

Immediate biomarker responses to benzo[a]pyrene in polluted and unpolluted populations of the blue mussel (*Mytilus edulis* L.) at high-latitudes

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Abstract

Immediate biomarker responses of two high-latitude populations of the blue mussel *Mytilus edulis* to benzo[a]pyrene (B[a]P) were evaluated. Mussels collected from a clean and a polluted site in southwest Iceland were exposed to the nominal dose of 100 µg B[a]P L⁻¹ for 3 h, after 4 days of acclimatization in clean seawater. To test the sensitivity to the toxicant and immediate biological responses, the following biomarkers were used: DNA single strand breaks, heart rate and feeding rate.

All the biomarkers revealed differences between the study sites. Irrespective of the origin of the organisms, the short time exposure to the high B[a]P concentration did not induce DNA single strand breaks or significantly affect the feeding rate. However, the heart rate results showed significantly different responses. The mussels from the polluted site (Reykjavík harbour) increased their heart rate when exposed to B[a]P, while no difference was observed between the heart rate values of the individuals from the clean site (Hvassahraun). The mussels seem to sense the pollutant they have been previously exposed to, and their acute response indicates physiological adaptation to the polluted environment. The results indicate limited sensitivity and temporal predictivity, i.e. transient measurable changes of these biomarkers, as well as showing that the background of the organisms should be considered when evaluating short-term biomarker responses to contaminants.

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1. Introduction

The effects of oil compounds, in particular polycyclic aromatic hydrocarbons (PAHs), on marine organisms have recently received much attention due to several large oil spills resulting in extensive environmental and economic effects (AMAP, 2002; Albaigés et al., 2006; Loureiro et al., 2006). The continued threat of oil spills may in the future reach new dimensions. Recent

climate change has resulted in decrease of the ice sheet of the Arctic Ocean (ACIA, 2004, 2005), and it can be foreseen that within several years large oil tankers can use the North-east passage between Asia and Europe, i.e. crossing the Arctic Ocean north of Russia (UNEP, 2004). This, in addition to increasing gas and oil exploration in the Arctic, increases the need to further understand the effects of oil compounds and PAHs on organisms at high-latitudes.

Effects of PAHs on organisms are extensive on various organism levels, including genotoxic, biochemical and physiological effects (Livingstone et al., 1985; Lowe et al., 2006). The responses or tolerance of organisms to PAHs can however vary considerably, even for the same species. Partly, this can be due

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to the fact that PAH mixtures can vary significantly within the geographical range of a species (Anderson and Lee, 2006; Skarphéðinsdóttir et al., 2007). Populations have also been shown to respond differently to PAHs, possibly reflecting the pre-exposure history and/or heritable, genetic changes in populations chronically exposed to mutagenic xenobiotics (Courtenay et al., 1999; Ma et al., 2000). Additionally, PAH levels and the subsequent effects on organisms, such as DNA adducts, may vary considerably between seasons at high-latitudes (Skarphéðinsdóttir et al., 2005). Little is still known of how responses, sensitivity and tolerance to PAHs are influenced by exposure to the contaminants. The estimation of PAH exposure and effects in invertebrate species remains problematic despite intensive research effort (Galloway, 2006).

The aims of this study were to evaluate immediate responses and sensitivity to benzo[a]pyrene of blue mussels (*Mytilus edulis* L.) coming from two different high-latitude environmental conditions: a polluted site (Reykjavík harbour) and a clean site (coastal site of Hvasshraun). Furthermore, to evaluate the heart rate as a potential monitoring tool at high-latitudes. The PAH compound benzo[a]pyrene (B[a]P) was selected as the test pollutant because it is one of the most harmful polycyclic aromatic hydrocarbons (IARC, 1987), and has also been reported in high concentrations in Reykjavík harbour (Ericson et al., 2002; Halldórsson et al., 2005). It is a potent genotoxic chemical, readily accumulated in the tissues of bivalve molluscs. Various effects of B[a]P on the blue mussel have been recorded, both at cellular and individual level (e.g. Eertman et al., 1995; Garcia and Livingstone, 1995; Large et al., 2002; Skarphéðinsdóttir et al., 2003; Moore et al., 2006).

