

Individual Variability in Reproductive Success Determines Winners and Losers under Ocean Acidification: A Case Study with Sea Urchins

Peter Schlegel^{1*}, Jon N. Havenhand², Michael R. Gillings¹, Jane E. Williamson¹

¹ Department of Biological Sciences, Macquarie University, Sydney, Australia, ² Department of Biological & Environmental Sciences, University of Gothenburg, Tjärnö Marine Biological Laboratory, Strömstad, Sweden

Abstract

Background: Climate change will lead to intense selection on many organisms, particularly during susceptible early life stages. To date, most studies on the likely biotic effects of climate change have focused on the mean responses of pooled groups of animals. Consequently, the extent to which inter-individual variation mediates different selection responses has not been tested. Investigating this variation is important, since some individuals may be preadapted to future climate scenarios.

Methodology/Principal Findings: We examined the effect of CO₂-induced pH changes (“ocean acidification”) in sperm swimming behaviour on the fertilization success of the Australasian sea urchin *Heliocidaris erythrogramma*, focusing on the responses of separate individuals and pairs. Acidification significantly decreased the proportion of motile sperm but had no effect on sperm swimming speed. Subsequent fertilization experiments showed strong inter-individual variation in responses to ocean acidification, ranging from a 44% decrease to a 14% increase in fertilization success. This was partly explained by the significant relationship between decreases in percent sperm motility and fertilization success at $\Delta\text{pH}=0.3$, but not at $\Delta\text{pH}=0.5$.

Conclusions and Significance: The effects of ocean acidification on reproductive success varied markedly between individuals. Our results suggest that some individuals will exhibit enhanced fertilization success in acidified oceans, supporting the concept of ‘winners’ and ‘losers’ of climate change at an individual level. If these differences are heritable it is likely that ocean acidification will lead to selection against susceptible phenotypes as well as to rapid fixation of alleles that allow reproduction under more acidic conditions. This selection may ameliorate the biotic effects of climate change if taxa have sufficient extant genetic variation upon which selection can act.

Citation: Schlegel P, Havenhand JN, Gillings MR, Williamson JE (2012) Individual Variability in Reproductive Success Determines Winners and Losers under Ocean Acidification: A Case Study with Sea Urchins. PLoS ONE 7(12): e53118. doi:10.1371/journal.pone.0053118

Editor: Sam Dupont, University of Gothenburg, Sweden

Received: April 24, 2012; **Accepted:** November 28, 2012; **Published:** December 27, 2012

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Funding: This work was funded by the Australian Research Council and a Linnaeus-grant from the Swedish Research Councils VR-Review and Formas (Swedish Research Council for Environment, Agricultural Sciences and Spatial Planning)(www.cemeb.gu.se). The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Competing Interests: The authors have declared that no competing interests exist.

* E-mail: peter.schlegel@mq.edu.au

Introduction

Environmental factors directly affect populations by selecting resilient individuals. Selection at the gametic level, or during early life, has strong and immediate effects at the population level, carrying over into subsequent life stages. Heritability of this resilience leads to cascading adaptive effects in subsequent generations. For example, in free-spawning marine organisms, sperm selection during fertilization plays a key role by determining the nature and diversity of genotypes in the subsequent generation [1,2] and thus their resilience to environmental change.

Rising atmospheric carbon dioxide levels are a key driver of environmental change, and will likely lead to rapid ocean acidification [3,4]. With gametes possessing no, or only limited, buffering capacities against CO₂-mediated pH changes in seawater, the dynamics of fertilization and subsequent development are likely to be affected in all free-spawning marine

organisms, with potentially severe implications [5,6]. Yet we know little about the relative fitness of individuals within species under the predicted acidification of the ocean.

The sensitivity of reproductive processes to ocean acidification has thus far been assessed from mean responses of mixtures of gametes and/or larvae obtained from multiple individuals [7–10] (but see [11]). However, the key determinant of reproductive success in a future ocean is not the average response, but the proportion of successful offspring contributed by each individual under the changed environmental conditions. Individual-level responses to ocean acidification have been examined to some extent in larval development processes [12,13], but not closely in fertilization processes. In this context, the importance of naturally high variability that is observed in fertilization success of individual pairwise crosses [14,15] becomes apparent: not all matings are equal. Consequently, acidification-mediated impacts on reproductive success and subsequent development might result in flow-on

consequences for genetic diversity and population demographics [12].

The early life history stages of echinoderms are particularly useful for studies of fertilization success, as these species are experimentally tractable and ecologically important, often acting as ecosystem engineers [16,17]. Here, we investigate the effects of CO₂-induced ocean acidification on the early life history stages in the Australasian sea urchin *Heliocidaris erythrogramma*, focussing on intra-specific variation in responses, which can be highly variable for this species [18]. Following the A1FI-scenario from the IPCC's 4th assessment report [3], we compared the effects of present day conditions for southeast Australia with the end-of-century scenario (year 2100; pCO₂ = 970 μatm ≈ 0.3 pH unit reduction) and a high-CO₂ scenario (year 2300; pCO₂ = 1600 μatm ≈ 0.5 pH unit reduction). Observed effects on sperm swimming behaviour were applied within an established fertilization kinetics modelling framework [19,20] to predict fertilization outcomes of single urchin pairs at each pCO₂ level. These were then compared to observed results from fertilization experiments conducted in the laboratory.

