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Assessment of swimming behavior of Pacific oyster D-larvae (*Crassostrea gigas*) following exposure to model pollutants

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15 This study describes an image analysis method that has been used to analyze early life stages behavior 16 of native oysters (Crassostrea gigas) from the Arcachon Bay (SW, France). In a second time, this study 17 evaluated the impact of Cu and S-metolachlor pollutants on D-larvae behavior and possible 18 relationship between developmental malformations and abnormal swimming behavior. Analyses in 19 wild and cultivated oyster D-larvae were investigated during two breeding-seasons (2014 and 2015) at 20 different sampling sites and dates. In control condition, the average speed of larvae was 144 µm.s⁻¹ and 21 the maximum speed was 297 μ m.s⁻¹ while the trajectory is mainly rectilinear. In the presence of 22 environmental concentration of copper or S-metolachlor, no significant difference in maximum or 23 average larval speed was observed compared to the control condition but the percentage of circular 24 trajectory increased significantly while the rectilinear swimming larvae significantly declined. The 25 current study demonstrates that rectilinear trajectories are positively correlated to normal larvae 26 while larvae with shell anomalies are positively correlated to circular trajectories. This aberrant 27 behavior could impacts the survival and spread of larvae and consequently the recruitment and 28 colonization of new habitats.

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Key words: early life stage; Image analysis; speed; trajectory; malformations; copper; S metolachlor, swimming behavior.

- 32
- 33 Introduction

34 Behavioral analysis is increasingly used to study the effects of chemicals and drugs on humans and 35 other mammals (XXX). In recent years, many biological early warning systems have been developed 36 that evaluate the behavioral responses of aquatic organisms to water quality (Melvin and Wilson, 37 2013; Garaventa et al., 2010; Van der Schalie et al., 2001). In contrast, effects of contaminants on aquatic invertebrate behavior are less frequently studied in comparison to developmental or 38 39 reproductive toxicology (Melvin and Wilson, 2013; Scott and Sloman, 2004). However, behavioral 40 indicators of toxicity appear ideal for assessing the effects of pollutants on aquatic organisms since 41 they link physiological function with ecological processes (Scott and Sloman, 2004). In aquatic 42 toxicology, the link between behavioral science and impacts of toxic substances only took importance 43 over the last decade (review in Faimali et al., 2017). The behavior can be the result of adaptations to 44 environmental variables. Therefore, a selective response is permanently adapted by direct interaction 45 with the physiological aspects of the chemical and physical social environment. Thus, the study of 46 behavioral parameters now represents valuable tools to identify and assess the effects of exposure to 47 environmental stressors. Until a few years ago, the development of tools facilitating image acquisition 48 and behavioral endpoints, could explain this phenomenon (Melvin and Wilson, 2013). In recent years, 49 much progress has been made in the technological tools available for quantifying behavior (Lv et al., 50 2013). Thus, the study of behavioral parameters now represents valuable tools to identify and assess 51 the effects of exposure to environmental stressors. Many studies have focused on the behavioral study 52 of fish larvae or adults (Caldwell et al., Handy et al., 1999; Kazlauskiene et al., 2010; Le Bihanic et al., 53 2015; Sommers et al., 2016; McCallum et al, 2017; Martin et al., 2017). Very few studies have focused 54 on mollusks and more particularly on bivalves (review in Faimali et al., 2017). However, bivalve 55 mollusks such as mussels, clams and oysters contribute significantly to world aquaculture production 56 and the same species have been used as sentinel for pollution monitoring of coastal marine waters 57 The Pacific oyster Crassostrea gigas (Thunberg) is ranked number one in terms of world aquaculture 58 production with 555 913 tons estimated in 2013 (FAO, 2015). Despite living attached for most of their 59 life stages, bivalves have a swimming larval stage, which play an important part for the colonization of the environment. In C. gigas embryos incubated at 20 °C, the first movements were observed in 60 61 trochophore at 6.5 hours post-fertilization (hpf) and most of individual (85%) can swim at 11.5 hpf 62 (Suquet et al., 2013). Up to now, only a few works has focused on the swimming behavior of early life stage of marine bivalves (Mileikovsky, 1973; Toost et al., 2008) and only a few ones have investigated 63 64 the abiotic factors as pH, salinity (Suquet et al., 2012; Suquet et al., 2013) and pollutants (Horiguchi 65 et al., 1998) on swimming capacities of bivalves. To our knowledge, none investigated the impact of pollutants on oyster larval swimming capacities. 66

67 To analyze the possible impact of pollutants on oyster larval behavior, two pesticides, copper and 68 metolachlor, frequently occurring in the marine coastal water were tested. Copper (Cu) is among the 69 most hazardous metal for bivalve larvae (His et al., 1999). Although Cu is an essential micronutrient for 70 living organisms (Festa and Thiele, 2011), it can be toxic above a certain concentration depending on the organism (Flemming and Trevors, 1989). All over the world, Cu is used both as fungicide in fruit 71 72 culture and as part of antifouling paint to prevent aquatic organisms from attaching themselves to the 73 hulls of vessels (Turner, 2010). Copper concentrations in the Arcachon Bay have been increasing for 74 several years and current levels exceed µg.L⁻¹ (Gamain et al., 2016). The impact of copper on aquatic 75 organism behavior has already been documented on fish (Sommers et al., 2016). Pesticide 76 contamination in Arcachon Bay is widely dominated by the herbicide S-metolachlor at around 10 ng.L⁻ 77 ¹ and its metabolites (Auby et al., 2007; Budzinski et al., 2011; Gamain et al., 2016). In fact, Arcachon 78 Bay is end recipient of several rivers draining a watershed of 4138 km² (Auby et al., 2014), which is 79 mainly dominated by agriculture and urban areas. Metolachlor is used in agriculture to control pre-80 emergent and early post-emergent broadleaf and grass weeds, and is regularly detected in surface 81 water and groundwater.

