

Gas Chromatographic-Mass Spectrometric Analysis of Polycyclic Aromatic Hydrocarbon Metabolites in Antarctic Fish (*Notothenia gibberifrons*) Injected with Diesel Fuel Arctic

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Abstract. Fish (*Notothenia gibberifrons*) collected in pristine Antarctic sites were injected with Diesel Fuel Arctic (DFA). Gas chromatography/mass spectrometry (GC/MS) was used to identify and quantify polycyclic aromatic hydrocarbon (PAH) metabolites obtained from hydrolyzed fish bile. The concentrations of naphthols (NPH), phenanthrenols (PHN), dibenzothiophenols (DBT), and total PAH metabolites (Σ PAH) vary with time. The response curve (production of PAH metabolites vs. time) resembled a Sigmoid curve, with an initial low response at 24 h followed by a rapid rise in production of PAH metabolites (55.74 μ g/g) 120 h after exposure to DFA. PAH metabolites identified include NPH, PHN, and DBT and their alkylated derivatives, reflecting the composition of the DFA to which the fish were exposed. The GC/MS technique is highly sensitive, particularly in the detection of multi-ring PAH metabolites. The results suggest that analysis of PAH metabolites is a valuable tool for environmental monitoring and assessment of exposure to petroleum.

Virtually all organisms possess biotransformation or detoxication enzymes which convert lipophilic xenobiotics to water-soluble and excretable metabolites (Livingston 1993). In the metabolic process, PAH are altered by phase I metabolism into various products such as epoxides, phenols, quinones, dihydrodiols, dihydrodiol epoxides, tetrahydrotriols, and tetrahydrodrotetrols (e.g., Varanasi and Gmur 1980). In phase II metabolism, these products are converted into highly water-soluble conjugates with a large water-soluble moiety, such as the tripeptide glutathione or sugar derivative glucuronic acid (Helou and Payne 1978). The composition and concentrations of PAH metabolites in fish bile would thus be expected to indicate

the extent of PAH exposure and the degree of exposure of fish to PAH (Krahn *et al.* 1992).

Diesel Fuel Arctic (DFA) is a refined oil that contains semi-volatile aromatic hydrocarbons such as naphthalenes and fluorenes (Kennicutt *et al.* 1991). In this study, DFA was administered by intramuscular (i.m.) injection to Antarctic fish (*Notothenia gibberifrons*). Biliary PAH metabolites were identified and quantified by gas chromatography/mass spectrometry techniques (Yu 1994). The objective of this study was to determine if the analysis of bile metabolites can be used as a reliable tool for monitoring and assessment studies of petroleum hydrocarbon contamination.

Materials and Methods

Collection of Fish

The fish (*Notothenia gibberifrons*) were caught at pristine sites in Low Island and Dallman Bay, Antarctic (McDonald *et al.* 1992), using an otter trawl, in March, 1991. Fish were transported in chilled sea water to the laboratory for further processing.

Exposure of Fish to Diesel Fuel Arctic (DFA)

Fish were maintained in running tap water at -1.5°C to 1.0°C in 5 ft diameter outdoor flow-through tanks. Fish were acclimatized for 1 week prior to initiation of the dosing experiment. The DFA (100 μ L) was administered by intramuscular injection (i.m.). The fish were not fed after injection. Fish were sacrificed for analysis at 0 (served as control point), 12, 24, 36, 72, 96, and 120 h after dosing. The gallbladder was removed immediately and stored at -60°C in amber vials until analysis.

Enzymatic and Acidic Hydrolysis and Extraction of Biliary Metabolites

Biliary conjugated (water-soluble) PAH metabolites were incubated with β -glucuronidase and aryl sulfates to cleave either glucuronide

Table 1. Mass fragments and intensities of ions of polycyclic aromatic hydrocarbon (PAH) metabolites

