

# Long term exposure of marine mussels to paracetamol: is time a healer or a killer?

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## Research Article

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# Abstract

Pharmaceuticals pose a major threat to the marine environment and several studies have recently described their negative effects on marine organisms. Pharmaceutical compounds are constantly being released into aquatic ecosystems and chronic exposure, even at low concentrations, may have impact on marine organisms. The purpose of the present study is to evaluate the biological changes induced by one of the most widely used pharmaceuticals – paracetamol – in the blue mussel *Mytilus edulis*, after a long-term exposure at environmentally relevant concentrations. We present our data alongside and in comparison with results from a previous short-term exposure, to demonstrate the significance of exposure period on the effects of paracetamol in adult blue mussels. After 24 days of laboratory exposure, seven potential target genes were selected to examine toxicological effects in mussels' gonads and possible disruptive effects on reproductive processes. The results show the modulation of some important reproduction-related genes: *estrogen receptor-2 (ER2)*, *vitelline envelope zona pellucida domain-9 (V9)* and *vitellogenin (VTG)*. Variations in mRNA expression of four other genes involved in apoptosis (*HSP70*, *CASP8*, *BCL2* and *FAS*) are also highlighted. Histopathological alterations caused by paracetamol, together with neutral red retention time response in mussels' hemocytes, are presented herein. Overall, this study highlights the exacerbated effects of low concentration of paracetamol after chronic exposure, similar to the damage induced by higher concentrations in a short exposure scenario, thus emphasizing the importance of length of exposure period when studying the effects of this substance. Additionally, this study also discusses the potential of paracetamol to inflict several major changes on the reproductive system of mussels and thus possibly affect the survival of populations.

## 1. Introduction

Global consumption of pharmaceutical products, prescribed or over the counter, is projected to rise in the coming years (Fabbri and Franzellitti, 2016; aus der Beek et al., 2016). Pharmaceutical consumption is estimated at more than 200,000 tonnes per year in Russia, China and India (Tijani et al., 2016), and is expected to be much higher for regions with high use of pharmaceuticals such as USA and Europe. Paracetamol or acetaminophen is an active ingredient in hundreds of prescriptions and over the counter (OTC) medicines (Roberts et al., 2016). At present, paracetamol is used as an analgesic and antipyretic, and there is a significant prevalence for self-medication worldwide (Tariq and Din, 2017), especially in the context of the COVID-19 pandemic. It is sold both as paracetamol only and, in many OTC medicines, is also combined with other active ingredients that treat cough, colds, flu, and pain-related conditions (Wood et al., 2010). Paracetamol is recognised as the most frequently used analgesic in the UK (Bertolini et al., 2006), and is also the world's most widely marketed OTC drug (Warwick, 2008), mostly because of its non-prescription availability and low cost (Jozwiak-Bebenista and Nowak, 2014; B rzsaga Arencibia and Choonara, 2012). Nevertheless, the consumption is set to rise significantly during the COVID-19 pandemic.

An increasing number of ecotoxicology studies show that pharmaceuticals pose a risk to aquatic organisms. This risk is evidenced, not only by the sustained persistence of pharmaceuticals in different

aquatic compartments, but also by their bioaccumulation in many species, as documented by Mimeault et al. (2005), Vernouillet et al. (2010), Wang & Gardinali (2013), Brodin et al. (2014), Du et al. (2015) and de Solla et al. (2016). Many studies focusing on the adverse outcomes of pharmaceuticals have been recorded in freshwater species (Flammarion et al., 2000; Hoeger et al., 2005; Mimeault et al., 2005), whereas only very limited data have been reported in marine organisms.

Pharmaceuticals in general can exert considerable pressure on reproductive associated mechanisms in aquatic organisms, leading to knock-on ecological effects on populations and communities. Franzellitti et al. (2013) reported that fluoxetine, a common antidepressant, was associated with many detrimental effects on reproduction and other major physiological systems (including reproduction) in Mediterranean mussels *Mytilus galloprovincialis*, even at concentrations below or approaching environmental levels. Fonseca et al. (2019) showed that tamoxifen, the oldest hormone therapy for breast cancer, can cause endocrine disruption in male *M. galloprovincialis* exposed only for 14 days. Niemuth & Klaper (2015) documented that metformin can cause intersex and a reduction of fecundity in fathead minnows *Pimephales promelas*. A recent study by Koagouw & Ciocan (2018) also recorded pathologies in the gonads and increased *vitellogenin* mRNA expression of blue mussels *Mytilus edulis* exposed to metformin.

As the ocean acts as the ultimate receptacle of a vast quantity of natural and anthropogenic waste that is continuously emitted from urban and industrial sources (Norse and Crowder, 2005; Pereira et al., 2016), marine organisms are potentially at critical risk. A review by Ebele et al. (2017) highlights the main routes of pharmaceutical pollution as: sewage treatment plants (STPs), wastewater treatment plants (WWTPs) and landfill leaching, while other sources include industrial manufacturing, agriculture, veterinary use and body excretion after consumption. The authors also emphasize that pharmaceuticals are often persistent and frequently found in surface water at concentrations ranging from ng/L to mg/L, whilst low concentrations of these pollutants are detected even in drinking water (Kasprzyk-Hordern et al., 2008; Caban et al., 2015). The persistence of pharmaceuticals in the aquatic environment allows for scenarios of chronic exposure, highlighting the potential importance of exposure time in determining the severity of pharmaceutical impact on non-target organisms. The relative importance of contaminant concentration versus exposure duration has so far been little studied, although some studies have taken exposure duration into consideration when considering the impacts of contaminants (Cope et al., 2008; Huang et al., 2019).

While paracetamol has been detected in various aquatic environments at concentrations ranging from 3.3 ng/L (Fairbairn et al., 2016) to 16 µg/L (Agunbiade and Moodley, 2014), the levels reported in seawater vary from 3.2 ng/L (Benotti and Brownawell, 2007) to more than 200 µg/L (Togola and Budzinski, 2008). The continuous high consumption and production of paracetamol as well as its evident occurrence in seawater give rise to concerns regarding the impact on marine organisms, especially filter feeders. This study explores the impacts of paracetamol on the gonads of marine mussels *Mytilus edulis* after a long-term exposure, and discusses the potential reproductive challenges that may arise. Here, we present our data alongside and in comparison with results from a previous short-term exposure

(Koagouw and Ciocan, 2019), in order to demonstrate the importance of the length of exposure on the potential biological and ecological damage inflicted by paracetamol.