Materials and Methods

CO₂ Treatment

Experimental CO₂ treatments were achieved by bubbling a mixture of air and CO₂ through filtered seawater (FSW; 0.22 μm filtered). pH (NBS scale) was maintained using microprocessor-controlled CO₂ micro-injection systems in separate FSW-tanks. Systems were set to maintain a pH change (ΔpH) of -0.5, -0.3 or 0 pH units (control, no addition of CO₂), resulting in treatment pH values of 7.6, 7.8 and 8.1. Dissolved oxygen levels were maintained by slow bubbling of filtered air. pH levels were checked prior to experiments. Total alkalinity was determined once by titration [21] (all seawater used in experiments was taken from a closed recirculating system in which alkalinity was controlled to be constant). Parameters of the CO₂ system (Table 1) were calculated with CO2SYS [22] using the dissociation constants of Dickson & Millero [23].

Experimental Animals

Heliocidaris erythrogramma (test diameter = 50.3 ± 1.3 mm, mean ± S.E.) were collected during their spawning season between February and March 2011, from shallow subtidal areas at Long Bay (33°57.5'S, 151°15.2'E) and Bare Island (33°59.2'S, 151°13.5'E), Sydney, Australia. Animals were immediately transported to Macquarie University and held in tanks with flowing seawater for up to one week maximum before being used for experiments. Individuals were collected fresh each week, alternat-

Table 1. Seawater parameters for the three different pH treatments.

pH _{NBS}	T (°C)	Sal	A _T (μeq kg ⁻¹)	pCO ₂ (μatm)	Ω _{Ca}	Ω _{Ar}
8.12 ± 0.06	20.5 ± 1	35.38 ± 0.06	2073 ± 4	413	3.674	2.392
7.80 ± 0.01	20.5 ± 1	35.38 ± 0.06	2073 ± 4	952	1.949	1.269
7.60 ± 0.01	20.5 ± 1	35.38 ± 0.06	2073 ± 4	1572	1.277	0.831

pH_{NBS}, temperature (T), salinity (Sal) and total alkalinity (A_T) were measured directly and used to compute partial pressure levels of carbon dioxide (pCO₂) and seawater saturation states for calcite and aragonite (Ω_{Ca} and Ω_{Ar} respectively) using CO2SYS. Means ± S.E.

doi:10.1371/journal.pone.0053118.t001

Table 2. Ocean acidification effects on sperm speed and percent sperm motility for each male *Heliocidaris erythrogramma*.

pH	Male	Site	Sperm speed (μm s ⁻¹)			Percent motility (%)		
			7.6	7.8	8.1	7.6	7.8	8.1
A	1	1	36.19	33.80	33.64	23.12	20.88	23.29
B	1	1	37.67	36.72	36.48	36.62	35.94	46.27
C	1	1	37.11	35.72	36.47	45.13	48.34	56.16
D	1	1	37.79	39.11	38.40	53.24	54.55	54.61
E	2	2	35.91	37.16	38.27	26.91	22.38	32.45
F	2	2	42.85	42.12	42.87	51.21	43.08	57.13
G	2	2	37.32	38.06	38.27	36.41	38.35	51.37
H	2	2	44.75	44.99	44.38	43.70	45.44	51.38
I	2	2	38.33	39.66	39.44	29.96	32.84	36.63
J	1	1	37.57	36.29	39.87	41.30	40.99	48.96
K	1	1	40.92	42.43	41.35	59.93	62.42	64.89
L	1	1	39.29	39.61	40.34	37.64	46.39	50.70
M	1	1	36.03	37.65	37.45	30.52	31.44	28.87
N	2	2	39.54	40.24	38.63	58.68	65.98	66.68
O	2	2	34.67	35.95	35.99	44.32	51.03	54.79
P	2	2	37.57	36.33	33.66	56.66	53.40	62.93
Q	2	2	41.18	41.32	39.19	53.23	60.35	57.81
R	1	1	35.76	34.14	34.64	70.00	68.86	74.13
S	1	1	41.88	41.54	38.40	45.40	53.60	56.64
mean			38.56	38.60	38.33	44.58	46.25	51.50
S.E.			0.32	0.32	0.36	1.08	1.17	1.10
P between sites			0.348	0.194	0.388	0.761	0.946	0.965

Significant differences in parameters between sites (P ≤ 0.05) are shown in bold. doi:10.1371/journal.pone.0053118.t002

ing between sites and those from different sites were never mixed. Three collections were done at Long Bay (males/pairs A–D on 7 Feb 2011, J–M on 21 Feb 2011, R and S on 14 March 2011; Table 2) and two at Bare Island (males/pairs E–I on 14 Feb 2011 and N–Q on 1 March 2011; Table 2).

