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Marine A ac It re in a Changing Environment

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Cover photo of historic canned oyster collection at Copps Island Oysters, LCC, in Norwalk, Connecticut. Photo credit: NOAA Fisheries

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Intro ction

The impacts and interactions between environmental change and aquaculture are diverse and many are likely unknown. What we do know is that nutrient pollution is driving eutrophication and dead zones, ocean acidification is changing water chemistry and climate change is already influencing our food supply, fresh water availability, weather and way of life. Aquaculture will be impacted by, and can also impact, these environmental changes over various scales. Aquaculture of finfish, shellfish and seaweed have different threats, benefits and opportunities related to environmental change, but in the right location, and with the right management they can be mostly beneficial to the environment. Because of the significant aquaculture industry impacts caused by environmental change, and the potential for aquaculture to provide some mitigation, research on this topic was a priority shared by the Fisheries Research and ducation Agency (FR A) of apan, the U. S. National Oceanic and Atmospheric Administration (NOAA), and the U.S. Department of Agriculture. These Proceedings represent the second of three consecutive years of meetings where the theme is Aquaculture in a Changing nvironment. This year, the research focused on the impacts to aquaculture production due to environmental change (e.g., ocean acidification impacts on shellfish aquaculture) and science to mitigate these impacts (counter measures). Proceedings from the previous Symposium were published by FR A (4 th U NR), where the focus was on the potential of aquaculture to mitigate impacts of environmental change (e.g., sequestering carbon, bioextraction of nutrients and CO₂, antacid, oxygen production). Looking forward, we will seek to explore science and technology that address these issues and will enable aquaculture to prosper in a changing environment and be a part of the solution to reducing the human impacts of environmental change.



NOAA Fisheries Office of Aquaculture

United States Panel Chair



The infl ences of Environmental Changes on a anese Nori Maric It re

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: Pyropia tenuipedalis, breeding, cell selection, low salinity tolerance

ABSTRACT

Nori mariculture has a long history of over one hundred years and is one of the most important fisheries industries in apan. Recently, nori production has been decreasing due to environmental changes, such as seawater temperature rise and low nutrient levels. As for the seawater temperature rise, the start of nori mariculture is delayed because of high water temperature staying above 23 C in autumn, thereby shortening the season of nori mariculture. Spores start to be released when the water temperature decreases below 23 C. Low nutrient levels in ambient waters cause bleaching of nori thalli, reducing the quality and price of nori. Currently, new culture species and strains of nori around the coasts of apan are being explored. The thallus has a simple structure and cells within the thalli sometimes have mutations. Mutant cells may have the potential for tolerance to high water temperature and low nutrient level, as well as other favorable characteristics. More recently, shortening of the thalli is observed frequently. One of the causes is predation by herbivorous fishes (e.g. Acanthopagrus schlegelii, Mugil cephalus cephalus) and ducks (Anas penelope, Anas acuta). In apan, it has been reported that herbivorous fishes cause the disappearance of marine forests, which is referred to as iso-vake. The duration of iso-yake caused by herbivorous fish has been prolonged because of increased water temperatures. While the activities of the iso-yake causing fish are reduced below 17 C, the herbivorous fishes eating

nori are active at low water temperatures. ven if some measures are taken to control herbivorous fish, birds would alternatively eat nori. It is urgently necessary to develop effective measures to control these herbivores.

INTRODUCTION



Figure 1. Photograph showing nori mariculture in Fu uo a Prefecture.

Nori mariculture has a long history and many farmers and researchers have developed innovative technologies which made nori mariculture one of the most important fisheries industries in apan (Fig. 1). However, nori production has decreased due to environmental changes, such as seawater temperature rise and low nutrient levels. At present, nori mariculture of *Pyropia ye oensis* is mainly used and *P. tenera* is used in limited areas*. Culture strains have been repeatedly selected within *P. ye oensis* for a long time and, as a result, the strains

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have lost genetic diversity. Dissolved inorganic nitrogen (DIN) levels started to decrease (i.e. oligotrophication) in the 1 80s in the western part of apan, and in the 1 0s in the eastern and central parts of the Seto Inland Sea (Abo and Yamamoto, 201). After the late 1 0s, disappearance of macroalgal beds, generally called iso-yake in apanese, has spread along the coast of western apan because of increasing sea water temperature (Yoshida et al., 201). Nori production had the highest value in 2001 and has been decreasing until present. However, in the nori mariculture industry, it was not until recent years, when the production fell below the level of demand of 8 billion sheets, that the effects of oligotrophication and water temperature rise became an issue. Therefore, many researchers are searching earnestly for measures to address reduced nori mariculture production. In the present paper, we introduce the history, problems and solutions of nori mariculture.

Bac gro n on nori maric It re in a an

In apan, the use of nori has a long history of over two hundred years. In the Bunka-Bunsei period (1804 1830), atsukawa Shunsen designed the ukiyo-e print of nori making (Miyata and Tomizuka, 7). People during this era harvested nori growing in the wild. The procedure for nori making then is similar to that in the present, such as chopping, forming and drying, except that everything was done by hand. It is thought that sights of nori-making were common scenery in the do period (1603 1867). In the early 1 00's, nori mariculture began and was cultured on sodahibi, which are made of tree branches and set around the coast (Okamura, 1 0). However, at that time, nobody knew where the spores came from, and nori was harvested, chopped and formed into sheets by hand like in the do period. In recent years, nori mariculture has changed. Dr. athleen Mary Drew Baker's discovery of the microscopic conchocelis phase within the life cycle of nori (Drew 1 4) made it possible for apanese researchers to establish methods of maintaining different strains of nori. Nori mariculture in the present uses nets instead of sodahibi and is operated on larger

scales using machinery for harvesting, chopping and drying of the nori. Nori mariculture has become one of the most important fisheries industries in apan with many innovations.



Figure 2. Photographs showing well grown nori thalli (a), bleached thalli (b) and products of well grown and bleached thalli (c). Red arrow shows product of bleached thalli.

ro lems of nori maric It re

Nori mariculture is one of the largest fisheries industries in apan. However, nori production has decreased for the past decade from 10 billion sheets in the 1 0's to below 8 billion sheets in recent years. Nori production further decreased to 6.3 billion sheets in 2018. The causes of the decrease in nori production are mainly seawater temperature rise and low nutrient levels.

Nori mariculture starts when the seawater temperature decreases below 23 C. The start of nori mariculture is delayed by increasing water temperatures, staying above 23 C in autumn, due to global warming. Nori mariculture used to begin in the first quarter of October in the 1 80's, but began in the second to the last quarter of October in the 2010's. On the other hand, the timing of spring seawater temperature rise is unchanged, which results in a shortening of the culture period for nori. In apan, the oligotrophication of coastal waters due to the regulation of wastewater discharge is becoming problematic. Low nutrient levels in the

ambient water cause bleaching of nori thalli, reducing the quality and price of nori. Generally, nori thalli grow with a dark black color (Fig. 2-a, c), but in an oligotrophic environment, thalli grow with a light brown color (Fig. 2-b, c). Bleached thalli are 2 to 3 PY cheaper per sheet than dark black thalli.

otential Sol tions to the ro lems

Many researchers are trying to develop new breeding strains of nori with tolerances to environmental changes, such as high water temperature and low nutrient levels. In addition to



Figure . Photograph showing e perimental mariculture of new culture species Pyropia tenuipedalis.

conventional selective breeding, cell selection within thalli and cell fusion using protoplasts and mutation breeding with heavy ion beams are also carried out at several institutes (Sakaguchi 2011; Niwa et al., 2011). In China, pigment mutants using ⁶⁰Co- ray irradiation were isolated (Yan and Aruga, 2000). Field use of the strains obtained by cell fusion and mutation breeding is not yet permitted in apan. However, strains obtained by cell selection within thalli have begun to be used in field culture (Sakaguchi, 2011; Abe et al., 201). It is thought that cell selection within thalli is a useful technique for development of new strains and its use in the field has no legal restrictions so far. In addition, many researchers are exploring alternative culture species in the genus Pyropia. Strains of P. ye oensis

used for mariculture have been screened by selective breeding for over 40 years, making genetic diversities of the strains poor. Moreover, breeding methods such as cell fusion and mutation mentioned above require special machines and techniques. Bangiaceae including the genus *Pyropia* contains 30 species, and many of them are used as local food resources. However, the distribution of nori species in many local areas is unclear. Different nori species growing in the wild have a wide variety of characteristics and they may be candidates for new culture species suited for different environmental conditions. In fact, *P. tenuipedalis* has been cultured for a few years on an experimental scale (Fig. 3).

Countermeasures for oligotrophication, fertilization, optimal management of wastewater, and plowing of the sea bottom have been attempted in several water bodies. Since nutrients are used not only by nori but also by phytoplankton, which could cause adverse effects to the environment, these methods must always be accompanied by a survey of the occurrence of phytoplankton (Mine, 201). Moreover, development of polyculture techniques combining bivalve aquaculture (e.g. oyster and clams) with nori mariculture has been attempted. Bivalves feed on phytoplankton, which compete with nori for nutrients, and their excreta is supplied to nori as nutrients. Although these attempts are in the trial phase, environmental improvement utilizing biological functions is promising.

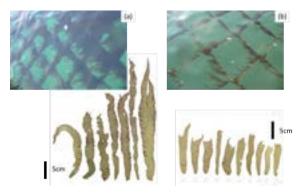


Figure . Photographs showing nori culture net with well grown thalli and the specimens (a) and culture net with shortened thalli and the specimens (b).

An emerging ro lem

In recent years, many incidents of shortening of nori thalli have been observed in apan (Fig. 4). Until now, it was thought that one of the ma or causes of the shortening was low salinity of ambient water, based on the symptom of the cell damage (irita and Matsui, 1 3). However, occurrences of the shortening have also been observed without low salinity conditions. Field observations at nori mariculture sites using video and time-lapse cameras have revealed another cause of the shortening: predation by herbivorous fishes (e.g. Acanthopagrus schlegelii and Mugil cephalus cephalus) and ducks (Anas penelope and Anas acuta). In apan, it has been reported that herbivorous fishes (e.g. Calotomus japonicus, Siganus fuscescens and Kyphosus bigibbus) cause the disappearance of marine forests, which is referred to as iso-yake. The duration of iso-yake caused by herbivorous fish has been getting longer because of seawater temperature rise because of global warming. The activities of fish causing iso-yake are reduced below 17 C, allowing macroalgae to thrive in the winter. However, the fishes and ducks causing the shortening of nori can actively graze nori even below 17 C. The seawater temperature typically ranges from 10 C to 20 C during nori culture season, and grazing pressure can cause an enormous loss of nori thalli. It is necessary to develop effective measures to lessen predation damage by these herbivorous fishes and ducks immediately.

* Pyropia tenuipedalis, P. ye oensis and P. tenera in this paper were replaced to eopyropia tenuipedalis, . ye oensis and . tenera by Yang et al. (2020), respectively.

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In this paper, the authors developed a PCR-RFLP method useful for discrimination of 16 apanese and two non- apanese Porphyra and Pyropia species. These species have simple morphological features, making morphological species identification difficult. The two primer sets on the mitochondrial DNA used in this study were able to amplify single fragments with PCR (partial mitochondrial DNA related to ATP6 gene and trnC rns). All 18 species were successfully distinguished using a combination of five restriction enzymes (Ta I, SspI, AciI, Cfr13I and AluI). It was concluded that PCR-RFLP analysis is a useful tool for discrimination of wild strains of Porphyra and Pyropia species for potential use in mariculture.

Nakayama, T., M. Abe, N. Murase, and Y. Shikano. 2017. Influence of salinity on growth of red alga *Pyropia tenuipedalis* and *Pyropia ye oensis* foliose thallus. Aquacult. Sci. 6: 321-330.

Pyropia tenuipedalis is a new culture species of nori

in Yamaguchi Prefecture, apan. The habitat of this species is more brackish as compared with that of *P. ye oensis*, which is the common species in apanese nori mariculture. In this paper, the authors investigated the relationship between salinity and growth of *P. tenuipedalis* and *P. ye oensis*. It was revealed that *P. tenuipedalis* tolerates lower salinity in comparison with *P. ye oensis*.

Abe, M., Y. Fu ita, M. obayashi, . Fu iyoshi, M. Tamaki, Y. Fukui, M. Satomi, and N. Murase. 201 . ffects of antibiotics on survival and growth for *Pyropia ye oensis* protoplasts. Aquacult. Sci. 63: 1-8.

The nori thallus sometimes has cell mutations. For the breeding of *Pyropia* species, it was necessary to develop an axenic cell culture method. In this paper, we investigated a way of isolating axenic protoplasts of P. ye oensis and the effects of twelve antibiotics on the survival and growth of the protoplasts. The axenic protoplasts were able to be isolated by a series of treatments with sodium citrate seawater. The growth rates of the protoplasts treated with FRM and ABPC were 1.1 2.7 times higher than that of the control, while PCG and SM suppressed growth. The authors succeeded in developing an axenic culture method for Pyropia species, but the survival, growth and morphogenesis of the cells in the axenic culture varied. Improvement of the methodological stability is necessary.

Yang L.- ., Y.-Y. Deng, G.-P. u, S. Russel, .- . Lu, and . Brodie. 2020. Redefining *Pyropia* (Bangiales, Thodophyta): four new genera, resurrection of *Porphyrella* and description of *Calidia pseudolobata* sp. nov. from China. . Phycol., 6: 862-87 .

Develo ment of Environmentall rien I Cost Effective an N tritionall Balance Alternative rotein Base Diets for igh all e Marine ish C It re in Recirc lating A ac It re S stems

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: Alternative protein sources, fish meal replacement, black sea bass, southern flounder, and cost-effective aquafeeds

ABSTRACT

With very little domestically grown seafood, along with increasing exploitation of wild stocks, the US annual seafood trade deficit now exceeds 1 billion. conomics, environmental issues and fish feed are factors vital to the sustainability of the aquaculture industry in the US and apan, and these factors increase the pressure on aquaculturists to develop eco-friendly and cost-effective aquaculture practices. One of the challenges faced by the aquaculture industry is to prevent the pollution of water by artificial feed. Fish meal is the main protein source used in aquafeed and is made from small pelagic species of forage fish. Two nutrients in fish meal, nitrogen and phosphorus, have a great impact on the environment, and their release can result in eutrophication. Fish meal could also contain polychlorinated biphenyls (PCBs), dioxins, and other harmful chemicals, and become a vector of contamination in farm-raised finfish. Alternate protein sources such as terrestrial animal and plant protein sources can reduce the amount of wild fish used as protein, and potentially reduce the nutrient levels in effluent waste. One of the goals of our research at the University of North Carolina Wilmington Center for Marine Science is to develop sustainable marine finfish feed with reduced inclusion of fish meal in recirculating aquaculture systems, which are widely considered to be an

environmentally friendly technology for producing farmed fish.

A series of experiments were conducted to test the effects of different dietary levels of soybean meal, ultra-low gossypol-based glandless cottonseed meal, and poultry meal in the diet of southern flounder and black sea bass reared in a recirculating aquaculture system for 8-10 weeks. Growth performance, feed utilization, digestibility, and the biochemical composition of fish tissues were evaluated.

Results suggest that about 68 and 3 % fish meal protein could be replaced by soybean meal protein and 82 and 100% of fish meal protein could be replaced by poultry meal protein in the diet of black sea bass and southern flounder, respectively. Results also suggest that fish meal protein could be replaced up to 100 and 7 % by glandless cottonseed meal protein in black sea bass and southern flounder diets, respectively. These findings reveal species-specific differences in effective substitution levels of alternative protein sources to fish meal protein, and provide a basis for developing environmentally- sound and cost-effective predominantly terrestrial plant and animal protein-based diets for high value marine fish culture in the USA and apan.

INTRODUCTION

In terms of dietary composition, protein is the single largest and most expensive component in fish feed. In aquaculture feed, fish meal is a popular source of high-quality protein and highly digestible essential amino and fatty acids (Cho and im 2011). By replacing fish meal in aquaculture diets, alternate terrestrial plant and animal protein sources can lower feed costs, reduce the amount of wild fish used as protein, and potentially reduce the nutrient levels in effluent waste (Trushenski et al. 2006). However, for most species, there is a limit to how much fish meal can be replaced by alternative protein sources without negatively affecting fish growth and feed efficiency (Gatlin et al. 2007).

So ean meal

Soybean meal is considered to be one of the most suitable ingredients for replacing fish meal in commercial fish feed. Compared with other plant protein sources, soybean meal has high protein content, very low carbohydrate and fiber, high digestibility, and a good amino acid profile (Hardy 2006; Gatlin et al. 2007). Soybean meal is used in fish feed not only because of its high protein content, but also owing to its worldwide availability.

o goss ol cottonsee meal

Cottonseed meal (CSM) has received limited attention as a potential fish meal replacement in agua feeds; however, CSM protein is an increasingly attractive alternative to fish meal protein due to improved plant production and processing methods. Due to the large global production of cotton, as well as the large quantities of cottonseed by-products, CSM is much cheaper per unit of protein than fish meal and other fish meal replacements. An important constraint to the use of CSM in animal feeds is gossypol, a terpene-based secondary metabolite that has an important role in the cotton plant's defense mechanisms against pests and possibly diseases (Romano and Scheffler, 2008). The utilization of CSM as an ingredient in the feeds of animals and

fish has been improved by reducing or eliminating gossypol through traditional genetic selection or through genetic modification (GMO). Regular high-gossypol CSM (R-CSM) protein has been used to replace fish meal protein at maximum levels of 3 % in grass carp (heng et al., 2012) and 30% in parrotfish feeds (Lim and Lee, 200). However, research on fish meal replacement by low-gossypol based CSM is very limited on marine fish species.

o ltr ro ct meal

Terrestrial animal protein sources have several advantages, including a similar amino acid profile to fish meal, availability, and relatively low cost. Poultry by-product meal (PBM) is a protein source produced from waste and by-products of processed chickens, possibly including heads and feet, but excluding feathers and intestines. Like other animal-based protein feedstuffs, it has a high protein content, but can vary in compositional quality and lacks certain essential amino acids (Tacon et al. 2006). PBM has been used successfully to replace fish meal at high levels of dietary inclusion for a number of finfish species. In gilthead seabream, 0% of the fish meal protein was successfully replaced with PBM protein without a reduction in growth (Nengas et al. 1 uvenile red drum, 67% of the fish meal protein was replaced by PBM protein with no reduction in growth (ureshy et al. 2000).

Blac sea ass

Black sea bass *Centropristis striata* is a commercially important marine finfish species that commands a high market price. This species inhabits the coastal waters of the eastern USA, from the Gulf of Maine to Florida. The abundance of black sea bass along the U.S. ast Coast has been declining since the 1—0s (NOAA 2012; ASMFC 2016). Potential for limited market supplies and for higher prices of ocean caught black sea bass in the future are important economic incentives to investigate the feasibility of black sea bass production via aquaculture to help meet market demand (Watanabe 2011; Watanabe et al. 2016). Hatchery-raised black sea bass uveniles originating

from captive wild- caught broodstock are routinely cultured through the market stage in a recirculating aquaculture system at the University of North Carolina Wilmington (UNCW) Aquaculture Facility (Watanabe et al. 2003; Watanabe 2011; Watanabe et al. 2021).

So thern lo n er

The southern flounder Paralichthys lethostigma is a flatfish in the family Paralichthyidae. It can be found in coastal waters from Albemarle Sound, North Carolina, through the South Atlantic states with the exception of South Florida. Southern flounder landings have declined, leading to more stringent fishery regulations and interest in culturing native flatfishes for stock enhancement or food fish production. The development of intensive culture methods for southern flounder in the southeastern U.S. is of great interest because of its euryhaline character and its status as a highly desirable food and recreational species and potential for commercial culture. Hatchery methodology for spawning and larval rearing is well investigated (Daniels and Watanabe, 2003; Watanabe et al., 2006; Wright- Moore et al., 201).

MATERIA S AND MET ODS

E erimental animals an s stem

uvenile black sea bass and southern flounder were cultured from eggs supplied by photothermally conditioned captive broodstock held at the UNCW Center for Marine Science, Aquaculture Facility (Wrightsville Beach, NC). Adults were induced to spawn using luteinizing hormone-releasing hormone analogue (LHRHa) implants (Watanabe et al., 2003). ggs were hatched and reared through uvenile stages in 1 0 L tanks using protocols established at UNCW (Watanabe, 2011; Carrier et al. 2011, Russo et al. 2017). All the experiments were conducted in a controlled environment laboratory consisting of twenty-four 7 -L rectangular (76 x 32 x 43 cm) glass tanks, supported by a recirculating aquaculture system.

Table 1. E perimental designs.

xperimental fish	Tested alternatives	Replacement levels	References
Black sea bass	soybean meal	0-100%	Alam et al. 2012
Southern flounder	soybean meal	0-100%	Alam et al. 2011
Black sea bass	low- gossypol cottonseed meal	0-100%	Anderson et al. 2016
Southern flounder	low- gossypol cottonseed meal	0-100%	Alam et al. 2018
Black sea bass	poultry by product meal	0-100%	Dawson et al. 2018
Southern flounder	Poultry by product meal	0-100%	Dawson, 2012

E erimental iets an esigns

A series of feeding trials were designed to test the substitution limits of soybean meal, CSM and PBM protein in replacement of menhaden fish meal protein (FMP) in the diet of uvenile black sea bass and southern flounder (Table 1) (Alam et al. 2011, 2012, 2018, Dawson, 2012, Dawson et al. 2018, Anderson et al. 2018).

St ies ith so ean meal

Three experiments were conducted to determine the extent to which FMP can be replaced by solvent-extracted soybean meal protein (SBP) in the diet of uvenile black sea bass. In all experiments, diets were formulated to replace FMP by SBP from 0 to 100% and supplemented with squid meal, krill meal, and attractants (Alam et al. 2012). For southern flounder, two feeding experiments were conducted replacing FMP with SBP at 0, 10, 20, 30, 40, 0, 60 and 70% with or without supplemental L-methionine and L-lysine (Alam et al. 2011).

St ies ith lo goss ol CSM

ight diets were formulated for uvenile black sea bass to replace FMP by three CSM proteins: a CSM prepared from glandless cottonseed meal (GCSM, 0.4% crude protein, CP), a CSM that had been solvent extracted with acidic ethanol to remove the gossypol (SCSM, 3.8% CP), and a CSM prepared from regular (glanded) cottonseed (RCSM, 4 % CP). Three diets replaced 0, 7 and 100% of FMP with GCSM, and three diets replaced 0, 7 and 100% of FMP with SCSM. One diet replaced 100% FMP with RCSM protein (Anderson et al. 2016). For southern flounder, CSM proteins from GCSM, genetically-modified low-gossypol seed (GMO-CSM, 6.0% CP) and R-CSM were evaluated to replace FMP. Six diets replaced 0, 7 and 100% FM protein with GCSM or GMO-CSM protein (Alam et al. 2018).

St ies ith BM

Two feeding trials were conducted to determine the maximum substitution limits of poultry by-product meal (PBM; 66% CP) protein for fish meal (Fish meal; % CP) protein in the diet of uvenile black sea bass and southern flounder. ight diets were formulated for each experiment to replace FMP with feed-grade PBM protein at 0 (control), 40%, 0%, 60%, 70%, 80%, 0%, and 100% in black sea bass and southern flounder diets (Dawson et al. 2018, 2012).

All experimental test diets were prepared at the UNCW Aquaculture Facility as described by Alam et al. (2011, 2012, 2018) and stored in a freezer (-20 °C). All diets were isonitrogenous and isolipidic for each experiment. A control diet (0% SBM, PBM and CSM) was formulated with high FMP and other practical protein sources. L-methionine and L-lysine were supplemented to the diets to equal the control diet.

ee ing trials

xperimental diets were fed to triplicate groups of fish twice a day (0 00 and 1 00 h) to apparent satiation (i.e., as much as they can consume with minimal wastage) for six to ten weeks. Fish in

individual tanks were lot weighed every two weeks to monitor weight gain throughout the experiment. Mortalities were also recorded for each tank throughout the experiment. Water quality was monitored throughout the experiment on a weekly basis and maintained at optimum levels (Watanabe et al. 2011). Temperature, salinity, dissolved oxygen, and pH were monitored using a multiparameter probe (YSI, Yellow Springs, Ohio, USA). Ammonia and nitrite were monitored using a HACH spectrophotometer (DR 2010, Loveland, Colorado, USA).

Biochemical an statistical anal sis

At the end of the feeding trial, five fish from each tank were collected for biochemical analysis to determine proximate composition (moisture, ash, lipid, and protein-AOAC, 2000) and fatty acid profiles of the whole bodies of the fish. Fatty acid analysis was conducted in triplicate as described in Dawson et al. (2018) and Alam et al. (2012). Total amino acid analysis of diets was conducted in duplicate (AAA Laboratories, Damascus, Oregon, USA). At the conclusion of the experiment, fish were reared for an additional 14 days on their treatment diets to determine the apparent digestibility coefficient (ADC) of crude protein in the experimental diets using 0. % chromic oxide as inert marker. Chromic oxide content was determined using a spectrophotometer through a modified Furukawa and Tsukahara (1 66) method. Statistical analysis was performed using the MP 7 statistical software (SAS Institute Inc., ver. 13, Cary, North Carolina).

RESU TS AND DISCUSSION

E eriments ith so ean meal rotein SB

No significant differences in body weight gain, feed efficiency, and survival were observed among fish fed diets replacing FMP with SBP at 0 to 70% (Alam et al. 2012). In addition, body weight gain was not significantly different in fish fed supplemental methionine and lysine in 70% SBP diets compared to fish fed 0% SBP diets. No

significant differences were observed in whole-body n-3 polyunsaturated fatty acids among the treatments. Broken-line regression of the specific growth rate data suggested that the maximum level of FMP replacement with SBP in black sea bass diets was 67.6 68.4% with 7. %

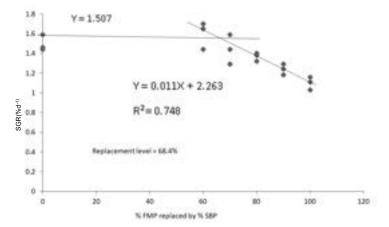


Figure 1. ro en line regression analysis between specific growth rate (SGR) and FMP replaced by S P with supplemental s uid meal, rill meal and attractants in the diets of juvenile blac sea bass (Alam et al. 2012).

squid meal and % krill meal in the diet (Fig. 1) and 7.2 8.0% without squid and krill meal supplementation (Alam et al. 2012). In black sea bass diets supplemented with squid meal, krill meal, and attractants, the maximum replacement level of FMP with SBP (67.6 68.4%) was higher than reported for other marine finfish species such as apanese flounder (4 %, ikuchi 1); gilthead seabream (4 %, Martinez-llorens et al. 2008); cobia (0%, hou et al. 200), and Atlantic cod (0%, hou et al. 200)Walker et al. 2010), but lower than that found for freshwater fishes such as common carp (100%, iola et al. 1 82) and Nile tilapia (100%, Deyab et al. 2002). In general, the fatty acid composition of the fish whole body reflected the composition of their diets. Linoleic acid (18:2 n-6) was higher in the SBP diets as well as in the whole body of fish fed those diets. Studies on yellow perch (asper et al. 2007), pike perch (Schultz et al. 200), and apanese flounder (Lee et al. 2000) also found higher levels of linoleic acid in fish fed diets containing SBP or soybean oil. As SBP content was increased in the diets, fish oil was increased to compensate for the low lipid content of SBP,

leading to higher n-3 PUFA levels in the SBP diets. All three feeding trials showed that black sea bass uveniles are able to utilize high levels of soybean meal in their diet without reducing growth performance.

In southern flounder, no significant differences were observed in body weight gain, specific growth rate, feed intake, feed conversion and protein efficiency ratios, and the whole-body proximate composition among fish fed diets replacing 0 40% of FMP with SBP (Alam et al. 2011). When compared with fish fed 0% SBP, weight gain decreased significantly for the fish fed more than 40% SBP. Broken-line regression showed that the optimum levels of FMP replacement with SBP in the diet of southern flounder without and with supplemental methionine and lysine were 3 .1% and 38. %, respectively, when diets contained % squid meal and % krill meal. Maximum replacement level of FMP with SBP without supplementing amino acids for southern flounder (40%) in this study was similar to what was found in cobia (40%; hou et al. 200) and slightly lower than reported in the gilthead sea bream (4 %; Martinez-Llorens et al. 2008). In both experiments (with or without amino acids). whole-body moisture, protein, and lipid contents were not affected by SBP replacement of FMP up to 40%, similar to what was reported in yellow croaker (Ai et al. 2006), rainbow trout (Bureau et al. 2000), and Indian ma or carp (han et al. 2003).

E eriments ith glan less cottonsee meal CSM

Growth performance of black sea bass uveniles fed diets replacing between 0 and 100% of FMP protein with low-gossypol GCSM, or replacing 0 to 7 % of FMP with low-gossypol SCSM protein was not impaired compared to fish fed a FMP-based control palatability of the RCSM diet was attributable to the anti-nutrient compound gossypol. No significant differences in survival, feed conversion ratio (FCR), protein efficiency ratio (P R), or whole body protein or lipid composition were observed among the fish fed the low-gossypol diets. Gossypol (2 . mg kg) was only detectable in

the livers of fish fed the high-gossypol RCSM diet. Fish whole body essential amino acid compositions did not differ significantly among treatments. Whole body n-3 PUFAs decreased, while n - 6PUFAs increased with increasing CSM protein in the diets. The apparent digestibility coefficient of protein was high (83.1 87.1%) for all treatments (Anderson et al. 2016). For uvenile black sea bass, 7 % FMP in the diet can be replaced with low-gossypol CSM protein prepared by solvent-extraction, and 100% of FM can be replaced with low-gossypol CSM protein prepared from glandless seed with no adverse effects on survival, growth or feed utilization. These results indicate that the CSM protein from glandless seed was utilized as efficiently as FMP by uvenile black sea bass. In comparison, low-gossypol CSM protein has been used to replace FMP at levels ranging from 20 to 100% for hybrid striped bass (Sullivan and Reigh, 1), channel catfish (Li et al., 2008), rainbow trout (Lee et al., 2006), Florida pompano (Riche and Williams, 2010; Cook et al., 2016), white shrimp (Siccardi et al., 2012; Richardson et al., 2016) and black sea bass (Anderson et al., 2016).

For southern flounder, growth performance (final weight and percent weight gain) of uveniles fed diets replacing from 0 to 100% FMP with GCSM-, GMO-CSM, or R-CSM protein was not different from fish fed a control FMP-based diet (Figure 2; Alam et al. 2018). However, in the present study, replacing FMP with 100% GCSM, GMO or R-CSM appeared to lower growth performance (albeit not significantly), suggesting that longer-term studies are needed. After eight weeks of feeding, survival of fish ranged from 80 to 1%, with no treatment differences. Apparent protein digestibility of diets was significantly higher for the fish fed 7 % and 100% GCSM and 100% GMO-CSM protein diets (83., 83. and 86. %, respectively) compared with

the control diet (7 .4%) (Alam et al. 2018). Arginine levels in the diets increased as CSM was increased, consistent with the high arginine concentrations found in CSM. Liver gossypol was only detectable in fish fed the 100% R-CSM diet (37 gg). Replacing up to 7 % FM protein by GCSM- or GMO-CSM protein did not affect whole body omega-3 PUFAs, or liver gossypol. The results suggest that up to 7 % of fish meal protein

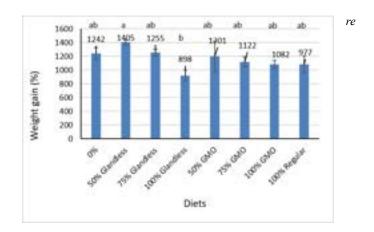


Figure 2. Effects of replacing FMP with CSM protein of different sources in southern flounder diets Glandless, GMO genetically modified (glandless), and Regular (glanded). A control fish meal based diet (0 CSM) was also tested. ars represent percent body weight gain of juvenile flounder after wee's of feeding. Values are mean SEM of triplicate tan s.Means with different letters differ significantly (Alam et al. 201).

may be replaced by GCSM or GMO-CSM protein in the diet of uvenile flounder without adverse effects on growth performance and body composition. In black sea bass uveniles, replacing 100% FM protein with GCSM protein did not alter growth performance or survival when diets were supplemented with lysine (Anderson et al., 2016), results similar to what was found in southern flounder in the present study. These findings suggest that CSM protein, regardless of source, was more digestible than menhaden FM in uvenile southern flounder.

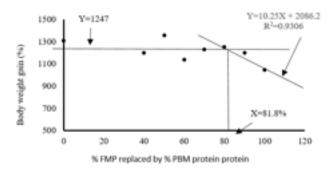


Figure . ro en line regression analysis between body weight gain and FMP replacement by P M protein in the diets of juvenile blac sea bass

E eriments ith o ltr ro ct meal BM

For black sea bass, no significant differences in body weight gain were observed in fish fed the 40 0% PBM protein diets compared with the control diet. However, body weight gain of fish fed the 100% PBM protein diet was significantly lower than in the control group. Regression analysis with body weight gain indicated that PBM protein can replace FMP in black sea bass diets at levels as high as 81.8%, with no reduction in fish growth performance (Fig 3) (Dawson et al. 2018). For fish fed diets with up to 0% PBM protein, feed conversion and protein efficiency ratios were not significantly different from fish fed a control 100% FM-protein-based diet (Dawson et al. 2018). Apparent digestibility coefficients of dietary protein remained high (81.6 87.0%) under all levels of FMP replacement with PBM protein. After the feeding trial, whole body and muscle protein content and the concentrations of whole body n-3 polyunsaturated fatty acids showed no significant differences among the treatments at FM protein replacement levels up to 0%. For the marine finfish totoaba, PBM protein was a good source of nutrients in uvenile diets at a FMP replacement level of 67%, whereas fish fed a 100% PBM protein diet showed the slowest growth and highest mortality (apata et al. 2016). In the marine fish, Florida pompano, growth performance was unaffected when PBM protein replaced 67% of the

FMP in the diet (Riche 201); however, growth performance was reduced when PBM protein replaced 100% of the fish meal protein (Rossi and Davis 2012). The results demonstrated that FMP can be replaced by feed-grade PBM protein in uvenile black sea bass diets at levels as high as 81.8% without adversely affecting survival, growth, feed utilization, fish biochemical composition, or ADC of protein or lipid. Poultry byproduct meal is a highly effective protein source for alternative protein-based feed formulation for black sea bass. uvenile southern flounder fed diets with PBM protein in substitution for FMP at levels up to 100% showed comparable growth (i.e., fish weight and percent weight gain) to fish fed the control FMP diet (Dawson, 2012). Many marine carnivorous species such as cobia (hou et al., 2011) and humpback grouper (Shapawi et al., 2007) have been reported to show no differences in growth when fed diets with substantial levels of PBM (60 and 7 %, respectively) in replacement of FMP, but reduced growth when FMP was completely replaced with a 100% feed- grade PBM diet (Shapawi et al., 2007). Feed utilization (feed intake, FCR, and P R) in uvenile southern flounder fed diets replacing FMP with up to 100% PBM protein was not significantly different from fish fed a control fish meal diet (Dawson, 2012). This suggests that palatability, digestibility, and assimilation of feed were not compromised by substitution of PBM protein for FMP in the diet at any replacement level (Dawson 2013). These results are very similar to those reported in African catfish (Abdel-Warith et al., 2001) and sunshine bass (Webster et al., 1 Whole body moisture content of uvenile southern flounder was not affected by level of PBM protein replacement for FMP. Dietary fatty acid composition was reflected in the whole body fatty acid profile (Dawson 2012). The ADC of protein by southern flounder was not affected by PBM protein substitution for FMP at levels of up to 100%, suggesting PBM protein is as digestible as FMP in uvenile southern flounder. In conclusion, PBM protein completely replaced FM protein in the diet of uvenile southern flounder without adverse effects on growth or feed utilization, indicating an

exceptional ability of this species to digest and to assimilate PBM protein.

CONC USION

The results demonstrate that uvenile black sea bass and southern flounder are able to utilize high levels of soybean meal, low-gossypol CSM, and PBM protein in their diet without reducing growth performance. About 68 and 3 % fish meal protein could be replaced by soybean meal, 82 and 100% of fish meal protein could be replaced by poultry meal in the diet and 100 and 7 % fish meal protein could be replaced by glandless cottonseed meal protein in the diet of black sea bass and southern flounder, respectively. These findings provide a basis for developing environmentally-sound and cost-effective predominantly terrestrial plant and animal protein-based diets for high value marine fish culture.

AC NOW ED EMENTS

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One of the biggest operational expenses in marine recirculating aquaculture systems (RAS) is high quality feed. Traditionally, fishmeal (FM) is used as the main source of protein in marine fish diets due to its high protein content, amino acid profile, digestibility, and palatability. Unfortunately, as the aquaculture industry continues to grow it is creating a higher demand for FM worldwide. Limited supplies along with increasing demand are causing the price of FM to continually increase. In an aim to investigate the replacement of FM by alternative protein sources in southern flounder diets, glandless cottonseed meal was tested at our University of North Carolina (UNCW)-Aquaculture Facility. Cottonseed meal (CSM) proteins from geneticallyimproved (glandless) low-gossypol seed (GI-CSM, 2.1% crude protein, CP), genetically-modified low-gossypol seed (GMO-CSM, 6.0% CP) and from an untreated regular (glanded) seed (R-CSM

4 . % CP) were evaluated to replace fish meal (FM) protein (. % CP) in uvenile southern flounder *Paralichthys lethostigma* diets. The results suggest that up to 7 % of fish meal protein may be replaced by GI- or GMO-CSM protein in the diet of uvenile flounder without adverse effects on growth performance and body composition. Cottonseed meal is a potentially cost- effective alternative plant protein source for use in aquafeeds. These ultra-low gossypol cottonseed flour proteins could be inexpensive protein sources for the commercial culture of southern flounder and other finfish species.

