Lessons from Japan marine stock enhancement and sea ranching programmes over 100 years

Shuichi Kitada

Tokyo University of Marine Science and Technology, Tokyo 108-8477, Japan.

E-mail: kitada@kaiyodai.ac.jp

Telephone: +81-297-45-5267

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Abstract

More than 26 billion juveniles of 180 marine species are released annually into the wild in over 20 countries, but the usefulness of this strategy remains unclear. Here, I analyse the effects of stocking by Japanese marine and salmon stock-enhancement programmes and evaluate their efficacy through a Bayesian meta-analysis of new and previously considered cases. The posterior mean recapture rate (± SD) was 8.3 ± 4.7%. Without considering personnel costs and negative impacts on wild populations, the mean economic efficiency was 2.8 ± 6.1, with many cases having values of 1 to 2. On the macro-scale, the proportion of released seeds to total catch was 76 ± 20% for Japanese scallop, 28 ± 10% for abalone, 20 ± 5% for swimming crab, 13 ± 5% for kuruma prawn, 11 ± 4% for Japanese flounder, and 7 ± 2% for red sea bream; according to these percentages, stocking effects were generally small, and population dynamics were unaffected by releases but dependent on the carrying capacity of the nursery habitat. All cases of Japanese releases, except for Japanese scallop, were probably economically unprofitable. Captive breeding reduces the fitness of hatchery fish in the wild. In addition, long-term releases replace wild genes and may cause fitness decline in the recipient population when the proportion of hatchery fish is very high. Short-term hatchery stocking can be useful, particularly for conservation purposes, but large-scale programmes may harm the sustainability of populations. Nursery habitat recovery and fishing pressure reduction often outperform hatcheries in the long run.

Keywords: Bayesian meta-analysis, ecological impacts, genetic impacts, marine stock enhancement, sea ranching, stocking effects

Introduction

The history of stocking fish larvae dates back to the 1870s (Blaxter 2000; Svåsand et al. 2000; Bell et al. 2005). Approximately 150 years later, seed production technologies has progressed, and the release of hatchery-reared animals into the wild is now a popular fisheries, forestry, and wildlife management tool (Laikre et al. 2010; Taylor et al. 2017). More than 26 billion juveniles of 180 marine species, including salmonids, are released into the wild every year in more than 20 countries (Kitada 2018). In coastal marine fisheries, uses of hatchery releases are classified into several categories with different objectives: stock enhancement ("stocking cultured organisms to replenish or increase abundance of wild stocks"), sea ranching ("stocking for put-grow-and-take food fisheries"; Leber et al. 2004) and restocking ("restoring depleted spawning biomass to a level of producing regular yields"; Bell et al. 2008). Moreover, marine ranching (basically, hatchery releases accompanied with artificial reef construction in Japan) is rapidly growing in China (Lee & Zhang 2018; Wang et al. 2018; Zhou et al. 2019) and Korea (Lee & Rahman 2018). Despite the huge number of seeds released every year, most studies have been conducted only at the experimental stage (Taylor et al. 2017). In a previous study evaluating the economic performance of 14 cases involving 12 species worldwide, most cases were found to be economically unprofitable because of the high cost of seed production compared with prevailing market prices (Kitada 2018). Empirical studies are still too limited, however, to determine the full effectiveness of hatchery-release efforts (Laikre et al. 2010), and basic questions, namely, "Do hatcheries produce extra fish for harvest, or do they simply replace natural fish with hatchery fish?" and "Are hatcheries cost-effective for producing fish?" have seldom been answered (Waples 1999). More empirical studies are thus obviously needed to settle the controversy of whether hatchery stocking is useful or harmful (Hilborn 1992; Blaxter 2000; Svåsand et al. 2000; Naish et al. 2007; Araki & Schmid 2010; Laikre et al. 2010).

The feasibility and risks of hatchery stocking cannot be tested solely by examining pilot-scale releases—full-scale releases must be considered (Hilborn 2004). Despite this principle, most marine stocking programmes worldwide are in pilot-scales and test enhancement scenarios and release strategies (Taylor et al. 2017). Among the world's countries, Japan is exceptional, having released the largest number of marine and salmonid species, often on a large scale. Focused and in-depth analyses of Japanese full-scale releases can thus provide one of the best information for improving understanding of the positive and negative effects of marine stock enhancement and sea ranching programmes.

In regard to Japanese marine stock enhancement and sea ranching programmes, several reviews have focused on the effects of stocking several fishes (Masuda & Tsukamoto 1998; Kitada 1999; Kitada & Kishino 2006), kuruma prawn (Hamasaki & Kitada 2006, 2013), abalone (Hamasaki & Kitada 2008a), and decapod crustaceans (Hamasaki & Kitada 2008b; Hamasaki et al. 2011). Attention has also been directed towards seed production (Fushimi 2001; Takeuchi 2001; Le Vay et al. 2007), seed quality and behaviour (Tsukamoto et al. 1999), broodstock management (Taniguchi 2003, 2004), and genetic effects on wild populations (Kitada et al. 2009). Reviews on salmonids have covered large-scale hatchery releases, mainly focusing on chum salmon and pink salmon; however, their emphasis was mainly on ecology, with stocking effects not fully evaluated (e.g., Kaeriyama 1999; Morita et al. 2006; Kaeriyama et al. 2012; Nagata et al. 2012; Miyakoshi et al. 2013; Kitada 2014; Morita 2014). My previous systematic review provided a perspective on economic, ecological, and genetic effects of marine stock enhancement and sea ranching programmes worldwide, including salmon (Kitada 2018), but its focus was global. No integrated review evaluating the results of both marine and salmonid stock enhancement in Japan has yet

appeared.

Here, I first outline the history and present status of Japanese salmon and marine hatchery programmes. Second, I evaluate the stocking effects of major full-scale projects using a novel Bayesian meta-analysis, with new cases added to those of the previous study. Third, I predict changes in the contribution of hatchery releases to commercial catches of representative species on a macro-scale. Finally, I summarize the consequences of the world's largest hatchery stocking programmes with three different types of broodstock management, typically used worldwide: (I) wild collection of larvae, (II) wild collection of parents (may include individuals of hatchery-origin), and (III) captive breeding (Kitada 2018)—namely, those of red sea bream in Kagoshima Bay (Type III, run for 45 years), Japanese scallop (Type I, 50 years), and chum salmon (Type II, more than 100 years). The results obtained here should benefit future fisheries management and conservation practices worldwide.

Marine and salmonid enhancement programmes in Japan

Hatcheries and seed release

Japan's marine stock-enhancement programme was initiated by the Fishery Agency in 1963. At that time, Japan aimed to become a fully developed country under its national industrialization policy. Many Japanese coastlines were reclaimed, mainly up until the 1970s (Fig. 1), and separate coastal industrial zones were created; thereafter, many coastal fishers moved to cities to earn higher incomes, and coastal fisheries failed to grow (Kitada 2001). The marine stock-enhancement programme was a mitigation policy designed to improve degraded habitats and thereby enhance coastal fisheries. Salmon hatcheries have a much longer history in Japan; they were started in the 1880s to increase fishery production of returning chum salmon and have continued for over 130 years (Miyakoshi et al. 2013). As

shown later, however, chum salmon stock enhancement failed during the early 1960s. Japan's marine stock-enhancement programme was therefore started with no successful case precedent. This situation was exactly the same as that described by Hilborn (1992): "Many believe that the future of fisheries lies with artificial propagation. Indeed, throughout the world most management agencies seem to be relying on some form of artificial propagation to rebuild fish stocks that are depleted due to poor fisheries management or poor habitat management."

To obtain a bird's-eye view of stocking activity, I first created maps of salmon and marine hatcheries. Approximate locations of salmon hatcheries were plotted on a map of Japan using 15 regional maps of salmon hatchery locations obtained from the website of the Japan Fisheries Research and Education Agency (FRA, salmon.fra.affrc.go.jp). I identified 262 salmon hatcheries in 11 prefectures of northern Japan, among which 242 are privately managed, mainly by fishers and cooperatives (Fig. 2a). The seven prefectural and 13 national hatcheries are primarily research facilities. The major target salmonid species are chum salmon (*Oncorhynchus keta*) and pink salmon (*O. gorbuscha*). The number of released chum salmon in Japan has increased remarkably since the 1970s, reaching a historical maximum of 2.1 billion per annum in 1991. Since then, however, releases of chum salmon have dropped—to 1.5 billion in 2018 (North Pacific Anadromous Fish Commission, NPAFC, 2019) (Fig. A1). A similar increasing trend occurred with pink salmon starting in the 1970s; however, the number of released pink salmon is much smaller, namely, 113 million in 2018, which is approximately 7% of that of chum salmon. The number of masu salmon (*O. masou*) released in 2018 amounted to 7 million. Sockeye salmon (*O. nerka*) are also released, but at even smaller numbers (NPAFC, 2019).

