

Bioaccumulation and metabolisation of 14C-pyrene by the Pacific oyster Crassostrea gigas exposed via seawater

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Paco Bustamante, Andrea Luna-Acosta, Sarah Clamens, Roberto Cassi, Hélène Thomas-Guyon, et al.. Bioaccumulation and metabolisation of 14C-pyrene by the Pacific oyster Crassostrea gigas exposed via seawater. Chemosphere, 2012, 87 (8), pp.944-950. 10.1016/j.chemosphere.2012.01.049 . hal-00682281

HAL Id: hal-00682281 https://hal.science/hal-00682281v1

Submitted on 24 Mar 2012 $\,$

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5	Bustamante P ^{1*} , Luna-Acc	osta A^1 , Clemens S^{1-2} , Cassi R^2 , Thomas-Guyon H^1 , Warnau M^2				
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7						
8	¹ Littoral Environnemen	t et Sociétés (LIENSs), UMR 7266, CNRS-Université de La				
9	Rochelle, 2 rue Olympe de Gouges, F-17000 La Rochelle, France					
10						
11	² IAEA – Environement I	Laboratories, 4 Quai Antoine Premier – MC-98000 - Principality of				
12	Monaco					
13						
14						
15	*Corresponding author:	Prof. Paco Bustamante				
16		Phone: +33-546-507-625				
17		Fax: +33-546-458-264				
18		E-mail: pbustama@univ-lr.fr				
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20 Abstract: The first objective of this study was to determine the bioaccumulation kinetics of 21 pyrene in the soft tissues of *Crassostrea gigas* (mantle, muscle, gills, digestive gland, and the 22 remaining soft tissues). As bivalves can biotransform hydrocarbons in more polar compounds 23 (metabolites) that are more easily excreted, the second objective was to investigate the oyster 24 capacity to metabolize pyrene into its metabolite, the 1-hydroxypyrene. To these ends, oysters were exposed 24h to waterborne ¹⁴C-pyrene then placed in depuration conditions for 15d. 25 26 Oysters efficiently bioaccumulated pyrene in their soft tissues and equilibrium was reached 27 within the exposure time. The metabolite1-hydroxypyrene was also detected in oyster tissues 28 but represented only 4 to 14% of the parent pyrene. At the end of the exposure period, the gills and the mantle showed the highest pyrene proportion of total soft tissue content, i.e. 47% 29 30 and 26%, respectively. After 15d of depuration, the mantle contained 32% and 30% of the 31 remaining pyrene and 1-hydroxypyrene, respectively. As C. gigas did not display a high capacity for metabolizing pyrene, it can be considered as a good bioindicator species to 32 33 survey and monitor pyrene contamination in the coastal marine environment.

34

Keywords: polycyclic aromatic hydrocarbons; bivalve; tissue distribution; bioaccumulation;
kinetics

38 1. Introduction

39 Polycyclic aromatic hydrocarbons (PAHs) are found in many common products, such as 40 petrol, fumes and household heating (e.g. using oil, gas or wood). Petroleum spills and 41 discharges, seepages, industrial and municipal wastewater, urban and suburban surface run-42 off, and atmospheric deposition contribute to aquatic contamination caused by PAHs (Eisler, 43 1987). During the last decades, many studies have monitored the inputs, fluxes and fate of 44 PAHs in the marine environment (Obana et al. 1983; Baumard et al. 1999). However, in order 45 to assess the state of the marine ecosystem, it is necessary to know the fraction of these 46 compounds which can be taken up by aquatic biota and their potential toxic effects (Escartin 47 and Porte, 1999).

The quality of aquatic environments can be assess through the analysis of organisms 48 considered as indicators of pollution, such as oysters, mussels and other bivalve molluscs 49 50 (Pereira et al. 1992; Jaffé et al. 1995; Lauenstein, 1995; Beliaeff et al. 1997; Gunther et al. 51 1999). Indeed, because of their biological (e.g. capacity of bioaccumulation, resistance to 52 physico-chemical stresses) and ecological characteristics (e.g. worldwide distribution, 53 abundance of their populations), bivalves are among the best candidates to be bioindicator 54 species (Phillips, 1976; Farrington and Tripp, 1993). These organisms can bioaccumulate a 55 large variety of pollutants at levels higher than those present in the surrounding waters or 56 sediments, and their behaviour can be recorded in short periods of time (Baumard et al. 1998; 57 Solé et al. 2000). In this way, mussels have been extensively used worldwide as sentinel 58 organisms to monitor the uptake and accumulation of PAHs in the coastal environments, in 59 the Mussel Watch Program in the USA (O'Connor 1996, O'Connor and Lauenstein 2006), the 60 Coordinated Environmental Monitoring Program (OSPAR 2010) and the Réseau National de 61 la Contamination CHimique (ROCCH 2008) in France. In this last country, oysters are also