To test the sensitivity and immediate metabolic responses of the blue mussel to B[a]P we used a suite of biomarkers, as widely recommended to get a holistic view on the subject (Hebel et al., 1997; Nicholson and Lam, 2005) and proposed for rapid assessment of marine pollution and coastal marine ecosystem health (Bowen and Depledge, 2006). We used the following biomarkers: heart rate, feeding rate (i.e. clearance rate; part of the biomarker scope for growth) and DNA single strand breaks, measured by means of the comet assay. These biomarkers have been widely applied to molluscs for ecotoxicological and monitoring purposes, both in laboratory and field studies (e.g. Widdows et al., 1997; Bolognesi et al., 2004; De Pirro and Marshall, 2005; Emmanouil et al., in press). They are non-specific biomarkers responding to a wide range of contaminants, which selection was based partly on the background information on the use of these biomarkers at high-latitudes (feeding rate, DNA strand breaks; Halldórsson et al., 2004, 2005). The high nominal B[a]P concentration of 100 µg L⁻¹ was chosen to stimulate biological responses to be evaluated in the time-span of 3 h.

To our knowledge, short or long term effects of B[a]P on the heart rate of mussels have not earlier been evaluated, but cardiac activity in bivalve molluscs has recently been suggested as a good biomarker of the metabolic status, reflecting general environmental stress (Galloway et al., 2004; Bowen and Depledge, 2006).

2. Materials and methods

2.1. Study area and sample collection

The blue mussels (*M. edulis* L.) were collected at two sites in Southwest Iceland, i.e. at a clean site and at a polluted one. The clean site is at Hvasshraun (64°01'11"N, 22°09'31"W) on the northern side of the Reykjanes peninsula. This site has been shown to be pristine and organisms from this site have frequently been used in monitoring programmes and as reference material in experiments (Yngvadóttir et al., 2006; Da Ros et al., 2007). No ship traffic or industrial activity is close to this site (<10 km).

The contaminated site is the inner side of Reykjavík harbour (64°09'01"N, 21°56'05"W). This harbour has extensive shipping activity and previous studies have shown high levels of PAHs in the tissues of the blue mussel. At the same location resident mussels had for example $\Sigma 16$ PAHs of 27 µg g⁻¹ lipid, and transplanted mussels had $\Sigma 32$ PAH levels as high as 56 µg g⁻¹ lipid (subtidal) and 114 µg g⁻¹ lipid (intertidal). These values correspond to 2.8, 4.8 and 9.7 µg g⁻¹ dry wt., respectively. (Halldórsson et al., 2005; Skarphéðinsdóttir et al., 2005).

Blue mussels (45–50 mm) were collected at low tide from the lower intertidal and transported moist and chilled to the laboratory in Sandgerði (<1 h transport time). Ten individuals from each site were immediately taken for evaluation of DNA single strand breaks, and the remaining mussels were placed in acclimation tanks upon arrival. The acclimation was held in running seawater (9.5 °C ± 0.3; salinity 32 ± 1) originating from a bore hole extending 50 m into the ground. The seawater, flowing through the porous rock (lava), is free from particles and anthropogenic material and was aerated upon arrival to the building. The blue mussels were allowed to acclimatize for 4 days before the measurements.

2.2. Cardiograph

Cardiographic traces of heartbeat were obtained by means of a non-invasive method derived from Depledge and Andersen (1990). The sensor consisted of an Infra-Red light emitting diode axially coupled with a phototransistor. For cardiographic measurements, sensors were glued externally to the mussel shell in a position corresponding to the heart. The signals from the phototransistor were filtered and amplified with appropriate circuits, passed on to a Fluke 105B portable oscilloscope and automatically saved on a computer at programmed intervals. Recordings of heartbeat traces (duration 20 s, sampling frequency 12.5 Hz) were obtained for each mussel at set intervals (60, 120 and 180 min).

