

Journal of Applied Ecology

MS VICTORIA DOMINGUEZ ALMELA (Orcid ID : 0000-0003-4877-5967)

DR ROBERT BRITTON (Orcid ID : 0000-0003-1853-3086)

Article type : Research Article

Editor : Brittany Mosher

Predicting the outcomes of management strategies for controlling invasive river fishes using individual-based models

Victoria Dominguez Almela¹, Stephen C. F. Palmer², Demetra Andreou¹, Phillipa K. Gillingham¹, Justin M. J. Travis², J. Robert Britton¹

¹Department of Life and Environmental Sciences, Faculty of Science and Technology, Bournemouth University, Poole, BH125BB, UK; ²School of Biological Sciences, University of Aberdeen, Tillydrone Avenue, Aberdeen, AB242TZ, UK.

Corresponding author:

Victoria Dominguez Almela

This article has been accepted for publication and undergone full peer review but has not been through the copyediting, typesetting, pagination and proofreading process, which may lead to differences between this version and the [Version of Record](#). Please cite this article as [doi: 10.1111/1365-2664.13981](https://doi.org/10.1111/1365-2664.13981)

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Accepted Article

dominguezalmela@gmail.com

ORCID:0000-0003-4877-5967

Abstract

1. The effects of biological invasions on native biodiversity have resulted in a range of policy and management initiatives to minimise their impacts. Although management options for invasive species include eradication and population control, empirical knowledge is limited on how different management strategies affect invasion outcomes.
2. An individual based model (IBM) was developed to predict how different removal ('culling') strategies affected the abundance and spatial distribution of a virtual, small-bodied, r -selected alien fish (based on bitterling, *Rhodeus sericeus*) across three types of virtual river catchments (low/intermediate/high branching tributary configurations). It was then applied to nine virtual species of varying life history traits (r - to K -selected) and dispersal abilities (slow/intermediate/fast) to identify trade-offs between the management effort applied in the strategies (as culling rate and the number of patches it was applied to) and their predicted effects. It was also applied to a real-world example, bitterling in the River Great Ouse, England.
3. The IBM predicted that removal efforts were more effective when applied to recently colonized patches. Increasing the cull rate (proportion of individuals removed per patch), and its spatial extent was effective at controlling the invasive population; when both were relatively high, population eradication was predicted.
4. The characteristics of the nine virtual species were the main source of variation in their predicted abundance and spatial distribution. No species were eradicated at cull rates below 70%. Eradication at higher cull rates depended on dispersal ability; slow dispersers required lower rates than fast dispersers, and the latter rapidly re-colonised at low cull rates. Optimum trade-offs between management effort and invasion outcomes were generally when intermediate effort was applied to intermediate numbers of patches. In the Great Ouse, model

predictions were that management interventions could restrict bitterling distribution by 2045 to 21% of the catchment (versus 90% occupancy without management).

5. **Synthesis and application:** This IBM predicted how management efforts can be optimized against invasive fishes, providing a strong complement to risk assessments. We demonstrated that for a range of species' characteristics, culling can control and even eradicate invasive fish, but only if consistent and relatively high effort is applied.

Keywords

Biological invasion, *RangeShifter*, River catchment, Simulation model, Dispersal

1. Introduction

Biological invasions are an important component of global change (Simberloff et al., 2013), which have resulted in the application of a range of policy and management initiatives to minimise their impacts (Larson et al., 2011). Optimising population control efforts within management programmes remains highly challenging, and considerable uncertainty remains in how to apply limited staff and financial resources to population control measures, especially as to how these measures should be applied spatially and temporally (Maguire, 2004).

Following the introduction of a new species, the probability of an invasive population developing depends on the interactions of a range of biotic and abiotic factors, including the species' dispersal abilities and life history traits, and the environmental conditions encountered (Catford et al., 2009). Dispersal rates are important and are influenced by habitat connectivity (Hastings et al., 2005), which is often more constrained in freshwaters than in terrestrial environments (Gozlan et al., 2010; Radinger et al., 2017). Management responses also influence invasion probabilities (Britton et al., 2011a) and their effectiveness generally increases when they occur soon after introduction, when the species is spatially constrained and of relatively low abundance (Rytwinski et al., 2019).

Effective strategies for controlling invasive species must consider two important aspects. First, the most efficient, cost-effective and safe means of removing individuals from a local population must be determined, which will vary by the invader's life stage and the ambient environmental conditions (Buhle et al., 2005). Second, the timing and location of the control measure needs determining. For example, in invasive plant management, there is debate about whether efforts should be focused at the range front, where the invader is of low abundance and patchy in distribution, or at the invasion core, where populations are established and usually more abundant (e.g. Hastings et al., 2006; Januchowski-Hartley et al., 2011). Invader dispersal rates and abundances can also be affected by habitat complexity, with terrestrial invaders generally spreading more easily in unfragmented than fragmented landscapes (Dewhurst & Lutscher, 2009). Therefore, understanding the ease with which

invaders spread in different environmental configurations needs consideration within management planning (Lurgi et al., 2016).

In managing freshwater invaders, the utility of individual-based models (IBMs) to predict rates of establishment and spread has recently been highlighted (Dominguez Almela et al., 2020), along with comparisons of the methods for controlling and/or eradicating local populations (Rytwinski et al., 2019). There is, however, less known on how invader dispersal rates affect the long-term efficacy of population control, especially at large spatial scales and in open, linear systems. In rivers, while controlling invasive fish can be a management priority, their removal usually relies on capture methods (e.g. nets, traps, electro-fishing; Rytwinski et al., 2019). Although these are effective at capturing (and culling) fish species across most of their size range (Davies & Britton, 2015), they are unlikely to remove all the population, and surviving individuals can then potentially compensate for losses (Berry et al., 2012). This can result in culled populations rapidly recovering to their previous abundances and management objectives not being met (Davies & Britton, 2015; Dominguez Almela et al., 2020).

Here, our aim was to develop an IBM to predict the effects of management control strategies on invasive river fishes. We applied it within factorial experiments to ‘virtual’ invasive fish with a range of life history traits in river networks of varying spatial complexity to predict how different culling strategies affected fish spread and abundance. The objectives were to predict: (1) how altering the application of culling (rate/location/timing/life-stage) affected the abundance and spatial distribution of an initial invasive fish, based on the demographic characteristics of bitterling *Rhodeus sericeus*, a small bodied invasive fish (Dominguez Almela et al., 2020); (2) how invader abundances and distributions varied across nine virtual invasive fish differing in life history traits and dispersal abilities; and (3) the trade-offs between management effort (cull rate and the number of patches culled) and the predicted invader abundances and distributions to optimize management responses. The final model was then applied to a real-world scenario to predict how management interventions could have constrained the real-world invasion of bitterling in the River Great Ouse, Eastern England (Dominguez Almela et al., 2020).

2. Material and methods

2.1. *Model configuration and virtual species*

The model was implemented in a customized version of the individual-based spatially explicit modelling platform *RangeShifter* (Bocedi et al., 2014, 2020), incorporating a new module for managing invasive species. River catchments of an overall mean extent of 74.5 ha and of three basic types were created using ArcGIS Pro, where the configuration varied by the extent of its tributary branching: high branching (12 tributaries), moderate branching (8 tributaries) and low branching (4 tributaries). Each catchment type was replicated three times in slightly different configurations (Fig. 1). All catchments were created in raster format at 50 m resolution, and each cell was assigned a habitat quality score proportional to stream width (Dominguez Almela et al., 2020). Catchments comprised a mean of 298 (SD: ± 20) cells divided into a similar number of non-overlapping patches (mean = 44.9 patches/catchment, SD = ± 0.3), which were delineated manually such that no patch overlapped a confluence. The ratio of good to poor quality habitat sections was similar in all configurations; in the main stem, it was kept consistent at 1:1, and in the tributaries, the ratio was maintained across replicates within the branching groups (low/moderate/high) at an overall mean of 8:15.