Mussels are excellent indicator organisms for environmental monitoring and have been intensively used worldwide to monitor marine pollution (Rittschof and McClellan-Green, 2005). Representatives of Mytilidae such as *M. edulis* and *M. galloprovincialis* are also widely used as indicators in several studies on the effects of pharmaceuticals, due to their well-known physiology and their wide geographical distribution (Świacka et al., 2019). In this study, we employed neutral red retention time assay to enable observation of the effects at cellular level. Histopathological examination was performed to determine sex and any pathological conditions observed in the gonad tissue. Three genes related to reproduction – *vitellogenin (VTG)*, *vitelline envelope zona pellucida domain-9 (V9)* and *estrogen receptor-2 (ER2)* – were investigated, as well as four genes involved in apoptosis: *heat shock protein-70 (HSP70)*, *caspase-8 (CASP8)*, *B-cell lymphoma-2 (BCL2)* and *Fas cell surface death receptor (FAS)*. This study is highly pertinent in the context of a potential increase of paracetamol in seawater following the COVID-19 pandemic

## 2. Materials And Methods

### 2.1. Sample collection

Blue mussels *M. edulis* were collected by hand from a single population located in Hove Beach, East Sussex, UK (50.823797, -0.173423) at low tide during April 2018. Mussels were placed on ice following collection and directly transported to the laboratory, where they were then washed and stored in an artificial seawater container (Instant Ocean® Sea Salt, USA) for acclimatisation purposes. During this time, the mussels were fed each day with 500 µL of green algae *Tetraselmis sp.* culture suspension (ReefBoost, UK) per 5 litres of artificial seawater. The starting water temperature (15 °C) was then slowly increased over the following 6 days to a steady experimental limit of  $20 \pm 2$  °C.

### 2.2. Experimental exposure

All procedures were performed in compliance with the ARRIVE guidelines and carried out in accordance with the UK Animals (Scientific Procedures) Act 1986 and EU Directive 2010/63/EU on the protection of animals used for scientific purposes. The procedures have been approved by the Animal Welfare and Ethics Review Bodies (AWERB), University of Brighton. Only mussels between 30 and 50 mm in length were used in exposure experiments. Artificial seawater was prepared in compliance with the manufacturer's instructions, and tanks used for exposures contained approx. 1 litre of artificial seawater per mussel. Exposures consisted of a control (artificial seawater only) and three separate treatments of paracetamol (40 ng/L, 250 ng/L and 100 µg/L). These nominal concentrations were based on concentrations detected in the marine environment by previous studies (Togola and Budzinski, 2008; Nödler et al., 2014; Bebianno et al., 2017). Twice weekly (at 72 and 96 hours) artificial seawater was renewed, and the exposed groups were treated with paracetamol (BioXtra, ≥99.0%, Sigma-Aldrich).

Physical characteristics of seawater (temperature, salinity, conductivity and resistance) were monitored daily and the exposure was suspended after 24 days. All experimental tanks were set up in duplicate.

A total of 10 mussels from each group were collected, measured and dissected following the completion of the exposure. Approx. 1 cm square of each gonad was fixed in neutral buffered formaldehyde in clean tubes and preserved at 4 °C for the purposes of histological analysis. For the molecular analysis, the tissues were immediately transferred to RNA $later$  (Invitrogen, UK) and kept at -80 °C.

### 2.3. Water analysis

The detailed protocol is outlined in Koagouw & Ciocan (2019). Water samples (1000 mL) were obtained from each group 15-30 minutes after paracetamol was added ( $t_0$ ) and immediately before artificial seawater was changed (72 and 96 hours). All samples were processed through solid phase extraction (Strata™-XL-C 100  $\mu$ m polymeric strong cation 2 g / 20 mL giga tube cartridge, Phenomenex, USA) after two filtrations using 1.2  $\mu$ m Whatman grade GF / C microfiber glass filter paper (GE Healthcare, UK) and 0.22  $\mu$ m nylon membrane filter (GE Healthcare, UK). The extract was evaporated by centrifugation under vacuum (Speedvac, Savant) and reconstituted with LC-MS grade water prior to analysis.

The concentration of paracetamol was determined by liquid chromatography-mass spectrometry (LC-MS) using a standard curve. Paracetamol separation was performed by ultra-high performance LC (Ultimate 3000, Thermo Scientific) using reversed phase chromatography (Kinetex XB-C18, 5  $\mu$ m, 100 Å, 100 x 2.1 mm, with trap column, Phenomenex, UK). Mass spectrometry (Orbitrap Q Exactive, Thermo Scientific) was conducted in positive mode using heated electrospray ionization (HESI) with a probe temperature of 200 °C. The area below the peak was defined by the reconstructed ion chromatogram of the fragment at 110.0602  $m/z$  and quantitation was determined using Quan Browser data processing software (Xcalibur V:4.1.31.9, Thermo Scientific). The detailed parameters of this procedure are described in Koagouw & Ciocan (2019).

### 2.4. Neutral red retention time (NRRT) assay

The NRRT procedure was adapted from Lowe & Pipe (1994) and Lowe et al. (1995) in Mamaca et al. (2005). At the end of the exposure, hemolymph of mussels ( $n = 3$ ) from each group was withdrawn using a syringe containing physiological saline solution (ratio 1:1), and then transferred to clean tubes. 30  $\mu$ l of hemolymph-saline mixture was transferred onto the poly-L-lysine coated microscope slide, followed by 30  $\mu$ l of the neutral red working solution, and incubated in a light-proof humid chamber for 15 minutes at room temperature ( $t_0$ ). Each slide was observed at 30-minute intervals for a total of 180 minutes using light microscopy (Leitz Wetzlar, Germany) (40x/100x); the slide was returned to the humid chamber after each observation. The observations were terminated, and the retention time recorded when 50% of the small granular hemocytes visibly leaked their dye into the cytosol.