Collection of Gametes

Gametes were obtained by injecting urchins with 0.8–1 ml of 0.5 M KCl through the peristomal membrane, followed by gentle shaking. This concentration of KCl reliably induced gamete release without being lethal. Individuals were used once (for convenience, the term “individual” will be used here to refer both to individual males [for sperm speed and motility measures] and to individual pairs [for fertilization success measures]). Eggs were collected in FSW, diluted to a concentration of 50 eggs ml⁻¹ and incubated in seawater at one of the three pH conditions for 10 min prior to use in experiments. Sperm were collected “dry” and held on ice until use to extend their lifespan. Experiments were conducted immediately after spawning. All experiments were done in a constant temperature room at a standard temperature of 20.5 ± 1°C (mean ± range).

Sperm Motility and Speed

Data were obtained for each of 19 male urchins. The motility assay followed that of Havenhand & Schlegel [24]. Briefly, 0.5–1 μl of freshly collected sperm from an individual male was diluted into 1 ml of seawater of each pH immediately prior to use (10 replicate sperm suspensions for each CO_2 treatment and male). Sperm concentrations across assays ranged consistently from $1\text{--}2 \times 10^4$ sperm μl^{-1} . A 60–70 μl drop of sperm suspension was placed between an albumin-coated microscope slide and coverslip, separated by a 0.75 mm thick O-ring. Sperm movement was recorded for 2s at the midpoint of the drop, at 25 frames s^{-1} , using a digital camera (Sumix SMX-160) mounted on a compound microscope (Olympus BX51). Pilot experiments showed illumination by the microscope lamp has no impact on the temperature inside the drop during videotaping (time of slide on microscope: approx. 10s). All recordings (one per sperm suspension) were done within 30s after creation of each sperm suspension. Videos were post-processed and analysed using CellTrak 1.3 (Motion Analysis Corporation) to determine sperm speed and percentage of motile sperm (i.e., sperm moving faster than $15 \mu\text{m s}^{-1}$ on average). Only sperm classified as motile were used for sperm speed analyses.

Fertilization Success

Data were obtained for 18 pairwise crossings using the same males used in the sperm motility experiments (above). Each male was crossed with one female (no female was spawned for the first male, hence $n = 18$). Each individual was used only once (18 males and 18 females in total). For each of the three CO_2 treatments, nine four-fold serially diluted sperm-concentrations and one control (FSW only) were prepared in 6-well plates holding 6 ml of seawater. Sperm concentrations and CO_2 treatments were distributed across multiple 6-well plates to preclude plate effects. Approximately 200 eggs were added to each well in filter dishes (25 mm diameter \times 20 mm height, 80 μm mesh floor). Eggs and sperm were mixed and left for 10 min to fertilize. Eggs were then rinsed twice (with water of the relevant pH) to remove sperm and left for 2 h to develop, (typically this was to the 4 cell stage). Fertilization success rates (FSR) were determined by counting the proportion of cleaved eggs *in vivo*. Sperm concentrations in the stock solutions were checked *post-hoc* using a haemocytometer (Neubauer improved). All fertilizations were conducted within 15 min of obtaining the gametes.

Modelling of Fertilization Kinetics

We used modelling [19,25] to combine the results from motility and fertilization experiments to investigate the importance of pH-induced changes in sperm motility in the fertilization process. Fertilization kinetics curves [20,26] were fitted to the measured fertilization data for each cross and CO_2 treatment. Fertilization efficiency (F_e) and polyspermy block efficiency (B_e) were estimated using least squares [27]. Sperm speeds and percentage of motile sperm for each male were taken from video analyses. Sperm concentration for modelling was defined as the product of the observed sperm concentration in the serial dilution and observed percent motility of that male in that treatment. For the control assay (pH 8.1) we identified the sperm concentration that yielded maximum fertilization success ($F_{\text{max Control}}$; Figure 1), the sperm concentration that yielded 50% of F_{max} ($F_{50 \text{ Control}}$ [28]) and the respective fertilization success rates at each of these sperm concentrations (“ $\text{FSR}_{\text{max Control}}$ ” and “ $\text{FSR}_{50 \text{ Control}}$ ”). For each of the CO_2 treatments fertilization success at $F_{50 \text{ Control}}$ was observed (“ FSR_{obs} ”) for each cross. These values were then compared to the fertilization success at $F_{50 \text{ Control}}$ that was