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Three experiments were conducted at our UNCW-Aquaculture Facility to determine the extent to which menhaden fish meal protein (FMP) can be replaced by solvent extracted soybean meal protein (SBP) in the diet of uvenile black sea bass Centropristis striata. Diets were formulated replacing FMP by SBP at 0, 10, 20, 30, 40, 0, and 60% (experiment 1) and 0, 60, 70, 80, 0, and 100% (experiment 2), with supplementation with squid meal, krill meal, and attractants in both experiments. xperiment 3 was designed to replace FMP by SBP at 40, 0, 60, 70, and 80% without supplemental squid and krill meal and at 60% and 70% with supplemental methionine and lysine. Diets were fed twice daily to triplicate groups of 1 per group) in 7 L tanks containing recirculating seawater. Fish were fed for 6, 10, and 8 weeks in experiments 1, 2, and 3, respectively. No significant differences in body weight gain, feed efficiency, and survival were observed among treatments in experiment 1. In experiment 2, no significant differences in percent weight gain were observed among fish fed diets replacing FMP at 0,

60, and 70%. In experiment 3, body weight gain was not significantly different for fish fed supplemental methionine and lysine in 70% SBP diets compared with fish fed 0% SBP diets. No significant differences were observed in whole body n 3 polyunsaturated fatty acids among treatments in experiment 2. Broken line regression of the specific growth rate data suggested that the maximum level of FMP replacement with SBP in black sea bass diets was 67.6 68.4% with 7 g kg squid meal and 0 g kg krill meal in the diet and 7.2 8.0% without squid and krill meal supplementation.

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Feeding trial was conducted to determine the maximum substitution of limits of poultry byproduct meal (PBM) protein for fish meal (FM) protein in the diet of uvenile black sea bass Centropristis striata (family Serranidae). ight isonitrogenous (44% CP) and iso-lipidic (13%, crude lipid) diets were formulated to replace FM protein with PBM protein at 0 (control), 40, 0, 60, 70, 80, 0, and 100% in black sea bass diets. Diets were fed twice daily to triplicate groups of uveniles to apparent satiation for 8 weeks in a recirculating aquaculture system. Final survival was excellent (-100%) in all diet treatments, with no significant differences. No significant differences on body weight gain were observed in fish fed the 40- 0% PBM protein diets compared to the control diet.

However, body weight gain of fish fed the 100% PBM protein diet was significantly lower than in the control group. Regression analysis with body weight gain indicated that PBM protein can replace FM protein in black sea bass diets at levels as high as 81.8%, with no reduction in fish growth performance. For fish fed diets with up to 0% PBM protein, feed conversion and protein

efficiency ratios were not significantly different from fish fed a control 100% FM protein-based diet. Apparent digestibility coefficients of dietary protein remained high (81.6-87.0%) under all levels of FM replacement with PBM protein. After the feeding trial, whole body and muscle protein content and

the concentrations of whole body n-3 PUFAs showed no significant differences among the treatments at FM protein replacement levels up to 0%. PBM is a promising alternative protein source for sustainable diet development in black sea bass.

Sea ee A ac It re in a Warming Environment B il ing Resilience aminariales Saccharina angustissima haeo h ceae

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: aquaculture, climate change, kelp, Saccharina angustissima

ABSTRACT

Locally sourced, high quality sea vegetables, in particular kelp, are sparking consumer interest and demand in the New ngland region, USA. The narrow-bladed sugar kelp, Saccharina angustissima, with one narrow (1.6 0.7 cm wide) and one long (average 1.8 0. 6 m, up to 4.4 m) blade is morphologically and genetically distinct from Saccharina latissima, which grows much wider (20-30 cm) and is usually shorter (up to 3. m) in length. In the intertidal, S. angustissima is adapted to withstand extreme hydrodynamic forces an order of magnitude higher than the subtidal populations of sugar kelp that are usually found in more sheltered locations. It is restricted in distribution in midcoastal Maine and occurs in the low intertidal, exposed to high ocean swells. In collaboration with kelp farmers from Maine Fresh Sea Farms and Maine Coast Sea Vegetables, we successfully domesticated this kelp, growing it to full maturity on longlines to be used commercially for its desirable culinary traits. Based on harvest results from two growing seasons, biomass yields of this kelp were up to 24.1 (6.3) kg m⁻¹ of line with a plant density of 400 plants m⁻¹ of line. The phenotypic characteristics that make this kelp exceptionally adapted to extreme sea conditions were preserved at the open-water farm sites. xperiments with the microscopic gametophyte stages of S. angustissima showed that it has the

capacity to withstand some temperature

fluctuations. Overall, this domesticated kelp species has great potential as an economically valuable sea vegetable in the Gulf of Maine. Nevertheless, because of its restricted geographic distribution, care must be taken to protect the donor population. Although additional work needs to be done to bring this kelp to market, growing it also offers a suite of ecosystem services, including nutrient bioextraction and temporary habitat formation for sea animals.

INTRODUCTION

Global aquaculture production continues to increase with wild fisheries stagnating. Sustainable cultivation can be a significant contribution for food security. Seaweeds, including those incorporated into integrated multi-trophic aquaculture (IMTA) and nutrient bioextraction, contribute to food and feed production. Currently, on a global scale, seaweed production is valued at 11 billion USD (FAO 2018) with % of cultivation taking place in developing countries where it provides income to millions of families. The global industry is dominated by a few main seaweed species with a predominance of Saccharina japonica, Undaria pinnafida, Pyropia/Porphyra spp., Gracilaria spp., Eucheuma/Kappaphycus spp. (im et al. 201; Buschmann et al., 2017). Seaweed aquaculture is also gaining interest in the US and urope. In New ngland, USA, kelp farming is developing as an environmentally and economically sustainable aquaculture activity that is revitalizing the working

waterfronts by providing economic opportunities that are ecologically sound (Augyte et al, 2017; Yarish et al. 2017).



Figure 1. Saccharina angustissima bed in the field showing long and narrow morphology.

The rare kelp, *Saccharina angustissima* (F.S. Collins) Augyte, Yarish Neefus, is an endemic narrow-bladed species found on localized and exposed islands and promontories on the contiguous mid-coastal areas of Maine, USA (Figure 1, Mathieson et al. 2008, Augyte et al. 2018). *S. angustissima*, has one blade that is very narrow (1.6

0.7 cm wide) and long (average 1.8 0. 6 m, up to 4.4 m) that is morphologically and genetically distinct from Saccharina latissima, which grows much wider (20-30 cm) and is usually shorter (up to 3. m) in length (Augyte et al. 2018). Many studies on kelps have identified phenotypic plasticity as a driver of the great range of morphological variation seen in thallus shapes as well as macroalgal production and physiology when exposed to various environmental parameters, specifically strong wave exposure (Hurd 2000, Blanchette et al. 2002, Fowler-Walker et al. 2006). Transplant experiments of kelp (in the order Laminariales) from exposed environments with rapid water movement to protected ones with slow flow, often result in blade morphology changing from narrow, thick, flat blades to wider, thinner, and more undulate blades

(oehl 2008). The research conducted here using both genetics and sea farming allowed us to test the assumption that this unique kelp had a genetic basis for its morphology and was not ust phenotypically plastic based on the exposed conditions of where it was found. It was also unknown if this species could be domesticated using traditional kelp cultivation methods.

MATERIA S AND MET ODS

Reproductive material was collected in Harpswell, Maine, USA, following low tides in October-November when the kelp was at peak sorus production. Following standard cultivation protocols, kelp was grown in nursery conditions at the UCONN Stamford Marine Biotechnology Laboratory on seedstring before being outplanted at two open-water sea farms in Maine. Sporophytes of the common sugar kelp, S. latissima, with parental meiospores obtained from subtidal populations from Casco Bay, Maine, were grown alongside the S. angustissima sporophytes to test how environment affects morphology. The wave energy at both farm sites was low compared to what the kelp population experiences in the field. The resulting lengths and widths of the two morphologies were compared at harvest. The kelp was grown to maturity on longlines for 6 months. In the lab, experiments were run to expose the microscopic gametophyte stages to different temperature and light levels to better understand the ecophysiological tolerance of these life stages.

RESU TS

The results of the common garden experiments at two sea farms revealed that the sporophyte *S. angustissima* blade retained its length to width ratio and did not become wide like the sporophytes of *S. latissima* (Figure 2). However, some characters, including thickness of the blade, were lost, while ruffles formed on blade edges.



Figure 2. Saccharina angustissima from the sea farm in Sorrento, Maine, USA. (Photo courtesy of Sarah Redmond).

These results confirmed that environmental cues alone were not wholly responsible for the unique morphology in *S. angustissima* and suggested a genetic basis. Peak meristematic growth rates for blade length occurred from March through April at 2.8 (0.34) cm day⁻¹. The kelp was harvested from May through une with yields of up to 17 (4.4) kg m⁻¹ of line and plant density of 330 plants m⁻¹ of line at the Bristol farm and yields of 13.3 (6.2) kg wet weight m⁻¹ line and a plant density of 400 plants m⁻¹ of line at the Sorrento farm. Second season yields at Sorrento were on average 24.1 (6.3) kg m⁻¹ of line.



Figure . Saccharina angustissima juvenile sporophyte grown in lab conditions.

The microscopic gametophytes and uvenile

sporophytes grew best at temperatures of 8-13 C at the lowest irradiance of 10- mol photons m⁻² s⁻¹ (Augyte et al. *Accepted*). Light had a significant effect on both male and female gametophyte grown only at the higher temperatures. Temperatures of 8-1 C and irradiance levels of 10- 100 mol photons m⁻² s⁻¹ were conditions for the highest uvenile sporophyte growth (Figure 3). Finally, calculations estimating the nutrient bioextraction capability of the cultivated *S. angustissima* kelp harvested in une reveal N removal of 88.7 kg ha⁻¹ and C removal of 1666.7 kg ha⁻¹ (Augyte et al. 2017).

DISCUSSION

Both farms grew significantly narrower blades of *S. angustissima* than of the sugar kelp, *S. latissima*. Common garden experiments with the two morphotypes identified trait stability for length and width, while blade ruffles and thickness varied with the environment. Overall, this unique kelp form has emergent potential as a crop for the Gulf of Maine, with several sea farmers growing it for commercial purposes. In terms of microscopic stages, high light intensity effects might pose stress on *S. angustissima*, although, as an intertidal species, it could also be better adapted to temperature and light extremes than its subtidal counterpart, *S. latissima*.

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Consumer interest and demand for North Atlantic sourced sea vegetables drives opportunities for aquaculture development in the northeast USA. The unique morphology and desirable culinary traits of the wild narrow-bladed kelp, Saccharina latissima forma angustissima, were successfully translated into a cultivated crop on two geographically distinct open-water farms in Maine, USA. nvironmental conditions, growth, and tissue analysis were quantified. Peak meristematic growth rates for blade length occurred from March through April at 2.8 (0.34) cm day⁻¹. The kelp was harvested from May through une with yields of up to 17 (4.4) kg m⁻¹ of line and plant density of 330 plants m⁻¹ of line at the Bristol farm and yields of 13.3 (6.2) kg wet weight m⁻¹ line and a plant density of 400 plants m⁻¹ of line at the Sorrento farm. Second season yields at Sorrento were on average 24.1 (6.3) kg m⁻¹ of line. Both farms grew significantly narrower blades of f. angustissima than of the sugar kelp, S. latissima. Common garden experiments with the two morphotypes identified trait stability for length and width, while blade ruffles and thick-ness varied with the environment. Calculations estimating the nutrient bioextraction capability of the cultivated f. angustissima kelp harvested in une reveal N removal of 88.7 kg ha⁻¹ and C removal of 1666.7 kg ha⁻¹ (combined farm site averages). Overall, this unique kelp form has potential as a new aquaculture crop for the Gulf of Maine while providing several coastal ecosystem services.

Augyte, S., L. Lewis, S. Lin, C.D. Neefus, and C. Yarish. 2018. Speciation in the extremely exposed intertidal: the case of *Saccharina angustissima* (Collins) Augyte, Yarish Neefus *comb. nov. et stat. nov.* (Laminariales, Phaeophyceae). Phycologia 7: 100-112.

Saccharina latissima is a perennial kelp with a circumboreal distribution from the North Pacific to the North Atlantic coasts. Our study clarified the taxonomy of the morphologically distinct Saccharina latissima forma angustissima (Collins) A. Mathieson from the low intertidal zone on exposed islands and ledges of Casco Bay, Maine, USA. To identify genetic divergence between the two morphotypes, S. latissima and S. latissima f. angustissima, we used a multilocus phylogenetic approach. Genetic analysis suggested low divergence between the two forms. However, there was as much or more genetic divergence between S. latissima and S. latissima f. angustissima as there were between other taxonomically accepted species of Saccharina. To investigate sexual compatibility between the two forms, we made reciprocal crosses of the gametophytes and observed sporophyte formation. All crosses were successfully grown to the uvenile sporophyte stage, suggesting that the two are reproductively compatible in vitro. It is unknown if the two populations freely hybridize in the field. Last, we compared wave action, the ecological factor most likely driving the unique morphology, at exposed sites with S. latissima f. angustissima and protected sites with S. latissima. The mean wave force at the exposed site was over 30 times higher in magnitude than at the protected site respectively, during the summer. The significant differences in morphology, the lack of specimens with intermediate morphologies, and the results of a common garden experiment suggest that the morphological differences in S. latissima f. angustissima are heritable with a genetic basis. Therefore, on the basis of our molecular evidence coupled with ecological studies, we are elevating S.

latissima f. angustissima (Collins) A. Mathieson to specific rank as S. angustissima (Collins) Augyte, Yarish Neefus comb. nov. stat. nov

. . im, G.P. raemer, and C. Yarish. 201. Use of sugar kelp aquaculture in Long Island Sound and the Bronx River stuary for nutrient extraction. Marine cology Progress Series 31:1 -166.

The present study revealed an even higher nutrient bioextraction capacity in the cold-water species Saccharina latissima at three sites - the mouth of the Bronx River stuary (Bronx, NY; BR), western Long Island Sound (Fairfield, CT; WLIS), and central Long Island Sound (Branford, CT; CLIS), during winter and spring of the 2012–2013 growing season. These sites differ in temperature CLIS WIS), salinity (BR (BR WLIS CLIS) and nutrients (BR WLIS CLIS). It was estimated that S. latissima could remove up to 180, 67 and 38 kg N ha^{-1} at BR , WLIS and CLIS respectively, in a hypothetical kelp farm system with 1. m spacing between longlines. In the same hypothetical kelp farm system, the estimated carbon sequestration values are 13 0 (BR),1800 (WLIS) and 1100 (CLIS) kg C ha⁻¹. The potential monetary values of N sequestration by the sugar kelp are up to 1600 (BR), 760 (WLIS) and 430 (CLIS) ha⁻¹, if incorporated into the State of Connecticut Nitrogen Credit Trading Program and a carbon-pricing scheme. The potential economic values of C sequestration are 30–300 (BR), 40-400 (WLIS), and 24-240 (CLIS) ha⁻¹. These results suggest that seaweed aquaculture is a useful technique for nutrient bioextraction in urbanized coastal waters, such as LIS and BR. Alternation of the warm- and cold- water species would maximize nutrient bioextraction and augment other ecosystem services, producing economic benefits for the region while helping to manage non-source eutrophication.

Scallo arming in Maine a ear Overvie

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: scallop, aquaculture, Maine, Aomori

ABSTRACT

Climate change has profoundly impacted capture fisheries in the Gulf of Maine. Notably, the largest fishery in the United States is in flux: landings of lobster (omarus americanus) have shifted northeast in recent years. These challenges which are not unique to Maine and are also experienced by apanese fisheries - have caused great uncertainty among fishermen in Maine. Consequently, some are turning to aquaculture to compliment fishing incomes. Maine's fishing and aquaculture industries have benefited from exchanges with apan's wellestablished scallop industry, and we continue to learn directly from the apanese. Coastal nterprises Inc. (C I), Maine Sea Grant (M SG) and others are working to introduce commercially-viable scallop aquaculture into the state of Maine, building on a 20- year-old foundation of expertise. Maine s sister-state relationship with Aomori Prefecture is central to this process, with an origin dating to the wreck of the Bath-built Cheseborough, which foundered off the coast of Aomori in 188. Currently, several pro ects are being carried out in Maine, developing the culture sector for scallops in coastal waters. An aquaculture cooperative in Maine focused on scallops is now in place. Optimization of spat collection has been ongoing work over several years. Biotoxin monitoring in scallop tissues enables some producers to access the live market. A

underway. Altogether, fishermen, farmers, economic development groups, scientists and regulators are engaged in pursuit of developing Maine's scallop aquaculture industry. These collaborations - combined with expertise from apanese colleagues - are a powerful aggregation of expertise to solve the many problems involved. x-vessel value for Maine scallops has risen for over a decade, though state and federal landings have increased, reflecting the value that consumers place on Maine scallops specifically; and that trend is expected to continue (Hale Group 2016). Trials by C I and collaborators indicate that the apanese production machinery appears to work well, and growth rates from ongoing studies and older works appear to fall within acceptable limits, often in excess of 2 mm year shell height (Cowperthwaite, unpublished; Morse, unpublished; Davidson et al. 2014; Pottle and Hastings 2001). Lastly, all indications point to a growout period between two and three years from spat (seed) to commercially viable harvest size. Aquaculture practitioners in Maine desire to incorporate apanese expertise so the industry can ramp up effectively and provide new sources of protein. We seek ongoing advice with lease site set-up and husbandry, growout techniques, biofouling and predators, equipment and machinery, handling, processing, food safety and packaging. This innovative and unique pro ect offers a one-of-a-kind exchange experience that is

dedicated license for scallop spat collection is

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helping to further strengthen ties between the United States and apan. The introduction of apanese technical knowledge and equipment to our Maine scallop industry is the first of its kind in the United States and is groundbreaking for the American aquaculture community.

Maine Aomori relationshi

Groups from Maine visited Aomori, apan in 1 2010 and 2016 with the primary focus on scallop aquaculture. Specific topics included: spat collection, growout techniques, product development, and environmental interactions. Aomori is the northernmost prefecture on the apanese mainland of Honshu (ust south of Hokkaido) and includes some of the most productive scallop grounds in all of apan.

Aomori has a population of 1. M people and when overlaid on the US (from a geographic standpoint) it's at the same latitude as the state of New York. There are several similarities Aomori shares with Maine, including rural communities with natural resource and tourism-based economies encompassing fishing, farming and forestry.

Maine has had a longstanding connection with Aomori; one that began with a shipwreck. On October 31, 188, the Maine-built Cheseborough, was caught by a typhoon off the coast of the Aomori Prefecture. The ship was destroyed and only four of the (23 member) crew survived. In 1, a Sister City agreement was signed between



Figure 1. The Cheseborough, built in ath, Maine, and lost off the coast of Aomori Prefecture, Japan, October 1, 1

Maine Aomori to remember and recognize the tragedy as an event that brought our cultures together, and eventually a sister state agreement was signed. Over the years there have been several cultural, education and trade exchange programs. This laid the foundation for the opportunity to learn about their scallop industry (Fig.1).

Following is a timeline of some important events in the development of scallop culture in Maine. In many cases, the scientific questions being asked are direct outputs from technology transfer from apan.

The delegation from Maine that visited Aomori in was an important event in the advancement of scallop farming and stock enhancement the US. Thirteen individuals from industry, science, management and education travelled for a week to Aomori Prefecture, where stops at processors, fishermen's cooperatives, and scientific offices provided the US delegates with critical first-hand information about spat collection and growout of scallops. quipped with this experience, there were a number of subsequent outreach efforts all along the coast, to introduce the materials and methods of seed collection to Maine s fishermen. Importantly, such events also involved scientists and regulators, which allowed the licensing process for such work to proceed smoothly, and to assist other scientists in oining the overall pursuit of diversifying the scallop industry in Maine. Another principal outcome was the publication of a report that covered the ma or findings of the trip, entitled: The culture of apanese scallops (Patinopecten yessoensis) (Beal et al.,).

In the years immediately after the 1 delegation, there were upward of 100 Maine fishermen who participated in scallop spat collection, along virtually the entire Maine coast, and several reseeding efforts - mostly of the informal kind - were undertaken. Of the quantifiable efforts that occurred, re-seeding success was undemonstrated

(Schick and Feindel, 200). Given this lack of clear proof that reseeding worked, most fishermen eventually stopped participating in spat collection. However, this period introduced spat collection to a wide segment of fishermen in Maine, and the processes and terminology involved became much more familiar territory for fishermen. In addition, spat collection provided a mechanism by which the early life stages of scallops became more clear, as well as the interplay between spawning populations, oceanographic conditions, substrate and larval behavior. Regular communications and interactions with Canadian scientists, fishermen and gear suppliers also grew during this time; all of which became resources for Maine participants. As a result, those in Maine involved in scallop spat collection and aquaculture became much more connected to efforts elsewhere, and in essence greatly broadened their professional networks.

In 2001, the consulting company Gardiner Pinfold completed a document entitled: Development Potential of the Maine Scallop Industry. The report included the potential for the development of an aquaculture industry, and identified farm production as an area of growth, though with caveats on sale of whole products and the size of the market.

2001 also saw a report from Tom Pottle and his collaborators on a demonstration pro ect in Cobscook Bay (Pottle and Hastings, 2001). The pro ect examined growth rates and density in caged scallops and was notable in that it provided a roadmap for the testing needed to sell whole scallops. This roadmap and regulatory process was invaluable for efforts that came later.

The mid- to late-2000 s saw relatively low activity in scallop reseeding and aquaculture, although involvement in spat collection continued each year with a small group of fishermen. Scallops were added to the species allowed within the Limited Purpose Aquaculture License (LPA) in 2008. This was important because the LPA was an easy way for new entrants to experiment with aquaculture in

Maine's coastal waters, and it opened the door for small trials. Consequently, familiarity with aquaculture increased for many fishermen, and the passing of time allowed fishermen to better understand the opportunities and mechanics of aquaculture, and not to see the new industry as merely a threat to their existing businesses.

Between 2011 and 201 , there were several significant events in both the relationship between Maine and Aomori, and the development of scallop farming in Maine.



Figure 2. 2016 Maine Delegation (front row) at the Aomori Prefecture Welcome Reception.

At the invitation of Shingo Mimura (Governor of Aomori Prefecture), Coastal nterprises, Inc. (C I), the Friends of Aomori (FOA), and Maine Sea Grant assembled a ten-member group of shellfish aquaculture industry professionals from Maine to visit Aomori, apan, in the fall of 2016. The purpose of the visit was to expose Maine aquaculturists to Aomori's well-established scallop aquaculture industry experts for an intense one-week tech transfer exchange trip to help advance the development of Maine's scallop aquaculture industry. The apanese are recognized world leaders in aquaculture and through the existing relationships Maine has with Aomori (Friends of Aomori, Maine-Aomori Sister-State Advisory Council) we strengthened our relationship with Aomori prefecture staff and private businesses through aquaculture. (Fig. 2)



Figure . Location and map Aomori Prefecture, Japan.

Our visit focused on Mutsu Bay, which is surrounded by Aomori Prefecture. The bay experiences three foot (1 m) tides and encompasses 6 0 square miles (1683 km²). Tsugaru Strait is to the north, the Sea of apan to the west, and the Pacific Ocean to the east. (Fig 3)

Aomori Agen a

Our agenda included visits to learn about scallop processing, the Fisheries Research Institute for a scientific and regulatory overview, a few different fishing cooperatives, observing aquaculture gear on land, a few trips out on scallop farms to witness ear-hung scallops being washed and stocking densities being reduced in pearl nets. Lastly, we spent two days on land viewing several demonstrations of specialized scallop aquaculture machinery including: scallop graders, an age pin automatic pin setter, scallop drilling machine, and scallop cleaning machines. We learned about the use of various equipment installed on scallop vessels to accommodate long line culture, as well as specific gear used in the water including: spat collection gear, pearl nets, lantern nets and ear hanging gear.

M ts Ba

A typical long line (in Mutsu Bay) is 200m (6 0

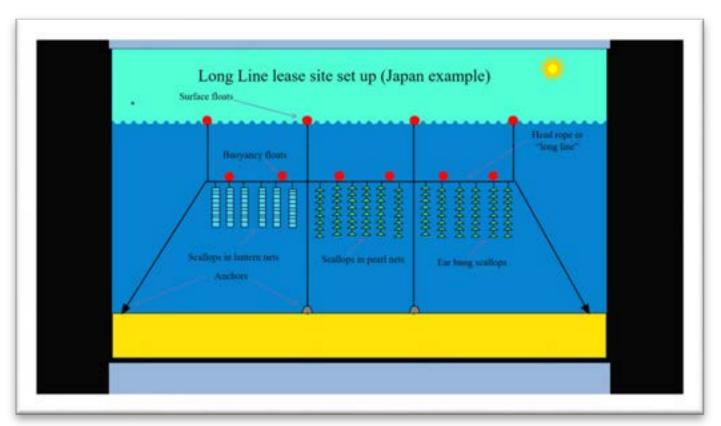


Figure 4. Schematic of a typical longline.



feet). Farms are sited in water that is typically at least 70 feet (23 m) deep or greater. All stages of collection and growout can occur on the same line including: spat collection bags, pearl nets, lantern nets and the ear-hanging technique. ach long line is held taut with tensioning buoys and suspended 10-20 feet below the surface of the water. Dropper lines with ear hung scallops or nets are kept 10 feet (3m) off the bottom to avoid predators.

Anchors weigh between 130-220 lbs. (-100 kg) and each long line has three anchors on each end (Fig 4). A long line is lowered and raised at a single point along the line to tend to the gear. The line is then secured on the bow and stern of the vessel with automated star wheels, which allows the vessel to slowly work its way along the line. Biofouling is partially mitigated by raising and lowering the line in the water column at different times of the year. High water temperatures are also a reason to lower the line to cooler waters.

Figure 5. The principal steps of Japanese scallop culture, and the times of year at which they occur.

Scallo C It re in M ts Ba

The basic process of scallop culture in Mutsu Bay is illustrated along a 12-month timeline. Spat collection bags are set in April-May. Aomori Prefecture and fishing co-ops collectively monitor environmental conditions in the bay to best predict approximate settlement and when to set bags.

In uly and August uveniles are transferred out of spat collection bags and put into pearl nets 80-100 animals per net chamber. In October, the uveniles are then transferred into clean pearl nets and the stocking density is reduced to 1 -20 animals per net chamber. In February and March, the scallops are then transferred to either lantern nets or the ear hanging technique for further growout. At the beginning of the second year, some half-mature scallops are harvested and sent to market, others are grown out for the remainder of year two and some for a third year for the largest meats. (Fig.)

Commercial Efforts in Maine

The effort to test apanese scallop equipment in Maine with Atlantic Sea Scallops was led by C I. C I has been working with two commercial farms: Bangs Island Mussel and Pine Point Oyster Co. This involved drilling and pinning on ear hanging lines to determine: how the shells hold up to drilling, the optimum time to drill, and to track mortality. Several months later we returned to the ear hung scallops to run them through the wash machine to clean off the biofouling organisms. Observations were made about the organisms encountered, the fragility of shells going through the machine, pins breaking off, mortality, cleaning frequency and growth rates. We've also been tracking labor costs and market price paid at final sale of the scallops.

S at ag materials:

Two seasons of spat collection experiments were undertaken to examine the efficacy of collection substrate on numbers of scallops retained, size frequency, and presence of other settlers. The rationale behind the experiments was that the standard substrate - Netron - has grown very expensive in recent years. We undertook two seasons of experiments to compare Netron to less expensive options.

In September and October of 2016, four lines of spat collectors were deployed in the coastal waters of Maine, approximately three miles southeast of Cape lizabeth, in approximately 0m of water. ach line supported 12 spat collectors; four at each depth of approximately 40m off bottom, 30 m off bottom, and 24 m off bottom. ach level contained one collector filled with the following substrates: Netron (serving as the control), and one each of Industrial Nettings 1 6 (O -7100), (O -7822), (O -1 81). ach collector contained

approximately 21 ft² of substrate. Collectors were hauled in une and uly of 2017, and comparisons were made via ANO A for the number of scallops retained, their size frequency, and the numbers of other bivalve settlers in the bags, by mesh type and depth. Depth significantly affected the number of scallops and other bivalves per collector (ANO A, p 0.001 p 0.027), but not scallop shell height (ANO A, p 0. 1).

Additionally, the number of mussels was significantly higher near the surface of the water (ANO A, p 0.001, Table 1). The mean number of scallops per collector (reported as the average, the standard error) was highest at mid depths with mesh (122 3 2671). The mean number of all other bivalves per collector was highest at mid depths (1084 1 0), and the mean number of mussels per collector was highest near the surface of the water column (3 3 4). Mussels were analyzed separately because they can cause substantial damage to scallop spat. The mean scallop shell height across all depths was .41 0.2 mm (Fig 1).

Retrieval date had no effect on the mean number of scallops per collector (t-test, p 0.24) nor the mean shell height of scallops (t-test, p 0.4). Retrieval date affected the mean number of other bivalves per collector (t-test, p 0.004), but because scallops grow slowly, the two-week difference in collection date did not have a large effect and we treated all data equally, with four replicates of each factor.

Two spat bags were lost during the experiment: one at the surface with mesh and one at mid depth with mesh. Notably, blue mussels (*Mytilus edulis*) were observed to settle preferentially on the bags nearest the surface (Fig 6).

Table 1. Summary of effects of mesh type and depth on scallop catches in spat collectors, 2016 201.

	7171 2068	6 47 73	3773 1 37
	10 80 2803	122 3 2671	4620 31
	77 8 1366	8626 100	3 23 1001
	6771 11	7 2 871	42 4 140

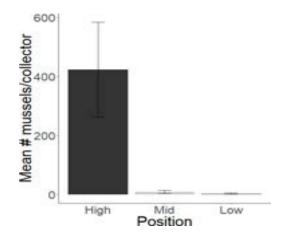


Figure 6. Mussel settlement on spat collectors by depth, 2016 201.

Given the results of the 2016-2017 experiment, we repeated the experiment, with slightly different variables. Collectors were set in the same locations, and set in Sept and October of 2017, and retrieved in une and uly of 2018. We repeated the same depths as previously, but our substrate types changed, to include: Netron (control), O -7822 (as in 2016 2017), and two similar agricultural nettings - Tenax Plurima and Tenax R- . All bags contained approximately 21 ft² of netting.

Although the data are not completely analyzed to date, we examined the same variables of depth and mesh type on the catches of scallop spat, size frequency and the presence of other settling bivalves. Data thus far indicates that for this site, the number of scallops collected goes down with

greater set depth and that the O -7822 appears to perform as well as the more expensive Netron (Table 2). In coming months, we will analyze the data using ANO A, to test for statistical differences.

Table 2. Mean number of scallops retained in collectors by depth and by substrate type, 201 201.

	7 4.64	3668. 7	3726.00	4 66.40
	7038. 7	2 .83	1831.67	470 .6
	287 .03	2326.2	24 . 0	2 66. 3
	44 .36	2 1.2	2862. 0	4237.70

E loring the feasi ilit of hole live scallo mar ets

Maine fishermen and scallop farmers have shown interest in pursuing markets for live and whole scallops, such as indicated by the Pottle and Hastings pro ect (2001) and the Gardiner-Pinfold report (2001), but biotoxins have presented a persistent threat to exploring this opportunity. Specifically, saxitoxin and domoic acid are differentially present in scallop tissues, and reside principally in the digestive gland and the mantle, where they can persist for weeks and months. Moreover, as the scallop metabolizes these compounds, they may transform from low-toxicity forms to high-toxicity forms over time, even in the absence of a harmful algal bloom (Bricel and Shumway, 1 8). Therefore, any meaningful effort will need to include a rigorous testing program to ensure the safety of any product other than adductor muscles.

Such work has been ongoing since May of 2017, on three sites along the coast of Maine. In brief, the testing and communications mechanisms created during the Pottle and Hastings pro ect have been revisited and updated and have allowed whole scallops to go to market in each winter between 2017 and 201. We have documented that toxicity will vary strongly by site, and that regular testing can provide the necessary confidence in product safety for regulators, producers and consumers. While testing using the mandated High-Performance Liquid Chromatography (HPLC) method is expensive, it has been helpful in supporting the market exploration of whole and live scallops, and has provided producers with additional income, while adding valuable information about the requirements of scale to be able to afford testing costs.

ANNOTATED BIB IO RA O E WOR S

Beal, B.F., M. Hastings, W. Hopkins, S. Inches, D. Morse, Porter, T. Pottle, S. Rappaport, L. Taylor, and Trenholm. 1 The culture of apanese scallops (*Patinopecten yessoensis*). Report of the Maine delegation to Aomori Prefecture, apan, May 14-21, 1 The Maine Aquaculture Innovation Center, 4 p.

This report summarizes the findings of the delegation from Maine to Aomori Prefecture in 1. It reviews the history of the scallop aquaculture industry and fishery in apan, and the technologies, management, science and to some degree the products and processing that have supported the development of this sector. In particular, there is attention paid to the spat collection process for scallops, underlaid by reviewing the biology of *Patinopecten yessoensis*, and the subsequent steps and equipment used in scallop farming in Aomori.

Bricel . M. and S. . Shumway. 1 8. Paralytic shellfish toxins in bivalve molluscs: occurrence, transfer kinetics and biotransformation. IN: Reviews in Fisheries Science 6(4): 31 -383.

This paper is a critical review of the biotoxin dynamics of several species of bivalve molluscs, including the sea scallop, *Placopecten* magellanicus. Specifically, the paper reviews the global distribution, sources of variation in toxicity, anatomical partitioning, metabolism and detoxification kinetics of toxins important in Paralytic Shellfish Poisoning (PSP). As it relates to the scallop aquaculture effort in Maine, the paper provides details showing that scallops can rapidly toxify, that toxins are spread differentially between tissues within the scallop, and that toxins can take an extended time (weeks or months) to be fully metabolized. Moreover, the paper notes that toxins change form during metabolism, and can go from less-toxic forms to more-toxic. The practical implication is that even in the absence of a toxic algal bloom, a scallop can test above regulatory limits, through this biotransformation process.

Davidson, L.A., R. Nowlan, and M. Niles. 2014. Open ocean sea scallop (*Placopecten magellanicus*) trials in Chaleur Bay: Comparing culture gears and husbandry practices. Canadian Technical Report of Fisheries and Aquatic Sciences 3121. 26p.

This report examined the use of several different gear types in raising sea scallops; some were suspended and some on the seabed. The main evaluation was a comparison of alternate gears versus the traditional lantern net. Lantern nets have low unit cost and good performance, but high labor cost for maintenance. The study identified cost and growth tradeoffs for the different methods used, and recommended cost analyses at given sites for the methods in question. Additionally, the study identified spat collection in the region as being high enough to support ongoing activities.

Gardiner-Pinfold Consulting conomists, Ltd. 2001. Development Potential of the Maine Scallop Industry. Report to the Maine Dept. of Marine Resources. 60p.

This report examines the status of the scallop

resource in Maine, and mechanisms by which the resource can be supported to gain maximum value. They address details of the product relative to the market, an examination of the wild fishery and its structure, and the possibility of cultured scallops.

Hale Group. 2016. Farmed Shellfish Market Analysis. A report to the Gulf of Maine Research Institute, Portland, M . 6 p.

The analyses for farmed product in this study addressed eastern oysters, blue mussels and sea scallops. Authors summarized the large domestic market for scallop products in the US, and the landings for US vessels as well as imports. They pro ect that supply over the next 1 years - the time of the planning horizon - will stay even or perhaps decline somewhat, and that Maine-produced scallops command a price premium. They also note that the industry is still in early phases of development, and much needs to be done to capture the market opportunity.

Pottle, T. . and M. Hastings. 2001. Sea Scallop Demonstration Pro ect, Final Programmatic Report. Submitted to National Fish and Wildlife Foundation. 10p, with appendices.

This pro ect brought forth important processes for the sale of whole scallops in Maine, and tested culture gear and density in a scientific setting. A purpose-built vessel was constructed for the pro ect, and growth was found to be acceptable at all densities tested. A protocol and Memorandum of Understanding was developed with the Maine Dept. of Marine Resources as a prerequisite for permission to sell whole scallop products. Bi-weekly testing of scallop tissues revealed that regulatory limits for saxitoxin and domoic acid were not exceeded over the duration of the study.

Schick, D. and S. Feindel. 200. Maine scallop fishery: monitoring and enhancement. Final report to the Northeast Consortium. Award 013-138-20. 1-30. 73p.

This pro ect created and refined a methodology for evaluating scallop enhancement efforts in coastal Maine, paired with information about the fishing fleet, and the spatial patterns of the scallop resource in terms of size, meat yield, relative abundance, recruitment and associated fauna. A GIS database of suitable spat collection areas was created, and spat collection field trials were undertaken. Field deployment of collected spat followed. Collections were generally positive, with some spat collectors containing over 10,000 individuals, although collections varied widely by location. Results of reseeding efforts were mixed, with an admission that more complete observation of re-seeded areas was warranted.

