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▶ To cite this version:

Baptiste Alglave, Youen Vermard, Etienne Rivot, Marie-Pierre Etienne, Mathieu Woillez. Identifying mature fish aggregation areas during spawning season by combining catch declarations and scientific survey data. Canadian Journal of Fisheries and Aquatic Sciences, 2023, 80 (5), pp.cjfas-2022-0110. 10.1139/cjfas-2022-0110. hal-04037065

HAL Id: hal-04037065 https://hal.science/hal-04037065

Submitted on 29 Jun 2023

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Canadian Journal of Fisheries and Aquatic Sciences May 2023, Volume 80, Issue 5, Pages 808-824 <u>https://doi.org/10.1139/cjfas-2022-0110</u> https://archimer.ifremer.fr/doc/00817/92910/

Identifying mature fish aggregation areas during spawning season by combining catch declarations and scientific survey data

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Abstract :

Identifying and protecting essential fish habitats like spawning grounds requires an accurate knowledge of fish spatio-temporal distribution. Commercial declarations coupled with Vessel Monitoring System provide fine scale information on the full year to map fish distribution and identify essential habitats. We developed an integrated framework to infer fish spatial distribution on a monthly time step by combining scientific and commercial data while explicitly considering the preferential sampling of fishermen towards areas of higher biomass. We developed a method to identify areas of persistent aggregation of biomass during the spawning season and interpret these as spawning areas. The model is applied to infer maps of relative biomass for three species (sole, whiting, squids) in the Bay of Biscay on a monthly time step over a 9-year period. Integrating several fleets in inference provides a good coverage of the area and improves model predictions. The preferential sampling parameters give insights into the temporal dynamics of the targeting behavior of the different fleets. Last, persistent aggregation areas reveal consistent with the available literature on spawning grounds, highlighting the potential of our approach to identify reproduction areas.

29 **1** INTRODUCTION

Integrating fisheries into Marine Spatial Planning (MSP) to preserve ecosystem functions and ensure a sustainable exploitation requires an accurate knowledge of fish spatiotemporal distribution and more specifically of fish essential habitats such as reproduction and nursery grounds (Janßen et al. 2018). However, such knowledge is still missing for many species due to a lack of data with sufficient spatial, temporal or demographic resolution (Delage and Le Pape 2016; Regimbart et al. 2018).

36 The available data to map fish distribution and identify essential habitats mainly rely on 37 either scientific survey data (fishery-independent data) or commercial data available 38 through on-board observer programs (fishery-dependent data) (Pennino et al. 2016). Both 39 data sources benefit of direct on-board recording of catches and are usually considered 40 as high quality data. Furthermore, both data sources were proved to be complementary 41 (Rufener et al. 2021). Scientific data benefit from a standardized sampling plan, a 42 standardized catchability and occur each year at the same period. Consequently, they 43 provide standardized data on a large spatial extent for most species and size classes 44 (Hilborn and Walters 2013; Nielsen 2015). Observer data potentially provide data over the 45 full year for all caught species, even though they do not follow a standardized protocol as 46 survey data. However, both scientific survey and onboard observer data are characterized 47 by a relatively low sampling intensity in space and time. Because of material limitations, 48 surveys occur only once or twice a year and provide a limited number of samples each 49 time (ICES 2005) and observer programs only cover a limited fraction of the entire fleet 50 (e.g. only 1% of all sea trips are covered by the French observer programs - Cornou et 51 al., 2021). The low sampling density of both data sources may lead to imprecise predictions (ICES, 2005; Alglave et al., 2022) and constrains to consider only rough temporal resolution (e.g. semesters, quarters or seasons – see for instance Kai et al., 2017; Pinto et al., 2019; Rufener et al., 2021) to ensure a satisfying spatial coverage of the data at each time step. However, the temporality of key biological events, such as the reproduction peak, may be much tighter than the temporal resolution of data (Biggs et al. 2021). Hence, those data alone are likely not sufficient to provide accurate inferences on essential fish habitats such as spawning grounds.