Compound	M+ (m/z) ^a	Other fragment ions (m/z)			
Naphthols					
C0 ^b	216 (100) ^c	201 (85)	185 (50)	137 (35)	
C1 ^d	230 (35)	215 (33)	141 (100)	73 (39)	
C2 ^e	244 (100)	229 (64)	214 (35)	199 (23)	73 (33)
C3 ^f	258 (14)	243 (12)	191 (11)	169 (100)	73 (13)
Phenylphenols					
C0	242 (100)	227 (88)	211 (65)	152 (27)	113 (27)
C1	256 (47)	241 (43)	211 (54)	167 (100)	
C2	270 (23)	255 (56)	225 (8)	181 (100)	73 (15)
C3	284 (59)	269 (98)	195 (100)	73 (74)	
Fluorenols					
C0	254 (48)	239 (15)	165 (100)	73 (37)	
C1	268 (15)	253 (10)	178 (100)	73 (42)	
C2	282 (100)	267 (27)	251 (12)	192 (39)	
C3	296 (100)	281 (27)	233 (33)	185 (21)	
Phenanthrenols/anthracenols					
C0	266 (100)	251 (44)	235 (28)	73 (50)	
C1	280 (65)	265 (31)	191 (100)	165 (21)	73 (29)
C2	294 (90)	279 (26)	205 (100)	189 (12)	131 (31)
C3	308 (100)	293 (80)	219 (70)		
Dibenzothiophenols					
C0	272 (23)	257 (8)	168 (100)	103 (30)	
C1	286 (44)	271 (46)	197 (100)	165 (15)	73 (33)
C2	300 (67)	286 (36)	269 (100)	254 (44)	
C3	314 (25)	299 (10)	283 (100)	225 (25)	73 (42)
Dibenzofuranols					
C0	256 (47)	241 (43)	211 (54)	167 (100)	
C1	270 (52)	225 (37)	181 (100)	73 (41)	
C2	284 (35)	268 (100)	195 (50)	179 (45)	73 (55)
C3	298 (25)	283 (19)	266 (75)	195 (27)	73 (100)
Pyrenols/Benzofluoranthenols					
C0	290 (100)	275 (29)	259 (30)	73 (49)	
C1	304 (100)	289 (26)	73 (57)		
C2	318 (31)	303 (100)	165 (66)		
Chrysenols					
C0	316 (40)	301 (12)	205 (20)	73 (100)	
C1	330 (100)	315 (35)			
C2	344 (100)				
BaP/Benzofluoranthenols					
C0	340 (100)				
C1	354 (100)				
C2	368 (100)	353 (52)	73 (77)		
C3	382 (100)				

^a: molecular ions of PAH metabolites

^b: unsubstituted PAH metabolites

^c: relative intensities of the ion as percentage of the primary ion (with 100% of the intensity)

^d: alkylated PAH metabolites with one methyl substituent

^e: alkylated PAH metabolites with two methyl substituents or one ethyl substituent

^f: alkylated PAH metabolites with three methyl substituents or one ethyl and two methyl substituents or one propyl substituent

and/or sulfate conjugates (Stein *et al.* 1984). Bile samples (100 μ l) were treated with 2,000 units β -glucuronidase (containing 20 units of sulfatase activity) dissolved in 1 ml of 0.4 M acetate buffer (pH 5.0). Surrogate standard containing 2,6-dibromophenol, 1-naphthol-d₈, phenanthrene-d₁₀, and chrysene-d₁₂ (10.0 ng/ μ l) was added, and the mixture was incubated in a DB-1 Dri-Block at 40°C for two hours. The sample was extracted with a solvent mixture consisting of 1 ml methylene chloride (CH₂Cl₂) and 100 μ l methanol (CH₃OH). Two additional extractions with 1 ml methylene chloride each were performed. The combined extract was dried with 0.1 g anhydrous sodium sulfate (Na₂SO₄), concentrated under a stream of nitrogen and transferred to an HPLC V-vial capped with Teflon-lined cap for GPC HPLC cleanup.

This fraction contained glucuronides and sulfates. Glutathion conjugates remaining in the aqueous phase were recovered by acidic hydrolysis described below.

The aqueous phase was treated with 0.1N hydrochloric acid (HCl) to adjust the pH to 1.0 and hydrolyze glutathion conjugates. The hydrolysis was conducted at ambient temperature. A surrogate standard containing 1-naphthol-d₈, phenanthrene-d₁₀, and chrysene-d₁₂ (10.0 ng/ μ l) was added before the sample was extracted three times with 1 ml methylene chloride each time. The combined extracts containing metabolites hydrolyzed from glutathion conjugates were then dried with 1 g anhydrous sodium sulfate (Na₂SO₄), concentrated, and further purified by GPC HPLC.