The initial virtual fish, whose demographic and dispersal parameters were drawn from bitterling (*cf.* Dominguez Almela et al., 2020), had the characteristics of an alien fish that followed a stage-structured population dynamic: juveniles (<1 year old), sub-adults (1 to 2 years) and adults (over 2 years). Its population parameters (Table 2, Table S1) were also similar to other small-bodied, invasive, pest fish species, such as topmouth gudgeon *Pseudorasbora parva* (Britton & Gozlan, 2013). It was set to reproduce once per year, exhibit density-dependent fecundity and have a sex ratio at birth of 1M:1F. Its survival probability was also density-dependent, applied and weighted per stage, so the effect of sub-adults on the survival of adults was 10 % of the effect of adults on each other and on sub-adults (Dominguez Almela et al., 2020). After reproduction, juveniles, sub-adults and adults could disperse according to a density-dependent emigration probability, limited to one dispersal event

per lifetime. Movement from the natal patch during the transfer phase of dispersal was modelled by the stochastic movement simulator (SMS; Palmer et al., 2011), which simulates movement from cell to cell on the basis of perceived costs within a limited perceptual range and a tendency to follow a correlated path (directional persistence). A relative cost of movement map was derived from the habitat map such that perceived costs were inversely related to habitat quality; thus, upon reaching a confluence, a disperser would more likely move into the wider of the two streams available. Individuals could settle in any non-natal patch subject to an inverse density-dependent settlement probability (Table 2, Table S1).

For all model simulations, initial populations were established in 10 patches (including in tributaries) at the upstream end of the catchment (Fig. 1). These were mainly in first and second order streams, the rationale being that this enabled the invasion front to spread mainly downstream into streams of higher order (Kim et al., 2021), although the direction of individual movement was stochastic in the model and could occur in either direction unless the fish was in a terminal patch. Propagule pressure by the random addition of further individuals into the system was not considered in the model. Each simulation (a single combination of culling parameters and catchment; Table 1) was run for 30 years and replicated five times. A ‘control’ simulation was run for each catchment in which no management was applied.

2.2. *Experimental designs*

The management of the modelled populations took the form of an annual cull, the format of which was controlled by a series of variable parameters. A specified number of patches was selected each year for culling in which individuals were subjected independently to a mortality probability determined as a stage-dependent logistic function of density and subject to a maximum culling rate. This means that the proportion of individuals removed from a patch per year (the ‘cull rate’) would depend on its density and low invader numbers would translate into low removals reflecting the difficulties of catching those individuals when the population is very low. In contrast, if the density were high, the logistic function would result in removal of a higher proportion of individuals up to a pre-defined maximum. Both the culling rate and the number of patches selected were kept constant across years. We modelled the cull as occurring after dispersal and its spatial application as biased toward recently colonized patches based on results from preliminary experiments (see Appendix S1-3 in Supporting Information). Then, the first experiment assessed the interaction of culling rate and number of patches culled by varying the cull rate on an initial virtual species across different numbers of patches (Experiment 1, Table 1). The second experiment varied the demographic (as type of demographic species, SpType) and dispersal (as type of dispersal species, SpDispType) characteristics of the model alien species that was being managed (Experiment 2, Table 1). Finally, in the third experiment, we applied the model to a real case study using bitterling as the model species and the Great Ouse as the model catchment, following Dominguez Almela et al. (2020).

Experiment 1: Interaction of culling and management effort

Six levels of the maximum number of patches culled (N), ranging from low to high numbers (4 to 24) of patches culled per year, were tested against 8 levels of maximum cull rate (CR, 0.2 to 0.9) (Table 1). The cull was applied to all three life stages simultaneously. The experiment was run for all 9 catchments (432 parameter combinations).

Following the model simulations, the results of Experiment 1 were analysed using two population-level summary statistics that were extracted from the model output data:

(1) Rate of population increase (P_1):
$$P_1 = \frac{Nind_{t+n} - Nind_t}{n},$$

where $Nind_t$ is the total number of individuals across the catchment at time t and n is the number of years during which there was active growth. The simulated population when no culling was applied was used to determine n , the number of years of approximately constant growth before the population growth trajectory began to decline.

(2) Rate of change in patch occupancy (Q_1): $Q_1 = \frac{NOccupPatches_{t+n} - NOccupPatches_t}{n}$,

where $NOccupPatches_t$ is the number of occupied patches at time t and n is the number of years for the period of active growth as above.

Experiment 2: Trade-off between the number of patches culled and the cull rate within each patch across a range of species

This experiment evaluated a potential resource-limited trade-off between the maximum number of patches culled and the maximum cull rate within each patch across a further eight contrasting virtual species (specific culling strategy; SCS), to identify whether patterns detected in the initial species were common across species with contrasting life history traits and dispersal abilities (Table 1). It was based on the assumption that the total management resource (finance, man-power) was fixed annually, and so managing more patches in a year would result in reduced effort per patch, reducing the effects of the culling on the invader. For example, culling all patches would spread resources thinly across the whole catchment, so the lowest maximum cull rate ($CR = 0.1$) would have to be applied per patch in each year in this scenario (Table 1). The initial virtual species (Experiment 1) was the reference species, having intermediate demographic traits and dispersal abilities, and 8 additional virtual species were developed that varied by their: (i) demographic traits and/or (ii) dispersal traits. (i) could go from being more intensely r -selected (i.e. higher fecundity, lower survival) to more intensely K -selected (lower fecundity, higher survival). (ii) varied from being better dispersers by increasing stage-dependent maximum emigration probabilities (more dispersers per generation) and decreasing maximum settlement probability (more likely to keep moving), to poorer dispersers by decreasing stage-dependent maximum emigration probabilities (fewer dispersers per generation) and increasing maximum settlement probability (less likely to keep moving) (Table 2). The experiment was applied to all 9 catchments (729 parameter combinations).

As there was substantial variation between population trajectories in the experimental predictions, fixing a specific year as the basis for calculating the rate of population increase (P_1) and change in patch occupancy (Q_1) was inappropriate. Therefore, for each replicate simulation trajectory, the rate of population increase in the first decade (P_{10}) was calculated, using:

$$P_{10} = \frac{pop[year = 10] - pop[year = 1]}{10}$$

where $pop[year=10]$ is the number of individuals at year 10, and $pop[year=1]$ is the number of individuals at year 1, and similarly for years 6 to 15 inclusive to give P_{15} ; 11 to 20 (P_{20}), 16 to 25 (P_{25}) and 21 to 30 (P_{30}). Finally, the maximum mean annual increase in population size achieved was determined as:

$$P_2 = \max(P_{10}, P_{15}, P_{20}, P_{25}, P_{30})$$

The same approach was used to calculate the maximum decadal rate of change in patch occupancy (Q_2):

$$Q_{10} = \frac{NOccupPatches[year = 10] - NOccupPatches[year = 1]}{10}$$

Where, Q_{10} is the rate of change in patch occupancy for the years 1 to 10 inclusive,

$NOccupPatches[year=10]$ is the no. of occupied patches at year 10, and $NOccupPatches[year=1]$ is the no. of occupied patches at year 1, etc.