59 Commercial catch declarations combined with their fishing locations available from VMS 60 (Vessel Monitoring System) were proven to be an interesting alternative to obtain landing 61 per unit effort (LPUE) data with fine spatial and temporal resolution (Pedersen et al. 2009; 62 Bastardie et al. 2010; Gerritsen and Lordan 2010; Hintzen et al. 2012; Murray et al. 2013; 63 Azevedo and Silva 2020). However, considering commercial fisheries data to infer fish 64 spatial distribution remains highly challenging. Among other challenges, this implies accounting for fisher sampling behavior. Fisher typically tend to preferentially sample 65 areas of higher biomass (a process referred to as preferential sampling, PS - Diggle et al. 66 67 2010). Hence, because data preferentially represent areas of highest biomass, ignoring 68 PS in the distribution of fishing effort when estimating spatial distribution on larger areas 69 can lead to overestimated biomass predictions (Conn et al. 2017; Pennino et al. 2019; 70 Alglave et al. 2022).

In a recent paper, Alglave et al. (2022) developed an integrated modeling framework to infer spatial distribution of fish abundance by combining scientific survey CPUE and commercial LPUE data while accounting for PS in the distribution of fishing effort. They applied their framework to commercial data of a single month to match with the scientific survey and did not consider any temporal dimension. 76 In this paper, we extend the modeling framework from Alglave et al. (2022) by adding a 77 temporal dimension to estimate fish spatio-temporal distribution at a monthly time step. 78 Our new model accounts for the variation over time (monthly time step) in the biomass 79 field as well as in the intensity of PS for distinct fishing fleets. To demonstrate the value of 80 the method, we selected and applied the model to 3 demersal species in the Bay of Biscay 81 (common sole, whiting and squids) characterized by contrasted configurations regarding 82 the available knowledge of their spawning grounds. We used those applications to reinforce results obtained in Alglave et al. (2022) demonstrating how the integrated 83 framework allows to combine the information from several fleets in order to produce 84 85 accurate maps of spatio-temporal biomass. To illustrate the capacity of the framework to identify areas of aggregation during the spawning season, we processed model outputs 86 87 to identify areas of recurrent aggregation occurring during the reproduction season and 88 compared these to the information available in the literature.

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90 2 MATERIAL AND METHODS

91 In this section, we first present the different species, the datasets and how we process 92 and combine them to produce LPUE data in space and time. Second, we extend the model 93 proposed by Alglave et al. (2022) to introduce a temporal dimension on a discrete monthly time step. In our applications, the models were fitted to data from 2010 to 2018 on a 94 95 monthly time step (108 time steps). Then, we illustrate how the PS component modifies 96 model predictions and can be interpreted, and how integrating several fleets in the analysis further improves model predictions. Last, we detail the method used to 97 98 investigate spatio-temporal dynamics from model outputs and identify reproduction 99 grounds based on the aggregation patterns of each of three species.

100 2.1 Case studies

101 Sole is a data-rich case. Direct information about reproduction grounds is available 102 through egg and larvae surveys (Arbault et al. 1986). Reproduction period fall between 103 January and April, but the peak of the reproduction fall in February. Discard rate is also 104 very low, which makes the landings data a good proxy of the catch (ICES 2019a).

105 By contrast, Whiting is a data-poor case study where only indirect information of 106 reproduction period exists through spring trawl surveys (Houise and Forest 1993).

107 Reproduction period fall between March and May. Discard rates can be high (about 30%)

and thus landing data may provide a biased picture of the real catches (ICES 2019b).

109 Our third case study is Squids that represent a mixture of several species declared under

110 a common denomination in the catch (Loliginidae here referred as squids): *Loligo Vulgaris*

111 (Lamarck, 1798), *Loligo forbesii* (Steenstrup, 1856) and *Alloteuthis sp* (Lamarck, 1798).

112 Overall scientific survey suggests that the predominant species in the Bay of Biscay is

Loligo Vulgaris (ICES, 2020a, p.17). All 3 species are data-poor: no information exists regarding their reproduction grounds but some information of the reproduction period exists for *Loligo Vulgaris* (Moreno et al. 2002). For this species, the reproduction period

- 116 fall between January and April.
- 117 2.2 Data

118 2.2.1 High spatial resolution catch per unit effort data for the mature component119 of the populations

We pre-processed the VMS and catch declaration (logbook) data to obtain high spatial resolution LPUE data for the mature component of those three stocks, for three different fishing fleet, and for each month of the 2010-2018 time series. In the text, we used the term Landing Per Unit of Effort (LPUE) to refer to commercial observations expressed in kg of (mature) biomass per hour fished. Discards are neglected in our approach, hence LPUE are considered as biomass indices.

