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OPEN Sperm motility of oysters from distinct populations differs in response to ocean acidification and freshening

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Species' responses to climate change will reflect variability in the effects of physiological selection that future conditions impose. Here, we considered the effects of ocean acidification (increases in pCO₂; 606, 925, 1250 µatm) and freshening (reductions in salinity; 33, 23, 13 PSU) on sperm motility in oysters (Crassostrea gigas) from two populations (one recently invaded, one established for 60+ years). Freshening reduced sperm motility in the established population, but this was offset by a positive effect of acidification. Freshening also reduced sperm motility in the recently invaded population, but acidification had no effect. Response direction, strength, and variance differed among individuals within each population. For the established population, freshening increased variance in sperm motility, and exposure to both acidification and freshening modified the performance rank of males (i.e. rank motility of sperm). In contrast, for the recently invaded population, freshening caused a smaller change in variance, and male performance rank was broadly consistent across treatments. That inter-population differences in response may be related to environmental history (recently invaded, or established), indicates this could influence scope for selection and adaptation. These results highlight the need to consider variation within and among population responses to forecast effects of multiple environmental change drivers.

Species' responses to climate change vary among levels of organisation. For example, differences have been found among Phyla¹, among populations of the same species²⁻⁴, and among individuals of the same populations⁵⁻⁷. Where variation in responses is large and heritable there is greater scope for selection^{8,9} and, thereby, the potential for fitness to be influenced 10,11. Consequently, understanding response variability among individuals is fundamental to identifying the potential for adaptation, and improving our ability to predict impacts of future change³.

Strong selective pressures on reproduction can carry over into subsequent life stages. For example, sperm selection plays a key role in shaping the subsequent generation in many broadcast-spawning marine organisms^{12,13}. Moreover, gametes of broadcast-spawning organisms are often particularly sensitive to selection as they are released into the environment and typically possess limited buffering capacities^{8,14,15}. Where environmental conditions modify gamete motility, selection may act upon individuals to result in the modification, or loss, of certain populations16

Ongoing ocean acidification will likely generate significant novel selection pressures. While many studies have indicated that ocean acidification could lead to reduced sperm motility^{8,17-19}, others have found it can remain unchanged²⁰, or even increase²¹. Yet ocean acidification does not operate in isolation, and many studies show that multiple simultaneous drivers can have very different effects to those of drivers operating alone²². Climate projections indicate than in many regions freshening - the decline in seawater salinity - will arise from altered patterns of precipitation and ice sheet melt^{23–25}. When experienced in isolation, freshening is reported to lower sperm motility in a range of species (e.g. Atlantic cod Gadus morhua²⁶, polychaete Galeolaria caespitosa, and

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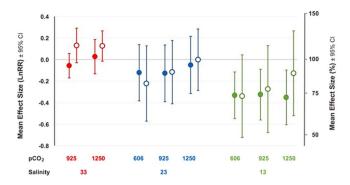


Figure 1. Mean effects (mean \log_e response ratio, LnRR, ±95% CI) of ocean acidification (pCO₂, μ atm) and freshening (salinity, PSU) on sperm motility (measured in terms of Sperm Accumulated At Surface, SAAS) in oysters from two populations (invasive, Sweden, closed circles and established, Guernsey, open circles; n = 14 per population). LnRR's calculated using 606 μ atm CO₂ × 33 PSU as reference. Note difference in scale on y-axis from Fig. 2.

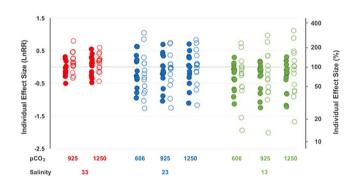


Figure 2. Individual effects (log_e response ratio, LnRR) of ocean acidification (pCO_2 , μatm) and freshening (salinity, PSU) on sperm motility (measured in terms of Sperm Accumulated At Surface, SAAS) in oysters from two populations (invasive, closed circles and established, open circles). LnRR's calculated using 606 μatm $CO_2 \times 33$ PSU as reference. Note difference in scale on y-axis from Fig. 1.

Source	df	MS	F	p	
Population	1	303.67	2.11	0.14	
Salinity	2	114.28	3.86	0.02	
pCO ₂	2	17.00	11.05	< 0.01	
Individual(Population)	26	144.04	106.51	< 0.01	
$Population \times Salinity$	2	12.64	0.43	0.65	
Population \times pCO_2	2	13.07	8.50	< 0.01	
Salinity $\times p$ CO $_2$	4	0.71	0.34	0.84	
$Salinity \times Individual (Population)$	52	29.62	21.90	< 0.01	
p CO $_2 \times$ Individual(Population)	52	1.54	1.14	0.25	
Population \times Salinity \times p CO $_2$	4	2.13	1.03	0.40	
Salinity \times p CO $_2$ \times Individual(Population)	104	2.06	1.53	< 0.01	
Residual	504	1.35			

Table 1. ANOVA on Sperm Accumulated At Surface (SAAS) of individual oysters from two populations (invasive, Sweden; established, Guernsey) exposed to ocean acidification (pCO_2) and freshening (salinity). $p \le 0.05$ shown in bold.