The main objective of this paper is to study the behavior of D-larvae of oysters, *Crassostrea gigas* in response to copper or S-metolachlor exposure. In order to do that, a method based on image analysis was developed. The open-source program ImageJ (https://imagej.nih.gov/ij/) and the ImageJ Plugin wrMTrck were used. This plugin was initially developed to study the motility of the nematoda *Caenorhabditis elegans* while swimming in liquids (Nussbaum-krammer et al. 2015) and it presents therefore an interesting similarity with swimming larvae. The method allows analyzing multiple parameters including the total travelled length, the distance and the speed of larvae.

This paper aims to: 1) describe an image analysis method that has been used to analyze D-larvae behavior, 2) apply this process to assess the impact of Cu and S-metolachlor pollutants on D-larvae behavior and 3) investigate possible relationship between developmental malformations and abnormal swimming behavior.

93 2. Materials and methods

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95 **2.1 Method based on imagery**

2.1.1 Image acquisition

To analyze the behavior of the native oyster, wild and cultivated oysters were collected during oyster
breeding period (July and August 2014 and 2015) from three different sites in the Arcachon Bay:
Comprian, Les Jacquets and Banc d'Arguin (Fig. 1). Comprian is located close to the River L'Eyre and

therefore is under the influence of fresh water inputs in contrast to Les Jacquets or to Arguin. The reference water was collected near the Banc d'Arguin. Adult oysters were brought back to the laboratory in tanks filled with seawater from the sampled site and were kept overnight in this water continuously aerated and maintained at 12 °C.

104 Mature oysters (male and female) were induced to spawn by thermal stimulation, alternating 105 immersion in FSW of 15 °C and 28 °C for 30 min. Spawning males and females were individually isolated 106 in beakers containing 500 mL of 0.2 µm FSW at the spawning temperature. They were left undisturbed 107 for 15 min and were then removed from beakers. Eggs and sperm from two individuals were selected 108 to give a single pairing. Sperms and eggs were sieved separately through a 50 µm and 100 µm meshes 109 (Sefar Nitex), respectively to eliminate debris and feces. Sperm mobility was checked and the number 110 of eggs was counted under microscope (LEICA DME) at a magnification of 100. Eggs were fertilized with 111 sperm in ratio of 1:10 (egg:sperm) homogenizing with an agitator to prevent polyspermy. Fertilization 112 success was verified under microscope, and embryos were then counted and transferred to 24-well 113 microplate (Greiner Bio-One, Cellstar free of detectable DNase, RNase, human DNA and pyrogens) for 114 embryotoxicity assays. The embryotoxicity assay has been described in details by His et al. (1999) and 115 Quiniou et al. (2005) and normalized (AFNOR 2009). Fertilized eggs (around 300 eggs) were exposed 116 in wells containing 2 ml of toxicant solution. The eggs density was slightly changed compare to the 117 recommended AFNOR document specifying densities between 20 000 and 50 000 embryos per liter. 118 These microplates were incubated in climatic chamber at 24 °C for 24 h in the dark.

After 24h incubation, a 2-minute film was recorded by means of a microscope (Nikon Inverted
 Microscope Eclipse TS 100 / TS100-F, TS100LED MV / MV F-TS100 LED) and an acquisition software
 (NIS Element D). Recorded films were on a MPEG format and included 100 frames per second.

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2.1.2 Description of the image analysis process

A freeware (VirtualDub) for video conversion has been used to subsample the film to 4 fps and convert it into AVI format. The AVI format is used within ImageJ. The entire flow of analysis as well as the expected results are schematized on Fig. 2. The films (.avi) are opened as a stack of images (Fig. 2A) and converted into grayscale (Fig.2B). The entire stack of images is then converted into a binary stack of images using the maximum entropy (Kapur et al., 1985) of their histogram to stablish a threshold (Fig. 2C). When running the wrMTrck plugin, proper settings to track oyster larvae were selected (Table 1).