133 The locations of marine hatcheries—so-called "saibai-gyogyo centres"—were also plotted by
134 using a list of their addresses compiled by the National Association for Promotion of
135 Productive Seas. Presently, 65 prefectural marine hatcheries operate in 43 prefectures (Fig.
136 2b). In addition, 12 FRA research stations are operating; these were once national marine
137 hatcheries managed by the Japan Sea-Farming Association, which merged with the FRA
138 following administrative reform in 2003. Because the FRA stopped seed production for
139 release, four previous national marine hatcheries have closed as of 2018. In 2017, 74 marine
140 species (excluding salmon) were being released in Japan, including 34 fishes, 10 crustaceans,
141 22 shellfishes, and 8 other marine species (e.g., sea urchin, sea cucumber, and octopus). I
142 organized the number of released seeds of the 16 major species whose annual release
143 exceeded million seeds for the period 1983–2017 according to annual seed production and
144 release statistics (Fishery Agency et al. 1985–2019). These species are abalone (*Haliotis* spp.),
145 Japanese scallop (*Mizuhopecten yessoensis*), Manila clam (*Ruditapes philippinarum*), green
146 tiger prawn (*Penaeus semisulcatus*), kuruma prawn (*Marsupenaeus japonicus*), offshore
147 greasyback prawn (*Metapenaeus ensis*), swimming crab (*Portunus trituberculatus*), black sea
148 bream (*Acanthopagrus schlegelii*), flatfish (e.g., *Limanda yokohamae*), Japanese flounder
149 (*Paralichthys olivaceus*), Pacific herring (*Clupea pallasii*), red sea bream (*Pagrus major*),
150 sailfin sandfish (*Arctoscopus japonicus*), tiger puffer (*Takifugu rubripes*), sea urchins (e.g.,
151 *Strongylocentrotus intermedius* and *Heliocidaris crassispina*), and sea cucumber
152 (*Apostichopus armata*). The number of releases of 10 species has been decreasing, including
153 iconic target species such as kuruma prawn, swimming crab, abalone, red sea bream, Japanese
154 flounder, and sea urchin (Figs. A2, A3; Supplementary Data). The number of releases of
155 Japanese scallop has slightly increased, while releases of tiger puffer, Pacific herring, and sea
156 cucumber have significantly increased. The changes in release practices reflect changes in
157 subsidies and budgets of the Fisheries Agency and prefectural governments, thus showing that

the Japan marine stock enhancement programme has been governmentally led. The only exception is Japanese scallop stocking, which is run by the fishers themselves.

Evaluation of stocking effects

In earlier surveys, the counting of external tags reported by fishers was the main method used to estimate recapture rates (Kitada 1999). This type of survey was applied to estimate migration, growth, and life history, including comparisons of seed quality (Kitada & Hirano 1987; Shiota & Kitada, 1992; Kitada et al. 1994; Okouchi et al. 1994; Takaba et al. 1995). Tag-reporting rates were often very small, however, and tag-shedding and tagging mortality caused biased estimates. Researchers became aware of bias in the results when they estimated stocking effectiveness. As an alternative approach, surveys of commercial landings at fish markets (SCFMs) have been used to estimate landings of released seeds and the hatchery contribution to the landings. By sampling landings at fish markets, researchers have aimed to comprehensively estimate the stocking effect. The SCFM approach was first applied to red sea bream in Kagoshima Bay. To identify released fish at this location, almost all red sea bream landed at the Kagoshima Fish Market were checked for a deformity of the internostril epidermis (Shishidou 2002; Shishidou & Kitada 2007; Kitada et al. 2019). To sample Japanese flounder in Miyako Bay, all flounder at the Miyako fish market were examined (Okouchi et al. 1999, 2004). Because such a census approach was generally difficult to apply, sampling surveys have been accordingly introduced throughout the country. The procedure used in these surveys is to check all fish landed and sold at a selected market on a selected day to differentiate released seeds from wild individuals. This latter approach, treated as a procedure, implemented in two stages to allow for the formulation of estimators and their variance (Kitada et al. 1992), has been applied to abalone (Kojima 1995), kuruma prawn (Yamaguchi et al. 2006), and masu salmon (Miyakoshi et al. 2001). When information on

total landing days is lacking, simple random sampling is assumed, and landings from releases are then estimated (Obata et al. 2008). Some other SCFM-based studies have assumed simple random sampling, but the precision was not evaluated in most cases. In yet another approach, genetic marking has been applied to mud crab (*Scylla paramamosain*) to estimate recapture rates using genetic stock identification (Obata et al. 2006).

A population dynamic model was developed to predict the effects of fishing regulations and hatchery releases on fishery production. Under the auspices of the Fisheries Agency, the Japan Sea-Farming Association tested this model using red sea bream and Japanese flounder data collected throughout the country (Kitada & Okouchi 1994). The model parameters were then adjusted to fit predicted catches with observed ones. Although the retrospective prediction was particularly sensitive to natural mortality coefficient values and parameters in the reproduction models, such models of simulated fishery production are convenient for ascertaining the relative effect of fishery management and release strategies. Indeed, similar simulation models are presently being used for Pacific herring, Japanese flounder, and tiger puffer. In many cases, however, the predictions have failed—such as with Japanese Spanish mackerel (*Scomberomorus niphonius*) in the Seto Inland Sea (Obata et al. 2007) and red sea bream in Kagoshima Bay (Shishidou et al. 2012).

To boost the effectiveness of hatchery releases, a marine-ranching project led by the Fishery Agency endeavoured to shape ‘natural’ farms to provide sound habitat, such as enhanced seaweed communities to promote nutrient enrichment in deep-sea upwelling systems (AFFRC 1989). The major technologies applied for marine fishes that can be reared and released as seeds around artificial reefs have been hatchery releases and artificial reef construction, and feeding systems with acoustic conditioning are also a popular tool (Kudo & Kimoto 1994;

Kayano et al. 1998). I could not find scientific literature evaluating the effectiveness of Japan's marine-ranching projects, and their usefulness thus remains largely unknown. Consequently, I excluded the effects of marine ranching in Japan from my subsequent analyses.

Meta-analysis of stocking effects

Indices of stocking effect and Bayesian summary statistics

I used recapture rates, yield-per-release (YPR), and economic efficiency as indices of the performance of hatchery releases. YPR is defined as the weight of fish caught (g) per individual released (Svåsand et al. 2000; Kitada & Kishino 2006; Hamasaki & Kitada 2008b) as follows:

$$YPR = r \times w$$

where *r* is the recapture rate of released juveniles, and *w* (g) is mean body weight per recaptured individual. Economic efficiency (*E*) was calculated as the ratio of net income to release costs:

$$E = YPR \times v/c.$$

Here, *v* is fish price per gram, and *c* is the cost of each seed; therefore, *v*/*c* is the cost performance of a seed (Kitada 2018). The true mean for the recapture rate, YPR, and economic efficiency for each study (y_i) was not observed but instead estimated ($\hat{y}_i, i = 1, \ldots, n$). When samples (i.e., cases within a study) were taken by random sampling, then $\mathrm{E}(\hat{y}_i) = y_i$, and the variance within each study (σ_i^2) was also estimated from the data (s_i^2).

The release variables, such as tags, sizes at release, release areas, and time periods of surveys, varied in each study; therefore, the summary statistics needed to account for this variation among studies. Here, I assumed a superpopulation for the recapture studies by applying a

Bayesian approach. The sample mean of the i-th case study ($\hat{y}_i$) may follow a normal distribution in accordance with the central limit theorem. Because studies were carried out in different areas and years, each study could be regarded as independent. For this condition, the approximate likelihood of the sample mean $\hat{y}_i\ (i = 1, \dots, n)$ can be written as:

$$L\left(\hat{y}_i\ (i = 1, \dots, n) \middle| y_i\ (i = 1, \dots, n)\right) = \prod_{i=1}^{n} \frac{1}{\sqrt{2\pi\sigma_i^2}} \exp\left(-\frac{1}{2\sigma_i^2}(\hat{y}_i - y_i)^2\right)$$

I assumed a prior distribution for $y_i\ (i = 1, \dots, n)$, where y_i is assumed to be normally distributed around the mean μ with variance σ^2. The mean and variance of the superpopulation are hyperparameters. The prior distribution of $y_i\ (i = 1, \dots, n)$ can be written as:

$$\pi(y_1, \dots, y_n | \mu, \sigma^2) = \prod_{i=1}^{n} \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{1}{2\sigma^2}(y_i - \mu)^2\right)$$

By integrating the product of the prior distribution and the likelihood function in terms of y_i, the marginal likelihood function was derived. In this case, it was explicitly obtained as:

$$\tilde{L}(\sigma^2, \mu\ | \hat{y}_i(1, \dots, n) = \prod_{i=1}^{n} \frac{1}{\sqrt{2\pi(\sigma^2 + \sigma_i^2)}} \exp\left(-\frac{1}{2(\sigma^2 + \sigma_i^2)}(\hat{y}_i - \mu)^2\right)$$

According to this equation, the sample mean $\hat{y}_i$ is distributed around the mean μ, with variances calculated for between cases and within cases.

The log marginal likelihood function is:

$$\log\tilde{L}(\sigma^2, \mu\ | \hat{y}_i(1, \dots, n) = -\frac{n}{2}\log 2\pi - \frac{1}{2}\sum_{i=1}^{n}\log(\sigma^2 + \sigma_i^2) - \frac{1}{2}\sum_{i=1}^{n}\frac{1}{\sigma^2 + \sigma_i^2}(\hat{y}_i - \mu)^2$$

The first derivatives on μ and σ^2 are then:

$$\frac{\partial \log\tilde{L}}{\partial \mu} = \sum_{i=1}^{n}\frac{1}{\sigma^2 + \sigma_i^2}(\hat{y}_i - \mu)$$

$$\frac{\partial \log \tilde{L}}{\partial \sigma^2} = -\frac{1}{2}\sum_{i=1}^{n}\frac{1}{\sigma^2 + \sigma_i^2} + \frac{1}{2}\sum_{i=1}^{n}\frac{1}{(\sigma^2 + \sigma_i^2)^2}(\hat{y}_i - \mu)^2$$

From $\partial \log \tilde{L} / \partial \mu = 0$, the maximum likelihood estimator (MLE) of μ as a weighted average of $\hat{y}_i$ $(i = 1, \ldots, n)$ is obtained as follows:

$$\hat{\mu} = \sum_{i=1}^{n}\frac{\hat{y}_i}{\sigma^2 + \sigma_i^2} \Big/ \sum_{i=1}^{n}\frac{1}{\sigma^2 + \sigma_i^2} \qquad (1)$$

By substituting σ^2 with $\hat{\sigma}^2$ from Eq. (2) and σ_i^2 with the sample variance s_i^2, the MLE of μ (posterior mean) is obtained.