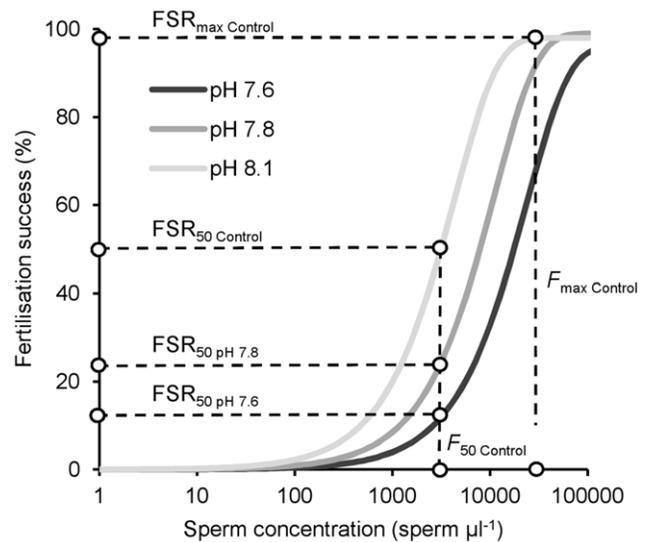


Figure 1. Schematic representation of the relationship between fertilization success and sperm concentration of *Helicidaris erythrogramma* at different pH levels, assuming a negative pH impact on fertilization. Determining fertilization success at an intermediate sperm concentration (here, the sperm concentration that generates 50% of maximum observed fertilization success in controls) yields maximum sensitivity in the assay. All data in Fig. 1 are theoretical. $\text{FSR}_{\text{max Control}}$ = maximum fertilization success in Controls; $\text{FSR}_{50 \text{ Control}}$ = 50% of maximum fertilization success in Controls; $F_{\text{max Control}}$ = sperm concentration that generates maximum fertilization success in Controls; $F_{50 \text{ Control}}$ = sperm concentration that generates 50% of maximum fertilization success in Controls. $\text{FSR}_{50 \text{ pH } 7.8}$ = observed fertilization success in pH 7.8 treatment at the sperm concentration that generates 50% of maximum fertilization success in Controls; $\text{FSR}_{50 \text{ pH } 7.6}$ = observed fertilization success in pH 7.6 at the sperm concentration that generates 50% of maximum fertilization success in Controls. Actual fertilization curves vary for each individual pair.

doi:10.1371/journal.pone.0053118.g001

obtained from the model using the sperm speed and percent motility values observed for each treatment (“ FSR_{mod} ”).

Data Analyses

All percentage data were arcsine transformed prior to statistical analyses and checked for normality [29]. Levene’s test was used to assess for homogeneity of variances among individuals and among treatments. Two-factor Analysis of Variance (ANOVA) was used to assess pH effects on sperm speed and percent motility across all males (pH fixed, male random). Tukey’s test was used to compare *post-hoc* differences among means. The magnitude of responses of percent motility and fertilization success to pH treatments was assessed using logarithmic responses ratios (LnRR ; natural log of treatment response divided by control response [30]). Mean LnRR s and 95% confidence intervals of fertilization success were determined by bootstrapping in R (100,000 iterations; [31]). Regression analysis was used to assess relationships between observed and modelled fertilization outcomes. All statistical tests were carried out using SPSSTM.

Results

Sperm Motility

We analysed over 141,000 sperm tracks (pH 8.1 $n = 43,271$; pH 7.8 $n = 50,135$; pH 7.6 $n = 48,330$).

Acidification significantly decreased the average percentage of motile sperm (by 7% at $\Delta\text{pH}=0.3$ and 9% at $\Delta\text{pH}=0.5$ pH, $P<0.001$; Fig. 2A, Table 2&3A). Responses of individual males differed significantly ($P<0.001$; Table 3A) with reductions in percentage of motile sperm ranging from -6.5% to -15.7% at $\Delta\text{pH}=0.3$, and -9.7% to -17.4% at $\Delta\text{pH}=0.5$ (Fig. 2C).

In contrast, average sperm speed was not significantly affected by acidification ($p=0.710$; Fig. 2B, Table 2&3B), although again there were significant differences in responses between individuals ($P<0.001$; Table 3B). Upper and lower bound 95% CIs for individual response ratios (LnRRs) of sperm speed were equivalent