As a result, each tracked larvae had a number assigned, each number is used to identify the tracked
larvae in the result file. The result file (Fig. 2E) include: the track number, the length which is the sum
of all movement vectors (pixel), the distance which is the distance between the start and the finish

position (pixel), the number of frames, the first frame, the total time (seconds), the maximum speed (pixel/second), the mean and the standard deviation of the surface of a larva during the tracked period (pixel²), the mean and the standard deviation of the perimeter of a larva during the tracked period (pixel), the average speed, the body length per second, the average X position of the larva during the tracked period and the average Y position of the larva during the tracked period. A drawing gathering all tracked larval paths detected throughout the video (Fig. 2D) is also obtained as a result. The gathering drawing allowed to identify three different types of traces of larval path: (1) Rectilinear,

defined as the trajectory in which the length and the distance are quite similar (Track 1 Fig. 3), (2)
Circular, defined as the trajectory in which the length is a lot bigger than the distance (Track 2 Fig. 3),
(3) motionless, in which the length has a low value (Track 3 from Fig. 3). It is important to note that a
single larva can be detected multiple times since they can exit and enter the field of view (Track 4 from
Fig. 3). Furthermore, two larvae may collide, if it is the case, their larval path may be altered (Track 5

- 146 from Fig. 3).
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Table 1: Optimized settings for wrMTrck plugin

Parameters for tracking oyster D-larvae in 'wrMTrck'	Value
Minimum size (pixel ²)	400
Maximum size (pixel ²)	1000
Maximum velocity (pixels/frame)	100
Maximum area change (%)	50
Minimum track length	20
Threshold for turn	2
Size of bin for speed histogram (pixel frame)	0
Show Path Length	Yes
Show Labels	Yes
Show Positions	Yes
Show Paths	Yes
Show summary	Yes
Smoothing	Yes
Raw Data (0:off, 5:bendcalc)	0
Ben detection (0:off, 3:AR+Histogram)	0
Frame per second	4
Background subtraction (0:off)	0

Threshold method	Otsu
Size of labeling font	18

151 **2.2 Developmental abnormalities**

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At the end of the 24 h incubation, 25 μL of 1% buffered formalin were added in each well and the percentage of abnormal larvae was recorded. Hundred individuals per well were directly observed under inverted microscope (Nikon eclipse TS100/TS100-F; TS100 LED MV/ TS100 LED-F MV) to determine the number of abnormal D-shell larvae according to the criteria described in His et al. (1999) and Quiniou et al. (2005). An important prerequisite for this test is the presence, in control condition (24 °C in the absence of contamination) of less than 20 % of abnormal larvae. In this experimentation, four different couples were used and four replicates were performed for each condition.

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161 **2.3 Chemical analyses**

Reference toxicants (CuSO₄ and S-metolachlor) and formalin were purchased from Sigma–Aldrich Chemical (St. Quentin Fallavier, France). Seawater was collected outside the Arcachon Bay (SW France) near the Banc d'Arguin. Immediately after sampling, seawater was filtered using membrane filter of 0.45 µm and then 0.2 µm (Xilab) to eliminate debris and microorganisms. FSW was stored at 4 °C in the dark and was used within 3 days. A few hours before experiment, FSW was filtered again at 0.2 µm. This reference water was chemically analyzed for pollutant concentration determination.

The range of test concentrations was chosen on the basis of preliminary studies (Gamain et al., 2016).
The metal and pesticide solutions were made up from analytical grade copper sulfate (CuSO₄, 5H₂O)
and S-metolachlor. Working solutions were obtained diluting the stock solutions (100 mg.L⁻¹ for copper
and 250 mg.L⁻¹ for S-metolachlor) in FSW and were chemically analyzed. Two concentrations of
exposure were selected for copper: 1 and 10 µg.L⁻¹ and three for -S-metolachlor: 10, 100 and 1000
ng.L⁻¹. The working solutions were chemically analyzed to confirm pollutant concentrations.

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175 2.3.1 Copper analysis

For chemical analysis of copper at 1 and 10 μg.L⁻¹, each seawater sample was acidified with 5% final of
nitric acid (Nitric acid 65%, Fluka). Samples were then analyzed by Inductively-Coupled Plasma Optic
Emission Spectrometry (ICP-OES, Vista Pro, Agilent Technologies) and by Inductively-Coupled Plasma
Mass Spectrometry (ICP-MS, Xseries2, Thermofisher Scientific). The standards solutions were prepared
from a multi-elementary calibration solution (Astasol-Mix M010, Analytika, Czech Republic), in a
seawater certified solution (NASS-6 from NRCC-CNRC, Ottawa, Canada). The samples were diluted in a

3% final nitric acid solution (made from a nitric acid 65% FisherScientific Trace Metal Grade solution)
1:2 (v/v) for ICP-OES analysis and 1:3 (v/v) for ICP-MS analysis. Quantification limit were 10 µg.L⁻¹ (ICP0ES) and 0.3 µg.L⁻¹ (ICP-MS).

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2.3.2 Metolachlor analysis

187 Chemical extraction

188 50-mL water samples (pH adjusted to 7.0 ± 0.1 with HCl 0.1 N) were filtered using GF/F glass microfiber 189 filters (0.7 µm pore size). Before analysis, pre-concentration of the analytes was performed using Solid-190 Phase Extraction (SPE) with Oasis HLB cartridges (Waters), according to the method described by 191 Lissalde et al. (2011). SPE was conducted using a Visiprep 12-port manifold (Supelco, France). The 192 conditioning, extraction and rising steps were carried out under a 53.33 kPa vaccum. The SPE cartridges 193 were successively washed with 5 mL of methanol, conditioned with 5 mL of ultrapure water, loaded 194 with 50 mL water samples, then rinsed with 5 mL of UPW containing 15% HPLC grade methanol. 195 Cartridges were then dried under a nitrogen stream for 30 min. Elutions were achieved with 3 mL of 196 methanol, followed by 3 mL of a mix of methanol: ethyl acetate (75:25 v/v). 2.5 µL of a solution of internal standard (metolachlor d6) at 1 ng μ L⁻¹ was then added to the 6-mL extracts, followed by a 197 solvent evaporation under a gentle stream of nitrogen and then dissolved in 250 µL of UPW containing 198 199 10% HPLC grade acetonitrile prior to analysis.