Assuming for simplicity that σ_i^2 does not depend on a particular case, $\sigma_i^2 = \sigma'^2$ for all i $(1, \ldots, n)$ and from $\partial \log \tilde{L} / \partial \sigma^2 = 0$, the MLE of σ^2 is:

$$\hat{\sigma}^2 = \frac{1}{n}\sum_{i=1}^{n}(\hat{y}_i - \mu)^2 - \sigma'^2$$

By estimating μ by the weighted average $\hat{\mu}$ given by Eq. (1) and σ'^2 by the mean of the sample variance s_i^2, the approximate variance estimator is:

$$\hat{\sigma}^2 = \frac{1}{n}\sum_{i=1}^{n}(\hat{y}_i - \hat{\mu})^2 - \frac{1}{n}\sum_{i=1}^{n}s_i^2 \qquad (2)$$

The MLE of μ used here had a different weight than the weighted mean (also the MLE) used in my previous study (Kitada 2018), while $\hat{\sigma}^2$ was the same. Eq. (1) provided a more realistic estimate than that obtained using the previous weighted average, particularly when very small $\hat{y}_i$ values ($\hat{y}_i < 1$) were included (i.e., when there was large variation in $\hat{y}_i$). The average and standard deviation (SD) with a 95% confidence interval (mean ± 2 SD) were visualized for each case using the 'forestplot' function in R. When the lower confidence limit took a negative value, it was replaced by 0.

Recapture rate

I summarized 21 full-scale programmes that reported recapture rates of hatchery individuals based on various marking methods (Table A1). Eighteen of these programmes were also included in my previous study (Kitada 2018). The mean recapture rate of chum salmon was newly calculated for Hokkaido, the main production area, and was based on simple return rates (number of fish returned after 4 years/number released) taken from the FRA website (salmon.fra.affrc.go.jp). Recapture rates from 17 of 21 studies of marine species were estimated using SCFM data or a combination of SCFM data and reported recaptures. In addition, recapture rates were summarized from previous reviews for the whole country for red sea bream (Kitada & Kishino 2006), kuruma prawn (Hamasaki & Kitada 2006, 2008b), and abalone (Hamasaki & Kitada 2008a). Recapture rates were calculated for black rockfish (*Sebastes schlegelii*) (Nakagawa et al. 2004), Japanese flounder (Kitada et al. 1992; Tominaga & Watanabe 1998; Ishino 1999; Atsuchi & Masuda 2004; Tomiyama et al. 2008), Japanese scallop (Kitada & Fujishima 1997), Japanese Spanish mackerel (Obata et al. 2008), mud crab (Obata et al. 2006), short-spined sea urchin (*Strongylocentrotus intermedius*) (Sakai et al. 2004), spotted halibut (*Verasper variegatus*) (Wada et al. 2012), swimming crab (Okamoto 2004), and tiger puffer (Nakajima et al. 2008). Moreover, I newly added four cases of three species: barfin flounder (*Verasper moseri*) in Hokkaido (Koya 2005; Murakami 2012; NPJSEC 2015), red spotted grouper (*Epinephelus akaara*) in Osaka Bay (Tsujimura 2007), and tiger puffer in the Ariake Sea (Matsumura 2005, 2006); these cases were included because stocking effects have recently been reported for the first two species and the number of tiger puffer individuals released has increased in recent years (Fig. A3). In particular, more than one million juveniles (~8 cm total length, TL) of barfin flounder in Hokkaido have been released annually along the Pacific coast since 2006.

In total, recapture rates were included for 15 species comprising eight marine fishes, one salmonid, three crustaceans, two shellfishes, and one sea urchin (22 programmes) (Table A1). After excluding from analysis any cases that reported only point estimates, the recapture rates of marine species from 20 programmes (Table A1) varied widely among species and cases, ranging from 0.9% to 34.5% (Fig. 3). Japanese scallop had the highest recapture rate (34.5 ± 10.2%), followed by sea urchin (18.2 ± 17.5%), both with relatively large variations. Most of the marine fishes, namely, barfin flounder, black rockfish, Japanese flounder, Japanese Spanish mackerel, red spotted grouper, spotted halibut, and tiger puffer (to age 0+), had recapture rates in the range of 11%–15%. Abalone likewise had a relatively high and highly variable recapture rate (12.2 ± 8.1%). Red sea bream had recapture rates of 7%–8% with only small variation. In contrast, the return rate of chum salmon was much smaller, 3.6 ± 1.1% on Hokkaido and 1.6 ± 0.6% on Honshu (figure not shown). Crustaceans generally had values smaller than 5%, with large variations in recapture rates observed for kuruma prawn (2.8 ± 4.5%) and mud crab (0.9 ± 0.7%). The posterior mean recapture rate was 8.3 ± 4.7%. The empirical distribution of the posterior mean showed that the recapture rates of any species could fall into that range with 95% probability.

YPR and economic efficiency

To analyse YPR and economic efficiency, I revisited 10 cases for which YPR and economic efficiency values were previously reported (Kitada 2018) and added six new cases: those of barfin flounder in Hokkaido, Japanese flounder at Fukushima, red spotted grouper in Osaka Bay, tiger puffer in the Ariake Sea, and swimming crab in Lake Hamana and in the Seto Inland Sea (Table A2). YPR and economic efficiency values were recalculated for kuruma prawn (Hamasaki & Kitada 2006, 2008b), swimming crab (Hamasaki et al. 2011), and

abalone (Hamasaki & Kitada 2008a) based on previous reviews. For the calculation of the YPR of chum salmon, mean body weights were revised from 1974–2017 catch statistics obtained from the NPAFC and 1974–2017 return rates in Hokkaido obtained from the FRA (http://salmon.fra.affrc.go.jp/zousyoku/sakemasu.html). The revised mean body weight of chum salmon was 3 197 ± 306 g. Mean body weights were also recalculated for Japanese scallop based on Kurata (1999), yielding a revised estimate of 177 ± 30 g. YPR values reported for pink salmon in Hokkaido by Ohnuki et al. (2015) were given in monetary terms (1.5–2.2 yen/released individual), which was calculated from the estimated proportion of Hokkaido-originated hatchery fish in the landings. Because the YPR of pink salmon in Hokkaido was not provided in terms of weight, recapture rate, or cost performance of a seed (v/c), I excluded this case from the analysis.

YPR and economic efficiency were ultimately evaluated for 16 cases involving 12 species (Table A2). After omitting cases with only point estimates for YPR and/or those without information needed for calculation of v and c, a meta-analysis was performed on 12 cases involving 9 species for YPR and 13 cases involving 10 species for economic efficiency. YPR values varied between species and cases. The empirical distribution of the summary statistics had a long tail to the right, with a posterior mean of 65 ± 74 g (Fig. 4a). YPR was highest in barfin flounder (182 ± 9 g) and Japanese Spanish mackerel (170 ± 8 g), followed by chum salmon (119 ± 45 g) and Japanese scallop (61 ± 18 g), both in Hokkaido. Red sea bream and Japanese flounder had YPR values of 30–59 g. Other YPR values were 34 ± 10 g for swimming crab, 26 ± 19 g for abalone, 4 ± 3 g for mud crab, and 0.9 ± 1.5 g for kuruma prawn.

The posterior mean of economic efficiency was 2.8 ± 6.1. The economic efficiency of several

cases ranged from approximately 1 (the break-even point) to 2, with the lower 95% confidence limit below 0 (Fig. 4b). The highest economic efficiencies were those of chum salmon (19 ± 7) and Japanese scallop (18 ± 5) in Hokkaido, where the seed cost for chum salmon was set at 2.5 yen/juvenile (Hokkaido Salmon Propagation Association 2017) (Table A2). The economic efficiency of red sea bream in Kagoshima Bay was also high (5 ± 3), with similar economic performances noted for abalone across coastal Japan (4 ± 2) and barfin flounder on the Pacific coast of Hokkaido (3 ± 0.1). Japanese flounder had a much smaller economic efficiency, 0.9–1.6. Crustaceans consistently exhibited economic efficiency values around 1; among them, kuruma prawn had the lowest value, 0.7 ± 0.9. Economic efficiency is a function of YPR (recapture rate × body weight) and the cost performance of a seed (v/c). As shown in Fig. 4c, which is a scatter plot of YPR vs. economic efficiency that depicts the stocking-performance characteristic of each species, chum salmon and Japanese scallop in Hokkaido had the highest economic efficiencies.

I did not analyse net present value (NPV) (Sproul &Tominaga 1992; Moksness & Støle 1997; Moksness et al. 1998; Svåsand et al. 2000) because various data, such as annual costs for harvest, management, and interest rates, were not available for every case. If the economic efficiency estimates obtained here had relied on NPV, they would have been smaller (Kitada 2018); in that case, the values of the estimates would have depended on interest rates and time duration (although interest rates in Japan are currently low, <0.05%). In addition, the seed cost used in the analysis did not include personnel expenses, facilities, monitoring, or administration costs; furthermore, it did not account for the cost of negative effects on natural populations and ecosystems (Waples 1991; Winton & Hilborn 1994; Hilborn 1998; Waples 1999; Waples & Drake 2004; Amoroso et al. 2017). The estimates of economic efficiency obtained here are thus too high. If all the costs were available and included in the analysis, all

cases might be unprofitable except Japanese scallop in Hokkaido, where fishers pay most of the programme costs. Chum salmon in Hokkaido had higher economic efficiency than Japanese scallop, but it has been suffered from continuous decline in the population size since around 1996 (see, later discussion).

