

Working title: Highly resolved spatiotemporal simulations for exploring mixed fishery dynamics

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Abstract

To understand how data resolution impacts inference on mixed fisheries interactions we developed a highly resolved spatiotemporal discrete-event simulation model incorporating: i) delay-difference population dynamics, ii) population movement using Gaussian Random Fields to simulate patchy, heterogeneously distributed populations, and iii) fishery dynamics for multiple fleet characteristics based on population targeting under an explore-exploit strategy. This is implemented via a mix of correlated random walk movement (for exploration) and learned behaviour (for exploitation) phases of the fisheries.

Fifty years of sub-daily fishing activity was simulated and used to draw inference on the underlying community structures. We compared inferences based on: commercial catch, a simulated fixed-site sampling survey design and the true underlying populations. We i) establish the potential limitations of fishery-dependent data in providing a robust picture of spatiotemporal distributions; and then ii) simulated an area closure based on areas defined from the different data sources at a range of temporal and spatial resolutions.

Our framework allows users to explore the assumptions in modelling observational data and evaluate the underlying dynamics of such approaches at a fine

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spatial and temporal scale. In application to a mixed fishery exploiting four different populations we found different spatial patterns were evident and that the effectiveness of the spatial closure reduced when data were aggregated across larger spatial areas. However, aggregation across time periods has less of a negative impact on the closure success and while not as effective as when based on the true population, closures based on high catch rates observed in commercial data could still reduce fishing on a protected species.

We conclude from our example framework application that commercial data, while containing bias, provide a useful tool for managing catches in mixed fisheries if applied at the correct spatiotemporal scale.

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1. Introduction

Fishers exploit a variety of fish populations that are heterogeneously distributed in space and time with varying knowledge of species distributions. As fishers do not have full control over what species they select when fishing in ‘mixed fisheries’ it can result in catch of low quota or protected species. If over-quota catch of a species for which they have no quota is discarded without being accounted for it limits our ability to control fishing mortality (Alverson et al., 1994; Crowder et al., 1998; Rijnsdorp et al., 2007) and the ability to manage fisheries for the biological and economic sustainability (Ulrich et al., 2011; Battseer et al., 2015).

There is increasing interest in technical solutions such as gear and spatial closures as measures to reduce unwanted catch (Kennelly and Broadhurst, 2002; Catchpole and Revill, 2008; Bellido et al., 2011; Cosgrove et al., 2019) and adaptive spatial management strategies have been proposed as a way of reduc-

16 ing over-quota discards (Holmes et al., 2011; Little et al., 2014; Dunn et al.,
17 2014). However, if fisheries are to reduce unwanted catch through spatial avoid-
18 ance, an in-depth understanding of spatiotemporal fishery dynamics is required.

19
20 Understanding the correct scale for spatial management measures to be effec-
21 tive is crucial as it enables implementation of effective solutions which minimise
22 economic impact (Dunn et al., 2016). For example, the problem can be to iden-
23 tify a scale that promotes species avoidance for vulnerable or low quota species
24 while allowing continuance of sustainable fisheries for available quota species.
25 Identifying the correct spatial scale remains a challenge because data on fish
26 location at high temporal and spatial resolutions is expensive and difficult to
27 collect and proxies are usually inferred from scientific surveys or commercial
28 catches with limited spatial and temporal resolution. Thus, implementation of
29 spatial measures is hampered by a lack of knowledge of fish and fishery spa-
30 tiotemporal dynamics and understanding of the scale at which these processes
31 become important for management.

32
33
34 Identifying appropriate spatial scales for fisheries closures has been a high-
35 lighted as crucial to their success (Costello et al., 2010; Dunn et al., 2016).
36 Inference on fisheries spatial dynamics is hampered where spatial information
37 is coarse due to low resolution reporting of fisheries catch which is aggregated
38 across larger gridded areas (Branch et al., 2005). Further, if data does not allow
39 identification of spatial features it may lead to poorly sited closures which are
40 ineffectual or have unintended consequences. For example, increased benthic
41 impact on previously unexploited areas from the cod closure in the North Sea
42 were observed without the intended effect of reducing cod exploitation (Rijns-
43 dorp et al., 2001; Dinmore et al., 2003)).

44
45 More refined spatiotemporal information has since become available through
46 the combination of logbook and Vessel Monitoring System (VMS) data (Lee

et al., 2010; Bastardie et al., 2010; Gerritsen et al., 2012; Mateo et al., 2016) and more real-time spatial management has been possible (e.g. Holmes et al., 2011). However, fishers establish favoured fishing grounds through an explore-exploit strategy (Rijnsdorp et al., 2011; Bailey et al., 2018) where they search for areas with high catches and then use experience to return to areas where they’ve experienced high catch in the past. This leads to an inherently biased sampling where target species are over-represented in the catch as fishers exploit areas of high abundance. There is a need to understand the influence of these biases on any spatial management measures which are implemented based on inference from commercial landings or catch data.

To understand the effect of spatiotemporal aggregation of data and fishery targeting on our perception of spatial abundance of different fish populations we ask two fundamental questions regarding inference derived from observational data:

1. Do different source of sampling-derived fisheries data reflects the underlying community structure?
2. How does data aggregation and data source impact on the success of spatial fisheries management measures?

To answer these questions we i) develop a simulation model where population dynamics are highly-resolved in space and time, using a Gaussian spatial process to define suitable habitat for different populations. As the precise locations of the fish are known directly rather than inferred from sampling or commercial catch, we can use the population model to validate how inference from fisheries-dependent and fisheries independent sampling relates to the real community structure in a way we could not with real data. We ii) compare, at different spatial and temporal aggregations, the real (simulated) population distributions to samples from fisheries-dependent and fisheries independent catches to test if these are a true reflection of the relative density of the populations.

76 We then iii) simulate a fishery closure to protect a species based on different
77 spatial and temporal data aggregations.

78

79 We use these evaluations to draw inference on the utility of commercial data
80 in supporting management decisions.

81 2. Materials and Methods

82 A Discrete-event simulation (DES) model of a hypothetical fishery was de-
83 veloped as a software package (*MixFishSim*). The modular approach enabled
84 efficient computation by allowing for sub-modules implemented on time-scales
85 appropriate to capture the characteristic of the different processes (Figure 1).
86 The following sub-modules were included to capture the full system: 1) Popu-
87 lation dynamics, 2) Recruitment dynamics, 3) Population movement, 4) fishery
88 dynamics.

89

90 Population dynamics operate on a daily time-step, while population move-
91 ment occurs on a weekly time-step, with the fishing module operating on a
92 tow-by-tow basis (i.e. multiple events a day).

93 2.1. Population dynamics

94 The basic population level processes were simulated using a modified two-
95 stage Deriso-Schnute delay difference model which models the fish populations in
96 terms of aggregate biomass of recruits and mature components rather than keep-
97 ing track of individuals (Deriso, 1980; Schnute, 1985; Dichmont et al., 2003). A
98 daily time-step was chosen to discretise continuous population processes on a bi-
99 ologically relevant and computationally tractable timescale. Population biomass
100 growth was modelled as a function of previous recruited biomass, intrinsic pop-
101 ulation growth and recruitment functionally linked to the adult population size.
102 Biomass for each cell c was incremented each day d as follows (the full parameter

list is detailed in Table 1):

$$\begin{aligned}
B_{c,d+1} = & \\
& (1 + \rho) B_{c,d} \cdot e^{-Z_{c,d}} - \rho \cdot e^{-Z_{c,d}} \quad \times \\
& (B_{c,d-1} \cdot e^{-Z_{c,d-1}} + Wt_{R-1} \cdot (\alpha_{d-1} \cdot R_{\tilde{y}(c)})) \quad + \\
& Wt_R \cdot (\alpha_d \cdot R_{\tilde{y}(c)})
\end{aligned} \tag{1}$$

where ρ is Brody's coefficient, shown to be equal to e^{-K} when K is the growth rate from a von Bertalanffy logistic growth model (Schnute, 1985). Wt_{R-1} is the average weight of fish prior to recruitment, while Wt_R is the average recruited weight. α_d represents the proportion of fish recruited during that day for the year, while $R_{c,\tilde{y}(c)}$ is the annual recruits in year y for cell c .