62 used as a sentinel species in the ROCCH monitoring program, which is particularly relevant63 in areas where mussels are absent.

The exposure of marine organisms to PAHs has often been evaluated by measuring tissue 64 65 contaminant contents (Varanasi et al. 1989). However, when considering biotransformation capacities of these organisms, this approach becomes less relevant. Indeed, biotransformation 66 67 refers to the entire modification of chemical molecules occurring in the organisms. 68 Metabolism of PAHs in marine invertebrates is apparently related to cytochrome P450 (EC 69 1.14.14.1). This enzyme converts parent hydrophobic and lipid-soluble PAHs, into water 70 soluble metabolites. However, the mechanisms by which the involved enzymes are regulated 71 are still poorly understood (Hahn, 1998). In marine mammals and birds, some studies have 72 shown that the low concentrations of PAHs in their tissues were due to a combination of 73 inefficient bioaccumulation from food and rapid metabolisation and excretion of accumulated 74 PAHs (Watanabe et al. 1989; Fossi et al. 1995). In the common sole Solea solea affected by a 75 strong PAH pollution, the liver produced metabolites that were released through the bile 76 (Budzinski et al. 2004). Previously, it was believed that molluscs possessed a weak to non-77 existent ability to metabolize PAHs (Lee et al. 1972a; Palmork and Solbakken, 1981). 78 Conversely, molluscs may have relatively strong metabolisation systems (McElroy et al. 79 2000) and it has been shown that molluscs collected in strongly polluted environments 80 contained often low concentrations of PAHs (e.g. Varanasi et al. 1989; Baumard, 1997).

The Pacific oyster *Crassostrea gigas* (Thunberg, 1793) has a high economic value in the world and especially in France. Indeed, most of French oyster-farming raise this species which production averages up to 128 000 tonnes per year (CNC, 2004). Many farmhouses exist all along the French coastline and they are very sensitive to marine contamination, especially during oil slicks. It is therefore fundamental to better understand the mechanisms of bioaccumulation of PAHs in this species.

87 Various studies have shown that pyrene and other PAHs with four benzene rings are among 88 the most predominant PAHs in bivalves (Obana et al. 1983; Varanasi et al. 1985; Wade et al. 89 1988; Kaag et al. 1997). Moreover, pyrene is considered as one of the 16 most toxic contaminants for the environment and the dominant PAHs in the marine environment 90 91 (Giessing et al. 2003). Therefore, the first aim of this study was to follow the kinetics of 92 uptake and depuration of pyrene in the organs and tissues of C. gigas exposed via seawater. In order to study environmentally realistic contaminant levels, the pyrene used was ¹⁴C-labelled 93 94 and measured using highly sensitive radiodetection technique. The second objective of this 95 work was to determine the presence and the kinetics of formation of 1-hydroxypyrene, i.e. the 96 pyrene metabolite previously demonstrated to be predominant in fish bile (e.g. Krahn et al. 97 1987; Ariese et al. 1993),

98

99 **2. Materials and methods**

100 **2.1. Biological material**

101 Oysters were purchased from a shellfish farm on the French Atlantic coast (La Rochelle). 102 Organisms were then transferred to the Environment Laboratories premises (IAEA, Monaco). 103 Prior to the experimentation, specimens were acclimated to laboratory conditions for two 104 months (constantly aerated open-circuit aquarium; salinity: 36 ± 1 p.s.u.; temperature: 19 ± 1 105 °C; pH: 8; light/dark cycle: 12h/12h). During acclimation, bivalves were fed phytoplankton 106 using the Prymnesiophyceae *Isochrysis galbana* (10⁴ cells ml⁻¹). Recorded mortality was 107 lower than 5% over the acclimation period.