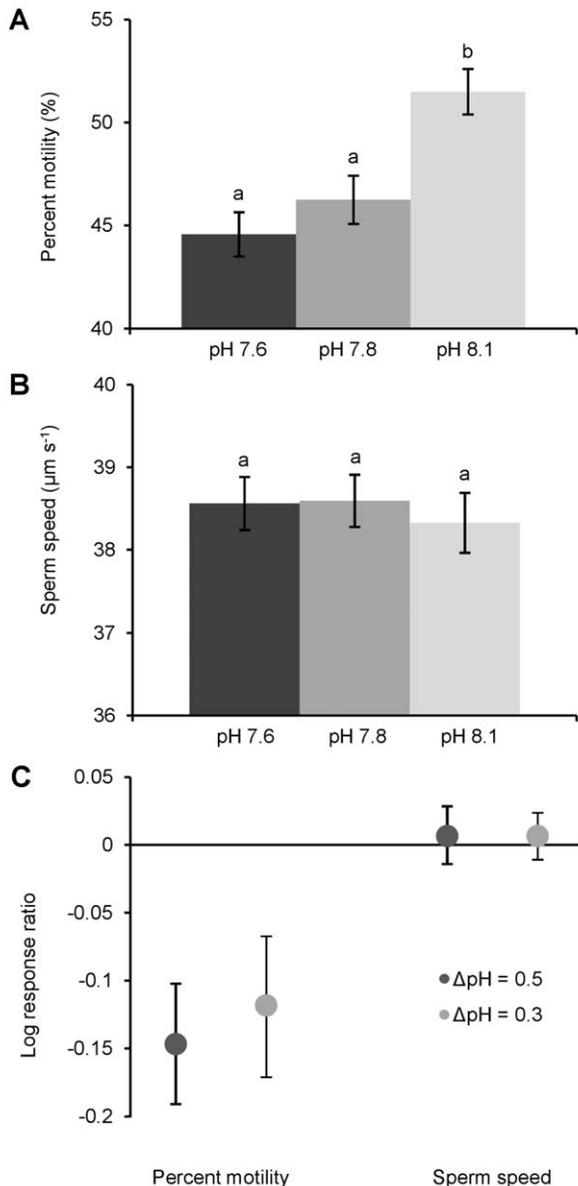


Figure 2. Impacts of ocean acidification on sperm motility and sperm swimming speed in *Heliocidaris erythrogramma*. Proportion of mean (\pm S.E.) motile sperm (A) and sperm speed (B) at different levels of ocean acidification (pH mediated by CO_2 addition). Lower case letters indicate significantly different groups at $p=0.05$ (Tukey's test). (C) Mean logarithmic response ratios ($\pm 95\%$ CI) of effects of ocean acidification on percent motility and sperm speed ($n=19$). doi:10.1371/journal.pone.0053118.g002

Table 3. Two-way ANOVA for percent motility (A) and sperm speed (B) of *Heliocidaris erythrogramma* across different pH treatments (fixed) and males (random).

A Percent motility				
	df	MS	F	P
pH	2	0.270	24.666	<0.001
Male	18	0.531	48.504	<0.001
pH * Male	36	0.011	1.226	0.176
Residual	509	0.009		
B Sperm speed				
	df	MS	F	P
pH	2	4.067	0.346	0.710
Male	18	218.547	18.617	<0.001
pH * Male	36	11.739	0.801	0.791
Residual	509	14.654		

Significant effects ($P\leq 0.05$) are shown in bold.
doi:10.1371/journal.pone.0053118.t003

to $+2.4\%$ to -1.1% ($\Delta\text{pH}=0.3$), and $+2.9\%$ to -1.4% ($\Delta\text{pH}=0.5$; Fig. 2C). There were no significant differences in sperm parameters between males from different sites at any pH level (Table 2).

Fertilization Success

Ocean acidification substantially increased the variance of observed (FSR_{obs} in Fig. 3A) and modelled fertilization success (FSR_{mod} in Fig. 3A).

The overall effect of ΔpH on observed fertilization success (Table 4) was not statistically significant ($P=0.9$, Table 5A), however bootstrapping showed that responses were highly variable between pairwise crosses (Fig. 3B, Table 4). The 95% CIs around the mean log response ratios (LnRR) varied from moderately positive (11% and 14% increases in fertilization success at $\Delta\text{pH}=0.3$ and $\Delta\text{pH}=0.5$, respectively) to strongly negative ($\leq 44\%$ and $\leq 79\%$ decreases at $\Delta\text{pH}=0.3$ and 0.5 , respectively; Fig. 3B). There was no significant difference in maximum fertilization success between pairs from different sites ($P=0.202$).

Modelling the effects of the observed changes in sperm percent motility and sperm swimming speeds on fertilization success yielded predictions that broadly mirrored the patterns seen in observed measurements (FSR_{mod} in Fig. 3B, Table 5B): for $\Delta\text{pH}=0.3$ the modelled LnRR 95% CI ranged from an equivalent of 0 to -34.8% , and for $\Delta\text{pH}=0.5$ from -3.4% to -36.1% . Regression analyses (Fig. 4) revealed that 34% of the observed change in fertilization success could be attributed to changes in sperm motility at $\Delta\text{pH}=0.3$, but only 4% at $\Delta\text{pH}=0.5$.

Discussion

Our finding that the effects of acidification on sperm swimming behaviour (Fig. 2C, Table 2), and fertilization success (Fig. 3, Table 4), vary significantly between individuals is biologically important. Differences in individual responses are the raw material for effective selection, and especially so at the critical life stage of fertilization [2]. However, inter-individual variation has been overlooked in most previous, group-mean based investigations.