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Mortality $Z_{c,d}$ can be decomposed to natural mortality, $M_{c,d}$, and fishing mortality, $F_{c,d}$, where both $M_{c,d}$ and $F_{c,d}$ are instantaneous rates with $M_{c,d}$ fixed and $F_{c,d}$ calculated by solving the Baranov catch equation (Hilborn and Walters, 1992) for $F_{c,d}$:

$$C_{c,d} = \frac{F_{c,d}}{F_{c,d} + M_{c,d}} \cdot \left(1 - e^{-(F_{c,d} + M_{c,d})}\right) \cdot B_{c,d} \tag{2}$$

where $C_{c,d}$ is the summed catch from the fishing model across all fleets and vessels in cell c for the population during the day d , and $B_{c,d}$ the daily biomass for the population in the cell. Here, catch is the sum of those across all fleets and vessels, $C_{c,d} = \sum_{fl=1}^{FL} \sum_{v=1}^V E_{fl,v,c,d} \cdot Q_{fl} \cdot D_{c,d}$ with fl and v the fleet and vessel respectively and $E_{fl,v,c,d}$ and Q_{fl} fishing effort and catchability of the gear, and $D_{c,d}$ is the density of the population at the location fished.

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2.2. Recruitment dynamics

Recruitment is modelled as a function of adult biomass. In *MixFishSim*, it can either take the form of a stochastic Beverton-Holt stock recruitment relationship, or a stochastic Ricker stock recruitment relationship. The Beverton-Holt

relationship is defined as (Beverton and Holt, 1957):

$$\begin{aligned}\bar{R}_{c,d} &= \frac{(\alpha \cdot S_{c,d})}{(\beta + S_{c,d})} \\ R_{c,d} &\sim \log N[(\log(\bar{R}_{c,d}), \sigma^2)]\end{aligned}\tag{3}$$

where α is the maximum recruitment rate, β the spawning stock biomass (SSB) required to produce half the maximum stock size, S current stock size and σ^2 the variability in the recruitment due to stochastic processes. The stochastic Ricker form (Ricker, 1954) is:

$$\begin{aligned}\bar{R}_{c,d} &= B_{c,d} \cdot e^{(\alpha - \beta \cdot B_{c,d})} \\ R_{c,d} &\sim \log N[(\log(\bar{R}_{c,d}), \log(\sigma^2))]\end{aligned}\tag{4}$$

where α is the maximum productivity per spawner and β the density dependent reduction in productivity as the SSB increases.

2.3. Population movement dynamics

Population movement is a combination of directed (advective) movement where at certain times of year the population moves towards spawning grounds by increasing the probabilities of moving into the spawning grounds from adjacent cells, and random (diffusive) movement, governed by a stochastic process where movement between adjacent cells is described by a set of probabilities.

To simulate fish population distribution in space and time a Gaussian spatial process was employed to model habitat suitability for each of the populations on a 2d grid. We first defined a Gaussian random field process, $\{S(c) : c \in \mathbb{R}^2\}$, where for any set of cells c_1, \dots, c_n , the joint distribution of $S = \{S(c_1), \dots, S(c_n)\}$ is multivariate Gaussian with a *Matérn* covariance structure, where the correlation strength weakens with distance. This enables us to model the spatial autocorrelation observed in animal populations where density is more similar in nearby locations (Tobler, 1970; F. Dormann et al., 2007; Poos and Rijnsdorp, 2007b) and we change the parameters to implement different spatial structures for the different populations using the *RandomFields* R package

(Schlater et al., 2015). We define a stationary habitat field with an anisotropic pattern (to simulate a depth gradient) and combine it with a temporally dynamic thermal tolerance field to imitate two key drivers of population dynamics. Each population was initialised at a single location, and subsequently moved across the entire space according to a probabilistic distribution based on habitat suitability (represented by the normalised values from the GRFs), temperature and distance from current cell:

$$Pr(C_{wk+1} = J | C_{wk} = I) = \frac{e^{-\lambda \cdot d_{I,J}} \cdot (Hab_{J,p}^2 \cdot Tol_{J,p,wk})}{\sum_{c=1}^C e^{-\lambda \cdot d} \cdot (Hab_{c,p}^2 \cdot Tol_{c,p,wk})} \quad (5)$$

Where $d_{I,J}$ is the euclidean distance between cell I and cell J , λ is a given rate of decay, $Hab_{c,p}^2$ is the squared index of habitat suitability for cell c and population p , with $Tol_{c,p,wk}$ the temperature tolerance for cell c by population p in week wk (see below).

During pre-defined weeks of the year the habitat suitability is modified with user-defined spawning habitat locations, resulting in each population having concentrated areas where spawning takes place. The populations then move towards these cells in the weeks prior to spawning, resulting in directional movement towards the spawning grounds.

A time-varying temperature covariate changes the interaction between time and suitable habitat on a weekly time-step. Each population p was assigned a thermal tolerance with mean, μ_p and variance, σ_p^2 so that each cell and population temperature suitability is defined that:

$$Tol_{c,p,wk} = \frac{1}{\sqrt{2\pi \cdot \sigma_p^2}} \cdot \exp\left(-\frac{(T_{c,wk} - \mu_p)^2}{2 \cdot \sigma_p^2}\right) \quad (6)$$

Where $Tol_{c,p,wk}$ is the tolerance of population p for cell c in week wk , $T_{c,wk}$ is the temperature in the cell given the week and μ_p and σ_p^2 the mean and standard deviation of the population temperature tolerance.

175 The final combined process results in a population structure and movement
176 pattern unique to each population, with population movement occurring on a
177 weekly basis. The decision to model population movement on a weekly timescale
178 was to reflect that fish tend to aggregate in species specific locations that have
179 been observed to last around one to two weeks (Poos and Rijnsdorp, 2007b).
180 Therefore this process approximated the demographic shifts in fish populations
181 throughout a year with seasonal spawning patterns (Figure S5).

182 *2.4. Fleet dynamics*

183 Fleet dynamics can be broadly categorised into three components: fleet tar-
184 geting - that determined the fleet catch efficiency and preference towards a
185 particular population; trip-level decisions, that determines the initial location
186 to be fished at the beginning of a trip; and within-trip decisions, that determines
187 movement from one fishing spot to another within a trip. An explore-exploit
188 type strategy was implemented in the model that combined these three compo-
189 nents for individual vessels to maximise their catch from an unknown resource
190 distribution (Bailey et al., 2018). The decision to use an individual based model
191 for fishing vessels was taken because fishers are heterogeneous in their location
192 choice behaviour due to different objectives, risk preference and targeting prefer-
193 ence (Van Putten et al., 2012; Boonstra and Hentati-Sundberg, 2015). Therefore
194 fleet dynamics are emergent from individual dynamics rather than pre-defined
195 group dynamics.

196 *2.4.1. Fleet targeting*

197 Each fleet of n vessels was characterised by both a general efficiency, Q_{fl} ,
198 and a population specific efficiency, $Q_{fl,p}$ which are each bound by $[0,1]$. The
199 product of these parameters $[Q_{fl} \cdot Q_{fl,p}]$ affects the overall catch rates for the fleet
200 and the preferential targeting of one species over another. This, in combination
201 with the parameter choice for the step-function defined below (as well as some
202 randomness from the exploratory fishing process) determined the preference of
203 fishing locations for the fleet.

204 *2.4.2. Decision about where to fish at the start of the trip*

205 Several studies (Girardin et al., 2016, for a review) have confirmed past
206 activity and past catch rates are strong predictors of fishing location choice.
207 For this reason, the fleet dynamics sub-model included a learning component,
208 where a vessel’s initial fishing location in a trip was based on selecting from
209 previously successful fishing locations. This was achieved by calculating an
210 expected revenue based on the catches from locations fished in the preceding
211 trip as well as the same month periods in previous years and the travel costs
212 from the port to the fishing grounds. Then a vessel chooses randomly from the
213 top 70 % of fishing events (defined as the ‘threshold’) in terms of expected profit
214 within that season.

