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Effects of *Alexandrium minutum* exposure upon physiological and hematological variables of diploid and triploid oysters, *Crassostrea gigas*.

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Abstract

The effects of an artificial bloom of the toxin-producing dinoflagellate, *Alexandrium minutum*, upon physiological parameters of the Pacific oyster, *Crassostrea gigas*, were assessed. Diploid and triploid oysters were exposed to cultured *A. minutum* and compared to control diploid and triploid oysters fed *T. Isochrysis*. Experiments were repeated twice, in
April and mid-May 2007, to investigate effects of maturation stage on oyster responses to *A. minutum* exposure. Oyster maturation stage, Paralytic Shellfish Toxin (PST) accumulation, as well as several digestive-gland and hematological variables, were assessed at the ends of the exposures.

In both experiments, triploid oysters accumulated more PSTs (approximately twice) than diploid oysters. Significant differences, in terms of phenoloxidase activity (PO) and reactive oxygen species (ROS) production of hemocytes, were observed between *A. minutum*-exposed and non-exposed oysters. PO in hemocytes was lower in oysters exposed to *A. minutum* than in control oysters in an early maturation stage (diploids and triploids in April experiment and triploids in May experiment), but this contrast was reversed in ripe oysters (diploids in May experiment). In the April experiment, granulocytes of oysters exposed to *A. minutum* produced more ROS than those of control oysters; however, in the May experiment, ROS production of granulocytes was lower in *A. minutum*-exposed oysters. Moreover, significant decreases in free fatty acid, monoacylglycerol, and diacylglycerol contents in digestive glands of oysters exposed to *A. minutum* were observed. Concurrently, the ratio of reserve lipids (triacylglycerol, ether glycerides and sterol esters) to structural lipids (sterols) decreased upon *A. minutum* exposure in both experiments. Also, several physiological responses to *A. minutum* exposure appeared to be modulated by maturation stage as well as ploidy of the oysters.

**Keywords:** oysters, ploidy, physiology, harmful algal bloom, *Alexandrium minutum*, PST accumulation.
1 Introduction

Among harmful algae, *Alexandrium* species are known to produce Paralytic Shellfish Toxins (PSTs), the most widespread shellfish-contaminating biotoxins, with outbreaks occurring worldwide (Huss, 2003). In France, *Alexandrium minutum* Halim (1960) has been known to bloom in coastal waters since the 1980’s (Lassus et al., 1992), especially in North Brittany (English Channel) during summer (Morin et al., 2000).

PSTs are comprised of approximately 20 naturally-occurring biotoxin derivatives that vary widely in specific toxicity (measured by standard mouse bioassay). The basic molecular structure is that of saxitoxin (STX). PSTs are neurotoxins, the mode of action of which involves a reversible and highly specific block of sodium channel transport, disabling the action potential of excitable membranes (nerves and muscle fibers) (Narahashi, 1988).

The current EU regulatory limit for human consumption of shellfish is set at 80 µg STX eq. 100g⁻¹ shellfish meat (SM). Considering, however, possible consumption of a large portion (400g) of shellfish, the European Food Safety (EFSA) recently established that the maximum concentration in shellfish meat should be less than 7.5 µg STX eq. 100g⁻¹ SM to avoid exceeding the acute reference dose (ARfD) of 0.5 µg STX eq. kg⁻¹ body weight (EFSA Journal, 2009). It is also estimated that 25% of EU samples compliant with the EU limit exceeded the concentration set by ARfD (EFSA Journal, 2009). The mouse bioassay (AOAC, 1990) protocol is the officially-prescribed method for the evaluation of STX-group toxin contamination. Although MBA sensitivity (37 µg STX eq. 100g⁻¹) allows quantification of STX-group toxins at the current EU regulatory limit, it is not within the range of ARfD concentrations (EFSA Journal, 2009). Only the HPLC-fluorescence detection method has sensitivity sufficient to quantify STX-group toxins at 1-8 µg STX eq. 100g⁻¹ SM.
Several commercially-harvested bivalve species, such as oysters, are known to accumulate PSTs by feeding on **PST-producing phytoplankton** (see review by Bricelj and Shumway, 1998). Bivalves show significant (up to 100-fold) inter-specific differences in accumulation of PSTs, which was inversely correlated with toxin sensitivity (Bricelj and Shumway, 1998). This variability in sensitivity to PSTs appeared to be related to nerve sensitivity in a dose-dependent manner. Indeed, Bricelj and Shumway (1998) reported that $10^{-7}$ g.ml$^{-1}$ STX was sufficient to block the action potential of nerves in eastern oysters *C. virginica* when $10^{-3}$ g.ml$^{-1}$ was insufficient to block action potential in blue mussels (*Mytilus edulis*). Some bivalve species possessing nerves insensitive to PST (*M. edulis*) readily feed on toxic cells and thereby accumulate high toxin levels. In contrast, species such as *Crassostrea virginica* are highly sensitive to PSTs, accumulating fewer toxins and exhibiting physiological and behavioral mechanisms to avoid or reduce exposure to toxic cells (Bricelj and Shumway, 1998). The Pacific oyster *C. gigas* and the soft-shell clam *Mya arenaria* (two PST-sensitive species) were reported to reduce filtration activity when feeding on PST-containing microalgae (Lassus et al., 2004; Bricelj and Shumway, 1998). Differences in toxin accumulation (up to five times) were also observed between different populations of the same species, *M. arenaria*, and this difference was surmised to be related to nerve sensitivity differences (Bricelj et al., 2005). Indeed, a natural mutation of a single amino acid residue decreasing affinity (1,000-fold) of the saxitoxin-binding site in the sodium channel pore, was found to be responsible for the difference in nerve sensitivity between two populations of *M. arenaria* exposed to PST-producing *Alexandrium fundyense* (Bricelj et al., 2005).

Toxin composition and content in toxigenic microalgae vary greatly according to species or strain and depend also environmental or culture conditions (Hégaret et al., 2009). An *A. minutum* strain isolated in France was found to produce 1.5 pg STX eq. cell$^{-1}$ (Lassus et al., 2004); whereas, the same species isolated in New Zealand produced 11 pg STX eq. cell$^{-1}$. 

(Chang et al., 1997). Chou and co-workers (2004) reported that toxin content of different clones of *A. minutum* isolated in Taiwan varied from 11 pg STX eq. cell\(^{-1}\) to 103 pg STX eq. cell\(^{-1}\). This variability in algal toxin content has consequences to feeding responses to and toxin accumulation of bivalves exposed to *Alexandrium* species. Bardouil and co-workers (1993) observed clearance rate in *C. gigas* decreased more drastically when oysters were exposed to *A. tamarense* (7.2 pg STX eq. cell\(^{-1}\)) than to *A. minutum* (0.5 pg STX eq. cell\(^{-1}\)). When exposed to less-toxic *Alexandrium* species or strains, sensitive bivalves such as *C. gigas* can feed on and accumulate PSTs (Lassus, unpubl. obs.).