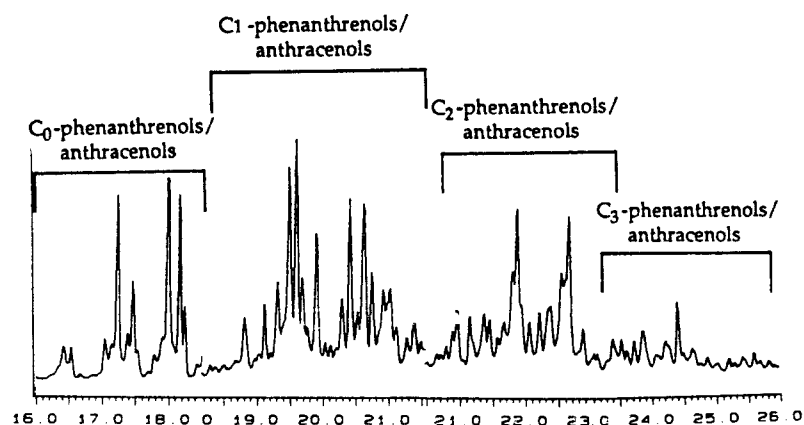
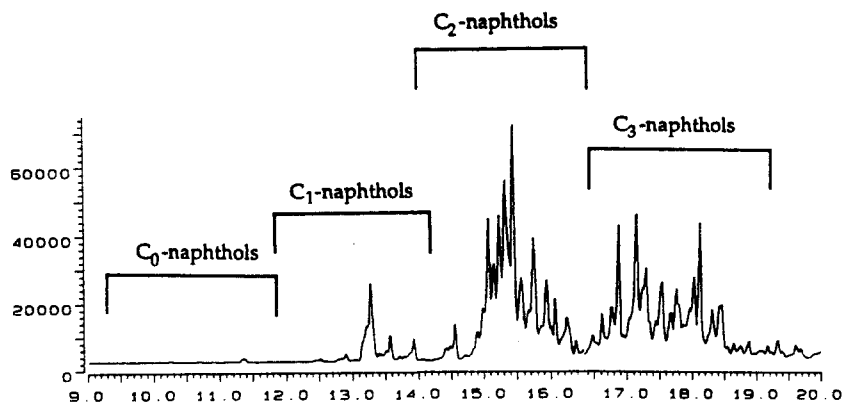


Fig. 1. Mass fragmentograms of naphthols (a) and phenanthrenols/anthracenols (b) in fish bile sampled 120 h after intramuscular injection of Diesel Fuel Arctic

GPC HPLC Clean-up of Extracts

The concentrated bile extracts from enzyme hydrolysis or from acidic hydrolysis were processed by GPC HPLC to remove lipids. The HPLC conditions used were modified from those reported by Krahn *et al.* (1992) for the separation of the PAH metabolites from biogenic material in the extracts after hydrolysis. A flow rate of 7 ml/min for the mobile phase (methylene chloride) was employed and the collection time for the metabolites started at the elution time of biphenyl and ended based on the elution time of 1-hydroxybenzo[a]pyrene. The HPLC analysis was performed at ambient temperature.

GC/MS Analysis of PAH Metabolites

Polycyclic aromatic hydrocarbon metabolites were converted to their corresponding trimethylsilyl (TMS) derivatives by silylating the PAH metabolites after the HPLC purification with bis(trimethylsilyl-fluoro)acetamide (BSTFA). The derivatization took place at 70°C for 1.5 h in a GC conical vial capped loosely. After completion of derivatization, the sample was cooled to ambient temperature and concentrated under a stream of nitrogen. GC internal standards (100 µl) fluorene- d_{10} and benzo[a]pyrene- d_{12} (0.50 ng/µl) were added. The final sample was evaporated to 100 µl with a gentle stream of nitrogen before GC/MS analysis.