### 2.5. Tissue preparation for histological examination

Histological examination was used both for the purposes of assessing the sex of individuals and for analysing any pathological conditions found in mussel gonads. The analysis was performed according to Koagouw & Ciocan (2019). The 7 µm slices cut from paraffin embedded blocks were stained with hematoxylin and eosin. Histological evaluation of tissue was conducted under light microscopy (Leitz Wetzlar, Germany) (40x/100x) and histopathological conditions were documented along with micrographs referring to each condition, using GXCam Hichrome-Lite (GT Vision, UK).

## 2.6. Gene expression analysis

The analyses were performed following the methodology of Koagouw & Ciocan (2019).

### 2.6.1. RNA extraction and cDNA synthesis

Total RNA from the gonads (n = 10 for each experimental group) was individually isolated using SurePrep™ TrueTotal™ RNA Purification Kits (Fisher Scientific, UK) and Monarch® Total RNA Miniprep Kit (New England Biolabs, UK) following manufacturer instructions. Qubit™ RNA HS Assay Kit (Invitrogen, UK) and Qubit® Fluorometer was used to quantify the extracted RNA concentration. The cDNA synthesis was carried out with Transcriptor High Fidelity cDNA Synthesis Kit (Roche, UK) as per manufacturer's instructions and the complementary DNA (cDNA) concentration in each sample was measured using Qubit™ dsDNA HS Assay Kit and Qubit® Fluorometer (Invitrogen™, UK).

### 2.6.2. Quantitative real time PCR

Molecular analysis was performed to investigate potential changes in the pattern of expression of selected transcripts, as described by Koagouw & Ciocan (2019): *vitellogenin (VTG)*, *vitelline envelope zona pellucida domain-9 (V9)*, *estrogen receptor-2 (ER2)*, *heat shock protein-70 (HSP70)*, *caspase-8 (CASP8)*, *B-cell lymphoma-2 (BCL2)* and *Fas cell surface death receptor (FAS)*. The same primers as in Koagouw & Ciocan (2019) were used for this study (Table S1).

## 2.7. Data analysis

The average cycle quantification (*C<sub>q</sub>*) of reference genes *18S rRNA* and *EF1*, as suggested by Cubero-Leon et al. (2012), was used as normalization factor. The computation of the relative changes in the target gene expression identified by real-time qPCR applied the comparative  $2^{-\Delta\Delta C_t}$  method, expressed as fold changes to the control group as defined by Livak and Schmittgen (2001).

Statistical analysis was performed using GraphPad Prism 8. One-way analyses of variance (ANOVA), followed by Tukey's post-hoc multiple comparison tests were performed to identify significant differences ( $p < 0.05$ ) between groups.

# 3. Results

## 3.1. Water analysis

The percentage reduction of paracetamol in treatment groups after 72 and 96 hours is depicted in Figure 1. Overall, the reduction ranged from 19% to nearly 100% after 72 hours of exposure, with an average of 61%. After 96 hours, the reduction was 63-83% with an overall average of 72%.

### 3.2. Neutral red retention time assay

The lysosome membrane integrity of hemocytes expressed as neutral red retention time is displayed in Figure 2. All exposed groups showed a significant decrease in the ability to contain the dye within the lysosomes, relative to control; however, there was no statistically significant difference between the paracetamol treatments.

### 3.3. Histopathology observation

Follicle dilatation was recorded as the most widespread pathology in the exposure groups (Figure 3). This pathological condition was observed in all exposed groups, at a prevalence of 70-80%, with the highest frequency in the 100 µg/L group. Gamete degeneration is another pathological condition that showed a high occurrence, being detected in 50-60% of individuals in all groups exposed to paracetamol. Inflammatory pathologies such as hemocytic infiltration and hemocytic aggregate were also observed in as many as 50% and 30% of individuals, respectively. Female mussels exposed to the lowest concentration of paracetamol 40 ng/L showed a high incidence of atretic condition in their gonads up to 40%. Parasitic infestation was observed in all treatment groups, although with low prevalence (10-20%).

The micrographs of histopathological conditions documented during observation are shown in Figure 4.

### 3.4. mRNA expression analysis

The mRNA expression of each target gene is shown in Figure 5 and is expressed here as fold changes to the control group (with control group standardised to 1); down regulation is represented by values below 1 and up regulation by values above 1. The data presented here are plotted alongside results recorded by Koagouw & Ciocan (2019), in order to ensure a direct comparison between variation in mRNA expression in short and long exposures of paracetamol.

*VTG* mRNA expression was down regulated in all exposed groups by 4-11 fold changes compared to the control group, although the responses were not significantly different between the treatment groups. A similar trend was previously observed in the short exposure data (Koagouw and Ciocan, 2019). A very drastic down regulation of expression was also observed in *V9* mRNA transcript following 24 days' exposure. Whilst the response recorded for this transcript followed a dose-dependent trend in the short exposure experiment, a longer exposure to paracetamol induced a more severe gene silencing, ranging from 8-17 fold changes, which was similar in all exposed groups.

The mRNA expression of *ER2* showed a more severe suppression in the long exposure experiment compared to the short exposure (up to 33 fold changes compared to 4-12), whilst the *HSP70*, *BCL2* and *FAS* data behaved in a similar manner: following long term exposure, gene expression was heavily

suppressed in all treatment groups. The transcription patterns were concentration dependent in short-term exposure data, however, the long term exposure to paracetamol seems to exacerbate the effects, with *FAS* transcript down regulated by 11-100 fold changes compared to the control group. For *CASP8*, all exposed groups in the short- and long-term exposures showed a similar pattern of down regulation.

## 4. Discussion

### 4.1. Reduction of paracetamol in artificial seawater

A substantial depletion in paracetamol content was observed after 72 hours, while after 96 hours the decrease is likely to have balanced to equilibrium, and reduction occurred at a steadier rate. The pattern of reduction here may be associated with the amount of paracetamol that could be absorbed by mussels per day. The data here, however, only show the presence and the reduction trend in the artificial seawater, and further analyses to validate the quantification of the contaminant in mussels are needed, to confirm the absorption of paracetamol.

At 72 hours of exposure, the reduction of paracetamol levels exhibited a monotonic response, before transitioning to a more U-shaped pattern of reduction after this point in time. U-shaped curves are very common in toxicological studies (Davis and Svendsgaard, 1990; Calabrese, 2008; Douron, 2010) and are a very important consideration in toxicological and environmental health risk assessments, especially in case of no-observed-effect levels. The transition in the reduction pattern of paracetamol here after 72 hours, therefore, may provide important information on the interpretation of ecotoxicology assessment of this pharmaceutical in mussels, when the observed effects have different patterns (monotonic vs non-monotonic) or do not display expected effects due to the different exposure time.