*Alexandrium* species are also known to produce other toxic compounds, such as ichthyotoxins (Emura et al., 2004) and allelochemicals (Arzul et al., 1999; Tillmann et al., 2008). Ford et al. (2008) tested effects of two *A. tamarense* strains, PST and non-PST producing, upon Manila clam *Ruditapes philippinarum* and *Mya arenaria* hemocytes. This study showed that the non-PST strain had more-negative impacts on hemocytes (decreased adhesion and phagocytosis) compared to the PST-producing strain of *A. tamarense* (Ford et al., 2008). Based upon biological effects of *Alexandrium* exposure clearly unrelated to PSTs, one can speculate that effects of PSTs and other toxic compounds/molecules may also result in damage to organs and physiological processes other than muscles. Considering this, *Alexandrium* effects upon nerves and muscles of exposed bivalves can be linked to PSTs, but there appear to be responses to other compounds not clearly identified.

Aside from toxin accumulation and associated human health issues, there is some concern about the impact of *A. minutum* exposure on the physiology and health of *C. gigas*. Furthermore, the physiological status of animals may also feed-back to rates of toxin accumulation and effects during HAB exposure.

Li et al. (2002) studied the effect of *Alexandrium tamarense* (PST-containing strain) on bioenergetics and growth rate of the clam *Ruditapes philippinarum* and the mussel *Perna viridis*. 
High concentrations of toxic *A. tamarense* (resulting in high PST burdens in the tissues) decreased clearance rate of the clam but not of the mussel. Absorption efficiency, however, decreased for both species with diets containing PST, which resulted in a reduction in energy budget. HABs occurring during specific stages of reproduction could be another major factor affecting oysters, in terms of energy budget. In *C. gigas*, energy balance (evaluated through scope for growth methods) has been demonstrated to decrease as gametogenesis progresses, resulting in a negative scope for growth (Lambert et al., 2008). The digestive gland plays an obvious, major role in nutrient digestion and assimilation, as digestive-enzyme activities in bivalves can be affected by nutritional condition. Indeed, changes in enzymatic activities are mechanisms used by bivalves to optimize energy gain when experiencing variation in dietary input (Fernández-Reiriz et al., 2001; Labarta et al., 2002). The mussel *M. chilensis* can, indeed, use toxic microalgae (*Alexandrium catenella*) as a food source by adjusting carbohydrase activities (amylase, laminarinase and cellulase) and absorption mechanisms (Fernández-Reiriz et al., 2008). It is, thus, pertinent to assess digestive-enzyme activities in bivalves exposed to toxic microalgae. Moreover, the digestive gland is also involved in energy storage, preferentially as lipids (Soudant et al., 1999). As HABs are likely to impact digestive-gland structure and functions, it appears prudent to assess how HABs could modulate quantities of individual lipid classes involved in energy storage in this organ. Moreover, the digestive gland is the organ accumulating the most toxins compared to other tissues in bivalves (Bricelj and Shumway, 1998).

In bivalves, one line of defense to noxious, harmful or pathogenic agents resides in circulating cells called hemocytes that are similar to white blood cells in vertebrates (Cheng, 1996). Concerning hemocyte variables, Hégaret et al. (2007a, 2007b, 2008) reported that harmful-algal exposure can modulate cellular immune components and functions of bivalves.
Moreover, Galimany et al. (2008) observed an inflammatory response in the stomach of *Mytilus edulis* exposed to *A. fundyense*. These findings indicate that the bivalve immune system can be activated by certain harmful algae, or conversely can be suppressed. The reproductive period is also associated with changes in hemocyte variables; some are depressed in *C. gigas* during gametogenesis, specifically hemocyte concentration, phagocytosis, and adhesion (Lambert et al., 2008). During gametogenesis, hemocytes in triploid oysters were found to have higher phagocytic, esterase and peroxidase activities than those of diploids (Gaignaire et al., 2006). These differences were attributed to the reduced gametogenic development of the triploids.

Indeed, triploid oysters are increasingly used for aquaculture because they can be marketed during summer when diploid oysters are fully-ripe and not appreciated by consumers. It is unknown, however, how triploidy may affect toxin accumulation.

The purpose of the present study was to determine the effects of an artificial bloom of the toxin-producing dinoflagellate, *Alexandrium minutum* (strain AM89BM), upon digestive parameters, and hemocyte and plasma variables of the Pacific oyster, *Crassostrea gigas*. Diploid and triploid oysters were compared to assess any differences in toxin accumulation and physiological responses to harmful-algal exposure. Experiments were conducted at two different periods (one month apart), using the same oyster stock, to obtain a gradient of gonad maturation. After 4 days of exposure to *A. minutum* or *Isochrysis* sp. (clone Tahitian T.Iso) as a non-toxic control, toxin accumulation, reserve lipid classes and amylase activities, humoral phenoloxidase variables, hemolyse/agglutination capacity, and hemocyte concentration, morphology, viability, phagocytosis activity, reactive oxygen species production and phenoloxydase activity, were measured.
2 Materials and Methods

2.1 Biological material

2.1.1 Oysters

Diploid and triploid Pacific oysters, *Crassostrea gigas*, used in the two experiments were obtained from an oyster producer at île de Kerner (Morbihan, FRANCE) and belong to the same commercial stocks (20-21 months old). For each experiment, we used 60 diploid oysters and 60 triploid oysters. In April, flesh dry weight was 1.22 ± 0.12 g and 0.90 ± 0.09 g in diploids and triploids, respectively. In May, flesh dry weight was 1.33 ± 0.14 g and 0.98 ± 0.09 g in diploids and triploids, respectively. At both collection times, three pools of four oysters (for each ploidy) were confirmed to be free of PST contamination (no detectable levels of PSTs by IP-HPLC analysis).

2.1.2 Algal culture

*Alexandrium minutum* (strain AM89BM) was grown in 10-liter batch culture using autoclaved seawater filtered to 1µm and supplemented with L1 medium (Guillard and Hargraves, 1993). Cultures were maintained at 16 ± 1°C and 100 µmol photon.m\(^{-2}.s\(^{-1}\), with a dark:light cycle of 12:12h. *A. minutum* was harvested after 12 days, still in exponential growth phase under our conditions. At this age, this strain produced 1.3 ± 0.1 pg STX eq. per cell (measured by the method of Oshima (1995)).