The Infl ence of Climate an Environment on the S a ning Con ition an arval Set of Nat rali e acific O sters rass str a gigas in a U S West Coast Est ar

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: Pacific oyster, condition index, Willapa Bay, spawning, climate

ABSTRACT

The Pacific oyster Crassostrea gigas was introduced to the US west coast in the early 1 00's where it replaced the native oyster Ostrea lurida and has become the mainstay for the shellfish aquaculture industry. Pacific oysters only regularly spawned and became naturalized in several discrete estuarine locations where conditions allowed for both adult oyster conditioning and spawning and larval survival, retention and settlement. The shellfish industry relied on natural set at these locations and or continued to import seed from apan until the advent of hatchery technology in the late 1 70's. Since that time, the industry has gradually shifted to hatchery production of oyster larvae, with large operators establishing their own in-house hatcheries and smaller companies incorporating remote setting facilities and purchasing product from hatcheries that specialize in producing late stage eyed larvae. Most operators continued to take advantage of wild set until a prolonged set of failures in larval production termed the seed crisis occurred in both hatcheries and in these estuaries like Willapa Bay, Washington, where wild set was and continues to be a feature of management on state established oyster reserves. The failures at hatcheries have been clearly linked to water chemistry and ocean

acidification, where aragonite saturation state is reduced and larval oysters are unable to deposit shell, but evidence for direct effects of carbonate chemistry in estuaries like Willapa Bay is more equivocal due to complex interactions with other factors. ariables like temperature, salinity, and phytoplankton as food, as well as other sources of larval mortality, are still related to the proximity of the ocean and cold upwelled water, but have been less studied in this estuary. Some of these factors also influence gametogenesis and spawning in adult oysters. We summarize historical records for spawning and setting in this naturalized Pacific oyster population, including a long term record of oyster condition. We propose a new effort to evaluate some of these additional factors and compare these records with data from locations in apan, where these oysters were originally sourced, and other locations with similar climate records where Pacific ovsters have been introduced. This effort will help understand and potentially mitigate for the effects of anthropogenic changes to these systems and an uncertain future climate.

INTRODUCTION

The Pacific oyster *Crassostrea gigas* was introduced to the US west coast in the early 1 00's where it replaced the native oyster *Ostrea lurida*

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and has become the mainstay for the shellfish aquaculture industry (Lindsay and Simons 1 Robinson 1 7; Shaw 1 7; Steele 1 64). Pacific oysters only regularly spawned and became naturalized in several discrete estuarine locations where conditions allowed for both adult oyster conditioning and spawning and larval survival, retention and settlement (Chapman and sveldt 1 43; uayle 1 88). The shellfish industry relied on wild set at these locations with the most consistent supply of US seed oysters produced in Dabob Bay in Washington state and or continued to import seed from apan until the development of consistent hatchery technology in the late 1 70's (Chew 1 84; Hi Im et al. 1 76; Langmo and Clark 1 7). Since that time, the industry has gradually shifted to hatchery production of oyster larvae, with large operators establishing their own in-house hatcheries and smaller companies incorporating remote setting facilities and purchasing product from hatcheries that specialize in producing late stage eyed larvae (Nosho and Chew 1 1). Most operators continued to take advantage of wild set until a prolonged set of failures in larval production termed the seed crisis occurred in both hatcheries and in these estuaries like Willapa Bay, Washington (Hales et al. 2017). Wild set, however, continues to be an important feature of management of Pacific oysters on state established oyster reserves in Willapa Bay (Dumbauld et al. 2011).

Due at least in part to the longstanding importance of the oyster industry in Washington state, a relatively substantial effort was devoted to understanding or at least monitoring and forecasting oyster spawning, larval abundance, and larval set, with water chemistry laboratories established at both the Washington Department of Fish and Wildlife (then Fisheries) field stations located on Willapa Bay (Nahcotta) and Dabob Bay (Pt. Whitney) (Lindsay et al. 1 ; Packer and Mathews 1 80). In addition to monitoring water chemistry and oyster larvae during the summer spawning season, a long term monitoring program was undertaken to examine the condition or fatness of adult oysters (Westley 1 ; Westley 1 61; Westley

1 64), with sampling in Willapa Bay beginning in and continuing with some gaps through present day.

Recent failures of Pacific oyster larval production at US West Coast shellfish hatcheries have been clearly linked to water chemistry and ocean acidification, where aragonite saturation state is reduced and larval oysters are unable to deposit shell (Barton et al. 201). vidence for direct effects of carbonate chemistry on larvae and wild set Pacific oysters in estuaries like Willapa Bay, however, is more equivocal due to complex interactions with other factors (Hales et al. 2017; Ruesink et al. 2018). Upwelled oceanic water, which already has a low aragonite saturation state, enters this coastal estuary during summer months (Banas et al. 2007; Roegner et al. 2002), and while this low aragonite saturation state has been enhanced by recent atmospheric CO₂ contributions (Hales et al 2017), other riverine drivers influence this chemistry up estuary (Ruesink et al. 2018). ariables like temperature, salinity, and phytoplankton as food have also been shown to vary along the estuarine gradient (Ruesink et al. 2003; Ruesink et al. 201) and potentially influence gametogenesis and spawning in adult oysters. This estuarine gradient is well known to the shellfish growers, in part as the fattening line in Willapa Bay (Banas et al. 2007; Dumbauld et al. 2011). Oyster growers typically plant or harvest uvenile seed oysters that are grown south of this imaginary line (from ust North of Nahcotta to ust north of the Nemah River on the ast side, Fig. 1) and transplant these oysters to better fattening beds located north of this line for eventual harvest. This line also serves as a rough boundary for the locations of successful spawning and setting of Pacific oysters in this estuary, with the ma ority of naturally set oysters occurring south of the line (Dumbauld et al. 2011; Ruesink et al. 2018). Thus older oysters north of the line have a high condition index or meat to volume ratio while those south of the line a much lower condition, in part because they spawn during the summer season and release gametes.

Condition index has been relatively extensively evaluated as a tool for examining the effects of environment on oysters and other bivalves elsewhere (Dutertre et al. 200; Filgueira et al. 2013; Rheault and Rice 1 6; Soniat et al. 1 8), and the Willapa Bay record was previously examined as well (Schoener and Tufts 1 87; Schumacker 1 ; Strickland and bbesmeyer). We are unaware, however, of any temporal records like this that now exceed six decades. We characterize spatial trends that are related to the slightly less complete record for spawning and setting of this naturalized Pacific oyster population in this estuary, and propose a new examination of the relationship between condition and similar records of ocean climate in the Northeast Pacific.

MATERIA S AND MET ODS

While all measures of bivalve condition relate some estimate of meat biomass to the volume of the shell cavity, multiple methods have been used to make these assessments (Abbe and Albright 2003; Crosby and Gale 1—0; Lawrence and Scott 1—82). Condition of oysters in Willapa Bay from 1—4—1—8 was measured using the Westley method (Westley 1—61) which uses Archimedes principle to obtain the shell cavity volume and is calculated as follows:

$$CI = \frac{\text{Dry body wt x } 100}{[\text{whole wt. in air - whole wt. in water}] - [\text{shell wt. in air - shell wt. in water}]}$$

Oysters were pooled to make these assessments so no estimate of error was obtained for these monthly values. After a detailed comparison with other methods, a consistent relationship was established between conditions collected using this technique and the gravimetric method, which is much easier to conduct on individual oysters and less prone to error (Schumacker 1). Thus, the gravimetric technique was adopted and all values from 1 present were calculated as follows:

$$CI = \frac{\text{Dry body wt x } 100}{[\text{whole wt.}] - [\text{dry shell wt.}]}$$

values are converted to gravimetric equivalents using the relationship developed by): Gravimetric Westley 1.180 -Schumacker (1 0. 0 (n 04, r^2 0. 126). Twenty oysters were sampled monthly at each of four sites in most years from 1 4 to present (Stony Pt., Nahcotta, Oysterville, and Stackpole; Fig. 1). xpanded surveys at six additional sites were conducted by WDFW biologists from 1 6 -1 78, with four of those notably located south of the fattening line described above, near the south end of the estuary (Refuge, Bear River, Smokey Hollow and Naselle, Fig. 1). We used these data to examine spatial and annual temporal patterns in condition index values and group data (selecting portions of the gametogenic cycle months) to average and obtain annual means and better examine the longer 60 year record. Some simple univariate analysis (ANCO A with month as a co-variate) were conducted to test for differences by site. We calculated and examined the long term anomaly deviation from the mean for each of these records. Finally, we examined two similar long term records of environmental data from the astern North Pacific: the 1-Nino Southern Oscillation (NSO; data accessed at https://www.ncdc.noaa.gov/teleconnections/enso in dicators soi) and the Pacific Decadal Oscillation (PDO; data accessed at http: research. isao.washington.edu pdo PDO.latest) and calculated similar anomalies to visually compare with the oyster condition index record and evaluated these relationships with simple multiple regression. We also used the condition index data from one site (Nahcotta) where oysters appear to spawn on a regular basis (e.g. condition drops dramatically between consecutive months) to determine whether these climate anomalies were correlated with the month(s) during which this apparent spawning occurred. Finally, we calculated a similar anomaly for the less continuous long term record of oyster larval set in Willapa Bay, collected by WDFW biologists and ourselves over time for comparison. We expect the mostly visual examinations presented here to instruct new quantitative time series analyses like those undertaken by previous authors where time lags and processes can be incorporated and further assessed (Schoener and Tufts 1 87).

locations for both *C. virginica* (Abbe and Albright 2003; Austin et al. 1 3; Galtsoff 1 64; Medcof

RESU TS AND DISCUSSION

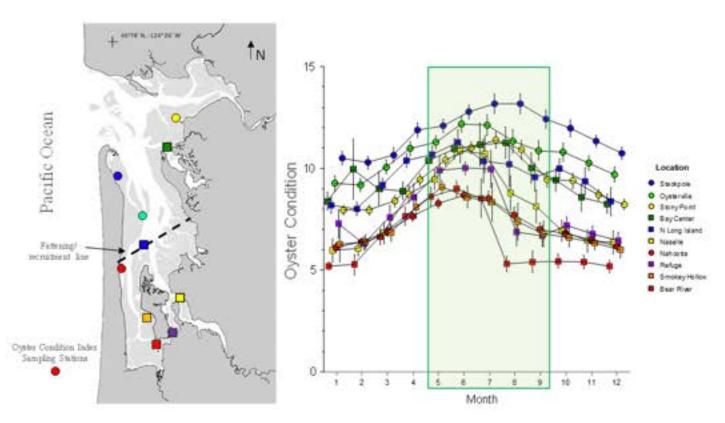


Figure 1. Average oyster condition at 10 sites in Willapa ay, Washington, monitored from 1 6 1 (bars SEs). ote estuarine gradient with higher values near the estuary mouth and lower values and seasonal drop during summer at sites south of an estuarine fattening line during late spring and summer (May Sept).

The seasonal pattern of oyster condition was similar, but average condition varied by site over the 10 year interval with data from 10 sites in Willapa Bay (1 68-1 7 , Fig. 1B). As might be expected, oyster condition increased from seasonal lows in anuary and February to highest values in summer (une August), and then declined again over fall months. This seasonal cycle has been well documented, and is related to changes in the reproductive cycle and quantity of associated nutrients including carbohydrates, proteins and minerals in the tissues and loss of meat weight when if the gametes are spawned at numerous

and Needler 1 41: Rainer and Mann 1 2: Rheault and Rice 1 6; Soniat et al. 1 8) in the US as well as C. gigas in Asia and multiple locations worldwide where it has been introduced (Chavez- illalba et al. 2003; nriquez-Diaz et al. 200; ang et al. 2000; Mason and Nell 1 Ubertini et al. 2017). While condition was significantly different amongst years (ANCO A, 2 .7, p 0.001), it was highest at sites near the estuary mouth and north of the fattening line (Stackpole, Ovsterville, Stony Point, and Bay Center) where it also fluctuated less, especially during the summer season. Oyster condition was significantly lower at the southern end of the estuary (Nahcotta, Smokey Hollow, Bear River, Naselle, and Refuge; ANCO A, Site F₁, 8 406.1, p 0.001) where there was often a more dramatic

drop during summer, presumably due to spawning events at these sites. This type of regional variation has also been well studied and related to environmental parameters like temperature, salinity and food (phytoplankton or chlorophyll) that influence oyster growth and reproduction (Brown and Hartwick 1 88; Dutertre et al. 200; Gasmi et al. 2017). We suspect that both temperature and food are the key factors in Willapa Bay, but that water residence time is also important. A north to south gradient in chlorophyll a, phytoplankton production, and phytoplankton species composition has been observed in Willapa Bay that is driven by intrusion of oceanic water into the estuary during spring and summer months (Newton and Horner 2003; Roegner et al. 2002; Ruesink et al. 2003). When a non-conservative tracer for Chl-a was added to a detailed oceanographic model created for this estuary (Banas et al. 2007), results suggested that there was a large difference in water residence time between areas north of a mid- estuary line, such that benthic grazers, like oysters south of this line, would not experience these intrusions of oceanic phytoplankton as often and potentially be sub ect to drawdowns of this production by grazers north of the line. Growers have recognized this feature for some time and traditionally plant small seed oysters to grow to harvest size and or place shell on their beds to encourage natural settlement of these oysters south of this line and then transplant them to areas north of the line before marketing them.

Based on the observed seasonal cycle, we divided the data for the extended 1 4-2017 time series into two separate time segments that reflect oyster physiology and ocean estuary conditions instead of calendar year: 1) winter and early spring months (October through April, termed winter) during which oysters resorb gametes and accumulate glycogen and other reserves and 2) late spring and summer (May through September, termed summer) during which oysters produce gametes and potentially spawn. The average condition during winter at the four sites measured over this longer time frame was highest at Stackpole (11.02)

0.13) and lowest at Nahcotta (6.64 0.0), and fluctuated over time declining on average from 1 4 1 7 and increasing since then (Table 1, Fig. 2).

Table 1. Average oyster condition inde (SE) for winter and summer months at each site. These values represent the entire time series $(1\ 5\ 201\)$ for the first four stations, but only the shorter period $(1\ 6\ 1\)$ for the latter $si\ sites$.

Site	Winter CI (Oct- April)	Summer CI (May-Sept)
Stackpole	11.02 (0.13), n 3	12.70 (0.18), n 237
Oysterville	10.02 (0.14), n 314	11. 7 (0.21), n 213
Stony Point	8. 6 (0.11), n 3 2	10.67 (0.17), n 266
Nahcotta	6.64 (0.0), n 3 1	8.06 (0.14), n 26
Bay Center	8. 1 (0.28), n 67	10.74 (0.31), n 48
North Long Island	.08 (0.20), n 62	10.42 (0.3), n 47
Naselle	6.76 (0.21), n 6	. (0.34), n 47
Smokey Hollow	6.66 (0.20), n 67	8.01 (0.32), n 47
Refuge	7.18 (0.21), n 67	8.63 (0.3), n 48
Bear River	.83 (0.17), n 6	7.32 (0.33), n 48

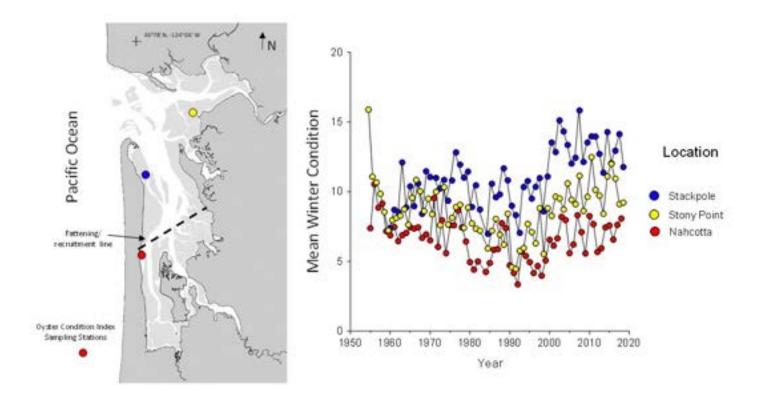


Figure 2. Average winter (Oct through April) oyster condition inde at sites in Willapa ay from 1 5 201. ote distinct differences between sites, but similar fluctuations and long term trends with a decline from 1 5 1 and subse uent increase.

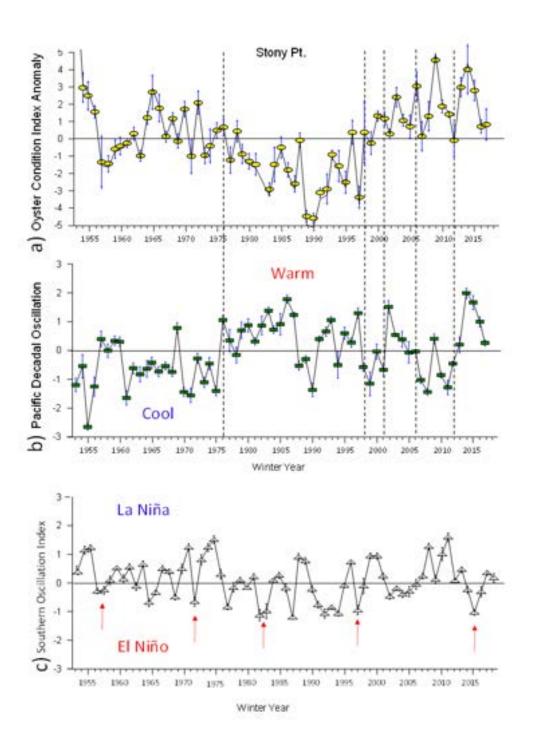


Figure . a) Oyster condition inde anomaly at Stony Pt. site for winter and spring months (October through April) compared with b) Pacific Decadal Oscillation anomaly and c) Southern Oscillation Inde anomaly for the same period. ote the a is has been inverted for the PDO and shifts are noted with vertical dashed lines and warmest El i o events with red arrows.

Summer values were higher but more variable, especially at sites closer to the southern end of the estuary, where rapid declines during these months indicated that the oysters likely spawned. Since the most complete data records were for the Stony Pt.

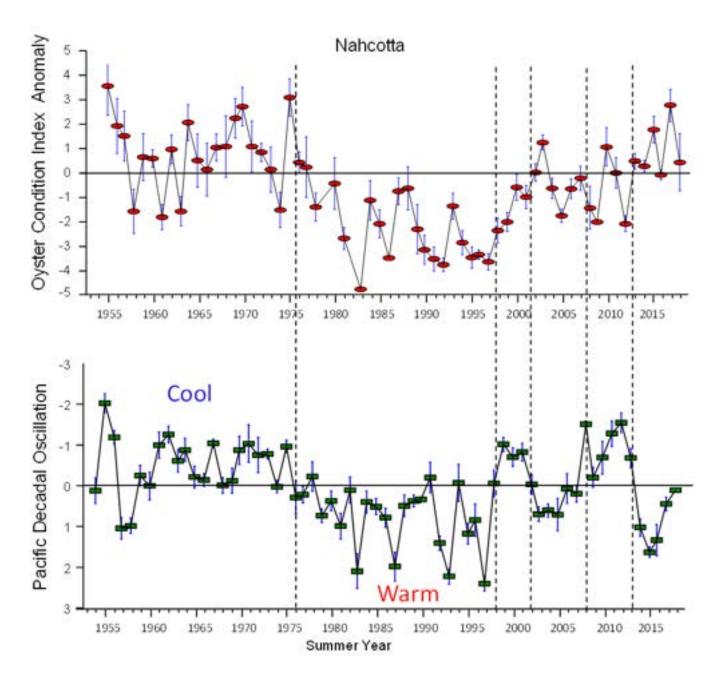


Figure . Oyster condition inde anomaly at ahcotta during summer months (May through September, top) compared with Pacific Decadal Oscillation anomaly for this same period (bottom). The a is for the PDO has been inverted and shifts are noted with vertical dashed lines.

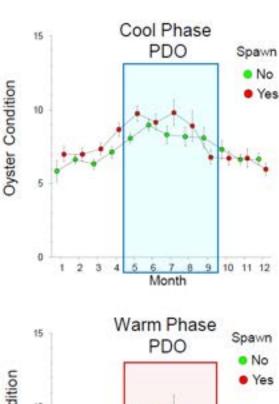
and Nahcotta sites, we used these records to calculate and evaluate the long term anomalies in condition and compare them with several indices of nearshore ocean climate. A general relationship was apparent between oyster condition and the Pacific Decadal Oscillation (PDO), with negative anomalies corresponding with warm phases and positive condition anomalies with cool phases (Fig. 3), but a breakdown in the strength of this pattern occurred after about 1 7. A general visual relationship was also evident between the winter

oyster condition anomaly and the southern oscillation index (SOI) anomaly, with positive SOI values (and 1 Ni o events) corresponding to lower condition and negative SOI (La Ni a events) corresponding with positive condition (Fig. 3). Though individual events, including those after 7, seem to be correlated, no statistical relationship emerged. Schoener and Tufts (1 87) found little relationship with sea surface temperature when examining the same data set, but a significant relationship with sea level height that best fit a phase lag with changes in sea level height preceding those in condition by several months. They documented the lowest condition index on record to that point in time corresponding with the 1 82-1 83 1 Ni o. That low value has now been surpassed by two even more negative condition anomalies, one of which corresponds with the even stronger 1 7 1 Ni o. The previous declining trend in oyster condition in Willapa Bay has also been correlated to the Pacific Northwest Index (a combined climate index derived from three more terrestrial parameters including temperature and precipitation) and the decline in Columbia River discharge (Strickland and bbesmeyer 1 we suspect the recent rebound in condition would not track at least the latter.

Table 2. Results of simple multiple regression relating the long term record of oyster condition inde anomaly at two sites to similar records for the Pacific Decadal Oscillation (PDO) and Southern Oscillation Inde (SOI). Results from two separate models for summer and winter averages are shown.

	stimate	S	df	F	P
Model			3	10.7	0.001
Summer PDO	-0. 7	0.1	1	2 .4	0.001
Summer SOI	0.01	0.28	1	0.002	0. 7
Site (Stony Pt.)	-0.0	0.24	1	0.0	0.81
rror Total			122 12		
Winter PDO	-0.12	0.22	3	2.48	0.12
Winter SOI	0.3	0.18	1	0.4	0. 0
Site (Stony Pt.)	-0.06	0.20	1	0.0	0.76
rror Total			123 126		

The relationship between PDO and the oyster condition anomaly was significantly negative in summer (Fig. 4, Table 2). Years when a significant drop in condition occurred during these summer months, signaling a spawning event, were clearly evident at the Nahcotta site, and we further separated this data into those years occurring during warm (PDO 0) and cool phases (PDO 0). Spawning occurred during both phases, yet during warm phases of the PDO, condition was significantly higher in years that these events occurred than when they did not (Fig. , ANCO A, 11.12, p 0.001), while there was no difference during the cool phase (ANCO A, F1,180 1.78, p 0.18). In those years when these spawning events occurred, they often occurred earlier (une and uly) during warm phase years and later (August) during cool phase years. This result suggests that oceanic conditions may affect the seasonal gametogenic cycle prior to spawning. More broadly, multiple triggers including temperature, but also food and even disturbance events, can be associated with spawning (Dutertre et al. 200; Ubertini et al. 2017).



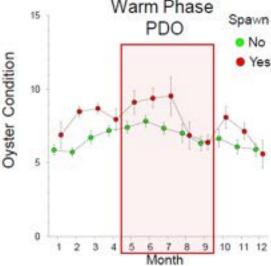


Figure 5. Average monthly oyster condition inde at ahcotta site for years where the average PDO anomaly in the summer (May through September) was negative (top) and positive (bottom). ears with a visually distinct drop in condition (presumably due to spawning) are contrasted with years when this was not apparent. ote the apparent seasonal difference in timing of spawn events with more and potentially earlier spawning events occurring during warm phase of the PDO.

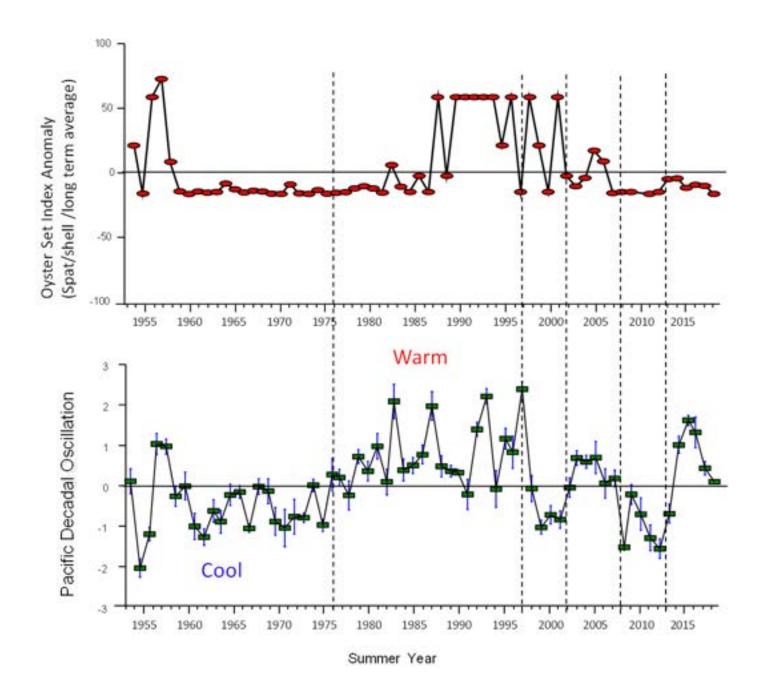


Figure 6. Anomaly of oyster sets (counts of oyster spat per shell) on seasonal shell strings deployed at three sites in Willapa ay (top) compared with the Pacific Decadal Oscillation anomaly for summer months (May October bottom) with shifts noted by dashed lines.

There appears to be a positive relationship between successful spatfall resulting from these spawning events and the PDO with larger spatfalls occurring during warm phases of the PDO (Fig. 6). While it may be that earlier or multiple spawning events are simply more likely to result in successful spatfall in this estuary, due to the typical three week period of larval development during which other factors (e.g. water retention, vertical migration, wind and weather) are involved, this relationship requires further investigation. The PDO has been related to numerous other biological endpoints, from zooplankton to coho salmon (Malick et al. 201; Menge et al. 2008; Shanks 2013). This included

oyster condition in one of the original characterizations (Hare and Mantua 2000), but defining the mechanism for this interaction with nearshore ocean conditions remains a useful research avenue. The PDO has recently also been related to phytoplankton community structure (Du et al. 201) with cool phases being connected to a diatom rich community and warm phase communities dominated by dinoflagellates. These authors have also shown that a similar community was present during the recent presence of an anomalous warm water mass (referred to as the blob, Bond et al. 201) in the nearshore coastal ocean off of the US Pacific Coast and that the chl a signature and overall biomass remained high. This could explain the breakdown we observed in the relationship between the PDO anomaly and oyster condition at about the same point in 2013, but more evidence should be gathered to test this hypothesis, and we propose that a new effort to evaluate some of these additional factors be initiated. This could include a comparison with data from locations in apan where these oysters were originally sourced and similar long-term records for other locations where Pacific oysters have been introduced and long term records are in place, in order to understand and potentially mitigate the effects of anthropogenic changes to these systems and an uncertain future climate.

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ANNOTATED BIB IO RA O E WOR S

Barton, A., G.G. Waldbusser, R.A. Feely, S.B. Weisberg, .A. Newton, B. Hales, S. Cudd, B. udeline, C. . Langdon, I. efferds, T. ing, A. Suhrbier, and . McLaughlin. 201 . Impacts of Coastal Acidification on the Pacific Northwest Shellfish Industry and Adaptation Strategies Implemented in Response. Oceanography 28: 146-1 .

This is the most recent review of the history and science underpinning the effects of changing seawater chemistry on bivalve shellfish larvae and the impacts that have already taken place to the commercial shellfish aquaculture industry on the

US West Coast. Multiple authors contributed to this review, which addresses a broad audience, but covers the leading research on direct effects to bivalve larvae, as well as monitoring seawater conditions and adapting to these changes.

Dumbauld, B.R., .L. Ruesink, A.C. Trimble, and B. . auffman. 2011. The Willapa Bay oyster reserves in Washington State: Fishery collapse, creating a sustainable replacement, and the potential for habitat conservation and restoration. ournal of Shellfish Research 30: 71-83.

The authors review the history of the state oyster reserves in Willapa Bay, Washington, where Pacific oysters were introduced in 1 28 and one of a few locations on the US West Coast where a naturalized population became established. They review the record of larval set in this estuary and attempt to link this to broad scale oceanographic conditions (but see Hales et al 2017 and Ruesink et al. 2018 below for a more detailed discussion of water chemistry within this estuary).

Hales, B., A. Suhrbier, G.G. Waldbusser, R.A. Feely, and .A. Newton. 2017. The carbonate chemistry of the Fattening Line, Willapa Bay, 2011-2014. stuaries and Coasts 40: 173-186.

The authors present detailed data on seawater chemistry (especially PCO₂ and aragonite saturation state) for Willapa Bay, Washington, where Pacific oysters have been the mainstay of the oyster aquaculture industry for almost 100 years and there is a long term record of spawning and setting. They reconstruct this record for a longer historical period and their data suggests that recent conditions provide a smaller window of optimal conditions (low aragonite saturation state and warm enough temperatures for oyster spawning) than occurred historically. While they did not sample larvae (see Ruesink et al 2018 below) and therefore can't

confirm effects, they substantiate the complexity of measuring these effects and attributing them to a single cause in an estuary where conditions are variable.

Ruesink, .L., A. Sarich, and A.C. Trimble. 2018. Similar oyster reproduction across estuarine regions differing in carbonate chemistry. Ices ournal of Marine Science 7: 340-3 0.

These authors measured seawater chemistry in Willapa Bay, Washington, but unlike Hales et al (2017), they also present simultaneously collected data on four cohorts of Pacific oyster larvae that were collected over three summers. The southern end of Willapa Bay has two arms which create distinctly different characteristics because one is much more affected by riverine conditions that cause reduced aragonite saturation relative to the other. Oyster settlement differed greatly between cohorts, but they did not find differences they could attribute to this different water chemistry, and instead found thermal conditions were perhaps more important.

Weisberg, S.B., N. Bednarsek, R.A. Feely, F. Chan, A.B. Boehm, M. Sutula, .L. Ruesink, B. Hales, .L. Largier, and .A. Newton. 2016. Water quality criteria for an acidifying ocean: Challenges and opportunities for improvement. Ocean Coastal Management 126: 31-41.

This review represents a call for developing ecologically relevant water quality criteria for acidification and augmenting coastal water quality monitoring efforts to characterize this at appropriate temporal and spatial scales. Historical data for oyster larval recruitment in Willapa Bay is one of two case examples presented that demonstrate the challenges involved in relating biological endpoints to appropriate water chemistry criteria in estuaries where conditions are variable.

A Revie of the otential of Macroalgae ee st ffs to Re ce Methane Emissions from R minant ivestoc

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: seaweed, climate change, methane, aquaculture, livestock

ABSTRACT

To address climate change in California beyond carbon-based gases, the state has recently passed legislation to reduce short-lived climate pollutants. In particular, the state aims for a 40% methane reduction by 2030. Several reductions are expected to come through existing technologies associated with landfills and energy producers. However, the state's largest contributor to methane production is cattle, specifically 1.4 million resident dairy cows and one million beef cattle. Methods to reduce gases produced from cattle have largely been unexplored in California. However, recent research in Australia has shown that some species of sub-tropical seaweed can reduce methane production from cattle by up to % when used as a feed additive at 2% inclusion rates of organic matter. Temperate species have not yet been assessed for methane reducing properties, but many of the species present in the California Current cosystem are closely related to these sub- tropical species. The nutrient-rich waters off the Californian coast are ideal for the development of seaweed aquaculture and this pro ect offers an excellent opportunity to address the underdevelopment of aquaculture in California. Seaweed aquaculture s

low environmental impact, coupled with the potential to mitigate climate change impacts and bio-remediate coastal eutrophication zones make this area of research a promising mitigation solution for climate change. Research is required to assess different species of local seaweed for total gas production and digestibility when added to cattle diets. Methane reducing candidate species also need to be assessed for optimal culture conditions and scalability to be used as a feed additive for ruminant livestock.

INTRODUCTION

Seaweed has been harvested by humans for more than 1700 years and has been used for a variety of purposes, including food, animal feed, and numerous industrial uses over this history. However, it is only in the last few decades that seaweed aquaculture has begun to develop. At last census, approximately 27 million tons of seaweed is cultivated per year, representing a quarter of the total aquaculture production worldwide (FAO, 2016). The vast ma ority of seaweed is produced in China and Indonesia, generating a combined total of 23 million tons. Cultivation of seaweed in the US is nascent and does not yet generate enough products

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to be quantifiable on a global scale. This imbalance is largely due to the traditional and common use of seaweed in Asian countries. However, a US market has begun to emerge based on powerful consumer preferences towards organic and sustainable products. The sustainable benefits of seaweed aquaculture are undeniable; it requires zero additional nutrient inputs for growth and does not displace terrestrial crops for arable land or require fresh water, representing an alternative biomass for nutrition, compounds and energy in the future.

Furthermore, seaweed aquaculture can provide a myriad of ecosystem benefits including: oxygenation and uptake of nutrients by enhancing primary production (squez et al., 2014); bioremediation of eutrophic zones, reducing harmful algal blooms; creating food, habitat and refuge for a diverse array of fish and invertebrates, thereby increasing biodiversity (Almanza and Buschmann, 2013); and mitigation of greenhouse-gas emissions by acting as a CO2 sink (Chung et al., 2011) and most recently by reduction of methane production from livestock (Li et al., 2018a; Maia et al., 2016).

Release of greenhouse gases into the atmosphere is a ma or environmental, social, and economic threat worldwide. Methane is a particularly potent greenhouse gas that is 2 times more efficient at trapping heat in the atmosphere than carbon dioxide (Solomon et al., 2007). It is well documented that ruminant livestock contribute significantly to the global release of methane via digestion. Methane is emitted from livestock because of methanogenesis (a byproduct) in the forestomach of ruminants; a similar process occurs in the decomposition of manure. The United States is the fourth largest cattle producer in the world and cattle contribute 26% of the total U.S. methane emission (PA, 2018). Dairy cattle are the largest methane producers among livestock and produce between 73 and 182 kg of methane per cow annually (Dong et al., 2006). California is the dairy industry's most productive state and has 1.4 million dairy cows that account for the ma ority of the enteric methane

emissions. Consequently, California recently enacted legislation in 2016 mandating a 40% reduction from present day methane emissions by 2030 (2016).

Several technologies and strategies currently exist to mitigate methane production from sources like landfills, yet solutions for the dairy industry are less developed. Dairy digesters, specialized structures used to capture methane generated from decomposing manure, are slowly being implemented on farms in California. However, this only accounts for about a quarter of the methane emitted from dairy cattle with the vast ma ority being released via burps as a result of enteric fermentation (PA, 2018; Powers et al., 2014). nteric methane is difficult to mitigate because it involves the natural process of a cow's digestion. However, research on dietary strategies have shown promise (Hristov et al., 201; Lee et al., 2017; Maia et al., 2016). Some dietary practices that have been shown to reduce methane include the addition of ionophores, fats, the use of high quality forages, and the increased use of grains (Caro et al., 2016). These nutritional strategies reduce methane through the manipulation of the ruminal environment, inhibition of methanogens and protozoa, or by redirection of hydrogen ions away from methanogens. Relatively new mitigation options have been investigated and include the addition of such additives as probiotics, acetogens, bacteriocins, organic acids, and plant extracts (i.e. condensed tannins) (Martin et al., 2010). More recently, researchers abroad have begun to investigate the effect of sub-tropical seaweed as feed additives for ruminants, discovering that a red macroalga called Asparagopsis ta iformis can reduce methane production by % at less than 2% inclusion in vitro and produced up to a 81% reduction in vivo sheep fed at 3% inclusion rates (Li et al., 2018a; Machado et al., 2016). The mode by which A. ta iformis reduces methane production is still unclear, but most recent research attributes the reductions to the bioaccumulation of halogenated methane analogues produced as algal secondary metabolites (Machado et al., 2018). The advantages

of seaweed over other dietary manipulations is that seaweed has shown the most significant methane reductions with minimal declines in digestibility (Li et al., 2018b; Maia et al., 2016), and seaweed production does not compete with terrestrial food production systems.

While an assessment of enteric methane production has been done for a number of tropical species (Machado et al., 2014), nothing is known about the methane reduction properties of temperate macroalgae. This presents an excellent opportunity for the development of the seaweed aquaculture industry in California, in temperate coastal waters that are highly productive compared to tropical latitudes, and thus provide superior conditions for seaweed culture.

: In concert with identification of methane reducing seaweed species, it is prudent to theoretically consider the scalability of seaweed production. The United States has the world's largest marine exclusive economic zone, an area equivalent to the total land area of all 0 states. The U.S. Department of nergy estimates that the U.S. has suitable conditions and geography for producing approximately 00 million tons (DW) of seaweed (ARPA-, 2017). California has the third largest coastline in the U.S., with state waters encompassing approximately 13,688 km². Based on a series of assumptions, including an average seaweed production value of 1,604 t km² year⁻¹ (DW) (iao et al., 2017) and a dairy cow consuming ,8 kg⁻¹year⁻¹ (DW), a total of 140,000 t (DW) seaweed year-1 would be required to feed the state's 1.4 million dairy cows at a 1% inclusion rate. Using these assumptions, a total of 86 km², or 0.6% of state waters, is all that would be required to culture the necessary seaweed volume to achieve approximately a 0% or more reduction of methane emissions from California's 1.4 million dairy cattle, the biggest methane contributor in the state. While this estimation is based on many assumptions, a total of 0.6% of state waters potentially being used for seaweed aquaculture seems relatively tolerable, especially when

considering the likely secondary ecosystem benefits resulting from seaweed cultivation. California has the potential to utilize this resource to build and grow the nascent seaweed aquaculture sector. Growing seaweed for livestock feed offers a unique opportunity to circumvent many challenges associated with terrestrial production systems, while also mitigating methane emissions. Such a large potential local market could help develop vital infrastructure, skills, and distribution systems for seaweed aquaculture, if realized.