108

109 2.2. Radiotracer and radioanalyses

110 The ¹⁴C-labelled 4, 5, 9, 10 pyrene was purchased from Sigma, USA. Specific activity was 111 $2.17 \ 10^9 \text{ Bq mmol}^{-1}$. Stock solutions were prepared in methanol and a final concentration of 112 $27 \ \mu g \ L^{-1}$ was used. Two mL of the mixtures containing pyrene and 1-hydroxypyrene or purified pyrene samples (see below) were transferred to 20 mL glass scintillation vials (Packard) and mixed with 10 mL of scintillation liquid (Ultima Gold, Packard). ¹⁴C-radioactivity was measured using a 1600 TR Liquid Scintillation Analyser (Packard). Activity was determined by comparison with standards of known activities and measurements were corrected for counting efficiency and quenching effect. Counting time was adjusted to obtain a propagated counting error less than 5%.

120

121 **2.3. Experimental procedure**

122 <u>2.3.1. Uptake phase</u>

123 Forty four oysters were placed in a 50L glass aquarium containing natural seawater (closed circuit) spiked with ¹⁴C-labelled pyrene. The initial pyrene concentration in the aquarium was 124 $0.27 \ \mu g \ L^{-1}$. This concentration matches with PAHs values found in strongly polluted areas 125 126 (Axelman et al. 1999). Every 30 minutes, seawater radioactivity was measured and pyrene 127 was added as required in order to keep its concentration constant during the whole 128 accumulation phase (24 h). Oysters were not fed during the exposure period. Four animals 129 were collected at different times (0, 2, 5, 9, 19 and 24h) in order to follow the uptake kinetics of ¹⁴C-pyrene and the formation of its metabolite. At the end of the exposure period (24h), 130 131 seawater was sampled to detect whether metabolites could have been released by exposed 132 animals.

At each sampling time, oyster soft tissues were dissected into five compartments: mantle, gills, muscle, digestive gland and remaining tissues (i.e., labial palps, gonad and heart). Each organ and tissue was weighed and crushed. Then, samples were treated with 200 μ L of β glucuronidase-aryl-sulfatase mixture containing 100 000 units mL⁻¹ of glucuronidase (Sigma) and 7500 units mL⁻¹ of sulfatase (Sigma) for enzymatic deconjugation. To this purpose, the samples were buffered to pH 5 with 3 mL of 5 M potassium acetate buffer. They were then
placed in an oven at 37°C during 20 hours.

140 Pyrene and 1-hydroxypyrene were extracted via two subsequent liquid/liquid extractions. The 141 first extraction was performed by adding 5mL of a hexane/dichloromethane solution (50:50) 142 and the second extraction was performed by adding 4mL of a methanol/dichloromethane 143 solution (10:90) to extract pyrene and 1-hydroxypyrene, respectively. For both extractions, 144 the samples were mechanically shaken for 30 minutes and then centrifuged at 6000 rpm for 5 145 minutes. In the first extraction, the organic phase (5 mL) was recovered in a flat bottom flask. 146 Following the second extraction, the organic phase was recovered and combined with the first 147 one. Two mL of the extracted solution was radioanalysed to determine the amount of pyrene 148 + 1-hydroxypyrene in each organ.

The rest of the mixture was concentrated under a gentle stream of nitrogen to 1 mL and then separated by using upti-clean SPE glass columns Si/Cn-S (Interchim, Montluçon, France).
Pyrene was eluted with 5 mL of a hexane/dichloromethane (50:50) solution which was radioanalysed.

153 The 1-hydroxypyrene content in each organ was calculated by comparing the results of the154 two radioanalyses.

155

156 <u>2.3.2. Depuration phase</u>

At the end of the exposure period, the remaining organisms were placed in an open circuit 50L seawater aquarium (salinity: 36 ± 1 p.s.u.; temperature: 19 ± 1 °C; pH: 8; light/dark cycle: 12h/12h). At different times of the depuration period (0, 6, 18, 36, 96,192 and 336 h) four oysters were collected and their soft tissues dissected in order to follow the variation in pyrene and its metabolite. The dissected tissues and organs were processed according to the same method as previously described. 163 **2.4. Data analyses**

164 <u>2.4.1. Uptake kinetics</u>

A first order model was used to assess changes in pyrene concentration in oyster tissues along time during the exposure to waterborne pyrene. In this model the change in tissue activity with time was calculated by:

168

$$dA_{\rm org} / dt = k_u A_{\rm SW} - k_e A_{\rm org}$$
 (Eq. 1)

170

171 where A_{org} = activity of pyrene in tissue (Bq g⁻¹ tissue)

- 172 A_{SW} = activity of pyrene in seawater (Bq g⁻¹ water)
- 173 $k_u = uptake rate constant (Bq g⁻¹ h⁻¹)$

174
$$k_e = elimination rate constant (Bq g^{-1} h^{-1})$$

176

177 As A_{SW} was maintained constant during the uptake experiment phase, Eq. (1) can be 178 integrated to estimate tissue activities at any exposure time by:

179

180
$$A_{org} t = A_{SW} (k_u / k_e) (1 - e^{-K_e t})$$
 (Eq. 2)

181

182 where $A_{org} t$ = tissue activity at time t

183

184 When steady-state tissue activities are attained (i.e. $dA_{org} / dt = 0$), the bioconcentration factor

185 (BCF) can be estimated as follows:

187
$$A_{org} / A_{SW} = BCF = k_u / k_e$$
 (Eq. 3)

188

In this study, the BCF is the ratio between ¹⁴C-labelled pyrene in the body and in the 189 190 surrounding seawater. 191 192 2.4.2. Depuration kinetics The depuration kinetics were best fitted using either a single-component exponential equation 193 194 (Eq.(4)) or a double-component exponential equation (Eq.(5)): 195 $A_t = A_0 (e^{-k_e t})$ 196 (Eq. 4) $A_t = A_{0s} (e^{-k_{es} t}) + A_{0l} (e^{-k_{el} t})$ 197 (Eq. 5) 198 where A_t = remaining activity at time t (Bq g⁻¹) 199 A_0 = activity at time 0 (Bq g⁻¹), i.e. at the beginning of the depuration period 200 k_e = depuration rate constant (Bq g⁻¹ h⁻¹) 201 202 For double-component exponential model (Eq. 5), a 'short-lived' component (s) and a long-203 204 lived' component (l) describe the radiotracer proportion that is depurated rapidly (s) and 205 slowly (1), respectively. For each exponential component (s and l), a biological half-life can be calculated ($T_{b1/2s}$ and $T_{b1/2l}$) from the corresponding depuration rate constants (k_{es} and k_{el} , 206 207 respectively) according to the relation: 208 $T_{b1/2} = \ln 2 / k_e$ 209 (Eq. 6) 210 211 212

213 <u>2.4.3. Statistical analysis</u>

214 Model constants and their statistics were estimated by iterative adjustment of the model using 215 the nonlinear curve-fitting routines in the Statistica 6 software. For depuration kinetics, best 216 fitting models were selected between single- and double-component exponential equation, 217 according to the highest determination coefficient and examination of residuals. The level of 218 significance for statistical analyses was always set at α =0.05.

219

220 **3. Results**

221 **3.1. Seawater exposure**

222 Uptake of pyrene in five body compartments (mantle, gills, muscle, digestive gland and 223 remaining tissues) of C. gigas exposed to spiked seawater for 24h is presented in Fig. 1. The parameters and statistics of the uptake kinetic of ¹⁴C-pyrene are summarized in Table 1. The 224 225 steady-state BCF of pyrene in the whole soft parts of the oysters was observed within 24h and 226 reached 1560 \pm 630 (Table 1). Among the tissues, the gills accumulated rapidly and strongly 227 the contaminant with a BCF of 2080 ± 860 . However, each organ accumulated the pyrene in a 228 different way (Fig. 1). For example, during the uptake phase, the mantle and the remaining 229 tissues efficiently accumulated waterborne pyrene but the state of equilibrium was not 230 reached during the time frame of the experiment (Fig. 1).

The final distribution of the contaminant in each compartment after 24h of exposure is presented in Fig. 2. Among tissues, gills contained half of the whole body burden radioactivity and consistently showed the highest rate of accumulation with a k_u of 433 Bq g⁻¹ h^{-1} (Table 1). In contrast, the muscle was the organ with the lower rate of accumulation (k_u = 38 Bq g⁻¹ h⁻¹) and thus, only contained a very low proportion (5%) of the total quantity of ¹⁴Cpyrene present in oyster soft tissues (Fig. 2). Quantification of the pyrene metabolites was carried out at each sampling time, and the
proportion of 1-hydroxypyrene represented between 4 and 14% of the total radioactivity (Fig.
3).