*Isochrysis* sp., clone Tahitian (T.Iso), cultures were obtained from the Argenton hatchery (IFREMER - FRANCE). Cultures were produced in 300-liter cylinders containing 1-µm filtered seawater enriched with Conway medium at 24 ± 1°C, air-CO\(_2\) (3%) mix aerated, and with continuous light. T.Iso was harvested in the exponential growth phase (6-8 days) for the feeding experiments.
2.2 Experimental design of *A. minutum* exposures

For each experiment (April and May), 120 oysters (60 diploids and 60 triploids) were placed haphazardly in twelve 15-L tanks (10 oysters per tank). Oysters were acclimated for 10 days with continuous flow of 14 ml.min\(^{-1}\) of seawater (filtered to 0.5 µm) with *T. Iso*, 5.10\(^5\) cells.ml\(^{-1}\) at 16 ± 1°C. After acclimation, diploid and triploid oysters were fed continuously for 4 days 14 ml.min\(^{-1}\) with 5.10\(^5\) cells.ml\(^{-1}\) of *T. Iso* (6 control tanks with diploids and triploids) and with 5.10\(^3\) cells.ml\(^{-1}\) of *A. minutum* (equivalent to 6.5 ng STX eq. - 6 treatment tanks with diploids and triploids).

2.3 Oyster sampling

At the end of the algal-exposure, all oysters were sampled and distributed as follow. For each tank (10 oysters), gonads of two oysters were used for histological analysis of maturation stage. Pooled digestive glands of four oysters were used to measure toxins, reserve lipid contents, and amylase activity. Four oysters were used for individual plasma and hemocyte variable measurements and condition index.

2.4 Qualitative analysis of maturation stages

Gonads were dissected and transferred into Bouin fixative (for 48 h). Fixed gonads were dehydrated in ascending ethanol solutions, cleared with xylene and embedded in paraffin wax. Five-micrometer thick sections were cut, mounted on glass slides, and stained with Harry’s hematoxylin-Eosin Y (Martoja et al., 1967). Slides were examined under a light microscope to determine gametogenic stage according to the reproductive scale reported by Mann (1979). In this scale, four stages are defined: stage 0 (inactive), stage 1 (early gametogenesis), stage 2 (late gametogenesis) and stage 3 (ripe).
2.5 Condition index

To assess oyster-flesh dry weight, soft tissues were removed from shells and placed in a pre-weighed aluminum cup. Shell and flesh were dried for 48h at 70°C and then weighed.

Condition index of individual oysters was then calculated as described previously (Lucas and Beninger, 1985), following the formula: (g dry flesh weight / g dry shell weight) x 100.

2.6 Digestive gland variables

Just after dissection, digestive glands were immediately frozen in liquid nitrogen, weighed, pooled (1 pool of 4 digestive glands per tank), and stored at -80°C until analysis. Later on, pools were ground with a “Dangoumau” homogenizer into liquid nitrogen and divided for three different analyses.

2.6.1 Toxin content

One gram of ground digestive gland was extracted in 2 ml of 0.1 N HCl (2 v/w) at 4°C. After centrifugation (3,000 × g, 15 min, 4°C), the pH of extracts was adjusted. If above 3.0, pH was adjusted to 3.0 with 12 N HCl. After half-dilution, supernatants were ultra-filtered (20 kDa, Sartorius Centrisart) and stored at 4°C until analysis. PSTs were analyzed by ion-pairing, high-performance liquid chromatography (IPHPLC) according to the method of Oshima (1995). The molar concentration (µmol.L⁻¹) was converted into µg STX eq. 100 g⁻¹ of digestive gland by using the conversion factors of Oshima (1995). Results were expressed in µg STX eq. 100 g⁻¹ of digestive gland wet weight.

2.6.2 Reserve lipid content

Ground digestive gland (250 mg) were extracted in 6 ml of Folch solution (chloroform:methanol 2:1). Lipid classes were analyzed by high-performance, thin-layer chromatography (HPTLC) on HPTLC glass plates (1,010 mm) pre-coated with silica gel 60
from Merck (Darmstadt, Germany). A preliminary run was carried out to remove possible
impurities using hexane:diethyl ether (1:1), and the plate was activated for 30 min at 110°C.
Lipid samples (4 µl) were spotted on the plates by the CAMAG automatic sampler. The
neutral lipids were separated using a double development with hexane:diethyl ether:acetic
acid (20:5:0.5) as first solvent system followed with hexane:diethyl ether (93:3) as a second
solvent system. Lipid classes appeared as black bands after dipping plates in a cupric-sulfate,
phosphoric-acid solution and heating for 20 min at 160°C (charring). Seven neutral lipid
classes (categorized as storage lipids: free fatty acids, sterol esters, glycerid ethers,
monoacylglycerol, diacylglycerol and triacylglycerol; considered as structural lipids: sterols)
were identified based upon standard (Sigma–Aldrich, France) and coloring techniques. The
charred plates were read by scanning at 370 nm, and black bands were quantified by Wincats
software. Results were expressed as mg of each identified neutral lipid class per g of
digestive gland wet weight.

2.6.3 Amylase activity

Ground digestive gland (200 mg) was homogenized in 1 ml of distilled water and 200 µl of
this solution were added to 10 µl of 0.5 M CaCl\textsubscript{2} solution before analysis to assess amylase
activity. Amylase activity was then assayed by determination of starch hydrolysis according
to the iodine reaction (Samain et al., 1977) modified by Le Moine et al. (1997). One unit of
alpha-amylase was defined as the amount of enzyme that degrades 1 mg.min\textsuperscript{-1} starch at 45°C.
To assess specific activities, total proteins were determined using the BCA Protein Assay
(Biorad). For protein extraction, 200 µl of the above solution were added to 200 µl of 2 N
NaOH solution. Protein analysis was carried out on 10 µl of 1/10 diluted samples according
to the manufacturer’s description. Briefly, 200 µl of dye reagent was added to 10 µl of
sample, incubated at 37°C for 1 hour and the absorbance was measured at 595 nm. Sample
ODs were compared to a standard curve of Bovine Serum Albumin (BSA), and results were
expressed as mg of protein.ml⁻¹. Amylase activity was expressed as UI of amylase activity per mg of total protein (specific activity).

2.7 Hemolymph variables

2.7.1 Hemolymph sampling

Hemolymph was withdrawn from individual oysters using a 1 ml plastic syringe fitted with a 25-gauge needle inserted through a notch made adjacent to the adductor muscle just prior to bleeding. All hemolymph samples were examined microscopically for contamination (e.g., gametes, tissue debris) and then stored in micro-tubes held on ice. As recommended by flow cytometer (FCM) manufacturer, all samples were filtered through 80 µm mesh prior to analysis to eliminate any large debris (> 80 µm) which could potentially clog the flow cytometer. Three hundred microliters (3 measures x 100 µl) of each hemolymph sample were used to measure hemocyte variables by flow cytometry. The remaining hemolymph was separated into cellular (hemocytes) and supernatant (plasma) fractions by centrifugation (800×g, 5 min, 4°C) prior to freezing (−20°C). These samples then were used to measure biochemical hemocyte and plasma variables (protein content, phenol-oxydase activity and hemolysis/agglutination titers). Methods for measuring cellular (hemocyte) and humoral (plasma) variables are described hereafter.