: The potential economic benefit of this research to California's aquaculture industry could be vast considering the immense value of the state's dairy and beef industry at 21 billion dollars annually. Beyond this, the nation's combined cattle industry is worth 78.2 billion, of which most cattle are fed formulated diets for a significant period of their life. Development and production of feed additives in California that could be exported and supplemented to cattle diets across the United States would be extremely beneficial to California's entire economy. Additionally, the culture of seaweeds for such purposes in California could help develop nascent aquaculture industries providing alternative economic growth for coastal communities that are significantly dependent mostly on commercial fisheries. Outside the potential economic value of this research, the state would also stand to benefit environmentally by reduction of methane and absorbing agricultural nutrient nonpoint source pollutants as they are incorporated into seaweed biomass. California is home to a significant number of coastally drained waterbodies that are impaired due to high levels of nutrients coming from irrigated agriculture (Anderson et al., 2003). Sixteen of these eutrophic water bodies drain into the Monterey Bay where Moss Landing Marine Laboratories is located. Seaweed aquaculture has been shown to be effective in bioremediation of coastal eutrophic areas (He et al., 2008). Development of seaweed aquaculture could aid in reducing the impact of eutrophication by absorbing nutrients from the water and incorporating them into seaweed biomass. This biomass could then be harvested for a

variety of uses, including the recycling of these nutrients back into agricultural production as a feed ingredient.

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ANNOTATED BIB IO RA O E WOR S

Machado, L., M. Magnusson, N.A. Paul, R. de Nys, and N. Tomkins. 2014. ffects of Marine and Freshwater Macroalgae on In itro Total Gas and Methane Production. PLOS ON . , e8 28 .

The authors for the first time assess an array of different sub-tropical seaweeds for methane reducing properties in cattle using in vitro techniques. They find that a red seaweed species called *Asparagopsis ta iformis* is the most effective seaweed tested, reducing methane production by up to %.

Maia, M. R., A. . Fonseca, H.M. Oliveira, C.

Mendonca, and A.R. Cabrita. 2016. The Potential Role of Seaweeds in the Natural Manipulation of Rumen Fermentation and Methane Production. Sci Rep. 6, 32321.

This research is important in that it assesses the effect of the basal feed source (substrate) on methane reduction for a number of different seaweed species. The authors find that there is a significant effect on methane reduction based on whether the basal feed is either grass or corn, highlighting the need to further examine the seaweed interaction with different cattle feeds.

Li, ., H.C. Norman, R.D. inley, M. Laurence, M. Wilmot, H. Bender, R. de Nys, and N. Tomkins. 2018. *Asparagopsis ta iformis* decreases enteric methane production from sheep. Animal Production Science. 8, 681-688.

The authors for the first time assess the effect of seaweed on methane reduction using a whole animal experimental design. The article reports that sheep fed a 3% diet of seaweed *Asparagopsis* ta iformis reduce their methane emissions by 81% without any decrease in weight gain compared to the control treatment.

Con ition In e an ec n it of Manila Clam *u ita s hi i inarum* Relate to a itat Environment

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: Ruditapes philippinarum, Manila clam, condition index, fecundity, habitat

ABSTRACT

Understanding reproductive traits is important for maintenance or recovery of Manila clam Ruditapes philippinarum fisheries production, which has dropped considerably in apan over the past half century. Relationships between condition index and number of eggs (fecundity) of the Manilla clam were assessed at several clam habitats in Ise Bay, apan, during spring and autumn spawning seasons in 201. Condition index was calculated from the shell size and soft body wet weight. Fecundity was quantified using an indirect enzyme-linked immunosorbent assay (LISA) with a monoclonal antibody specific to a Manilla clam yolk protein. Land elevation (i.e. subtidal or intertidal), association with river input (river mouth or not), closeness of the shore, and chlorophyll a concentration in the water column were the habitat parameters analyzed in this study. A

correspondence analysis revealed that the condition index during both spawning seasons was positively correlated (r + 0.48 to +0.80) with the 1st and 2nd correspondence analysis scores on various habitat parameters. On the other hand, the fecundity, which is summed for each spawning season (three sampling days) and normalized to 30 mm shell length, was negatively correlated with the 2nd habitat score (spring: r -0.64, autumn: r -0.80). These results show that the nutritional condition as determined by the condition index is not the sole factor affecting fecundity of the clam. Actually, the fecundity was high in the intertidal zones and enclosed river mouth areas, whereas the condition index was higher in the subtidal zones and open non-river mouth areas. The information obtained about asari clam reproduction may be useful for stock enhancement by improving reproduction in the populations. This study was conducted as part of a research pro ect for recovery of asari populations by the Fisheries Agency of apan.

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INTRODUCTION

Manila clam, Ruditapes philippinarum, alias name short neck clam, or asari in apanese, is an important bivalve species for commercial and recreational fisheries in apan. According to the statistics released by the Ministry of Agriculture, Forestry and Fisheries of apan, fisheries production of suspended and bottom culture of bivalve species, such as scallop and oyster, are slightly decreasing or staying level in apan (Fig. 1). However, capture fisheries production of Manila clam has dropped considerably in apan over the past half century (Matsukawa et al., 2008). The production peaked at 170,000 tons in 1 83 and went down to ,000 tons % decrease. The Fisheries in 2016, which is a Agency of apan and local governments have undertaken some pro ects for clam stock enhancement, but the decreasing trend is continuing.

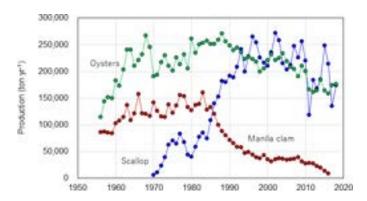


Figure 1. Fisheries production of bivalve species Manila clam, Japanese scallop and oysters, from the statistics released by the Ministry of Agriculture, Forestry and Fisheries, Japan. The production of Manila clam includes a small amount of purplish Washington clam, Sa idomus purpurata, production.

We need more basic knowledge on the population dynamics to improve or create new countermeasures to mitigate the population decline of the clam. Reproductive traits may be important for executing stock enhancement programs for the clam.

One of the most important reproductive traits to be studied is variation in fecundity (the number of eggs) of the clam associated with environmental factors. Some previous studies reported that individual fecundity is simply proportional to the clam's body weight; in other words, larger clams spawn more eggs than smaller clams (Toba Miyama 1 4) and robust clams have more fecundity than thin clams (Park et al. 2006, Hasegawa et al. 2014). Therefore, individual fecundity depends on the body weight and the nutritional condition, and fecundity per unit body weight is relatively constant among clams of different sizes. These statements sound intuitively true, however, these results were obtained for clams collected within the same habitat or cultured under similar rearing conditions except food level (abundance). ariation in the clam's growth depending on the habitat was evaluated in various studies such as akino (1 6). The effects of various environmental factors on clam fecundity should be further studied.

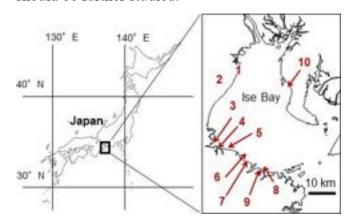


Figure 2. Study sites in Ise ay, Japan. The ID 1 on the map represents the Manila clam, Ruditapes philippinarum, habitat in Kusu, o aichi (2 Shiro o, Su u a, Miwatari, Matsusa a, Kushida, Matsusa a, 5 igashi urobe, Matsusa a, 6 Offshore Muramatsu, Ise, Intertidal Muramatsu, Ise, Kawaguchi flat, Ise, Isshi imae flat, Ise, 10 Kosugaya, To oname).

Habitat
ID

ID										
Land elevation	inter	sub	sub	sub	sub	sub	inter	sub	inter	inter
River input	none	River mouth	River mouth	River mouth	none	none	none	River mouth	River mouth	none
Chl. a g L ⁻¹	12.8	6.	7.6	4.	6.1	6.8	6.8	4.0	3.2	8.7
Openness of the shore	1 0	0	0	0	1 0	210	210	0	10	180

In apanese coastal areas, Manila clams can be found in various habitats. In Ise Bay, the clams are distributed in habitats ranging from intertidal to subtidal, exceeding 10 m depth, and from river mouth to non-river offshore areas (Hanyu et al. 2017). River mouth areas have large salinity fluctuations, while non-river areas typically have stable and high salinity water, ranging from closed and calm areas to more open areas under the influence of strong wave action, and from poor to good food availability. Therefore, we need to evaluate the reproductive capacity of the clam under a wide variety of environmental conditions.

MATERIA S AND MET ODS

St area an environmental characteristics of ha itat

Condition index and the fecundity was assessed in Manila clams collected from 10 locations in Ise Bay, central apan, in 201. The environmental characteristics of the studied clam habitats are

shown in Table 1. Here we focused on the land elevation, river water input, chlorophyll a level as an index of food abundance, and openness of the shore to offshore water. Chlorophyll fluorescence at around 1-m depth from the sea surface was measured by a CTD (Rinko profiler, F Advantech Co. Ltd., or DS, Hydrolab Co. Ltd.) and chlorophyll a concentration was estimated from these at each clam collection site. The mean chlorophyll a concentration in 201 (n 4 to 20) was used as the chlorophyll a level at each habitat. The openness was expressed as the angle between two lines connecting the habitat and two points on the coastline 1 km away from the habitat (i.e. a modified method of that described by Ruuskanen et al. 1).

Meas rement of con ition in e an fec n it of the clams

Manila clams were collected approximately monthly for monitoring of nutritional condition and fecundity from 10 habitats in Ise Bay between April and November 201. Approximately 10. 30 clams

with a shell length (SL) of 16 1 mm (mean 31.2 mm) were haphazardly collected. After measuring the shell length, shell height (SH), shell width (SW), and the total wet weight, the clams were dissected, and soft body tissue wet weight was measured. The soft body tissue of female clams (identified by the presence of oocyte in cut out tissue) was stored at -80 C for fecundity assessment. Condition index (CI) of the clams was calculated as follows;

$$CI = \frac{Soft\ tissue\ weight\ (g\ WW)}{SL\ (mm)\ x\ SH\ (mm)x\ SW\ (mm)}\ x\ 10$$

The number of eggs in the whole soft body tissue of individual clams was determined using an indirect enzyme-linked immunosorbent assay (LISA) with monoclonal antibodies specific to Manila clam yolk protein following Park and Choi (2004). Frozen whole tissue was freeze dried and homogenized using a sonicator in 10 1 ml buffer (20 mM Tris-HCl, 1 0 mM NaCl, pH7.) containing a protease inhibitor cocktail (04080-11 for General use; Nacalai Tesque Inc., yoto, apan). The homogenate supernatant was separated by centrifugation (1,000 g for 20 min at 4 C) and stored at -80 C until analysis. The supernatants were diluted 10-100 times with 0. % BSA-PBS buffer before LISA. A 100 1 sample was added to a 6-well polystyrene microplate and incubated at 4 C overnight. After incubation, the plate was washed twice with PBS containing 0.0 % detergent (Tween20; Promega Co. Ltd., Madison, WI, USA); 2 0 1 of 1% BSA and % skim milk in SBB was added as a blocking agent. After incubation at room temperature for 1 hour and washing times, the monoclonal antibody (Hamaguchi and Usuki, 2006) was added to 100- 1 aliquots in each well and incubated at room temperature for 2 hours. After washing times, a 100- l aliquot of polyclonal goat anti-mouse immunoglobulins peroxidase con ugate (diluted 1: 4000) was added to each well. Following incubation at room temperature for 2 hours, 100 1 of substrate- chromogen containing 3,3,, tetramethyl-benzidine (TMB+; Dako Co. Ltd.,

Glostrup, Denmark) was added as a color agent. Finally, 0 1 of 2-N sulfuric acid was added as the stop solution. The optical density of the end product was measured at 4 0 nm using a microplate reader. The tissue homogenate supernatant of the clams and the yolk protein content per egg obtained by Hasegawa et al. (2014) was used for calculation of the number of eggs. Clam fecundity was represented as the number of eggs per unit of soft body weight (1 g WW).

In south-western parts of apan including Ise Bay, the clam has two spawning seasons: spring and autumn, and the clam spawns repeatedly during each spawning season (Matsumoto et al. 2014). Therefore, as the representative fecundity in each spawning season, we summed the egg number that was normalized to the size of the clam, 30 mm SL (for removing the effect of size), for three months in each spawning season. However, the histological observation of the gonadal development showed that small numbers of Manila clam in Ise Bay were sexually undifferentiated even during the spawning seasons (Matsumoto et al. 2014). Therefore, the sum of the egg number might be overestimated in our study.

Anal sis of the relationshi s et een Manila clam an ha itat

To relate the condition index to habitat parameters, and fecundity to habitat parameters, correspondence analysis was performed using the computing environment R (R Development Core Team, 2017). Habitat parameters were summarized in two-dimensional graphical form (1st and 2nd axis). Correlation analysis was conducted between condition index or fecundity and habitat scores of these two axes in each spawning season. As the representative fecundity in each of the two spawning seasons, the egg number was summed and normalized to 30 mm SL for three months. The mean value of condition index for three months in each spawning season was used as the representative nutritional condition of the clam.

RESU TS AND DISCUSSION

Fecundity and condition index of the clam had seasonal and spatial variations (Fig. 3). Condition index and fecundity were higher in spring and autumn, coinciding with the spawning season as reported by Matsumoto et al. (2014). However, there were large variations in each month. These variations imply that there are spatial variations in condition index and fecundity among the habitats, although the variation of fecundity among individuals was also large. Correspondence analysis revealed that canonical correlations of the 1st and 2nd axis were 0.48 and 0.31, respectively, and the cumulative contribution ratio up to the 2nd axis was over 78%. The habitat could be summarized by these two axes. By summarizing the set of habitat data in a two-dimensional graphical plot, the 10 habitats were separated into 4 groups (Fig 4). Habitats in

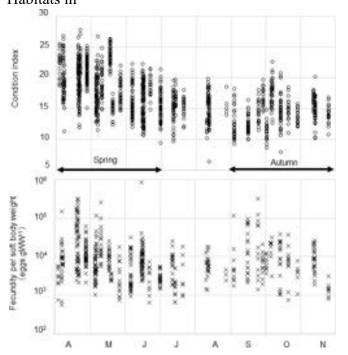


Figure . Seasonal variations in condition inde (a) and fecundity (b) of Manila clam, Ruditapes philippinarum, during 2015 in Ise ay, Japan.

non-river areas and open shore with strong wave action had positive mean scores of the 1st axis (right side) and food availability is considered to be higher than that in the river mouth and closed

habitats. Moreover, the river mouth and intertidal habitats had a negative score of the 2nd axis (below part) and the variation of environmental conditions, such as salinity and temperature, are considered to be larger than those in subtidal and non-river habitats. Therefore, it is considered that the 1st axis is related to food availability and the 2nd axis is related to the magnitude of environmental stress.

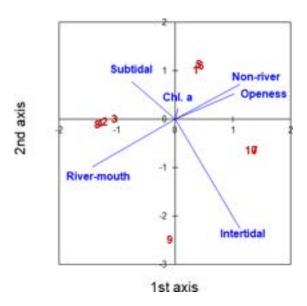


Figure . Two dimensional graphical plot t of Manila clam, Ruditapes philippinarum, habitats and habitat parameters in Ise ay, Japan by correspondence analysis. The ID 1 on plot represents the Manila clam habitat in Kusu, o aichi (2 Shiro o, Su u a, Miwatari, Matsusa a, Kushida, Matsusa a, 5 igashi urobe, Matsusa a, 6 Offshore Muramatsu, Ise, Intertidal Muramatsu, Ise, Kawaguchi flat, Ise, Isshi imae flat, Ise, 10 Kosugaya, To oname).

Relationships between habitat scores of the 1st and 2nd axis and condition index or fecundity in two spawning seasons were different between the condition index and fecundity and between spawning seasons (Fig.). In both spring and autumn spawning seasons, the average condition index of the 10 habitats was positively correlated with the habitat scores of the 1st axis (r +0.48 in spring and +0.7 in autumn) and the 2nd axis (r +0.80 in spring and +0.24 in autumn). On the other hand, fecundity during the two spawning seasons negatively correlated with the habitat score of the 2nd axis (r -0.64 in spring and -0.80 in autumn). There was only a limited relationship between fecundity and habitat scores of the 2nd axis (r

+0.07 in spring and +0.11 in autumn). Different relationships between condition index or fecundity to habitat scores show that the nutritional condition as determined by the condition index is not the sole factor affecting the fecundity of the clam, contrary to the results in previous studies (Hasegawa et al. 2018).

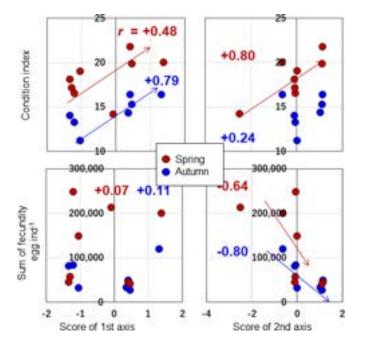


Figure 5. Relationships between condition inde or fecundity of Manila clam, Ruditapes philippinarum, and the score of 1st or 2nd a is of correspondence analysis.

Moreover, if the 1st and the 2nd axes are related to food availability and magnitude of environmental stress, respectively, condition index of the clam, which is positively correlated with the 1st axis, is considered to reflect the food availability. These relationships may be intuitively acceptable. Otherwise, fecundity, which is negatively correlated with the 2nd axis, is considered to reflect the magnitude of environmental stress. The fecundity was higher in the intertidal zones and river mouth areas than in subtidal and non-river (offshore) areas. This trend may be explained by clams exposed to stressful and hazardous habitats exerting more energy to reproduction than those under mild and reliable environmental conditions.

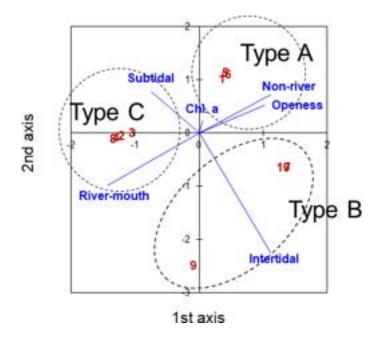


Figure 6. abitat types of Manila clam, Ruditapes philippinarum, on two dimensional graphical plot by correspondence analysis.

In conclusion, the nutritional condition as determined by the condition index is not the sole factor affecting the fecundity of the clam. The condition index was higher in the subtidal zones and open non-river mouth areas, where food availability was good. On the other hand, the fecundity was high in the intertidal and enclosed river mouth areas, which are stressful and life-threatening to the clams. These findings on clam reproductive traits may be useful in improving the efficiency of stock enhancement programs by maximizing the reproductive potential of the clam populations. For example, a habitat categorized as type A in Fig. 6 can be considered as a useful fishing ground in which the clam nutritional condition is good therefore, survival rate and growth are also good but the fecundity per body weight is not high. In habitat type A, enforcement of catch size regulation may improve the reproductive contribution through increasing the number of large clams. Habitat type B has ideal conditions for the clam to thrive because both nutritional condition and fecundity are high. Creation of this type of habitat may be helpful for sustainable fishing of the clam. The clam in habitat type C has a short life with high contribution to reproduction because fecundity per unit body

weight is high. Improvement of food availability in this type of habitat may increase the reproductive contribution of the clam.

AC NOW ED EMENTS

We are grateful to C. Hiro and . Seko for assistance with analysis. Also, the members of aiyo ngineering Co., Ltd, Diving Techno Co., Ltd and Fisheries Cooperative Association around Ise Bay for assistance with field research. This study was supported by the Ministry of Agriculture, Forest, and Fisheries and the Ministry of ducation, Culture, Sports, Science, and Technology of apan.

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Toba, M. and Y. Miyama. 1 4. Relationship of size to gonadal maturation and spawning in

artificially conditioned Manila clams. Nippon Suisan Gakkaishi, 60: 173-178.

In this study, the authors observed sexual maturation and measured the number of eggs released per spawned female sub ected to thermal spawning stimuli in breeding experiments for evaluating the size dependent batch fecundity of asari clam. Smaller clams at 0 yr old (groups A and B) produced as many eggs per soft tissue weight as larger clams at 1 yr (groups D and). No significant differences in egg diameter, percent fertilization, and larval survival from straight-hinge to pediveliger stage were observed among the clam size groups.

akino, . 1 6. Growth and survival of apanese littleneck clam *Ruditapes philippinarum* in cage test related to the sharpness index. Nippon Suisan Gakkaishi, 62: 376-383.

The author evaluated the variation in asari clam growth in a tidal flat in Tokyo Bay, apan, by using the sharpness index defined as width length of shell. The author concluded that uvenile clams offshore grew faster than those that were shoreside, but the uvenile survival rate in the intermediate zone of the flat was higher than those in the offshore and shoreside areas. uveniles having a high sharpness index tended to show a high survival rate. At the same age, the uveniles with a high sharpness index grew faster than or equal to those with a low sharpness index

Ocean Aci ification E eriment in Seagrass O ster Ecos stems Using a Novel S stem to Con ct In Sit CO Enrichment

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: Oyster aquaculture, ostera marina, Ocean acidification, Free ocean CO_2 enrichment, Non-environmental risk deterrents

ABSTRACT

Coastal ecosystems are under the influence of global environmental changes that threaten their ecological status and the sustainability of seafood production. Seagrass beds, which are widely distributed along coastlines around the world, are highly important ecosystems, offering various types of ecosystem services to human beings. Seagrass beds are important for environmental change mitigation and adaptation measures, via carbon storage, acidification mitigation, water quality improvement, as well as providing a nursery for coastal marine organisms. Therefore, seagrass beds are essential for harmonization of coastal fisheries and aquaculture with environmental conservation, resulting in the sustainable use of ecosystem services. Here we developed a novel system of insitu manipulation of CO2 and nutrient enrichment to demonstrate the function of seagrass in mitigating environmental changes. The system is called the Free-Ocean Real-Time xperimental System (FORT S). In addition, we developed nonenvironmental risk deterrents for invertebrates made from natural algae, specializing in arthropod and molluscan species (PCT P2018 038204). Two types of deterrents were made: liquid and drypowdered chemicals. The liquid deterrent can be

manipulated by FORT S. For instance, the deterrent against the arthropod species prevents settlement and recruitment of sessile crustaceans such as barnacles at low concentrations and also removes shrimps and amphipods from the target area at higher concentrations. Since the algal species used as a deterrent is also taken as an oral herbal medicine in Asian countries, the deterrent is safe for humans.

Recently, there has been a focus on the interaction between oyster aquaculture and seagrass vegetation as a common practice to achieve better coastal management with the framework of the sustainable development goals (SDGs) in a changing environment. Oysters have been grown using the original oyster seagrass interactions, including facilitation of spat survival rate, and especially trophic support from eelgrass beds to oysters over an extended period of time. This is typically regarded as the indigenous and local knowledge (IL) of apanese oyster aquaculture. Using the FORT S, we can demonstrate not only the direct effects of environmental changes such as pCO₂ increase pH decrease, eutrophication and agrochemical inputs on oysters, but also the mitigative effects of seagrass beds, which benefit oyster aquaculture.

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Campbell, . . and .W. Fourqurean. 2013. ffects of in situ CO₂ enrichment on the structural and chemical characteristics of the seagrass *Thalassia testudinum*. Mar. Biol. 160: 146 -147 .

Seagrasses commonly display carbon-limited photosynthetic rates. Thus, increases in atmospheric pCO₂, and consequentially oceanic CO₂(aq) concentrations, may prove beneficial. While addressed in mesocosms, these hypotheses have not been tested in the field with manipulative experimentation. This study examines the effects of in situ CO₂(aq) enrichment on the structural and chemical characteristics of the tropical seagrass, Thalassia testudinum. CO2(aq) availability was manipulated for six months in clear, open-top chambers within a shallow seagrass meadow in the Florida eys (USA), reproducing the forecasted CO₂ environment for the year 2100. Structural characteristics (leaf area, leaf growth, shoot mass, and shoot density) were unresponsive to CO₂(aq) enrichment. However, leaf nitrogen and phosphorus content declined on average by 11 and 21%, respectively. Belowground, non-structural carbohydrates increased by 2 %. These results indicate that increased CO₂(aq) availability may primarily alter the chemical composition of seagrasses, influencing both the nutrient status and resilience of these systems.

Gattuso, .P., W. irkwood, .P. Barry, . Cox, F. Gazeau, L. Hansson, I. Hendriks, D.I. line, P. Mahacek, S. Martin, P. Mc lhany, .T. Peltzer, . Reeve, D. Roberts, . Saderne, . Tait, S. Widdicombe, and P.G. Brewer. 2014. Free-ocean CO₂ enrichment (FOC) systems: present status and future developments. Biogeosciences 11: 40 7-407 .

Free-ocean CO₂ enrichment (FOC) systems are designed to assess the impact of ocean acidification on biological communities in situ for extended

periods of time (weeks to months). They overcome some of the drawbacks of laboratory experiments and field observations by enabling (1) precise control of CO₂ enrichment by monitoring pH as an offset of ambient pH, (2) consideration of indirect effects such as those mediated through interspecific relationships and food webs, and (3) relatively long experiments with intact communities. Bringing perturbation experiments from the laboratory to the field is, however, extremely challenging. The main goal of this paper is to provide guidelines on the general design, engineering, and sensor options required to conduct FOC experiments. Another goal is to introduce xFOC, a community-led initiative to promote awareness, provide resources for in situ perturbation experiments, and build a user community. Present and existing FOC systems are briefly described and examples of data collected presented. Future developments are also addressed as it is anticipated that the next generation of FOC systems will include, in addition to pH, options for oxygen and or temperature control. FOC systems should become an important experimental approach for pro ecting the future response of marine ecosystems to environmental change.

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Ocean acidification poses multiple challenges for coral reefs on molecular to ecological scales, yet previous experimental studies of the impact of pro ected CO₂ concentrations have mostly been done in aquarium systems with corals removed from their natural ecosystem and placed under artificial light and seawater conditions. The Coral Proto Free Ocean Carbon nrichment System (CP-

FOC) uses a network of sensors to monitor conditions within each flume and maintain experimental pH as an offset from environmental pH using feedback control on the in ection of low pH seawater. Carbonate chemistry conditions maintained in the -0.06 and -0.22 pH offset treatments were significantly different than environmental conditions. The results from this short-term experiment suggest that the CP-FOC is an important new experimental system to study in situ impacts of ocean acidification on coral reef ecosystems.

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Antifouling activity is one poorly investigated property of seaweed natural products. To determine whether seaweeds contain chemicals able to influence the settlement of fouling organisms in the field, crude organic extracts from Stypopodium onale, Dictyota menstrualis (Phaeophyceae) and Laurencia obtusa (Rhodophyceae) were incorporated at natural volumetric concentrations, into hard stable gels that served as substrata for fouling in the experiments. Fouling organisms settled at a significantly higher rate on plates treated with S. onale extracts than on control gels, while settlement was strongly inhibited on gels containing K. obtusa extracts. Fouling on gels treated with the D. menstrualis extract was not significantly different from the fouling found on control gels. The findings suggest that the broad antifouling properties of the crude extract of L. obtusa inhibit the settlement of fouling as well as hinder the development of settled fouling species, thereby reducing the richness of species. The results imply that L. obtusa possibly harbors powerful agents that can be explored for the development of antifouling technology.

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Since early 2008, an increasing number of countries have ratified an international treaty to ban the application of antifouling (AF) coatings based on organotin compounds (e.g. tributyltin (TBT) and triphenyltin). As a result, the demand for environmentally-friendly, non-toxic or low-toxic AF compounds and technologies (green AF agents) has become an urgent reality. Marine coatings based on Cu2O and various other biocides have a negative impact on the environment, and they must eventually be replaced by new, effective, and environmentally-friendly AF compounds. This mini-review describes important AF compounds discovered from a variety of organisms from 2004 until mid-200, and discusses recent and general trends in the discovery of AF compounds. Finally, a perspective on the future of AF compound development is presented. The discussion is aimed at updating scientists and engineers on the current challenges facing AF research.

Stark, .S., .T. Peltzer, D.I. line, A.M. ueiros, T. . Cox, . Headley, . Barry, F. Gazeau, .W. Runcie, S. Widdicombe, M. Milnes, N.P. Roden, . Nlack, S. Whiteside, G. ohnstone, . Ingels, . Shaw, L. Bodrossy, .D. Gaitan- spitia, W. irkwood, and . Gattuso. 201 . Free Ocean CO₂ nrichment (FOC) experiments: scientific and technical recommendations for future in situ ocean acidification pro ects. Progress in Oceanography 176: 8 -107.

Free Ocean CO2 nrichment (FOC) experiments are a relatively recent development in ocean acidification research, designed to address the need for in situ, long-term, community-level experiments. FOC studies have been conducted across different marine benthic habitats and regions, from Antarctica to the tropics. Based on this previous research we have formed some core operating principles that will aid those embarking on future FOC experiments. FOC studies have

potential to provide important insight into the effects of ocean acidification that can add to or refine conclusions drawn from laboratory or single species studies, because they are conducted in situ on intact assemblages. Scaling up from suborganismal and individual effects to also include indirect impacts on the ecosystem and ecosystem services, make FOC experiments essential to filling in current knowledge gaps in the understanding of ocean acidification. While FOC systems are complex, relatively costly, and somewhat difficult to operate, the challenges they pose are tractable and they have proven to be a useful approach in ocean acidification research. The aim of this paper is to draw from the experiences of past FOC experiments and provide practical

advice for designing, building and operating a FOC experiment. Some of the most important recommendations include: field testing the system design; having a backup power supply; using replicate treatment enclosures; monitoring and maintaining the chemistry appropriately; allowing sufficient time to achieve near CO2 equilibrium conditions; and having a scientific focus with a core set of hypotheses. Future FOC experiments could focus on longer durations, multiple factors, and testing more intact benthic marine communities and ecosystems. The authors hope this paper will enable further FOC deployments and experiments, as well as provide some guidelines to improve future FOC studies and advance ocean acidification research.

Effects of Ocean Aci ification on the Earl Develo mental Stages of the Commerciall Im ortant astro o s E o A alone an orne T r an in a an

Ryo imura^{1*}, Hideki Takami², Toshihiro Onitsuka³, Tsuneo Ono⁴, and Yukihiro No iri

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: Atlantic salmon, eastern oyster, breeding program, disease resistance

ABSTRACT

Commercially valuable shellfish species will be affected more seriously by ocean acidification (OA) in con unction with overfishing. We investigated the effects of OA on fertilization, larval development, metamorphosis, and post-larval growth of valuable gastropods; zo abalone aliotis discus hannai and turban shell Turbo cornutus, using a CO2 manipulation system (AICAL pCO2 control system, imoto lectric Co., Ltd, apan). There was no effect of exposure to 1000 atm pCO₂ seawater observed in fertilization, malformation, or mortality rates in either gastropod. However, fertilization and hatching rates decreased with exposure to 1 00 atm pCO₂ seawater and the malformation rates at 1000 atm increased significantly in zo abalone. The diameter of larval shells in 800 atm seawater was significantly smaller compared to those in control seawater (400-4 0 atm pCO_2) for both species. Scanning electron microscope images of larval shells of zo abalone from 1000 atm showed abnormal development, suggesting problems with shell deposition and or increased shell dissolution. The composition of developmental stages of embryos in horned turban differed between the experimental levels of pCO₂ seawater. These results indicate that pCO_2 1000 atm

reduced developmental performance of these gastropods in early life stages.

The natural habitat of these gastropods is kelp beds where photosynthesis and respiration by macroalgae lead to marked pCO₂ diel fluctuations. These fluctuations are larger than the pCO_2 change pro ected to be caused by OA for open ocean waters by 2100. We also conducted laboratory investigations on the effects of ocean acidification, in relation to diel cycles of seawater pCO₂ levels pro ected for the near and distant future, on the early development of aliotis discus hannai. Shell length, abnormality and mortality of larval abalone were examined under different pCO₂ levels with mimicked diel cycle using the AICAL pCO₂ control system. Adult abalone, with fully matured gonads, were induced to spawn and gametes were artificially fertilized in concentrations of ca. 400 (atmosphere-balanced, control treatment), 800 or 1200 pCO₂ atm. Sixteen hours post-fertilization, hatched larvae were exposed to running seawater ad usted to a constant level ca. 400, 800 or 1200 atm pCO₂ or fluctuating on a diel cycle 800 400 (400-1200) or 1200 400 (800-1600) atm pCO_2 pCO₂ in 4-day experimental periods. There were no significant differences in the mortality rate, abnormality rate and shell length of larvae among the 400, 800, and 400-1200 atm pCO_2 treatments.

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In contrast, the abnormality rate was significantly higher in the 1200 and 800-1600 atm pCO_2 treatments than in the control treatment. Larval shell length in the 800-1600 atm pCO_2 treatment was significantly smaller than in the other pCO_2 treatments. The exposed duration time exceeding the aragonite saturation state was a critical factor determining the negative impacts on the larval development of abalone.

INTRODUCTION

Anthropogenic elevation of the partial pressure of CO₂ (pCO₂) in seawater has resulted in a lowering of seawater pH (ocean acidification) and most calcifying organisms investigated showed reduced calcification in response to increased pCO₂ (Fabry et al., 2008). urihara (2008) reviewed ocean acidification (OA) effects on the early developmental stages of marine invertebrates including bivalves (oysters and mussels); however, no reports on larval gastropods were included. In order to clarify impacts of OA on fishery-important marine gastropods, we investigated the OA effects on the early development of zo abalone and the horned turban by constant elevated pCO₂ levels. We also investigated the effects of diel fluctuating pCO₂ levels on the survival and larval shell formation of zo abalone

MET ODS

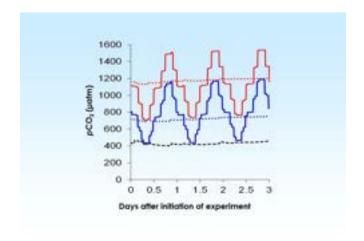


Figure 1. Changes in partial pressure of CO_2 in e-perimental seawater during the diel cycle treatment e-periment conducted 21-2 December 2010 in the Shiogama laboratory. Lines represent the values monitored by the AICAL system every hour. Dotted lines represent values in the constant treatment (1200, 00, 50 (atmosphere balanced) atm) and line represents those in the diel cycle treatment (00 1200, 00 1600 atm), respectively.

We used the AICAL pCO₂ control system, described in imura et al. (2011), for making controlled seawater of specific pCO₂ levels. The system showed stable and good separation of pCO_2 between tested groups through the gas monitoring and water sample (imura et al., 2011). We investigated fertilization rate, hatching rate, malformed rate, developmental stage retardation, and maximum larval shell length. These methods for rearing and testing were described in imura et al. (2011) for zo abalone and in Onitsuka et al. (2014) for horned turban. In the experiment testing diel-fluctuated pCO2 level on zo abalone, we set the pCO₂ levels as follows: 4 0 (atmosphere balanced), 800, 800, 400, 1200, 1200, 400, atm (Fig. 1).

RESU TS

There was no effect of exposure to 1000 atm pCO₂ seawater observed in fertilization, malformation, or mortality rates in either species. The fertilization rate and hatching rates decreased with exposure to 1 00 atm pCO₂ seawater in zo abalone. The malformation rates at 1000 atm increased significantly compared to those in control seawater in zo abalone. The maximum larval shell length of larvae from 800 atm seawater treatments were significantly smaller compared to those in control seawater in both species. Scanning lectron Microscope (S M) images of larval shells of zo abalone in 1000 atm seawater showed abnormal development with shell deposition and or increased shell dissolution. The proportions of embryos in horned turban that attained the 4-cell stage and 8-cell stage were significantly lower at the higher pCO₂ seawater treatments than those at the control levels 2 h after fertilization. Similar trends were observed in the ratio of veligers at 24-26 h after fertilization. These results are described in imura et al. (2011) for abalone, and Onitsuka et al. (2014) for horned turban, respectively.

In the diel fluctuation of pCO_2 level experiment, significantly higher malformation rates and smaller maximum shell lengths were detected in the 1200 and 1200+-400 atm pCO₂ treatments than in the 4 0 atm pCO₂ treatments. These negative impacts were greater in the 1200+-400 atm than in the 1200 atm. To test cumulative effects of elevated pCO₂, the index Integral pCO₂ over 1100 atm was used because the aragonite saturation state of seawater around the omega value of 1.0 was equivalent to 1100 atm pCO_2 . The index was defined by the following formula: Integral pCO₂ over 1100 atm (P-1100)i P: pCO2 over 1100 atmi: exposed hours to pCO₂ over 1100 atm

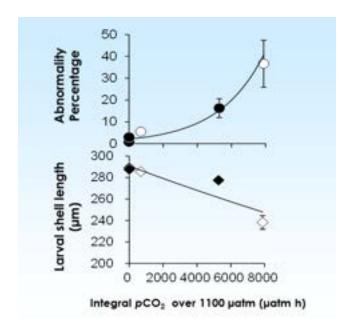


Figure 2. Relationship between larval malformation rates (upper), and shell length (lower) of aliotis discus hannai 2 h after initiating the cycle treatment e periment and the integral pCO2 over 1100 atm. White and blac colored symbols represent the results of the treatments with a constant and cyclically fluctuating partial pressure of CO2, respectively. Each bar represents standard error.

The abnormality rate increased with the increment of Integral pCO₂ over 1100 atm . Furthermore, the larval shell length decreased as the index value increased (Fig. 2). (The results of the diel fluctuation of pCO₂ experiments were described in Onitsuka et al., 2018 and were included with another experiment's results).

DISCUSSION

In these experiments, our results show that $p\text{CO}_2$ 1000 atm reduced developmental performance of these gastropods in early life stages (imura et al. 2011; Onitsuka et al. 2014). The mortality abnormality rate, and larval shell length were significantly different between the 1200 and 1200 400 atm $p\text{CO}_2$ treatments, indicating $p\text{CO}_2$ diel fluctuation can produce greater negative impacts on the larval life of abalone. The exposed duration time exceeding the aragonite saturation state was a critical factor determining the negative impacts on the larval development of abalone (Onitsuka et al. 2018: In the paper, the authors introduced a new index CPC, the effective

cumulative value of seawater pCO₂ over a specific criterion value, which was calculated using the threshold pCO₂ level defined as the pCO₂ value corresponding to 1.1 of the aragonite saturation state in seawater).