Two microliters of derivatized PAH metabolites were injected in the splitless mode into the HP 5985 GC/MS linked with an HP1000 (RTE-6) data system. A J&W Scientific Durabond DB-5MS fused silica capillary column (30 m × 0.25 mm i.d.) was used for all analyses with the oven temperature programmed from 60°C at a rate of

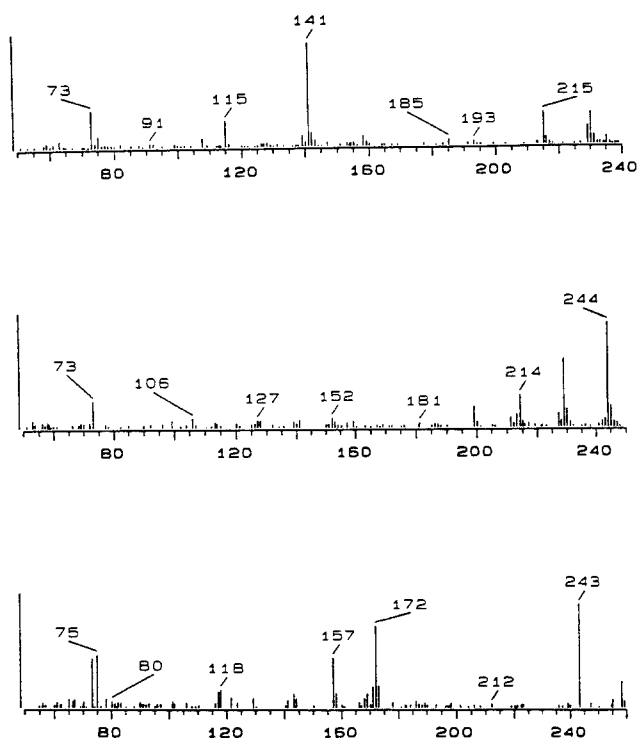


Fig. 2. Mass spectra of naphthols: (a) C₁ naphthols ($m/z = 230$); (b) C₂ naphthols ($m/z = 244$); and (c) C₃ naphthols ($m/z = 258$)

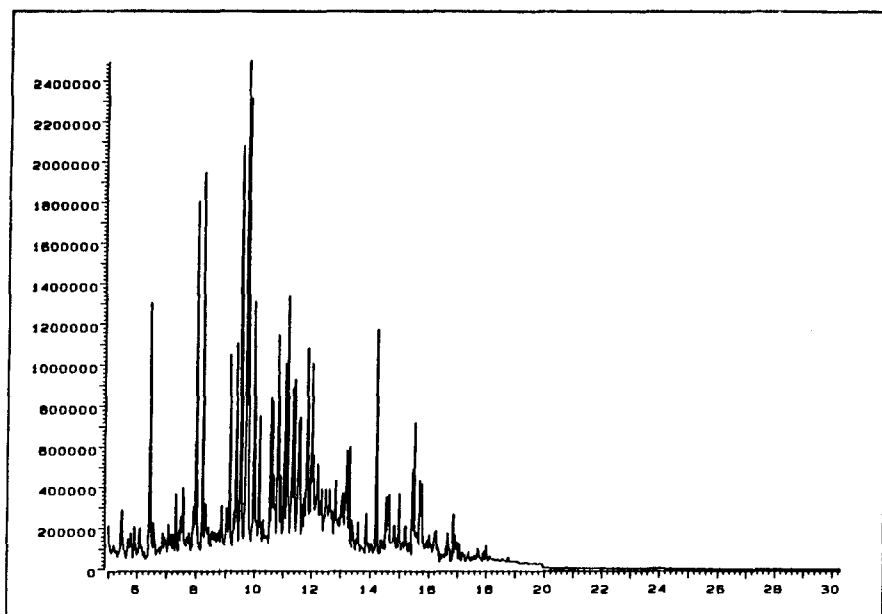


Fig. 3. Total ion chromatogram (TIC) of Diesel Fuel Arctic (DFA)

15°C/min to 130°C with initial holding at 60°C for 1 min, then from 130°C to 300°C at a rate of 5°C/min holding at 300°C for 5 min. The total running time was 45 minutes. The injector temperature was 300°C and the interface temperature was 280°C. The effluent from the GC capillary column was fed directly into the ion source (220°C) of the mass spectrometer. Selective ion monitoring (SIM) was routinely employed to acquire higher sensitivity. A full scan run was performed on most samples to confirm the SIM analysis results.