Pacific oyster $Crassostrea\ gigas^{27}$). However, the combined effects of freshening and ocean acidification on sperm motility remain unassessed.

We, therefore, measured the effects of ocean acidification and freshening on sperm motility at both the population and individual levels. To study these effects we used the model oyster species *C. gigas*. *C. gigas* is an important aquaculture and invasive species²⁸, whose persistence into the future may be modified as its reproduction can be influenced by both ocean acidification²⁹ and salinity³⁰. Here, we tested: (1) the effects of ocean acidification and

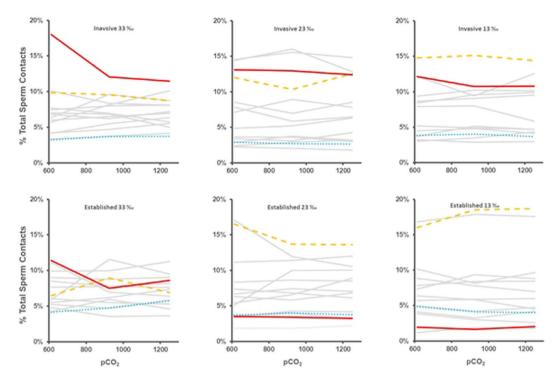


Figure 3. Change in male performance rank with ocean acidification (pCO_2 , μ atm) and freshening (salinity, PSU). Data are sperm contacts for each male as a percentage of total number of sperm contacts (sum of all contacts for all males) in each treatment. Lines represent individual males. For each population: the highest performing male in the ambient (control) treatment (606 μ atm CO_2 , 33%) is indicated by a solid/red line; the lowest performing male in the control treatment is indicated by a dotted/blue line; the highest performing male in the most extreme treatment (1250 μ atm CO_2 , 13%) is indicated by a dashed/orange line.

freshening on sperm motility; (2) if (and how) these effects varied between populations and among individuals; and (3) the impacts of acidification and freshening on male performance rank. We investigated these responses in C. gigas individuals sampled from two populations: one that invaded the low-salinity west coast of Sweden recently (within ≤ 10 years of this study; termed "invasive"), and one that has been established for 60+ years in the stable full-salinity waters of Guernsey, British Channel Islands (termed "established"). This work contributes to developing a broader understanding of the extent to which populations, and male performance broadly, will be modified in the future.

Results

Across populations, sperm motility declined with freshening (Fig. 1). Responses to acidification, however, differed; sperm motility increased with acidification in males from the established population, whereas sperm motility did not change substantively with acidification for those from the recently invaded population (open vs closed circles respectively, Fig. 1). Consequently, for established population males, the positive effects of acidification offset the negative effects of freshening such that their combination resulted in sperm motility that was equivalent to that in control conditions (i.e. LnRR \approx 0; Fig. 1). Similar antagonistic effects of freshening and acidification were not seen in males from the recently invaded population, which therefore showed freshening-induced reductions in sperm motility under future climate scenarios. These findings were reflected in the formal PERMANOVA results (Table 1), which showed that the two populations responded differently to pCO_2 (Population $\times pCO_2$, p=<0.01), and that responses to pCO_2 and freshening varied significantly among individuals (Salinity $\times pCO_2 \times Individual$).

Sperm from different males responded differently to the treatments in terms of their motility; some males showed improvements in sperm motility under acidification and freshening, whereas others were negatively affected (Fig. 2). Freshening caused a clear increase in response variance among males, especially in the established population (Fig. 2). In contrast, acidification had no effect on response variance.

