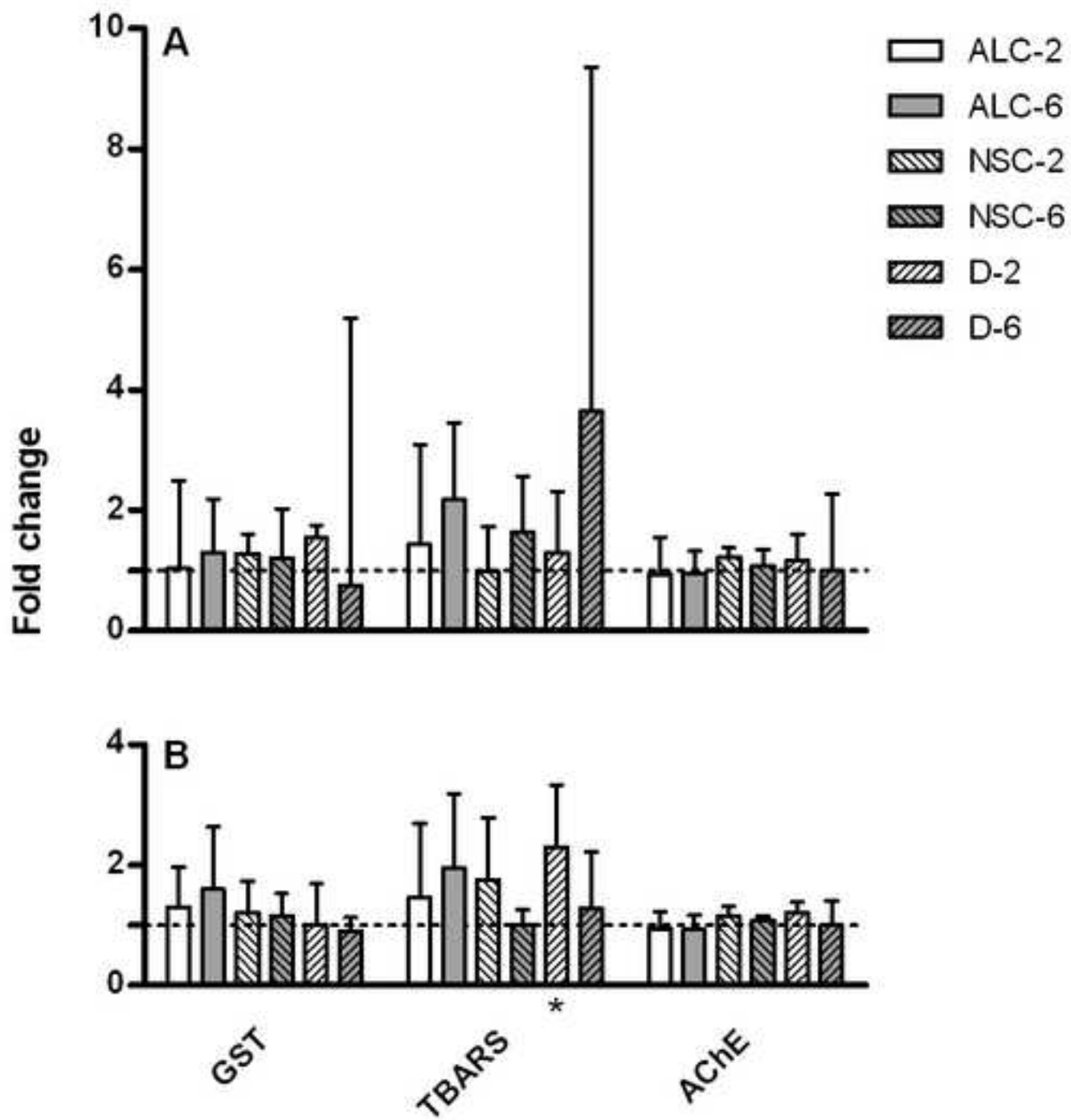




Figure 1. Composition of the three different oils included in the current study. Saturates include iso- and cyclic alkanes. Aromatics include monoaromatics, PAHs and hetero-polycyclic compounds. Resins are heterogeneous compounds, generally of high polarity. Asphaltenes are condensed aromatic macromolecules. Data from Radović et al. (2012).

Figure 2. Experimental setup of one of seven exposure groups: for each treatment, four tanks received effluents from separate columns containing gravel with same oil type and concentration (no oil for the control group). Seawater inlet was at bottom of columns (A), column effluent at top of columns (B), effluent inlet at bottom of aquaria (B) and aquaria outlet at top of aquaria (C). The exposure groups were randomly distributed.

**Figure 3.** Differences (relative to control) in hepatic CYP1A mRNA, hepatic CYP1A protein, hepatic EROD activity and gill EROD activity in cod after A) one week and B) three weeks of exposure to WAF of 2 and 6 g kg<sup>-1</sup> gravel of Arabian Light crude oil (ALC), North sea crude oil (NSC) and ship-diesel (D; average and 95% confidence intervals). Significant differences to the control group have been indicated below x-axis (\*; cf tables 3 and 4). Group size (n) as presented in tables 3 and 4.

**Figure 4.** Differences (relative to control) in gill GST activity, gill LPO level and muscle AChE activity in cod after A) one week and B) three weeks of exposure to WAF of 2 and 6 g kg<sup>-1</sup> gravel of Arabian Light crude oil (ALC), North sea crude oil (NSC) and ship-diesel (D; average and 95% confidence intervals). Significant

differences to the control group have been indicated below x-axis (\*; cf tables 3 and 4). Group size (n) as presented in tables 3 and 4.

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- Atlantic cod were exposed to the water-accommodated fraction of three different oils
- PAH metabolites in bile confirmed an environmentally realistic exposure scenario
- Responses in CYP1A and oxidative stress pathways differed among the oil types
- Biomarkers that could differentiate among oil types were recognized

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