Identification and Quantification of PAH Metabolites

Polycyclic aromatic hydrocarbon metabolites were tentatively identified by comparing retention times and mass spectra for each metabolite to (a) retention time of reference standards where available or (b) the published spectra (Takahashi *et al.* 1979; Jacob and Grimmer 1985; Krahn *et al.* 1992) or (c) interpretation of the mass spectrum. In addition, identification of some metabolites, particularly those of the C₂-C₃ alkylated PAHs, high molecular weight PAHs, was based on the molecular ion and on fragmentation patterns. The identification of the metabolites was based on the characteristic fragment ions of the compounds (Table 1).

The absolute quantitation of individual metabolites was based on GC/MS response relative to that of surrogate standard. The GC/MS was initially calibrated by duplicate injections of standard at different concentrations. The concentration of each compound was calculated based on d₈-naphthol, a surrogate standard. The molecular ions of PAH metabolites were used as the quantification ions. In a few cases where the molecular ion has a low intensity, a distinct fragment ion was used for quantification. For instance, fragment ion (m/z = 208), rather than the molecular ion of d₈-naphthol (m/z = 223), was used for quantification due to the very high intensity of fragment ion m/z = 208.

Results and Discussion

Mass Chromatogram of PAH Metabolites

The trimethylsilyl (TMS) derivatives of PAH metabolites were well resolved. Mass fragmentograms of homologue series (C₀-C₃) of naphthols and phenanthrenols/anthracenols are presented

Table 2. Polycyclic aromatic hydrocarbon (PAH) composition of Diesel Fuel Arctic determined by GC-MS

Compound	concentration (ng/g)	$\Sigma C_x / \Sigma C_0$
naphthalene	995.78	7.3
1-methylnaphthalene	1395.73	
2-methylnaphthalene	1267.90	
C ₂ -naphthalene	2461.47	
C ₃ -naphthalene	1525.78	
C ₄ -naphthalene	652.44	
biphenyl	541.58	
fluorene	472.74	3.4
C ₁ -fluorene	618.86	
C ₂ -fluorene	607.45	
C ₃ -fluorene	372.29	
phenanthrene	570.61	1.9
anthracene	19.82	
C ₁ -phenanthrene/anthracene	625.06	
C ₂ -phenanthrene/anthracene	359.66	
C ₃ -phenanthrene/anthracene	105.97	
C ₄ -phenanthrene/anthracene	37.51	
dibenzothiophene	79.32	3.6
C ₁ -dibenzothiophene	126.36	
C ₂ -dibenzothiophene	100.92	
C ₃ -dibenzothiophene	58.41	
pyrene	18.45	0.2
fluoranthene	5.62	
C ₁ -pyrenes/fluoranthenes	4.24	
chrysene	3.21	0.9
benz[a]anthracene	0.58	
C ₁ -chrysene/benzanthracene	1.88	
C ₂ -chrysene/benzanthracene	1.54	
ΣC_0	2707.71	
ΣC_x	10324.47	
$\Sigma C_x / \Sigma C_0$	3.8	

in Figure 1. The structure of various isomeric hydroxyl metabolites, due to limited availability of reference standards, could not be established based solely on their mass spectra. However,