### 4.2. Neutral red retention time of mussel hemocytes

Lysosomal membrane stability is a very sensitive indicator of cellular damage, because lysosomes represent the main cellular site for sequestration and detoxification of contaminants (Dailianis et al., 2003). The principle of NRRT assay is simple: neutral red dye enters the cell and is taken up and retained by lysosomes in healthy cells. In stressed mussels, this process is measured using NRRT, and depicts the integrity and capacity of the lysosomal membrane to retain the dye for a period of time. Therefore, the assay measures the ability of cytological processes to adjust to stress conditions (Lowe and Pipe, 1994). In other words, mussels that have low immunity will respond by showing a lower retention time of the toxicant, in this case neutral red, as a result of the lysosomal membrane destabilization.

In this study, exposed mussels showed a shorter retention time compared to a control, however no significant difference was recorded between different treatment groups. Parolini et al. (2009) employed neutral red retention assay in investigating the effects of three NSAIDs on the hemocytes of zebra mussel (*Dreissena polymorpha*) and demonstrated that paracetamol was the lowest in the toxicity scale compared to diclofenac and ibuprofen. Our results suggest that although paracetamol exerts an effect on

mussel (*M. edulis*) hemocytes, a higher dose of this particular contaminant does not necessarily increase the biological response, probable due to its lower toxicity as suggested by Parolini et al. (2009).

Also, worth noting here is the relationship between retention time and the reduction of paracetamol in exposed groups. The percentage reduction of paracetamol in artificial seawater at 96 hours was observed to be highly correlated with the retention time in each treatment group. Additionally, the lower retention time in mussels groups 40 ng/L, 250 ng/L and 100 µg/L can be associated with the parasitic incidence in each treatment group, suggesting a strong link between the lower immune response recorded here and the susceptibility of mussels to parasitic infection.

### **4.3. Histological alterations in the gonads of mussels after exposure**

In mussels exposed to paracetamol for 24 days, follicle dilatation was observed as the most commonly occurring pathological condition in the gonads. Such a prevalence raises concerns regarding mussels' reproductive health, as suggested by Sunila (1987). It is worth mentioning that even exposures at low concentrations of paracetamol, within the range frequently reported in the marine environment, have the capacity to induce widespread follicle dilatation in mussel gonads (Koagouw and Ciocan, 2019).

In mammals, more than 90% of follicles undergo a degenerative process as part of their developmental cycle (Kerr et al., 2013), with some indications that this degenerative stage is induced by apoptosis of granulosa cells, which are influenced by a precarious balance of pro-survival factor withdrawal and pro-apoptotic factors (Manabe et al., 2004; Hatzirodos et al., 2014; Zhang et al., 2018). A study by García-Gasca et al. (2010) also suggested that this condition might be a useful predictor of environmental stress for coastal ecosystems, as it is directly related to the reproductive system and the success of reproduction. The high prevalence of follicle dilatation in the results presented here implies a considerable potential for paracetamol to disturb mussel reproduction and possibly to interfere with population sustainability.

Gamete degeneration was also one of the most prevalent histological conditions in mussels exposed to paracetamol. Boumela et al. (2009) suggest that the quality of gametes is important not only for gamete survival rates but also for the early stages of embryo development, thus representing a key to successful reproduction. Rouabhi et al. (2019) studied the reproductive cycle of Mediterranean mussels *M. galloprovincialis* from a contaminated coast in Algeria, and inferred that gamete degeneration and spawning cessation due to coastal pollution and global warming could endanger recruitment of mussels and, eventually, the shellfish industry as a whole. In this study, a high incidence of gamete degeneration occurred after 24 days' exposure to paracetamol (Fig. 3). This raises concerns regarding the possibility of impaired development of mussels, and eventually the survival of populations themselves, when this pharmaceutical is present in seawater for at least that period of time.

In bivalves, hemocytic infiltration has been linked with the immune responses to stress-related events, especially infectious diseases (Allam and Raftos, 2015). Several studies have shown that hemocytic infiltration is usually associated with inflammatory responses in organisms, and elicits profound

detrimental effects in some species. Recent research (Gornati et al., 2016) suggested that titanium dioxide nanoparticle exposure led to hemocytic infiltration in *Mytilus galloprovincialis*. Hemocytic infiltration was also prevalent in *Mytilus galloprovincialis* used in field monitoring to assess the effects of an oil spill along the North coast of the Iberian Peninsula (Garmendia et al., 2011). These results imply that environmental contaminants are able to trigger immunological responses, and can also cause immunological changes by affecting cellular energy metabolism. Hemocytic infiltration shows a very interesting pattern in our study, whereby the occurrence displays the opposite trend to the concentration (Fig. 3). A previous 7-day exposure experiment (Koagouw and Ciocan, 2019) showed a dose-related trend, with a higher incidence of hemocytic infiltration in higher concentration groups, while our longer exposure data suggest a decline in the prevalence of this condition. While the adaptability of mussels to tolerate the contaminant might be a factor, this result is more likely linked to the fact that mussels exposed to the highest concentration of paracetamol are more susceptible to hemocytic aggregate condition (30%), a more severe pathology characterized by the formation of hemocyte clusters (Auffret and Oubella, 1997; Garmendia et al., 2011).

An interesting result worth noting here is that all the pathological conditions were detected in almost all exposure groups, from the lowest to the highest concentration, confirming the potential damaging effect of paracetamol even at lower levels of contamination. Overall, the results here present a concerning picture, suggesting paracetamol concentrations as low as 40 ng/L can induce almost the same adverse effects caused by a concentration 2,500 times higher, given a long exposure scenario.