2.7.2 Measurements of hemocyte variables by flow cytometry

Characterization of hemocyte sub-populations, number and functions were performed using a FACScalibur (BD Biosciences, San Jose, CA USA) flow cytometer (FCM) equipped with a 488 nm argon laser. Two kinds of hemocyte variables were evaluated by FCM: descriptive variables (hemocyte viability and total and hemocyte sub-population counts), and functional variables (phagocytosis and reactive oxygen species (ROS) production). Analyses were done as described below.
Descriptive variables: Hemocyte viability, total and hemocyte sub-population counts

These variables were measured individually on hemolymph samples (4 individuals per tank).

An aliquot of 100 µl of hemolymph from an individual oyster was transferred into a tube containing a mixture of Anti-Aggregant Solution for Hemocytes, AASH (Auffret and Oubella, 1995) and filtered sterile seawater (FSSW), 200 µl and 100 µl respectively.

Hemocyte DNA was stained with two fluorescent DNA/RNA specific dyes, SYBR Green I (Molecular probes, Eugene, Oregon, USA, 1/1000 of the DMSO commercial solution), and propidium iodide (PI, Sigma, St Quentin Fallavier, France, final concentration of 10 µg.ml⁻¹) in the dark at 18°C for 120 minutes before flow-cytometric analysis. PI permeates only hemocytes that lose membrane integrity and are considered to be dead cells; whereas, SYBR Green I permeates both dead and live cells. SYBR Green and PI fluorescences were measured at 500-530 nm (green) and at 550-600 nm (red), respectively, by flow-cytometry.

Thus, by counting the cells stained by PI and cells stained by SYBR Green, it was possible to estimate the percentage of viable cells in each sample. All SYBR Green-stained cells were visualized on a Forward Scatter (FSC, size) and Side Scatter (SSC, cell complexity) cytogram. Three sub-populations were distinguished according to size and cell complexity (granularity). Granulocytes are characterized by high FSC and high SSC, hyalinocytes by high FSC and low SSC, while agranulocytes have low FSC and SSC. Total hemocyte, granulocyte, hyalinocyte, and agranulocyte concentrations estimated from the flow-rate measurement of the flow-cytometer (Marie et al., 1999) as all samples were run for 30 sec.

Results were expressed as number of cells per milliliter of hemolymph.

Functional variables

These variables were measured individually on hemolymph samples, for each condition.
Phagocytosis

An aliquot of 100 µl hemolymph, diluted with 100 µl of FSSW, was mixed with 30 µl of YG, 2.0-µm fluoresbrite micro-spheres, diluted to 2% in FSSW (Polysciences, Eppelheim, Germany). After 120 minutes of incubation at 18°C, hemocytes were analyzed at 500-530 nm by flow cytometry to detect hemocytes containing fluorescent beads. The percentage of phagocytic cells was defined as the percentage of hemocytes that had engulfed three or more beads (Delaporte et al., 2003).

Reactive oxygen species production

Reactive oxygen species (ROS) production by untreated hemocytes was measured using 2’7’-dichlorofluorescein diacetate, DCFH-DA (Lambert et al., 2003). A 100-µl aliquot of pooled hemolymph was diluted with 300 µl of FSSW. Four µl of the DCFH-DA solution (final concentration of 0.01 mM) was added to each tube maintained on ice. Tubes were then incubated at 18°C for 120 minutes. After the incubation period, DCF fluorescence, quantitatively related to the ROS production of untreated hemocytes, was measured at 500-530 nm by flow-cytometry. Results are expressed as the geometric-mean fluorescence (in arbitrary units, AU) detected in each hemocyte sub-population.

2.7.3 Biochemical hemocyte and plasma variables

Hemocyte and plasma phenoloxidase activities

Plasma samples were thawed on ice, and 100 µl of each was transferred in ninety-six-well plates. For hemocytes, cells were suspended in 100 µl of FSSW and frozen and thawed on ice three times successively. Phenoloxidase activity was measured as described by Reid (2003). Briefly, 50 µl of Tris–HCl buffer (0.2M, pH = 8) and 100 µl of l-DOPA (20 mM, L-3,4-
dihydrophenyl-alanine, Sigma D9628) were added to each well. The micro-plate was rapidly mixed for 10 s. The reaction was then measured at ambient temperature, with color change recorded every 5 min, at 492 nm, over a period of 1 h. The micro-plate was mixed prior to each measurement. Two controls, without sample but containing l-DOPA and Tris–SDS buffer, were measured in parallel, and these values were subtracted from test values to correct for possible auto-oxidation of the l-DOPA.

To access phenoloxidase specific activity, protein analysis was carried out as described for digestive glands (see paragraph 2.6.3), except that proteins were not extracted with NaOH and samples were not diluted. Results were expressed as phenoloxidase-specific activity.

Agglutination and hemolysis titers in plasma

Agglutination titer (indicative of the presence of lectins) and hemolysis titer (indicative of red blood cell lysis factors) were measured on a sub-sample of plasma (supernatant) fraction. Quantification of agglutination titer was performed according to the protocol from Barracco et al. (1999), using horse red blood cells. Briefly, 50-µl plasma samples were added to U-shaped wells of 96-well microtiter plates, and a two-fold, serial dilution (pure solution to 1/2 dilution) was prepared using Tris-buffered saline (containing 0.15 M NaCl). The same volume of a 2% suspension of horse red blood cells in TBS was added to each well and incubated for 3 h at room temperature. In controls, oyster plasma was replaced with TBS. Agglutination titer and hemolysis titer were expressed as the log (base 2) of the reciprocal of the highest dilution showing a positive pattern of agglutination or hemolysis of red cells, respectively.
2.8 Statistical analysis

Differences between experiments (April and May) were assessed using Student's T-test. Results of each experiment were analyzed statistically using Multifactor-ANOVA (MANOVA) for each physiological parameter and hemocyte variable as the dependent variable, and feeding treatment and ploidy as independent variables. Whenever a clear trend appeared on the graphs, a Student's T-test was also used within ploidy groups to assess differences linked to dietary treatment for a dependent variable. We used Statgraphics Plus statistical software (Manugistics, Inc, Rockville, MD, USA). Results were considered significant when the P-value was < 0.05.

3 Results

3.1 Gonad maturation stages

Oysters fed T. Iso and A. minutum were combined to assess oyster gonad maturation (Fig. 1). Maturation of both diploid and triploid oysters was more advanced during the May experiment than during April. In both experiments, triploids were less mature than diploids. Oyster groups in both experiments can be classified according to gonad maturation stage (from less mature to more mature) first were triploids in April, triploids in May, then diploids in April, and at last diploids in May. Triploids in the April experiment were mostly at the undifferentiated stage, and one third were in early and late gametogenesis; triploids in the May experiment were in early and late gametogenesis, diploids in the April experiment were dominated by late gametogenesis with 20% mature oysters; and finally, diploids in the May experiment were mainly (60%) observed to be sexually mature while 40% were in late gametogenesis.
3.2 Wet weight of digestive gland and condition index

There was no significant difference, in term of condition index (CI), attributable to diet or ploidy in both experiments (Table 1). Whole-oyster dry weight (DW) was significantly higher in diploid oysters than in triploid oysters for both experiments (in April p=0.0045 and in May p=0.0021, MANOVA). DW of both diploid and triploid oysters did not change significantly between the two experiments. Wet weights (WW) of digestive glands were similar in the April experiment regardless of diet or ploidy. In the May experiment, mean WW of digestive gland was significantly higher in diploids than in triploids. Exposure to A. minutum in this experiment resulted in a significant decrease in digestive gland WW compared to T.Iso feeding. A significant interaction between ploidy and diet was also noted; lower digestive gland WW was found in triploid oysters exposed to A. minutum.