Differential sensitivity of males to acidification and freshening was reflected in relative male performance, which was calculated for each treatment as the percentage of total sperm contacts (SAAS) for that population that were obtained by any given individual (Fig. 3). In the recently invaded population, males that performed best (i.e. had a high performance rank) in control conditions tended to retain performance rank under acidification and freshening (e.g. small changes in performance rank in solid-red, dotted-blue, and dashed-yellow lines in Fig. 3, upper panels). In the established population, however, the highest ranked male under control conditions was one of the poorest-performing males when exposed to acidification and freshening, and *vice versa* (change from 1st to 13th in solid-red line, and from 7th to 1st in dashed-yellow line, Fig. 3, lower panels). This population-wise difference in effect of acidification and freshening on male performance was seen clearly in correlations of performance

Freshening (salinity, PSU)	33		23			13		
Acidification (pCO ₂ , μatm)	925	1250	606	925	1250	606	925	1250
Invasive, Sweden	0.80	0.61	0.74	0.72	0.70	0.70	0.64	0.56
Established, Guernsey	0.60	0.66	0.20	0.27	0.36	0.35	0.16	0.27

Table 2. Correlation of sperm performance (SAAS rank) in the ambient (control) treatment (606 μatm $CO_2 \times 33$ PSU) with performance under acidification and freshening (Spearman's rank correlation coefficient ρ). ρ values with $p \le 0.05$ shown in bold.

rank among the treatments (Table 2). Rank order of performance in males from the recently invaded population under control conditions was significantly correlated with that under all combinations of acidification and freshening ($p \le 0.05$, Table 2), whereas performance rank in males from the established population were only significantly correlated between control conditions and acidification at 33 PSU. No negative rank-order correlations were observed.

Discussion

Our finding that different populations, and individuals, of oysters showed distinctly different responses to ocean acidification and freshening could have pervasive implications for this species under marine climate change. We measured sperm motility using a metric (SAAS²⁷) that integrates sperm swimming speed and percent motility, two key determinants of fertilisation success in free-spawning organisms^{31,32}. Consequently, the large changes in motility we observed under acidification and freshening imply that these drivers could cause correspondingly large shifts in fertilisation success for individuals³³. If sperm responses to acidification and freshening are heritable, these findings indicate there is substantial, and population-dependent, potential for selection and adaptation of this species under future climate change.

At the population level, clear patterns of response in oyster sperm motility emerged. Average sperm motility for the individuals we sampled from an established population increased with acidification, but decreased with freshening, such that these two drivers cancelled each other out at low salinity and high pCO_2 ; a combination that is likely to manifest regionally in the future. In contrast, sperm motility of oysters sampled from an invasive population was unaffected by acidification (a result also obtained previously²⁰), but decreased with freshening irrespective of pCO_2 . Such population-level differences in the effects of acidification have also been reported in another oyster (i.e. Sydney rock oyster, *Saccostrea glomerata*³). It is possible the different responses we observed were driven by parental environmental history³⁴ and/or local adaptation³⁵, although in that context it is surprising that oysters from the invasive population, which has recently undergone strong selection for tolerance to low salinities²⁸, generally responded more negatively to freshening than those oysters from the established population that has lived many decades in \approx 34 PSU. Equally, it is possible that some form of bottleneck or founder event has resulted in reduced genetic diversity of the invasive population (relative to the established population)^{36,37}. Irrespective of the mechanism, such population-specific findings highlight the difficulties – and dangers – of forecasting species responses to climate drivers based on investigations of only one population.

Each of the populations considered showed substantial inter-individual variation in sperm motility responses. While some oysters responded negatively to treatments, others showed no response, with some responding positively. Such patterns of strong inter-individual variability in responses can reflect differences among parental genotypes and environments that then influence gametes exposed to different environmental conditions^{38,39}. Alternatively, differences among individuals could result from non-genetic paternal effects such as differential gamete maturity. Variability between individuals has been reported previously in the context of organism responses to ocean acidification^{7,8,20,33,40}. We are, however, unaware of any other study that has investigated the effects of multiple climate change-related drivers on individual sperm performance (but for the combined effects of ocean acidification and a toxicant – i.e. copper – see⁴¹). The growing body of literature showing that inter-individual variability in responses may be common⁴², argues for shifting focus away from understanding "mean" responses and toward investigating changes in variance and (hence) the potential for differential individual success in a future ocean.

Importantly, we found that variability of individual responses itself varied among populations and treatments. Freshening led to a clear increase in inter-individual variance in sperm motility (SAAS), particularly in the established population. In contrast, acidification had no effect on response variance, which – within each salinity – remained more or less constant in both populations. As noted earlier, shifts in variance of these populations may have been influenced by the environment of the parental organism. That is, the narrower response variance seen in oysters from the invasive population (Fig. 2) could reflect the strong salinity-driven selection this invasive population has recently experienced²⁸, potentially reducing phenotypic variance. Alternately, this could be due to bottleneck and/or founder events that caused reductions in genetic diversity of these populations^{36,37}. Nonetheless, recent work has shown local selection in this species can be pervasive³⁵. Despite differences in magnitude of responses, the general pattern of increased variance under freshening indicates that differences among males will be amplified, a pattern which could have implications for male competitiveness during broadcast spawning.