Experiment 3: Case study on specific management to control bitterling in the Great Ouse

The predictive performance of our approach was evaluated using bitterling as the model invader and the River Great Ouse in Eastern England as the model river basin. The map of the catchment (including its division into 272 patches) and the key model parameters were taken from Dominguez et al., (2020), where the parameter estimates used were from their posterior distributions obtained by approximate Bayesian computation. The full posterior parameter distributions (from 250 simulations) were used to determine the level of culling needed for reducing the bitterling spatial range, using the culling scenarios applied in Experiment 2 (SCS; Table 1), giving a total of 2,250 parameter

combinations. The simulations started in 1983 when the species was first recorded in fisheries monitoring surveys (Dominguez Almela et al., 2020) and ran for 100 years, producing 5 replicates per set of parameter combinations. Values of rate of population increase (P_2) and change in patch occupancy (Q_2) were calculated, with the mean of each set of 5 replicates providing a single prediction for each posterior parameter combination per SCS, allowing the confidence intervals for variation only between posterior parameter sets.

2.3. *Statistical analyses*

For both P_n and Q_n , factorial linear models for each experiment were fitted using R version 3.6.3 (R Core Team, 2020) to partition the variance in the response variable. The models incorporated, as appropriate, the management scenarios that were applied (N, CR, SCS), the three sets of river catchments (low, medium and high branching; B, ID) and species (SpID, SpType, SpDispType) (Table 1). The effect of the factor(s) that had the greatest influence on model outcomes were investigated further using posterior marginal means analyses (package ‘emmeans’, Lenth, 2020).

Table 1 Factorial design of simulated management experiments. Factors: number of patches culled (N); maximum culling rate (CR); specific culling strategy (SCS).

Experiment 1									
N	4	8	12	16	20	24			
CR	0.9	0.8	0.7	0.6	0.5	0.4	0.3	0.2	
Experiment 2									
SCS	CR=0.9 N=5	CR=0.8 N=9	CR=0.7 N=14	CR=0.6 N=18	CR=0.5 N=23	CR=0.4 N=28	CR=0.3 N=32	CR=0.2 N=37	CR=0.1 N=41
Variation of species	3 strongly <i>K</i> -selected species (1 'fast-disperser', 1 'intermediate-disperser', 1 'slow-disperser')								
	3 with similar demographic traits as for experiment 1 (1 'fast-disperser', 1 'intermediate-disperser', 1 'slow-disperser')								
	3 strongly <i>r</i> -selected species (1 'fast-disperser', 1 'intermediate-disperser', 1 'slow-disperser')								
Experiment 3									
SCS	CR=0.9 N=5	CR=0.8 N=9	CR=0.7 N=14	CR=0.6 N=18	CR=0.5 N=23	CR=0.4 N=28	CR=0.3 N=32	CR=0.2 N=37	CR=0.1 N=41

Table 2 The model parameters of the nine virtual species used in Experiment 2. See Table S1 for parameters common to all species. *Species 5 was the same as the single species used in Experiment 1 (and in preliminary experiments 0a, 0b in Supplementary Information). **1/b is the rate of density dependence, i.e. the rate at which mean fecundity decreases with increasing local density.

Parameter	Strongly <i>K</i>- selected species	Intermediate demography	Strongly <i>r</i>- selected species
Fast disperser	Species 1	Species 2	Species 3
Fecundity	30	63.77	180
1/b** (inds/ha)	823.73	1750.96	4942.34
Survival probability in juveniles	1	0.93	0.8
Survival probability in sub-adults	1	0.89	0.2
Survival probability in adults	0.5	0.4	0.1
Max. emigration probability in juveniles	0.4	0.4	0.4
Max. emigration probability in sub-adults	0.7	0.7	0.7
Max. emigration probability in adults	0.9	0.9	0.9
Max. settlement probability	0.6	0.6	0.6
Per-step mortality	0.01	0.01	0.01
Intermediate dispersal	Species 4	Species 5*	Species 6
Fecundity	30	63.77	180
1/b (inds/ha)	823.73	1750.96	4942.34
Survival probability in juveniles	1	0.93	0.8
Survival probability in sub-adults	1	0.89	0.2
Survival probability in adults	0.5	0.4	0.1
Max. emigration probability in juveniles	0.1	0.1	0.1
Max. emigration probability in sub-adults	0.18	0.18	0.18
Max. emigration probability in adults	0.18	0.18	0.18
Max. settlement probability	0.84	0.84	0.84

Per-step mortality	0.01	0.01	0.01
Slow disperser	Species 7	Species 8	Species 9
Fecundity	30	63.77	180
1/b (inds/ha)	823.73	1750.96	4942.34
Survival probability in juveniles	1	0.93	0.8
Survival probability in sub-adults	1	0.89	0.2
Survival probability in adults	0.5	0.4	0.1
Max. emigration probability in juveniles	0.01	0.01	0.01
Max. emigration probability in sub-adults	0.05	0.05	0.05
Max. emigration probability in adults	0.14	0.14	0.14
Max. settlement probability	1	1	1
Per-step mortality	0.01	0.01	0.01

3. Results

Experiment 1: Interaction of culling and management effort

The predicted effects of culling on both P_1 and Q_1 increased as both the maximum cull rate (CR) and maximum number of patches (N) increased, which together accounted for most of the variation (Tables 3, S2b, S3b). When CR was low, it had only minor effects on P_1 and Q_1 , except when N was high (Fig. 2). When CR was increased to a medium level (0.5, 0.6), then values of P_1 and Q_1 suggested that culling could contain the spread of the invader, but when CR was ≥ 0.7 , eradication of the species was possible if N was also high (at least 16 patches culled per year) (Fig. 2). All replicates led to eradication when CR was at least 0.8 and N was ≥ 12 patches per year, and also when CR was 0.9 and N = 8 patches per year.

Table 3 Principal sources of variance explained (%) in the summary statistics rate of population increase (P_1) and rate of change of patch occupancy (Q_1) for experiment 1. Factors: maximum number of patches culled (N); maximum cull rate (CR); catchment branching (B); catchment ID number (ID).

	Catchment related factor		Management related factor		
	B	ID	N	CR	CR*N
P_1	4.5	4.2	22.4	44.3	13.5
Q_1	1.4	0.7	23.2	51.7	17.5

Experiment 2: Trade-off between the number of patches culled and the cull rate within each patch across a range of species

In general, the major source of variation in P_2 and Q_2 was from differences between the characteristics of the nine species (SpType; SpDispType; SpID) (Tables 4, S2b, S3b). Marginal means analysis between species type (SpType; SpID) and the SCS revealed that P_2 and Q_2 could be substantially reduced when N was set at a relatively high number of patches, despite culling being at less than maximum efficiency (Fig. 3, S4). Five out of the nine species presented at least two scenarios in which the virtual species was eradicated in some of the replicates (Table 5). The scenario where 0.8 cull rate (CR) was applied in 9 patches (N) had the highest number of replicates across species in which eradication occurred (152/729 replicates). No species was eradicated at $CR < 0.7$. Of the more r -selected species group, 82% of slow and 28% of intermediate dispersers were successfully eradicated during simulations at $CR \geq 0.7$ (Table 5). In particular, the scenario where 0.8 cull rate was applied to 9 patches predicted that for the r -selected slow dispersers, there were no replicates where the species was not eradicated. In r -selected fast dispersers, P_2 and Q_2 were reduced, but their populations were only predicted to be eradicated in a single replicate (Fig. 3, S4). Conversely, 18% of intermediate and 61% of slow dispersers within K -selected species were successfully eradicated in simulations at $CR \geq 0.7$, while fast disperser abundances were reduced but were never fully eradicated at any of the 9 CR scenarios (Table 5; Fig. 3, S4).

There were also marked differences in the response of P_2 and Q_2 in relation to the different dispersal abilities of the nine species (SpDispType) (Fig. 4). Species with slow dispersal abilities required less effort to control, enabling lower numbers of patches to be culled to achieve similar outcomes as intermediate or fast dispersers at higher numbers of patches. Moreover, the intermediate and fast dispersers revealed some compensatory responses to the culling, suggesting their invasion could actually benefit from some culling scenarios through increased P_2 and Q_2 .