215 *2.4.3. Decision about where to fish within the trip*

216 Fishing locations within a trip are initially determined by a modified ran-
217 dom walk process. As the simulation progresses the within-trip decision become
218 gradually more influenced by experience gained from past fishing locations (as
219 per the initial trip-level location choice), moving location choice towards areas
220 of higher perceived profit. A random walk was chosen for the exploratory fishing
221 process as it is the simplest assumption commonly used in ecology to describe
222 optimal animal search strategy for exploiting heterogeneously distributed prey
223 about which there is uncertain knowledge (Viswanathan et al., 1999). In a ran-
224 dom walk, movement is a stochastic process through a series of steps. These
225 steps have a length, and a direction that can either be equal in length or take
226 some other functional form. The direction of the random walk was also cor-
227 related (known as ‘persistence’) providing some overall directional movement
228 (Codling et al., 2008).

229

230 For our implementation of a random walk directional change is based on a
231 negatively correlated circular distribution where a favourable fishing ground is
232 likely to be “fished back over” by the vessel returning in the direction it came
233 from. The step length (i.e. the distance travelled from the current to the next

234 fishing location) is determined by relating recent fishing success, measured as
 235 the summed value of fish caught (revenue, Rev);

$$Rev_{c,d} = \sum_{p=1}^P L_{c,d,p} \cdot Pr_p \quad (7)$$

236 where $L_{c,d,p}$ is landings of a population p , and Pr_p price of a population. All
 237 population prices were kept the same across fleets and seasons. Here, when
 238 fishing is successful vessels remain in a similar location and continue to exploit
 239 the local fishing grounds. When unsuccessful, they move some distance away
 240 from the current fishing location. The movement distance retains some degree
 241 of stochasticity, that can be controlled separately, but is determined by the
 242 relationship:

$$Le = e^{\log(\beta_1) + \log(\beta_2) - \left(\log\left(\frac{\beta_1}{\beta_3}\right)\right) \cdot Rev} \quad (8)$$

243 Where β_1 , β_2 and β_3 are parameters determining the shape of the step function
 244 in its relation to revenue, so that, a step from (x1,y1) to (x2, y2) is defined by:

$$\begin{aligned} (x2, y2) = & x1 + Le \cdot \cos\left(\frac{\pi \cdot Br}{180}\right), \\ & y1 + Le \cdot \sin\left(\frac{\pi \cdot Br}{180}\right) \end{aligned} \quad (9)$$

$$with \quad Br_{t-1} < 180, Br_t = 180+ \sim vm[(0, 360), k]$$

$$Br_{t-1} > 180, Br_t = 180- \sim vm[(0, 360), k]$$

245 where Le is the step length, k the concentration parameter from the von Mises
 246 distribution that we correlate with the revenue so that $k = (Rev + 1/RefRev) \cdot$
 247 max_k , where max_k is the maximum concentration value, k , and $RefRev$ is
 248 parametrised as for β_3 in the step length function.

249 2.4.4. Local population depletion

250 Where several fishing vessels exploit the same fish population competition is
 251 known to play an important role in local distribution of fishing effort (Gillis and
 252 Peterman, 1998). If several vessels are fishing on the same patch of fish, local
 253 depletion and interference competition will affect fishing location choice of the

254 fleet as a whole (Rijnsdorp, 2000; Poos and Rijnsdorp, 2007a). To account for
255 this behaviour, the fishing sub-model operates spatially on a daily time-step so
256 that for future days the biomass available to the fishery is reduced in the areas
257 fished. The cumulative effect is to make heavily fished areas less attractive as a
258 future fishing location choice as reduced catch rates will be experienced.

259 *2.5. Fisheries independent survey*

260 A fisheries-independent survey is simulated where fishing on a regular grid
261 begins each year at the same time for a given number of stations (a fixed station
262 survey design). Catches of the populations at each station are recorded but not
263 removed from the population (catches are assumed to have negligible impact
264 on population dynamics). This provides a fishery independent snapshot of the
265 populations at a regular spatial intervals each year, similar to scientific surveys
266 undertaken by fisheries research agencies.

268 *2.6. Software: R-package development*

269 The simulation framework is implemented in the statistical software package
270 R (R Core Team, 2017) and available as an R package from the author's github
271 site (www.github.com/pdolder/MixFishSim).

273 **3. Parameterisation**

274 We parameterise *MixFishSim* to investigate the influence of data aggregation
275 on spatial inference.

276 *3.1. Population models*

277 We parametrised the simulation model for four example populations with
278 different demographics, growth rates, natural mortality and recruitment pa-
279 rameters (Table 4). Habitat preference (Figure S1) and temperature tolerances

(Figures S3, S4) were defined to be unique to each population resulting in differently weekly distribution patterns (Figures S5-S7). In addition, each of the populations was assumed to have two defined spawning areas that result in the populations moving towards these areas in pre-defined weeks (Figure S2) with population-specific movement rates (Table 4).

3.2. Fleet parametrisation

The fleets were parametrised to reflect five different characteristic fisheries with unique exploitation dynamics (Table 5). By setting different catchability parameters ($Q_{fl,p}$) we create different targeting preferences between the fleets and hence spatial dynamics. The learned random walk process implies that within a fleet different vessels have different spatial distributions based on individual experience. The step function was parametrised dynamically within the simulations as the maximum revenue obtainable was not known beforehand. This was implemented so that vessels take smaller steps when fishing at a location that yields landings value in the top 90th percentile of the value experienced in that year so far (as defined per fleet in Table 5).

Fishing locations were chosen based on random search and, with increasing proportion as time progressed, experience of profitable catches built up in the same month from previous years and from the previous trip. 'Profitable' in this context was defined as the locations where the top 70 % of expected profit would be found given revenue from previous trips and cost of movement to the new fishing location. This probability was based on a logistic sigmoid function with a lower asymptote of 0 and upper asymptote of 0.95, and a growth rate that ensures the upper asymptote (where decisions are mainly based on past knowledge) is reached approximately halfway through the simulation.

3.3. Survey settings

The survey simulation was set up with a fixed gridded station design with 100 stations fished each year, starting on day 92 and ending on day 112 (5

stations per day) with same catchability parameter ($Q_p = 1$) for all populations p . This approximates a real world survey design with limited seasonal and spatial coverage.

3.4. Example research question

To illustrate the capabilities of *MixFishSim*, we investigate the influence of the temporal and spatial resolution of different data sources on the reduction in catches of a population given spatial closures. To do so, we set up a simulation to run for 50 years based on a 100×100 square grid (undetermined units), with five fleets of 20 vessels each and four fish populations. Fishing takes place four times a day per vessel and five days a week, while population movement is every week.

How does sampling-derived fisheries data reflect the underlying population structure?

To answer this question we compare different spatial and temporal aggregations of the true population distributions to:

- a) **fisheries-independent data:** the inferred population density from a fixed-site sampling survey design as commonly used for fisheries monitoring purposes;
- b) **fisheries-dependent data:** the inferred population density from our fleet model that includes fishery-induced sampling dynamics.

We allow the simulation to run unrestricted for 30 years, then implement spatial closed areas for the last 20 years of the simulation based on data (either derived from the commercial catches, fisheries-independent survey or the true population used at different spatial and temporal scales.

The following steps are undertaken to determine closures:

- 338 1. Extract data source (true population, commercial or survey),
- 339 2. Aggregate according to desired spatial and temporal resolution,
- 340 3. Interpolate across entire area at desired resolution using simple bivariate
- 341 interpolation using the *interp* function from the R package *akima* (Akima,
- 342 2006). This is intended to represent a naive spatial model of catch rates,
- 343 without knowledge of the spatial population dynamics.
- 344 4. Close area covering top 5 % of catch rates

345 In total 28 closure scenarios were run that represent combinations of:

- 346 • **data types:** commercial logbook data, survey data and true population,
- 347 • **temporal resolutions:** weekly, monthly and yearly closures,
- 348 • **spatial resolutions:** 1 x 1 grid, 5 x 5 grid, 10 x 10 grid and 20 x 20 grid,
- 349 • **closure basis:** highest 5 % of catch rates for the protected population

350 Survey closures were on an annual basis only, as this was the most temporally
351 resolved survey data available.

352 4. Results

353 4.1. Emergent simulation dynamics

354 Individual habitat preferences and thermal tolerances result in different spa-
355 tial habitat use for each population (Figure 2) and consequently different sea-
356 sonal exploitation patterns (Figure 3).