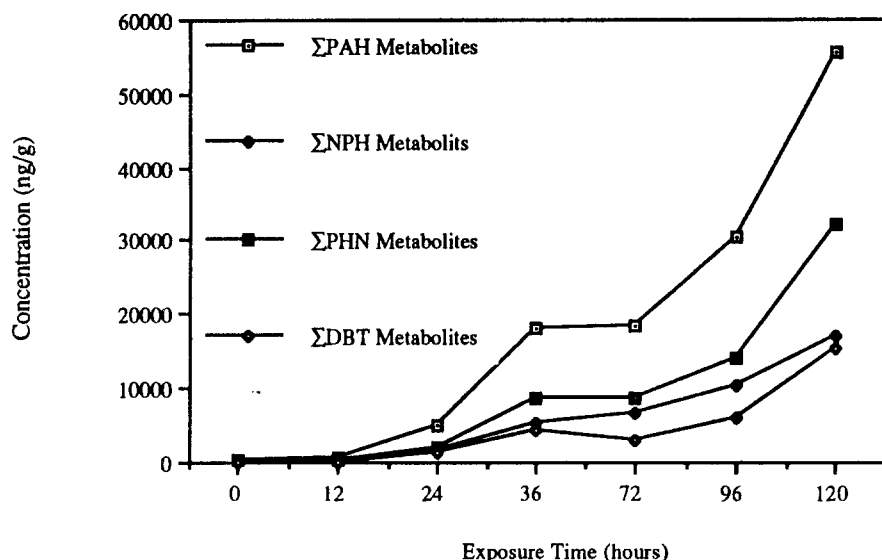


Fig. 4. The response curve: concentrations of total PAH metabolites (Σ PAH), phenanthrenols/anthracenols (Σ PHN), naphthols (Σ NPH), and dibenzothiophenols (Σ DBT) detected in fish bile are plotted against exposure time (hours). A Sigmoid curve is the result. See text for discussion

Table 3. Concentration of polycyclic aromatic hydrocarbon (PAH) metabolites detected by GC/MS in Antarctic fish (*Notothenia gibberifrons*) injected with Diesel Fuel Arctic

Exposure time (h)	Metabolite concentration (ng/g)							
	Σ NPH*	Σ DBT	Σ PHN	Σ PAH	ΣC_0	ΣC_x	$(\Sigma C_x/\Sigma C_0)_{met.}$	$(\Sigma C_x/\Sigma C_0)_{met./oil}$
0	195.3	23.0	249.0	467.2	66.6	400.7	6.0	1.6
12	176.5	111.6	379.0	667.1	39.1	628.0	16.1	4.2
24	1544.4	1291.0	2154.7	4990.0	167.8	4822.2	28.7	7.6
36	5331.6	4226.8	8582.4	18140.8	682.0	17458.8	25.6	6.7
72	6777.2	2865.2	8839.9	18482.3	401.6	17080.7	42.5	11.2
96	10413.9	5976.4	14110.4	30500.7	2142.1	28358.6	13.2	3.5
120	16941.6	15428.4	32225.3	55737.9	5287.1	50450.9	9.5	2.5

* Σ NPH: total C_0 - C_3 naphthols; Σ DBT: total C_0 - C_3 dibenzothiophenols; Σ PHN: total C_0 - C_3 phenanthrenols/anthracenols; Σ PAH: total PAH metabolites; ΣC_0 : total unsubstituted PAH metabolites; ΣC_x : total alkylated PAH metabolite; $(\Sigma C_x/\Sigma C_0)_{met.}$: ratio of total alkylated PAH metabolites over total unsubstituted PAH metabolites; $(\Sigma C_x/\Sigma C_0)_{met./oil}$: comparison of the ratio of total alkylated PAH metabolites over unsubstituted PAH metabolites to that of the oil to which the fish were exposed

the identity of each metabolite series (C_0 , C_1 , C_2 , and C_3) was readily assigned from interpretation of their mass spectra.

The TMS-ethers of PAH metabolites are characterized by a prominent molecular ion (Table 1, Figure 2) which appears as either a base peak, or peak with over 40% abundance compared to the base peak on mass spectrum. Other major fragment ions include $M^+ - 15$ and $M^+ - 31$. A strong $M^+ - 89$ ion [-OTMS, i.e., -O-Si(CH₃)₃] indicates the presence of a hydroxyl group.

Diesel Fuel Arctic (DFA)

Diesel Fuel Arctic (DFA) is a refined oil that is enriched in naphthalene and its alkylated (C_x) derivatives, with lesser amounts of C_0 (parent compounds) and C_x phenanthrene/anthracene and dibenzothiophene (Kennicutt *et al.* 1991, 1992; Figure 3).

Polycyclic Aromatic Hydrocarbon Biliary Metabolites from *Notothenia gibberifrons*

Concentrations of PAH metabolites measured in fish by the GC/MS method after i.m. injection of DFA at a dose level of