Table 4 Principal sources of variance explained (%) in summary statistics rate of population increase (P_2) and rate of change of patch occupancy (Q_2) for Experiment 2. Factors: type of demographic species (SpType); type of dispersal species (SpDispType); species ID number (SpID) and specific culling strategy (SCS).

	SpType	SpDispType	SpID	SCS	SpType*SCS	SpDispType*SCS
P_2	33.21	34.13	6.37	11.09	3.61	1.71
Q_2	4.62	56.87	1.12	20.42	0.95	3.69

Table 5 Counts of replicates per scenario in which virtual species were eradicated. The maximum number of replicates possible per scenario and species is 45. Slow dispersers: Species 7, 8 and 9; Intermediate dispersers ('Inter.'): Species 4 and 6; Fast disperser: Species 3. All other species had no scenarios where they were eradicated

Scenario	Strongly <i>K</i> -selected		Intermediate demography	Strongly <i>r</i> -selected		
	Species 4	Species 7	Species 8	Species 3	Species 6	Species 9
Dispersal ability	Inter.	Slow	Slow	Fast	Inter.	Slow
0.7 cull rate (CR), 14 patches (N)	4	2	0	1	18	23
0.8 cull rate (CR), 9 patches (N)	19	43	25	0	20	45
0.9 cull rate (CR), 5 patches (N)	1	37	7	0	0	43

Experiment 3: Case study on specific management to control bitterling in the River Great Ouse

Our IBM predicted a significant response in this bitterling population to the different culling scenarios in both P_2 ($F_{8,2241} = 165.23$, $p < 0.001$) and Q_2 ($F_{8,2241} = 1953.5$, $p < 0.001$) (Fig. 5) analogous to that predicted in Experiment 2 for intermediate demography/dispersers (*cf.* Fig. 4). Where Dominguez Almela et al. (2020) predicted bitterling would occupy 90% of patches in the river in 2045 (Fig. 6a), the application of a yearly SCS of $CR = 0.7$ and $N = 14$ (5 % of patches) was predicted here to result in the population occupying only 21% of the area, which was similar to their spatial extent recorded in 1984 (Fig. 6b).

4. Discussion

Our model predictions provide a series of novel insights that should assist decision-making for managers dealing with invasive fish specifically and invasive species more generally. Our IBM identified the optimal management effort (as specific culling strategies involving the cull rate and its spatial application) for invaders across a range of demographic traits and dispersal abilities. The model predictions demonstrated that while eradication of invasive fish is possible by culling, it requires relatively high cull rates to be used, with the highest probability of eradication achieved in fish of low dispersal abilities. For fast-dispersing fish with both r - and K -selected characteristics, eradication was predicted to be difficult to achieve irrespective of the cull rate used, although the predictions did indicate that the spatial spread and abundance of the invader can be constrained. Correspondingly, for fast dispersing invasive fishes, the predictions suggest it is imperative for management control efforts to be implemented rapidly and utilise high cull rates where feasible. Such management insights have, so far, been unobtainable from meta-analyses of empirical data (e.g. Rytwinski et al., 2019). Moreover, when the IBM was applied to invasive bitterling in the River Great Ouse, England, the spatial extent of their invasion by 2045 (90% of the catchment) was predicted to be constrained to their 1984 distribution (21% of the catchment) using an annual cull rate of 0.7 in 14 patches (around 5% of the catchment). This indicates that a long-term management control strategy could substantially constrain the invasion of an alien fish in a relatively open system, even if the species could not be eradicated.

Our IBM predicted that different culling strategies result in a range of effects on the population growth and spread of invasive species. If the annual specific culling strategy was held at a fixed level across all years, then increasing the number of patches culled per year must be matched by a reduction in the cull rate per patch. The predictions indicated that a reduced cull rate, even when applied over larger spatial areas, was generally ineffective in constraining the fish dispersal and population growth rates. Correspondingly, the application of low cull rates across large spatial areas is not an effective management option. In contrast, relatively high cull rates often predicted population eradication, even

when the number of patches culled were not necessarily high (e.g. a cull rate of 0.9 in only 8 patches). These predictions are thus important in the context of management planning of how specific culling strategies could be applied spatially and in relation to the ability of capture methods to remove high proportions of the target species within managed patches. They also demonstrated the compensatory responses that will occur within the fish population when the cull rate is too low, whereby the reduction in population size results in subsequently higher reproductive rates and abundances, and then faster dispersal (Berry et al., 2012).

Although the IBM was effective at predicting how the cull rate affects the abundance and dispersal of the invader, determining the extent of a fish population that can be removed effectively from a population using capture methods, such as electric fishing, can be difficult. While electric fishing is considered an effective fish capture technique, it has inherent issues relating to species detectability and detection bias (Beaumont, 2016). Its probability of capture is species-dependent with, for example, it varying between 0.35 and 0.64 across 15 stream fish species (Reid et al., 2009). It can also be relatively ineffective at capturing early life-stages of fish, where alternative methods might be more effective, such as micro-mesh seine nets and/or traps (Nunn et al., 2001; Britton et al., 2011a). Nevertheless, electric fishing is regularly and successfully used to remove alien fish from invaded waters when chemical treatments cannot be applied, with the meta-analysis of Rytwinski et al. (2019) reporting a 58% success rate for population eradication and 56% for population control over a range of alien fish species.

The predicted outcomes of the specific culling strategies on the invading populations were also strongly influenced by the invader's demographic traits. The initial virtual species, based on bitterling, had a suite of demographic traits that were similar to those of small-bodied invasive fish more generally that are considered pests in many parts of the world and so often receive considerable management attention (e.g. Britton & Brazier, 2006; Britton et al., 2010). However, invasive fish with demographic traits that are less intensively *r*-selected or even *K*-selected are also considered undesirable in many parts for the world. For example, relatively large-bodied alien fish of the Salmonidae family are often targeted for management in North America (Rytwinski et al., 2019) and

common carp *Cyprinus carpio* have also received considerable management attention in many areas of the world, including Australia (Pinto et al., 2005) and South Africa (Davies et al., 2020). It was thus important to understand how these initial predictions varied according to demographic traits and dispersal abilities. The predictions revealed that population eradication was achievable in five of the nine virtual species, including species with both *r*- and *K*-selected traits, but only when the cull rate was relatively high (≥ 0.7) and when at least 14 patches were culled. However, cull rates below 0.7 never resulted in eradication. That invasive fish with *r*-selected demographic traits were predicted to be the most challenging to control and eradicate is arguably an intuitive result, as these traits, common in many invasive fish, enable the rapid development of highly abundant populations where individuals can then disperse (Gozlan et al., 2010). The eradication of two strongly *K*-selected virtual species was predicted, but only when they were intermediate or slow dispersers, and at a high cull rate. The real-world simulation of bitterling in the River Great Ouse indicated that although population eradication would not occur under any of the specific culling strategies that were simulated, strong containment was possible.