3.3 Toxin content

PST content in digestive gland was significantly higher in May than in April (p=0.003, T-test). In both experiments, triploid oysters accumulated more toxin -- about twice -- than diploids (Fig. 2); April experiment p=0.032 and May experiment p=0.047, T-test).

Concomitantly, A. minutum cells were observed in digestive gland and bio-depots.

3.4 Digestive gland parameters

3.4.1 Reserve lipid content

Neutral lipid classes detected in digestive glands were free fatty acids, sterol esters, ether glycerides, sterols, monoacylglycerols, diacylglycerols and triacylglycerols (Table 2). A reserve/structure ratio was determined as the ratio between reserve lipids (sterol ester + glycerid ether + triacylglycerol content) and a structural lipid (sterol content).
In both experiments, monoacylglycerol, diacylglycerol and free fatty-acid contents (Fig. 3) were significantly lower in both diploid and triploid oysters fed A. minutum as compared to those fed T.Iso (Table 2). In May, contents of sterols, triacylglycerols, and sterol esters, as well as the reserve/structure ratio, were higher in diploids than in triploids.

In April, the reserve/structure ratio was significantly lower (Fig. 4) in triploid oysters exposed to A. minutum as compared to triploids fed T.Iso (p=0.0427, T-test). Exposure to A. minutum similarly resulted in a significant decrease of this ratio in diploids in May (p=0.0067, T-test).

3.4.2 Amylase-specific activity

Amylase-specific activity (ASA) was significantly higher in May than in April (p=0.0222, T-test) (Fig. 5). In both experiments ASA was higher in triploids than in diploids (April experiment p=0.0263; May experiment p=0.0134, MANOVA). ASA was higher in A. minutum-exposed than in control, diploid oysters in April (p=0.0467, T-test), but was similar in exposed and non-exposed triploid oysters. In May, only triploid oysters showed a significant increase of ASA upon A. minutum exposure (p= 0.0337, T-test).

3.5 Hemocyte and plasma variables

Overall, ploidy had more significant impacts on hemocyte and plasma variables than algal exposure (Table 3).

3.5.1 Hemocyte characteristics and functions analyzed by flow cytometry

Total hemocyte concentration (THC) was significantly higher in May than in April (p=0, T-test). THC increased significantly upon A. minutum exposure in diploid oysters in the April experiment (p=0.013, T-test) and triploid oysters in the May experiment (p=0.042 with α=0.05, T-test) (Fig. 6). This increase was mainly attributable to variation in granulocyte
counts, especially in April when granulocyte counts drastically increased upon *A. minutum* exposure.

Sizes (FSC) of granulocytes and hyalinocytes (Fig. 6) of oysters in the April experiment were significantly higher in triploids than in diploids, but hemocyte size was not affected by *A. minutum* exposure. In May, size and complexity (SSC) of both granulocytes and hyalinocytes were higher in triploids than in diploids. In the same experiment, *A. minutum* resulted in a significant increase in granulocyte and hyalinocyte size and in hyalinocyte complexity.

ROS production in granulocytes and hyalinocytes (Fig. 6) was significantly higher in May than in April (p=0.0002 for granulocytes and p=0.0011 for hyalinocytes, T-test). In both experiments, granulocytes of triploids produced more ROS than granulocytes of diploids (p=0.0034 in April and p=0.0012 in May, MANOVA). The same difference was observed for hyalinocytes but was only significant in April (p=0.0215, MANOVA).

*A. minutum* exposure resulted in opposite effects in the two experiments. In the April experiment, granulocytes of oysters fed *A. minutum* produced more ROS than those of control oysters (p=0.0119, MANOVA). In the May experiment, granulocytes and hyalinocytes of oysters fed *A. minutum* produced less ROS than those of control oysters (p=0.0067 and p=0.0358 respectively, MANOVA).

Neither phagocytosis nor percentage of dead cells was affected by algal exposure or ploidy.

### 3.5.2 Hemocyte and plasma phenoloxidase (PO) activities

PO in plasma was higher in April than in May (p=0.001, T-test). In April, PO in plasma decreased in *A. minutum*-exposed oysters, significantly only for diploids (p=0.008, T-test). There were no significant variations in PO in plasma in May. In hemocytes, PO (Fig. 7) was significantly higher in May than in April (p=0.0152, T-test). PO in hemocytes was higher in triploids than in diploids in both April and May experiments (respectively p=0.0108 and p=0.046, MANOVA). In April, PO in hemocytes was lower in oysters fed *A. minutum* than in
control oysters (p=0.0312, MANOVA). In May, PO in hemocytes was higher in diploids fed
A. minutum as compared to control diploids (p=0.0189, T-test) and lower in triploids
(p=0.0458, T-test).

3.5.3 Agglutination and hemolysis
There were no significant differences in agglutination or hemolysis titer according to algal
exposure or ploidy.

4 Discussion
The effects of Alexandrium minutum exposure for 4 days, specifically on toxin accumulation
and several physiological parameters, were evaluated in diploid and triploid Pacific oysters,
Crassostrea gigas. Although triploid oysters are increasingly used for aquaculture, it was
unknown how triploidy may affect toxin accumulation and physiological responses to
harmful algal blooms. Two experiments were conducted during two consecutive months to
assess the possible impact of reproductive stage on toxin accumulation and physiological
responses to A. minutum exposure. As A. minutum toxins are released in oyster digestive
glands, this organ could be expected to be impacted. The second physiological compartment
expected to be impacted by A. minutum exposure was the circulatory system, containing
hemocytes.
The most striking result in these two experiments was the difference in PST accumulation in
digestive glands between diploid and triploid oysters. In both experiments, triploid oysters
accumulated twice the toxin of diploids. One could be quick to attribute higher toxin content
in triploids to lower gametogenesis, but this hypothesis can be rejected as oysters in the May
experiment had both more-advanced gonad development and higher toxin contents than
oysters in the April experiment. Gametogenetic stage is not, however, the only trait that
distinguishes triploid from diploid oysters. The augmentation in genetic material and gene
copies in triploids also has physiological implications by changing heterozygosity. This increased heterozygosity in triploids can lead to additive and non-additive effects upon gene expression (Riddle et al., 2006; Johnson et al., 2007). Indeed, in parallel to toxin accumulation, amylase activity, ROS production, and phenoloxidase activities were found to be higher in triploid oysters than in diploids. Esterase and peroxidase activities were also found to be higher in triploid oysters than in diploid oysters (Gagnaire et al., 2007). This tends to support the hypothesis that triploids accumulate more toxin than diploids because they are metabolically more active. As triploid heterozygosity is higher, it is thought to have positive influences on feeding rate, absorption efficiency, and growth efficiency (Magoulas et al., 2000). Thus, one can speculate that the difference in toxin accumulation may reflect differences in metabolic and/or feeding activities between diploid and triploid oysters.