#### **4.4. The variation of mRNA expression of target genes in mussel gonads**

After a long-term exposure to paracetamol, the mRNA expression of *VTG* was equally down regulated in all exposed groups although in a non-monotonic dose response pattern, suggesting that the lowest concentration of paracetamol can induce a similar effect to a concentration 2,500 times higher (100 µg/L). *Vitellogenin (VTG)* mRNA expression is a sensitive marker for early assessment of contamination by endocrine disrupting chemicals (EDCs) in vertebrates (Hutchinson et al., 2006; Barucca et al., 2006; Sugawara, 2011; Kim et al., 2012). In invertebrates, although its mechanism of action, synthesis and function are still undefined and require further study (Matozzo et al., 2008; Porte et al., 2006), several studies have identified and recorded induction of vitellogenin following exposure to EDCs (Ciocan et al., 2010; Jubeaux et al., 2012). Similar to short exposure data (Koagouw and Ciocan, 2019), the long term exposure results indicate that even the lowest concentration of paracetamol (40 ng/L) can induce extreme downregulation in *VTG* irrespective of the length of exposure to paracetamol. This sensitivity indicates the potential for this target gene to be further explored as a biomarker for ecotoxicological studies on paracetamol.

A different pattern of the *vitelline envelope zona pellucida domain-9 (V9)* down regulation was recorded in mussels exposed to paracetamol for 24 days when compared to a shorter 7-days exposure (Koagouw and Ciocan, 2019). The *V9* mRNA expression results in all 24-days treatment groups were significantly lower compared to the control ( $p < 0.05$ ). The results presented here indicate that longer exposure to low

concentration of paracetamol has the potential to inflict the same level of biological responses as brief exposures to high concentrations. Considering that pharmaceuticals are constantly released into the aquatic environment and therefore the organisms are in prolonged contact with the contaminants, our results can be considered environmentally relevant.

Vitelline envelope or zona pellucida is known to play a fundamental role in various aspects of fertilisation, such as mediating the sperm binding process (Snell and White, 1996) and protecting against polyspermy (Coy et al., 2008). In addition, this protein has been advanced as a potential biomarker for environmental estrogens in fish (Celius and Walther, 1998; Celius et al., 1999). The paracetamol modulated *V9* expression presented herein suggests that reproductive impairment in mussels can potentially result from exposure to low concentrations similar to those detected in the natural environment.

In bivalves, since sex-related alterations may primarily be mediated by sex steroid receptors, expression variability in estrogen receptors indicates possible consequences that may occur in gametogenesis and reproductive processes (Croll and Wang, 2007). In this study, *ER2* mRNA expression was down regulated in all treatment groups, in a non-monotonic dose response pattern (Fig. 5c). The data suggest that a long exposure to paracetamol intensifies the effects by approximately 13-29 folds in all long term exposure tanks, compared to short exposure. As estrogenic activities are facilitated through estrogen receptors by regulating the target gene expression (Gao and Dahlman-Wright, 2011), the results presented here thus raise concern over the potential of paracetamol to induce disruption of estrogen modulated transcripts. In the 7-day exposure, male mussels were highly affected and displayed a monotonic trend of response in the *ER2* gene expression (Koagouw and Ciocan, 2019). The long term exposure data suggest similar levels of downregulation regardless of the sex of mussels and paracetamol concentration, thus a more widespread effect. Several studies have documented the expression of estrogen receptors in different invertebrates (Keay and Thornton, 2009; Jones et al., 2017), in addition to mussels (Puinean et al., 2006; Agnese et al., 2019; Balbi et al., 2019). However, it is also worth considering that *ER2* might display natural variation during different stages of gametogenesis (Ciocan et al., 2010), hence further investigation is required.

The expression of several target genes involved in apoptosis was investigated after long and short exposure to paracetamol. In this study, the mRNA expression of *CASP8* was down regulated by even very low concentrations of paracetamol present in the seawater, in both short- and long-term exposures (Fig. 5e). Reduced mRNA expression of *caspase-8* (*CASP8*) may bring about changes in the deterioration process of cellular components, acting to inhibit apoptosis (Kruidering and Evan, 2000; Romero et al., 2011). Ruocco et al. (2016) documented the activation of caspase-8 and changes in the expression level of *CASP8* that led to apoptosis in sea urchin embryos after exposure to oxylipins, diatom secondary metabolites. *CASP8* may also play an important role in programmed cell death, meaning its downregulation could contribute to cancer-related pathologies (Aghababazadeh et al., 2017). This suggests that paracetamol can potentially be involved in carcinogenic pathways or in any pathological condition caused by the down regulation of *CASP8* in mussels. As *CASP8* is one of the important genes

involved in apoptosis, its extreme sensitivity to paracetamol poses a real threat, possibly further highlighted by apoptosis-related pathologies recorded during histological examination (Fig. 3 and 4) of the reproductive system in mussels. Moreover, the high sensitivity of *CASP8* as demonstrated here makes this transcript recommended as a potential biomarker for the monitoring of paracetamol contamination in the environment.

The expression of *HSP70*, *BCL2* and *FAS* in this study were down regulated in all treatment groups following non-monotonic responses after 24 days' exposure, while a shorter exposure elicited a dose-dependent response (Fig. 5). Here, the different patterns of expression highlight the significance of exposure period on the modulation of these regulatory genes: they suggest a longer period of paracetamol exposure can induce similar levels of down regulation in mussels exposed to both low and high concentrations of this substance. The dramatic changes in response of all three transcripts in the low concentration group again suggest that long-term effects are even more deleterious than short-term effects. These results should be of great concern considering the presence of paracetamol in the environment will be very likely to persist in the long term, as a result of urban waste agglomeration, and especially in the very recent special case of the COVID-19 pandemic.

As the healthy cell death balance is maintained by the modulation of apoptosis-regulatory gene transcripts through the activation or inhibition of apoptosis (Kiss, 2010), alterations to this process may either lead to the progression of cell death or the survival of defective cells, which can contribute to carcinogenesis. The significant changes in expression of *HSP70*, *BCL2* and *FAS* in this study indicate the possible threat that might be faced by mussel populations, in particular through apoptosis-related mechanisms in the reproductive organs. While the suppression of *HSP70* in paracetamol-exposed mussels as observed in this study may lead to a higher risk of apoptosis and cell death, the modulation of both *BCL2* and *FAS* presented here (Fig. 5) can induce either apoptosis or cancer-related conditions in gonad cells. This particular apoptosis-related alteration is perfectly depicted in the histopathology result, where degeneration in follicles and gametes is evident in mussels' gonads (Fig. 3 and 4c-d). The presence of the apoptosis related pathologies, with high prevalences across all levels and durations of exposure, is consistent with the pattern of downregulation observed in target genes involved in apoptosis.