These invasion management predictions should be applied to informing real-world invasion risk assessment processes (Vilizzi et al., 2019), which are fundamentally important in prioritising the species and habitats for management (Roy et al., 2018). Best-practice guidance on risk assessments for alien species indicates that two of the four main assessment components are their probable spread and impact (Roy et al., 2018). In our summary analysis of the IBM output data, we directly predicted invader spread, and the parameter P_n can potentially predict impact due to the relationships of invader abundance with impact (Jackson et al., 2015). Consequently, provided some basic knowledge of the demographic and dispersal traits is available for a specific species, our IBM predictions should inform the risk assessment responses on their spread and impact, especially where there is strong understanding of the invader's demographic traits and dispersal abilities. This is because there were some strong differences in the responses of *r*- and *K*-selected species, and across varying dispersal abilities, to the different culling strategies. For example, 82 % of slow dispersing *r*-selected species were eradicated at cull rates above 0.7 versus 61 % of *K*-selected species, while eradication of fast dispersing species was generally predicted to be unlikely. Although detailed information on the

dispersal abilities of alien fish is often lacking, dispersal patterns are also likely to be catchment specific, varying according to, for example, the flow regime, river network complexity and habitat suitability, and the extent of human modification of them (Caiola et al. 2014). Correspondingly, following the detection of a new alien fish within a river catchment, practitioners should rapidly assess its life history traits (either directly or through literature review), its ability to disperse in the catchment (in relation to both the species' dispersal characteristics and the complexity of the river network), and its current spatial extent (Britton et al., 2011b). Concomitantly, the resource available for the control effort needs to be quantified, with identification of the desired management outcome (eradication versus control). In combination with the predictions outlined here across the different fish life histories and dispersal abilities, and the culling gradients, this information should then enable more informed decision making on the actual strategy to be implemented.

Our predictions thus provided considerable insights into how fish removal efforts and the traits of the target species interact to alter the outcomes of invasive fish culling strategies. Such insights are not possible from empirical studies due to their context-dependencies, and often poor experimental planning (e.g. lacking baseline data and control sites; Rytwinski et al., 2019). However, we acknowledge that our modelling processes have some inherent limitations. For example, while GIS techniques can accurately represent a linear river system, the division of the river into discrete patches, as required by *RangeShifter*, has some subjectivity and does not faithfully represent the natural heterogeneity of actual rivers (Dominguez Almela et al., 2020). While our general results should be qualitatively robust to this discretization, future work exploring the consequences of model choices related to the resolution of the spatial grid would be valuable. Also, model initialisation always introduced the fish to the catchment's upper reaches, enabling the invasion front to disperse downstream into higher order streams, where established populations of invasive fishes are often abundant (Kim et al., 2021). In reality, the release points of alien species are more stochastic, such that some dispersing aliens move from downstream to upstream areas (Vitule et al., 2012). The model also prohibited an individual from dispersing more than once in its life (Bocedi et al., 2014), an assumption potentially violated by some riverine fishes (Fausch et al., 2002; Radinger & Wolter, 2014). Moreover, the model was based on a discrete introduction of fish, and whilst invasions can

occur from a single release event, multiple releases can also occur, which can increase the probability of invasion success as it overcomes issues such as founder effects (Lockwood et al., 2005). However, a key component of eradication attempts of invasive species is preventing their re-introduction into the treated area, and so where management efforts are on-going to control invaders, these efforts should also include increased regulation and surveillance that aim to prevent further releases (Britton et al. 2011b). Finally, throughout the model, the total specific culling strategy was fixed across all years, as this enabled the model to account consistently for the relationship between cull rate and the number of patches culled. However, it is acknowledged that in reality, resource availability might vary by year and, for high priority species, be increased in the short-term. Moreover, it has already been discussed that the actual effort required to achieve the higher cull rates might exceed the levels of effort used in the model due to the increased degree of difficulty of capturing fish from small populations (Britton et al., 2011a). Consequently, while there is high confidence in our predictions of how the different cull rates applied across varying number of patches culled will affect the population of the target species, it is suggested that some caution is applied when considering that these strategies will all be of equal cost. Despite these issues, we argue strongly that the strength of our model is its ability to simulate the outcome of management interventions to control populations of invasive species that cannot be provided empirically (Dominguez Almela et al., 2020). It thus represents a major step forward in understanding how to develop strategic approaches for managing alien species in the environment.

In summary, this work provided insights into the outcomes of different control efforts on invasive fishes, and highlighted that, depending on the species characteristics, and the specific culling strategy, these outcomes can vary in target populations, but can include eradication when the target species is of low or intermediate dispersal ability and when a high cull rate is applied. These predictions strongly complement existing invasion risk assessments, and demonstrate that individual-based models are powerful tools for predicting optimal management interventions for high-risk invaders.

Conflict of interest

None of the authors have any conflicts of interest. J. Robert Britton is an Associate Editor of Journal of Applied Ecology but took no part in the peer review and decision-making processes for this paper.

Author contributions

All authors contributed to the study design. VDA led analyses and writing, assisted by SCFP, PKG, DA, JMJT and JRB. All authors contributed to revising the manuscript and approved its submission.

Acknowledgements

The *RangeShifter* software and manual can be downloaded from <https://rangeshifter.github.io>. VDA was supported by an iCASE studentship from the Natural Environment Research Council (NE/R008817/1) and the Environment Agency. SCFP and JMJT were supported by the Latin American Biodiversity Programme as part of the Newton Fund (NE/S011641/1), with contributions from the Natural Environment Research Council.

Data availability statement

Bitterling/ Ouse data were taken from this open data: <https://data.gov.uk/dataset/f49b8e4b-8673-498e-bead-98e6847831c6/freshwater-fish-counts-for-all-species-all-areas-and-all-years>. All other data used are available in the main text and supporting information.

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Figure Legend Section

Figure 1 Virtual catchments having a low number of tributaries (a1-3), medium (b1-3) and high (c1-3). Grey thick lines highlight the initialized patches in the upper areas of the catchment, and the arrows indicate the direction of flow along the main stem.

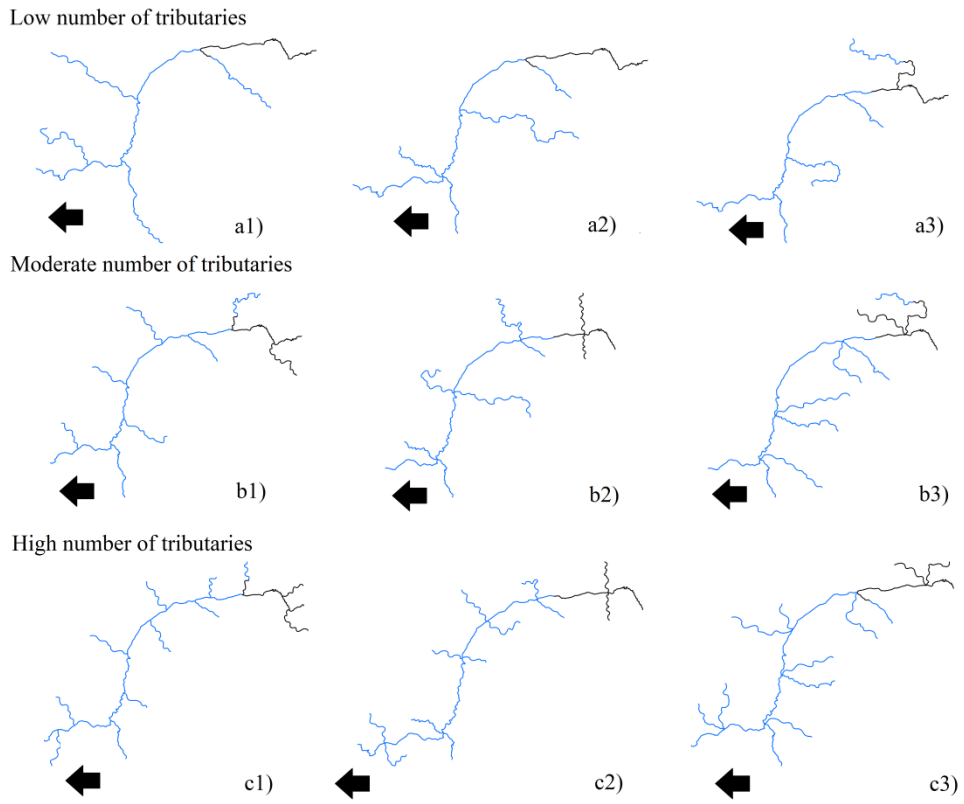
Figure 2 Interaction effects of the maximum cull rate (CR) and maximum number of patches culled (N) on the rate of population growth (P_1) and change of patch occupancy (Q_1) in Experiment 1.