The increase in toxin accumulation between April and May experiments is also possibly a result of an increase in feeding and digestive activities. Considering together A. minutum-exposed and non-exposed oysters, amylase activity, ROS production, and PO activity were found to be higher in May than in April. Even though oysters in both experiments originated from the same stock and were acclimated for 7 days prior to exposure to A. minutum, an additional month in field rearing conditions appeared to impart subsequent physiological status, including reproductive processes in the conditioned oysters. Indeed, during this additional month, temperature increased from 11 to 15°C and photo-period increased from 13h06 per day on 5 April to 14h38 per day on 2 May. These temperature and photo-period changes allowed diploid oysters to develop from early gametogenesis to a large proportion of mature gonads within one month. This is in good agreement with the study of Fabioux et al. (2005). The percentage of gonad occupation in C. gigas, in field conditions, increased from 15% to 50% between April and May (Fabioux et al., 2005). Gonad development and energy
allocation in the scallop *Pecten maximus* were shown to be modulated by temperature and photoperiod (Saout et al., 1999). Similarly Fabioux et al. (2005) experimentally demonstrated that the gametogenic cycle of *C. gigas* can be controlled by coupled modifications of temperature and photoperiod. In marine invertebrates, it is postulated that the nervous system under environmental influences has an effect on the endocrine regulation of reproduction (Olive, 1995, Lafont, 2000). In bivalves, fluctuations of neurotransmitters (monoamines) appeared to be related to seasonality and reproductive cycle (Lopez-Sanchez et al., 2009).

Although specific functions of monoamines are not clearly identified yet, they are thought to be involved in meiosis re-initiation in bivalve oocytes (Guerrier et al., 1993), control of ciliary movement (Carroll and Catapane, 2007), or spawning induction (Velez et al., 1990, Fong et al. 1996, Velasco et al., 2007).

In the present study, the general increase in measured physiological parameters is certainly determined by environmental conditions (temperature, salinity and food in the field) and we speculate that metabolism is further maintained during the experiment possibly through endocrine control. From all the above, we infer that increasing toxin accumulation paralleled increasing metabolic and physiological activities determined by ploidy and preceding field conditions.

Major changes in free fatty acids (FFA), monoacylglycerols (MAG) and diacylglycerols (DAG) contents in the oyster digestive gland were clearly attributable to algal exposure. In both experiments, concentrations of FFA, MAG and DAG were reduced upon *A. minutum* exposure in both diploids and triploids. These compounds are generally absent in gonad and muscle, and only transiently observed in digestive glands of oysters (Soudant et al., 1999).

The biological significance of FFA, DAG and MAG contents is still unclear as little has been
published on the subject. These lipid classes are thought to be intermediate products in the
synthesis or catabolism of both structural and reserve lipids.

Upon exposure to *A. minutum*, the ratio of reserve lipids (TAG, EGLY and SE) to structural
lipids (sterols) decreased in both experiments. This depletion of reserve lipids upon *A.
minutum* exposure, however, appeared to partially depend upon maturation stage. The most-
drastic decrease upon *A. minutum* exposure occurred when oyster gametogenesis was almost
absent (April experiment triploids) or predominantly terminated (May experiment diploids).
In these physiological conditions, more reserve lipids resided in the digestive gland, where
these energy reserves are potentially available to respond to stressful conditions. In contrast,
when oysters were in late gametogenesis (April experiment diploids and May experiment
triploids), reserve lipids were likely intensively transferred to the gonad and may thus be only
slightly affected by *A. minutum* exposure.

As FFA, MAG and DAG concomitantly decreased with the reserve/structure ratio, we
speculate that *A. minutum* negatively affects digestion of dietary lipids and/or synthesis of
storage lipids. Also, it has to be noted that contents of TAG and EGLY in *A. minutum-
exposed oysters decreased as toxin accumulation increased according to oyster groups.

Nevertheless, at this stage, it is difficult to establish clear relationships between reserve lipid
changes and toxin accumulation. To further progress on these aspects, it would be interesting
to combine this biochemical information with histological analyses.

Amylase-specific activity (ASA) was also modulated upon *A. minutum* exposure. Overall,
ASA was higher in oysters exposed to *A. minutum* than in *T. Iso*-fed oysters, but also higher
in triploids than in diploids. As mentioned earlier, ASA, along with other metabolic activities,
could partially explain the differences in toxin accumulation between diploids and triploids
and between April and May experiments. Digestive-enzyme activities are important in
maximizing absorption and food conversion efficiencies (Huvet et al., 2003). In *C. gigas*, a
positive correlation has been established between high specific amylase activity and high
food assimilation (Prudence et al., 2006). In the present study, digestive-enzyme activities
were likely controlled by physiological status, which varied with ploidy and times of
experiments. Thus, we speculate that higher amylase activity would result in higher A.
*minutum* digestion and toxin release and assimilation.

These results highlight the complexity of relationships between oyster physiology and toxin
accumulation. Toxin accumulation certainly depends upon the physiological status and
metabolic activities of the oysters; concomitantly, toxin accumulation can interfere with the
same physiological processes.

Regarding hemocyte variables, changes upon *A. minutum* exposure were mainly observed in
hemocyte cell density, phenoloxidase, and ROS production. Except in mature oysters
(diploids in experiment 2), *A. minutum* exposure resulted in increases in numbers of
circulating hemocytes. Increase in total circulating hemocyte counts is generally considered
to be an immune response to pathogens (Chu et al., 1993; Chu and La Peyre, 1993; Ford et
al., 1993; Anderson et al., 1995). Many toxic chemicals can modulate densities of circulating
hemocytes, which may increase or decrease according to chemical characteristics and
concentrations (Auffret et al., 2002; Gagnaire, 2005; Auffret et al., 2006). Further,
modulation of hemocyte counts upon *A. minutum* exposure appeared to vary according to
maturation stage of the oysters. Indeed, mature diploid oysters responded in an opposite
manner compared to other oyster groups by decreasing hemocyte count upon *A. minutum*
exposure. Ripe oysters are known to be especially sensitive to stress and to summer mortality
(Samain et al., 2007). *A. minutum* exposure, serving as an additional stress to reproductive
effort, may have re-enforced the decrease in hemocyte counts occasionally observed in fully-
mature oysters (Delaporte et al., 2006). This makes changes of hemocyte concentration difficult to interpret. The strongest increase in hemocyte concentration occurred in the oyster groups showing the strongest increase in amylase activity upon A. minutum exposure. Hemocytes have been thought for a long time to be involved in digestion processes and digestive activities (Cheng, 1996). The observed parallel between hemocyte counts and amylase activities may simply reflect involvement of hemocytes in microalgal digestion and nutrient assimilation.