The relative performance of sperm from different males shifted under the combined effects of acidification and freshening. For the established population, males that performed well under control (ambient) conditions performed poorly when exposed to acidification and freshening (Fig. 3, Table 2). This is the first such result for multiple climate drivers and extends a recent observation that acidification can change the male fitness landscape in broadcast spawners, and that this response can be modified by a co-factor (specifically a heavy metal⁵). In the invasive population, however, the relative performance of male oysters from the invasive population showed no substantive shift under freshening and acidification: males that performed well under control conditions

Treatment (nominal salinity, pCO ₂)	Salinity (PSU)	pН	Temp (°C)	A _T (μmol kg ⁻¹)	pCO ₂ (µatm)	Ω_{Ca}	$\Omega_{ m Ar}$
33,606	32.9 (<0.1)	8.02 (<0.01)	20.0 (0.2)	2261	620	3.23	2.09
33, 925	32.9 (<0.1)	7.86 (<0.01)	20.0 (0.2)	2261	938	2.34	1.51
33, 1250	33.0 (<0.1)	7.74 (<0.01)	20.0 (0.2)	2265	1270	1.82	1.18
23, 606	23.0 (<0.1)	8.03 (<0.01)	20.0 (0.2)	2004	604	2.44	1.52
23, 925	23.0 (<0.1)	7.86 (<0.01)	20.0 (0.2)	2004	927	1.71	1.07
23, 1250	23.0 (<0.1)	7.74 (<0.01)	20.0 (0.2)	2004	1245	1.32	0.82
13,606	13.0 (<0.1)	8.04 (<0.01)	20.0 (0.2)	1743	596	1.73	1.01
13, 925	13.0 (<0.1)	7.86 (<0.01)	20.0 (0.2)	1743	927	1.17	0.69
13, 1250	13.0 (<0.1)	7.73 (<0.01)	20.0 (0.2)	1743	1268	0.88	0.52

Table 3. Seawater chemistry of experimental treatments of ocean acidification (pCO_2) and freshening (salinity). Direct measures of salinity, pH, and temperature (Temp) were made on each experimental day (n=9 measures). Total alkalinity (A_T) was estimated from an established salinity:alkalinity relationship for the sea for this region (r=0.94). Partial pressure of CO_2 (pCO_2), and saturation states for calcite (Ω_{Ca}) and aragonite (Ω_{Ar}) using CO2calc (details in text). Data are mean (SE).

continued to perform well under freshening and acidification (Fig. 3, Table 2). Given that sperm motility is a key determinant of fertilisation success 31,32 – and hence adaptive capacity – continuing to explore how this feature can be modified by climate drivers and by the environmental history of a given population should be a focus of future research.

In conclusion, despite the burgeoning number of studies reporting effects of ocean acidification on the early life history of marine organisms, few have considered the importance of response variability, particularly for multiple climate drivers. Here, we show that the influence of ocean acidification on sperm motility and performance rank of individual male oysters can be strongly modified when combined with freshening, an effect that differs between populations and among individuals. This finding shows that quantifying inter- and intra-population response variability will be essential if we are to reliably project species' responses to future climate drivers.

Methods

Experimental design. We investigated responses of sperm from multiple male oysters from two populations (n = 14 individuals from each population), to three levels of pCO_2 and three levels of salinity in a fully-factorial design.

Experimental organisms. Pacific oysters were collected in the summer of 2015 from an "established" population, which has been exposed to oceanic seawater for over 60 years at Guernsey Sea Farms, British Channel Islands. This population is open to some immigration, and is genetically representative of farmed populations from which feral populations have established and spread. These oysters were conditioned to reproductive maturity by Guernsey Sea Farms Ltd, Vale, Channel Islands, and shipped to the Tjärnö Marine Laboratory, Sweden, immediately prior to experimentation. Oysters were also collected haphazardly from a recent "invasive" population (established \leq 10 years prior to this study²⁸) in western Sweden. This environment is characterised by highly variable salinity and pH (13–30 PSU, 7.7–8.1 pH)⁴³. Following collection, oysters from the invasive population were conditioned to reproductive maturity by Ostrea Sverige AB, South Koster, Sweden, with the same methods used by Guernsey Sea Farms Ltd. Conditioned oysters were held in temperature-controlled flowing seawater at the Tjärnö Marine Laboratory at 33 PSU and 17 °C (just below the temperature that elicits spawning) until experiments were run.