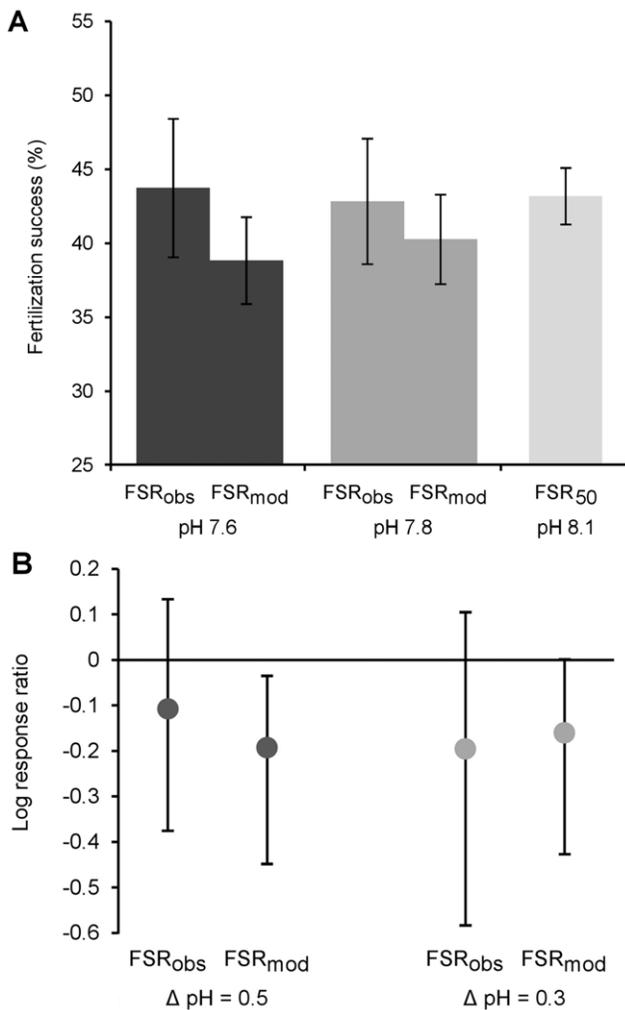


Figure 3. Effects of ocean acidification on fertilization success (FSR) in *H. erythrogramma*. (A) Mean (\pm S.E.) observed (FSR_{obs}) and modelled fertilization success (FSR_{mod}) for pHs 7.6 and 7.8, and mean (\pm S.E.) FSR₅₀ (50% of maximum FSR) for pH 8.1. (B) Bootstrapped mean logarithmic response ratios (\pm 95% CI) of effects of ocean acidification on FSR_{obs} and FSR_{mod}. FSR_{mod} shows change in fertilization success expected due to ocean acidification's influence on sperm swimming behaviour (Fig. 2C). (n = 18 replicate trials). See text for details. doi:10.1371/journal.pone.0053118.g003

The shortcomings of analysing group means are demonstrated by comparing the overall non-significant effect of ocean acidification on mean fertilization (Table 5) with the substantial inter-individual variation in response we observed (FSR_{obs} in Fig. 3B). The majority of individual pairs had reduced fertilization success under acidified conditions, however, some pairs showed increased fertilization success (Fig. 3B, Table 4). This illustrates the importance of examining individual responses: it is individuals that contribute differentially to the next generation – not group means. Consequently, approaches that assess the group mean response ignore evolutionarily important effects of rare individuals that may contribute disproportionately to the next generation. This example also emphasizes the need for adequate sample sizes in order to capture the variety of individual responses to ocean acidification, particularly in species with high inter-individual variation.

The effects of acidification on fertilization success have often been tested using inappropriately high sperm concentrations. Using a sperm concentration that yields maximum fertilization success in controls (F_{max} in Fig. 1) can generate misleading or uninformative results, because assays may be saturated and therefore unresponsive to subtle, but biologically important, differences in fertilization [32], and because such assays cannot detect possible positive effects of the treatment. Had such high concentrations been used in our experiments, the observed increases in fertilization success seen in some pairs would not have been detected (Fig. 3). Choosing 50% of the maximum fertilization success as the response variable (F_{50} in Fig. 1; [28]) allowed us to detect both negative and positive pH impacts on fertilization success, while ensuring maximum sensitivity in our assays.

Previous investigations of the effects of ocean acidification on fertilization success in *H. erythrogramma* have found contradictory results [11,33,34]. These differences may partly be explained by the use of saturated assays in some studies. Previous studies also used smaller sample sizes than used here (5 M \times F pairs [11], 3 replicates [33,34], vs 18 M \times F pairs in this study). The use of gametes mixed from multiple individuals [33,34] also precludes observation of intra-specific variation. Apparently conflicting results may also be explained by the intra-specific variation demonstrated in our experiments. Fertilization success in some pairs was negatively impacted by acidification (confirming [11]) whereas other pairs showed no (or little) net response (confirming [33,34]). Thus, we suggest that much of the controversy around the response of fertilization success in *H. erythrogramma* to ocean acidification can be attributed to a combination of the factors discussed above.

Our observation that lowered seawater pH did not affect the speed of motile sperm (Fig. 2B) but rather decreased the proportion of motile sperm (Fig. 2A) contrasts partly with earlier reports [11]. Sea urchin sperm are stored immotile in an acidic environment inside the testis after development, which inhibits respiration and metabolic processes prior to release [35–39]. Upon spawning, the difference between intra-cellular pH in the testis and extracellular pH in the seawater triggers mitochondrial activity and thus motility. Ocean acidification may reduce the pH gradient upon spawning to a point where it is insufficient to activate the sperm mitochondrion. Since we observed a decrease in the proportion of motile sperm under acidified conditions (Fig. 2A), but the swimming speed of those motile sperm did not decrease (Fig. 2B), this strongly suggests an effect on activation of the sperm mitochondrion but not on mitochondrial function once activated. As sperm have no actively transcribing nuclear genes or biochemistry, the most parsimonious explanation for our results is that genetic variation in mitochondrial membrane protein genes explains some of the observed inter-individual variation in sperm swimming behaviour, and hence fertilization success.