2.3. Feeding rate

Measurements of the feeding rate of the mussels (the volume of water cleared of algal cells per hour) were made using static system according to the procedure described by Widdows and Staff (1997) with minor modification. Each individual mussel was placed in a container holding 5 L of seawater with known algal concentration, and the exponential decline in cell concentration (>3 µm) was recorded.

An algal culture of *Isochrysis galbana* was added to each container reaching an initial concentration of ~25,000 cells mL⁻¹. After the addition of phytoplankton 3 min were allowed for the algal cells to be thoroughly mixed into the water column using magnetic stirrers. Care was taken not to disturb the mussels by placing the stirrers at the opposite side of the tanks which were kept in water bath to keep the temperature constant at 9.5 °C. A 20 mL sample from the water column of each tank was taken to measure its initial cell concentration. Later, six samples were taken from each tank at 30 min interval over a period of 3 h. The concentration of algal cells in each sample was measured using an electronic particle counter (Coulter Counter, model Z1) with a 100 µm orifice tube, counting in a 0.5 mL sub-sample each time. Three replicate counts were made on each sample and their mean calculated. Individual feeding rate was then estimated using the following equation:

$$\text{Feeding rate} = \text{Vol} * (\log C_1 - \log C_2) / \text{time}$$

where Vol is the volume of water in each tank, C₁ is the cell concentration at the start of each time increment, and C₂ is the cell concentration at the end of each time increment.

2.4. Comet assay

The comet assay (single cell gel electrophoresis) was based on the protocol of Singh et al. (1988) with some modifications. Ten mussels (length 46 ± 4 mm) from each group were used for analyses of DNA single strand breaks in haemocytes. Haemocytes were used because they are rapidly sampled, demanding short slide preparation time, and cell dissociation is not required, keeping possible cell damage to a minimum (Klobucar et al., 2003). Furthermore, haemocytes circulate in the open vascular system and thus, are closely exposed to environmental agents through transportation of toxicants and various defence mechanisms (Mersch et al., 1996). They are currently used for monitoring PAH exposure in the field (Thomas et al., 2007).

Hemolymph (~ 0.2 mL) was collected from the posterior adductor muscle, mixed with CMFS and centrifuged at 2000 rpm for 3 min. After discarding the supernatant, the pellet was resuspended in 0.5% low melting point agarose (LMP) and two 85 μ L drops were placed on fully frosted slides, precoated with 0.75% normal melting point agarose. The samples were lysed for 2 h at 4 °C before they were randomly placed in the electrophoresis chamber and covered with chilled freshly made alkaline buffer (pH > 13). The DNA was allowed to unwind for 20 min before electrophoresis (0.65 V/cm, 300 mA) for 30 min at 12 °C. All work was carried out in dim light to avoid UV damage to the DNA. As positive controls, four samples from Hvassahraun were exposed to 50 μ M H₂O₂ for 1 h at 4 °C. The DNA was stained with 50 μ L of ethidium bromide (20 μ g ethidium bromide/mL distilled water) and examined with a fluorescence microscope (Leitz) coupled to a CCD camera and an image analysis package (Komet 5, Kinetic Imaging Ltd., Liverpool, U.K.). All slides were independently coded prior to examination and 100 (2×50) randomly selected cell nuclei were scored per slide. The number of DNA strand breaks are expressed as percentage of DNA in the tail (% Tail DNA). Highly damaged nuclei (“clouds”) were excluded since they presumably represent dead cells (Hartmann and Speit, 1997).

2.5. Benzo[a]pyrene exposure

Benzo[a]pyrene (CAS: 50-32-8, cat. no: 159069, purity >99%) was purchased from ICN Biomedicals Inc. (Aurora, Ohio, USA). The mussels were exposed to the nominal dose of 100 μ g B[a]P L⁻¹ dissolved in 0.1% v/v acetone, or as controls to either 0.1% v/v acetone alone or clean seawater.