At the end of the exposure period, the concentration of 1-hydroxypyrene in seawater wasbelow the detection limit of the method.

242

243 **3.2. Depuration phase**

244 At the end of the exposure time, non-contaminating conditions were restored and depuration 245 kinetics of the pyrene were followed in the organs and tissues of the oysters for 15d. The loss of incorporated ¹⁴C-labelled pyrene followed a single or a double exponential model in the 246 247 different body compartments (Fig. 4 and Table 2). In the muscle, gills and mantle, depuration 248 was best described by a double-component exponential equation ($R^2 = 0.42$, 0.52 and 0.48, 249 respectively). The resulting biological half-lives $(T_{b1/2})$ ranged from 1.17 (mantle) to 9.35 250 hours (gills) for the short-lived compartment and from 2.75 (muscle) to 7.27 days (gills) for 251 the long-lived compartment (Table 2). In contrast, a single-component exponential equation 252 better fitted the depuration kinetics in the digestive gland and in the remaining tissues (R² 253 =0.37 and 0.30, respectively). They were characterized by a relatively strong retention of 254 pyrene: the resulting $T_{b1/2}$ were 87.4 and 97.4 hours, respectively, (i.e., 3.64 and 4.05 days) 255 (Table 2, Fig. 4).

The distribution of ¹⁴C-labelled pyrene among the oyster soft tissues was determined at the end of the depuration period (Fig. 5). It differed from the distribution observed at the end of the exposure period (Fig. 2), with a lower fraction associated to the gills (24 ± 10 vs. $47 \pm$ 13%) and a higher fraction associated to the digestive gland (19 ± 10 vs. $6 \pm 4\%$) and the remaining tissues (21 ± 18 vs. $16 \pm 3\%$). At the end of the depuration period, the digestive gland displayed the highest pyrene activity (data not shown). In contrast, the gills had lost 50% of their activity during the depuration phase. Figure 5 also shows the distribution of 1hydroxypyrene between the body compartments at the end of the depuration period. 1hydroxypyrene was distributed in similar proportion as pyrene in the oyster body
compartments.

266

267 **4. Discussion**

Oysters accumulated very efficiently the ¹⁴C-labelled waterborne pyrene following a 268 269 saturation model and after a short exposure period (24h), organisms reached the state of 270 equilibrium. Among soft tissues, the gills accumulated rapidly and strongly the contaminant 271 with a BCF of 2080 (Table 1) likely because of the high filtration rate of oysters which could be as high as $3.9 \text{ L} \text{ h}^{-1} \text{ g}^{-1}$ dry weight (Bougrier et al. 1995). Consequently, the gills displayed 272 273 the highest activities at the end of the exposure phase. Absorption of pyrene onto gills might 274 be facilitated in oysters, as it occurs in blue mussels Mytilus edulis which have a micellar 275 layer which absorbs hydrocarbons (Lee et al. 1972a). Nevertheless, the accumulated pyrene 276 was relatively rapidly lost from this tissue (Table 1). Indeed, pyrene proportion in the gills at 277 the end of the depuration phase was half than at the end of the exposure period (Figs. 2 and 278 5). This decrease is due to a fast depuration rate of pyrene in the gills (Table 1). At the same 279 time, the increase of pyrene proportions in the digestive gland and remaining tissues suggest 280 that it was transferred towards from the gills to these tissues as previously reported in other 281 organisms (Neff, 1979). Therefore, the decrease of pyrene observed in the gills was also due 282 to a relative increase of the activity in other compartments, such as the digestive gland (data 283 not shown).