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201 Instrumentation and data treatment

202 Metolachlor analyses were performed by liquid chromatography ACQUITY UPLC H-Class coupled to a 203 Xevo G2-S TOF-MS (Waters). The electrospray source was operated in positive mode at 0.7 kV and the 204 sample cone voltage set at 30 V. Nitrogen was used as nebulizer (flow rate 50 L h⁻¹, 150 $^{\circ}$ C) and 205 desolvation gas (flow rate 1200 L h⁻¹, temperature 600 °C). Data was acquired in the range from 50 to 1200 m/z and acquisition speed was set to 0.2 s. The resolving power full width at half maximum 206 207 (FWHM) was 30,000 at m/z 556.2771 (leucine encephalin used as lockmass compound). Data was 208 acquired using MSE in order to obtain both protonated molecular ions at low collision energy (CE=6 209 eV) and/or adducts and fragment ions with a collision energy ramp (CE=10-30 eV). Separation was 210 performed on an ACQUITY BEH C18 column (100 × 2.1 mm, 1.7 µm) from Waters (Milford, MA, USA) with a column temperature of 45 °C and using a binary gradient of water (A) and methanol (B) both 211 212 containing ammonium acetate (10 mM) at pH 5.0. A flow rate of 0.45 mL min⁻¹ was used and the 213 gradient ranged as follows: 98% A (0–0.25 min), 1% A (12.25–13 min), 98% A (13.01–17.00). Injection 214 volume was 20 µL. Data treatment was performed with MassLynx v4.1.

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216 Method validation and quality controls

Analytical method was validated in terms of calibration linearity, extraction recoveries, and limits of quantifications (LOQ) according to the French standard NF T90-210. Recovery, LOQ and LOD are showed in Table for the metolachlor into the different samples. For the quality controls, SPE was carried out routinely controlled, and the recoveries of two levels of spiked mineral water (e.g. 40 and 200 ng L⁻¹) were evaluated for each batch. The periodic control of two calibrating standards (e.g. 2 and 25 µg L⁻¹, every 10 samples) and analytical blanks were performed as well.

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Table 2. Acquisition and validation data for the metolachlor.

Compound	Retention time (min)	Quantification ion (m/z)	Confirmation ion (m/z)	Internal standard	% Recovery (±1 SD)	LOQ ^a (ng.L ⁻ ¹)	LOD ^a (ng.L ⁻ ¹)
Metolachlor	9.52	284.142	176.144	Metolachlor d6	110 (22)	0.4	0.1
Metolachlor d6	9.49	289.172	182.181	N/A	N/A	N/A	N/A

- ^a Limits of quantification and detection, respectively.
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- 228
- 229

230 2.4 Statistical analysis

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All data is expressed as means ± standard deviation (S.D.). Data was first processed using the 232 transformation: $p' = \arcsin \sqrt{p}$. P corresponds to the raw data (frequency of abnormalities) 233 specified in p values from 0 to 1 (Legendre and Legendre 1998). Homogeneity of variance (Levene's 234 235 test) was verified and statistical analysis was performed by the Kruskal-Wallis tests. Differences among 236 data from different conditions were tested using Kruskal post hoc test (equivalent to the Tuckey HSD 237 test for non-parametric data). 238 Principal component analysis (PCA) was performed using XXX to analysis both the spatial distribution 239 pattern of the different studied parameters and the relative relationship between treatment groups. 240

241 **3. Results**

- 243 3.1 Chemical analysis
- 244

245 The copper and S-metolachlor concentrations measured in the reference seawater and the different contamination solutions are shown in Table 3. Analyzes revealed the presence of copper in the 246 reference seawater at concentrations of 3.6 µg.L⁻¹. At the nominal concentration of 1 µg.L⁻¹, the 247 measured concentration was closed to that of control (3.75 µg.L⁻¹). At 10 µg.L⁻¹, the measured 248 249 concentration was 18% above expected concentration. Analyzes revealed the presence of S-250 metolachlor in the reference seawater at concentrations lower than 5 ng.L⁻¹. At the nominal 251 concentration of 10 ng.L⁻¹, measured concentration was 18 ng L⁻¹. At 100 ng.L⁻¹ and 1000 ng.L⁻¹ 252 measured concentrations were closed to expected one with a range of variation of the order of 8.1% 253 and 8.4% respectively.