2.6. Experimental setup

On arrival to the laboratory blood samples were taken from 10 blue mussels from each site for evaluation of the background levels of DNA single strand breaks with the comet assay. Other mussels to be used in the experiments were placed in acclimation tanks, without feeding.

Prior to the start of the first experiment the blue mussels were equipped with the IR sensors and individually positioned in 5 L acclimation tanks. After 60 min, cardiac activity of each mussel (three consecutive traces of 20 s each) was recorded, and the specimens then randomly assigned to control or experimental groups. In the first experiment 10 individuals from the clean site Hvassahraun were submerged in the two different solutions; clean water as control and solution of clean water and 0.1% v/v acetone, and their cardiac activity (three consecutive traces of 20 s each) was individually recorded after 60, 120 and 180 min.

In the second experiment 40 individuals were similarly maintained in the control or test solutions ($n=10$ for each site and solution; 100 μ g B[a]P L⁻¹ and 0.1% v/v acetone) for 3 h. After the addition of phytoplankton three minutes were allowed for the algal cells to be thoroughly mixed into the water column and a 20 mL sample was taken from each tank to measure its initial cell concentration. Additionally six samples were taken from the water column of each tank at 30 min intervals over a period of 3 h for feeding rate measurements. The cardiac activity of the mussels was recorded (three consecutive traces of 20 s each) after 60, 120 and 180 min.

After the last measurement of the cardiac activity, the blue mussels were removed from the containers and blood extracted from the posterior adductor muscle of each mussel and prepared for the comet assay (described above).

2.7. Data elaboration and statistical analysis

All data from the experiments were analysed by means of two factor ANOVA, with origin and treatments as fixed factors. Data were checked for normality, and homogeneity of variances was tested with Cochran's test. Post hoc comparisons were performed with the Student-Newman-Keuls test. No transformations were needed for the heart rate and feeding rate data. The comet assay data (mean % tail DNA) were log transformed to achieve normality and homogeneity of variances and one-way ANOVA was used to compare all group means (field and experiments). The level of significance was set as $p < 0.05$.

3. Results

3.1. Cardiac activity

The heart rate and consequently the metabolism of the mussels from the two sites differed considerably after the acclimatization period, before the addition of B[a]P (Fig. 1). Mussels from the clean site had lower heart rate (0.29 beats s⁻¹) than those from the polluted site (0.35 beats s⁻¹). These differences were highly significant (Table 1).

Secondly, the individuals from the two sites responded differently to the presence of B[a]P. The heart rate of the mussels from the clean site (Hvassahraun) did not change when exposed to the pollutant, while the specimens from the polluted site (Reykjavík harbor) responded significantly to B[a]P by increasing their heart rate (Fig. 1).

3.2. Feeding rate

Feeding rates (clearance of algal cells) were standardized to 1 g dry weight of soft tissue. Mussels from the clean site (Hvassahraun) had higher feeding rate (4.8 L h⁻¹ g⁻¹) than the mussels from the polluted site (Reykjavík harbour; 1.6 L h⁻¹ g⁻¹) and significant differences were explained only by the origin of the molluscs (Fig. 2 and Table 1).

The mussels from the clean site apparently showed a small decline in the feeding rate when exposed to B[a]P but it was 7% lower than for the control mussels of the same origin. On the contrary, mussels from Reykjavík harbour had, in presence of the toxicant, feeding rate 30% higher than the control mussels from the same site. These differences were, however, not significant when compared within sites.