It can be seen from a single vessels movements during a trip that the ves-
sel exploits three different fishing grounds, each of them multiple times (Figure
4(A)), while across several trips fishing grounds that are further apart are fished
(Figure 4 (B)). These different locations relate to areas where the highest rev-
enue were experienced, as shown by Figure 4 (D), where several vessels tracks

are overlaid on the revenue field, i.e.

$$Rev_c = \sum_{p=1}^P B_{c,p} \cdot Q_{fl,p} \cdot Pr_p$$

Vessels from the same fleet (and therefore targeting preference) may exploit some shared and some different fishing grounds depending on their own personal experience during the exploratory phase of the fishery (Figure 4 (C)). This results from the randomness in the correlated random walk step function, with distance moved during the exploitation phase and the direction stochastically related to the revenue experienced on the fishing ground (Figure 4 (D)).

4.2. How does sampling-derived fisheries data reflect the underlying population structure?

The aggregated catch composition from each of the data sources over a ten-year period (which shows average seasonal patterns) at different spatial resolutions highlights different patterns in perceived community structure depending on the data source and aggregation level (Figure 5). The finer spatial grid for the true population (top left) and commercial data (top middle) show visually similar patterns, though there are large unsampled areas in the commercial data from a lack of fishing activity (particularly in the lower left part of the sampling domain). The survey data at this spatial resolution displays very sparse information about the spatial distributions of the populations. The slightly aggregated data on a 5 x 5 grid shows similar patterns and, while losing some of the spatial detail, there remains good consistency between the true population and the commercial data. Survey data starts to pick out some of the similar patterns as the other data sources, but lacks spatiotemporal coverage. The spatial catch information on a 10 x 10 and 20 x 20 grid lose a significant amount of information about the spatial resolutions for all data sources, and some differences between the survey, commercial and true population data emerge.

Different perceptions of the proportion of each stock in an area are seen when we aggregate the data over a smaller geographical region at different timescales,

with weekly (top), monthly (middle) and yearly (bottom) catch compositions from across an aggregated 20 x 20 area (Figure 6). In the true population, the monthly aggregation captures the major patterns of composition seen in the weekly data with the percentage of different populations in the catch having similar mean and standard deviations. In the weekly data population 1 = 9.36 (3.99), population 2 = 83.2 (5.60), population 3 = 3.57 (1.23), population 4 = 3.91 (1.59); in the monthly data population 1 = 9.23 (3.87), population 2 = 83.3 (5.52), population 3 = 3.62 (1.15), population 4 = 3.86 (1.52). While means were similar some of the variation was lost when aggregated to an annual level; population 1 = 9.90 (0.173), population 2 = 82.2 (0.308), population 3 = 3.82 (0.119), population 4 = 4.03 (0.0502).

The commercial data on a weekly basis shows some of the same patterns as the true population, though the population 1 (in red) is less well represented and some weeks are missing catches from the area. Here, weekly and monthly compositions were nearly identical, with monthly composition of population 1 = 0.0472 (0.0139), population 2 = 94.4 (1.47), population 3 = 3.12 (1.47), population 4 = 2.40 (0.444). Again, yearly values head a similar mean but smaller standard deviation.

The survey data was only available on an annual basis, and showed again a slightly different composition; population 1 = 0.372 (0.00473), population 2 = 87.7 (0.193), population 3 = 0.729 (0.0200), population 4 = 11.2 (0.172).

4.3. How does data aggregation and source impact on spatial fisheries management measures?

We implemented a spatial closure using the different data sources and spatial and temporal aggregations as outlined in the protocol in Section 3.4. We used this to assess the efficacy of a closure in reducing fishing mortality on population 3, given availability of data and its use at different resolutions in order to evaluate the trade-offs in data sources.

414 The trend in fishing mortality for each species show that in most cases the
 415 fishery closure was successful in reducing fishing mortality on the species of in-
 416 terest (population 3; Figure 7), though interestingly the largest reductions in
 417 fishing mortality happened immediately after the closures, following which the
 418 fisheries “adapted” to the closures and fishing mortality increased again some-
 419 what. The exception to the success was the closures implemented based on the
 420 coarsest spatial (20 x 20) and temporal resolution (yearly) that was ineffective
 421 with all data sources. As expected, closures based on the “known” population
 422 distribution were most effective, with differing degrees of success using the com-
 423 mercial data. Fishing mortality rates on the other species changed in different
 424 proportions, depending on whether the displaced fishing effort moved to areas
 425 where the populations were found in greater or lesser density.

426
 427 A regression tree (using the R package REEMtree (Sela and Simonoff, 2012))
 428 highlights that the factor most contributing to differences in fishing mortality
 429 before and after the closure was the population (72 % showing that the closures
 430 were effective for population 3), followed by data resolution (21 %), data type
 431 (7 %) with the least important factor the timescale (< 1 %). In general the finer
 432 the spatial resolution of the data used the greater reduction in fishing mortality
 433 for population 3 after the closures (Figure 8). The notable outliers are the com-
 434 mercial data at the coarsest spatial resolution (20 x 20) at a yearly and weekly
 435 timescale, where closures were nearly as effective as the fine-scale resolution. In
 436 this case the closures were sufficiently large to protect a core area of the habitat
 437 for the population, but this was achieved in a fairly crude manner by closing a
 438 large area - including area where the species was not found (Figure 9) that may
 439 have consequences in terms of restricting the fishery in a much larger area than
 440 necessary.

441

442 5. Discussion

443 Our study presents a new highly resolved fisheries simulation framework,
444 evaluates the importance of data scaling and considers potential bias introduced
445 through data aggregation when using fisheries data to infer spatiotemporal dy-
446 namics of fish populations. Understanding how fishers exploit multiple hetero-
447 geneously distributed fish populations with different catch limits or conservation
448 status requires detailed understanding of the overlap of resources; this is difficult
449 to achieve using conventional modelling approaches due to species targeting in
450 fisheries resulting in preferential sampling (Martínez-Minaya et al., 2018). Of-
451 ten data are aggregated or extrapolated which requires assumptions about the
452 spatial and temporal scale of processes. Our study explores the assumptions
453 behind such aggregation and preferential sampling to identify potential impacts
454 on management advice. With modern management approaches increasingly
455 employing more nuanced spatiotemporal approaches to maximise productivity
456 while taking account of both the biological and human processes operating on
457 different time-frames (Dunn et al., 2016), understanding assumptions behind
458 the data used - increasingly a combination of logbook and positional informa-
459 tion from vessel monitoring systems - is vital to ensure measures are effective.

461 5.1. Simulation dynamics

462 We employ a simulation approach to model each of the population and fish-
463 ery dynamics in a hypothetical ‘mixed fishery’, allowing us to i) evaluate the
464 consequences of different aggregation assumptions on our understanding of the
465 spatiotemporal distribution of the underlying fish populations, and ii) evaluate
466 the effectiveness of a spatial closure given those assumptions.

467
468 Our approach is unique in that it captures fine scale population and fish-
469 ery dynamics and their interaction in a way not usually possible with real data
470 and thus not usually considered in fisheries simulations. While other simulation

frameworks seek to model individual vessel dynamics based on inferred dynamics from VMS and logbook records (Bastardie et al., 2010), or as a system to identify measures to meet particular management goals (Bailey et al., 2018), our framework allows users to explore the assumptions in modelling observational data and evaluate the underlying dynamics of such approaches at a fine spatial and temporal scale. This offers the advantage that larger scale fishery patterns are emergent properties of the system and results can be compared to those obtained under a statistical modelling framework.

Typically, simulation models that treat fish as individuals are focussed on exploring the inter- and intra- specific interactions among fish populations (e.g. OSMOSE; Shin et al. (2004)) in order to understand how they vary over space and time. Our focus was on understanding the strengths and limitations of inference from catch data obtained through commercial fishing activity with fleets exploiting multiple fish populations and realising catch distributions that may differ from the underlying populations, as identified by Gillis et al. (2008). As such, we favoured a minimum realistic model of the fish populations (Plagányi et al., 2014) taking account of environmental but not demographic stochasticity, while incorporating detailed fishing dynamics that take account of different drivers in a mechanistic way.