Since mitigation measures against ocean acidification are limited to reducing CO₂ emissions, small scale *p*CO₂ monitoring in coastal areas is thought to be important to properly assess OA effects on marine invertebrates.

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No iri. 2011. ffects of elevated *p*CO₂ on the early development of the commercially important gastropod, zo abalone, *aliotis discus hannai*. Fish. Oceanogr. ol 20: 3 7-366.

This is one of the first papers that investigated the effect of ocean acidification on commercially important gastropod species. This is also a very early paper using an AICAL system that precisely controls the pCO_2 concentration during the experiment.

urihara H. 2008. ffects of CO₂-driven ocean acidification on the early developmental stages of invertebrates. Mar. col. Prog. Ser. ol 373: 27 -284.

This review paper describes the effects of ocean acidification on the early development of many marine invertebrates including bivalves (oyster and mussel). Levels of carbon dioxide that affect the development of marine invertebrates were reviewed.

Onitsuka, T., H. Takami, D. Muraoka, Y. Matsumoto, A. Nakatsubo, R. imura, T. Ono, and Y. No iri. 2018. ffects of ocean acidification with *p*CO₂ diurnal fluctuations on survival and larval shell formation of zo abalone, *aliotis discus hannai*. Mar. nv. Res. ol 134: 28-36.

This paper assessed the effects of constant and diurnally fluctuating pCO_2 (mimicking the rhythm of the ocean) on development and shell formation of larval abalone using the AICAL system, which can precisely control pCO_2 . It is suggested that the effects of ocean acidification on development and shell formation of larval abalone can be determined by intensity and time of exposure to pCO_2 over the threshold associated with aragonite saturation state (-aragonite).

Chemical Changes in the Environment: What Does This Mean for Shellfish?

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: Bivalves, ocean acidification, coastal zones, aquaculture

ABSTRACT

Coastal areas, estuaries, and river deltas are experiencing an increase in carbon dioxide resulting from human activities. Approximately 30% of atmospheric carbon dioxide (CO₂) dissolves in the oceans. As this CO₂ dissolves, it reacts with water and produces carbonic acid (H₂CO₃), which dissociates into bicarbonate (HCO₃⁻), carbonate (CO₃²⁻), and hydrogen ions (H⁺). With increased CO₂ in the water, there are multiple ways that shellfish may be affected: (1) food availability; (2) physiological responses, and (3) sediment acidification.

Food availability may be influenced by changes in phytoplankton community structure and nutritional value. For example, increased carbon dioxide may cause a shift in phytoplankton biomass and community composition depending upon species' efficiency of inorganic carbon acquisition. Shifts in phytoplankton communities may favor phytoplankton with more- or less- nutritional content for support of human food production.

In addition to potential changes in food supply, coastal acidification may be causing physiological changes in marine bivalves that affect how they feed. Bivalve gills are constantly exposed to ambient water, with the gill cilia moving water through the shell for respiration and feeding. Under environmental stress, physiological changes to the

heart, gills, gonads, and general metabolism and regulation may occur. Physiological changes could result in slower growth, higher mortality, or inability to grow. The influence of ocean acidification on bivalve physiological functions is important to understand so that aquaculture may be modified to respond to a changing environment.

Finally, the same reaction that occurs in water during ocean acidification occurs at the sediment-water interface, where bivalves settle, following bacterial decomposition of settling organic matter. As particles settle, aerobic and anaerobic reactions, combined with redox processes, result in the production of CO2 in sediment porewater. Corrosive sediment can impede successful settlement, recruitment, and abundance in bivalve populations. Nutrient-enriched coastal areas in the United States are already experiencing sediment acidification, and this might be limiting bivalve recruitment at the sediment-water interface. The synthesis presented here will focus on the myriad of ways that ocean acidification may affect marine bivalves.

INTRODUCTION

Increased atmospheric carbon dioxide (CO₂) is being absorbed by the world's oceans and is thought to be a concern for some groups of marine organisms, especially those that contain structures composed of calcite or aragonite (Andersson et al., 2008; Dickinson et al., 2013; Gazeau et al., 2013).

In the ocean, CO₂ reacts with water to form carbonic acid (H₂CO₃), which dissociates into bicarbonate (HCO₃⁻), carbonate (CO₃²⁻), and hydrogen ions (H⁺), resulting in acidification. Currently average ocean pH is 8.1 and by 2100 ocean pH is predicted to drop to 7.7. (Caldeira and Wickett, 2003; Feely et al., 2004; Raven et al., 200 ; Turley and Findlay, 2016).

nvironmental variations (e.g., tidal, daily, and season changes) can influence metabolism, growth, and survival of marine shellfish (Norkko et al., 2006). Most bivalves can be classified as euryhaline organisms (Fl gel, 2013) and live in areas which experience changes in environmental parameters that are natural (e.g., temperature) and or a result of anthropogenic pollution (e.g., excess nutrients). Marine bivalves tolerate a range of environmental conditions; however anthropogenic pollution may affect bivalve energy balance such that a change in physiological processes (growth, reproduction) could result. Increasing our understanding of how organisms physiologically respond to specific changes in the environment will help in assessing how a community functions (Menge et al., 2002; Niemi et al., 2004).

In this study, we synthesize data from our ocean acidification experiments with phytoplankton growth and nutrition to examine physiological effects on bivalves exposed to ocean acidification, and discuss how sediment acidification may influence settlement.

h to lan ton

The correct nutritional components of phytoplankton are a key component regulating physiological performance of marine bivalves. The assimilation of key nutrients from phytoplankton, especially polyunsaturated fatty acids (PUFA), is required for growth and can only be acquired through trophic transfer (zgeta- Bali et al., 2012; Milke et al., 2004). Fatty acids are critical for growth, survival, neural development, stress resistance, membrane fluidity, and enzyme activity with enhanced growth and reproductive rates when fatty acids, especially PUFA and carbon (C),

nitrogen (N), and phosphorus (P), are optimized (lser et al., 2003; lser et al., 2000). Ocean acidification effects on phytoplankton may include changes in cellular carbon quotas (Burkhardt et al., 2001; Burkhardt et al., 1), changes in phytoplankton species composition (Riebesell et al., 2013), and nutritional quality to other trophic levels (Rossoll et al., 2012). To better understand what happens to marine phytoplankton following increased carbon dioxide, we conducted a series of seven experiments with temperate phytoplankton species, grown under pre-glacial pCO₂ conditions to predicted 2100 levels of CO₂. Table 1 summarizes the growth rates, C:P, N:P, and PUFA, MUFA, and SFA analysis from the seven species studied (ing et al., 201). This study found no significant difference in nutritional composition for any of these species, however some species experienced a change in growth rate. This research highlighted that although there were CO₂ driven changes in growth rate, it did not reflect changes in elemental composition or fatty acid composition. This research highlights that the complexing of ocean acidification on marine phytoplankton, will clearly depend on how the phytoplankton community changes. Understanding what drives community succession will help determine how changes in carbon dioxide might influence nutritional quality in the future.

Bivalve h siolog

Future ocean acidification conditions could result in decreased fertilization success, changes in growth rate, reduced calcification rates, and increased mortality (Melzner et al., 200). Our current understanding of how shellfish physiology is affected by ocean acidification is incomplete, but there is a growing amount of research suggesting that bivalves may be highly sensitive to ocean acidification (Dickinson et al., 2013; McCorkle et al., 201; Sugano et al., 201; Widman r et al., 2012). Understanding what physiological traits are responsible for sensitivity to ocean acidification may provide useful information for optimizing aquaculture practices with regard to bivalves.

Preliminary research suggests that under ocean

acidification, growth rates and survival of surfclams and sea scallops may decrease (McCorkle et al., 201; Sugano et al., 201; Widman r et al., 2012). Furthermore, shell deformities increased as carbon dioxide levels increased. Change in filtration rates is one possible mechanism for reduced growth. Preliminary short term experiments (two weeks) in the laboratory tested how ocean acidification affects the filtration rate of blue mussels, Mytilus edulis, and the eastern oyster, Crassostrea virginica, under current conditions and increased carbon dioxide. Feeding rates changed with increased carbon dioxide in both blue mussels and eastern oysters. Measurements of cilia beat function for blue mussels found that as carbon dioxide levels increased cilia beats decreased, suggesting that changes in feeding rates might be related to changes in cilia beat function (Meseck at al. 2020).

Dynamic energy budget (D B) models may be a useful tool to understand how changes in physiological rates would impact marine bivalves under ocean acidification. By understanding the metabolic costs of ocean acidification, we can model how biovolume, development, and reproduction might change. Further research on how bivalve physiology is affected under ocean acidification is necessary to understand how aquaculture practices may be adapted to the changing environment.

Se iment aci ification

Acceptance or re ection of habitat by bivalves as they settle from the water column to the benthos is not well understood, but research suggests that environmental, chemical, and physical variables at the sediment-water interface promote settlement and abundance of bivalves.

Recently, sediment porewater carbonate chemistry has been linked to larval settlement and survival in life stages of many marine bivalves (Clements and Hunt, 2014, 2017; Clements et al., 2016; Green et al., 2004; Green et al., 2013). Sediment acidification can occur at the sediment surface, as particulate material undergoes aerobic and anaerobic reactions, combined with redox processes, which produces CO₂. Once CO₂ is produced, it reacts with water to form carbonic acid (H2CO3), and dissociates into bicarbonate ions (HCO₃-), carbonate ions (CO₃²-), and hydrogen ions (H+) resulting in sediment acidification (Green and Aller, 2001; Tang and ristensen, 2007; Wenzh fer and Glud, 2004). These corrosive sediments (aragonite, in low pH conditions in the upper millimeters, where bivalves settle, and may reduce settlement, recruitment, and abundance in bivalve populations (Aller, 1 82; Green and Aller, 2001; warts and Wanink, 1 8).

Over the last three decades, bottom water in Long Island Sound (LIS) has been documented to have low pH (acidified sediments) and to be undersaturated with respect to carbonate minerals

Table 1. Relative difference in specific growth rate, carbon phosphorus ratio (C P), nitrogen phosphate ratio (P), PUFA, MUFA, and SFA for each of the seven phytoplan ton species using a one way A OVAP 0.05. The sign means no change between present day levels and ear 2100 pCO2 levels. Adapted from data presented in King et al. (2015).

Species	Growth rate	С:Р	N:P	PUFA	MUFA	SFA
Thalassiosira pseudonana						
Thalassiosira rotula	Increase	Decrease				
Thalassiosira weissflogii (CCMP2)						
Thalassiosira weissflogii (CCMP1010)						
Thalassiosira oceanica	Decrease					
Chlorella autotrophica	Increase					
Dunaliella salina						

(Aller, 1 82; Green and Aller, 2001; Green et al., 1 3). In Connecticut, the seafloor is leased to shellfish growers and they rely on natural reseeding of northern quahog *Mercenaria mercenaria* to occur. Understanding the influence that sediment acidification may have on natural settlement, recruitment, and abundance of commercial bivalves is essential for sustainable harvesting of shellfish. During 2013, we studied the relationships between abundance of bivalve molluscs and physical (grain size) and chemical variables (carbonate parameters) at the sediment-water interface on a leased shellfish bed during the bivalve settlement period when in LIS tends to be low (Meseck et al., 2018).

Briefly, Meseck et al. (2018) found that throughout the season, grain size in combination with carbonate chemistry was influencing bivalve settlement. When the settlement season was narrowed down to two weeks to remove temporal effects, grain size was no longer a significant contributor, but carbonate

chemistry still was. The contribution of carbonate chemistry parameters in predicting bivalve settlement explained 44% of the total bivalve benthic community, but the contributions of carbonate chemistry to settlement of individual species varied. For the two dominant species, carbonate chemistry explained 33% of M. arenaria and 71% of *ucula* spp. abundance (Table 3). These findings were consistent with previous research in LIS which reported increased mortality of bivalves (. annulata and Tellina agilis) in sediments that 1) with respect to were undersaturated (carbonate (Green et al., 1 3, 1 8). Our study did not focus on determining whether mortality or dissolution was occurring because of undersaturated sediments, however mean values in LIS were undersaturated during most of the sampling season. Results from the Meseck et al (2018) study suggest that increased pH and alkalinity may enhance bivalve community recruitment. These findings were similar to those reported by Clements and Hunt (2016) suggesting that carbonate chemistry may influence bivalve settlement.

DISCUSSION

nvironmental conditions of ocean acidification may influence bivalve survival, reproduction, growth, and overall health in both direct and indirect ways. Ocean acidification may influence multiple modes for marine bivalves. This summary illustrates how ocean acidification may influence phytoplankton production, shellfish physiology, and seafloor chemistry, and the importance of understanding these interactions for sustainable marine aquaculture.

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The authors conducted a field study assessing the relationship between uvenile soft-shell clam abundance and spatial, temporal, and environmental variables in the Bay of Fundy. Sediment pH, water temperature, location, and date was monitored in 2012 during the settlement season for *Mya arenaria* at four different study sites along the shore. Using Akaike's information criterion models that incorporated environmental variables alone, sediment pH, and minimum air temperature best predicted bivalve recruitment. By removing temporal effects, there was a significant positive relationship between sediment pH and grain size. These results suggest that carbonate geochemistry might influence bivalve settlement.

Meseck, S.L., R. Mercaldo-Allen, C. uropat, P. Clark, and R. Goldberg. 2018. ariability in sediment- water carbonate chemistry and bivalve abundance after bivalve settlement in Long Island Sound, Milford, Connecticut. Marine Pollution. 13. Pp. 16-17.

The authors conducted field experiments in 2013 to assess the relationship between porewater sediment carbonate chemistry (pH, Alkalinity, dissolved inorganic carbon), grain size, and bivalve abundance at two sites in Long Island Sound (LIS), Connecticut, USA. Akaike's linear model was used to predict total bivalve community abundance and

specific species abundance when appropriate. Over the entire season, 2 % of bivalve abundance at the sites could be explained by grain size, salinity, and pH. When temporal effects were removed, up to 71% of *ucula* spp. abundance could be accounted for by pH, phosphate, and silica concentrations in the porewater. These results suggest that bivalve abundance in LIS may be influenced by carbonate geochemistry and grain size.

Berm dez, R., M. Winder, A. Stuhr, A. Alm n, . ngstr m- st, and U. Riebesell. 2016. ffect of ocean acidification on the structure and fatty acid composition of a natural plankton community in the Baltic Sea. 2016. Biogeosciences. 13. Pp. 662 -663.

The authors conducted mesocosm perturbation experiments in coastal waters to investigate the response of phytoplankton to increased carbon dioxide with respect to community composition and fatty acid composition (nutritional statues). Their results found little change in plankton community composition from the natural condition (347 atm) to predicted levels for the year 2100 (1333 atm). The fatty acid composition was influenced by community composition, which was driven by phosphate and silica availability. The results presented here suggest that phytoplankton community composition is influenced more by nutrient availability than by CO₂, with little change in nutritional composition.

ing, A. L., B.D. enkins, .R. Wallace, Y. Liu, G.H. Wikfors, L.M. Milke, and S.L. Meseck. 201 . ffects of CO2 on growth rate, C: N: P, and fatty acid composition of seven marine phytoplankton species. Marine cology Progress Series. 37. Pp. -6 .

The authors conducted laboratory-based experiments on seven species of phytoplankton common in coastal estuaries of the North Atlantic, where carbon dioxide was manipulated to four levels ranging from the glacial minimum (2 0 ppm) to geological maximum levels (2 00 ppm).

Results indicated that carbonate system-driven changes in growth rate did not result in changes to elemental composition of the species studied. Fatty acid composition was not affected by elevated CO₂.

The results showed little sensitivity of common coastal marine phytoplankton species to elevated CO₂ that will result in changes in nutritional status.

C rrent Tren s in an Ocean Aci ification in A ac It re Sea Areas in o asho Ba a an

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: pH, AOU, ocean acidification, carbonate saturation state

ABSTRACT

Monthly variation in pH and carbonate saturation state () of aragonite and calcite were investigated in Gokasho Bay, Mie, apan, from November 2016 to October 201 . During the winter season, pH was almost constant (8.02 - 8.13) in the whole bay area, which was approximately the same level as those reported in surface waters off the south coast of apan. However, pH varied greatly during summer seasons (7.7 - 8.60). The lower value was thought to be caused by the remineralization of organic matter in the seabed, and the higher value was likely derived from high phytoplankton photosynthetic productivity. xcluding variation of pH caused by apparent oxygen utilization (AOU), acidification in the bay water was calculated to be 8.13 pH at present. The pH values currently measured are generally lower than those recorded in the 1 80s, suggesting the advance of ocean acidification (OA) in the bay water with a reduction rate of 0.08 pH decade. The calculated average values of 3) were relatively high in the whole bay area, indicating calcium carbonate is supersaturated in the current environment of the bay. If OA advances in surface water around apan in the near future, the degree of in the bay water would drop with the reduction in pH, and some negative impacts on

calcareous organisms may be brought about by this change.

INTRODUCTION

Coastal environments are influenced by various human-induced stressors. Recently, the global reduction of pH in surface seawater, which is referred to as ocean acidification (OA), has arisen as a consequence of oceanic uptake of anthropogenic CO₂. It is thought that OA affects the biogeochemical dynamics of calcium carbonate in the oceans (Doney et al., 200; awano, 2010; Garrard et al., 2013). OA is suggested to damage not only marine phytoplankton production but fisheries and seafood industries (Fu ii, 2018). Shellfish aquaculture occurs in many parts of coastal and bay areas in apan and there is a possibility that shellfish aquaculture will also be damaged by OA in the near future since the shells are composed of calcium carbonate whose deposition is controlled by carbonate species and the pH of seawater (Poach et al., 201). While there are many research studies on pH in open ocean and coastal areas (Fu ii et al., 2001; Ishii et al., 2011: hai et al., 2014; Lipuma, 2016; ubota et al., 2017), the effects of OA are not well understood in sea aquaculture areas due to large variations in pH

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associated with many biogeochemical processes. To obtain basic information for water quality of the coastal sea in apan, monthly hydrographical observations were conducted in a sea aquaculture area in Gokasho Bay, Mie, apan from November 2016 to October 201. This paper provides recent, high-quality pH values and carbonate saturation states () of aragonite and calcite in the bay water. Spatial and temporal variations in pH values will be discussed in relation to ocean acidification in apanese aquaculture sea areas.

MET ODS

St area

Gokasho Bay is located on the Pacific coast of apan, and consists of 3 coves (Gokasho, Funakoshi and Hazamaura) and the main bay which opens to the western North Pacific (Fig. 1). The Bay has a relatively small area of 22.2 km² and a water volume of 0.11 km³. Mean depth of the Bay is 12.7 m, and maximum depth is 30 m at the central part of the bay (Toda et al., 1 0). Hydrographical features of the bay are reported in detail by Toda et al. (1 0) and Abo (2000).

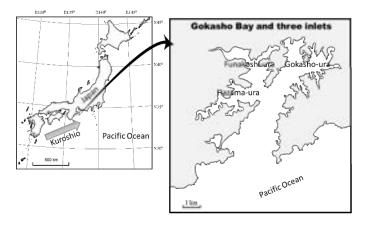


Figure 1. Map of Go asho ay.

Aquaculture is flourishing in Gokasho bay. Pearl aquaculture used to be the main industry in this area but has decreased significantly (Yokoyama et al., 1 6). Now, the aquaculture of red seabream (*Pagrus major*) in Hazamaura-Cove, Pacific bluefin

tuna (*Thunnus orientalis*) in the main bay, and oysters (*Crassostrea gigas* and *C. nippona*) and green alga (*Monostroma nitidum*) in areas close to the shore are operating in the bay.

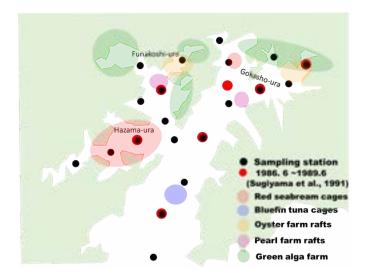


Figure 2. Sampling stations in Go asho ay. lac dots show the sampling stations from ovember 2016 to October 201 (this study), and Red dots from June 1 6 to June 1 (Sugiyama et al., 1 1). Major a uaculture area in the bay is also shown.

Sam ling an anal sis

Monthly hydrographic surveys in Gokasho Bay were conducted from November 2016 to October 2018 from the research vessel *amabou*. Seawater samples were collected in the bay at three depth layers using a 1.3-L itahara's water sampler at 1 stations from November 2016 to une 2017 and 20 stations from uly 2017 to October 2018 (Fig. 2). Samples were collected in 2 -ml co-stoppered glass test tubes for measurement of pH to avoid the contamination of CO₂ from the atmosphere and in 2 0 ml polyethylene bottles for measurements of total alkalinity. A CTD system (RIN O profiler, F Advantech) was also used to obtain hydrographic data (Watanabe et al., 201). Dissolved oxygen (DO) concentration obtained from the CTD system was calibrated in the laboratory with seawater analysis data from a DO meter (MM-60R, TOA-D) with dissolved

The pH of seawater was measured more precisely by potentiometric method using a pH meter

oxygen electrodes (O -470AA, TOA- D

(MM-60R, D -TOA Co.) with a glass electrode (LP-0320, D -TOA Co.). The glass electrode was standardized against the IS buffer scale (4.008, 6.86 and .180 pH at 2) which adapted the NBS buffer scale. Any drift in pH was corrected against the seawater buffer scale (Ono et al., 1 8;

of pH between NBS and total scales, the systematic error between these scales was ignored. To calculate carbonate systems, total alkalinity was measured from May 2016 to October 2018 using a

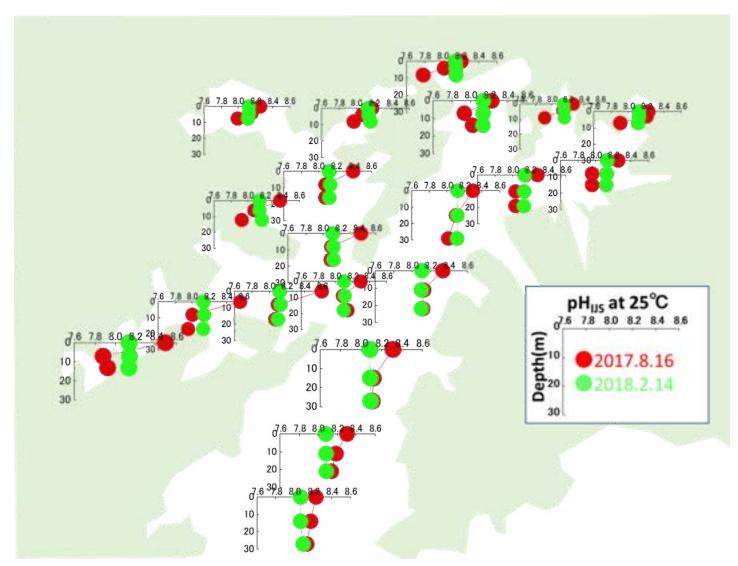


Figure . Vertical profile of p at each station on August 201 and February 201 in Go asho ay, Mie, Japan.

Dickson et al, 2007) and the precision of pH was 0.003. For measurement of pH, different pH scales were used. It is reported that the pH of IS (NBS) buffer scale are about 0.1 pH higher than that of the total scale (Gattuso and Hansson, 2011). In this study, because the large variation of pH value observed was much higher than the systematic error

one-point titration method (Culberson et al., 1 70; Tsunogai and Noriki, 1 83; Ono et al., 1 8) modified by using a small sample method (Suga et al., 2013). An aliquot (1 ml) of seawater sample was used to measure total alkalinity. In this research, systematic error of total alkalinity values was not corrected. The precision of total alkalinity was 0.006 meq L.

Calci m car onate sat ration state

Calcium carbonate saturation state of seawater () is one of the indexes for health of the ocean calcareous organisms. was calculated using the equation as follows;

$$\Omega_{cal_{,}arag} = \frac{[Ca^{2}][CO_{3}^{2}]}{K_{sp}^{cal'arag}}...(1)$$

where Ca²⁺ is the calcium concentration calculated from salinity (Millero, 2013), CO₃²⁻ is the carbonate concentration determined from observed pH and TA (Dickson at al., 2007), and sp^{cal,arago} is the solubility product for calcite and aragonite, calculated from salinity, temperature and water pressure (Mucci, 1 83; Millero, 1 83), respectively.

RESU TS AND DISCUSSION

An example of vertical profile of pH was shown at each station in August 2017 and February 2018 (Fig. 3). High pH of more than 8.3 was observed for surface water in August 2017 while pH decreased toward the sea bottom to less than 8.0.

ertical variation of pH tended to be greater at stations closer to the shore. However, pH in winter (February 2018) stayed constant around 8.1 because the bay water was homogenized due to winter vertical and horizontal mixing.

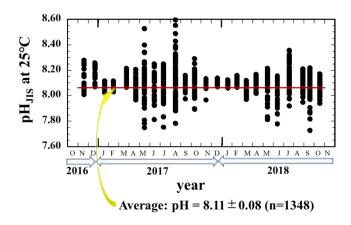


Figure . Monthly trend of p at all data from ovember 2016 to October 201 in Go asho ay, Mie, Japan. The average of p value is also shown.

Monthly trends of pH are shown in Fig. 4 from November 2016 to October 2018. pH values with a large variation ranging from 7.7 to 8.6 were found from spring to summer seasons, whereas the variation was relatively small, centered around 8.1, from autumn to winter. During phytoplankton blooms in surface water, particularly high pH of around 8.6 was observed in August 2017. Low pH values were seen in the suboxic bottom water during these periods. Seasonal average pH was calculated to be 8.11 0.08 (1, n 1384). This level was similar to those reported for apan's coastal waters and the western North Pacific surface water (awano, 2010; Ishii et al., 2011; apan Meteorological Agency).

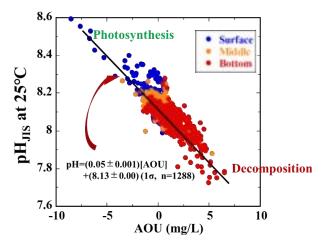


Figure 5. Relationship between AOU and p from ovember 2016 to October 201, e cept the data on January 201, in Go asho ay, Mie, Japan. The regression line is also shown.

Large seasonal variations of pH and dissolved oxygen levels come from biogeochemical and physical processes in the bay water. The relationship between pH and apparent oxygen consumption (AOU) at all stations are shown in Fig. . In general, pH is affected by photosynthesis and decomposition of organic matter. When photosynthesis is active, CO2 is taken up from the seawater to synthesize organic matter and to release oxygen. As a result, AOU decreases, and pH increases in the water column. On the contrary, decomposition of organic matter consumes oxygen and releases CO2 to seawater. As a result, AOU rises, and pH decreases. The observed correlation between AOU and pH suggests that the large seasonal variation of pH was at least partially caused by photosynthesis in the surface layer and decomposition of organic matter near the bottom.

Marine phytoplankton have a stoichiometric composition of carbon to oxygen (Anderson 1 The slope of the regression line between pH and AOU was calculated to be -0.0 0.001 pH (mg L) (or -0.0016 0.00003 pH (mol L)) (1, n 1288). The pH of surface water is dominated by photosynthesis, decomposition of organic matter and uptake of atmospheric CO2. When AOU is in condition of 0 mg L in sea water, the pH is apparently affected by only uptake of atmospheric CO₂. Therefore, the vertical intercept of the regression line can be defined as the acidification degree in the bay water under the conditions of absence of biochemical processes and AOU being 0 mg L. In Fig. , the pH of acidification degree was calculated to be 8.13 0.00. These results showed that the pH in the bay water changed with the AOU at the ratio of - 0.0 0.001 pH (mg L) around pH of 8.13.

A similar study was conducted in the bay from une 1 86 to May 1 8 by Sugiyama et al. (1 1).

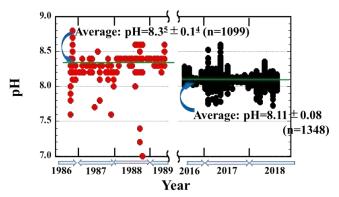


Figure 6 Comparison with p values between the trend of p from June 1 6 to ovember 1 (Sugiyama et al., 1 1) and ovember 2016 to October 201 (this wor) in Go asho ay, Mie, Japan. Since the p data from 1 6 to 1 was measured to be up to 2 digits after decimal point, the overall uncertainty of the p could be regarded to be 0.01 p units. Each average value of p is also shown.

Compared with the seasonal variation of pH values at present, the seasonal variation reported from 1 86 to 1 8 was larger, ranging from 7.0 to 8.8 (Fig. 6). If the ratio of pH AOU was constant in the bay water, the seasonal variation of pH could have also been larger than that of AOU at present. This suggests that the production and decomposition of organic matter were more active in the 1 80s than now. The average value of pH from 1 86 to 1 8 was calculated to be 8.3 0.14, which was 0.24 0.16 higher than the average value of 8.11 0.08 from 2016 to 2018. Assuming the pH decreased by 0.24 in the 30 years from 1 88 to 2018, the pH reduction rate is calculated to be -0.08 0.00 pH decade. This reduction value is greater than -0.02 0.007 pH decade of the south west coast apan (Ishii et al., 2011) or 0.0082 0.00 7 pH decade of the shelf break of the ast China Sea (Lui et al., 201). This suggests that OA may be progressing more rapidly in Gokasho Bay, though a longer-term data set is required to be more conclusive.

The monthly trends of omega calcite (cal) and aragonite (arago) were calculated for all data from May 2017 to October 2018 (Fig. 7). In summer, high value was observed in surface water and decreased with depth. The vertical variation of tended to be greater at the stations closer to the shore.

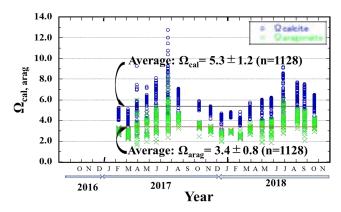


Figure . Monthly trends of calcium carbonate saturation states of seawater for calcite (cal) and aragonite (arago) from May 201 to October 201, e cept the data on October 201, in Go asaho ay, Mie, Japan. Each average value of calcium carbonate saturations is also shown.

On the other hand, hardly changed vertically nor horizontally in winter because the water was well mixed. These characteristics of were similar to those of pH. Under thermodynamic conditions of carbonate chemistry, when CO₂ dissolves in seawater, pH decreases and the following chemical reaction is taking place:

$$H_2O + CO_2 + CO_3^{2-} \rightleftharpoons 2HCO_3^{-} \cdot \cdot \cdot \cdot (2)$$

CO₂ reacts with CO₃²⁻ to form HCO₃⁻ in seawater and as a result, CO3²-concentration decreases. On the other hand, when CO2 is taken up by photosynthesis, pH increases and CO32- is supported by dissociation of HCO3². In that case, CO3² concentration increases. Since is a function of CO₃²-concentration as shown in equation (1), changes with a change of CO2 concentration and pH in seawater. That was a reason why the characteristics of were similar to those of pH. These results suggest that the seasonal variations of cal and arago were affected by photosynthesis in surface water and decomposition of organic matter in bottom water. The lowest values of cal and were found to be 2.4 and 1., respectively. The average cal and arago was calculated to be .3 1.2 and 3.4 0.8 (1, n 1128), respectively. These results show that the bay water is supersaturated for calcite and aragonite at present, although ocean

acidification may be advancing slowly in the bay. If the reduction rate of pH remains at 0.08 0.00 pH decade, will be less than 1 in 0 years. In the near future, the degree of in the bay water would drop with reduction of pH and could bring negative impacts to the calcareous organisms of the bay.

AC NOW ED EMENTS

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Ishii, M., N. osugi, D. Sasano, S. Saito, T. Midorikawa, and H.Y. Inoue. 2011. Ocean acidification off the south coast of apan: A result from time series observation of CO₂ parameters from 1 4 to 2008. . Geophys. Res. 116, C06022. apan Metrological Agency. Long- term trend of pH in surface water of the Pacific Ocean.

http://www.data.ma.go.pkaiyou.shindan.a.go
pHpac pH-pac.html

The authors determined that a significant trend in ocean acidification is superposed on the large seasonal and interannual variabilities of acidity in surface waters off the south coast of Honshu Island. apan, based on their observation of the carbonate system (partial pressure of CO₂, total inorganic carbon and pH). Multiple regression analysis of these parameters showed that pH and the aragonite saturation state (arago) decrease at a rate of -0.020 0.007 pH decade, and -0.12 0.0 arago decade, respectively. If future atmospheric CO2 levels keep increasing as predicted by IPCC scenario A1FI, which postulates tentative fossil fuel use associated with very rapid economic growth, a further reduction of -0.8 pH and -0.1 arago is likely in the next 0 years. The authors suggest that such a rapid reduction of arago could have negative impacts on a variety of calcareous organisms.

ubota, ., Y. Yokoyama, T. Ishiakwa, A. Suzuki, and M. Ishii. 2017. Rapid decline in pH of coral calcification fluid due to incorporation of anthropogenic CO₂. Nature, 7, 76 4, DOI:10.1038 s41 8-017-07680-0.

with the pH of the ambient seawater (pHsw) near these islands, estimated from a large number of shipboard measurements of seawater CO₂ and atmospheric CO₂, indicated that pHcF is sensitive to change in pHsw. The authors suggested that the calcification fluid of corals will become less supersaturated with respect to aragonite by the middle of this century (pHcF 8.3 when pHsw 8.0 in 20 0), earlier than previously expected, despite the pHcF-unregulating mechanism of corals.

Suga, H.P, S. Sakai, T. Toyofuku, and N. Ohkouchi. 2013. A simplified method for determination of total alkalinity in seawater based on the small sample one-point titration method. AMST C Rep. Res. Dev., 17, 23-33.

Measurement of pH and total alkalinity is the most convenient way to describe the carbon system in a seawater sample. In this paper, the authors introduced a modified method for determination of total alkalinity focused on reduced volume seawater samples (1 mL) based on the one-point titration method. The precision of this method is 0.1 - 0.2% (relative standard deviation). On the other hand, with no correction, the precision was 0.1 - 1.0%.

ven though values showed a systematic error, precision was maintained through the correction based on concurrent measurement of commercially available standards. This method allows us to describe the carbonate system in a small amount of water with sufficient precision and accuracy to investigate the mechanism of calcification using only a pH meter.

Fu ii, T., Y. omai, and T. Fu iwara. 2001. Carbon dioxide dynamics in coastal regions of Osaka Bay. apan Sci. Civil ng. Papers B2, 67, 11-1.

In coastal regions, the biological production is far greater than that of the open ocean and the short-term change of hydrographical conditions is significant. The authors conducted continuous measurements of salinity, pH, and dissolved oxygen

(DO) at three stations in Osaka Bay. Also, they calculated the values of carbonate species in the bay water by using the values of pH and total alkalinity. The results showed that dissolved inorganic carbon and DO fluctuated synchronously with high correlation (R² 0. 7), and the records of DO and

partial pressure of CO₂ (pCO₂) indicated the prominent diurnal variations which correspond to diurnal irradiation variations due to high productivity in the bay.

Ecological Interactions Among Wil life an Shellfish arms: A Case St from Ne erse USA on orseshoe Cra s an O ster arms

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: Oyster aquaculture, *Crassostrea virginica*, horseshoe crab, *Limulus polyphemus*, wildlife interactions

ABSTRACT

Globally, shellfish aquaculture is growing rapidly. For the first time in history, global seafood supply from aquaculture exceeded that of wild fisheries. Although shellfish culture is viewed as an ecologically sound industry, as farm production expands it faces key challenges in assuring ecological and social sustainability. Understanding the ways shellfish farms interact with coastal wildlife, particularly species of special concern, is among those challenges.

Farms for astern oyster (Crassostrea virginica) are commonly located in shallow coastal areas along the U.S. east coast, use a range of farm equipment, and involve regular access to care for and harvest livestock. In some cases, these farms are located in areas used by American horseshoe crabs (Limulus polyphemus) as they come ashore during spring to spawn along Atlantic Coast beaches. The sandy shores of the Delaware Bay, bordered by the states of New ersey to the north and Delaware to the south, host the largest spawning aggregations of the species in the world. Horseshoe crabs are economically and ecologically important; their blood is highly valuable in the medical industry and migratory shorebirds including the red knot (Calidris canutus rufa), a threatened migratory shorebird, feed on crab eggs when stopping in Delaware Bay.

Limited studies have been done to examine the

interactions among horseshoe crabs and intertidal oyster farms, and recently concern has been raised about the horseshoe crab's ability to traverse oyster farms to reach spawning habitat. A case study examining potential farm interactions with horseshoe crabs was done in Delaware Bay, New ersey, during the 2018 crab spawning season. The investigations included a range of controlled experiments and surveys during high and low tide using sonar and walking respectively, to observe crab behavior at farm and non-farm sites. In all cases, results indicate that crabs can successfully traverse farms and reach spawning beaches, and crabs do not differentially use farm versus non-farm areas of the Delaware Bay mudflats. These results provide important context for developing frameworks for managing ecological interactions among farms and wildlife species of concern.

INTRODUCTION

For the first time in history, global seafood supply from aquaculture exceeded that of wild fisheries (FAO, 2018). As aquaculture continues to develop globally (FAO, 2018), as well as domestically in the United States (National Marine Fisheries Service, 2018), the industry faces key challenges assuring ecological and social sustainability (Billing 2018). Understanding the ways shellfish farms interact with coastal wildlife, particularly species of special concern, is among those challenges (Barrett et al., 2018; Collier et al., 2018). The ways in which molluscan aquaculture interacts with fundamental

ecosystem processes such as particle depletion, nutrient cycling, and benthic-pelagic coupling has been relatively well studied (Newell 2004, Dumbauld et al. 200, Rose et al. 201). Likewise, attractiveness of farms in marine habitats, such as fish net pens and shellfish farm structures, to mobile fish and crustaceans is well documented (Callier et al, 2017). However, central to sustainability is appreciating the nature of the interaction among farms and wildlife that may use habitat near to or occupied by farms (Price et al., 2017; Barrett et al., 2018), interactions that are as of yet poorly studied.