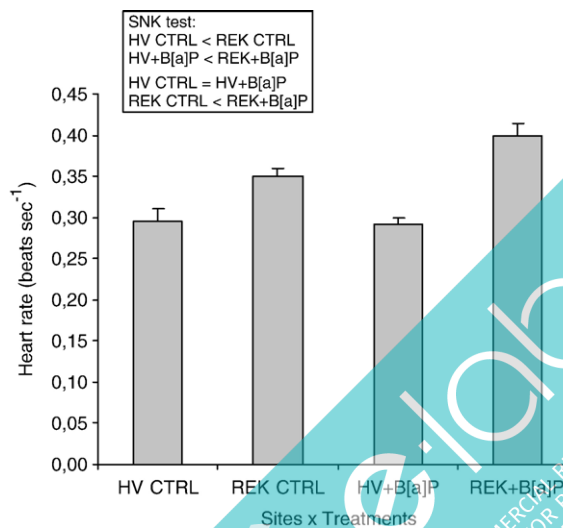


Fig. 1. Heart rate of the blue mussel (*Mytilus edulis*) during 3 h exposure to B[a]P. Values are expressed as beats s⁻¹ (mean \pm SD, $n=10$); HV = Hvassahraun clean site; REK = Reykjavík harbour; CTRL = 0.1% v/v acetone; B[a]P = 100 μ g benzo[a]pyrene L⁻¹ + 0.1% v/v acetone.

Table 1

Analysis of variance on the a) heart rate, b) feeding rate and c) DNA single strand break [$\log_e(x)$] results of the blue mussel (*Mytilus edulis*; $n=10$) from Hvassahraun (HV) and Reykjavík harbour (REK), with and without B[a]P treatment after 4 days of acclimatization

Source	df	MS	F	P
a) Heart rate				
Origin	1	0.065	40.39	***
Treatment	1	0.005	3.26	ns
Origin x Treatment	1	0.007	4.48	*
Residuals	36	0.002		
b) Feeding rate				
Origin	1	776.46	26.15	***
Treatment	1	0.05	0.02	ns
Origin x Treatment	1	18.45	0.62	ns
Residuals	36	29.69		
c) DNA single strand breaks				
Origin	1	0.014	0.085	ns
Treatment	1	0.238	1.404	ns
Origin x Treatment	1	0.229	1.349	ns
Residuals	3996	0.170		
Field vs. experiment				
Groups	5	1.492	8.432	***
Residuals	5994	118.2		

Post hoc comparisons were performed with the Student-Newman-Keuls test. * = $p < 0.05$; *** = $p < 0.001$; ns = not significant.

3.3. Comet assay

The amount of DNA single strand breaks (mean % tail DNA) in haemocytes was significantly higher in mussels from the polluted site (Reykjavík harbour; 12.26% tail DNA) compared to the mussels from the clean site (Hvassahraun; 9.65% tail DNA) when measured directly from the field (Fig. 3 and Table 1). After 4 days of acclimatization, and at the end of the experiment, the amount of DNA strand breaks was

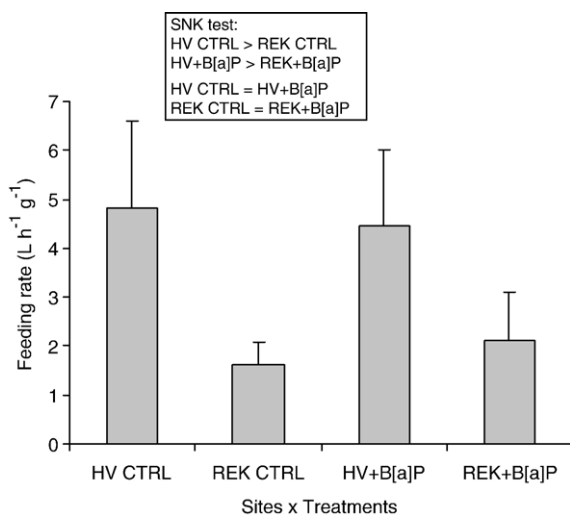


Fig. 2. Feeding rate of the blue mussel (*Mytilus edulis*) during 3 h exposure to B[a]P. Values are in $L h^{-1} g^{-1}$ (mean \pm SD, $n=10$). HV=Hvassahraun clean site; REK=Reykjavík harbour; CTRL=0.1% v/v acetone; B[a]P=100 μg benzo[a]pyrene L^{-1} + 0.1% v/v acetone.