100 μ l are shown in Table 2. The results of the dose-response study showed that PAH metabolites were produced within 24 h after i.m. injection of DFA into *Notothenia gibberifrons*. The variations of concentrations of NPH, PHN, DBT, and total PAH are presented in Figure 4. The plot resembles a Sigmoid curve, consisting of an initial low-level response at 24 h followed by a rapid rise in production of PAH metabolites up to 120 h (Figure 4). The rise phase was steep; for instance, the production of PAH metabolites increased from 30.50 μ g/g in Σ PAH at 96 h, to 55.74 μ g/g at 120 h, an increase of 1.8-fold. The PAH composition of the DFA was predominated by naphthalene and its alkylated components (Kennicutt *et al.* 1991, 1992). However, the PAH metabolite analysis results (Table 3, Figure 4) suggest that the production rate of metabolites was higher for phenanthrene compared to naphthalene. The cause for this phenomenon is not year clear. English sole exposed simultaneously to naphthalene and BaP metabolized BaP to a greater extent than naphthalene; the ratios of BaP metabolites detected to BaP and naphthalene metabolites to naphthalene were, respectively, 49:1 and 6:1 at 24 h after exposure (Varanasi and Gmur 1981). Phenanthrenes/anthracenes, as a higher ring PAH (3-ring) than naphthalene, could be metabolized more extensively than naphthalene by the Antarctic fish, simi-

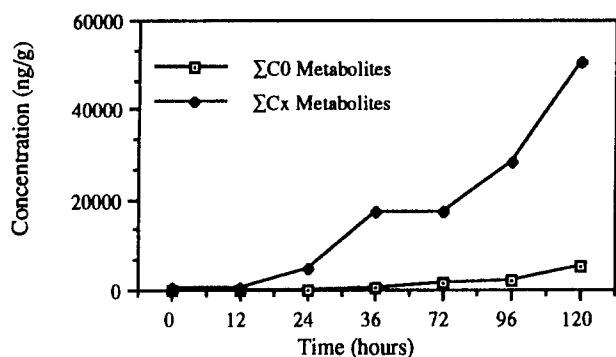


Fig. 5. Plot of the concentration of C_0 (total unsubstituted PAH metabolites) and C_x (total alkylated PAH metabolites) against exposure time (hour). C_x increased rapidly after 12 h, whereas C_0 increased only slightly

lar to BaP, where higher ring PAH (5-ring) were metabolized more extensively than naphthalene by English sole (Varanasi and Gmur 1981).

Alkyl substituted PAH metabolites (C_x) are much higher in concentration than unsubstituted ones (C_0) (Figure 5). The ratio of C_x/C_0 of metabolites detected in the fish bile ranged from 6.0 to 42.5 (Table 2), compared with DFA, whose C_x/C_0 ratio is 3.8; the C_x metabolites detected in fish were enriched from 1.8 to 12.9 fold. The concentrations of C_x metabolites show a rapid increase, whereas those of C_0 increased only slowly during the exposure period (Figure 5). This phenomenon may reflect the nature of the exposed petroleum (DFA). The C_x PAH of the DFA are 3.3 times higher than the C_0 PAH. It is also likely that the distribution pattern reflects the selectivity of metabolism and excretion (DiGiovanni and Juchau 1980), i.e., fish may preferentially metabolize alkyl-substituted aromatic hydrocarbons. Examination of Table 2 indicates that ratios of $\Sigma C_x/\Sigma C_0$ metabolites are much higher than the ratios of $\Sigma C_x/\Sigma C_0$ of parent hydrocarbons, particularly during the period of rapid rise in the production of PAH metabolites (Figure 4). This indicates the preferential metabolism of alkylated hydrocarbons by fish.

Krahn *et al.* (1992) were successful in using HPLC/fluorescence and GC/MS methods to detect PAH metabolites in fish exposed to petroleum after the *Exxon Valdez* oil spill. Hellou and Payne (1987) demonstrated in laboratory studies with trout exposed to water-soluble fractions or petroleum that detection of conjugated metabolites in the gallbladder bile of fish can provide a rapid investigative tool for assessment of petroleum and fuel oil pollution. The present study provides further support for the use of PAH metabolites in bile as monitoring indicators of PAH exposure in fish.

Conclusions

Antarctic fish (*Notothenia gibberifrons*) that were administered petroleum hydrocarbons (DFA) produced PAH metabo-

lites within 24 h of exposure. Production of PAH metabolites was generally proportional to exposure time. After an initial lag period (0–12 h), the production of PAH metabolites started at 24 h and maintained a steady concentration from 36 h to 72 h, followed by a rapid rise in concentration at 72 h. Fish bile was not obtained after 120 h. However, the findings in this study suggest that fish respond to PAH contamination by producing PAH metabolites. Therefore, the detection of PAH metabolites by GC/MS can provide a ready tool for monitoring and assessment studies.

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