In other studies, apoptosis in molluscs (Kiss, 2010) and in particular mussels (Estévez-Calvar et al., 2013; García-Gasca et al., 2010) has been proposed as a potential biomarker to monitor environmental stress in marine ecosystems. Moreover, Estévez-Calvar et al. (2013) specifically confirmed that expression of the genes involved in apoptosis can be harnessed to assess the stress related biological responses of coastal species. Characterizing the expression of these apoptotic regulatory genes is therefore crucial as a first step towards the development of potential biomarkers.

Overall, our study demonstrates the importance of exposure duration, in that longer exposure to paracetamol could magnify its effects in adult mussels. Different responses following short- and long-term exposures to contaminants have also been recorded by several studies, in vertebrates and invertebrates. Saravanan et al. (2014) reported different responses of thyroxine in carp *Cirrhinus mrigala*

after short- and long-term exposure to diclofenac and clofibrac acid. The authors showed that thyroxine level decreased only in two treatment groups exposed to clofibrac acid (and not in the diclofenac groups) following short-term exposure, but longer duration of exposure decreased thyroxine levels in all concentrations of both contaminants. In invertebrates, Oliveira et al. (2017) investigated physiological and biochemical alterations in mussels *Mytilus galloprovincialis* after short- and long-term exposure to carbamazepine. The authors revealed that among all physiological parameters studied, the condition and gonadosomatic indices were mostly adversely affected by long-term exposure to carbamazepine, and further concluded that this alteration could compromise the reproductive potential of organisms with implications for the survival of the population.

As the duration of exposure plays a significant role in determining the severity of effects, it is important to consider the length of contact time in assessing the effects of a given contaminant, in our case paracetamol. Our study would suggest that the assessment of a contaminant's effects should take into account not only its concentration or level in the environment, but also the length of exposure encountered by organisms.

## 5. Conclusion

Paracetamol does not appear to have a consequential impact on hemocytes, especially in terms of lysosomal membrane stability. However, more dramatic results are observed in the gonad tissue, where paracetamol is seen to induce major adverse changes, such as degeneration in follicles and gametes. Such changes pose a risk to the reproductive ability of this organism, and therefore a potential impact on population survival.

Paracetamol induced damage was also recorded at the molecular level. Patterns observed in mRNA expression after long exposure to paracetamol as observed in this study demonstrate the importance of length of exposure on the biological responses elicited in blue mussels. Overall, the expression of all transcripts investigated herein suggest that longer exposure to a contaminant may result in much greater effects, potentially equivalent to those registered after exposure to much higher concentrations. These findings indicate that the presence of paracetamol in the environment even at low concentrations has the potential to cause several major changes related to the reproductive system of mussels. Further investigations exploring and confirming some related aspects, especially the mechanism of action, are encouraged.

Over recent years, there has been growing concern regarding chronic pharmaceutical pollution of the aquatic environment. As rising human populations coincide with a higher demand for pharmaceuticals, their continuous emission is very likely to persist. The situation is likely to deteriorate further in the current pandemic, as pharmaceutical companies have seen an enormous demand for pain relief medicine, particularly paracetamol. Some of these will be ingested and then excreted, but it is predicted that a large amount will be disposed of, and will most likely end up in the natural environment in vast quantities.

Long exposure studies are therefore environmentally relevant, and our results demonstrate that at least for paracetamol, the effects are exacerbated in a chronic exposure scenario.

## Declarations

### Ethics approval and consent to participate

All procedures in this paper were performed in compliance with the ARRIVE guidelines and carried out in accordance with the UK Animals (Scientific Procedures) Act 1986 and EU Directive 2010/63/EU on the protection of animals used for scientific purposes. The procedures have been approved by the Animal Welfare and Ethics Review Bodies (AWERB), University of Brighton.

### Consent for publication

Not applicable

### Availability of data and materials

All data generated or analysed during this study are included in this published article and its supplementary information files.

### Competing interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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### Author contribution statement

Wulan Koagouw : Conceptualization, Methodology, Formal analysis, Investigation, Resources, Writing - Original Draft

Nicolas A. Stewart : Investigation, Resources, Writing - Original Draft

Corina Ciocan : Conceptualization, Methodology, Resources, Writing - Original Draft, Supervision

All authors read and approved the final manuscript.

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## Figures

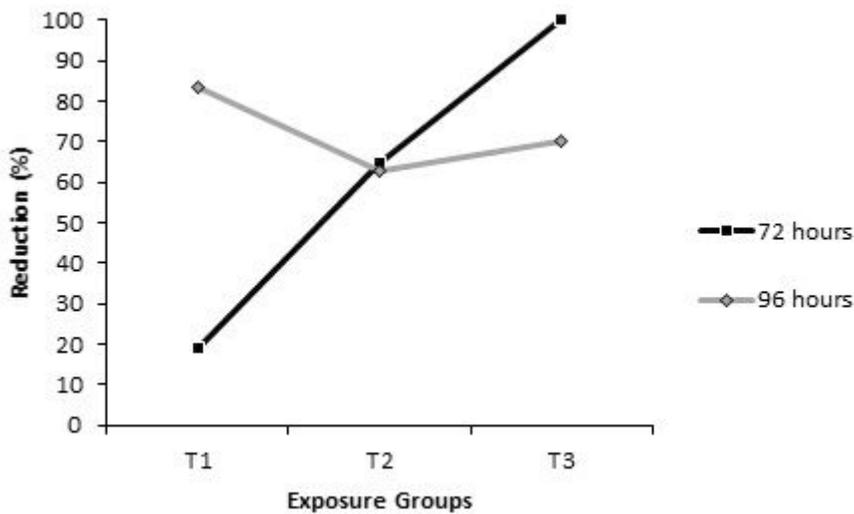
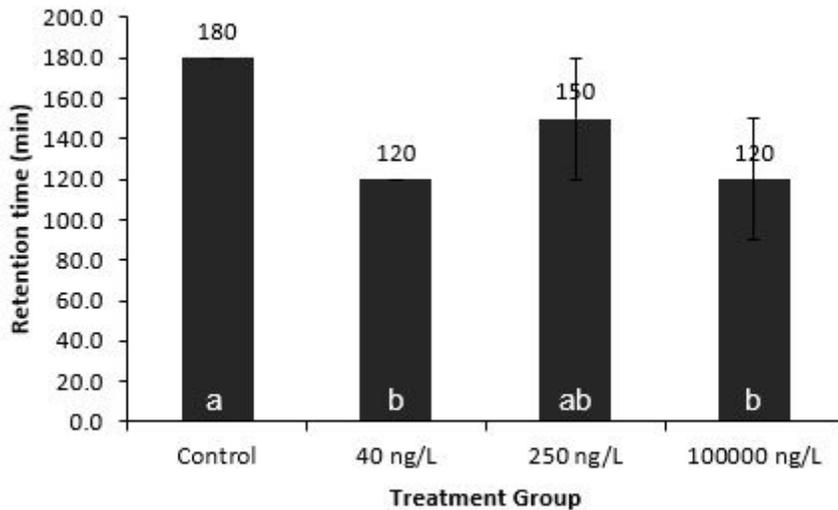