Variation in fertilization success may also have been influenced by parental environmental history, although our results suggest this was not likely to have been a significant driving factor. Individuals were collected from small areas of uniform habitat, and variance within single populations was as large as differences between populations (the converse would be expected if environmental history effects were more influential than genetic diversity). Early cleavage stages (2 h post-fertilization) are also largely independent of transcription of paternal DNA [40]. Consequently we suggest that variation in sperm swimming behaviour and sperm-egg binding compatibility [41,42] are the most likely explanations for the observed variance in responses to acidification.

Table 4. Modelled (FSR_{mod}) and observed (FSR_{obs}) fertilization success for each urchin pair under acidified conditions (pH 7.6 and 7.8), and parameters from Control observations (pH 8.1) used in modelling FSR_{mod} at lowered pH levels.

pH	7.6		7.8		8.1	
	FSR _{mod} (%)	FSR _{obs} (%)	FSR _{mod} (%)	FSR _{obs} (%)	FSR _{50 Control} (%)	F _{50 Control} (sperm μl ⁻¹)
A	no data					
B	26.46	63.92	25.33	50.38	28.95	1258.93
C	36.21	47.93	55.81	47.19	47.76	1584.89
D	23.25	10.46	24.45	12.95	24.94	4466.84
E	41.34	45.80	43.21	62.59	37.03	3162.28
F	51.59	24.21	50.45	37.27	49.74	7943.28
G	41.55	44.83	42.74	46.93	48.98	2511.89
H	38.78	61.78	39.09	51.80	47.37	22387.21
I	41.52	9.05	43.45	7.11	42.03	251.19
J	3.74	30.15	3.50	2.47	31.94	398.11
K	38.95	71.08	41.06	53.18	48.64	22387.21
L	40.78	64.17	41.26	60.09	48.06	10000.00
M	28.60	40.79	30.54	35.49	34.61	2511.89
N	52.12	47.55	53.29	53.32	48.81	3981.07
O	35.89	43.26	37.83	43.39	42.95	10000.00
P	43.72	28.96	41.66	52.20	49.88	50118.72
Q	49.49	22.29	49.71	41.97	48.26	8912.51
R	50.57	54.19	47.54	63.00	46.58	15848.93
S	54.44	76.84	53.87	49.54	50.77	2238.72
mean	38.83	43.74	40.27	42.83	43.18	9442.43
S.E.	2.93	4.68	3.02	4.24	1.91	2898.53

FSR_{50 Control} = 50% of maximum fertilization success in Controls; F_{50 Control} = sperm concentration that generates 50% of maximum fertilization success in Controls. Sperm data from each male in Table 2 were used in modelling FSR_{mod}. No females were spawned for male A. doi:10.1371/journal.pone.0053118.t004

More broadly, for free spawners such as *H. erythrogramma* the chance of successful fertilization depends on gamete concentra-

tions, gamete life span and (hence) on the distance between spawning individuals [14,43]. The density and abundance of *H. erythrogramma* varies substantially *in situ* – from <1 to >50 individuals.m⁻² [44,45], occurring in a range of even to patchy distributions. Low fertilization success due to sperm limitation may be common between widespread clusters of urchins because sperm longevity is short [46–48]. If many individuals within populations display decreased sperm motility due to ocean acidification, sperm limitation downstream of a spawning site may be exacerbated [49,50], further decreasing reproductive success and ultimately reducing the number of individuals that contribute to future generations [15,51].

Our results support the concept of ‘winners’ and ‘losers’ of climate change; a concept often proposed over the last decade [52–55], and applied to many organisational levels such as alleles within genes, individuals within populations, and species within ecosystems. Here we apply the ‘winners’ and ‘losers’ concept at the individual level. The substantial variation in sperm motility and fertilization success we observed in response to predicted ocean acidification (Figs 2, 3) shows that some individuals (here, males or male x female pairs, see above) are better equipped than others to cope with acidification. Future ocean acidification will likely reduce the proportion of fertilizations by acidification-sensitive gametes (‘losers’), and increase the proportion of fertilizations by acidification-resistant gametes (‘winners’). Whether these ‘winners’ are selected due to genetic traits (such as sperm swimming ability (Fig. 2A) and sperm-egg binding compatibilities [41,42]) or due to non-genetic maternal traits (such as egg condition [56]) will be

Table 5. Two-way ANOVA for observed (FSR_{obs}; A) and modelled fertilization success (FSR_{mod}; B) of *Heliocidaris erythrogramma* across different pH treatments (fixed) and males (random).