Shellfish aquaculture in the Delaware Bay, Cape Shore region of New ersey has a long history as a low impact, sustainable food production system (Hilborn et al., 2018; van der Schatte Olivier et al. 2018). Oyster farms currently occupy approximately 10 acres in this area of New ersey and produce over 1.8 million market-sized oysters annually (Calvo, 2016); however, during the first half of the 20th century, oyster farming was much more expansive, using large wooden intertidal racks to cultivate oysters over wide stretches of this region (Ford and Haskin 1 82). The vast intertidal mudflats of the lower bay, with rich oyster food resources, make this region a desirable location to grow high quality oysters; it is also a region used by important wildlife species.

Horseshoe crabs (*Limulus polyphemus*) are an economically, medically (Novitski, 200) and ecologically important species. nown to be the largest spawning aggregations in the world (Shuster and Botton, 1 8), hundreds of thousands come ashore during the spring to mate and lay eggs along sandy beaches of the Delaware Bay (Smith et al., 2002). The rufa subspecies of the red knot (Calidris canutus rufa), a migratory shorebird that uses stopover areas along the Atlantic coast of the U.S., including the Delaware Bay, was designated threatened' in 201 under the ndangered Species Act in the U.S. by the U.S. Fish and Wildlife Service (0 C.F.R. 17 2014). An important food source for red knots are lipid-rich horseshoe crab eggs, deposited on beaches by mating crabs during

the spring northward bird migration (Castro Myers, 1 3). The eggs become available to the transitory bird flocks as they are exhumed from nests by sediment disturbance such as crab burrowing and wave action (raeuter Fegley, 1 4; Smith, 2007) and become concentrated in the upper intertidal zone as the tide rises (Nordstrom et al. 2006).

A small portion of the total Delaware Bay shoreline is collectively used by crabs for spawning. This is a nexus of shorebird migratory stopover habitat, and is home to oyster farming. The overlap in both time and space of these two iconic and ecologically important species, and oyster farm activities, presents a unique opportunity to examine potential ecological interaction among wild species stocks



Figure 1 Photo of a typical oyster rac with bag on top at a Cape Shore farm. orseshoe crabs can be seen among the rac s and in the flooding tide waters. Crab spawning habitat (sandy beach slope) can be seen in the bac ground.

and shellfish farms. Little data has been collected specifically addressing the ability of horseshoe crabs to traverse intertidal rack-and-bag oyster farms to reach their spawning habitat (Figure 1). If crab spawning migrations are impeded by farms, it is possible that crab populations themselves may suffer, and in turn shorebird foraging patterns and opportunities may be altered. In this case study, our primary goal is to characterize the ways that horseshoe crabs interact with farm structures while passing through farms en route to inshore spawning habitat, and determine if the farms themselves present a barrier to crabs reaching beaches to lay eggs. Ultimately, this information is important for both wildlife species conservation and sustainable farm management.

MET ODS

All experiments and surveys were carried out during the horseshoe crab spawning season (Smith and Michels 2006; Shuster and Botton 1 8), from May through uly of 2018, at the Rutgers University Cape Shore Laboratory and nearby oyster farms located along the lower Delaware Bay. First, an experiment was conducted in which crabs were placed in a tank, along with oyster farm equipment (rack-and-bag) to observe the ways that crabs are, or are not, able to move past this equipment. Second, surveys were conducted on the flats during low and high tide conditions to assess crab presence absence relative to farm footprints, and to evaluate whether crabs use intertidal habitats differently when farm gear is present.

For the observation of crab movement around racks, a large fiberglass tank (3.7 m long x 1. m wide x 0. 1 m deep) was filled with filtered baywater to a depth of 30 cm. Mature horseshoe crabs were collected by hand from the ad acent Cape Shore mudflats. xperiments began May , 2018, and continued through une 21, 2018, during the spawning period for horseshoe crabs; thus, all animals used in the experiment were actively coming ashore to spawn and assumed mature. Twenty crabs, collected at random, were used for

each experimental trial. Before being placed into the tank, each crab was measured (widest distance across the prosoma) and sexed, then assigned an identification number that was written on both sides of the shell in yellow waterproof crayon (this mark was removable and was wiped off before returning these crabs to the flats after the experiment). The size, sex, and id number of each crab was recorded and marked animals were placed into the tank and allowed to acclimate for 1 minutes before the experimental treatment was placed in the tank with them. During this acclimation period, many of the females in the tank were attached to by males, forming amplexus pairs.

A total of eleven treatments were tested. Treatments included a control, in which the footprint of an ovster rack was drawn on the tank bottom but no physical structure was put in the tank, and 10 farm gear treatments of varying heights and configurations. The suite of treatment types included three rack heights (3, and 8 above the bottom) with and without oyster bags attached, an oyster bag on the bottom of the tank (no rack), a floating oyster bag tethered to the bottom with braided sinking line, oyster bags leaning on the side of a rack and a rack on its side (no bag). All racks were rebar racks that are used by farmers, had elastic cords with metal hooks attached (farmers use these to hold bags onto the racks), and bags contained oyster shells to mimic adult oysters. The designated treatment was placed in the tank with 20 crabs and a timer started. The crabs were observed continuously for 1 minutes as they moved about the tank and interacted with the treatment structure. A note was recorded for every crab each time that individual crab passed to the side, beneath, or over a structure. This 1 minute observation period was considered one replicate for the given treatment. ach treatment was replicated a minimum of nine, and maximum of 14 times, for a total of 13 trials.

Low tide surveys were conducted using paired transects, laid out and marked on four intertidal farms in the Cape Shore region. At each farm, paired transects (1m wide) were oriented

perpendicular to the shore with one transect intersecting a farm, and a parallel control transect passing through ad acent un- farmed intertidal habitat following methods described in Munroe et al. (2017). During daytime low tides, starting on May 8, 2018, and continuing through May 2, 2018, transects on all four farms were walked and all crabs encountered along the transect were documented, and their location (inshore of or within farm) was noted. Walks were repeated a minimum of weekly, for a total of nine repeat surveys during the 2018 spawning season. A ruskal-Wallis test was used to test if differences were observed in the number of crabs observed inshore, and within farm gear among paired farm:control transects.

Waters in the Delaware Bay are sufficiently turbid, that even in shallow water and over short distances, visual observation of crabs on the bottom during high tide is not possible. Therefore, specialized sonar (DIDSON sonar) mounted to the front of a small aluminum boat was used to observe crabs on the bottom during high tide. Sonar video was recorded as the boat motored slowly along parallel transects set in oyster farm and non-farm habitats, laid out as described for the low tide surveys. In addition to the transect videos, sonar video was taken at two paired mooring locations for durations of 10-1 minutes. One mooring point allowed observation within a farm; the other in a comparable control location with no farm gear. A total of nine paired mooring videos were taken.

RESU TS

In total, 330 male and 12 female crabs were used in the experimental trials, ranging in size from 16 to 28 cm prosoma width for males (mean 20.1 cm), and from 21 to 30. cm for females (mean 2.8 cm). Across all eleven rack treatments tested, all categories of crabs (single male, single female and amplexus pairs) were observed moving around and under over through the farm gear treatment. Interestingly, this includes single female crabs and crabs engaged in amplexus successfully passing both under and over the two shortest rack heights.

In zero of the 128 trials involving oyster racks and or bags (i.e. all non-control treatments) was a crab observed to be stuck or impeded from moving past or through the oyster gear.

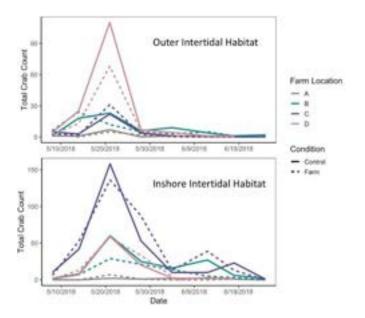


Figure 2 Crab counts along transects made during low tide. Upper panel shows crabs observed within the outer intertidal region, among the farm gear. Lower panel shows crabs observed inshore of the farm gear. ote that the length of transects varies among farms, and differs between outer and inshore habitats. Line colors depict each of the four farms, and dashed lines are counts along the transect bisecting the farm, solid lines show counts made along the control transect.

Numbers of crabs observed during the low tide surveys varied through the season, and among farms (Figure 2). Counts ranged from a low of zero at the beginning and end of the observation period, to a high of 13 per transect during the observation made on May 21st, 2018; this period fell between the New and Full moon in late May. In total, over all transects and across the entire observation period, 1,176 crabs were observed on the four farms, zero of which were impinged in farm gear. No difference exists among the number of crabs counted at habitat inshore of farm gear compared to controls (ruskall-Wallace test, p 0.73), nor at habitat within the farm footprint compared to controls (ruskall-Wallace test, p 0.4). The numbers of crabs counted during low tide differed significantly among farm sites (ruskall-Wallace

test, p 0.001 inshore habitat, p 0.004 within farm habitat).

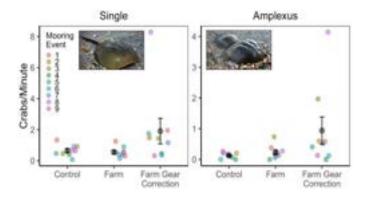


Figure Single and ample us crab counts made using sonar from all paired mooring events. Gray points represent the mean standard deviation. Colored points show data per observation event, with each event colored the same.

Sonar video surveys of crabs during high tide showed large aggregations of crabs in sloughs (muddy depressions), and sparsely distributed crabs moving independent of other crabs in seemingly random directions outside of sloughs. On many occasions, crabs were observed to move under and out from farm gear unimpeded. The number of crabs observed during high tide along transects varied depending on the day of observation, the farm, and whether counts were inshore of the farm gear versus in the area of the farm gear. For the region inshore of the farm gear, no significant difference was detected among control and farm counts for single crabs (p 0. 4) nor amplexus pairs (p 0. 3). In the intertidal area within the footprint of the farm gear (outer intertidal), no significant difference was detected among control and farm counts made for single or amplexus crabs, (all p values 0.13). Counts of crabs made during mooring events, standardized by effort to crabs minute, ranged from zero to nearly two crabs per minute (Figure 3). No significant difference was detected among paired control and farm counts for single crabs (paired T-test, p 0.37), nor amplexus pairs (Wilcoxon signed rank, p 0.33). When counts were corrected for view obstruction by farm gear, no significant difference among single crab counts at control vs. farm was detected (Wilcoxon signed rank, p 0.16); however, significantly more

amplexus pairs were observed at the farm mooring (Wilcoxon signed rank, p = 0.0).

DISCUSSION

xperiments and surveys conducted during the horseshoe crab spawning season in 2018 tested the ability of horseshoe crabs to move among and through oyster farm gear and evaluated the ways that crabs use habitat around oyster farms in the Delaware Bay region. Across all of the surveys and experiments, our results indicate that crabs can successfully traverse farms and reach spawning beaches, and that crabs do not avoid farm gear when accessing spawning beaches. These results provide important context for developing frameworks for ecological interactions among farms and wildlife species of concern, specifically with respect to concerns about the potential for changes in abundance of horseshoe crabs or their behavior due to the presence of oyster farm gear.

Mature crabs ranging in size and including single male and female crabs, as well as amplexus pairs, were observed to move beneath, over and around oyster racks ranging in height from three to eight inches off the bottom when underwater. Likewise, sonar data collected during high tide at farm and control locations on the Cape Shore intertidal flats documented crabs (both single crabs and amplexus pairs) moving unobstructed among farm gear in daylight and moonlight conditions. Crabs were able to access habitat inshore of farms at the same rate as those inshore of comparable control sites, suggesting that the presence of farms does not impede crabs from accessing upper intertidal spawning sites; consequently, the reproductive ability of the crabs is not impacted by oyster farms. When sonar counts of crabs at farm sites were corrected for obstruction of the view of the bottom, there in fact appears to be more crabs moving within farms at high tide compared to control sites without gear. Attractiveness of farm structures in marine habitats, such as fish net pens and shellfish gear, to mobile fish is well documented (Callier et al. 2017). It is possible that horseshoe crabs also

find oyster farm gear attractive due to increased foraging opportunities, shelter or other cues; future research may address this possibility. Concepts of conservation aquaculture highlight the important contributions shellfish farms and other forms of aquaculture can make in achieving local and global conservation goals (Froehlich et al 2017). When evaluating aquaculture impacts and interactions with wildlife, it is important to (1) establish appropriate reference sites at which to compare wildlife abundance to farms, and (2) collect data on potential differences in behavior and reproductive capacity for wildlife at farm sites (Barrett et al, 2018). In the studies documented in this report, appropriate controls were used in an effort to best illuminate changes, if any exist, in abundance of horseshoe crabs or their behavior due to the presence of oyster farm gear, with the intention to serve the needs of an adaptive management process aimed at balancing wildlife species conservation and sustainable farm management. Adaptive approaches that allow for inclusion of new data are critical to coastal management, and particularly so when considering appropriate ecosystem-based decisions that ensure protection of wildlife and viability of important coastal industries.

AC NOW ED EMENTS

I am indebted to the fearless field crew that helped count crabs at all hours of day and night, and in particular to hard-working interns osh Daw and Niki Cleary who were instrumental in data collection and analysis. I extend further gratitude for the continued cooperation of the oyster farmers of Delaware Bay who allow us access to their farms. Support for part of this pro ect was provided by Cape May County. This white paper is the result of research sponsored by the New ersey Sea Grant Consortium (N SGC) with funds from the National Oceanic and Atmospheric Administration (NOAA) Office of Sea Grant, U.S. Department of Commerce, under NOAA grant Award NA10OAR417008 and the N SGC. The statements, findings, conclusions, and

recommendations are those of the author and do not necessarily reflect the views of the N SGC or the U.S. Department of Commerce. N SG-1 - 64.

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ANNOTATED BIB IO RA O E WOR S

In this annotated bibliography, I have selected recent works that focus on interactions among wildlife - in particular horseshoe crabs or *abutogani* - and shellfish farms. Limited research has focused on this issue; therefore, I have included two experimental studies, and two broader reviews. Horseshoe crabs are an important and iconic species along the eastern United States, as well as in apan.

Barrett, Luke T., Stephen . Swearer, and Tim Dempster. 2018. Impacts of marine and freshwater aquaculture on wildlife: a global meta-analysis. Reviews in Aquaculture. 1-23. doi: 10.1111 raq.12277

In this review, authors from the University of Melbourne in Australia demonstrate evidence that supports higher wildlife biomass and diversity found around aquaculture farms. They argue that to properly understand the possible impacts of farms on wildlife, studies have thus far failed to address issues of impacts on behavior, reproduction or fitness. If animals that tend to aggregate around farms suffer poor fitness relative to those occupying non-farm habitats, the farm becomes an ecological trap'.

Callier, M.D., C. . Byron, D.A. Bengtson, P. .
Cranford, S.F. Cross, U. Focken, H.M. ansen,
P. amermans, A. iessling, T. Landry, and F.
O beirn. 2017. Attraction and repulsion of
mobile wild organisms to finfish and shellfish
aquaculture: a review. Reviews in
Aquaculture.

The ways in which finfish and shellfish farms act as attractant or repulsive areas for mobile marine species is reviewed. At finfish farms, food inputs and physical structure of the farm tend to drive aggregating effects. Likewise, at shellfish farms, the farm structure (both farm gear and shellfish themselves) and increased feeding opportunities via biodeposition, the shellfish crop or fouling organisms tend to attract wildlife to farms. In general, the authors note that a great deal of variability in attractiveness or repulsiveness exists when considering the suites of habitats, farms, and mobile fauna that have been studied thus far.

wan, Billy .Y., Chan Hoi in, and Siu Gin

Cheung. 2018. Habitat use of globally threatened uvenile Chinese horseshoe crab, *Tachypleus tridentatus* under the influence of simulated intertidal oyster culture structures in Hong ong. Aquatic Conservation: Marine and Freshwater cosystems. 28: 124-132.

The authors conducted this study to test if uvenile Chinese horseshoe crabs differ in abundance and if their feeding trails differ at oyster farm spat collection locations. They performed their test at a shallow mudflat in Hong ong which is home to a large population of uvenile horseshoe crabs. Addition of oyster spat collectors (bricks on the mudflat) lead to a significant reduction in the number of uvenile horseshoe crabs, and a reduction in the length of feeding trails.

Munroe, D., D. Bushek, P. Woodruff, and L. Calvo. 2017. Intertidal rack-and-bag oyster farms have limited interaction with Horseshoe Crab activity in New ersey, USA. Aquaculture nvironment Interactions. : 20 -211. https://doi.org/10.33/4/aei00227

In this study, the authors sought to assess whether intertidal oyster farms altered the distribution of mature horseshoe crabs as they came ashore to spawn in the lower Delaware Bay. In total, they used a combination of field surveys of crabs on mudflats and stranded along the upper shore, and controlled tests to determine if crabs can move past farm gear. Results show no evidence that horseshoe crab distribution within this area is altered by the presence of oyster farms.

Use of oint of ie i eo Cameras to Assess ish Interactions ith O ster A ac It re Cages

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: Aquaculture, Shellfish, Habitat, cosystem Services, eDNA

ABSTRACT

Oyster bottom cages are an increasingly popular style of aquaculture gear in the Northeastern United States. Cages are used to contain and protect shellfish during grow-out but also likely provide three- dimensional structure that may benefit the local wild fish community. Understanding the ecosystem services provided by aquaculture gear may help to inform regulatory decisions and inform a broader public discussion of shellfish farming and how farms interact with the local environment.

A low-cost camera mounting system was developed and tested to allow visual assessment of fish activity near an oyster cage farm and in a surrounding low seafloor structure environment. Minimal-structure t-platform stands were created to mount cameras ad acent to boulders, enabling comparison of fish interactions with cages to fish activity observed on natural structured rock reef habitat. Seawater samples were collected monthly during pilot camera deployments for environmental DNA (eDNA) analysis to detect fish species that may be present but not observed in video.

ideo collected by cameras attached to oyster cages and near boulders facilitated visualization and quantification of the wild fish communities along with documentation of behavioral interactions between animals and these habitats. Analysis of eDNA detected significant differences in fish community composition across the three months sampled, and significant differences between sites in two of the three months sampled.

*The Federal Government does not endorse the use of GoPro cameras

INTRODUCTION

Bottom cages are an increasingly common type of aquaculture gear for cultivating oysters in the Northeastern United States (Archer et al, 2014; Flimlin et al, 2010). Cages are used to contain and protect shellfish during grow-out but also provide three-dimensional structure that may attract the local wild fish community and provide habitat services (Dumbauld et. al, 200). A trapping and mark- recapture study in Narragansett Bay, Rhode Island, found the seafloor with oyster grow-out cages provided valuable habitat for uvenile and adult temperate reef fish (Tallman and Forrester, 2011). Habitat offered by oyster aquaculture gear may provide food resources, shelter, refuge from currents, and protection from predation for commercially and recreationally important species

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of fish. Documenting and quantifying the ecosystem services provided by aquaculture gear may help inform regulatory decisions and inform a broader public discussion about shellfish farming and how farms interact with the local environment.

Several structure-oriented species common to Long Island Sound are known to support important recreational and commercial fisheries in the Northeast. Demersal species including Black Sea Bass and Scup are federally managed under the Summer Flounder, Scup, and Black Sea Bass Management Plan for New ngland stocks (Atlantic States Marine Fisheries Commission, 2018). Tautog is a temperate reef fish also highly valued by fishermen, with 0% of landings contributed from recreational harvest (Atlantic States Marine Fisheries Commission, 201). An understanding of habitat preferences and use, particularly among species of commercial and recreational interest, may aid in regional management decisions related to aquaculture leasing.

Small self-contained point-of-view video cameras can enable fine spatial-scale studies to quantify the interactions between fish and oyster aquaculture gear. Here we report on a pilot study using these cameras and other commercially-available accessories to create an optimized and transferable system for the deployment of underwater video cameras across multiple sampling periods and geographic study locations. The information gained from this pilot program is presented in this report. The ultimate goal of our research program is to utilize this system to gain information on species composition and abundance of fish interacting with oyster cages, to quantify and document details of how fish species interact with oyster cages, to

compare how the habitat services provided by oyster cages compare with those provided by natural structured habitats, and to pilot the use of eDNA metabarcoding to characterize finfish communities associated with our study habitats over time and space.

MET ODS

We studied a style of commercial oyster bottom cage commonly used in the Northeast (Figure 1). Cages measure 4 x 3 x 2 ft (1.2 x 0. x 0.6 m) and consisted of three shelves, with two bags of oysters per shelf. We stocked each bag with 1 0 seed oysters (1 in; 2. cm) based on information from industry partners. Bricks were added to the base of the cage for ballast. Cages were deployed from May-September, 2017, at two nearshore sites off Milford, Connecticut; one ad acent to a high-density cage farm, and one in a low-structure environment. Water depths at the study sites ranged from 1 -20 feet at high tide.



Figure 1. Oyster cage used in this study. Full view showing three shelves, each with two bags of oysters. Feet of cages show added bric s for ballast.

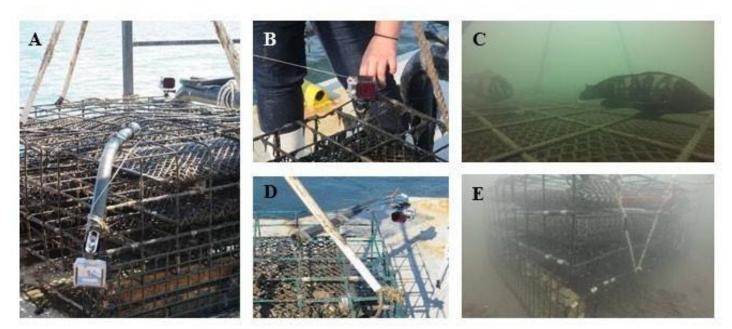


Figure 2. On bottom cage with camera views. (A) Close up of cage fitted with cameras, showing side view in the front of the photo and top camera at the rear. () Close up of top camera loo ing across the hori ontal surface of the cage. (C) The resulting view of the top camera. (D) View of the side camera loo ing at two sides of the cage and where the cage meets the sea floor. (E) Resulting view of the side camera.

We experimented with several methods for attaching point-of-view (GoPro) cameras to oyster cages. Initially a spring attachment was placed at a corner of the cage to provide flexibility to protect cameras during deployment and retrieval, but unfortunately resulted in too much camera motion during deployment and retrieval of cages. Camera mounts were instead constructed with flexible marine hose and inch P C, which provided stability during video recording but allowed some flexibility for camera protection. We found a combination of two time-synced cameras to provide adequate visual coverage of most of the cage and immediate surroundings. The first camera was affixed to a cage corner by 2 feet (0.6 m) of 7 8' flex marine wet exhaust and water hose, allowing the camera to hang off the side of the cage, enabling a view of two cage sides as well as the area where the cage feet met the seafloor. We initially experimented with a second camera positioned to look down at the top of the cage, but the poor water clarity prevented viewing the entire top of the cage. The final configuration had the second camera

positioned in a periscope like mount, which was able to capture the top horizontal surface of the cage, including viewing the top of the oyster bags proximal to the camera. (Figure 2).

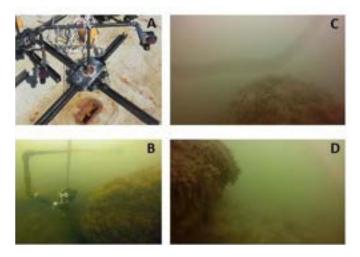


Figure . T platform construction to mimic camera views on natural structured habitat. (A) Full view of t platform with cameras attached. () Image of t platform deployed on a natural roc reef adjacent to boulder. (C) Resulting top view of the boulder. (D) Resulting side view of the boulder which also captures where the boulder and sea floor meet.

To compare fish interactions with cages to interactions with natural structured rock reef habitat, t-platform stands were devised to mount cameras ad acent to boulders while adding minimal additional structure (Figure 3). T-platforms were

built using threaded metal pipe and were designed to provide a field of view similar to that on the cage, capturing the top of the boulder and one side where the boulder meets the seafloor. The platforms are intended to be deployed at the beginning of the field season and left in place, using divers to attach and remove cameras. This should minimize the disturbance associated with camera deployment retrieval, and ensure consistent fields of view across multiple deployments.

Intervalometer Blink timers (CamDo, ancouver, Canada) were used with GoPro cameras to delay the start of video recording by 24-hours, to minimize any deployment-related disturbance prior to recording. ideo was recorded for eight minutes every hour from 7AM to 7PM to cover a complete tidal cycle and most daylight hours. A Polar Pro magenta filter, with a 0. stop reduction in exposure, was affixed to each camera lens to reduce green coloration of video footage caused by the natural attenuation of light in water, as well as phytoplankton growth in Long Island Sound. The addition of any filter reduces total light capture by the cameras, which limited the depth of our deployments, but the importance of this deployment restriction was outweighed by the large reduction in time required for video post-processing. One TCM-1 tilt current meter (Lowell Instruments LLC, North Falmouth, MA, USA) was deployed at each site to measure current speed and direction during trials. HOBO pendant temperature and light meters (Onset Computer Corporation, Bourne, MA, USA) were also deployed to document light penetration and seawater temperature.

nvironmental DNA (eDNA) analysis was used to detect fish species that may have been present but not observed on video. Following camera retrieval, seawater was collected near the seafloor using a Niskin Bottle during one deployment each in the months of une, uly, and August 2017. Samples were kept on ice until filtration (0.4 m) and then frozen (-20 C) until analysis. DNA was extracted and PCR was performed on the 12S mitochondrial rDNA region. Next generation sequencing was then

conducted on PCR amplicons, followed by bioinformatics analysis.

Pilot-scale deployments over the May-September 2017 field season yielded 20+ hours of video on weeks with the cage density comparisons. During weeks where t-platforms were deployed, 18+ hours of video was recorded on a continuous recording scheme. ideo was downloaded immediately following trips and duplicate copies stored in two separate locations for data management. ach file was renamed with a unique identification to ensure proper organization. Observer T software (v14.0; Noldus Information Technology, Wageningen, Netherlands) was used to analyze videos. Advantages of this software platform included multi-video playback, inter-coder analysis and flexible coding scheme, making it an ideal tool for our analysis.

ideo analysis is currently underway. Fish abundance will be reported using the MaxN metric (Watson et al. 200). The likelihood of doublecounting the same fish within each interval of video recording was increased by the typical behavior of fish associated with a structured environment. The MaxN metric avoids double-counting by limiting abundance to the maximum number of fish observed at any one time, and is thus likely a conservative estimate of total fish abundance associated with a structure. For our study, MaxN was defined as the maximum number of individuals of a given species present in a single frame within each 1-minute segment of video (Watson et al. 200). MaxN was calculated separately for each species observed. A behavior matrix under development will quantify and describe specific behaviors observed in video records. Precise definitions allow distinct behaviors to be identified and reduce the sub ectivity in behavior analysis and identification among coders. For example, the current working definition for station keeping above the cage is small fin movements to maintain a position both in a period of foraging and while in the water column or above cage.

RESU TS AND DISCUSSION

ideo analysis is ongoing, so only initial observations are reported here. The four most frequently observed fish species thus far were: cunner (Tautogolabrus adspersus), black sea bass (Centropristis striata), tautog (Tautoga onitis) and scup (Stenotomus chrysops), which are all commonly found in and around structurally complex habitat in Long Island Sound. These four species have been observed both on boulders and on oyster cages. Other species of fish captured on video include: butterfish (Peprilus triacanthus), banded rudderfish (Seriola onata), hake (Urophycis chuss), sea robin (Prionotus carolinus), striped bass (Morone sa atilis), summer flounder (Paralichthys dentatus), windowpane flounder (Scophthalmus a uosus), smallmouth flounder (Etropus microstomus) and yellow ack (Carangoides bartholomaei). Animals observed inside cages during camera deployment or retrieval, but not in video, include: naked goby (Gobiosoma bosci), oyster toadfish (Opsanus tau), conger eel (Conger sp.), and rock gunnel (Pholis gunnellus). Some of these species are more active at night, which may account for their absence from video to date. Invertebrates associated with cages (either in video or observed during deployment retrievals) include oyster predators such as oyster drill and sea star, as well as prey species for fish such as black-fingered mud crab. Other invertebrate species noted ad acent to a cage in video include: spider crab, blue mussel, slipper shell, horseshoe crab, blue crab, and channel whelk.

The results of eDNA metabarcoding efforts yielded 17 ma or and 20 rare finfish species from samples collected monthly during une, uly, and August deployments in 2017. Fish communities obtained by eDNA metabarcoding were significantly different across the three months, and communities between sites were different during the months of uly and August (Liu et al. 201). eDNA was detected for all of the fish species observed thus far in the videos. Using ordination analysis, the axis associated with sampling month represented the highest variation

(73.4%) among samples while the axis associated with habitat type represented minor variation (1 .6%) among samples.

Outreach is an important component of this pro ect. Public perception and understanding of aquaculture practices can be enhanced by news articles and other types of outreach (Froehlich *et al.* 2017). Heightening awareness of aquaculture practices and positive outcomes associated with aquaculture gear can increase support for aquaculture in the scope of the growing demand for shellfish. Our methodology was based on low-cost, readily-available cameras and mounts. An advantage of this design is that it can easily be replicated by aquaculture industry members, extension specialists, and citizen scientists. We have provided our methods in detail, with photos, within a .pdf document that is publicly available online

(https: www.fisheries.noaa.gov webdam download 8 62100). Our intention is to broaden video collection to other geographic locations, shellfish aquaculture industry practices, and seasons, to ultimately enable a more comprehensive understanding of the habitat services provided by shellfish aquaculture. Selected video clips collected from oyster cages are posted on our pro ect website (https: www.fisheries.noaa.gov aquaculture-habitat -N). There has also been traditional news and social media coverage of this pro ect via local news outlets as well as Twitter and Facebook coverage from both local and national NOAA Fisheries accounts.

This pro ect has yielded a low-cost, easily replicated system for observing fish activity associated with oyster aquaculture gear and natural structured habitat. ideo analysis generated during pilot-scale deployments will provide data on fish abundance and community composition in these two habitats, as well as insights into fish behavior. The continued investigation of fish interactions with aquaculture gear may help inform regulators, policy makers and fishery managers who make decisions about aquaculture practices. ideo generated by this study could additionally contribute to a broader public

understanding of interactions between shellfish aquaculture and the local environment.

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Watson D.L., .S. Harvey, M. . Anderson, and G.A. endrick. 200 . A comparison of temperate reef fish assemblages recorded by three underwater stereo-video techniques. Marine Biology 144: 41 -42 .

The authors deployed underwater stereo-video cameras to sample the relative density and species richness of temperate reef fishes in Southwest Australia. This study compared diver deploy, unbaited remote and baited remote cameras. They defined MaxN as species presence and the maximum number of individuals belonging to each species in the field of view at one time (MaxN).

This metric avoids repeated counting of fish within a given time frame and gives a conservative estimate of relative density relating to the area of survey.

DeAlteris .T., B.D. ilpatrick, and R.B. Rheault. 2004. A comparative evaluation of the habitat value of shellfish aquaculture gear, submerged aquatic vegetation and a non-vegetated seabed. ournal of Shellfish Research 23(3): 867-874.

The authors set out to evaluate habitat value of shellfish aquaculture gear (SAG), submerged aquatic vegetation (SA), and shallow non-vegetated seabed (N SB) over one year in Port udith, RI, USA. Oyster cages were sampled using lift nets that were diver deployed under oyster cages and left to soak for two weeks prior to retrieval. Submerged aquatic vegetation and non-vegetated seabed sites were sampled with quadrats, drop nets, and a venturi- driven suction dredge deployed from a skiff. Sessile invertebrate growth was documented on eelgrasses and on oysters and oyster cages. All mm were collected. The authors documented fish that the physical habitat of shellfish aquaculture gear had greater surface area compared with the other two habitat types as well as a significantly higher abundance and species richness of organisms per meter squared throughout the year. The authors concluded that shellfish aquaculture gear had greater habitat value compared to their control sites.

Tallman .C. and G. . Forrester. 2007. Oyster grow-out cages function as artificial reefs for temperate fishes. Transactions of the American Fisheries Society 136 (3): 7 0-7 .

The authors compared fish habitats within Narragansett Bay, RI, USA using trap surveys. They

compared three oyster grow-out sites, six natural rock reefs, and one artificial reef built for fish habitat, looking to identify patterns in fish density, growth and disappearance rates. Their traps were designed to sample both uvenile and adult fish and were deployed in the summer and fall. The study showed that oyster cages provide habitat for fish associated with hard bottom habitats including scup, tautog, and black sea bass. Dumbauld, B.R., .L. Ruesink, and S.S. Rumill. 200 . The ecological role of bivalve shellfish aquaculture in the estuarine environment: A review with application to oyster and clam culture in West Coast (USA) estuaries. Aquaculture 2 0:1 6-223.

This review article focuses on bivalve shellfish aquaculture on the US West Coast. Much like natural rock reefs, cages become colonized by epibenthic, emergent, and encrusting organisms and may provide sheltering fish with camouflage, refuge from predation, respite from high current flow, and a source of food. Multi-tiered off-bottom cages, an increasingly popular method for growing oysters on a smaller footprint, offer vertical three-dimensional structure and surface area similar to that afforded by natural reefs. In traditional on-bottom shellfish culture, where live oysters and shell cultch are distributed on sediments for grow-out and spat collection, oysters act as ecosystem engineers creating hard bottom irregular substrate on otherwise featureless seafloor. valuation of habitat services provided by oyster aquaculture can be challenging. Shellfish farms vary in spatial size and bottom area covered, variety and quantity of aquaculture gear, and have a seasonally variable footprint as gear is relocated or harvested. Aquaculture of oysters, whether in cages or grown on the seafloor, may represent valuable estuarine habitat.

i ri c ra ii ticus In ction of ir lence To ar arval O sters an Corals at Elevate Sea ater Tem erat res an otential Mitigation Strategies

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: Vibrio coralliilyticus, oyster, larvae, coral, mortalities

ABSTRACT

Recent research has advanced our understanding of the role pathogenic vibrios play in disease of various aquaculture species. Vibrio corallilyticus, formerly thought to infect ust corals, is now known to infect larval oysters, causing ma or losses in hatchery settings. The negative effect of V. corallilyticus on wild-type oyster production remains uncertain, but circumstantial evidence suggests it may be substantial. We identified eight strains of *V. corallilyticus* that infect both astern oyster (Crassostrea virginica) larvae and Pacific oyster (Crassostrea gigas) larvae in U.S. ast and Pacific coast hatcheries. West Coast hatcheries have experienced ma or production losses over the years due to the presence of *V. corallilyticus*. Some of the largest hatcheries have reported losses of up to 80%, resulting in shortages in seed oysters needed for commercial oyster production. Losses can exceed 10 billion larvae per year in the larger hatcheries. Most U.S. ast Coast hatcheries are considerably smaller, but also experience losses in astern oysters, possibly from V. corallilyticus, but also from the known shellfish pathogen V. tubiashii. Interestingly, V. tubiashii only infected astern oyster larvae, unlike V. coralliilyticus which causes acute mortalities with astern and Pacific species.

The infection and death of corals by *V. corallilyticus* is known to be enhanced at seawater

temperatures 27 C. In our studies, multiple strains of *V. corallilyticus* produced high larval oyster mortalities with LD o's ranging from 3.8 10³ to 4 10^4 CFU ml of seawater, depending on the V. corallilyticus strain and the oyster species (astern vs. Pacific oysters). Studies with knockout mutations demonstrated that the transcriptional regulator ToxR and the outer membrane protein OmpU were important in larval oyster (and coral) infections caused by V. corallilyticus. Gross pathological changes occur in V. corallilyticus-infected larvae, initially in the velum and cilia and then in the internal organs, which become liquefied. Infection of larval oysters is likely induced by stress and a lowering of resistance to these pathogens. Stressors, particularly in hatcheries, likely include incorrectly ad usted seawater temperatures, salinities, pHs, and dissolved oxygen levels or inadequate nutrition, over stocking, high overall bacterial loads, etc.

To begin to mitigate vibrios in hatcheries, we identified a variety of predatory bacteria which in nature help to modulate *Vibrio* levels in seawater and or shellfish. They include several predatory bacteria such as *Pseudoalteromonas piscicida*, which secrete digestive enzymes that inhibit and kill many kinds of bacteria including vibrios; and *alobacteriovora*, a *dellovibrio* and Like Organism (BALO), which infect Gram-negative bacteria, like vibrios, and replicate within them, killing the vibrios in the process. Together these

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bacteria may serve as probiotics in reducing hatchery mortalities and in killing vibrios and other bacterial pathogens in a variety of other aquaculture settings. Bacteriophages (phages) are another means to inactivate many undesirable pathogens in the environment, and are increasingly being used in aquaculture. To date, we isolated multiple phages against *V. corallilyticus* and *V. tubiashii* and showed the practical application of phage therapy to reduce or eliminate mortalities from *V. corallilyticus* and *V. tubiashii* in larval oysters.

INTRODUCTION

Naturally-occurring *Vibrio corallilyticus* is a wellrecognized pathogen of some corals. More recently, it has been found to be highly pathogenic toward larval Pacific oysters (Crassostrea gigas) and astern oysters (Crassostrea virginica) (Richards et al., 201; Ushi ima et al., 2018). These two oyster species are the primary commercially-harvested oysters in the United States. Another pathogen of larval astern oysters is Vibrio tubiashii, which is better known for disease outbreaks in ast Coast hatcheries. Although literature has commonly referred to *V. tubiashii* as a pathogen in West Coast shellfish hatcheries, some of the more prevalent strains associated with the mortalities were misidentified and were confirmed by genomic sequencing to be *V. corallilyticus* (Richards et al., 2014, 2018). Together, these two vibrios have contributed to episodic losses in hatchery production and shortages in seed oysters needed for commercial shellfish aquaculture. Interestingly, V. corallilyticus and V. tubiashii are only known to infect and kill oysters while in the larval stage, which is only two to three weeks in most cases. Once the larvae undergo metamorphosis, they appear resistant to infection by these vibrios.