Demographic stochasticity arises due to individual-level variability in time to reproduction and death. This form of stochasticity is often modelled by drawing random time intervals from a given distribution (Gillespie, 1977). The impact of demographic stochasticity depends on the population size, with the effects expected to decrease with increasing population size (Lande et al., 2010). This contrasts with environmental stochasticity, which affects all population sizes and is present at the population level in our model by variability in recruitment.

We take account of heterogeneity in fleet dynamics due to different preferences and drivers similarly to other approaches (Fulton et al., 2011), but at an

individual vessel rather than fleet level. We do not explicitly define fleets as rational profit maximisers at the outset, but consider there are several stages to development of the fishery; information gathering through search where the resource location is not known, followed by individual learnt behaviour of profitable locations. This provides a realistic model of how fishing patterns are established and maintained to exploit an uncertain resource through an explore-exploit strategy (Mangel and Clark, 1983; Bailey et al., 2018).

5.2. *How does sampling-derived fisheries data reflect the underlying population structure?*

Our results demonstrate the importance of considering data scale and resolution when using observational data to support management measures. We find that understanding of the community composition dynamics will depend on the level of data aggregation and its important to consider the scale of processes; including population movement rates, habitat uniformity and fishing targeting practices if potential biases in data are to be understood and taken into account (Figures 2, 4).

Our simulation shows that, despite biases introduced through the fishing process, the commercially derived data could still inform on the key spatial patterns in the community structures where the fisheries occurred, which was spatially limited due to the “hotspots” of commercially valuable species being fished. Similarly, despite the even spatial coverage the survey was able to capture some of the same spatial patterns as the true population, but missed others due to gaps between survey stations limiting spatial and temporal coverage (Figure 5). This provides a challenge when modelling unsampled areas in inferring species distribution maps, though these limitations may be overcome by understanding the relationship between the species and habitat covariates where these are known at unsampled locations (Robinson et al., 2011).

531 *5.3. How does data aggregation and source impact on spatial fisheries manage-*
532 *ment measures?*

533 From our simulations spatial disaggregation was more important than the
534 temporal disaggregation of the commercial data. This reflects the fact that there
535 was greater spatial heterogeneity over the spatial domain than experienced in
536 individual locations over the course of the year (Figure 2).

537
538 The yearly data assumes the same proportion of each population caught at
539 any time of the year due to the data aggregation. This assumption introduces
540 ‘aggregation bias’ as the data may only be representative of some point (or no
541 point) in time. The monthly data shows some consistency between the real pop-
542 ulation and commercial data for population 2 - 4, though population 1 remains
543 under-represented. On an annual basis, interestingly the commercial data un-
544 der represents the first species (in red) while the survey over represents species
545 1. This is likely due to the biases in commercial sampling, with the fisheries
546 not targeting the areas where population 1 are present and the survey sampling
547 areas where population 1 is more abundant than on average. This indicates that
548 fixed closures, at the right resolution, when based on commercially derived data
549 have the potential to reduced fishing mortality. The likely cost of poor spatial
550 and temporal resolution is associated with reduced effectiveness and potentially
551 closing fishing opportunities for other fisheries.

552
553 Two contrasting real world approaches in this respect were the spatial clo-
554 sures to protect cod in the North Sea. In one example, large scale spatial closures
555 were implemented with little success due to effort displacement to previously
556 unfished areas (Dinmore et al., 2003), while in another small scale targeted
557 spatiotemporal closures were considered to have some effect in reducing cod
558 mortality without having to disrupt other fisheries significantly (Needle and
559 Catarino, 2011). These examples emphasise the importance of considering the
560 right scale and aggregation of data when identifying area closures and the need
561 to consider changing dynamics in the fisheries in response to such closures.

562

563 Our study showed that fishing rates on other populations also changed (both
 564 up and down) as a side-effect of closures to protect one species. This indicates
 565 the importance in considering fishing effort reallocation following spatial clo-
 566 sures, and our simulation allows us to consider the spatiotemporal reasons for
 567 these changes.

568 5.4. Model assumptions and caveats

569 We model the population and fleet dynamic processes to draw inference on
 570 the importance of data scale and aggregation in understanding and managing
 571 mixed fisheries and their impact on multiple fish populations. In doing so, we
 572 have necessarily had to make a number of simplifying assumptions.

573

574 Fish populations in our simulations move in pre-defined timescales and ac-
 575 cording to fixed habitat preferences and temperature gradients (Figures S1, S3).
 576 Our assumptions in parameterising the model (movement rates, temperature
 577 tolerances) will have a direct impact on our conclusions on the relative impor-
 578 tance of spatial and temporal processes. These assumptions could be explored
 579 in a future study by varying the parameters and assessing the robustness of our
 580 conclusions. For our example application we have chosen movement rates to re-
 581 flect aggregation periods observed in past studies (Poos and Rijnsdorp, 2007b).

582

583 In addition, we have assumed that fishing vessels are not restricted by quota
 584 and therefore discarding of species for which vessels have no quota or that are
 585 unwanted is not taken into account. This is likely to be a significant source of
 586 bias in any inference using commercial data and should also be explored. For
 587 example, *MixFishSim* could be altered to allow for spatiotemporal appraisal of
 588 the impact of discarding on fisher behaviour and underlying populations via in-
 589 clusion as discarding behaviour, or through move-on rules or cessation of fishing
 590 activity when quota is exhausted.

591

592 5.5. Future applications of *MixFishSim*

593 We consider that the increased availability of high resolution catch and lo-
594 cational information from commercial fisheries will require it to be a key source
595 of data for ensuring management is implemented at the right scale in future.
596 For example, identifying hot-spots for bycatch reduction or identifying spatial
597 overlaps in mixed fisheries (Dolder et al., 2018; Gardner et al., 2008; Little et al.,
598 2014; Dedman et al., 2015; Ward et al., 2015). Our simulation model has the
599 potential to test some of the assumptions behind the modelling approaches in
600 identifying such hotspots and indeed behind spatiotemporal modelling in gen-
601 eral (e.g. comparing GAMs, GLMMs, Random Forests and geostatistical mod-
602 els under different data generation processes as exemplified by Stock et al. (2019)).

603
604 Other novel applications of our framework could be: testing different sur-
605 vey designs given multiple species and data generating assumptions (Xu et al.,
606 2015); commercial index standardisation methods and approaches and under-
607 standing of appropriate scales and data aggregations and non-proportionality
608 in catch rate and abundance (Harley et al., 2001; Maunder and Punt, 2004);
609 exploring assumptions about the distribution of natural mortality and fishing
610 mortality throughout the year and importance of capturing in-year dynamics
611 in estimating stock status (Liu and Heino, 2013); at sea sampling scheme de-
612 signs to deliver unbiased estimates of population parameters (Cotter and Pilling,
613 2007; Kimura and Somerton, 2006); adaptive management (Walters, 2007; Dunn
614 et al., 2016); testing the ability of commonly employed fleet dynamics models
615 such as Random Utility Models to capture fine scale dynamics and understand
616 their importance (Girardin et al., 2016); and as a detailed operating model in a
617 management strategy evaluation (Mahévas and Pelletier, 2004).

618

619 6. Conclusions

620 *MixFishSim* provides a detailed simulation framework to explore the inter-
621 action of multiple fisheries exploiting different fish populations. The framework
622 enables users to evaluate assumptions in modelling commercially derived data
623 through comparison to the true underlying dynamics at a fine spatial and tem-
624 poral scale. Understanding these dynamics, the limitations of the data and any
625 potential biases that may be introduced when making inference on spatiotempo-
626 ral interactions will enable users to identify weaknesses in modelling approaches
627 and identify where data collection is needed to strengthen inference.

628
629 Our application shows that inference on community dynamics may change
630 depending on the scale of data aggregation. There is an important balance in
631 ensuring that the data are sufficiently spatially and temporally disaggregated
632 that the main features of the data are captured, yet maintaining enough data
633 coverage that the features can be distinguished. We found in our application
634 that there was greater spatial heterogeneity than temporal heterogeneity and
635 that when using aggregated data to define spatial closures coarser temporal reso-
636 lution (months instead of weeks) could still achieve the same results in reducing
637 exploitation rates of a vulnerable species at the highest temporal resolution
638 data. Conversely, reducing the spatial resolution had a negative effect on the
639 effectiveness of the measures (though importantly, there was still some benefit
640 even with coarse spatial resolution).