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Table 3: Nominal and measured copper and S-metolachlor concentrations (mean values ± SD) at the
 beginning of the embryotoxicity test

	Nominal concentrations	Control (FSW)	1	10	100	1000
Measured	Copper (µg L ⁻¹)	3.63	3.75	11.8	-	-
concentration	S-metolachlor (ng L ⁻¹)	4.6	-	18	108.1	916

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261 3.2 Swimming activity

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In the control, the average speed of larvae was 144.4±34.6 µm.s⁻¹ and the maximum speed was 263 264 297.3 \pm 85.2 µm.s⁻¹ (Fig. 4) while the trajectory is mainly rectilinear (80.8 \pm 15.6 % of larvae) and to a 265 lesser extent circular (17.5±13.3 %). (Fig. 5). Motionless larvae are poorly observed in the control group 266 (1.9±3.1 %). In the presence of copper, no significant difference in maximum or average larval speed was observed at the concentration of $1 \mu g.L^{-1}$ or $10 \mu g.L^{-1}$ compared to the control condition (Fig. 4A). 267 268 After exposure to S-metolachlor, no significant difference in maximum or average larval speed was observed at the concentrations of 1 μ g.L⁻¹ and 10 μ g.L⁻¹ compared to the control condition (Fig. 4B). 269 270 Following copper exposure at both concentrations, the percentage of circular trajectory increased 271 significantly while the rectilinear swimming larvae significantly declined and a dose-dependent effect 272 was observed (Fig. 5A). In the presence of S-metolachlor, the percentage of circular trajectory tends 273 to increase with increasing S-metolachlor concentrations but it was only significant at 100 ng.L⁻¹ (Fig. 274 5B). In the meanwhile, rectilinear trajectories were reduced at all concentrations tested but it was only 275 significant at 10 and 1000 ng.L⁻¹.

277 3.3 Malformations

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279	Table 3: Abnormal larvae and	malformation	frequency in	oyster	larvae	exposed	to copper	or S-
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280 Metolachlore

Conditions	Abnormal	Development	Mantle	Shell anomalies
	larvae	arrest	anomalies	
Control	17.0 ± 2.42	4.05 ± 1.85	7.4 ±	5.55 ±
Cu 1	36.98 ± 3.30	10.94 ± 2.49	14.5 ±	11.79 ±
Cu 10	60.31 ± 6.6	19.67 ± 1.61	25.89 ±	14.75 ±
Met 10	32.48 ± 11.51	14.38 ± 3.21	12.58 ±	5.52 ±
Met 100	47.75 ± 12.58	18.70 ± 5.52	12.15 ±	16.9 ±
Met 1000	59.15 ± 5.24	28.60 ± 9.89	16.20 ±	14.35 ±

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Abnormal larvae average frequency was 17.0 \pm 2.4 % in control condition (Fig. 6 and Table 3). The percentage of developmental arrest increase significantly at 10 µg.L⁻¹ of copper while a dosedependent increase of malformed larvae was observed at all copper concentration tested (Fig. 6A and Table 3). After exposure to S-metolachlor the percentage of developmental arrest was significantly increased from 100 ng.L⁻¹ whereas the percentage of abnormalities increased significantly only at 1000 ng.L⁻¹ (Fig. 6B and Table 3).

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289 3.4 Relationship between malformations and swimming behavior

PCA featured 60.88 % of the selected variables considering the two first axes. The first axis accounted
for 38.23% while le second axis accounted for 22.65% of the variability.

For the plot of variables (Fig 7A), the PC1 was positively loaded by both circular trajectories and shell abnormalities, and negatively loaded by rectilinear trajectories. PC2 was negatively loaded by maximum and average speeds. Rectilinear trajectories appeared positively correlated to normal larvae (r=0.47, p=0.012) while larvae with shell anomalies were positively correlated to circular trajectories (r=0.60, p=0.001). In addition, motionless larvae were obviously anti-correlated to maximum and average speeds.

For the plot of treatment groups (Fig. 7B), the PC1 and PC2 allowed a good separation of the control group versus copper 1 µg/L and copper 10 µg/L groups. The separation of the control group and

- 300 metolachlor 100 ng/L and metolachlor 1000 ng/L groups was well defined. In contrast, the metolachlor
- 301 10 ng/L group was partly overlapping with the control, metolachlor 100 and 1000 ng/L groups.
- 302 Le groupe contrôle est bien associée à des larves normales et à des trajectoires rectilignes (Figure 7
- 303 A&B). Les malformations du manteau sont corrélées principalement à l'exposition au cuivre à 10 μg/L
- 304 et au S-metalchlor à 1000 ng/L.
- 305

306 4. Discussion

307 4.1 Normal swimming behavior of oyster D-larvae

The herein described video tracking system represents an operational tool to evaluate different swimming parameters of D-larvae of the Pacific oyster *Crassostrea gigas*. A lot of others automated systems have been described to analyze swimming activity, motility or frequency of pulsation in numerous invertebrates (for review see Faimali et al., 2017) but to our knowledge, this study is the only one analyzing both swimming speed and trajectory.

Our semi-automated video tracking system was able to identified three major types of traces of larval path: rectilinear, circular and motionless. It is to be noted that the identified three traces of larval path do only consider two dimensions and that larvae oysters moving in their environment can move in three dimensions. A study on three dimensions would not affect rectilinear paths but circular paths might split between real circles and 3D spirals.

In our experiment, under optimal development conditions (24 °C, salinity of 33 usi), oyster larvae mainly adopted rectilinear trajectories (81% ± 15). Thus, the most common behavior of the larvae would correspond to rectilinear swimming. These results confirm the early work of His et al. in 1989 on the mussel *Mytilus galloprovincialis*, which suggested that a circular or spiral swimming denoted an erratic behavior of the larvae.