Figure 3 Interaction effects of the species dispersal type (SpDispType) and specified culling strategy on the rate of change of patch occupancy (Q_2) during Experiment 2. No data-point shown at 80% (9 patches) in the *r*-selected species as all replicates were extirpated.

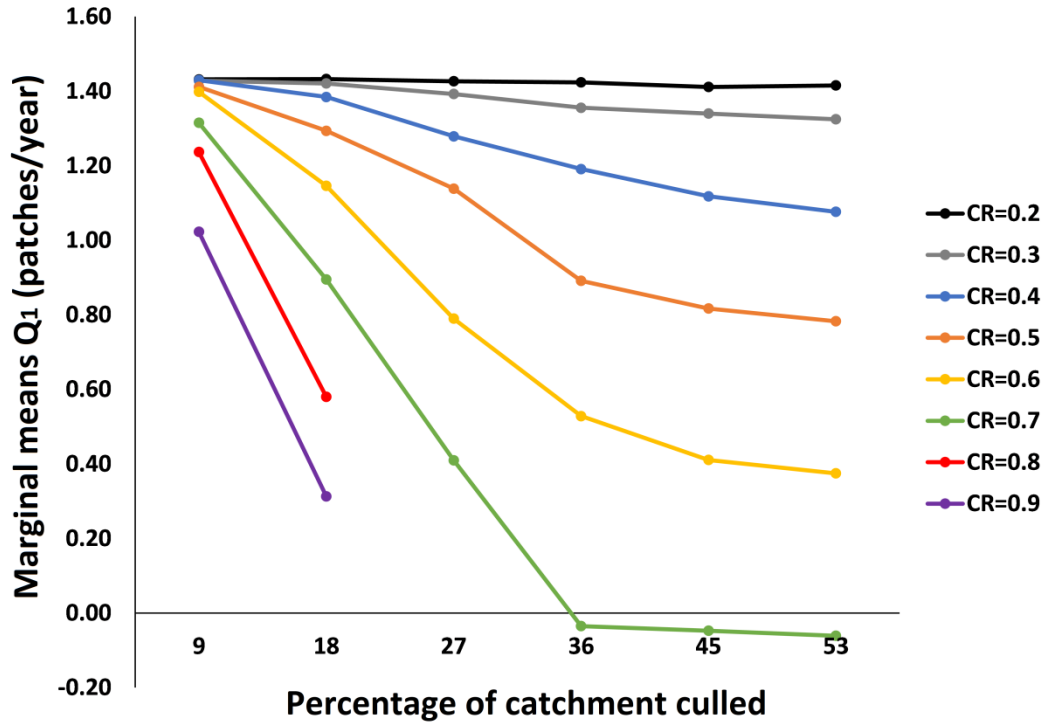
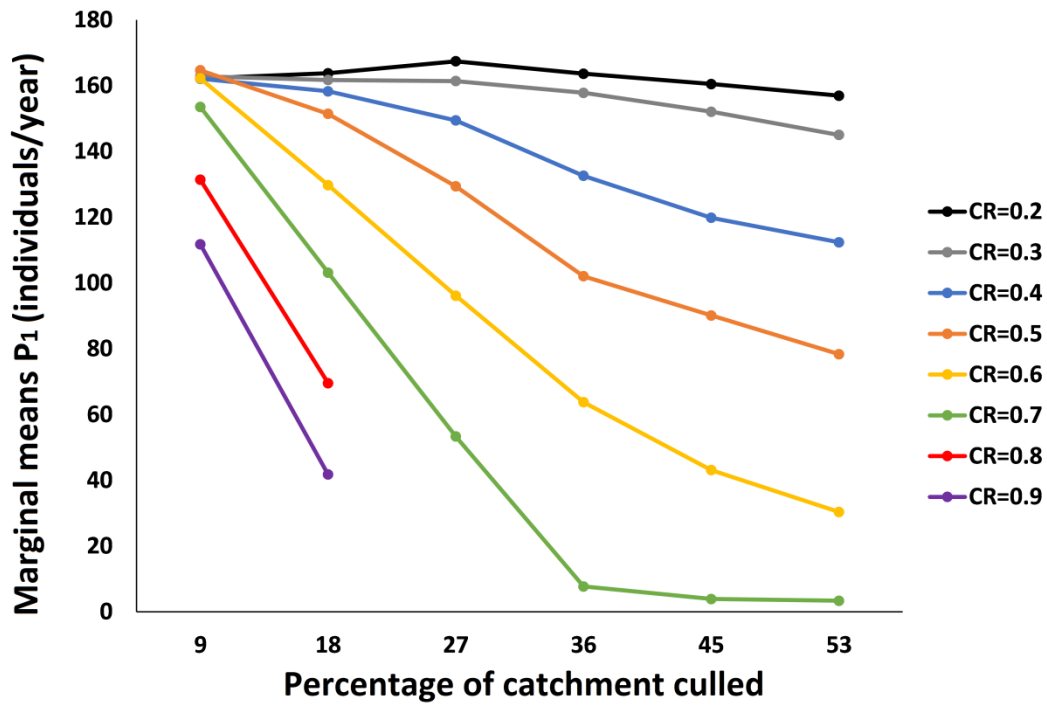
Figure 4 Interaction effects of the species dispersal type (SpDispType) and specified culling strategy on the rate of population increase (P_2) and change of patch occupancy (Q_2) for populations which were not eradicated during Experiment 2.

Figure 5 Effects of specified culling strategy on the rate of population increase (P_2) and change of patches occupancy (Q_2) of the bitterling species. Grey bars are 95% confidence intervals for the marginal means.

Figure 6 Comparison map with the predicted occupancy (by 2045) of the catchment (a) without management control (Dominguez Almela et al., 2020) and (b) after applying a yearly SCS of CR = 0.7 and N = 14. Colour-scale: probability of occupancy ranging from low (light grey) to high (dark blue). © Crown copyright and database rights 2018 Ordnance Survey (100025252).

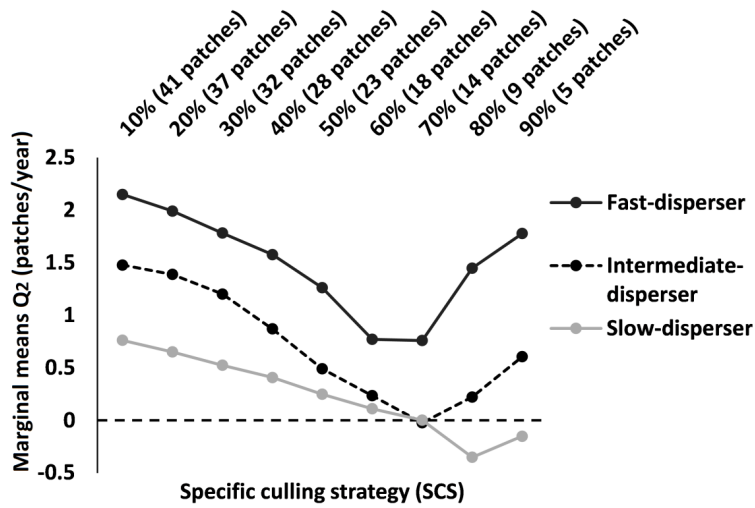


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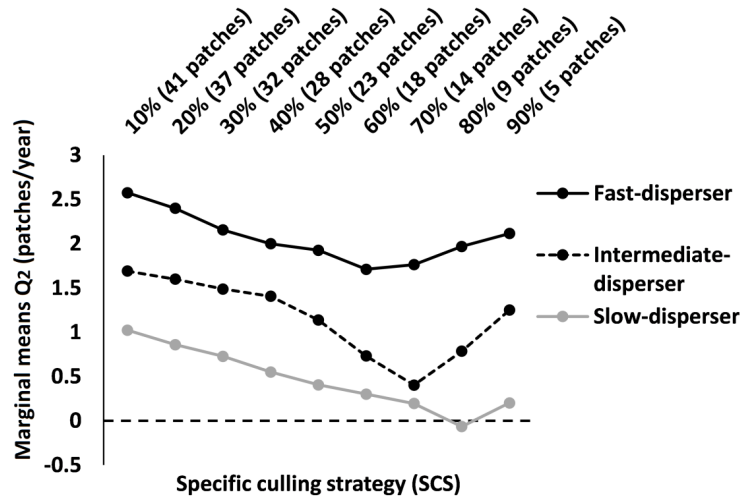