641
642 While any findings are likely to be case specific, our findings emphasise the
643 need to understand population demographics, habitat use and movement rates
644 in designing any closure scenario based on observational sampling. This infor-
645 mation can then be used to set the bounds on data aggregation used in modelling
646 studies aimed at informing the management measures.

647
648 *MixFishSim* has numerous potential additional applications as it enables

649 the user to apply methods to a fisheries system where there is detailed under-
650 standing of underlying spatiotemporal dynamics. This enables identification of
651 weaknesses or limitations which would not be possible otherwise. In future, we
652 recommend use of the framework to test hypothesis that are otherwise unable
653 to be analysed using real world data due to limitations of data collection. That
654 way the knowledge gained through simulation can inform the future design of
655 management measures.

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660 **Appendices**

Table 1: Description of variables for population and recruitment dynamics sub-modules.

Variable	Meaning	Units
Population dynamics		
<i>Delay-difference model</i>		
$B_{c,d}$	Biomass in cell c and day d	kg
$Z_{c,d}$	Total mortality in cell c for day d	-
$R_{c,\bar{y}}$	Annually recruited fish in cell	yr ⁻¹
ρ	Brody's growth coefficient	yr ⁻¹
Wt_R	Weight of a fully recruited fish	kg
Wt_{R-1}	Weight of a pre-recruit fish	kg
α_d	Proportion of annually recruited fish recruited during day d	-
<i>Baranov catch equation</i>		
$C_{c,d}$	Catch from cell c for day d	kg
$F_{c,d}$	Instantaneous rate of fishing mortality in cell c on day d	-
$M_{c,d}$	Instantaneous rate of natural mortality in cell c on day d	-
$B_{c,d}$	Biomass in cell c on day d	kg
Recruitment dynamics		
$\tilde{R}_{c,d}$	is the recruitment in cell c for day d	d^{-1}
$B_{c,d}$	biomass in cell c for day d	d^{-1}
α	the maximum recruitment rate (Beverton Holt) or maximum productivity per spawner (Ricker)	kg
β	the stock size required to produce half the maximum rate of recruitment (Beverton Holt) or density dependent reduction in productivity per capita of SSB	kg

Table 2: Description of variables for population movement sub-module.

Variable	Meaning	Units
<i>Thermal tolerance</i>		
$T_{c,wk}$	Temperature for cell in week	$^{\circ}\text{C}$
μ_p	Mean of the thermal tolerance for population	$^{\circ}\text{C}$
σ_p^2	Standard deviation of thermal tolerance for the population	$^{\circ}\text{C}$
<i>Population movement model</i>		
λ	Decay rate for population movement	-
$Hab_{c,p}$	Habitat suitability for cell c and population p	-
$Tol_{c,wk,p}$	Thermal tolerance for in cell c at week wk for population p	-
$d_{I,J}$	Euclidean distance between cell I and cell J	-

Table 3: Description of variables for fleet dynamics sub-module.

Variable	Meaning	Units
Rev	Revenue from fishing tow	€
L_p	Landings of population p	kg
Pr_p	Average price of population p	€ kg^{-1}
Le	Step length for vessel	-
Br	Bearing	degrees
k	Concentration parameter for von mises distribution	-
β_1	shape parameter for step function	-
β_2	shape parameter for step function	-
β_3	shape parameter for step function	-

Table 4: Population dynamics and movement parameter settings.

Parameter	Pop 1	Pop 2	Pop 3	Pop 4
Habitat quality				
Matérn ν	1/0.015	1/0.05	1/0.01	1/0.005
Matérn κ	1	2	1	1
Anisotropy	1.5,3,-3,4	1,2,-1,2	2.5,1,-1,2	0.1,2,-1,0.2
Spawning areas (bound box)	40,50,40,50; 80,90,60,70	50,60,30,40; 80,90,90,90	30,34,10,20; 60,70,20,30	50,55,80,85; 30,40,30,40
Spawning multiplier = 10				
Movement $\lambda = 0.1$				
Population dynamics				
Starting Biomass	1e5	2e5	1e5	1e4
Beverton-Holt Recruit α	6	27	18	0.3
Beverton-Holt Recruit β	4	4	11	0.5
Beverton-Holt Recruit σ^2	0.7	0.6	0.7	0.6
Recruit week	13-16	12-16	14-16	16-20
Spawn week	16-18	16-19	16-18	18-20
$K = 0.3$				
$wt = 1$				
$wt_{d-1} = 0.1$				
M (annual)	0.2	0.1	0.2	0.1
Movement dynamics				
μ	12	15	17	14
σ^2	8	9	7	10

Table 5: Fleet dynamics parameter setting.

Parameter	Fleet 1	Fleet 2	Fleet 3	Fleet 4	Fleet 5
Targeting preferences	pop 2/4	pop 1/3	-	pop 4	pop 2/3
Price $Pr_p1 = 100$					
Price $Pr_p2 = 200$					
Price $Pr_p3 = 350$					
Price $Pr_p4 = 600$					
Q_p	0.01	0.02	0.02	0.01	0.01
Q_p	0.02	0.01	0.02	0.01	0.03
Q_p	0.01	0.02	0.02	0.01	0.02
Q_p	0.02	0.01	0.02	0.05	0.01
Exploitation dynamics					
step function β_1	1	2	1	2	3
step function β_2	10	15	8	12	7
step function β_3	Q90	Q90	Q85	Q90	Q80
step function $rate$	20	30	25	35	20
Past Knowledge = TRUE					
Threshold	0.7	0.7	0.7	0.7	0.7
Fuel Cost	3	2	5	2	1

Table 6: Fishing mortality effects of the closure scenarios on population 3 (ordered by most effective first). The fishing mortality rate before the closure was 1.08.

Scenario No	F after closure	% F change	data type	timescale	resolution
9	0.29	-73.47	true Population	weekly	1.00
10	0.29	-72.94	true Population	monthly	1.00
11	0.35	-68.04	true Population	yearly	1.00
45	0.58	-46.70	commercial	yearly	20.00
1	0.58	-46.21	commercial	weekly	1.00
23	0.59	-45.27	true Population	weekly	5.00
2	0.59	-45.06	commercial	monthly	1.00
7	0.60	-44.48	survey	yearly	1.00
24	0.61	-43.20	true Population	monthly	5.00
3	0.64	-40.82	commercial	yearly	1.00
25	0.65	-39.94	true Population	yearly	5.00
17	0.67	-38.11	commercial	yearly	5.00
15	0.71	-34.38	commercial	weekly	5.00
43	0.71	-34.31	commercial	weekly	20.00
16	0.73	-32.58	commercial	monthly	5.00
51	0.78	-27.92	true Population	weekly	20.00
37	0.78	-27.76	true Population	weekly	10.00
39	0.79	-26.98	true Population	yearly	10.00
38	0.81	-25.47	true Population	monthly	10.00
21	0.81	-25.21	survey	yearly	5.00
35	0.81	-25.05	survey	yearly	10.00
44	0.87	-19.91	commercial	monthly	20.00
52	0.88	-18.39	true Population	monthly	20.00
30	0.96	-11.06	commercial	monthly	10.00
29	0.98	-9.80	commercial	weekly	10.00
31	1.03	-4.36	commercial	yearly	10.00

53	1.06	-1.64	true Population	yearly	20.00
49	1.07	-1.01	survey	yearly	20.00

Table 7: Mean and standard deviation of proportions of each species at different levels of temporal aggregation

Data type	Timescale	Population 1	Population 2	Population 3	Population 4
commercial	monthly	0.047(0.014)	94.435(1.47)	3.122(1.468)	2.396(0.444)
commercial	weekly	0.047(0.016)	94.426(1.514)	3.117(1.563)	2.411(0.498)
commercial	yearly	0.051(0.001)	94.388(0.205)	3.021(0.175)	2.539(0.046)
True Population	monthly	9.225(3.872)	83.287(5.522)	3.624(1.151)	3.864(1.519)
True Population	weekly	9.358(3.992)	83.165(5.596)	3.567(1.233)	3.91(1.592)
True Population	yearly	9.899(0.173)	82.25(0.308)	3.821(0.119)	4.031(0.05)
survey	yearly	0.372(0.005)	87.667(0.193)	0.729(0.02)	11.232(0.172)



Figure 1: Schematic overview of the simulation model. Blue boxes indicate fleet dynamics processes, the green boxes population dynamics processes while the white boxes are the time steps at which processes occur; $t = \text{tow}$, t_{max} is the total number of tows; (Rec), (Pop Movement), (Pop Dynamics) logic gates for recruitment periods, population movement and population dynamics for each of the populations, (Past Knowledge) a switch whether to use a random (exploratory) or past knowledge (exploitation) fishing strategy.