323 In the same laboratory conditions (24 °C, salinity of 33 usi), the average speed and the mean maximum velocity of oyster D-larvae were 144 and 297 µm.s-1, respectively. Suquet et al., (2012) reported an 324 average swimming speed of 105 μ m.s⁻¹ for C. gigas D-larvae in the absence of contamination. This 325 326 value is slight lower than the one calculated in our study. This difference can be explained, at least in 327 part, by the temperature difference between the two experiments, e.g. 19 °C for Suquet's study and 328 24°C for this study. 24 °C correspond to the ideal temperature of development for C. gigas (AFNOR, 329 2009, Gamain et al., 2016). Incubation of C. giggs embryos at temperature less than or equal to 20 ° C 330 give rise to significant increase of developmental abnormalities (Gamain et al., 2016). Those 331 abnormalities have likely consequences on the behavior and the speed of swimming of larvae. Indeed,

Horiguchi et al., (1998) reported unusual swimming behavior, low swimming activity and irregular
 movement of cilia due to atrophy of velum in larvae of different mollusk species exposed to organotin
 compounds.

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336 4.2. Effect of model pollutants on swimming behavior of oyster D-larvae

337 In our experimental setup (24 °C, salinity of 33 usi), there are no significant differences of the average 338 and maximum swimming speeds up to 10 μ g/L of copper and 1 μ g/L of S-metolachlor. In contrast, the 339 percentage of rectilinear larval paths decreased with the concentration of copper and S-metolachlor 340 (except for the 1000 ng/L) while the percentage of circular larval paths increases with the 341 concentration of copper and S-metolachlor (except for the 1000 ng/L). Absence of effect on swimming 342 speed means that the two compounds did not targeted the metabolism and/or the neuro-muscular 343 system of oyster D-larvae. Besides, a significant change of the larval trajectory was observed at the 344 lowest tested concentrations of copper (1 µg/L) and S-metolachlor (10 ng/L) with a striking shift of 345 rectilinear paths to circular ones. This trajectory shift was likely associated to the appearance of 346 morphological anomalies in developing oyster embryos since a significant positive correlation was 347 noted between circular trajectory and shell abnormality of D-larvae.

348 No data is available about effect of S-metolachlor on swimming activity of aquatic organisms. Besides, 349 a few works have documented effects of copper on fish and invertebrates. LaBretech et al. (2002) 350 observed a decrease of motility in larvae of Mercenaria mercenaria exposed to concentration of Cu 351 above 10 µg/L. Kwok and Ang (2013) also reported a significant inhibition of swimming activity of larvae 352 of the coral *Platigyra acuta* at concentration above 40 μ g/L. After 48h of depuration the larval motility 353 was not recovered indicating a possible persistent effect of copper. In zebrafish larvae exposed to Cu 354 concentrations from 9-20 µg/L, the mean and the maximal velocity were significantly increased, but 355 not effect was observed at higher concentrations (Da Silva Acosta et al., 2016).

356 Toxicity of copper and S-metolachlor on early life stage of aquatic species is well document. It has been 357 shown that both compounds induced significant developmental anomalies and DNA damage in 358 bivalves at environmentally realistic concentrations (Mai et al., 2012; 2013; Gamain et al., 2017). 359 Behavior is an integrated and whole-organism response (Faimali et al., 2017). Punctual and/or limited 360 modifications of this behavior is assume to be an immediate and adaptive response to stress. At the 361 contrary, permanent and deep perturbations of behavior are the results of irreversible changes at the 362 sub-individual level that can have dramatically impact on fitness and survival of individuals. The 363 significant positive correlation between circular trajectory and anomalies of the shell in oyster larvae 364 exposed to copper or S-metholachlor demonstrate the deep impact of these molecules on the embryolarval development of this specie. This study demonstrates, once again, the very high sensitivity of the
early development stages of oyster to chemical contamination.

367

368 **4.3.** Application of swimming activity monitoring of oyster larvae

Since a decade, researchers are using swimming activity in model vertebrates (Le Bihanic et al., 2015; XXX) and invertebrates (Chevalier et al., 2015; XXXX) as new endpoints to evaluate toxicity of chemicals. This kind of endpoint is increasingly used for environmental risk or water quality assessment in non-model organism such as crustacean and sea urchin (Garaventa et al., 2010; Morgana et al., 2016). Although mussels and oysters are widely used as sentinel species for pollution monitoring, to date very few works focused on swimming behavior of bivalves exposed to pollutants (Horiguchi et al., 1998).

In the present study, effect of pollutants was clearly detected on swimming behavior of oyster larvae at concentrations as low as 1 μ g/L of copper and 10 ng/L of S-metolachlor. These levels of contamination are environmentally relevant and are currently detected in seawater of numerous European coastal areas (XXXX) and notably in the Arcachon Bay (Gamain et al., 2017b). The swimming activity of oyster larvae could be used as an early warming indicator of the quality of water and/or of the toxicity of pollutants.

Abnormal swimming behavior of D-larvae can directly affect their survival or fitness. In addition, abnormal trajectories can impair dispersion of larvae and colonization of new habitats. Both phenomenon could impair recruitment and at terms impact survival of wild population and viability of production of oysters. In this respect, swimming behavior could be used as a suitable tool to analyze the health status of larvae from wild or farmed oyster populations.