Macro-scale contribution of released seeds to commercial landings

I estimated the contribution of hatchery releases to the commercial catch of eight species (Fig. 5). These iconic species of Japan's marine stock enhancement and sea ranching programmes are intensively released within the range of Japanese waters. Trends in catches were tested using Mann-Kendall trend test with R function MannKendall. After releases of these eight species, catches of Japanese scallop ($\tau = 0.81, p = 2.2 \times 10^{-16}$) and Japanese Spanish mackerel increased ($\tau = 0.80, p = 9.5 \times 10^{-7}$) (Fig. 5). Catches of Japanese flounder ($\tau = 0.06, p = 0.614$) were stable and those of red sea bream were slightly increasing ($\tau = 0.33, p = 0.005$) . In contrast, catches of kuruma prawn ($\tau = -0.95, p = 3.9 \times 10^{-15}$) and swimming crab ($\tau = -0.66, p = 1.4 \times 10^{-8}$) have decreased continuously since the mid-1980s. Continuous declines in abalone ($\tau = -0.84, p = 8.4 \times 13$) and sea urchin ($\tau = -0.89, p = 2.2 \times 10^{-14}$) catches have also been observed since the early 1970s. Different reasons have been advanced to explain these various trends. Cases showing increasing catch levels are consistent with the frequent claims of management agencies and hatchery advocates that the practice of hatchery release should be successful. A popular explanation for cases displaying a stable catch is that hatchery release stabilizes recruitment. Instances of decreasing catches have been attributed to decreased numbers of releases; if more seeds were released, catch levels would increase. To test these ideas, estimates of the contribution of hatchery-released seeds to the catches would be helpful.

To evaluate hatchery contributions to major species for all of Japan, I computed the expected catch from released seeds by multiplying the average YPR (listed in Table A2) by the number of fish released every year (Supplementary Data). The contribution to Japanese Spanish mackerel was calculated solely from Seto Inland Sea data because hatchery releases of this species in Japan are made only at that location. This simple analysis assumed that YPR was constant over years and that released seeds created the catch in the same year. The analysis thus did not account for a time lag; however, it allowed macro-scale comparisons between species of the approximate contribution of hatchery releases (Kitada & Kishino 2006).

The largest proportion of released seeds, 76 ± 20%, was for Japanese scallop, with released spat having a stable contribution (Fig. 5, vertical bars; Fig. A4). Wild scallop created by natural reproduction also contributed to the total catch. The catch decreased substantially during 2015–2017 following a bottom disturbance off the Okhotsk coast caused by a low-pressure bomb in December 2014 (Kitada 2018); according to the estimates for that time period, almost all of the catch comprised released spat, thus indicating that scallop populations in the fishing grounds were heavily damaged by the bottom disturbance. Although the analysis was simple, the results demonstrate that this approach was able to describe the population dynamics of Japanese scallop in Hokkaido.

The hatchery contribution to the increased catch of Japanese Spanish mackerel was very small, 2 ± 2%. The catch of this fish continued to recover after releases; however, the number of released juveniles was reduced 10 years after the beginning of the release project. These results clearly indicate that the population dynamics of this species were unaffected by the releases. A previous study found 35% variation in the biomass of age-0 Japanese Spanish mackerel, an observation that could be explained by the biomass of a prey fish, Japanese

anchovy (Nakajima et al. 2013). In another investigation, genetic stock identification following releases in 2001 and 2002 found admixture proportions of hatchery-origin fish at 8%–15% (Nakajima et al. 2014). Interestingly, the genetic admixture contribution of hatchery fish in the Seto Inland Sea was much higher than estimated contribution rates (2 ± 2%) in the present study, similar to findings for red sea bream in Kagoshima Bay (Kitada et al. 2019), again implying a trans-generational genetic effect. The hatchery contribution of Japanese flounder was 11 ± 4%, and that of red sea bream was 7 ± 2%. Over two decades, the number of released juveniles decreased by ~50% for flounder and ~63% for red sea bream, but catches of wild fish remained stable and/or slightly increased; this indicates that the hatchery releases did not boost the population size of either species on a macro-scale. On the basis of carrying capacity, natural reproduction should have supported the recruitment.

Among crustaceans, the hatchery contribution of kuruma prawn was 13 ± 5%. These results are in agreement with estimates for kuruma prawn from previous research (Hamasaki & Kitada 2013), namely, ~10% throughout Japan from 1977 to 2008. In that study, the decline in kuruma prawn catches was potentially attributed to warming ocean conditions, decreased fishing efforts due to fewer fishers, and reduced hatchery releases. A continuously decreasing trend in the catch implies a decline in natural recruitments, which suggests an environmental effect is responsible (Fig. 5). Reduced fishing efforts would likely work to increase population size, as demonstrated in the case of red sea bream (Kitada et al. 2019). A reduction in fishing efforts can therefore be excluded as a cause of the decreasing catches of kuruma prawn. For swimming crab, the hatchery contribution was 20 ± 5%; the catch continuously decreased, again showing a decline in natural recruitment. Juvenile crabs are also found on tidal flats after the C4 stage (~16 mm carapace width, CW) and/or C5 stage (~22 mm CW) (Hamasaki et al. 2011). The catches of swimming crab were positively correlated with those of kuruma

prawn ($r = 0.52,\ t = 4.49, \mathrm{df} = 53, p = 3.9 \times 10^{-5}$), suggesting a common environmental effect on juveniles of both species.

Abalone displayed a relatively high and stable hatchery contribution rate (28 ± 10%), yet the catch of this species continued to decrease, indicating decreases in natural recruitment. A negative correlation has been found between the Aleutian Low-Pressure Index, winter sea surface temperatures, and catches of Ezo abalone (*Haliotis discus hannai*) in northern Japan (Nakamura et al. 2005; Hayakawa et al. 2007), and cold winter seawater temperatures (<5 °C) affect the survival of young Ezo abalone (Takami et al. 2008). Seaweed community richness (carrying capacity) is the key factor for successful abalone stocking in Japan (Hamasaki 2008). I could not calculate the hatchery contribution rates of sea urchin because no published information for calculating YPR values was found. Surprisingly similar to abalone, however, the catch of sea urchin exhibited a decreasing trend ($r = 0.96, t = 24.63, \mathrm{df} = 53, p = 2.2 \times 10^{-16}$), thereby indicating that sea urchin abundance likewise heavily depends on the richness of the seaweed community.

Economic, ecological and genetic impacts of the world's largest programmes

Broodstock management for hatchery stocking typically involves three different approaches: (I) wild collection of larvae, (II) wild collection of parents (may include individuals of hatchery-origin), and (III) captive breeding (Kitada 2018). Among the 22 cases examined above, 15 were type III, 6 were type II, and 1 was type I (Table A1). To summarize the impacts of the world's largest stocking programmes, I analysed three programmes with contrasting management approaches and with the highest economic efficiency values: red sea bream in Kagoshima Bay and Japanese scallop and chum salmon, both in Hokkaido. The broodstock management approaches used in these programmes are as follows: for Japanese

474 scallop, collection of wild larvae from the sea (type I); for chum salmon, collection of parents
475 from the wild (type II); and for red sea bream, captive breeding in a concrete tank (type III).
476 Because all three broodstock management approaches are thus represented, the three case
477 studies analysed here may be useful for predicting the long-term impacts of hatchery stocking
478 programmes worldwide.
479
480 **Red sea bream in Kagoshima Bay (type III)**
481 The hatchery-release programme for red sea bream in Kagoshima Bay is one of the world's
482 largest programmes for marine fish species and the best-monitored one in Japan. Since the
483 programme's beginning in 1974, ~27 million hatchery-reared red sea bream (6–7 cm TL) have
484 been released in the bay. Starting in 1974, the broodstock of red sea bream intended for
485 release in Kagoshima Bay have been reared in a 100-m^3 concrete tank. In particular,
486 approximately 130 non-local red sea bream broodstock are maintained in the concrete tank for
487 natural spawning and have been repeatedly used as broodstock. In 1999 and 2014, 395 wild,
488 33 farmed, and numerous 2-year-old hatchery fish produced from the broodstock were added
489 to the broodstock. The number of parent fish used for seed production has varied from
490 approximately 45 to 187 annually. During 1989–2015, 1.6 million red sea bream were
491 examined at fish markets to identify hatchery fish caught in the bay. The catch of hatchery
492 fish reached 126 tonnes by 1991 but thereafter consistently decreased, dropping to 3 tonnes
493 by 2016. This decrease was due to a decline in fitness of the hatchery-reared fish in the wild,
494 which was caused by the repeated use of parent fish reared in captivity (captive breeding)
495 since 1974 (~nine generations). In contrast, the catch of wild fish increased after 1991 and
496 reached a maximum in 2016 following releases amounting to 168 tonnes. Denser seaweed
497 communities and reduced fishing efforts were the primary factors leading to the recovery of
498 the wild population of red sea bream. These results clearly show that the recovery of nursery

habitats and reductions in fishing efforts were more effective than hatchery stocking for recovering depleted populations (see Kitada et al. 2019 for detailed information).