Methods are needed to mitigate larval oyster diseases in shellfish hatcheries. Several potential probiotic approaches include the use of predatory bacteria, like *alobacteriovora* (Richards et al., 2012, 2013, 2016) and *Pseudoalteromonas piscicida* (Richards et al., 2017a, 2017b). In

addition, bacteriophages (phages) against both *V. corallilyticus* and *V. tubiashii* offer some promise as a therapeutic treatment for reducing hatchery-associated larval mortalities. Phage therapy is increasingly being used to combat bacterial disease in aquaculture (reviewed by Richards 2014).

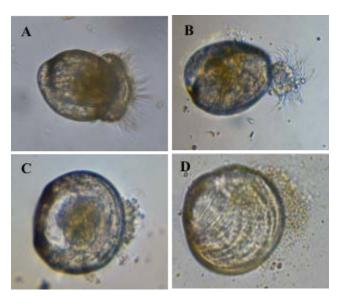


Figure 1. ormal and diseased Pacific oyster larvae. A) ormal, healthy larva and (D), larvae infected with V. corallilyticus.) Early stage infected larva with tangled cilia and lumpy velum. C) Later stage infection with loss of cilia and velar cells. D) Final stage showing dead larva and release of digested/li uified tissues from within the valves.

We conducted studies of larval oysters to compare disease pathologies associated with both *V. coralliilyticus* and *V. tubiashii* (Ushi ima et al., 2018). We identified the infectious doses of these pathogens in larval oysters. It is well known that elevated seawater temperatures enhance coral disease from *V. coralliilyticus*, so we also determined if similar high temperatures enhanced larval oyster mortalities. We also explored genes potentially responsible for the virulence of *V. coralliilyticus* in knock-out experiments (Ushi ima et al., 2018) and are currently investigating mitigation strategies to reduce larval oyster mortalities using bacteriophages and predatory bacteria.

RESU TS AND DISCUSSION

Disease atholog

Normal, uninfected, Pacific oyster larvae had a smooth velum and ordered cilia (Fig. 1A). They remained actively motile. Larvae infected with *V. coralliilyticus* showed tangled cilia and a lumpylooking velum within a day or two post infection (Fig. 2B), which greatly restricted their motility. Shortly thereafter, the cilia were either digested or released into the surrounding milieu while individual velar cells were also released into the milieu. Usually within a day, the tissues within the valves appeared to liquefy by microbial digestion and were released from the valves (Fig. 1C).

astern oysters exhibited the same pathology from *V. coralliilyticus*. Pathological signs of *V. tubiashii* infection were the same as for *V. coralliilyticus* infection in astern oysters, but *V. tubiashii* did not infect larval Pacific oysters in our studies.

Table 1. Increased larval oyster mortality from four V. corallilyticus strains due to elevated seawater temperatures

Vibrio coralliilyticus			
ATCC BAA- 4 0	23.4	38.8	1 .4%
OCN008	4.	78.8	23. %
OCN014	0.	3.7	42.8%
R 8	3.3	7 .2	2.%

Infectio s Dose of *c ra ii ticus* an *tu iashii* in Eastern an acific O sters

Our studies showed that the dose of pathogens causing 0% mortality (the LD 0) for both astern and Pacific oysters varied somewhat based on the strain of *V. corallilyticus* used. In general, LD 0's ranged from approx. 1 10³ to 1 10⁴ CFU ml of

seawater. In the case of *V. tubiashii*, LD ₀'s for strains ATCC 1 106 and ATCC 1 10 were 3.8 10^3 and 1.2 10^4 CFU ml, respectively, in astern oysters; however, these strains were not infectious to Pacific oysters. These ATCC strains are obtained from the American Type Culture Collection, Manassas, irginia (USA).

Effects of Elevate Sea ater Tem erat re on arval Mortalities

We also evaluated and determined that elevated seawater temperatures (27 C) increased larval oyster mortalities by 1 .4% to 42.8%, depending on the *V. corallilyticus* strain, compared to larval oysters maintained at 23 C (Table 1).

ir lence enes in c ra ii ticus

We also evaluated the ToxR transcriptional regulator and the outer membrane protein OmpU to determine if they were virulence factors in V. corallilyticus infection of larval Pacific oysters. Studies with knockout mutations demonstrated that ToxR and OmpU in V. corallilyticus were important in larval oyster (and coral) infections (Table 2). Mutants lacking ToxR had significantly reduced virulence compared to the wild type strain, while the mutant lacking OmpU had completely attenuated virulence at doses 10⁶ CFU ml. Larvae exposed to the OmpU mutant did not experience significantly more mortalities compared to replicates exposed to filtered sea water or a non-pathogenic bacterium, suggesting OmpU is essential for V. corallillyticus infection of Pacific oyster larvae. Furthermore, both the ToxR and OmpU mutant are avirulent towards coral, suggesting these proteins play similar roles for infections of multiple hosts (Ushi ima et al. 2016, 2018). In contrast, the mannose-sensitive hemagglutinin (MSHA) type I pili that are required for coral infection (Ushi ima et al. 2016) were not required for larval oyster infection (Ushi ima et al. 2018).

The protein ToxR is believed to positively respond to various environmental signals, including elevated water temperatures, which is then involved with upregulating the expression of various genes, including the gene encoding OmpU. Therefore, ToxR is not believed to be directly involved with shellfish mortalities, however, represents a molecular messenger that translates environmental signals, most of which have yet to be identified, to physiological responses. Hence, understanding the conditions conducive to increased *V. coralliilyticus* virulence (i.e. the regulators of virulence) would complement, and potentially enhance, the various mitigation efforts described here.

Table 2. Larval mortalities after deletion () mutation of various genes believed to be important for V. coralliilyticus virulence.

Vibrio coralliilyticus				
Wild type	4.	78.8		Yes
to R mutant	1 .8	2 .4	Yes	Yes
ompU mutant	4.	6.1	Yes	No
MSHA mutant	44.1	72.4	No	Yes

Mitigation atcheries Strategies for c ra ii ticus in atcheries

We have conducted considerable research over the past 10 years to isolate, characterize and test the efficacy of phages against *V. corallilyticus* and *V. tubiashii* strains in astern and Pacific oyster larvae. From over 20 phages isolated and characterized, several have been shown to be lytic phages (based on genomic sequencing). Such lytic phages are desirable for use in mitigation efforts because their replication cycle consistently results in host death. Currently, we have phages against eight strains of *V. corallilyticus*, including the principal strains associated with hatchery outbreaks on the West Coast (Table 3), and against *V. tubiashii* strain ATCC 1 106. Proof-of-principle testing has shown

that these phages can be combined in a cocktail to provide a safe treatment for *V. corallilyticus* in hatcheries. Testing of a preliminary phage cocktail at a West coast hatchery at Oregon State University showed nearly a 100% reduction in mortalities after challenging larval oysters with a lethal dose of V. corallilyticus. ast coast hatchery trials are planned in the spring. The goal is to market a phage cocktail for use in hatcheries world-wide. Probiotic means to eliminate V. coralliilyticus, V. tubiashii and other pathogens in hatcheries are also being explored using the predatory bacteria alobacteriovora (a genus of dellovibrio and Like Organisms BALO's) and Pseudoalteromonas piscicida. Both predators have broad host specificity, with the alobacteriovora targeting Gram-negative bacteria and P. piscicida targeting both Gram-positive and Gram-negative bacteria.

Table . Eight strains of V. corallilyticus illed by phage treatment.

Vibrio coralliilyticus	Vibrio coralliilyticus	
atccATCC BAA-4 0*	R 8	
ATCC 1 10 *	R 0 -10 -8	
R 22	OCN008	
R 0	OCN014	

^{*}Indicates strains known to infect both corals and oyster larvae

CONC USION

There are many obstacles to successful hatchery operations. Since *V. coralliilyticus* and *V. tubiashii* appear to be opportunistic pathogens, anything that stresses larval shellfish makes them more susceptible to infection. High seawater temperature is one factor that enhances *V. coralliilyticus* virulence and may be managed by maintaining temperatures under 27 C in the hatcheries.

Likewise, close monitoring of other hatchery parameters (like salinity, dissolved oxygen, pH, food supply, larval densities, etc.) and adequate hatchery sanitation will reduce larval stresses to enhance productivity. With new treatments on the horizon to combat vibrios, the future is looking brighter for hatcheries and the commercial oyster industry as a whole.

AC NOW ED EMENTS

The authors thank Ralph lston, Aquatechnics, Sequim, WA, for the R strains of *V. corallilyticus*; Chris Langdon and David Madison, Oregon State University, Hatfield Marine Science Center, Newport, OR, for phage efficacy testing; Alexander Sulakvelidze and acob Chamblee, Intralytix Inc., Baltimore, MD, for upscaling and preparing the phage cocktail; the staff of the Aquaculture Innovation Center, Rutgers University, Cape May, N, for larval astern oysters; oan Hendricks, Taylor Shellfish Farms, uilcene, WA, for larval Pacific oysters; eith Olson, Natural nergy Laboratory of Hawaii Authority, ailua- ona, HI, for seawater; Ronald Lau, ona Coast Shellfish LLC, ailua- ona, HI, for seawater and larvae; Claudia H se, Oregon State University, Corvallis, OR, for technical assistance, and Michael Watson, USDA, ARS, Dover, D, for conducting many of the assays.

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Richards, G.P., .P. Fay, .A. Dickens, M.A. Parent, D.S. Soroka, and .F. Boyd. 2012. Predatory bacteria as natural modulators of *Vibrio parahaemolyticus* and *Vibrio vulnificus* in seawater and shellfish. Appl. nviron. Microbiol. 78:74 –7466.

Levels of the human pathogenic Vibrio parahaemolyticus and Vibrio vulnificus are increasing in shellfish harvesting areas in many parts of the world due, in part, to elevated seawater temperatures. A means to reduce these vibrios in shellfish is needed. When seawater and oysters were spiked with *V. parahaemolyticus* or *V. vulnificus*, the Vibrio levels rapidly diminished over a 3-day period when natural seawater was used, but counts quickly climbed when sterile seawater was used. In an effort to determine what was inhibiting the vibrios in natural seawater, tests showed the presence of Vibrio predatory bacteria, later classified as alobacteriovora species. The alobacteriovora in natural seawater readily eliminated vibrios that were added to the seawater. In contrast, oysters spiked with vibrios showed some initial increase in Vibrio counts within the first 24 h, followed by a rapid decline to baseline levels. Thus, it appears that alobacteriovora are one of nature's tools to modulate bacterial levels in shellfish. Harnessing this tool may provide a means to reduce or eliminate pathogens in aquaculture operations and in the seafood industry.

Richards, G.P., M.A. Watson, D.S. Needleman, .M. Church, and C.C. H se. 201 .

Mortalities of astern and Pacific oyster

larvae by the pathogens *Vibrio coralliilyticus* and *Vibrio tubiashii*. Appl. nviron. Microbiol. 81:2 2–2 7.

Vibrio tubiashii has been a well-known pathogen in astern oyster (Crassostrea virginica) larvae on the United States (U.S.) ast Coast for many years and has also been associated with Pacific oyster (Crassostrea gigas) larvae on the U.S. West Coast. Vibrio coralliilyticus, a well-known coral pathogen and a cause of coral bleaching, has also been associated with disease of Pacific oyster larvae, particularly in ovster hatcheries. It became evident that some of the outbreaks on the West Coast that were thought to be from *V. tubiashii* were actually caused by *V. corallilyticus*. In this study, we evaluated two strains of *V. tubiashii* and four strains of *V. corallilyticus* in both astern and Pacific oyster larvae to determine their ability to cause larval mortalities. We showed that *V. tubiashi*i strains infected primarily the astern oyster larvae while all four of the *V. corallilyticus* strains infected the astern and Pacific oyster larvae. The LD ovalues for the *V. corallilyticus* strains ranged from 1.1 10⁴ to 3 10⁴ CFU ml of seawater in astern oyster larvae. In Pacific oyster larvae, the LD orange was 1.2 10⁴ to 4 10⁴ CFU ml of seawater. Together, these studies defined the host specificity of these Vibrio strains with regard to astern and Pacific oyster larvae and demonstrated the ability of coral- associated pathogens to also infect and kill larval oysters. Subsequent research (Ushi ima et al., 2018, see abstract below) showed similar results with other strains of *V. corallilyticus* and an up-regulation of mortalities at elevated seawater temperatures.

Richards, G.P., .P. Fay, . Uknalis, M. Olanya and M.A. Watson. 2016. Purification and host specificity of predatory *alobacteriovora* isolates from seawater. Appl. nviron. Microbiol. 82: 22–27.

alobacteriovora are small predatory bacteria found in the marine environment and may serve as biocontrol agents against human, fish and shellfish pathogens. They are within a group of predatory

bacteria known as the *dellovibrio* and Like Organisms (BALO). alobacteriovora were isolated from seawater and had broad specificity toward five strains of Vibrio parahaemolyticus, while two additional alobacteriovora strains isolated from low salinity seawater readily infected and killed the human pathogens E. coli O1 7:H7 and Salmonella Typhimurium DT104. The replication rate of alobacteriovora against E. coli and Salmonella increased as salinities decreased. It is likely that alobacteriovora could readily be isolated against a broad range of Gram-negative seafood pathogens. Improved methods were also developed to purify infectious alobacteriovora from their host cells. Overall, the use of alobacteriovora in various aquaculture applications may serve as an environmentally friendly, non-antibiotic treatment to reduce pathogens in fish and shellfish.

Richards, G.P., M.A. Watson, D.S. Needleman, . Uknalis, .F. Boyd, and .P. Fay. 2017. Mechanisms for *Pseudoalteromonas* piscicida- induced killing of vibrios and other bacterial pathogens. Appl. nviron. Microbiol. 83:e0017 -17, https://doi.10.1128/10.1128/10.117

Pseudoalteromonas are marine bacteria that are known to secrete antimicrobial compounds which inhibit competing bacteria in the marine environment. We identified a second method by which some Pseudoalteromonas (P. piscicida) kill competing bacteria. It involves the direct transfer of digestive vesicles from the surface of the Pseudoalteromonas to the surface of competitors, digestion of holes in the competitor's cell wall by proteolytic enzymes associated with the vesicles, and apparent feeding of the Pseudoalteromonas off the nutrients released by the digested bacterium in a predatory fashion. Among the P. piscicida enzymes

identified were aminopeptidase B, a trypsin-like serine protease, a chymotrypsin-like serine protease, and a cysteine protease. *Pseudoalteromonas piscicida* inhibited and killed the pathogens *Vibrio parahaemolyticus, V. vulnificus, V. cholerae, Photobacterium damselae, Shewanella algae*, and *Staphylococcus aureus*. Together, this data indicates that *Pseudoalteromonas piscicida* produce important antibacterial compounds that have a potential role in the probiotic treatment of aquaculture products and in reducing biofilm formation.

Ushi ima, B., G.P. Richards, M.A. Watson, C.B. Schubiger, and C.C. H se. 2018. Factors affecting infection of corals and larval oysters by *Vibrio coralliilyticus*. PLoS One. https://doi.org/10.1371/0urnal.pone.01/47

Vibrio corallilyticus is a naturally occurring marine bacterium which infects and kills corals and larval shellfish. It is best known as the cause of coral bleaching, which has contributed to the loss or damage to coral reefs worldwide. We determined that *V. corallilyticus* strains also infect and kill larval oysters and other shellfish, particularly in shellfish hatcheries, causing shortages of seed oysters needed for commercial shellfish operations. We determined that larval oyster mortalities were significantly higher at an elevated seawater temperature (27 C) compared to a lower temperature (23 C). Thus, multiple coral pathogens were found to infect larval oysters in a temperatureand dose-dependent manner. Also identified were virulence factors that promote the infection of both coral and ovster larvae. This work demonstrates for the first time that elevated seawater temperatures enable *V. corallilyticus* to more readily infect oyster larvae. It also serves to warn hatchery operators to maintain seawater temperatures below 27 C.

C st Distri tion atterns of the aral tic Shellfish oisoning lan ton S ecies an rium cat n a an aci icum off the acific Coast of Eastern a an

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: paralytic shellfish poisoning, cyst, Ale andrium catenella, Ale andrium pacificum

ABSTRACT

Mainly two dinoflagellate plankton species, Ale andrium catenella and A. pacificum, cause paralytic shellfish poisoning in apan. To examine how the two Ale andrium species' cysts distribute off the Pacific coast of eastern apan, we measured cyst abundance of the two species by quantifying their specific gene abundance in DNA extracted from marine sediments. Both species were detected in some ria-type inner bays in the Sanriku region. However, only A. catenella was detected from the oban area, which is located to the south of Sanriku region, opening widely to the Pacific Ocean. In addition, only A. pacificum was detected in the Sotobo area, which is located to the south of the oban area and is under the strong influence of the warm uroshio current. These results suggest that the distribution patterns of the two toxic phytoplankters differ depending on topography and the ocean current system.

INTRODUCTION

Paralytic shellfish poisoning (PSP) is caused by eating bivalves contaminated with a marine biotoxin that is produced by microalgae. In apan, it is a serious problem for fishermen because many bivalves cause PSP, and a voluntary shipment regulation is carried out when a PSP is detected

from their products. PSP occurs in the north-east Pacific coast and western part of apan, Seto inland sea, and yushu area. However, PSP does not occur in the middle parts of the Pacific coastal area and apan sea coastal area except in a few inner bays. Moreover, PSP occurrence has been shown to increase suddenly, for example, after the Great ast apan arthquake in 2011 (Ishikawa et al., 201). It is, therefore, essential to study the risks of PSP outbreaks in shellfish aquaculture areas without previous reports. One of the important precautions is monitoring of the toxin producing organisms. There are four PSP producing phytoplankton species in apan. Ale andrium catenella and A. pacificum are the ma or causative organisms. These phytoplankton species used to be called A. tamarense and A. catenella, respectively; however, recent molecular work has indicated a new nomenclature for species within the Ale andrium tamarense species complex (Litaker et al., 2018). A. catenella prefers cold water and often causes PSP in the northern areas of apan. On the other hand, A. pacificum prefers warm water. It is found mainly in the western part of apan, and also in north-eastern apan during summer. The toxicity level of A. pacificum is relatively low compared to A. catenella and is not generally considered to cause PSP in the north-eastern apan areas.

These phytoplankton produce a cyst after blooming

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stimulated by nutrient depletion. The germinating plankton become a seed population in the year of PSP outbreaks. To assess risks of PSP outbreaks, cyst abundances are often investigated in the area. The fluorescent dye method is commonly used to determine Ale andrium spp. cyst abundance (Yamaguchi et al., 1). However, it is difficult to determine whether the cysts belong to A. catenella or A. pacificum species (Nagai et al., 2012). There are some reports where molecular techniques have been used for identification of A. catenella and A. pacificum cysts (amikawa et al., 2007; rdner et al., 2010). In this study, we measured cyst abundances of A. catenella and A. pacificum by quantifying their specific gene abundance in the DNA extracted from marine sediments to examine how the cysts of the two Ale andrium species are distributed off the Pacific coast of eastern apan.

MET ODS

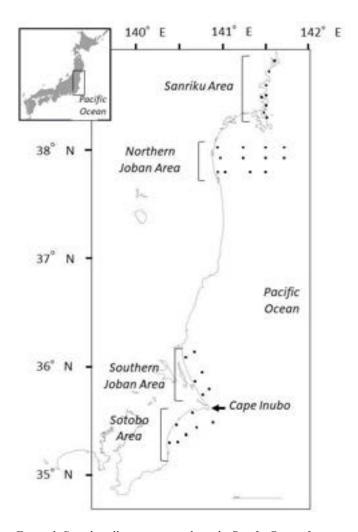


Figure 1. Sample collection points along the Pacific Coast of eastern Japan in 2015, located in the Sanri u rias coastal area, northern part of the Joban area, southern part of the Joban area, and the Sotobo area.

Sediment samples were collected from the Sanriku area, northern and southern parts of the oban area, and the Sotobo area (Fig. 1). The Sanriku area is a rias coast where mountains approach the sea and the coast line is very complex. The oban and Sotobo areas are geographically flat and are widely open to the Pacific. The Sanriku and oban areas are regions where the cold Oyashio current mixes with the warm uroshio current. The Sotobo area is mainly under the influence of the uroshio current. PSP generally occurs in the Sanriku and oban areas but not in the Sotobo area.

Sediment samples were collected using a grab or core type bottom sampler during the summer of 201 when *Ale andrium* vegetative cells were not observed. In the Sanriku area, samples were collected at 3-12 points for each examined bay. A surface layer sediment of approximately 2 cm depth was taken and kept in cold and dark conditions immediately after collection. Samples were preserved at -30 until DNA extraction.

DNA was extracted from marine sediments using a commercial kit, FAST DNA-SPIN kit for soil (MP Biomedicals). DNAs of *A. catenella* and *A. pacificum* were amplified by qPCR from the sediment DNA using specific primer sets reported by Nagai (2011). PCR reactions were performed using SYBR Premix x Taq II (Tli RNaseH Plus) (Takara Bio) with ABI StepOne System (Applied Biosystems). Plasmid DNA (pTaq2, Dyna xpress co. ltd) into which the PCR target fragment was inserted was used as a standard DNA. To estimate the density of cysts, one cyst was picked under the microscope, its DNA extracted, and the copy number determined using qPCR.

RESU TS

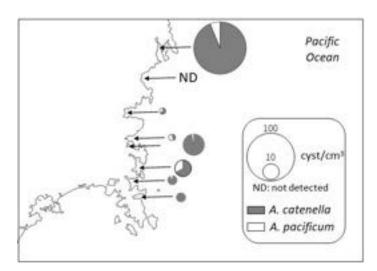


Figure 2. Cyst abundances of Ale andrium catenella and A. pacificum determined by PCR in sediments collected in the Sanri u rias coastal area

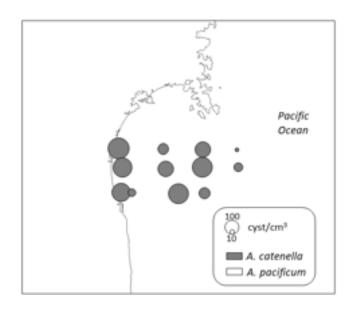


Figure . Cyst abundances of Ale andrium catenella and A. pacificum determined by PCR in sediments collected in the northern part of Joban area.

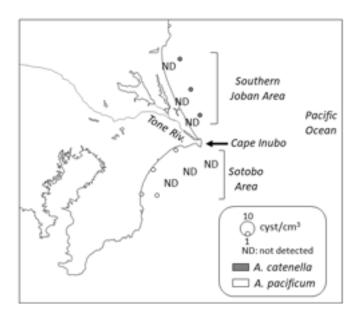


Figure Cyst abundances of Ale andrium catenella and A. pacificum determined by PCR in sediments collected in the southern part of Joban area and the Sotobo area.

In the Sanriku rias coastal area, both *A. catenella* and *A. pacificum* cysts were detected in the sediment samples, except in one small bay located

at the southernmost part of the area (Fig. 2). The proportion of *A. catenella* cysts was found to increase with the total cyst abundance. In the northern oban area, only *A. catenella* cysts were detected (Fig. 3). Cyst abundance was high in this area compared to the rias coastal area. Also in the southern part of the oban area, only *A. catenella* cysts were detected, although the density observed was very low (Fig. 4). In the Sotobo area, only *A. pacificum* cysts were detected and the density was as small as that observed for *A. catenella* cysts in the southern oban area.

DISCUSSION

Both Ale andrium species cysts were detected in the Sanriku rias coastal area, whereas only A. catenella was detected in the oban area. It is presumed that A. pacificum can survive in the Sanriku area because of high water temperature, which allows it to grow in summer in the small shallow bays. Although A. pacificum is not considered to cause PSP, recently PSP is found to occur during high temperature periods in the Sanriku area. The relationship between PSP occurrence and population dynamics of the two Ale andrium plankton should be studied, as these species concomitantly occur in some areas (Sakami unpublished data) and their vegetative cells have very similar forms. Therefore, it is important to monitor both species of PSP-causing phytoplankton to assess PSP outbreak risks in the Sanriku Rias areas, where cysts of both species are present.

In the oban area, only cysts of *A. catenella* were detected. A high cyst density of more than one hundred per square centimeter was observed at the offshore area, more than 0 km away from the coastline with a water depth greater than 100 m. This result suggests that not only coastal areas, but offshore areas should be investigated for the presence of PSP-causing plankton. The cyst density was found to be much higher in the northern part than in the southern part of the oban area. One of the reasons for the lower cyst density in the southern part is that the sediment in the southern

part is sand, in which cyst densities are generally low (Yamaguchi *et al.*, 2002; Shimada and Miyazono, 200; Yamamoto *et al.*, 200). In the past fifteen years, PSP occurred in the northern and southern part in thirteen years and seven years, respectively. Therefore, the difference in cyst abundance seems to indicate the difference in PSP outbreak risk in these areas.

In the Sotobo area, cysts of only A. pacificum were detected. The Sotobo area, located south to Cape Inubo, is under the strong influence of the strong and warm uroshio current which runs along apan's south coast. The lowest water temperature is to the north of Cape Inubo, and 14-1 the Sotobo area. The water temperature may be too high for A. catenella in the Sotobo area. There is a report that A. catenella cysts seldomly germinate at temperatures higher than 1 (Yamamoto et al., 200). In fact, the lowest water temperatures are in Hiroshima and Osaka Bays in around 10-13 the western part of apan, where A. catenella causes PSP. In addition, the river mouth of the Tone River. the largest river in apan, opens to the north of Cape Inubo. The large freshwater discharge can be a barrier for the distribution of the phytoplankton. Considering that PSP occurrence has not been reported in the Sotobo area, it seems that distribution of A. catenella is limited to the south of Cape Inubo.

In conclusion, we determined cyst abundances of two PSP-causing *Ale andrium* species by using a qPCR method from samples collected off the Pacific coast of eastern apan. We observed that: 1) both *A. catenella* and *A. pacificum* cysts are found in the Sanriku rias coastal area, 2) only *A. catenella* cysts are found in the oban flat coastal area, extending to the southern tip area where the warm uroshio current mixes with the cold Oyashio current and 3) only *A. pacificum* cysts are found in the Sotobo area under the strong influence of the uroshio current. These results suggest that the cyst distributions of the two species are influenced by the coastal geographic features, the water temperature and or the current systems in different

ways. Although more data are required to clarify relationships between cyst distribution and environmental factors, our study seems to indicate that cyst abundances and or fluctuation patterns are useful parameters for forecasting PSP outbreak risks in shellfish aquaculture areas.

AC NOW ED EMENTS

This study was supported financially in part by a grant of regulatory research pro ects for food safety, animal health and plant protection from the Ministry of Agriculture, Forestry, and Fisheries of apan.

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This study investigates the role of epigenetics (DNA methylation) in migration-related life history traits in *Oncorhynchus my iss*. The authors used reduced representation bisulfite sequencing to perform comparative DNA methylation analysis between uvenile resident and smolt F2 siblings

generated from a cross between steelhead (migratory) and rainbow trout (nonmigratory). Fifty-seven differentially methylated regions, many of which were in gene regulatory regions, were identified between residents and smolts, suggesting a relationship between epigenetic variation and variation in migration-related phenotypes.

Gavery, M.R. and S.B. Roberts. 2013. Predominant intragenic methylation is associated with gene expression characteristics in a bivalve mollusc. Peer 1: e21.

This dataset, generated using methylation-enriched high-throughput bisulfite sequencing, represents the first high-resolution methylome in any mollusc. DNA methylation data were compared to gene expression datasets and a positive relationship between intragenic methylation and gene expression levels was identified. These data suggest that DNA methylation patterns may play a role in regulating gene expression in molluscs.

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The authors report that in uropean sea bass (Dicentrarchus labra), which exhibits temperature-dependent sex determination, exposure to high temperature in early development was associated with increased DNA methylation in the promoter of the aromatase gene (cypl ala) and a higher proportion of phenotypic males. Furthermore, in vitro methylation of the aromatase promoter was sufficient to suppress transcription of the gene, supporting a role for DNA methylation as a mechanistic link between temperature and sex ratios in species exhibiting temperature-dependent sex determination.

Potok, M. ., D. A. Nix, T. . Parnell, and B. R. Cairns. 2013. Reprogramming the maternal

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Genome-wide DNA methylation patterns in zebrafish gametes, various stages of embryos and a somatic tissue (muscle) were analyzed using whole genome bisulfite sequencing. This high-resolution approach identified dynamic and unique patterns of DNA methylation during development in zebrafish. Results suggest that the functional significance of sperm DNA methylation patterns in fish is to provide transcriptional competency to the early embryo, which inherits' the DNA methylation pattern in the sperm.

Shao, C., . Li, S. Chen, P. hang, . Lian, et al. 2014. pigenetic modification and inheritance in sexual reversal of fish. Genome Research 24: 604-61.

The half-smooth tongue sole (*Cynoglossus* semilaevis) was used as a model to investigate the role of epigenetic regulation in species with environmental sex determination. Using genomewide bisulfite sequencing of normal male, female and pseudomale fish (generated by exposing genetic females to high temperature during a sensitive developmental window), the authors showed that sex-reversed genetic females exhibit methylation patterns consistent with genetic males, both of which differ from the methylome of normal females. Furthermore, it was reported that global methylation patterns are inherited by F1 pseudomale offspring generated by crosses between temperature-induced sex-reversed pseudomales and normal females, suggesting transgenerational epigenetic inheritance of sex reversal in this species.

Gavery, M.R. and S.B. Roberts. 2014. A context specific role for DNA methylation in bivalves. Briefings in Functional Genomics doi:10.10 3 bfgp elt0 4.

A review of current knowledge of DNA methylation in bivalves. A primary conclusion is

that the functional role of the gene could influence the role of DNA methylation in influencing expression.

Olson, C. . and S.B. Roberts. 2014. Indication of family-specific DNA methylation patterns in developing oysters. bioRxiv doi:

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This study provides the first single-base pair resolution DNA methylomes for both oyster sperm and larval samples from multiple crosses. While

sample sizes are very low, this work suggests DNA methylation patterns could be inherited.

Roberts, S. 201 . Compilation of DNA Methylation Genome Feature Tracks (*Crassostrea gigas*). figshare https://dx.doi.org/10.6084/m/.figshare.14/626/.v2

Genome feature tracks and accompanying IG session file to visualize DNA methylation data for the Pacific oyster (*Crassostrea gigas*).

Tren s in A ac It re ro ction in a an

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: demographic problems, earthquake, tsunami, declining industry

ABSTRACT

A wide variety of aquatic organisms are commercially cultured in apan. For marine species, national statistical data is published by the Ministry of Agriculture, Forestry and Fisheries on the aquaculture production of seven species of teleost fish (Oncorhynchus isutch, Seriola spp., Trachurus japonicus, Pseudocaran dente, Pagrus major, Paralichthys olivaceus, Tetraodontidae spp., Thunnus orientalis), two species of bivalves (Mi uhopecten vessoensis and Crassostrea spp.), one species of prawn (Marsupenaeus japonicus), one species of tunicate (alocynthia roret i), and four species of aquatic plants (Saccharina spp., *Undaria* spp., *Pyropia* spp. and *emacystus* spp.). Four freshwater species of teleost fish (Oncorhynchus my iss, Plecoglossus altivelis, Cyprinus carpio and Anguilla japonica) are also included in the statistics. Despite the historically strong affinity of apanese consumers to seafood, production of aquaculture and capture fisheries are on a long-term moderate decreasing trend largely due to socio-economic factors in apan. The total marine aquaculture production decreased from 1.28

10⁶ t in 1 6 to 1.03 10⁶ t in 2016. Bivalve aquaculture production was relatively constant around 4. 10 t. The production of oysters (*Crassostrea* spp., mainly Pacific oyster) and apanese scallop (*M. yessoensis*) was heavily damaged by the Great ast apan arthquake and subsequent tsunamis in 2011, and it has not recovered to the level prior to the earthquake. Teleost fish and seaweed production are on a

continuous decreasing trend. Teleost fish production was not severely affected by the earthquake, except for Coho Salmon (*Oncorhynchus isutch*), whose production level quickly recovered. Although the production of some seaweeds such as kelps (*Saccharina* spp.), wa ame (*Undaria* spp.) and laver or nori (*Pyropia* spp.) was devastated by the earthquake, not only in the Tohoku region but in many parts of apan, the long-term decreasing trend is not attributed to the earthquake. One of the causes for the production decline is considered to be the reduced number of management bodies due to the aging of the operators and shortage of successors.

The number of management bodies of *nori* culture, for instance, decreased from 1,3 4 in 1 63 to 3,81 in 2013. Mergers of the management bodies is one of the reasons for the decrease, but the number of farmers is certainly decreasing. On the other hand, aquaculture production of teleost fish and bivalves has increased in value in recent years. The increase in teleost fish production is attributable to the recent inclusion of Pacific bluefin tuna (T. orientalis), which was added to the MAFF statistics list in 2012. For bivalves, production of the common scallop leaped from PY 2 .7 10 in 2012 to PY 62.4 10 in 2016. In freshwater aquaculture, the apanese eel (A. japonica) has by far the largest production in volume and value. The seeds for eel aquaculture are all wild caught, and the low availability of glass eel has been problematic in recent years. Prompt establishment of mass production technologies for eel seed is desired.

INTRODUCTION

Being surrounded by the sea, fish and seafood have been staple foods for the apanese since ancient times. apan has a long history of aquaculture, which mythically dates back to the era of mperor eiko's (BC13 AD103, archeologically estimated to be in the 4th century) reign (Higurashi, 1 12). The mperor had a recreational fishing pond to which cultured fish were released for his amusement. Goldfish were introduced to Osaka from China in 1 02, and they have been cultured in apan ever since. Aquaculture of laver or *nori* (*Pyropia tenera*) is thought to have started in Shinagawa Bay, Tokyo, in the Bunki era (1684-1688) (Higurashi, 1 12).

The history of aquaculture in apan was summarized in a textbook (Tanigawa et al., 1 66). For example, Gorozaemon onishiya started pole (bouchot) aquaculture of oysters (*Crassostrea* spp) in Hiroshima Bay during the npo era (1673 -1681), and it had been commonly operated in coastal intertidal areas with appropriate environmental conditions in Hiroshima Bay, Matsushima Bay, and Ariake Sound until the suspended culture method was established by Seno and Hori in 1 23. The suspended culture grew significantly because of its high productivity and applicability in various environmental conditions. apanese eel (Anguilla japonica) aquaculture was established by ura iro Hattori in Tokyo in 187, and commercial scale culture was started in an 80,000 m² pond in Shizuoka Prefecture in 18 7.

apanese eel has by far the largest production by volume and value in inland aquaculture in apan today. Yellowtail (*Seriola uin ueradiata*) aquaculture was started by Sakichi Amino in agawa Prefecture in 1 27. It drastically expanded in western apan after the end of World War II, owing to the introduction of synthetic fibers for net cages and the establishment of wild seed collection methods. uruma prawn (*Marsupenaeus japonicus*) aquaculture techniques are the fruit of 30-years of

research and development by Motosaku Fu inaga (1 03-1 73). Some salmonids cultured or that used to be cultured in apan were introduced from the U.S.: rainbow trout (Oncorhynchus my iss, from Shasta, CA, in 1877), brook trout (Salvelinus fontinalis, in 1 01-1 02) and brown trout (Salmo trutta, accidentally introduced with rainbow trout). In addition to these, various other aquatic organisms are commercially cultured in apan. Despite the historically strong affinity of the apanese for seafood and fish, aquaculture production is on a long-term moderate decreasing trend largely due to socio-economic factors in apan. This report describes the present status of aquaculture in apan, and some of the problems confronting apan's aquaculture industry.

MATERIA S

The data analyzed and presented in this report are based on the national statistical data for aquaculture production published by the Ministry of Agriculture, Forestry and Fisheries of apan (MAFF). The statistical data on the aquaculture production by volume and value of ma or species are available online in the apanese language (http://maff.go.p/ tokei kouhyou kensaku bunya6.

For marine aquaculture, the MAFF statistical data contain seven species of teleost fishes (Oncorhynchus isutch, Seriola spp., Trachurus japonicus, Pseudocaran dente , Pagrus major, Paralichthys olivaceus, Tetraodontidae spp., Thunnus orientalis), two species of bivalves (Mi uhopecten yessoensis and Crassostrea spp.), one species of prawn (Marsupenaeus japonicus), one species of tunicate (alocynthia roret i) and four species of seaweeds (Saccharina spp., Undaria spp., Pyropia spp. and emacystus spp.). Four freshwater species of teleost fishes (Oncorhynchus my iss, Plecoglossus altivelis, Cyprinus carpio and Anguilla japonica) are also included in the MAFF statistics.

There are also miscellaneous groups, such as other

teleost fish, other molluscan shellfish and other seaweeds; production of each species is relatively small and pooled together for convenience. Some freshwater species were dropped from the statistical list after 2000 because of the relatively minor importance to the industry and the budgetary constraints of the related agencies. These species include the apanese silver crucian carp (*Carassius auratus langsdorfii*), tilapia, Chinese soft-shelled turtle (*Pelodiscus sinensis*), and other miscellaneous groups.

Statistical data of the Fisheries Census is available online at the e-Stat website in apanese and nglish languages (https: www.e-stat.go. p en). The statistical data are made available by the Minister of Internal Affairs and Communications to reveal the situation surrounding the fisheries industry in apan, such as the production and employment structures of fishing communities, and the distribution and processing of fishery products. The data in the census include the number of fishery cooperatives, fisheries management entities, fisheries workers and fishing vessels.