Experimental treatments. pCO $_2$ treatment levels were selected to reflect ambient inflowing seawater pCO $_2$ at the Tjärnö Marine Laboratory (606 μatm), late-century atmospheric pCO $_2$ (1250 μatm) under prevalent global climate change scenarios⁴⁴. Treatment levels were achieved by bubbling CO $_2$ -air mixtures into the treatment water until a given target pH_{NBS} was obtained. pH_{NBS} set-points equivalent to treatment pCO $_2$ levels were determined by calibrating pH_{NBS} against inflowing seawater equilibrated for 24 h with custom mixed gases of the relevant CO $_2$ concentration (AGA, Sweden AB). The response slope of the pH meter was calibrated daily using NBS buffers. Total alkalinity was estimated from salinity using a long-term salinity:alkalinity relationship for this location (r = 0.94)⁴³. This method has been shown to generate very low uncertainties in estimates of carbonate system parameters (uncertainties \leq ±0.006 pH_{NBS} and ±0.08 Ω_{Ar}) (for more details see⁴⁵). pCO $_2$ and saturation states for calcite (Ω_{Ca}) and aragonite (Ω_{Ar}) were calculated using CO2calc⁴⁶, with constants from Mehrbach *et al.*⁴⁷ adjusted by Dickson and Millero⁴⁸ (Table 3).

Salinity treatment levels were chosen to reflect the full-strength seawater typical of Guernsey (33 PSU), local sea surface salinity at Tjärnö, western Sweden (23 PSU), and sea-surface salinity in south-western Sweden (13 PSU), toward which the invasive oysters are spreading²⁸. Salinity was manipulated by diluting filtered sea water with filtered fresh water. Treatment levels were verified using a conductivity meter (WTW, Cond 3210, Germany) calibrated against laboratory salinity standards (Table 3).

Response variable: sperm motility (SAAS). Sperm motility for each replicate male (n = 14 per population), was quantified by measuring Sperm Accumulated Against Surface (SAAS)²⁷ under all combinations

of pCO₂ and salinity. For each male oyster, concentrated sperm were extracted directly from the testis using a Pasteur pipette inserted through a hole drilled in the shell above the gonad. Previous work has shown that any non-motile sperm released by this technique will have had minimal impact on the results²⁷. Namely, SAAS is almost exclusively caused by actively swimming sperm, and the sinking rate of dead/inactive sperm is so slow that it does not bias results obtained using the technique. Concentrated sperm were stored in an Eppendorf tube on ice. Sperm concentration in each suspension was verified by hemocytometer counts of an aliquot of sperm that had been immobilised and stained with Lugol's solution. For each male, independent sub-samples of sperm from the stock suspension were diluted into each of three replicate wells of a multi-well plate containing treatment seawater (three replicates per treatment combination and male). Volumes of sperm transferred were adjusted to ensure a final concentration of 2×10^6 sperm ml⁻¹ in the treatments. Sperm were exposed to the treatment seawater for 10 min, after which 1.5 ml of the suspension was pipetted to a new multi-well plate from which the pattern of accumulation of sperm on the bottom surface of the well over time was quantified. Sperm counts were determined using a phase-contrast inverted microscope (Leica, DMIL, Germany) equipped with a digital camera (PixeLINK, PL-D725CU, Canada). To quantify accumulation, still images of different (central) areas of the lower surface of the wells were taken 10 min after the addition of the sperm suspension to the well (n = 3 images per well). Digital images were post-processed and the number of sperm that had accumulated at the lower surface of the well (SAAS) counted manually.

Statistical analysis. The response of sperm motility (SAAS) to pCO_2 and salinity was determined by calculating logarithmic response ratios, LnRR = \log_e ([response in treatment]/[response in control]), for each treatment combination and male, using the 606 μ atm $CO_2 \times 33$ PSU treatment as the control. Unlike raw ratio data, \log_e response ratios are approximately normally distributed⁴⁹, and therefore we also calculated mean LnRR \pm 95% CI. We determined the statistical significance of sperm responses to the treatments using a three-way permutational analysis of variance (PERMANOVA) using the PRIMER package⁵⁰ with population, pCO_2 , and salinity as fixed factors, and individuals as a random factor (n = 14 per population, 3 replicates per male) nested within population. To reduce heterogeneity of variances, data were square root transformed before analysis.

Data Availability

The datasets generated during and/or analysed during the current study are available from the corresponding author on reasonable request.

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Author Contributions

L.J.F., C.A.S. and J.N.H. developed the project; L.J.F. conducted the experiments; L.J.F. and J.N.H. analysed and plotted the data; L.J.F., C.A.S. and J.N.H. prepared and edited the manuscript.

Additional Information

Competing Interests: The authors declare no competing interests.

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