As seen in this example, the hatchery releases of red sea bream into Kagoshima Bay substantially increased fisheries production for the first 15 years, a period during which the programme came to be regarded as representative and successful in Japan; importantly, however, the catch of hatchery fish then steadily decreased and remained very low. The declining catch of red sea bream in Kagoshima Bay was attributed to genetic effects (Kitada et al. 2019) through unintended domestication/selection by captive breeding (Ford 2002; Araki et al. 2007, 2008). Ample evidence exists that captive breeding of salmonids can reduce the fitness of hatchery salmon in the wild (Reisenbichler & McIntyre 1977; Fleming et al. 2000; McGinnity et al. 2003; Araki et al. 2007; Christie et al. 2014). Cumulative recapture rates of red sea bream until age 8+ decreased 15.6 ± 1.7% (standard error, SE) per year. In addition, the rate of fitness reduction in hatchery-reared populations was cohort-specific; it was constant over time within the cohort but exponentially decreased with the duration of captivity (Kitada et al. 2019). Indeed, the proportion of hatchery fish in landings in inner and central parts of Kagoshima Bay and outside of the bay were highest in 1990, at 83.3%, 33.5%, and 7.4%, respectively, but the proportion in 2015 was only ~1.0% (even in the inner part) (Kitada et al. 2019). This observation suggests that once a fitness decline arises in a hatchery population, the reduction in fitness may continue until the broodstock are completely replaced by wild fish.

Gene flow between red sea bream populations was very high, thereby expanding the area affected by the hatcheries. Although more areas would thus be affected, the influence on the meta-population was not very strong. A genetic diversity analysis of fish collected from

Kagoshima Bay suggested that the genetic effects of hatchery releases were gradually diluted by backcrossing with wild populations (Kitada et al. 2019), which would diminish the genetic effects of captive breeding if such effects were additive (Roberge et al. 2008). The increasing population of wild fish in Kagoshima Bay showed no fitness decline attributable to the small proportion of hatchery fish in the meta-population. The genetic effects of captive breeding can be gradually diluted if the proportion of hatchery fish in the recipient population is not substantial and/or hatchery releases stop.

Care should be taken, however, when speculating about cases with very high proportions of hatchery fish in the recipient population, particularly cases that use captive breeding. Captive-reared parents have been repeatedly used for abalone, barfin flounder, black sea bream, Japanese flounder, and tiger puffer (Table A1). In the case of barfin flounder, one million juveniles (~8 cm TL) were released annually on the Pacific coast of Hokkaido. The catch increased markedly, from 0.01 tonnes in 1996 to over 100 tonnes in 2015, and almost all catches in the North Pacific, from Ibaraki Prefecture to Hokkaido (170 tonnes), consisted of hatchery fish (NPJSEC 2015). In the case of barfin flounder, ~500 parent fish have been used for seed production for 1–3 generations in captivity, and the expected heterozygosity is very high, 0.87 (Andoh et al. 2013). More generations of barfin flounder are needed before the impacts of this case can be determined.

Japanese scallop (type I)

Japanese scallop accounts for ~27% of landings at the Hokkaido fishery (www.pref.hokkaido.lg.jp), where all scallop landings are obtained from releases of wild-born spat or from naturally-reproduced spat of released individuals (Kitada et al. 2001; Nishihama 2001; Uki 2006). Initiated by fishers, who pay most of the programme costs (Uki 2006), the

549 scallop-ranching programmes in Hokkaido are managed by cooperatives. The management
550 system for Japanese scallop used by the cooperatives has four components: (1) mass-releases
551 of wild-born spat (wild-born sea-collected larvae are reared in net cages for 1 year prior to
552 release); (2) removal of predators, such as starfish, before release; (3) monitoring of the
553 density of scallops in the fishing ground; and (4) rotation of fishing grounds chosen for
554 harvesting (Goshima & Fujiwara 1994; Kitada et al. 2001; Uki 2006). The fishing grounds are
555 generally partitioned into four areas, and 1-year-old wild-born spat (~4.5 cm) reared in net
556 cages are released into a given area after removal of starfish. After 3 years, the 4-year-old
557 released scallops are harvested. The following year, spats are released into another area. This
558 fishing-ground rotation system enables complete prohibition of the fishery for 3 years after
559 release.
560
561 Catches of Japanese scallop in the release areas, which have increased remarkably since the
562 first releases in the 1970s, reached a historical maximum of 359 000 tonnes in 2014 (Fig. 6a,
563 updated from Kitada 2018). In 2015, the catch dropped substantially, to 232 000 tonnes
564 (65%), because of a bottom disturbance on the Okhotsk coast caused by a low-pressure bomb
565 on 16–17 December 2014 (Kitada 2018), but recovered to 305 000 tonnes in 2018, as
566 expected. The catch recovery in 2018 would have been created by spat released in 2015. The
567 long-term trends in these release and catch statistics demonstrate that sea ranching of scallop
568 in Hokkaido has been successful, with the highest observed economic efficiency (18 ± 5).
569
570 **Chum salmon (type II)**
571 Chum salmon hatchery-stock enhancement in Japan is one of the world's largest salmon
572 stocking programmes (Amoroso et al. 2017). Hokkaido produces ~80% of chum salmon
573 returning to Japan, and fishery production accounts for 21% of the catch at the Hokkaido

fishery (www.pref.hokkaido.lg.jp). Because most landings are created by releases of hatchery-born fish (Kaeriyama 1999), fishers pay ~7% of their landings of chum salmon to the hatcheries in Hokkaido (Kitada, 2014). Long-term trends in releases and catches indicate that the maximum carrying capacity of natural rivers is ~10 million fish and that a large increase in the fish population was created by hatchery stocks (Fig. 6b, updated from Miyakoshi et al. 2013). Total production in Hokkaido reached a historical maximum of 61 million returns in 2004 and then substantially decreased to 16 million in 2017. Chum salmon in Japan are at the southern margin of the species range. My previous study found that 30% of the variation in decreasing catches could be attributed to an increasing sea surface temperature (SST) anomaly, and 62% was explained by SST and catches by Russians after 1996 (Kitada 2018). Clear differences have been found, however, between the run timing of chum salmon populations in Russia (Jun–Aug) and Japan (Sep–Nov), and migration routes are also different (Kondo et al. 1965; Morita 2016). These results suggest that the negative correlation between Japanese and Russian catches reported in the previous study was spurious.

No evidence of fitness decline has been reported among hatchery-born and wild-born chum salmon in Japan. The observed genetic effect is instead an altered population structure, with some populations nested across seven and/or eight regional groups (Beacham et al. 2008; Kitada 2014; Sato et al. 2014; Kitada 2018). The run-timing distribution of Hokkaido chum salmon has been altered by the preferential enhancement of early-running fish (returning in September/October), and the late-running population has almost disappeared (Miyakoshi et al. 2013). A lower reproductive success has been observed for early-spawning sockeye salmon in Washington State, USA, and early-emerging juveniles have had relatively low survival in recent years. These observations suggest that the skewed distribution of early spawning in Japan could reduce population fitness during a warming climate (Tillotson et al. 2019).

Almost all chum salmon returning to Japan are hatchery-reared fish (Kaeriyama 1999). These hatchery-reared fish are produced every year from returning adults, and the number of maintained parent fish has been very large (i.e., 15 000–85 000) (Kitada 2018). Even so, naturally spawning chum salmon have been detected in 31%–37% of 238 non-enhanced rivers surveyed in Hokkaido (Miyakoshi et al. 2012) and in 94% of 47 enhanced rivers and 75% of 47 non-enhanced rivers on the northern coast of Honshu Island (Sea of Japan) (Iida et al. 2018). In addition, a study using otolith thermal-marking estimated the proportion of naturally spawned chum salmon to total production at 16%–28% in eight rivers in Hokkaido, with large variation (0%–50%) (Morita et al. 2013). Despite variations in the proportion of natural spawning in rivers, gene flow facilitates the genetic admixture of hatchery-released fish, hatchery descendants, and wild fish in the entire Japan population. To visualize the magnitude of gene flow between populations, I re-examined microsatellite data from 26 chum salmon populations in Japan (14 loci, n = 6,028; Beacham et al. 2008) and computed F_{ST} values between population pairs based on the bias-corrected G_{ST} (Nei & Chesser 1983) using the GstNC function in the R package FinePop1.5.1. This G_{ST} estimator provides an unbiased estimate of F_{ST} when the number of loci becomes large (Kitada et al. 2017). I then superimposed a diagram in which population pairs with pairwise F_{ST} values < 0.01 were connected by lines onto a map of hatchery locations (Fig. 7). The mean pairwise F_{ST} was very small (0.007 ± 0.003). The F_{ST} threshold of 0.01 was based on the relationship $4N_e m = 99$, where N_e is effective population size and m is migration rate, corresponding to 99 effective parents migrating between each pair of populations per generation (see Waples & Gaggiotti 2006). Figure 7 depicts substantial gene flow between rivers. The causal mechanisms of population structuring are migration and genetic drift, and differentiation depends on the number of migrants ($N_e m$) (Waples & Gaggiotti 2006; Hauser & Calvalho 2008). Even in

624 Atlantic cod (*Gadus morhua*), a species with high gene flow, temporally stable but
625 significantly differentiated structure can be detected among populations (Hauser & Calvalho
626 2008). Constant gene flow among populations can create a stable genetic mixture in a meta-
627 population, such as that observed in Pacific herring populations in northern Japan (Kitada et
628 al. 2017). These results suggest that the nested population structure observed across the seven
629 and/or eight regional groups was caused by past translocations (Beacham et al. 2008;
630 Kaeriyama & Qin 2014).
631
632 Hatchery practices might increase the likelihood of chum salmon to stray (Quinn, 1993). I
633 summarized the results of marking studies of chum salmon juveniles in Hokkaido, where 2
634 028 thousand hatchery-reared salmon juveniles (3.5–4 cm BL, 0.45–5 g) were fin-clipped
635 and/or operculum-clipped and released without rearing between 1951 and 1955 (Sakano
636 1960). According to the results, 50 ± 22% of 2 085 recoveries were recaptured in the river of
637 their release, and 84 ± 12% were found when recoveries along nearby coasts were included
638 (Supporting Data); hence, the spawning fidelity of hatchery-reared chum salmon was
639 moderate. Estimates of straying can vary largely between specific hatchery releases and
640 rivers, but the genetic integrity of a population can be altered by straying regardless of the
641 strength of the native population's spawning fidelity (Quinn 1993). Early theoretical work
642 predicted that >99% (50%) of wild genes with additive effects are replaced by hatchery genes
643 in 12 generations (2 generations for 50% of wild genes) in the case of equal fitness between
644 hatchery and wild fish at a stocking rate of 0.5 (Matsuishi et al. 1995; see also Fig. 5 in Kitada
645 et al. 2019). Taking all of these results in consideration, I speculate that most chum salmon
646 returning to Japan are hatchery-released fish or wild-born hatchery descendants. This situation
647 is similar to that of hatchery-reared red sea bream in Kagoshima Bay. The significant decline
648 in the number of returns of Japanese chum salmon may therefore be caused by a fitness

decline in populations induced by long-term hatchery stocking.