Marine a ac It re

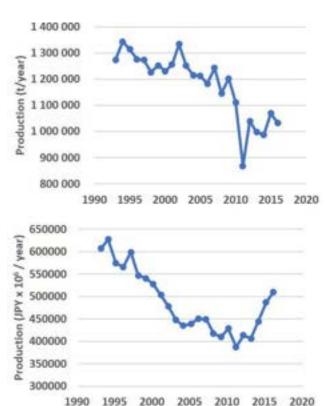


Figure 1. Total marine a uaculture production in Japan by volume (a) and value (b).

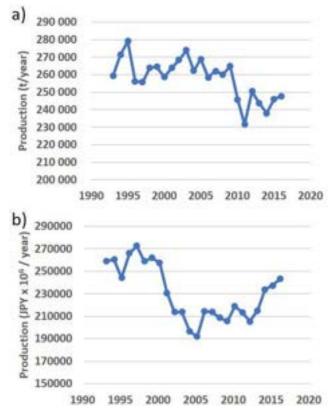


Fig. 2. A uaculture production of marine teleost fish in Japan by volume (a) and value (b).

Total marine aquaculture production gradually and continuously decreased from 1 6 (1.28 metric tons) to 2016 (1.03 10^6 t), i.e. a 1 .6% decrease over the 20 years in apan (Fig. 1a). It showed a sharp drop in 2011 (8.6 10 t) due to the damage caused by the Great ast apan arthquake and consequent tsunamis, and it went back to the long-term decreasing track in 2012. The total marine aquaculture production by value, on the other hand, continuously decreased until hitting the bottom in 2011. It has been on an increasing trend since 2012 (Fig. 1b). The production value in 2016 was PY .10 10¹¹ (approx. US 4.72 10). A similar trend is seen in the production of marine teleost fish. The total aquaculture production volume of teleost fish is on a long-term moderate decreasing trend, with a drop in 2011 due to the earthquake and tsunamis (2. 6 10 t and 2.48 10 t in 1 6 and 2016, respectively; Fig. 2a).

Teleost fish aquaculture was not severely affected by the earthquake, except for Coho Salmon (*O*.

isutch), whose production level drastically decreased from 1.48 10⁴ t in 2010 to 116 t in 2011 because it was conducted mostly along the north-east pacific coast close to the epicenter. However, it quickly recovered to .73 10³ t in 2012.

Yellowtail (Seriola uin ueradiata, S. dumerili and S. lalandi) and red seabream (P. major) are the two ma or marine teleost species cultured in apan (1.41 10 t and $6.70 10^4$ t, respectively in 2016), and they had relatively minor damage to production. Yellowtail production was greater in 2011 (1.46 10 t) than in 2010 (1.3 10 t). Damage to yellowtail aquaculture was relatively minor since the main production area is in western apan distant from the epicenter. About 12% of the yellowtail produced in apan is exported, and the US is the largest importer (8 % by value). The production of red seabream was on a long-term decreasing trend before the earthquake, the extent of which made the damage by the earthquake appear insignificant. The production, however, began an increasing trend after 2013.

Contrary to production volume, production value of teleost fish decreased from 1 7 to 200 (PY 1. 2

10¹¹) and thereafter turned into an increasing trend regardless of the earthquake (Fig. 2b). Production value was PY 2.43 10¹¹ in 2016. The recent increase in teleost fish production is in part attributable to the new inclusion of Pacific bluefin tuna (*T. orientalis*) to the MAFF statistics list in 2012. indai University established the complete aquaculture technique of Pacific bluefin tuna in 2002 and started providing hatchery produced uveniles to the private sector in 2007.

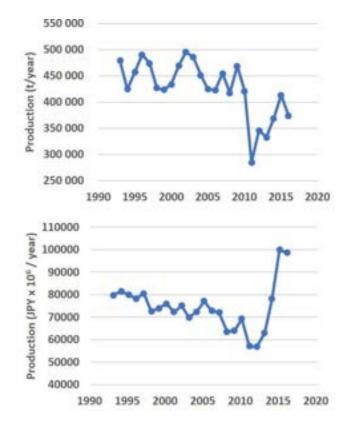


Fig. . A uaculture production of marine molluscan shellfish in Japan by volume (a) and value (b).

The production of marine molluscan shellfish aquaculture was relatively constant around 4. 10 t from 1 6 to 2010 (Fig. 3a). However, production was severely impacted by the earthquake in 2011 (2.8 10 t) and has not recovered to the level prior to the earthquake as of 2016 (3.74 10 t).

The production value of marine molluscan shellfish, however, showed a big leap after the earthquake (Fig. 3b) increasing by 1.73 times from 2012 (PY .6 10¹⁰) to 2016 (PY .86 10¹⁰). The production of oysters (mainly Pacific oyster, *C. gigas*) and apanese scallop (*M. yessoensis*) were both heavily affected by the earthquake. apanese scallop production was around 2.3 10 t before it declined to 1.18 10 t in 2011. It recovered to 2.48 10 t in 201 , exceeding the value in 2010 (2.20 10 t). The production value of the scallop showed a big leap after 2012 (PY 2. 7 10¹⁰), reaching PY 6.24 10¹⁰ in 2016.

Oyster production, on the other hand, was on a long-term decreasing trend before the earthquake (2.2 10 t and 2.0 10 t in 1 6 and 2010, respectively) and the trend persisted after the earthquake (1.66 10 t and 1. 10 t in 2011 and 2016, respectively). The production value of oysters also showed an increase after the earthquake but to a lesser extent compared to that of the scallop (PY 3.04 10¹⁰ t and 3. 4 10¹⁰ in 2012 and 2016, respectively).

Seaweeds are important commodities in apan as human food. Laver or *nori* (*Pyropia* spp, recategorized from *Porphyra*), for instance, has the second largest aquaculture production value (PY 1.00 10¹¹ in 2016) following yellowtail (PY

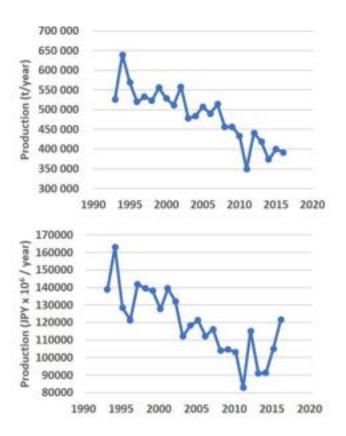


Fig. . A uaculture production of seaweed in Japan by volume (a) and value (b).

1.18 10¹¹) in apan. The aquaculture production of seaweeds is on a long-term decreasing trend both in volume and value (Fig. 4). The production volume

and value of seaweed were 3. 1 10 t and PY 1.22 10¹¹, respectively in 2016. These were 2 .6% and 12.3% less than those in 1 3.

Aquaculture of kelps (*Saccharina* spp,) and *wa ame* (*Undaria* spp.) are operated mainly in the Tohoku and Hokkaido regions and were damaged by the earthquake in 2011. Production of *wa ame* decreased from .24 10⁴ t in 2010 to 1.88 10⁴ t in 2011 (i.e. 64% decrease), and it quickly increased to 4.83 10⁴ t in 2012. Production of kelp was reduced by 42% from 2010 (4.33 10⁴ t) to 2011 (2. 1 10⁴ t). Damage to *nori* aquaculture was relatively minor since the main production area is in western apan.

Inlan a ac It re

apan's inland (freshwater) aquaculture production is small compared to marine aquaculture, only about 3% and 20% by volume and value, respectively in 2016. apanese consumers generally choose seafood in preference to freshwater fish. apanese eel (A. japonica) may be the only exception, although it is catadromous and not a freshwater species in a strict sense. It is not common to culture freshwater molluscs, crustaceans and algae in apan. Ma or freshwater aquaculture species in the global market, such as carp, catfish, and tilapia are not consumed in large quantities in apan.

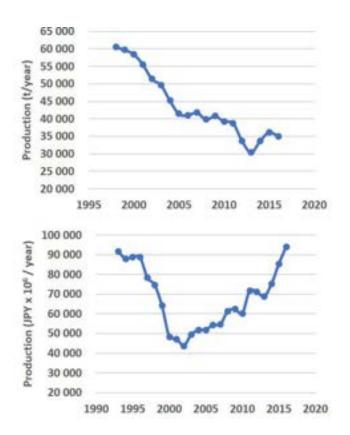


Fig. 5. Total inland a uaculture production in Japan by volume (a) and value (b)

The production volume of freshwater teleost fish has been on a continuous decline (Fig. a). The production decreased by 42% from 1-8 (6.0 10^4 t) to 2018 (3. $0-10^4$ t). The production value of freshwater teleost fish decreased from 1-3 (PY .20 10^{10}) to 2002 (PY 4.38 10^{10}) and has thereafter been on an increasing trend (Fig. b). The production value in 2016 (PY .40 10^{10}) exceeded that in 1-3.

apanese eel has by far the largest aquaculture production, comprising about 70% of the total freshwater aquaculture production value. The seeds for eel aquaculture are currently all wild caught, and the low availability of the glass eel has been problematic in recent years. apanese eel is listed as an endangered species in the IUCN Red List of Threatened Species. The unit price for the glass eel is soaring, marking the highest value (PY 3.0 10^6 kg) in 2018 (Fisheries Agency, 201), while it was PY 1.6 10 kg in 2003. Studies on artificial

spawning of apanese eel were started in the 1 60s, and the National Research Institute of Aquaculture, apan Fisheries Research and ducation Agency finally succeeded in the complete aquaculture of apanese eel at laboratory scale in 2010. However, technical difficulties remain to be overcome in scaling up the seed production in laboratory tanks to full industrial scale. Prompt establishment of mass production technologies for eel seeds is desired.

Declining a ac It re activities in a an

As above, aquaculture production volume of many species has been on a decreasing trend in apan for the past two decades. The earthquake and tsunamis had a big impact on the production of some of the species in 2011, but the long-term decreasing trend is not attributed to natural disasters. Although production values of some species are increasing due to increasing unit price, it can be said that aquaculture activities are generally dwindling in apan. This is also true for fisheries activities as a whole in apan.

One of the causes for the production declines is considered to be the reduced number of aquaculture management bodies due to the aging of the operators and a shortage of successors. The number of management bodies of yellowtail aquaculture, for instance, decreased 81.8% from 3,473 in 1 78 to 632 in 2013. Soaring feed prices are putting pressure on the profits of fed aquaculture of teleost fish. Similarly, the number of management bodies of *nori* culture decreased by 2.6% from 1 63 (1,3 4) to 2013 (3,81). The number of aquaculture farmers is decreasing in apan. As of 2016, the fishing industry employed 166,000 people, which was about half the number in 1 (32,000 people). The average age of fishing industry employees is 6.7 years old as of 2016. Merger is one of the reasons for the decrease in the number of the management bodies, but the ma ority of them are still small-scale, self-owned businesses, which are less competitive than enterprises within a large capitalist economy.

This is not simply a chicken-and egg-situation. apan is facing a labor shortage due to demographic problems; apan's population is shrinking and the average age is rising because of a low fertility rate. The rising education level of workers makes them reluctant to be engaged in menial obs. It seems that the socio-economic reconstruction of the fisheries society is needed for future development of aquaculture and fisheries in apan.

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IUCN Red List of Threatened Species (apanese eel, *Anguilla japonica*)https: www.iucnredlist.org species 166184 11 177 1)

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Despite the increasing global demand for aquaculture products, aquaculture business management is not necessarily sustainable in apan.

Some aquaculture companies producing pufferfish and red seabream went bankrupt in 2010 due to low fish prices and increasing feed prices. The author analyzed the supply and demand and price change trends of fishmeal and discussed their effects on

aquaculture business in apan. The author urged the importance of counter measures for insufficient fishmeal supply, such as use of unutilized fish and development of low fishmeal feed.

reliminar E eriments on Using the Sea ee E tract AM E to Enhance Thermal Tolerance an ro th of the S gar el Saccharina atissima from Ne Englan

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: Ascophyllum nodosum, biostimulants, thermal stress, suboptimal temperature

ABSTRACT

Seaweed extracts are widely used as biostimulants to enhance the performance of land crops. Acadian Marine Plant xtract Powder (AMP P), an extract derived from *Ascophyllum nodosum*, that has also been reported to enhance growth of *Kappaphycus alvare ii*, a red seaweed crop. Here we assessed if AMP P can enhance the thermal tolerance of the kelp *Saccharina latissima*. To assess the effects of AMP P, we first exposed early sporophytes to different concentrations (0.001, 0.00 , 0.0 , 1 and mg L⁻¹) of the product and allowed individuals to grow for 20 days at an optimal temperature of 12

1 . After determining that there were no significant differences in sporophyte growth related to the concentration of AMP P, we transferred the sporophytes to the suboptimal temperature of 18 where they remained for 14 days. Throughout the experiment, the photoperiod was maintained at 12:12 L:D with a photosynthetically active radiation (PAR) of 0 10 mol m⁻² s¹. We found that sporophytes previously exposed to AMP P showed a higher percentage of survival and growth than control sporophytes never exposed to AMP P. Sporophytes also showed thicker blades and a darker brown color than control sporophytes. These preliminary results indicate that AMP P may enhance growth of the sugar kelp, *Saccharina*

latissima, when exposed to suboptimal temperatures.

INTRODUCTION

The kelp farming industry is undergoing a fast global expansion, becoming one of the fastest growing industries in western countries (Cottier-Cook et al. 2016, im et al. 2017). In the United States for example, kelp farming, mainly of the sugar kelp Saccharina latissima is emerging as a sustainable aquaculture industry that is revitalizing the working waterfronts of New ngland by offering new economic opportunities for seafood production while at the same time removing excess nutrients from the coastal marine environment (Yarish et al. 2017). To keep its momentum, kelp aquaculture in the region needs to overcome challenges that may result in slowing its growth. For instance, it is necessary to invest in the development of strains that can resist diseases, biofouling, and increasing seawater temperature, while maintaining rapid growth rates and higher yields (im et al. 2017).

Rising seawater temperatures are already impacting most temperate marine ecosystems, affecting wild biomass production as a whole (Wernberg et al. 2016, 2018). Because kelp farming is a key component of aquaculture production (De Silva and

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Soto 200 , im et al. 2017), rapid adaptive measures should be considered to support expanding operations. In the United States, there are several ongoing selective breeding programs as part of a national effort to domesticate and improve (e.g., enhance thermal tolerance) native kelp cultivars, including *Saccharina spp.* (MARIN R, 2017).

xperiments conducted principally on the economically important red alga, Kappaphycus alvere ii, highlight what could be a faster alternative to enhance seaweed strains (Loureiro et al. 2014b, Marroig et al. 2016, Souza et al. 2018). These studies have shown that commercial extracts obtained from certain seaweeds can be used to improve the resistance and vigor of other non-related seaweeds and terrestrial plants. For example, similar to the results obtained in agriculture (han et al. 200), applying AMP P (Acadian Seaplants Ltd., Canada), a seaweed extract from Ascophyllum nodosum, to cultivars of Kappaphycus alvare ii resulted in growth enhancement (Hurtado et al. 200; Tibubos et al. 2017; Souza et al. 2018), reduction in epiphyte loads, and higher thermal tolerance (Loureiro et al. 2014b; Hurtado and Critchley 2018). Saccharina latissima, as other kelp species, is vulnerable to changes in water temperature. It shows an optimal growth between 10 and 1 C (Lee and Brinkhuis 1 88, gan et al. 1 8, Yarish et al. 1 0, Augyte et al. 2017), which decreases with temperatures exceeding 16 C, while 100% mortality is observed for temperatures exceeding 23 C (Lee and Brinkhuis 1 86). Here we report on the effects of applying AMP P to S. latissima as a potential method to increase thermal tolerance and enhance growth under optimal and suboptimal temperature conditions.

MET ODS

The experiment consisted of testing the effect of AMP P on the thermal tolerance and growth of uvenile sporophytes (00 m in length) of *Saccharina latissima*. Sporophytes were produced

by fertilizing gametophytes (SL18-UCONN-CC) isolated and cultivated in laboratory conditions from parental sporophytes collected from Cape Cod Canal (lat. 41.773818N; long. 70.4 448W).

A stock of AMP P solution was prepared by dissolving 10 g of AMP P in 1L of filtered and sterilized seawater. We then prepared five dilutions $(0.001, 0.00, 0.1, 0., and 1 g L^{-1})$ and a control solution consisting of 100 ml filtered and sterilized natural seawater (30 ppt). ach of the AMP P solutions and the control was then used to dip 36 kelp uvenile sporophytes (00 m in length, 1 days old); 6 kelp sporophytes per treatment plus 6 controls, where they remained for 4 minutes. Immediately after the 4 -minute dipping time, kelp sporophytes were rinsed with seawater and two sporophytes (pseudoreplicates) per treatment and control were transferred to 0 ml rlenmeyer flasks containing nutrient-enriched seawater (half strength Provasoli nriched Seawater (Provasoli 1 68) plus germanium dioxide (GeO₂) following Lewis (1 66) to inhibit the development of diatoms. ach pair of blades (N 3 per treatment) was allowed to grow for 20 days at 12 1 with a PAR of 0 10 M photons m⁻² s⁻¹. Blades were photographed every five days using a PixeLIN camera mounted onto an Olympus (S H) dissecting scope. Photographs were processed as binary images using Fi i by Image (Schindelin et al. 2012) to calculate the area occupied by each blade as a proxy to measure growth.

After the 20 days, the sporophytes were transferred to 12 ml rlemeyers flasks containing media as described above. They were allowed to grow for 1 additional days inside an incubator set at the suboptimal temperature of 18 0. . This allowed us to measure if the exposure to AMP P promoted an increase in thermal tolerance by facilitating a higher percentage of survival and growth. Sporophytes were photographed at the end of the 1 - day period and final blade area was calculated. After photographing, all of the survivors were cross-sectioned using a razor blade and photographed again to examine what appeared to be potential differences in the thickness of treated and non-treated blades.

Data anal sis

The specific growth rate (SGR, expressed as percent increase day⁻¹) of uvenile kelp sporophytes was calculated as:

$$SGR = (Ln(A1) - Ln(A2)) \div (T1 - T2) \times 100$$

where A1 and A2 are the area (mm⁻²) at time T1 and T2, respectively (im et al. 200).

Differences in SGR were analyzed using repeated measures ANO A as a function of concentration of AMP P $(0, 0.001, 0.00, 0.1, 0., and 1 g L^{-1})$ and time. For all of the analyses, significance was P 0.0 unless otherwise noted.

RESU TS

arly uvenile sporophytes of *Saccharina latissima* showed the highest specific growth rates (SGR) at day ten post-treatment with AMP P (Fig.1). Results show no significant difference in the SGR as a function of the concentration of AMP P when growing at 12 C(P - 0.0). Nonetheless, when sporophytes were exposed to 18 C, treated samples did show a higher percentage of survival (Fig. 2), darker brown color, and increased overall growth (Fig. 3) compared with control sporophytes, which looked pale.

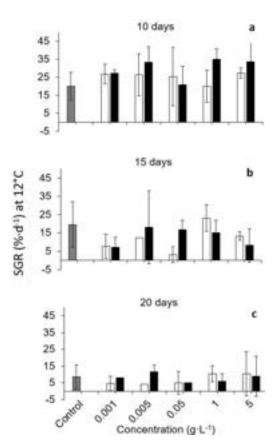


Figure 1. Specific growth rates (mean values SE,) of Saccharina latissima as a function of AMPEP concentration and time. The absence of error bars indicates only one survivor per treatment

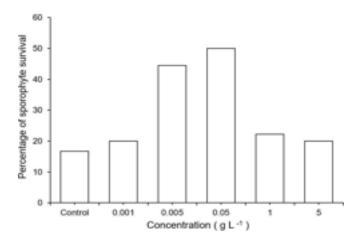


Figure 2. Percentage of survival of juvenile sporophytes of Saccharina latissima growing at 1 C after e posure to different AMPEP solutions.

Cross-section measurements also revealed that treated sporophytes showed thicker blades than non-treated individuals. Blade thickness of treated sporophytes (avg. 0.8 cm in length) was on average 1. 0.2 mm, while thickness of control sporophytes was 0.4 0.03 mm. We could not confirm significance for either survival, growth, or thickness due to a reduced number of survivors across treatments.

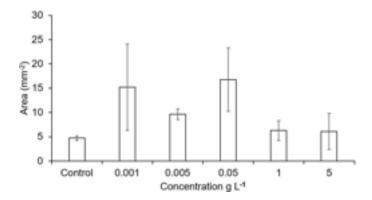


Figure 3. Final area of treated and control blades of Saccharina latissima e posed to 1 C for 15 days.

DISCUSSION

The commercial extract AMP P derived from *Ascophyllum nodosum* has been used widely in agriculture and horticulture for the beneficial effects of enhancing yields and reducing damage caused by external biotic and abiotic stressors in a variety of

land crops (refer to Craigie 2011 for a review). Similarly, studies conducted on red seaweeds have revealed a variety of benefits with an overall increase in seaweed vigor (Hurtado et al. 2012, Loureiro et al. 2014a, Marroig et al. 2016, Souza et al. 2018). To date, we are not aware of any study evaluating the effect of a brown seaweed extract (intertidal, Fucales) on kelp (Laminariales).

Hurtado and colleagues (2012) found that the use of AMP P on *Kappaphycus alvare ii* growing in optimal conditions triggered higher growth rates than control thalli never exposed to AMP P, and that low concentrations of the product (0.1 g L⁻¹) resulted in higher growth rates than higher concentrations (i.e. 1.0 g L⁻¹). In our study, we found that applying AMP P to *Saccharina latissima* uvenile sporophytes growing at ideal 12 C did not result in increased growth rates.

However, similar to our observations related to the darker color exhibited by treated sporophytes, other experiments conducted on *K. alvare i* also report that at suboptimal temperatures samples without any AMP P treatment showed bleaching of the thallus (Loureiro et al. 2014b). The dark brown coloration exhibited by the surviving kelp sporophytes exposed to AMP P could be related to varying concentrations of fucoxanthin, however further analyses are required to establish this association.

Furthermore, Loureiro et al. (2014) also found that applying AMP P to Kappaphycus alvare ii increased its tolerance to survive when exposed to suboptimal colder temperatures. Here we detected that by applying AMP P to sugar kelp, individuals seem to increase their probability of thriving at the sublethal higher temperature of 18 C. Additionally, as in land plants, seaweed extracts applied to other seaweeds stimulate the increase of phytohormones, which in turn promote cell division and elongation (han et al. 200, 2017, Hurd et al. 2014, Li and Mattson 201, Hurtado and Critchley 2018), possibly related to the increased thickness observed on our treated blades. Nevertheless, although a

higher number of treated sporophytes survived and these exhibited overall thicker and bigger blades than non-treated samples, a larger number of replicates is required to confirm significance. Also, field experiments are required to test how treated samples will perform over time in open ocean conditions and if there are any inheritable responses. However, our data suggests that applying AMP P to uvenile kelp sporophytes while in the nursery could enhance thermal tolerance of *Saccharina latissima*, allowing treated blades to grow larger than non-treated counterparts over the same period.

AC NOW ED EMENTS

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This contribution derives from the author's presentation at the International Seaweed Symposium. Craigie provides an overview of the utilization of seaweeds by humans, and more relevant, a summary of seaweeds used in agriculture. Furthermore, this paper details with examples how the application of seaweed as biostimulants enhances land crops in different ways. The author includes brand names, countries of origin, and nutritional content of various commercially available products.

Tibubos, .R., A. . Hurtado, and A.T. Critchley. 2017. Direct formation of axes in new plantlets of *Kappaphycus alvare ii* (Doty) Doty, as influenced by the use of AMP P

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This is a recent contribution describing the effect of AMP P + on the formation of vegetative shoots of Kannanhycus alvare ii. AMP P ⁺ is a Canadian commercial extract of the brown seaweed Ascophyllum nodosum with enhanced levels of potassium. The authors tested this extract in combination with colchicine or oryzalin, in addition to synthetic plant growth regulators. They found that $\operatorname{mg} \operatorname{L}^{-1} \operatorname{of} \operatorname{AMP} \operatorname{P}^{-1} \operatorname{with} \operatorname{plant} \operatorname{growth}$ regulators produced the longest direct axis shoots $(.6 \quad 0.33 \text{ mm}), \text{ followed by } 0.1 \text{ mg L}^{-1} \text{ of}$ AMP P + combined with 1.0 mg L⁻¹ oryzalin and plant growth regulators (8.7 0.00 mm). Their results suggest that using AMP P + could improve shoot formation of *K. alvare ii* plantlets in land-sea-based nursery cultivation.

Hurtado, A. . and A.T. Critchley. 2018. A review of multiple biostimulant and bioeffector benefits of AMP P, an extract of the brown alga *Ascophyllum nodosum*, as applied to the enhanced cultivation and micropropagation of the commercially important red algal carrageenophyte *Kappaphycus alvare ii* and its selected cultivars. ournal of Applied Phycology https://doi.org/10.1007/s10811-018-1407-4

The ma ority of publications related to using seaweed extracts on the aquaculture of other seaweeds have assessed the effect of *Ascophyllum nodosum* extracts on *Kappaphycus alvare ii*. This review mainly summarizes studies made on *K. alvare ii* exposed to different treatments of *A. nodosum* extracts. *A. nodosum* extracts are utilized to mitigate biotic and abiotic stressors of *K. alvare ii* sub ect to farming conditions. The authors provide examples related to micropropagation, field cultivation, endophyte mitigation, and impacts on the resulting carrageenan qualities of different *K. alvare ii* cultivars. In addition, the authors also mention alternative seaweed extracts that have been

applied to boost the growth of other cultivated seaweeds. Altogether, this review paper provides evidence of the favorable results of using

Ascophyllum extracts as biostimulants for Kappaphycus aquaculture.

Challenges to Esta lishment of Tetra loi Bree ing Stoc s for Shellfish A ac It re

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: tetraploid, shellfish, aquaculture

ABSTRACT

In the Gulf of Mexico region, oyster farming has been rapidly increasing in recent years, in part, to help meet the seafood market demand that has resulted from a decline in fishery landings since 2012. For the oyster farming industry, triploids are recognized as the preferred products because of their fast-growth, better meat quality, and year-round harvest. With the rapidly growing oyster farming industry, there has been a significant increase in demand for oyster triploid seed. Seed shortages have occurred in the past several years and are becoming a ma or constraint to the industry. To address the demand for triploid oyster seed, establishment of tetraploid breeding stocks are needed because tetraploids are key for commercial all-triploid seed production. With support from the Gulf States Marine Fisheries Commission in cooperation with the National Oceanic and Atmospheric Administration's National Marine Fisheries Service, tetraploid induction research was initiated. It is expected that establishment of tetraploid breeding stocks will be useful for triploid seed production with diversified genetic resources to support the oyster farming industry. In this publication, challenges for tetraploid production were discussed to provide insights and better understanding of triploid-tetraploid technology for shellfish aquaculture.

INTRODUCTION

Molluscan shellfish aquaculture has traditionally been a ma or component of aquaculture production worldwide and occurs mostly in marine water environments (8.3% of the total production). In 2016, the total molluscan aquaculture production worldwide was 17.1 million metric tonnes (2.2 billion) and the cultured molluscan species included oysters, clams, scallops, and mussels (FAO, 2018).

Sustainability of this large-scale aquaculture industry requires considerable efforts to increase product quantity and quality, improve aquaculture environments, and enhance the well-being of the community. Use of genetic breeding is one of the important strategies because genetically selected broodstock can produce well-adapted offspring or seed. For shellfish aquaculture, triploid-tetraploid technology has been widely recognized by the industry and utilized in many farming species, especially oyster species (Guo, 2004). Triploids grow fast, have better meat quality in summers, and probably resist diseases in aquaculture bivalve species (Guo et al., 200, Yang et al., 2018). So far, farming of triploids has become an important part of the global oyster industry. For Pacific oysters, Crassostrea gigas, triploids account for about 0% of the production on the west coast of the United States, 100% of the hatchery seed production in France (Degremont et al., 2016), and over 2.3

billion triploid seed in China (Yang et al., 201). For eastern oysters, *Crassostrea virginica*, triploids account for nearly 100% of the seed production in the Chesapeake Bay (Peachey and Allen, 2016) and a ma ority of oyster seed in the Gulf of Mexico (Wadsworth et al., 201). For Sydney rock oysters, *Saccostrea glomerata*, triploids account for about 1% of the production in Australia (Peachey Allen 2016).

Triploids can be produced by directly inhibiting the first or second polar body or crossing of diploids and tetraploids (Yang et al., 2018). Commercially, production of triploids is usually achieved by crossing (mating) oocytes from diploids with sperm from tetraploids in most hatcheries worldwide (Guo et al., 1—6) because this method can produce 100% triploids and ease to operation by commercial hatchery staff (Yang et al., 2018). Therefore, establishment of tetraploid breeding stocks is becoming a critical factor for the application of triploid-tetraploid technology in shellfish aquaculture.

Esta lishment of Tetra loi Bree ing Stoc s

Tetraploid induction in shellfish was initiated in the 1 80s and has been a challenge due to the poor survival of tetraploid larvae to beyond the metamorphosis stage (Guo et al., 200, Yang et al., 201). In 1 4, a unique methodology for tetraploid induction was reported in Pacific oysters and the induced tetraploid larvae survived to the adult stage (Guo and Allen, 1 4c) and were used for all-triploid seed production by crossing with normal diploids (Guo et al., 1 6). This innovative method used oocytes from triploid females to fertilize with sperm from diploid males and was followed by inhibition of the first polar body (PB 1) (Guo and Allen, 1 4c). So far, this method has been applied to several other shellfish species and produced viable tetraploids in Pearl oysters Pinctada margaritifera (He et al., 2000), eastern oysters Crassostrea virginica (Guo et al., 2002), Sumino oysters Crassostrea aria ensis (Allen et al., 200), bay scallops Argopecten irradians (Surier et al.,

2012), and Catarina scallops *Argopecten ventricosus* (Maldonado et al., 2003). Tetraploid stocks have been mostly established in oyster species by using this method and applying it to commercial triploid seed production.

Challenges for Tetra loi ro ction

The process to produce tetraploids using this innovative method includes the following steps: 1) produce triploids from normal diploids and culture to adult stage; 2) identify triploid individuals and find fecund triploid females; 3) collect oocytes from triploid females; 4) induce tetraploids by fertilizing oocytes from triploids with sperm from diploids and following polar body inhibition;) culture putative tetraploid larvae to beyond metamorphosis and adult stage, and 6) confirm tetraploid individuals by non-lethal ploidy determination as founders for breeding stocks. ach of these steps is connected to one another and problems which could occur at any step would result in failure of the tetraploid production.

During this process, the big challenges for successful tetraploid production are the infrequent occurrence of fecund triploid females, limited availability of oocytes from triploid females, and poor survival of putative tetraploid larvae to beyond metamorphosis (Yang et al., 201).

ccurr nc cun tri i s

Triploids possess three sets of chromosomes and could have abnormal chromosome synapsis and segregations during gametogenesis. Therefore, triploids generally exhibit abnormal gonad development. To obtain fecund triploid females for tetraploid induction, a large number of triploids need to be produced and screened. For oyster species, occurrence of fecund triploids differed depending on species. For example, about one female out of 1,600 triploid eastern oysters (i.e., 0.06%) was observed (Supan, 2000), but for Pacific oysters, higher proportions (8%) of fecund females were observed in triploids (Guo and Allen, 1—4a). For *Crassostrea hong ongensis*, most of the triploids exhibited polymorphic sterility with

atrophic gonads and abnormal gametogenesis (hang et al., 2017). For Soccostrea commercialis, triploids were found highly retarded (Cox et al., 1 6). In other shellfish groups, the occurrence of fecund triploid females also varied. For soft-shell clams, Mya arenaria, 77% of triploids were females (Allen et al., 1 86), for Dwarf surfclams, Mulina lateralis. % of triploids were females (Guo and Allen, 1 4b), and for Noble scallops, *Chlamys* nobilis, 2 % of triploids were females (omaru and Wada, 1 8). Ongoing research on eastern oysters in my laboratory showed the female occurrences in one-year old triploids were 0.6% (out of 840), 3. % (23 out of 630), and 1.2% (14 out of 1142) in three populations originating from different geographic locations.

Additionally, for the same species (eastern oysters), different cohorts of triploids may have significantly different occurrences of fecund triploids (Dr. ohn Supan, personal communication). ven for the same cohort of triploids (eastern oysters), the occurrences of fecund triploids varied greatly in different culture locations (Yang, personal observation in 201). A comparison of diploid and triploid gametogenesis in Pacific oysters indicated that high levels (42 to 0%) of gametogenesis occurred in triploids in all nine triploid batches in different culture locations, and could be the reason for the summer heavy mortality of these triploids (Houssin et al., 201).

For the mechanism of triploid female gametogenesis, a specific gene (Nanos-like gene) was reported to be perhaps responsible because this gene was expressed in the gonads of diploid females matching with gonad seasonal development, and was expressed significantly lower in the gonads of sterile triploids (u et al., 2018). Another two genes were also found to be correlated with triploid gametogenesis (iang et al., 2017). Observation of gametogenesis in triploids of Pacific oysters identified two types of gametogenesis: -pattern, which corresponded to animals displaying numerous proliferating gonia and resulted in abundant gametes, and p-pattern, which was associated with locked gametogenesis (only few

mature gametes at sexual maturity) with accumulation of abnormal gonia (ouaux et al., 2010).

imit a ai a i it ctsrm tri i mas

Oocytes from triploid organisms usually have abnormal meiosis due to the three sets of chromosomes. Thus, the number of oocytes from triploid females is usually low (Gong et al., 2004).

fforts have been made to improve gonad development in triploids, such as applying estradiol hormones to eastern oyster triploids (uintana, 200). The availability of oocytes varied from 100 to 1.6 million per female in one-year-old triploids (n 41, Yang, unpublished data), and could be related to culture locations and seasons sampled.

h rsuria inuc t tra i ara

This is a ma or challenge for tetraploid production. In Pacific oysters, the survival of putative tetraploid larvae to beyond metamorphosis was 0% in two replicates and 0.073 % in one replicate (Guo and Allen, 1 4c). Thus, to obtain viable tetraploid uveniles (after metamorphosis), it is necessary to have enough oocytes (at least 10,000) from triploid females. Additionally, tetraploid mussel larvae were found smaller than diploids and triploids in the same cohort (Scarpa et al., 1 3) and the same situation was observed in Pacific oysters (Benabdelmouna and Ledu, 201). Therefore, smaller larvae may be disposed of during larval culture. xtensive care is needed for culture of putative tetraploid larvae by saving slow growers and maintaining good culture conditions to ensure they can survive beyond metamorphosis.

SUMMAR

Triploid-tetraploid technology has been applied to commercial oyster aquaculture in many countries for years. stablishment of tetraploid breeding stocks is the key for the application of this technology, but many challenges exist. For successful tetraploid production, great efforts are

needed to address the key challenges, especially further investigations on triploid gametogenesis, which can increase the occurrence of triploid females for oocyte collection for tetraploid production.

AC NOW ED EMENTS

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Tetra loi ro ction for all tri loi see ro ction

Tetraploid production has been a challenge in shellfish due to their poor survival to spat and adult stages despite strong efforts on this research topic since the 1 80s. In 1 4, Guo and Allen reported a unique methodology for tetraploid induction in the Pacific oyster. The tetraploids survived to an adult stage and were used for all-triploid seed production by crossing with normal diploids (Guo et al., 1 6). This innovative method for viable tetraploid spat

and adult production uses oocytes from triploid females to fertilize with haploid sperm, followed by inhibition of the first polar body (PB 1) (Guo and Allen 1 4). Later, this method was applied to other oyster species and led to the production of viable tetraploids in the Pearl oyster Pinctada margaritifera, eastern oyster Crassostrea virginica, and Somonie oysters Crassostrea aria ensis (He et al. 2000, Guo et al. 2002, Allen et al. 200). This unique methodology has been patented by Rutgers University (US 824841A) and adopted by a private company. So far, tetraploid breeding stocks have been successfully established mostly in oyster species (including Pacific oysters, eastern oysters, Sydney rock oysters, and Suminoe oysters) by using this method and applied for commercial triploid seed production.

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An alternative metho for tetra loi ro ction sing tetra loi fo n ers

Complementarily, another novel approach has been reported in the Pacific oyster using tetraploid founders (McCombie et al. 200) to produce viable tetraploids by fertilizing oocytes from diploids with sperm from tetraploids followed by inhibition of the polar body 2 (PB2). This method offers an alternative way for producing more tetraploids after production of initial founders and avoids the use of oocytes from triploid females, which is always a big challenge. This is an approach to diversify, change, and enrich the genetic background of tetraploid populations (0% from mother in the 1st generation, and probably increased to 100% from mother in the second generation, depending on the chromosome segregation in sperm formation).

Guo, ., Y. Wang, . u, and H. Yang. 200 . Chromosome Set Manipulation in Shellfish. Pages 16 -1 4. in G. Burnell G. Allen editors. New Technologies in Aquaculture: Improving Production fficiency, uality and nvironmental Management. Woodhead Publishing Limited, Abington, Cambridge.

Revie san s mmaries

This book chapter systematically reviewed shellfish polyploid breeding, including triploid and tetraploid production, concepts, mechanisms, history, updates, and future application for the aquaculture industry. In this publication, two comprehensive tables are included to summarize triploid production, growth, and performance, and tetraploid induction and breeding. In addition, there are several figures included in this review to describe the chromosome segregation mechanism for triploid, tetraploid and aneuploid production.

Yang H., N. Simon, L. Sturmer. 2018 Production and performance of triploid oysters for aquaculture. lectronic Data Information Source - UF IFAS xtension, http://dx.html.edu.fa208.

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U ate revie s in e tension st le lications for o ster tri loi tetra loi technolog

These two publications summarized the history, mechanisms, procedures, and updates about oyster triploid-tetraploid technology. Different from other ournal publications, these two publications were written in extension style with glossaries as appendixes that are understandable for general public and industry communities. Download of these two publications is free through the website links stated above from the lectronic Data Information Source, Institute of Food and Agricultural Sciences xtension, University of Florida.

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