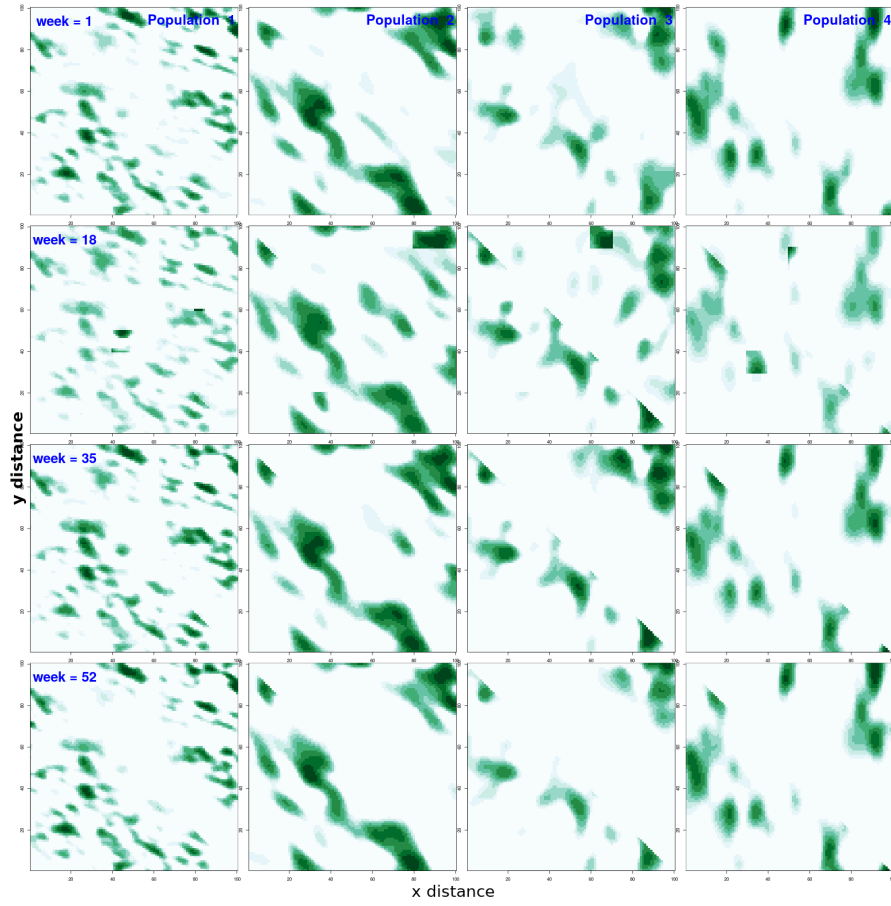


Figure 2: Spatial density (log abundance) for each of the four populations at four time steps. The darker the colour the greater the density of the population. Note that a diagonal anisotropic pattern representing a depth gradient can be clearly seen in populations 2 and 3. The concentrated spawning areas are also visible in the second row of the panels ($t=18$).

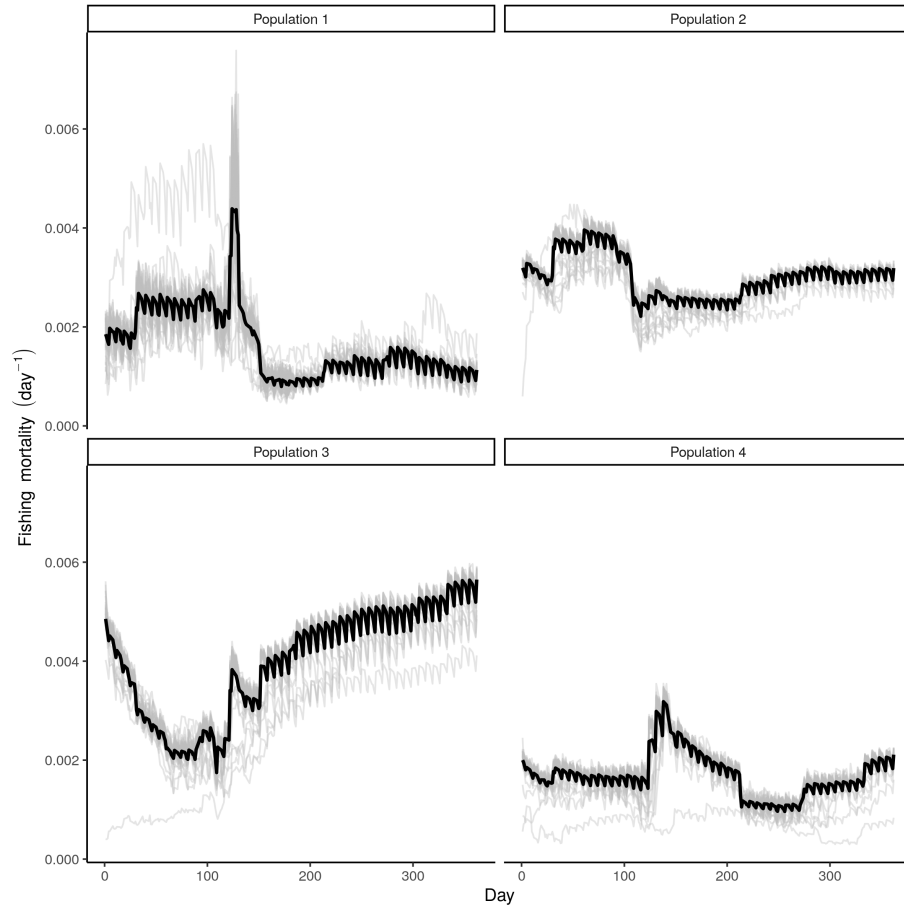


Figure 3: Fishing mortality dynamics - the daily fishing mortalities across the entire spatial domain showing weekly and seasonal patterns in exploitation. Individual years and the light grey lines, the mean of all years the thick black line.