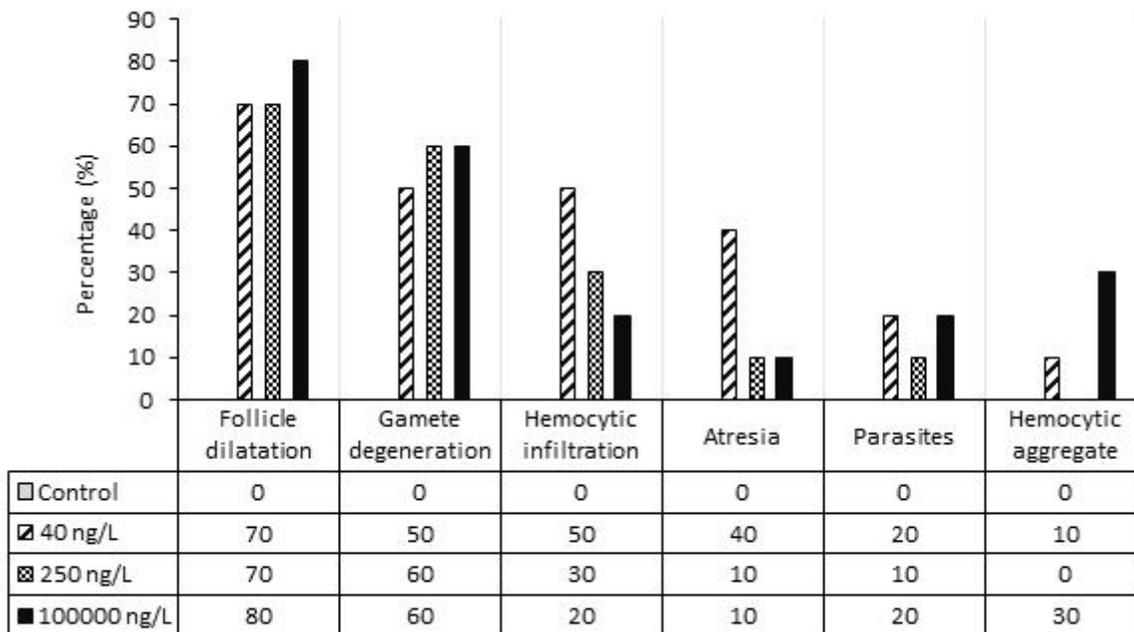
Figure 1

Percentage decrease of paracetamol level in the water after 72 and 96 hours, in three exposure tanks: T1 (40 ng/L), T2 (250 ng/L) and T3 (100 µg/L).



**Figure 2**

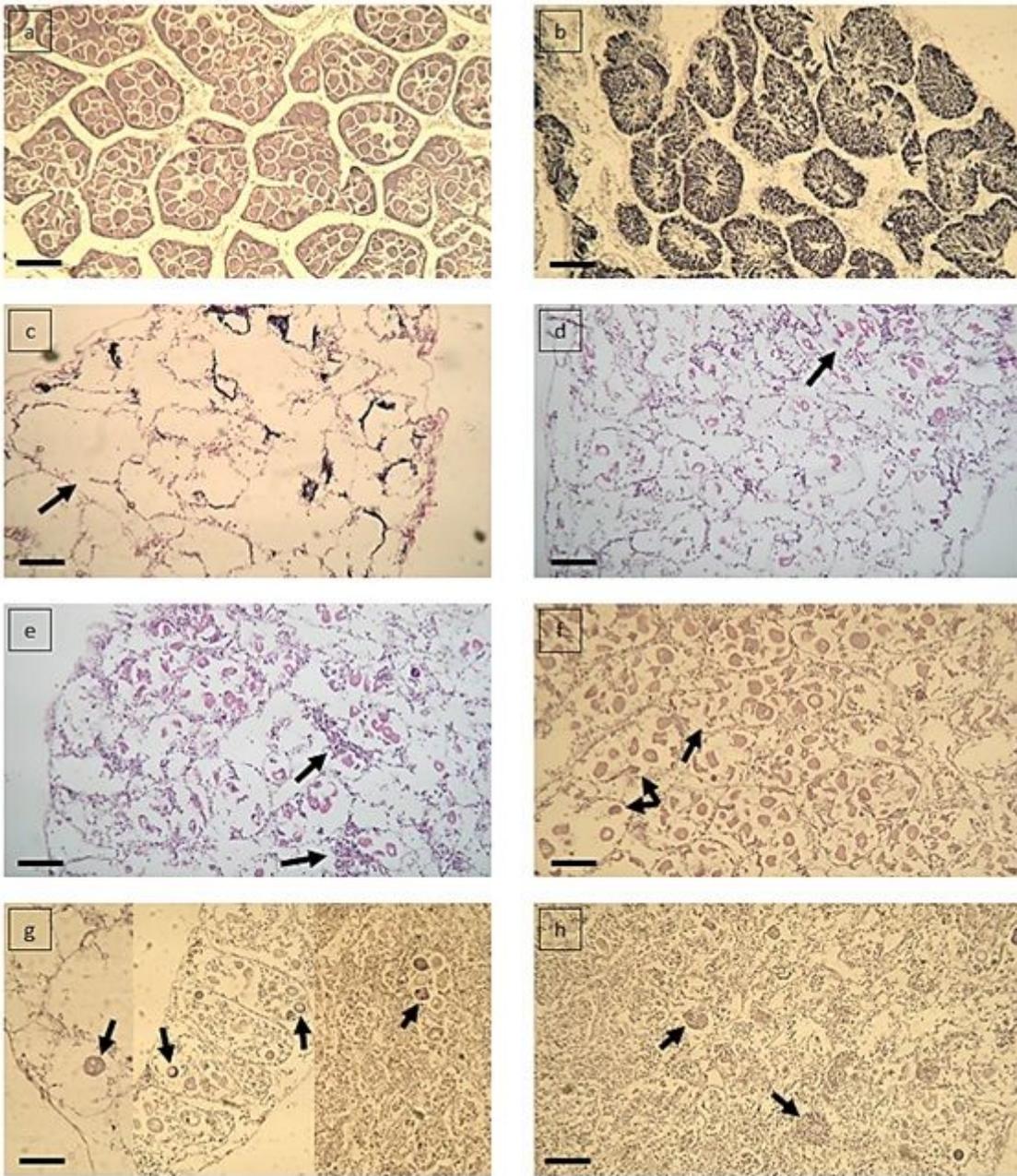
Neutral red retention time of hemocytes from mussels exposed to paracetamol for 24 days (n = 3). Different letters represent statistically significant differences between groups. Bars represent SD (one-way ANOVA, followed by Tukey's post hoc test, p < 0.05).