A FSR _{obs}				
	df	MS	F	P
pH	2	0.002	0.103	0.902
Male	17	0.065	3.711	0.001
pH * Male	34	0.018	18.794	0.181
Residual	1	0.001		
B FSR _{mod}				
	df	MS	F	P
pH	2	0.013	3.097	0.058
Male	17	0.045	11.005	<0.001
pH * Male	34	0.004	4.308	0.367
Residual	1	0.001		

Significant effects (P≤0.05) are shown in bold. doi:10.1371/journal.pone.0053118.t005

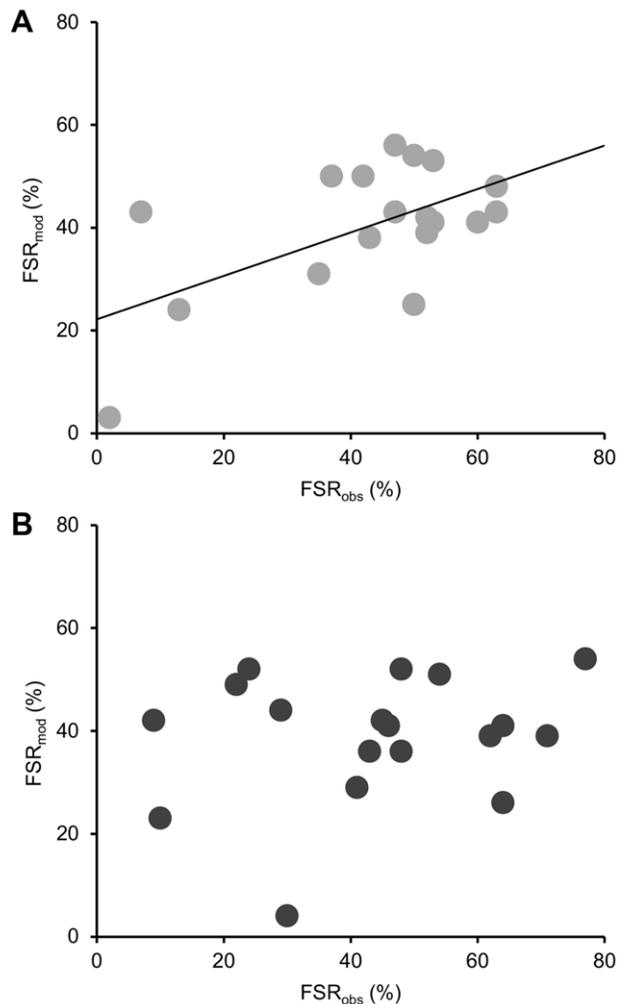


Figure 4. Scatterplots for observed (FSR_{obs}) versus modelled (FSR_{mod}) fertilization success for pH 7.8 (A) and 7.6 (B). Regression analyses revealed a significant relationship between observed (independent) and modelled fertilization (dependent) for pH 7.8 ($P=0.012$, $r^2=0.336$), but not for pH 7.6 ($P=0.413$, $r^2=0.042$). doi:10.1371/journal.pone.0053118.g004

immaterial to the selection of the ‘winners’ *per se*. It also remains unclear whether ‘winners’ of climate change-induced selection for

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fertilization success will actually remain ‘winners’ across the entire life-cycle. Nonetheless, it should be remembered that the heritability of the selected traits will of course strongly influence the future adaptation potential of the progeny and hence the long-term adaptive benefits of ‘winner’ status.

If this observed variability is heritable, it will have important implications for urchin populations in a near-future acidified ocean. Loss of less competitive gamete genotypes will reduce overall genetic diversity, at least initially. This loss may be counteracted by genetic drift and new mutations, although in the short term these are likely to be trivial compared to the effects of selection. The long-term fitness consequences, however, will depend on the fitness benefits of traits that covary with acidification-resistance in gametes, and the extent to which recombination in second and subsequent generations gather advantageous alleles in some individuals. Any selection that reduces available genetic diversity leaves future populations less capable of tolerating further perturbations [57,58], and this underscores the need for analyses of the fitness of the descendants resulting from experiments such as those we have described here.

In summary, despite an increasing number of studies focusing on the effects of ocean acidification on the early life history of marine organisms [59,60], very few studies have investigated individual-level responses to changing oceanic conditions. This is perplexing in light of the growing evidence for a degree of inter-individual variation that exceeds the noise of baseline variability [24]. Conducting adequately replicated studies to investigate inter-individual variability in response to marine climate change is imperative if we are to understand the capacity for selection and adaptation of marine species.

Acknowledgments

We thank Laura Elia and Larissa Trompf for assisting with the experiments, Christopher McRae for conducting the alkalinity analyses, and Andrew Beattie for insightful discussions regarding this research. The reviewers are thanked for helpful comments on the manuscript. This work was partly performed within the Linnaeus Centre for Marine Evolutionary Biology (<http://www.cemeb.science.gu.se>). Animals were collected under scientific collection licences from Primary Industries, Industry and Investment, New South Wales.

Author Contributions

Conceived and designed the experiments: PS JH JW. Performed the experiments: PS JH JW. Analyzed the data: PS JH. Contributed reagents/materials/analysis tools: PS JH JW. Wrote the paper: PS JH JW MG.

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