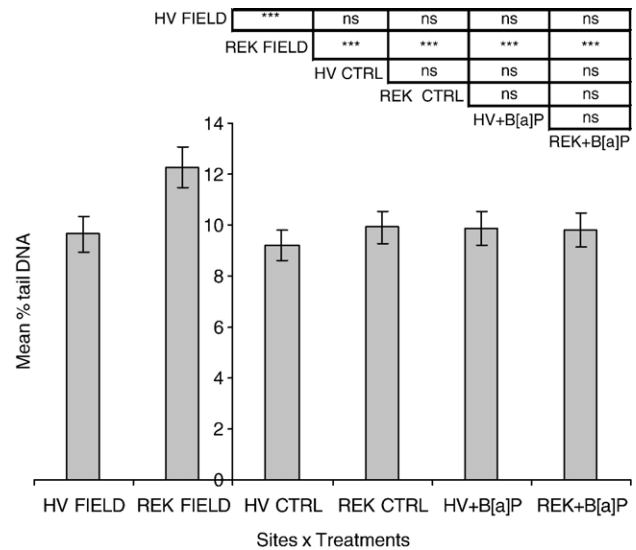


Fig. 3. DNA single strand breaks in haemocytes of the blue mussel (*Mytilus edulis*) measured before, and after acclimatization and 3 h exposure to B[a]P. Values are expressed as mean % tail DNA (mean \pm 95% C.I., $n=10$). HV=Hvassahraun clean site; REK=Reykjavík harbour; FIELD=No laboratory acclimatization before measurements; CTRL=0.1% v/v acetone; B[a]P=100 μg benzo[a]pyrene L^{-1} + 0.1% v/v acetone. SNK post hoc comparison between all samples. *** = $p < 0.001$; ns = not significant.

similar for both populations/treatments and similar to the initial level at the clean site.

4. Discussion

The clear difference in background pollution, as well as the short distance (~ 20 km) between the sites, provided the basis for comparing immediate responses of natural mussel populations to a PAH compound. The mussels from Reykjavík harbour were apparently under considerable stress, observed both at the cellular level (DNA damages) and physiological level (feeding rate and heart rate). The elevated levels of DNA single strand breaks in mussels sampled directly from the field indicate recent oxidative damages, presumably caused by PAHs and other toxic hydrocarbons in the harbour. In concordance, previous studies on mussels from the same sites revealed significant pollution (PAHs and Organotins) in Reykjavík harbour (Ericson et al., 2002; Halldórsson et al., 2004, 2005; Skarphéðinsdóttir et al., 2005; Da Ros et al., 2007). After 4 days in clean seawater the amount of detectable strand breaks had reduced and was similar as in mussels from the clean site. Although DNA repair can occur in matter of hours in bivalves (Pruski and Dixon, 2002; Villela et al., 2006), the recovery we observed could be explained by the repair of DNA damages and/or by replacement of damaged cells through apoptosis or cell death (Pruski and Dixon, 2002; Hook and Lee, 2004). These results support previous findings on rapid recovery of DNA strand breaks when organisms are allowed to acclimatize in a clean environment, stressing the need for rapid sampling when used in field monitoring. Increased heart rates and lowered feeding rates were still observed after the acclimation period for the harbour

mussels, suggesting ongoing removal of toxicants from the body.

The short time exposure to the nominal dose of 100 μg B[a]P L^{-1} did not exert DNA single strand breaks in haemocytes or lead to significant changes in the feeding rate of mussels from either site, although noteworthy trends in feeding rates were observed. The feeding rate of the mussels from the clean site decreased by 7%, while the mussels from the contaminated site showed about 30% increase in feeding rate in presence of the pollutant, suggesting elevated physiological activity. The most noticeable difference between the two sites was, however, seen in the cardiac activity of the blue mussels. The heart rate of the mussels from the clean site did not change when exposed to the high B[a]P concentration, while mussels from the contaminated site increased their heart rate significantly when exposed to B[a]P.