In the future, we can expect the use of video-tracking of oyster larvae as a convenient tool to guide the shellfish farmers towards the choice of the best batch of larvae and environmental agencies for the assessment of coastal water quality.

390

391 Conclusion

This study describes a semi-automated image analysis tool to analyze D-larvae swimming behavior of bivalves. Oyster larvae exposed to environmental concentrations of copper or S-metolachlor showed aberrant swimming trajectories but average and maximum speeds were not impacted. Rectilinear trajectories appeared positively correlated to normal larvae while larvae with shell anomalies were positively correlated to circular trajectories. This study showed that video-tracking of oyster swimming behavior is a sensitive and easy to perform tool for ecotoxicological testing. The use of this tool at
 large scale could need some improvements in particular for temperature control and automating of
 trajectory reading.

400

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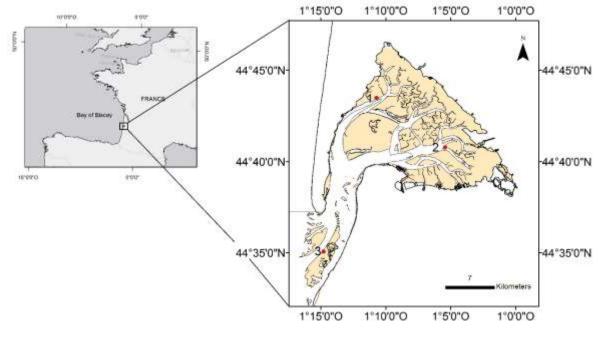
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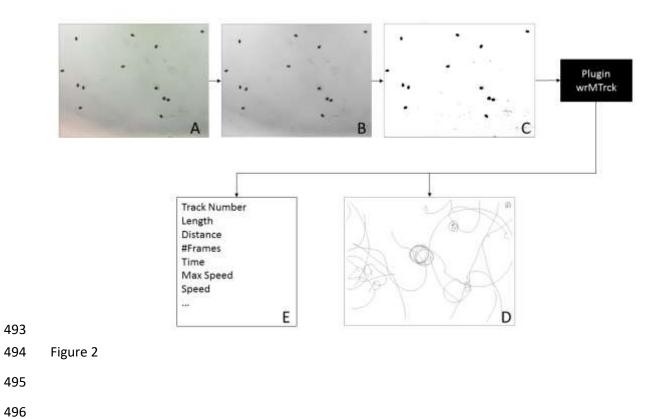
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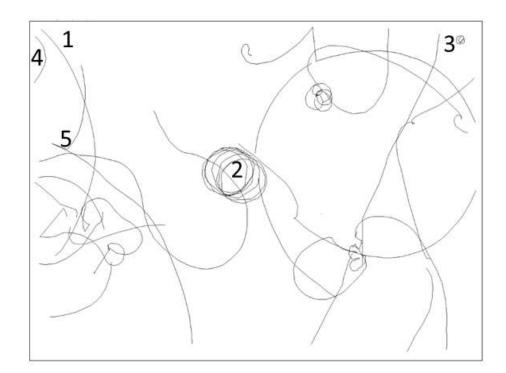
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- 462 Figures
- Figure 1: sampling site for mature oysters in the Arcachon Bay (SX France): 1) Les Jacquets, 2) Comprian
 and 3) Banc d'Arguin.
- Figure 2: Image Analysis: A) Original image, B) Grayscale image, C) 'Binarized' image, D) Traces of larval
 path, E) Quantified parameters.

- 467 Figure 3: Gathering drawing with different traces of larval paths: 1) Rectilinear, 2) Circular, 3)
 468 Motionless, 4) larval enters and exits field of view, 5) collision between two larvae.
- 469 **Figure 4:** Average and maximum speed (μm.s⁻¹) of D-larvae exposed to different concentrations of (A)
- 470 copper (μg.L⁻¹) or (B) S-metolachlor (ng.L⁻¹). Data points represent mean ± SD with 3-5 replicates per
 471 concentration and 12-74 larvae for each replicate.
- 472 **Figure 5:** Trajectories of D-larvae exposed to different concentrations of (A) copper (μ g.L⁻¹) or (B) S-473 metolachlor (ng.L⁻¹). Data points represent mean ± SD with 3-5 replicates per concentration and 12-74 474 larvae for each replicate. Different letters indicated significant differences between exposure 475 conditions (p<0.05).
- 476 **Figure 6:** Percentages of developmental arrest (DA) and abnormal D-larvae exposed to different 477 concentrations of (A) cooper (μ g.L⁻¹) or (B) S-metolachlor (ng.L⁻¹). Data points represent mean ± SD. 478 N=4 replicates per condition, and 100 larvae for each replicate. Different letters indicated significant 479 differences between exposure conditions (p<0.05).
- 480 Figure 7: Principal component analysis of the different types of abnormalities and trajectories of D-
- 481 larvae. Analysis represents normalized coefficients on the first two axes (axis 1: 38.23 %; axis 2: 22.65%)
- 482 for three D-larvae abnormalities (normal, shell abnormalities, mantle abnormalities), three trajectories
- 483 (R : Rectilinear ; S: Motionless, C: Circular) two speed (Vit Moy: average speed, Vit Max: maximum
 484 speed). The variables factor map is shown in (A) and the individual factor map in (B). Pearson
 485 correlation analysis between shell abnormalities and circular trajectories (C), and rectilinear
 486 trajectories and normal D-larvae (D).
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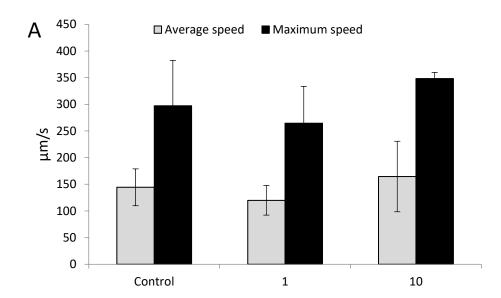