Reactive oxygen species (ROS) production and phenoloxidase (PO) specific activity in circulating cells were both affected by A. minutum exposure. ROS production is associated with internal chemical destruction of engulfed pathogens or foreign particles within hemocytes (Cheng, 1996; Cheng, 2000; Chu, 2000). Hemocyte ROS production may also be activated by high reproductive effort, leading some to consider reproductive activity as a physiological stress (Delaporte et al., 2006; Delaporte et al., 2007). The elevated energy demand for gamete production leads to a marked increase of whole-animal oxygen consumption during gametogenesis, corresponding to an elevated basal metabolism (Shumway et al., 1988). ROS production was thus hypothesized to reflect an oxidative stress during periods of high energy expenditure, such as active gametogenesis. ROS production was previously observed to increase in ripening oysters, especially in oysters known to be genetically more sensitive to summer mortalities (Delaporte et al., 2007). In the present study, ROS production was two times higher in oysters in the May experiment than in the April experiment. This suggests that ROS production may reflect increases in metabolic activities and energy expenditure formerly “programmed” by temperature and photo-period conditions of their rearing site. Temperature and photo-period may accelerate metabolic activities, even for triploid oysters which produced only few mature gametes and remained on average at intermediate stages of gametogenesis.
Upon *A. minutum* exposure, hemocyte ROS production significantly increased in the April experiment and significantly decreased in the May experiment, similarly in diploid and triploid oysters. In the April experiment, the increase in ROS production may reflect an increase in metabolic activities responding to *A. minutum* exposure. On the contrary, in the May experiment, we speculate that higher toxin accumulation exceeded the “tolerance” of oysters, affecting directly and more profoundly physiological and metabolic activities leading to reduced ROS production. At this point, we can only confidently conclude that changes in ROS production upon *A. minutum* exposure do not depend directly upon reproductive activities, as near fully-matured diploid oysters responded similarly to early-maturing triploid oysters. Also, as oysters in the May experiment accumulated twice the toxin of those in the April experiment, we cannot disprove the possibility that released toxins may act as a stimulant when present at low levels, while resulting in inhibitory effects when accumulated at higher concentrations. Similar observations were made in ecotoxicological studies assessing impacts of toxic chemicals upon immune functions in bivalves (Auffret and Oubella, 1997; Fournier et al., 2001; Auffret et al., 2002; Gagnaire et al., 2006). Several heavy metals were demonstrated to have a stimulatory effect upon hemocyte counts, chemotaxy, and mobility in field study. In contrast, in experimental studies, high concentrations of the same heavy metals were inhibitory to the same or related immune parameters (Auffret et al., 2002).

Phenoloxidase is activated by microbial substances and is thought to have a role in host defense in *C. gigas* (Hellio et al., 2007). PO specific activity was drastically reduced upon *A. minutum* exposure in both diploid and triploid oysters in the April experiment, but PO increased in diploids and decreased in triploids in the May experiment. Thus, mature diploid oysters respond in an opposite manner, compared to other oyster groups, to *A. minutum* exposure. *A. minutum* challenge seems to decrease the activity of PO in hemocytes from
oysters at early and intermediate maturation stages, and to increase activity in oysters with advanced maturation. Notably, PO activity decreased upon *A. minutum* exposure when concomitantly hemocyte counts increased. This suggests that hemocytes newly mobilized in the circulatory system may not be fully functional and are less able to produce PO activity.

Triploid oyster hemocytes were significantly larger (higher FSC) than those of diploid hemocytes, possibly because of their higher nuclear DNA content. Additionally, triploid oyster hemocytes were also more complex (higher SSC) than diploid hemocytes. As cell complexity is related to granule content, higher granule content in triploid hemocytes may parallel the higher hemocyte activities (ROS production as well as PO activity).

Triploid oysters were previously reported to have statistically-higher hemocyte phagocytosis, esterase, and peroxidase activities than diploid oysters (Gagnaire et al., 2007). If higher hemocyte chemical activities can confer a better tolerance to toxin accumulation, then triploid oysters may have a better ability to respond to *A. minutum* exposure than diploid oysters. As low toxin sensitivity results in high toxin accumulation (Bricelj and Shumway, 1998), triploids would be able to accumulate more toxin than diploids. Triploid hemocyte size and complexity were reduced upon *A. minutum* exposure in the May experiment but not in the April experiment. We therefore speculate that toxin accumulation was high enough to affect cell physiology, possibly through cell degranulation, upon toxin exposure.

To summarize, hemocyte responses to *in vivo* *A. minutum* exposure depended upon reproductive status, toxin accumulation, but also upon oyster physiological status in the field prior to collection for experiments. When considering only *A. minutum*-exposed oysters, positive linear correlations between PST accumulation and both hemocyte ROS production (granulocytes: $R^2 = 0.55$, $p<0.01$; hyalinocytes: $R^2 = 0.6$, $p<0.005$) and PO activity (total hemocytes: $R^2 = 0.74$, $p<0.005$) were found (data not shown). These relationships indicate
that these activities can be considered as good “markers” of metabolic activity useful in interpretation of physiologically-dependent differences in toxin accumulation. As stated above, however, these oxidative activities also may be modulated by other sources of stress, such harmful algae. Thus, it is difficult to unravel the respective influences of physiological status vs stress encountered by oysters.

Bivalve physiological responses to *Alexandrium* spp. exposure often are thought to be related to toxins (saxitoxin and derivates) affecting human health. It is not known, however, if these compounds are the ones affecting bivalves, or if other compounds produced by harmful algae can affect shellfish at a greater extent. Lush et al. (1997) reported that juvenile greenback flounder (*Rhombosolea taparina*) exposed to an *A. minutum* whole cell suspension showed gill damage characterized by severe epithelial swelling that was not related to PSTs. In addition to this ichthyotoxic effect of *A. minutum*, it has been reported that *A. minutum* showed potent toxic effects upon brine shrimp (*Artemia salina*) (Lush et al., 1996) and a harpacticoid copepod (*Euterpina acutifrons*) (Bagoien et al., 1996), independently of paralytic-toxin effects. Moreover, Ford et al. (2008) found no measurable effect of a PST-producing strain of *Alexandrium tamarense* on hemocytes of two bivalve species. Instead, extract from a non-PST-producing strain had a strong and consistent negative effect on hemocytes from two clam species, resulting in significantly-lower adherence and phagocytosis compared to a PST-producing strain and filtered seawater controls. These studies allow us to suggest that PSTs are not the only compounds responsible of *A. minutum* effects upon oyster physiology, but other active compounds are likely bio-active as well.