During the uptake phase, the mantle and the remaining tissues also accumulated pyrene but the steady-state was not reached for these compartments. This is probably due to the fact that, even if these tissues are in contact with seawater, their surface is much smaller than that of the

287 gills, and the cell types and the thickness of the epithelium differ completely (Auffret, 2003). 288 The resulting uptake rates (k_u) were lower for both tissues than for gills. During the 289 depuration phase, the mantle released pyrene quickly. In contrast, the remaining tissues 290 showed a slow increase of the pyrene proportion against the total pyrene content in oysters as 291 well as of its metabolite proportion (against the total metabolite content in oysters) during the 292 depuration period. It is well-known that tissues rich in lipids, e.g. gonads, accumulate 293 preferentially PAHs because of the highly hydrophobic nature of the latter (Berthelin et al. 294 2000, Meador et al. 1995). Moreover, a strong increase of the activity in the remaining tissues 295 was noticed during the third sampling in the depuration period (t = 36h). The four oysters 296 studied at this moment showed the particularity of being in the reproduction stage and of 297 having gonadic tissues more abundant than the average of other individuals. For this reason, 298 having acknowledged the unlikely hypothesis that these observations could come partly from 299 an error of manipulation, it seems that this increase of activity was rather due to a stronger 300 retention in mature gonadic tissues. This conclusion is consistent with results of Ellis et al. 301 (1993), where gonads from C. virginica displayed PAH concentrations five times higher than 302 in somatic tissues.

303 In contrast, the digestive gland and the muscle which are not truly in direct contact with 304 seawater displayed lower BCF than the gills, the mantle and the remaining tissues (Table 1). 305 Nevertheless, the uptake rate for the digestive gland was much higher than the one of the 306 muscle. The steady-state of pyrene in the digestive gland was reached very quickly, i.e. after 307 9h while it took 19h in the muscle (Fig. 1). The results strongly suggest that during the 308 exposure period pyrene was transferred from tissues in contact with seawater such as gills, 309 towards the digestive gland. Moreover, the digestive gland showed a higher percentage of 1-310 hydroxypyrene (19%) in relation to total radioactivity (pyrene + 1-hydroxypyrene) than the 311 other organs (9-12%) and the proportion of pyrene and its metabolite in the digestive gland 312 increased from 6-10% at the end of the accumulation period to 19-24% at the end of 313 depuration period (Figs. 2 and 5). This suggests that the digestive gland had a stronger 314 metabolisation capacity of pyrene than the other tissues.

315 Among all the tissues studied, the digestive gland and the remaining tissues displayed the 316 strongest retention capacity with $T_{b_{1/2}}$ of 87.4 and 97.4h, respectively. Indeed, just like almost 317 all organic contaminants, PAHs best store up in some tissues, namely in the hepatopancreas in 318 invertebrates and in the liver in vertebrates (Meador et al. 1995). Because of their highly 319 hydrophobic nature, PAHs are mostly accumulated in tissues with high lipid contents. The 320 transformed PAH metabolites generally accumulate in the hepatopancreas (Lee et al. 1976; 321 Neff et al. 1976; Meador et al. 1995). It has been proved that for some invertebrates and for 322 several fish species exposed to PAHs either via water, food or sediments, the cytochrome 323 P450 enzymatic system connected to an oxygenase function (MFO) is activated in their 324 hepatic structure (Andersson and Forlin, 1992). In the case of fish, this enzymatic system 325 allows excreting most of the PAHs bioaccumulated via the bile and the urine (Pritchard and 326 Bend, 1991). This particular function was difficult to bring to light concerning bivalves. For a 327 long time, it was admitted that bivalves did not possess a P450 system (Lee et al. 1972b; 328 Vandermeulen and Penrose, 1978). However, more recent studies have shown that bivalves 329 do possess a P450 system (Lake et al. 1985; McLeese and Burridge, 1987), allowing them to 330 metabolize PAHs. As it is the case for the majority of marine invertebrates, such ability to 331 metabolize PAHs is weaker than for vertebrates (Livingstone, 1994; Stegeman and Hahn, 332 1994).

Data obtained on 1-hydroxypyrene in the tissues and organs of *C. gigas* seem to confirm that metabolisation of pyrene in oysters is relatively weak and therefore would not be the driving mechanism responsible for its elimination. It is however possible that because of the large volumes of water they filter continuously, water-soluble metabolites would be rapidly

excreted and, therefore, not accumulated in the tissues (James, 1989). If the event lasts only
for few days, it is very likely that most of the accumulated PAHs are going to be eliminated.
In contrast, Meador et al. (1995) suggested that, during a chronic exposure, a non negligible
fraction of the PAHs could be stored in lipids and become less subject to the elimination by
diffusion or by metabolisation.