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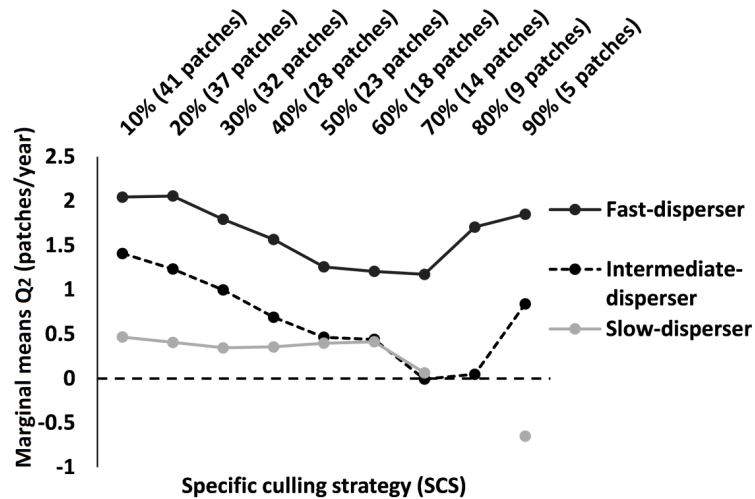
Strongly *K*-selected species

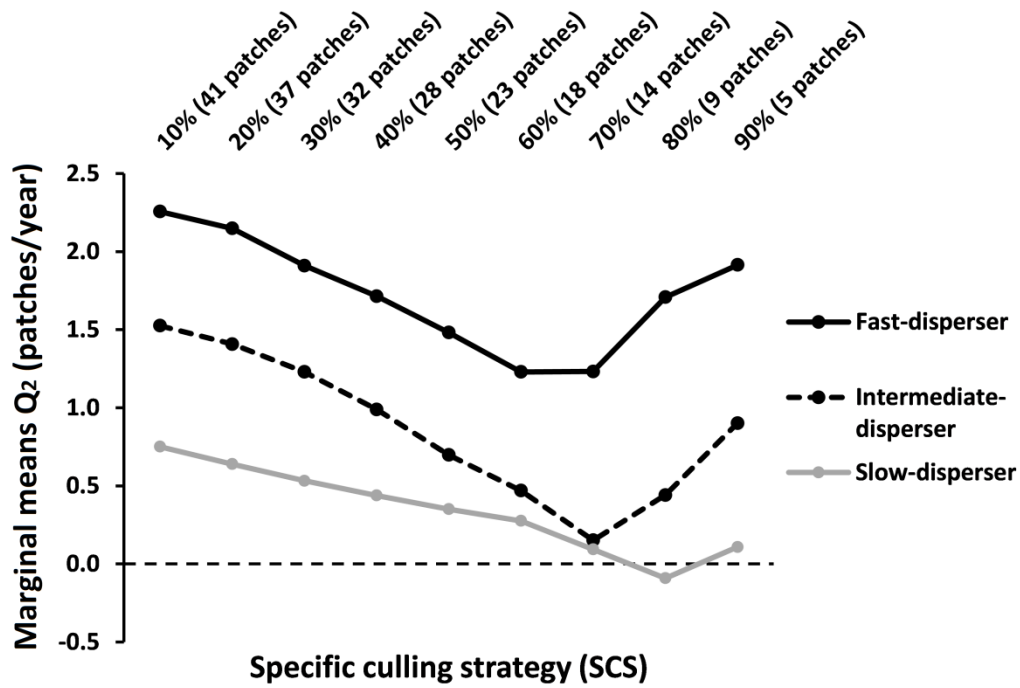
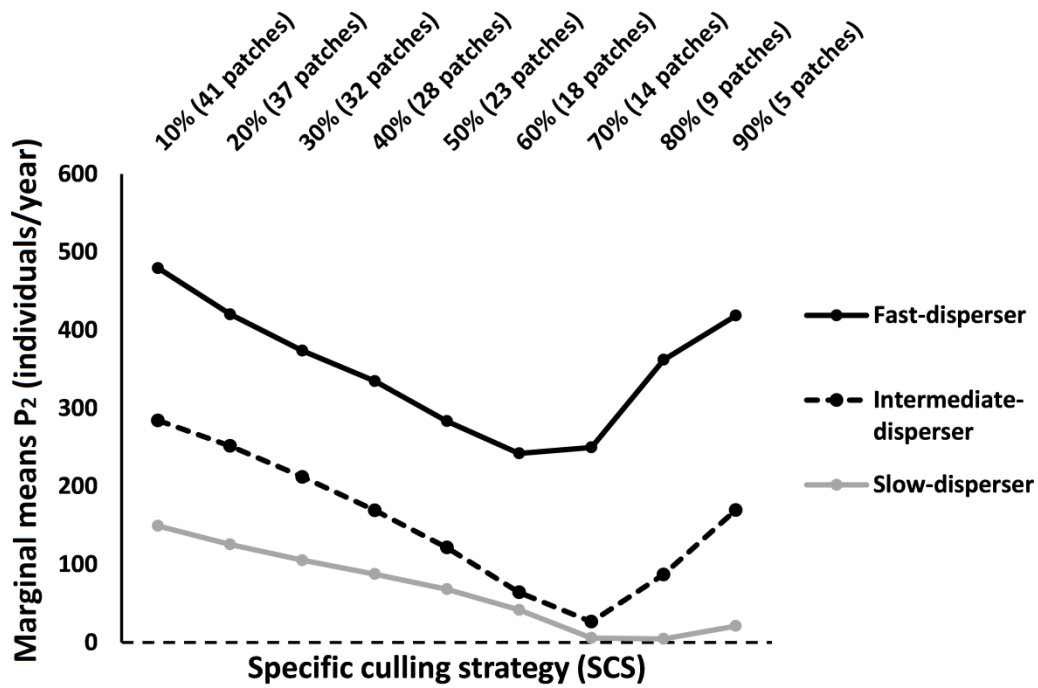