Conclusions

All cases of Japanese hatchery releases, except Japanese scallop, are probably economically unprofitable if the costs of personnel expenses, facility construction, monitoring, and negative impacts on wild populations are taken into account. Stocking effects are generally small, while the population dynamics are unaffected by releases but instead essentially depend on the carrying capacity of the nursery habitat. Hatchery rearing can reduce the fitness of hatchery fish in the wild, and long-term hatchery stocking can replace wild genes and cause fitness decline in the recipient population when the proportion of hatchery fish is very high. Short-term uses of hatchery stocking can be helpful, particularly for conservation purposes, but long-term programmes may harm the sustainability of populations. Recovery of nursery habitats and reduction in fishing efforts often outperform hatcheries in the long run.

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Data Availability Statement

All source data used are in the public sector, and links to their online sources are specified in the text or in Supplementary Data.

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Supporting Information

Additional Supporting Information may be found in the online version of this article:

Supporting Data

Catch and release data for major species in Japanese marine stock enhancement and sea ranching programmes and the results of marking experiments with chum salmon conducted in Hokkaido in the 1950s.

Figure Legends

Figure 1 Changes in tidal flat and eel grass (*Zostera marina*) areas (1945–1996). From the Ministry of Environment, www.env.go.jp and www.biodic.go.jp, accessed July 2019.

Figure 2 Maps showing the locations of marine and salmon hatcheries in Japan operated by different sectors. From FRA (salmon.fra.affrc.go.jp, accessed August 2019) and National Association for Promotion of Productive Seas (http://www.yutakanaumi.jp/, see text).

Figure 3 Forest plot of recapture rates from large-scale hatchery releases in Japan. Thin lines indicate 95% confidence intervals, with arrows (in the case of Japanese scallop and sea urchin) indicating that the confidence intervals penetrate the scale. Areas of the squares are proportional to the weight of the mean.

Figure 4 Performance of large-scale hatchery releases in Japan. Forest plots of **(a)** yield-per-release (YPR), **(b)** economic efficiency, and **(c)** YPR vs. economic efficiency. Note: the seed cost used in the analysis did not include personnel expenses, facilities, monitoring, or administration costs. SIS = Seto Inland Sea.

Figure 5 Total catch and recovery from releases of representative species in Japanese stocking programmes for all of Japan. Vertical lines depict recovery (expected catch from releases), which were estimated by multiplying values of yield-per-release (YPR) (see Table A2) and the numbers released (Supplementary Data). For sea urchin, no YPR data were available.

Figure 6 Long-term release and catch (return) statistics for **(a)** Japanese scallop (updated from Kitada 2018) and **(b)** chum salmon in Hokkaido (updated from Miyakoshi et al. 2013) (see Supplementary Data).

Figure 7 Gene flow between Japanese populations of chum salmon. Populations with pairwise $F_{ST} < 0.01$, as estimated from microsatellite data of 26 populations (14 loci, $n = 6\,028$; Beacham et al. 2008), are connected by lines.

Appendices

Table A1 Recapture rates of 21 major marine stock enhancement and sea ranching programmes in Japan

Species (Broodstock, Type)	Geographical region	Years for calculating recapture rate	Size at release size (cm)	Number released	Marking methods †	Estimation methods ‡	Recapture rate ± SD (%)	Source
Barfin flounder (Captive, III)	East-western Hokkaido	1987–	8	Million/year 2006–	EAT, ALC, no marking since 2000	SCFM, all landings were regarded as released fish	12.1 ± 0.6	Koya (2005) Murakami (2012); NPJSEC (2015)
Black rockfish (Captive, III)	Yamada Bay, Iwate	1995–1997	8.7–12.9	447,394	Removal of ventral fins	SCFM	11.8 ± 0.9	Nakagawa et al. (2004)
Chum salmon (Returned adults, II)	Hokkaido	1974–2017	5 g	1.0 billion/year	Otolith thermal marking, ALC	No. retuned (4 years after)/ No. released	3.6 ± 1.2	FRA, http://hnf.fra.affrc.go.jp/
Japanese flounder (Captive, III)	Fukushima	1987	10	246,000	PES	SCFM, a two-stage sampling method	15.0 ± 3.3	Kitada et al. (1992)
Japanese flounder (Captive, III)	Fukushima	1994–2002	10	8,260,000	PES	SCFM	12.1 ± 4.8	Tomiyama et al. (2008)
Japanese flounder (Captive, III)	Ishikari Bay, Hokkaido	1989	6.0–7.8	149,555	PES, EAT, FC	SCFM Reported recapture	5.7 ± 3.5	Tominaga & Watanabe (1998)
Japanese flounder (Captive, III)	Kagoshima Bay	1989–1995	8.6–10.5	2,189,000	PES	SCFM	2.4 ± 0.7	Atsuchi & Masuda (2004)

Japanese flounder (Captive, III)	Miyako Bay, Iwate	1986–1992	7.9 (7.0–9.3)	611,000	Ratex and brand	SCFM with fish census	14.5 ± 7.3	Okouchi et al. (1999); Okouchi et al. (2004)
Japanese flounder (Captive, III)	Southern Hokkaido	1987–1993	7–15.5	1,069,000	EAT, PES	SCFM	10.4	Ishino (1999)
Japanese Spanish mackerel (Wild, II)	Eastern Seto Inland Sea	2002–2003	10.6 ± 1.7	160,122	ALC	SCFM	15.0 ± 0.7	Yamazaki et al., 2007; Obata et al. (2008)
Red sea bream (Captive, III)	Kagoshima Bay	1974–	6.0–7.0	0.5–1.3 million/yr	DIE	SCFM	8.0 ± 4.2	Shishidou (2002); Kitada & Kishino (2006)
Red sea bream (Captive, III)	Sagami Bay, Kanagawa	1978–	6.0–7.0	0.8–1.2 million/yr	EAT, DIE	Reported recapture and SCFM	7.1 ± 2.9	Kitada & Kishino (2006)
Red spotted grouper (Captive, III)	Osaka Bay, Osaka	2000–2007	10	4,000/yr	EAT	Reported recapture	2.2 ± 1.0 (1.5–3.4)	Tsujimura (2007)
Spotted halibut (Captive, III)	Fukushima	1993–2007	7.5–51.0	426,704	Dart tag, ALC	SCFM	11.1 ± 11.4	Wada et al. (2012)
Tiger puffer (Captive, III)	Ariake Sea, Kyushu Island	1991–2003	0.3–10.2	1,313,450	ALC, TC	SCFM (for matured fish)	11.6 ± 7.0 (Age0+) 0.2 ± 0.1 (spawners)	Matsumura (2005; 2006)
Tiger puffer (Captive, III)	Mie, Aichi, Shizuoka	2001–2005	7.7 ± 1.6 (5.6–10.0)	452,839	VIE	SCFM	5.1 ± 6.5	Nakajima et al. (2008)

Kuruma prawn (Wild, II)	Western Japan coasts	1980–1991	2.3 ± 0.6	1,261,039	CWT, UC	SCFM	2.8 ± 4.5	Hamasaki & Kitada (2006)
Mud crab (Wild, II)	Urado Bay, Kochi	1997–2001	0.9–1.5 (C3–C5)	475,300	GEN	Genetic mixing proportion	0.9 ± 0.7	Obata et al. (2006)
Swimming crab (Wild, II)	Lake Hamana, Shizuaoka	1998	2.2	3,300	CWT	SCFM	1.2	Okamoto (2004)
Abalone (Captive, III)	Over Japan coasts	1980–1991	2.3 ± 0.6	1,261,039	GM	SCFM	12.2 ± 8.1	Hamasaki & Kitada (2008a)
Japanese scallop (Wild spat collected, I)	Okhotsk Sea coast, Hokkaido	1870s–	4.5	Over 3 billion/yr	No marking	Regression analysis	34.5 ± 10.2	Kitada and Fujishima (1997)
Short-spined sea urchin (Wild, II)	Tomari, and Akkeshi, Hokkaido	1987–1998	0.8–1.8	1,961,000	Width of the first ring (FR) in the genital palte	SCFM, Discrimination from FR size distributions	18.2 ± 17.5	Sakai et al. (2004)

† EAT, external anchor and T-bar tag; ALC, alizarin complexone on otolith; TC, oxytetracycline on otolith; GEN, genetic marking; PES, pigmentation of eyeless side; FC, fin clipping; VIE, visual implant elastomer; DIE, deformity of internostril epidermis; UC; uropod cutting. GM; green mark on shells due to hatchery diet.

‡SCFM, sampling survey of commercial landings at fish markets.