491 Figure 1

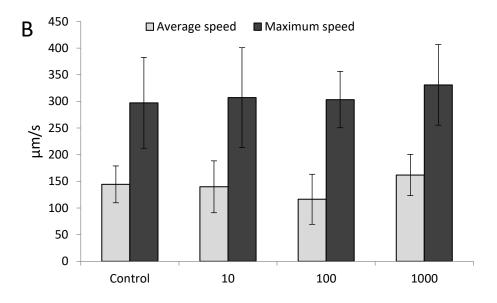




498 Figure 3

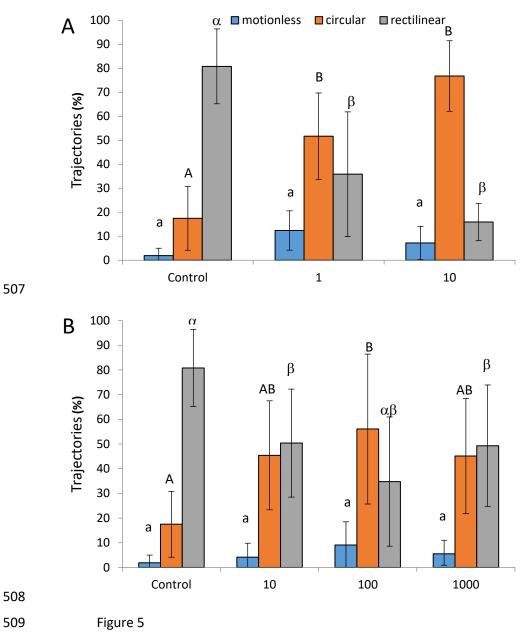


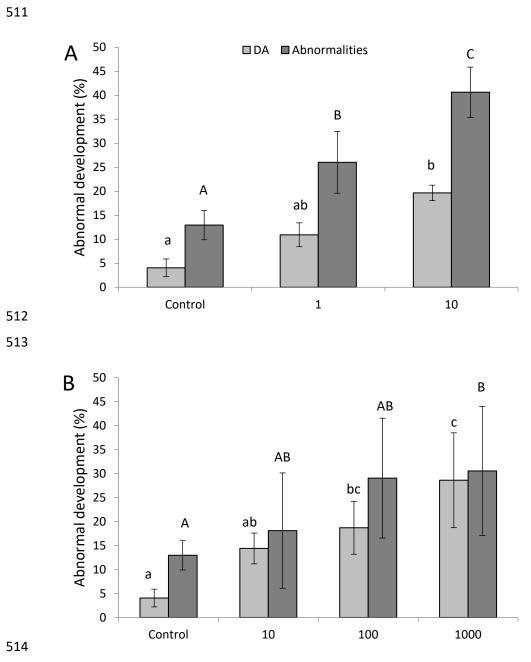




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Figure 4







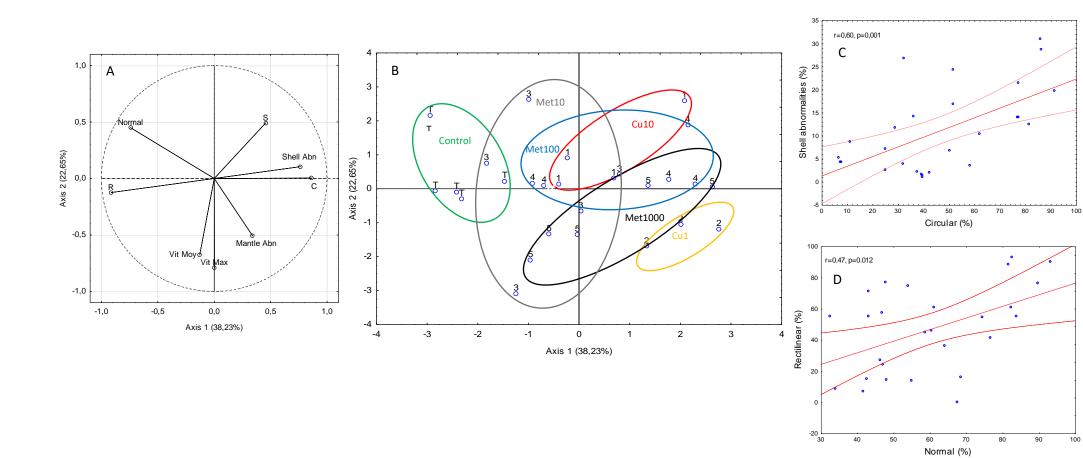


Figure 7