5 Conclusion
The most striking result of this study was the difference in PST accumulation between diploid and triploid oysters: triploids accumulate about twice the toxin as diploids. This difference may be attributable to differences in physiology linked to ploidy, especially during reproduction. This finding can have important implications, in terms of oyster production and risk management. Despite the finding that *A. minutum* exposure was not lethal to oysters, exposure to a toxin-producing microalga can significantly impact oyster physiology, as compared to non-toxic algae (*T. Iso*). *A. minutum* exposure affected several digestive and hematological parameters, and these responses were modulated by ploidy and maturation stage. Indeed, for some physiological parameters such as phenoloxidase activity and hemocyte concentration, ripe oysters responded in an inverted manner as compared to maturing oysters. We highlight, however, that observed effects of *A. minutum* were not only related to PSTs, but likely also to other bioactive compounds produced by *A. minutum*. Results of the present study showed that analyses of digestive-gland activities and composition (neutral lipids for example) can provide information on effects of *A. minutum* exposure upon oysters. Also, it could be productive to investigate other lipid classes, such as cell membrane constituents (polar lipids). Finally, biochemical approaches developed here can be complementary to histo-pathological observations as such methods have been successfully applied to assess impacts of toxic microalgae upon mollusks (Galimany et. al, 2008).

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**Figure captions**

**Table 1**: Oyster condition index, body dry weight and digestive gland wet weight according to ploidy and microalgal exposure, in April and May experiments. This table also includes the results of the MANOVAs testing ploidy and microalgal exposure effects separately in both experiments.

**Table 2**: Neutral lipid class contents (expressed as mg.g$^{-1}$ of tissue wet weight) in oyster digestive glands according to ploidy and microalgal exposure, in April and May experiments. This table also includes the results of the MANOVAs testing ploidy and microalgal exposure effects separately in both experiments.

**Table 3**: Effects of ploidy and microalgal exposure on oyster hemocyte and plasma variables, tested by MANOVA in April and May experiments.

**Fig. 1**: Oyster maturation stages (expressed as %, n = 12 oysters) according to ploidy in April (A) and May (B) experiments, regardless of dietary conditioning (*T. Iso* or *A. minutum*). Stages 0, 1, 2 and 3 correspond to reproductively inactive, early gametogenesis, late gametogenesis and ripe, respectively.

**Fig. 2**: PST content (mean of 3 pools of 4 oysters each, as µg STX equiv. 100 g$^{-1}$ of tissue wet weight, ± CI) in digestive gland of diploid and triploid oysters exposed to *A. minutum* in April (A) and May (B) experiments.

**Fig. 3**: Sum of monoacylglycerol (MG), diacylglycerol (DG) and free fatty acid (FFA) contents (mean of 3 pools of 4 oysters each, as mg.g$^{-1}$ of tissue wet weight, ± CI) in oyster digestive gland according to ploidy and microalgal exposure, in April (A) and May (B) experiments. □ indicates statistically significant difference according to microalgal exposures (MANOVA).

**Fig. 4**: Ratio between reserve (sterol ester + glycerid ether + triacylglycerol content) lipids and structural (sterol content) lipids (± CI) in oyster digestive gland according to ploidy and microalgal exposure, in April (A) and May (B) experiments. # and □ indicate statistically significant differences according to ploidy or microalgal exposure, respectively (MANOVA); * indicates statistically significant difference according to microalgal exposure (T-test).

**Fig. 5**: Amylase-specific activity expressed as amylase activity as IU per mg of total protein (mean of 3 pools of 4 oysters each, ± CI) in oyster digestive gland according ploidy and microalgal exposure, in April (A) and May (B) experiments. # indicates statistically
significant difference according to ploidy (MANOVA); * indicates statistically significant
difference according to microalgal exposure (T-test).

Fig. 6: Total hemocyte count (cells.ml\(^{-1}\)), ROS production in granulocytes (AU), and size of
hyalinocytes (AU) according to ploidy and microalgal exposure in April (A) and May (B)
experiments (mean of 12 individual oysters, ± CI). # and □ indicate statistically significant
differences according to ploidy and microalgal exposure, respectively (MANOVA); *
indicates statistically significant difference according to microalgal exposure (T-test). AU:
Arbitrary unit.

Fig. 7: Specific activity (SA) of phenoloxidase (PO) expressed as PO activity (IU) per mg of
total protein in hemocytes (mean of 12 individual oysters ± CI) according to ploidy and
microalgal exposure, in April (A) and May (B) experiments. # and □ indicate statistically
significant differences according to ploidy and microalgal exposure, respectively
(MANOVA); * indicates statistically significant difference according to microalgal exposure
(T-test).
Fig. 1
Fig. 2
Figure(s)

Fig. 3
**Figure(s)**

**A (¤)**

![Bar chart A](chart_A.png)

- **Diploids**
- **Triploids**

**B (#)**

![Bar chart B](chart_B.png)

- **Diploids**
- **Triploids**

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**Fig. 4**
Fig. 5

A (#)

Amylase specific activity

0.0 0.2 0.4 0.6 0.8 1.0

diploids triploids

B (#)

T. Iso
A. minutum

*
Fig. 6
Fig. 7
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P = ploidy; D = diet; P/D; interaction ploidy and diet; Significant differences are indicated by * when p<0.05, ** when p<0.01, *** when p<0.001; NS non-significant.
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P = ploidy; D = diet; P/D = interaction ploidy and diet; Significant differences are indicated by * when p<0.05, ** when p<0.01, *** when p<0.001; NS = non-significant.
Table 3:

<table>
<thead>
<tr>
<th>variables</th>
<th>April experiment</th>
<th>May experiment</th>
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<tbody>
<tr>
<td></td>
<td>ploidy</td>
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<tr>
<td>concentration of granulocytes</td>
<td>*</td>
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<tr>
<td>concentration of agranulocytes</td>
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<td>NS</td>
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<td>concentration of agranats</td>
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<tr>
<td>size of granulocytes</td>
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<tr>
<td>size of hyalinocytes</td>
<td>**</td>
<td>NS</td>
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<td>complexity of granulocytes</td>
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<td>complexity of hyalinocytes</td>
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<td>arcin (% phagocytic hemocytes)</td>
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<tr>
<td>ROS production of granulocytes</td>
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<tr>
<td>ROS production of hyalinocytes</td>
<td>*</td>
<td>NS</td>
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<tr>
<td>hemocyte phenoloxidase SA</td>
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</tbody>
</table>

Significant differences are indicated by * when p<0.05, ** when p<0.01, *** when p<0.001; NS non-significant.

The following variables were not presented in this table as these were not significantly affected by ploidy or microalgal exposure or the interaction (MANOVA): concentration of total hemocytes, concentration of hyalinocytes, % of dead hemocytes, and plasma variables (phenoloxidase specific activity, agglutination titer and hemolysis titer).