Table A2 Performance of 15 major marine stock enhancement and sea ranching programmes in Japan

Species	Geographical region	Marking methods †	Estimation Method ‡	Recapture rate (%)	YPR § (g)	Economic efficiency ¶	Source
Barfin flounder	East-western Hokkaido	-	YPR	12.1 ± 0.6	181.5 ± 9.0	2.7 ± 0.1	This study (*w*=1500, *v*=1.2, *c*=81)
Chum salmon	Hokkaido	-	YPR	3.6 ± 1.2	118.5 ± 45.1	18.9 ± 7.2	Kitada (2018); This study (*w*=3310, *v*=0.4, *c*=2.5)
Japanese flounder	Kagoshima Bay	PES	SCFM	2.4 ± 0.7	29.7 ± 2.6	1.1 ± 0.1	Atsuchi & Masuda (2004); Kitada & Kishino (2006)
Japanese flounder	Miyako Bay	PES	CS	13.5 ± 6.4	51.8 ± 24.2	1.6 ± 0.7	Okouchi et al. (2004); Kitada & Kishino (2006)
Japanese flounder	Fukushima coast	PES	SCFM	12.1 ± 4.8	n.a	0.9 ± 0.4	Tomiyama et al. (2008)
Japanese Spanish mackerel	Eastern Seto Inland Sea	ALC	SCFM	15.0 ± 0.7	169.7 ± 8.3	1.0 ±0.1	Obata et al. (2008)
Red sea bream	Kagoshima Bay	DIE	SCFM	8.0 ± 4.2	59.0 ± 27.2	5.0 ± 2.7	Shishidou (2002); Kitada & Kishino (2006)
Red sea bream	Sagami Bay	DIE	SCFM	7.1 ± 2.9	54.9 ± 30.4	1.4 ± 0.3	Kitada & Kishino (2006)
Red spotted grouper	Osaka Bay, Osaka	EAT	MR	2.2 ± 1.0	7.7 ± 0.4	0.3 ± 0.01	Tsujimura (2007); This study (*w*=350, *v*=5, *c*=150)

Tiger puffer	Ariake Sea, Kyushu Island	ALC, TC	YPR	11.6 ± 7.0 (Age0+) 0.2 ± 0.1 (spawners)	n.a	2.4 (Age0+ spawners)	Matsumura (2005; 2006)
Kuruma prawn	Western coasts	UC, CWT	SCFM	2.8 ± 4.5	0.9 ± 1.5	0.7 ± 0.9	Hamasaki & Kitada (2006, 2008b)
Mud crab	Urado Bay, Kochi	GEN (mtDNA)	SCFM	0.9 ± 0.7	3.7 ± 3.0	1.9 ± 1.5	Obata et al. (2006), Hamasaki et al. (2011)
Swimming crab	Lake Hamana, Shizuaoka	CWT	SCFM	1.2	1.5	n.a	Okamoto (2004)
Swimming crab	Seto Inland Sea	-	REG	17	33.6 ± 9.5	1.0 ± 0.3	Hamasaki et al. (2011) This study (*v*=0.15, *c*=5)
Abalone	Over coast of Japan	GM	SCFM	12.2 ± 8.1	25.6 ± 19.1	3.5 ± 2.4	Hamasaki & Kitada (2008a)
Japanese scallop	Okhotsk coast, Hokkaido	-	YPR	34.5 ± 10.2	60.9 ± 18.0	17.9 ± 5.3	Kitada & Fujishima (1997); Kurata (1999); This study (*w*=176.5, *v*=0.9, *c*=3)

1083 †EAT, external anchor and T-bar tag; ALC, alizarin complexone on otolith; TC, oxytetracycline on otolith; GEN, genetic marking; PES,
1084 pigmentation of eyeless side; DIE, deformity of internostril epidermis; UC, uropod clipping; CWT, coded-wire tags; GM, green mark on the
1085 shell.
1086 ‡SCFM, sampling survey of commercial landings at fish markets; CS, census of commercial landings; MR, mark-recapture; REG, regression
1087 analysis; YPR, yield per release.
1088 § Grams of fish caught per individual released.
1089 ¶ Ratio of net income to release cost, excluding personnel expenses, expenditure for hatchery facilities, and monitoring costs;
1090 n.a, not analysed.

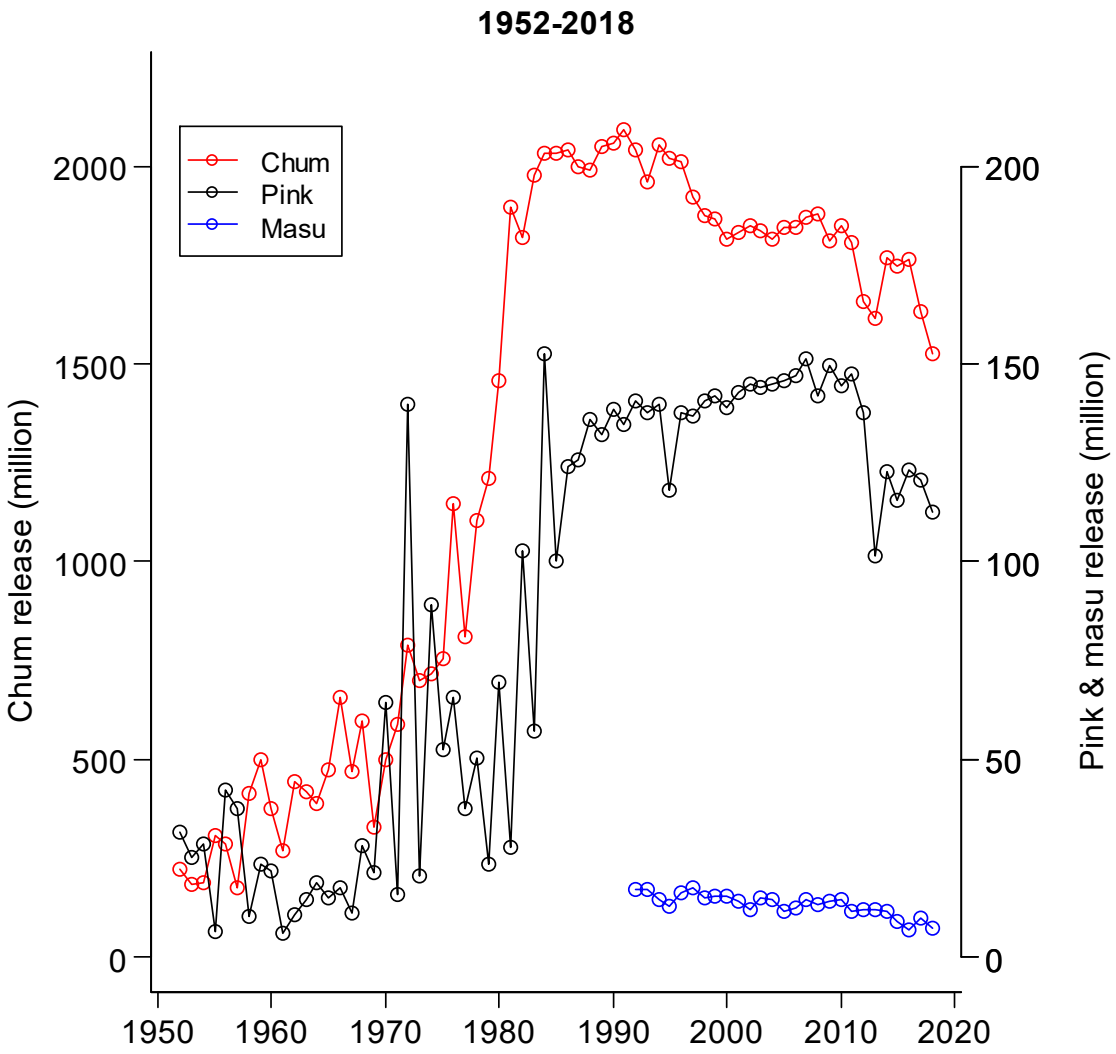

Figure A1 Number of released juveniles of chum, pink, and masu salmon in Japan (1952–2018). Data from the North Pacific Anadromous Fish Commission (www.npafc.org, accessed August 2019).