**Figure 3**

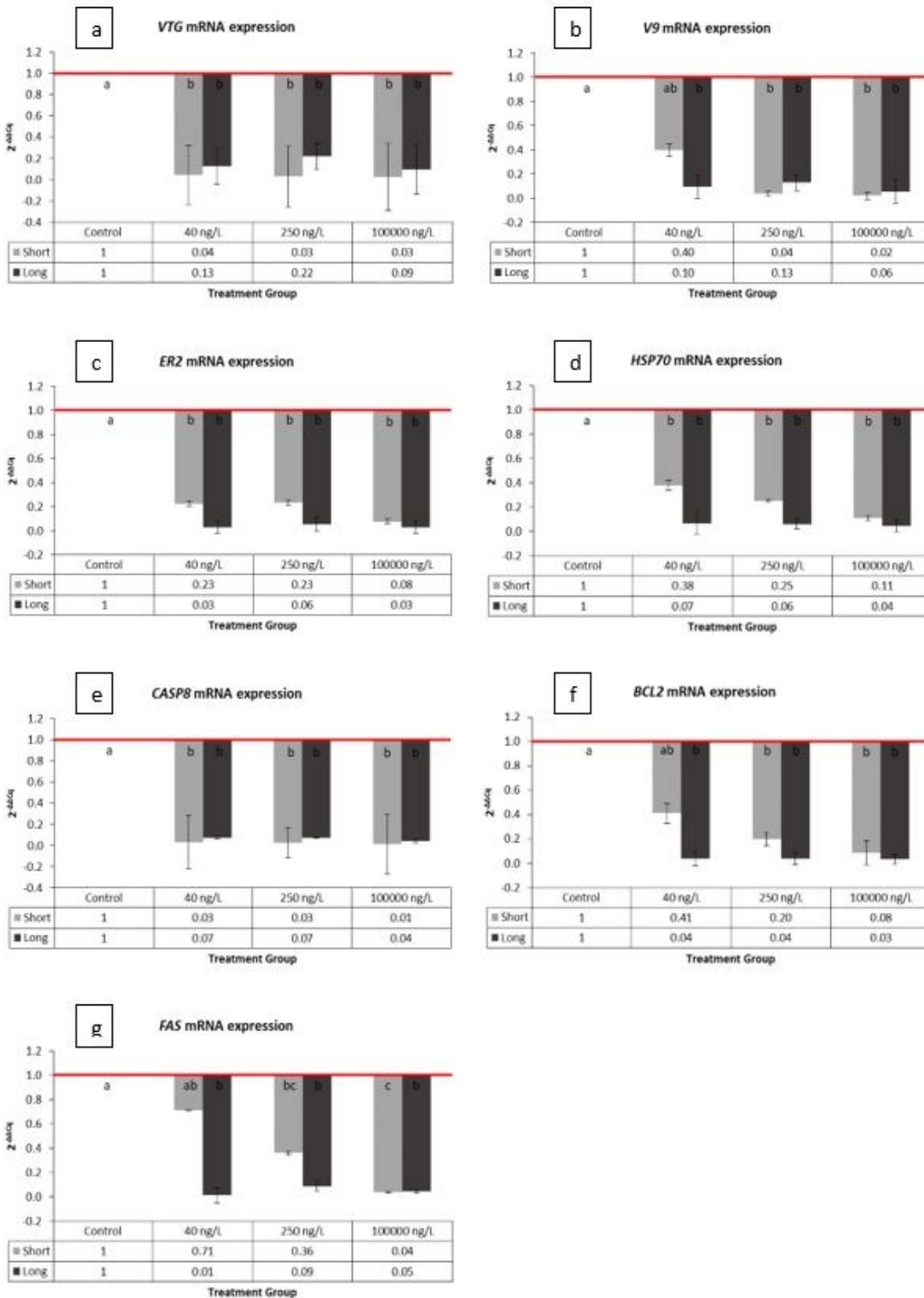
The occurrence of histopathological conditions observed in the gonad tissue of mussels exposed to paracetamol for 24 days (n=10). The treatments were as follows: Control, 40 ng/L, 250 ng/L and 100

µg/L.



**Figure 4**

Histopathological conditions in mussel gonads. Sections of 7 µm, stained with hematoxylin and eosin. (a) Normal female, (b) Normal male, (c) Follicle dilatation, (d) Gamete degeneration, (e) Hemocytic infiltration, (f) Atresia, (g) Parasites, (h) Hemocytic aggregate. Arrows point to each pathological condition. Scale bar = 100 µm.



**Figure 5**

Summary of mRNA expression of VTG, V9, ER2, HSP70, CASP8, BCL2, and FAS (a – g respectively) in fold changes compared to control group in mussel gonads (n=10). Data plotted alongside data from short exposure experiment published by Koagouw & Ciocan (2019). Different letters represent statistically significant differences between groups, bars represent SD (one way ANOVA, followed by Tukey's post hoc test,  $p < 0.05$ ).

## Supplementary Files

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- [TableS1.docx](#)