342 In conclusion, C. gigas presents a good potential of bioaccumulation of the pyrene and seems 343 to have a low metabolisation capacity for this compound. Such a bioaccumulation capacity 344 suggests that C. gigas could be used as a valuable bioindicator for pyrene. Following a marine 345 contamination, oysters are thus able to rapidly accumulate the contaminant in detectable 346 concentrations. Besides, as this species is very common in several regions of the world, 347 samples can be collected at any time and in many countries. This makes the situation easier 348 for comparing data. However, its speed of depuration is also quick, thus it does not allow the 349 recording of long-term pollution. On the contrary, it is very interesting for the monitoring of 350 the pollution variations on short-term periods.

351

352 **5. Acknowledgments**

353 MW is an Honorary Senior Research Associate of the National Fund for Scientific Research 354 (NFSR, Belgium) and held a 2008-2009 Invited Expert position at LIENSs (CNRS-Université 355 de La Rochelle), supported by the Conseil Régional de Poitou-Charentes. This work was 356 supported by the IAEA and LIENSs. The IAEA is grateful for the support provided to its 357 Environment Laboratories by the Government of Monaco.

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Figure 1: Uptake kinetics of 14 C-labelled pyrene in five body compartments and in the whole soft parts of the oysters (n=4) during the 24h seawater contamination period. Parameters and statistics of the uptake kinetics are given in Table 1.



Figure 2: Distribution of ¹⁴C-labelled pyrene (%) among the body compartments of oysters (n=4) at the end of the exposure period (24h).



Figure 3: Proportion (%) between¹⁴C-labelled pyrene (in white) and 1-hydroxypyrene (in black) in the whole soft parts of the oysters (n=4) at the end of the exposure period (24h).



Figure 4: Loss kinetics of 14 C-labelled pyrene in five body compartments and in the whole soft parts of the oysters (n=4) after a 24h seawater exposure. Parameters and statistics of the loss kinetics are given in Table 2.



Figure 5: Distribution (%) of ¹⁴C-labelled pyrene and 1-hydroxypyrene among the different body compartments of the oysters (n=4) at the end of the depuration period (15d).

Table 1. Parameters of the equation describing the uptake of pyrene in the body compartments and in the whole soft parts of the oysters (n=4) after 24h seawater contamination (k_u =uptake rate constant; k_e = elimination rate constant; BCF=Bioconcentration factor)

Body compartment	$(Bq g^{-1} h^{-1})$	k_{e} (Bq g ⁻¹ h ⁻¹)	BCF (k _u / k _e)	\mathbb{R}^2	
Mantle	102 ± 10	0.056 ± 0.006	1820 ± 530	0.45	-
Gills	433 ± 62	0.208 ± 0.030	2080 ± 860	0.28	
Muscle	38 ± 6	0.051 ± 0.008	750 ± 380	0.38	
Digestive gland	234 ± 50	0.137 ± 0.030	1710 ± 970	0.26	
Remaining tissues	100 ± 12	0.045 ± 0.006	2220 ± 810	0.35	
Whole body	158 ± 20	0.101 ± 0.014	1560 ± 630	0.39	

Table 2. Parameters of the equations describing the loss kinetics of pyrene in the different oyster body compartments in the whole soft tissues (n=4) after a 24h exposure from seawater. O and T: 1- and 2-exponential loss equations, respectively; A₀: remaining activity at time 0; k_e: depuration rate constant; *s*: short and *l*; long-lived elimination, respectively; R²: determination coefficient; T_{b½}: biological half-life in hours (h) or days (d). For abbreviation definitions, see 'Data and statistical analyses'

Compartment	Model	A_{0s}	k _{es}	$T_{b1/2s}\left(h\right)$	A_{01}	k _{el}	$T_{b1/2l}\left(d\right)$	R ²	р
Mantle	Т	443	0.591	1.17	769	0.006	4.92	0.42	< 0.001
Gills	Т	1578	0.074	9.35	368	0.004	7.27	0.52	< 0.001
Muscle	Т	291	0.087	7.93	166	0.011	2.75	0.48	< 0.001
Digestive gland	0	1360	0.008	87.4				0.37	< 0.001
Remaining tissues	0	1203	0.007	97.4				0.30	< 0.001
Whole soft parts	Т	600	0.195	3.55	721	0.007	4.29	0.53	< 0.001