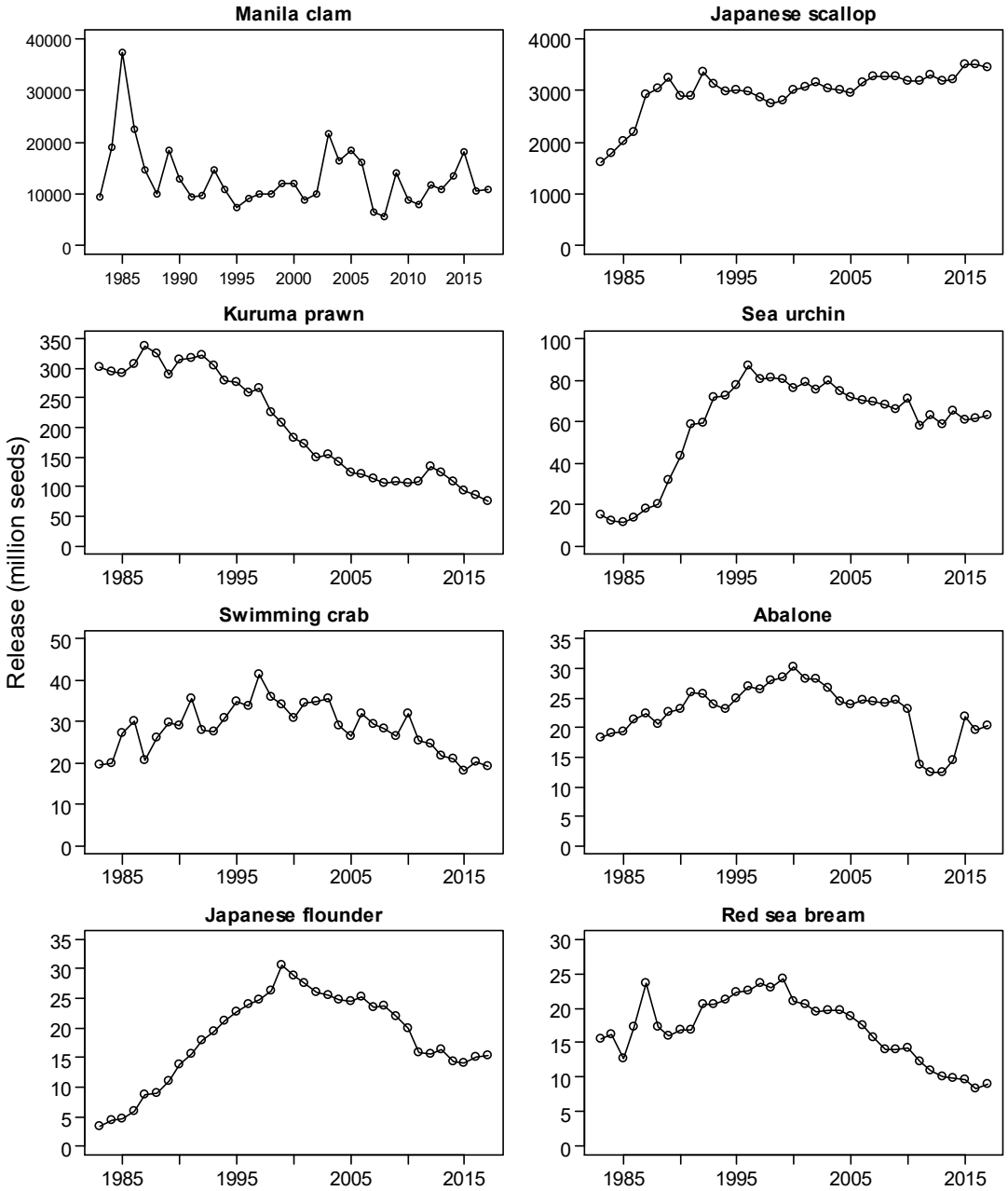

Figure A2 Number of released seeds of the top eight species of Japanese marine stock enhancement and sea ranching programmes (1983–2017). Data from the Fisheries Agency, Fisheries Research and Education Agency, and National Association for Promotion of Productive Seas (1985–2019).

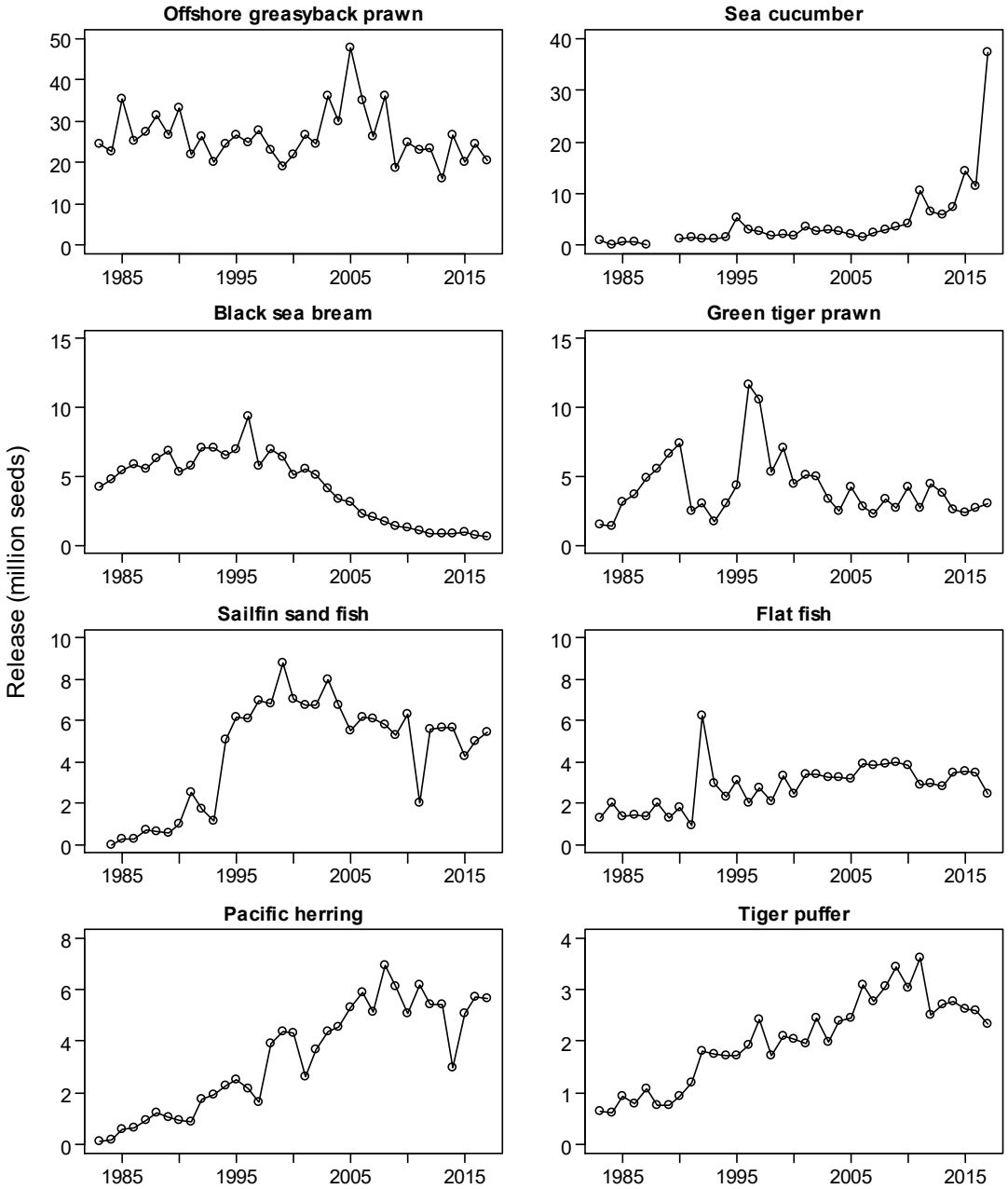

Figure A3 Number of released seeds of 9th–16th-ranked target species of Japanese marine stock enhancement and sea ranching programmes (1983–2017). Data from the Fisheries Agency, FRA, and NAPPS (1985–2019).

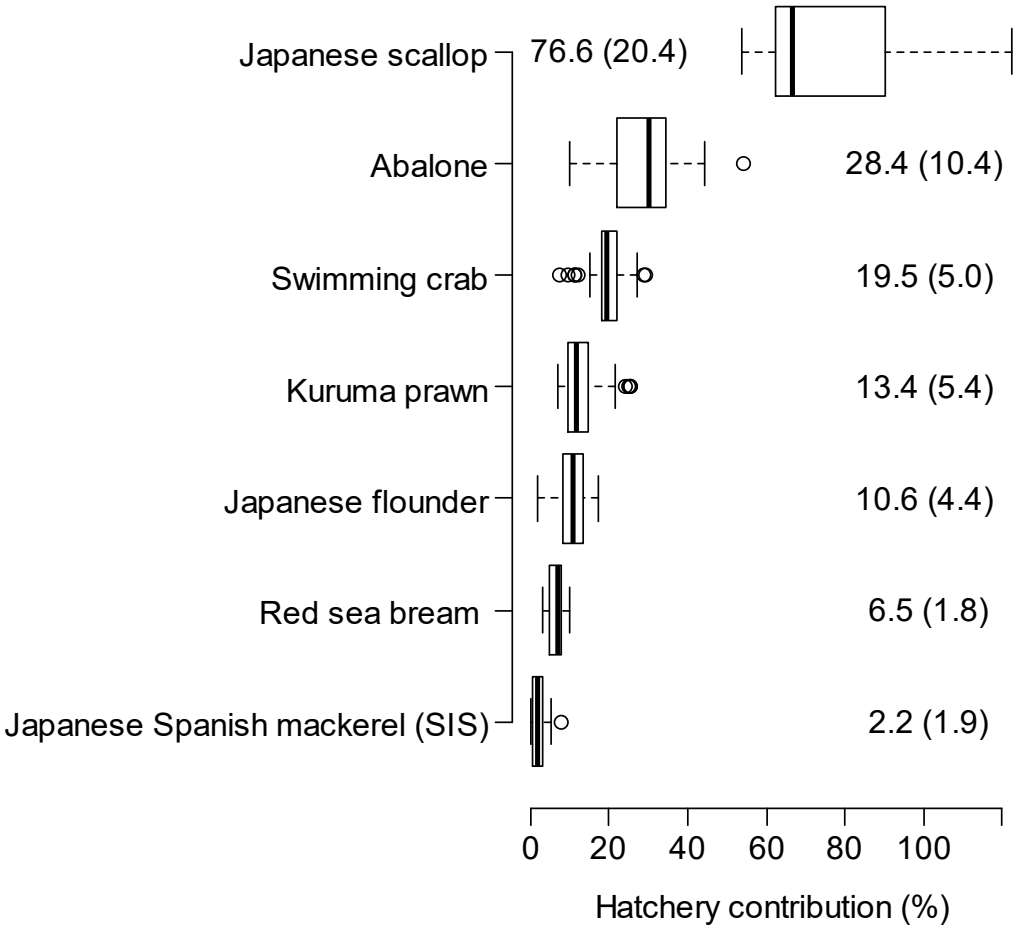

Figure A4 Percent contribution of hatchery-reared individuals to the commercial catch, calculated from Figure 5.