Intermediate demography

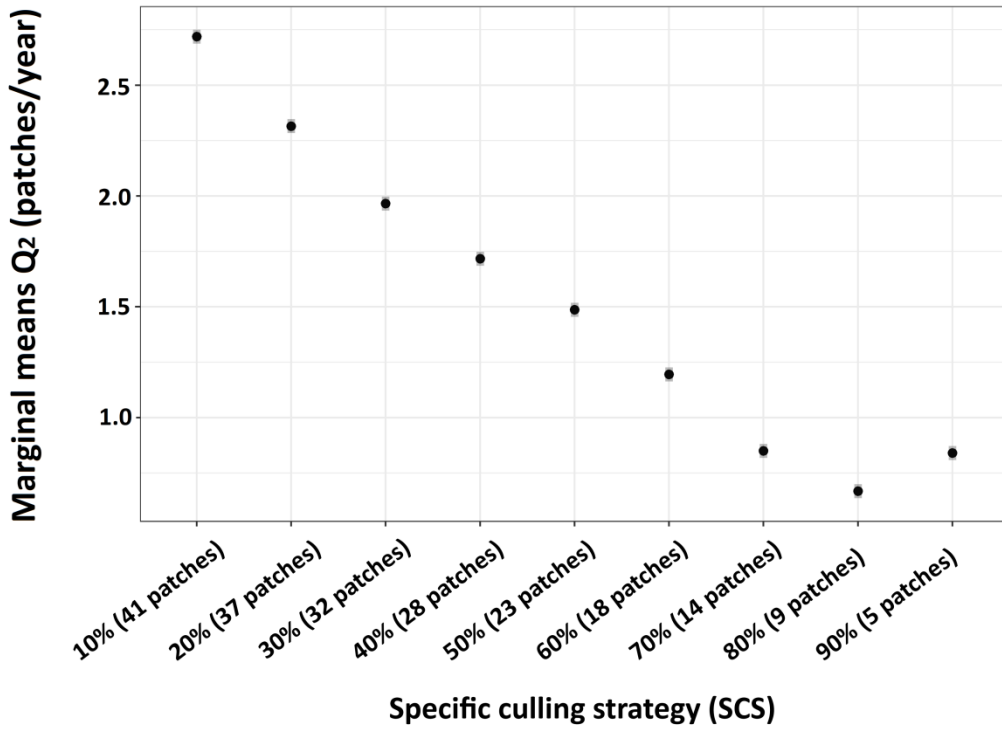
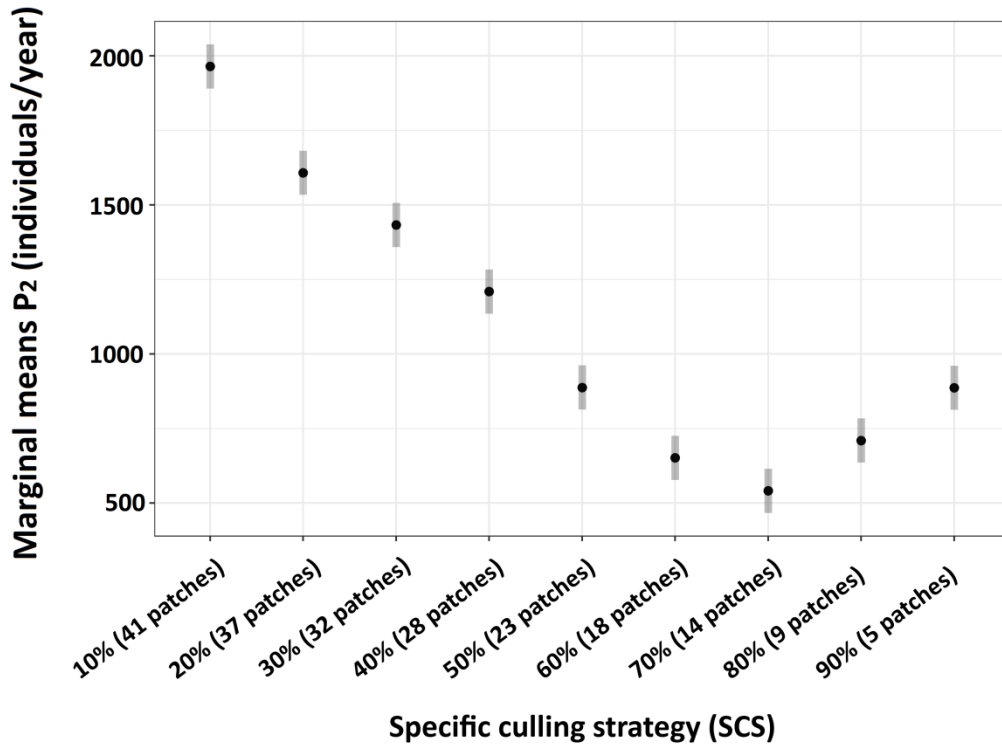


Strongly *r*-selected species

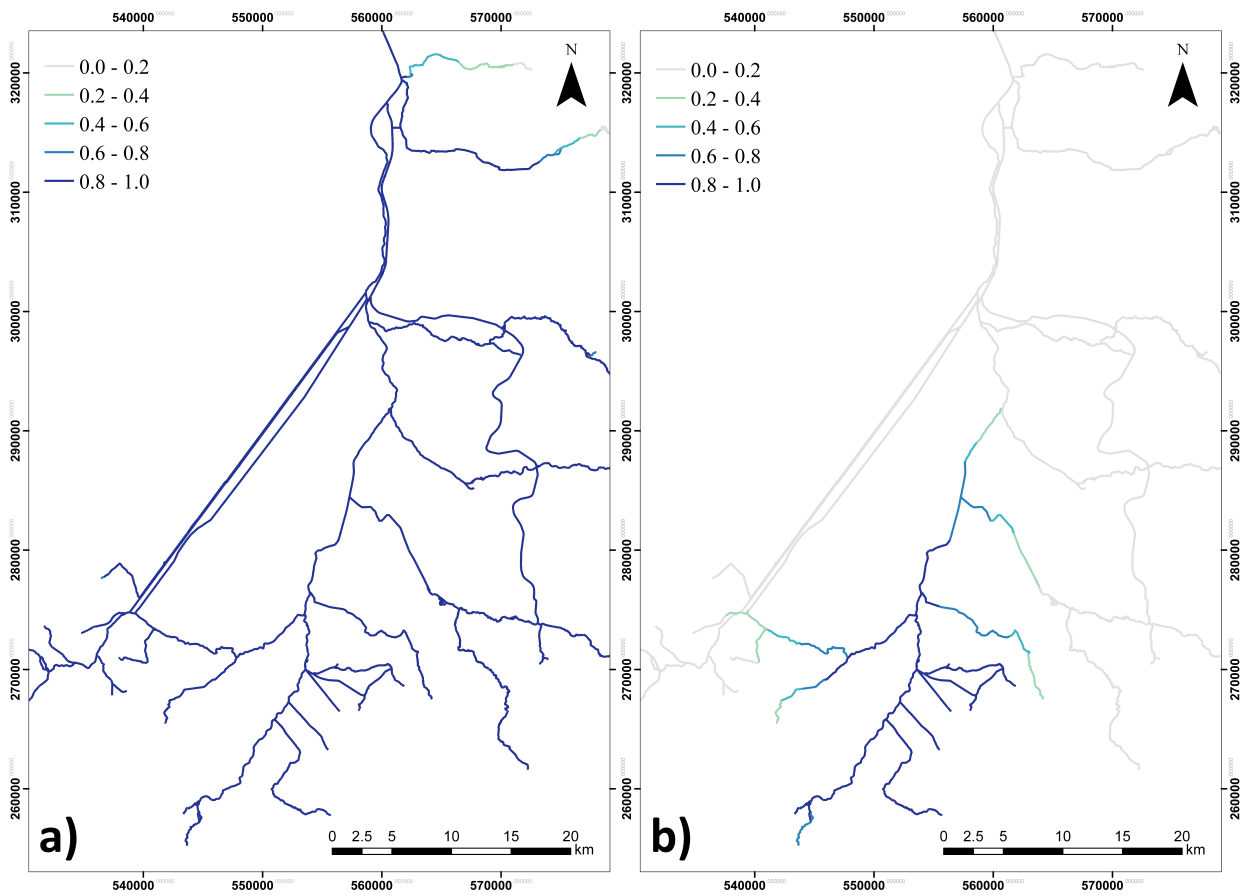




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