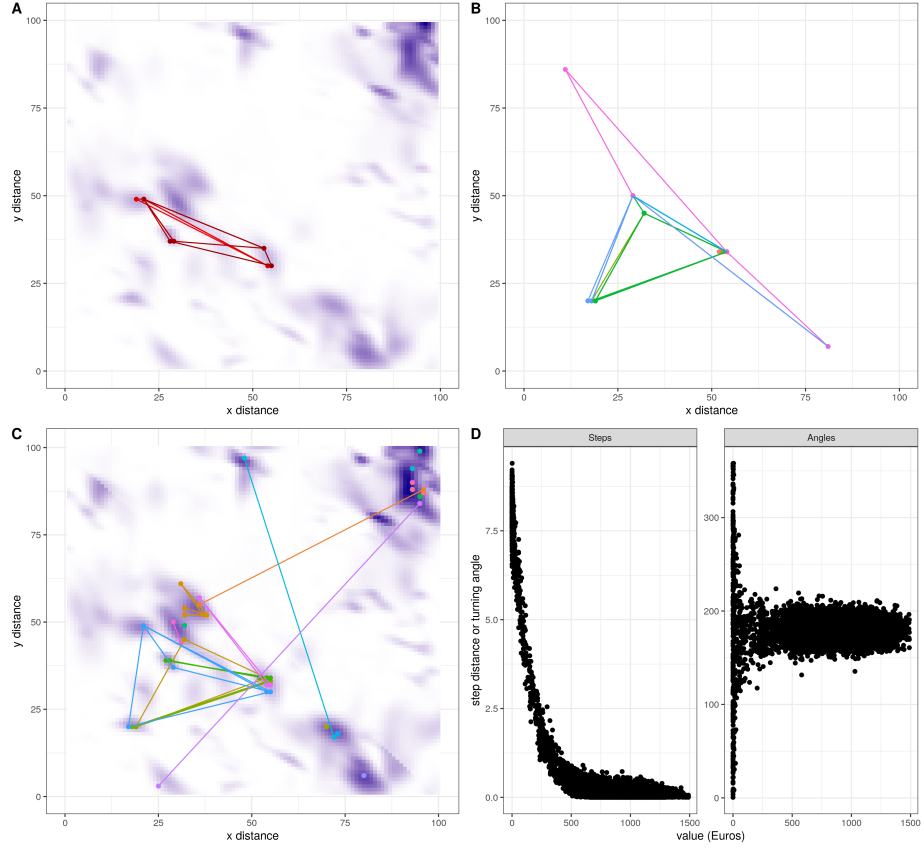


Figure 4: (A) The fishing locations (points) and movements (lines) of a single vessel during a trip overlaid on the revenue of a fishing site (landings \times price); (B) the fishing locations of the vessel over several trips (value field changes over the period so not shown). Note that movements are a mixture of correlated random walk and experience-based, and that the field is wrapped on a torus so that distances between opposite side of the spatial domain are considered next to each other. For these reasons there are moves larger than the step length in D; (C) the locations of multiple vessels from the same fleet overlaid on the value field, (D) the realised step distance and turning angles for a single vessel over the simulation.

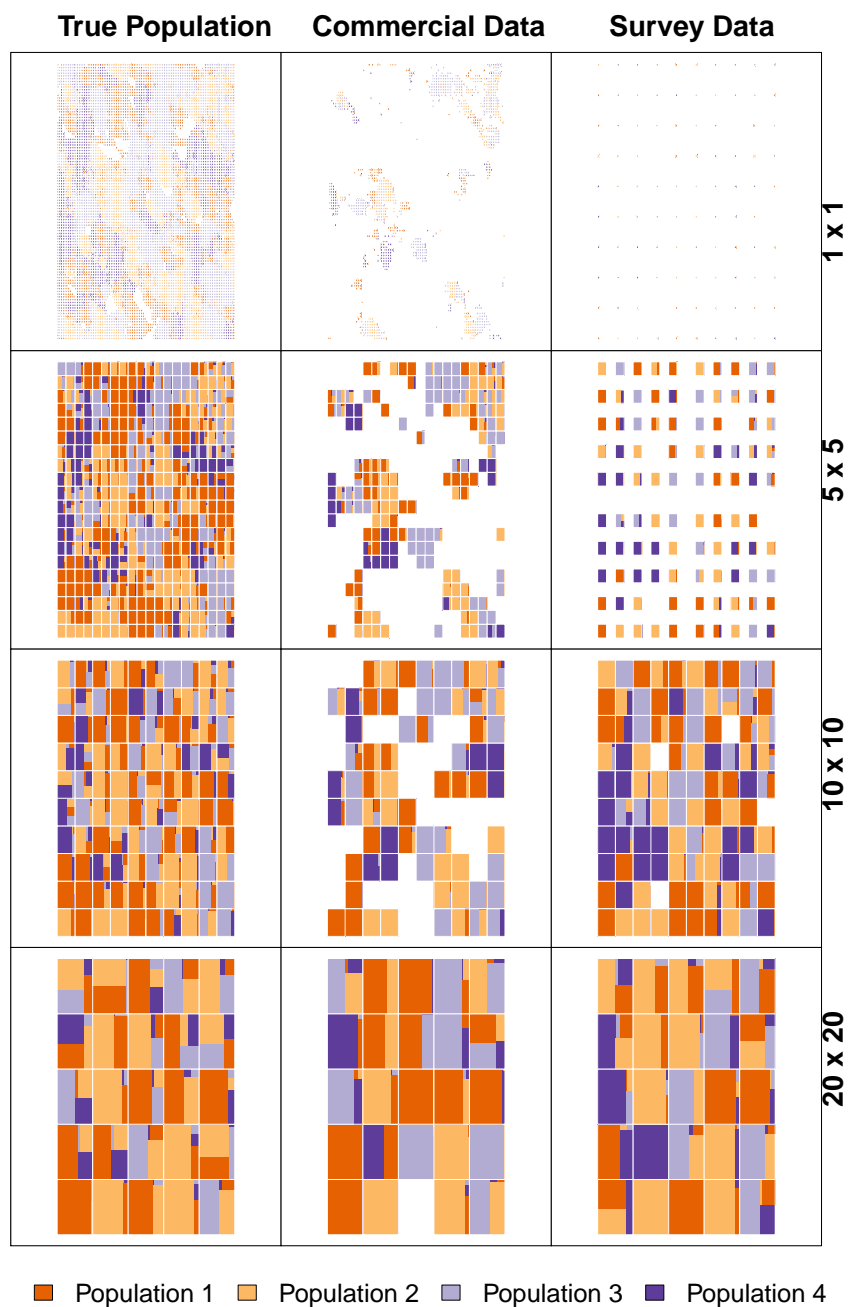


Figure 5: Data aggregation at different spatial resolutions over a ten year period.



Figure 6: Proportion of each population (y axis) for data aggregated at different temporal resolutions. Data is aggregated over a ten-year period for an area 20 x 20. Each bar represents either a week, month or year respectively.

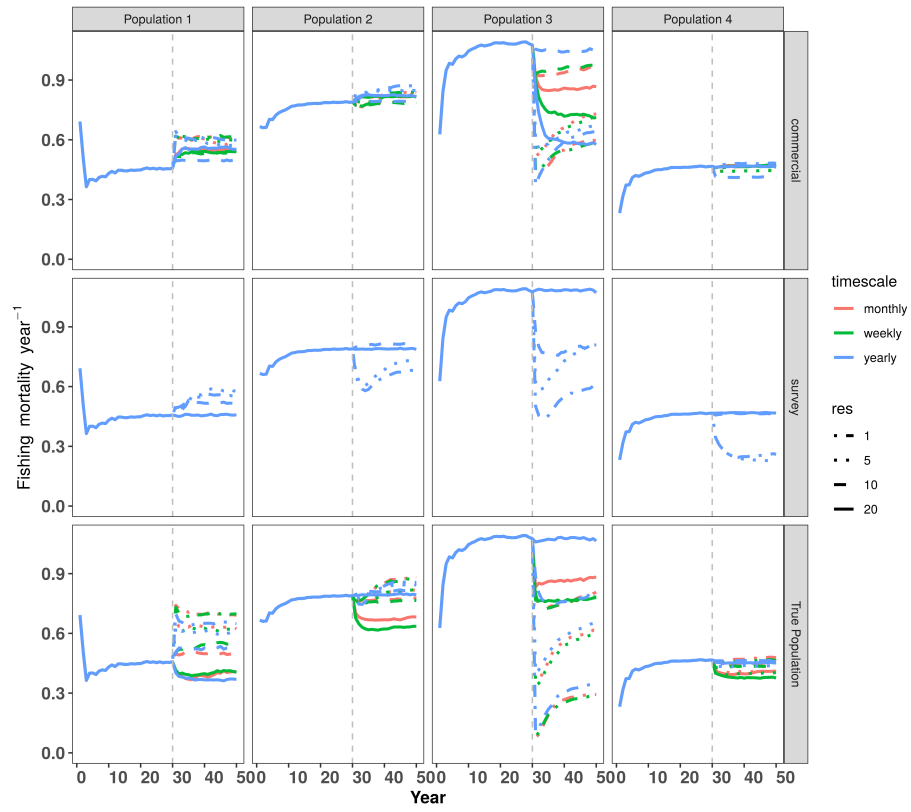


Figure 7: Comparison of closure scenarios effect on fishing mortality trends. Line colour denotes timescale, while linestyle denotes spatial resolution. The vertical dashed line indicates the onset of the spatial closures.



Figure 8: Comparison of closure scenario effectiveness based on different spatial and temporal resolutions.



Figure 9: Closure fishing locations based on annual closures with a coarse spatial resolution. Closure location can be seen in red box in relation to a) before the closure fishing locations, b) after the closure fishing locations, c) population 3 habitat distribution.

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