The accumulation rate of B[a]P by the blue mussel can be very rapid. Blue mussels exposed to 1 ppb B[a]P for 6 h, followed by 6 h in clean seawater, had tissue concentrations 70–80% of the initial B[a]P concentration in the water (Durand et al., 2002). Several studies have also shown acute effects of B[a]P on the DNA integrity of bivalves (Bihari and Fafandel, 2004; Siu et al., 2004; Kim and Hyun, 2006). These effects are mainly through metabolism of B[a]P to reactive intermediates (e.g. epoxides, quinones, cation radicals) leading to oxidative lesions (Mitchellmore et al., 1998). Siu et al. (2004) observed for example significantly increased DNA strand breaks in haemocytes of the green-lipped mussel (*Perna viridis*) after 1 day exposure to 30 μg B[a]P L^{-1} (measured by means of the comet assay). The exposure to B[a]P in the present study did, however, not generate DNA strand breaks in haemocytes despite the high nominal concentration. The exposure time was shorter in this experiment (3 h) compared to other short-term *in vivo* genotoxic studies using B[a]P (≥ 24 h), suggesting longer exposure time needed to exert genotoxic effects.

Reduced scope for growth of the blue mussel has been related to PAH exposure in the field, primarily due to reduction in feeding rate (i.e. clearance rate) (Widdows et al., 1997; Halldórsson et al., 2005). PAHs can exert narcotic effects on the gill ciliary activity, leading to decreased feeding rate, although many of the high molecular weight PAHs do not exert direct toxic effects (Donkin et al., 1989, 1991). However, the high molecular weight B[a]P has been shown to induce blood cell lysosomal membrane damages in the blue mussel, possibly due to direct toxic effect of B[a]P (Okay et al., 2000). Only minor effects were observed on the feeding rate in the present study, not indicating immediate toxic effects on the gill ciliary activity.

This study revealed an interesting difference in the heart rate response, depending on the background of the organisms. Firstly, mussels coming from the polluted site had higher baseline heart rates relative to the ones from the clean site. Similar results were found for other molluscan species and in a different ecological context (Nicholson, 1999; Wedderburn et al., 2000; Chelazzi et al., 2004). Secondly, when exposed to B[a]P the heart rate of the polluted mussels increased

significantly while no effects were seen in mussels from the clean site. Little information is available on the effects of PAHs on the heart rate of other marine molluscs. However, Galloway et al. (2002) found strong correlation between increased heart rates of the ribbed mussel (*Geukensia demissa*) and PAHs and PCBs in a field study. Effects of copper on the heart rate of the blue mussel are well documented but opposite to our findings the heart rate generally decreases in mussels exposed to unrealistically high levels of copper (e.g. Grace and Gainey, 1987; Curtis et al., 2000, 2001; Brown et al., 2004). Copper has been suggested to stimulate an inhibitory cholinergic neuronal pathway to the heart in the blue mussel, leading to reduced heart rate (Curtis et al., 2001). However, elevated heart rate at high copper levels has been observed for the green-lipped mussel (*P. viridis*) and increased heart rate of this species when transplanted to polluted coastal sites in Hong Kong was suggested to reflect enhanced metabolism during detoxification process (Nicholson, 2003; Nicholson and Lam, 2005).

In the present study the higher baseline heart rate of the blue mussels collected from the polluted site suggests ongoing detoxification process, but the fact that only these mussels responded to B[a]P does not indicate immediate toxic effects, but rather adaptive response to the contaminant, i.e. reflecting the pre-exposure history. The lack of heart rate response of the mussels from the clean site indicates a limited sensitivity of this biomarker, demonstrating that further studies are required for its application to pollution monitoring at high-latitudes. Although all the biomarkers revealed a significant difference between the study sites, the results from the laboratory experiments indicate limited sensitivity and temporal predictivity of these non-specific biomarkers. Thus, further investigation on the temporal predictivity, i.e. transient measurable changes of the heart rate and DNA strand breaks in haemocytes, is suggested. The ability of the mussels to modulate metabolism depending on the environmental background pollution demonstrates as well that the background of the organisms should be considered when evaluating short-term biomarker responses to contaminants.

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