126 Our model can integrate data of different fishing fleets. For the purpose of our application, 127 we selected three different metiers that belong to the same fleet of trawlers (in this case, 128 the 'métier' term refers to a combination of gear and of a set of species that are targeted 129 by the vessels – see the Data Collection Framework and EC (2008)): OTB DEF (bottom 130 otter trawl targeting demersal species), OTB CEP (bottom otter trawl targeting cephalopods) and OTT DEF (multi-rig otter trawl targeting demersal species). In the 131 132 following those three métiers are referred as three fleets. These three fleets were selected 133 as they offer three main advantages. First, their targeting behaviors and technical 134 characteristics are similar. Second, catch per unit effort of trawlers are generally good 135 indicator of fish relative abundance while other gears (longline, gillnet) may face saturation 136 effects leading to non-linear relationship between catches and fishing time (Hovgêrd and Lassen 2008). Third, the combination of the three fleets cover the full spatial domain(Figure 1).

139 Because one of our primary goal is to identify spawning grounds, we filtered only the 140 mature fraction of the landings (i.e. the fraction of the individuals that can potentially 141 reproduce, not per se the fraction of the population that are spawning – this is detailed further in the discussion). This was done by crossing the landings data with length class 142 143 and maturity data. For most of the landings, information on the commercial size categories 144 is available from the sales notes. These commercial categories are regularly sampled to 145 derive length structure of each commercial category. This allows us to estimate the 146 proportion of potentially mature fish in each commercial category by applying maturity 147 ogives and in turn estimate the proportion of mature fish for each landing declaration. See 148 SM1 for more detail. Note that this procedure was not possible for squids, as there are no data on maturity and size classes for this species group. 149

Landing data were then combined with VMS data to finally obtained high spatial resolution
LPUE data discretized on a 0.05° x 0.05° grid (i.e. 5.5 km x 3 km) on a monthly time step
(see SM2). This combination requires:

(1) to identify the fishing locations within the VMS data. This is realized individually
 for each fishing vessel trajectory based on a speed threshold similarly as in
 common data processing methods (Hintzen et al. 2012).

(2) to reallocate the logbooks declaration on the related VMS fishing locations. This
 reallocation is realized individually for each fishing vessel trajectory by uniformly
 reallocating the landings on all fishing locations. The link between both data
 sources is realized through the combination 'vessel identifier x statistical

rectangle x fishing trip x day'. LPUE are then computed by simply dividing the
 reallocated landings by the related fishing time.

162 2.2.2 Scientific data

We also integrated scientific data in the analysis. For whiting and squids we used the survey data from the EVHOE survey. The Orhago survey was used for sole (ICES, 2020 - see SM3, Figure S3). The data were extracted from the DATRAS database on the period 2010 - 2018. Only the mature fraction of the survey catches were kept in the analysis to make it comparable with commercial data.

Orhago is an annual beam trawl survey occurring in November and designed to assess sole stock status in the Bay of Biscay. Each year 50 stations are sampled within 4 strata all along the Bay of Biscay. Note that this survey is mainly coastal and does not sample offshore areas. EVHOE is an annual bottom trawl survey occurring in late October, November and early December with a stratified sampling plan. It is designed for demersal fishes in the Bay of Biscay and in the Celtic Sea. In the Bay of Biscay, 80 to 90 sampling hauls are recorded each year.

175 2.3 Spatio-temporal integrated model

Alglave et al. (2022) developed a hierarchical integrated statistical model to infer spatial
distribution of fish density through scientific survey data and commercial data. It is
structured in 4 layers:

the latent field that represents biomass spatial distribution, and that is the
main target of the inferences;

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the observations from scientific surveys and commercial declarations that
 are considered as direct zero-inflated observations of the latent field at the
 registered fishing locations;

the fishing sampling intensity that relates fishing locations to the latent field
 and model explicitly the PS of commercial fleets towards areas of higher
 biomass;

187 - unknown parameters that control the shape of the biomass latent field and
188 the sampling process.

189 This first model was purely spatial as no temporal dimension was included in the model.

190 In this paper, we extend the framework by incorporating a temporal component to model

191 the evolution of the latent field of biomass across the monthly time steps (Figure 2).

192 2.3.1 Biomass field

As a notable extension of Alglave et al. (2022), the biomass field (eq.1) is modeled as a
spatio-temporal Gaussian Random Field (GRF) through a log link as:

195
$$\log(S(x,t)) = \alpha_S(t) + \delta(x,t) \quad (1)$$

196 where $x \in D \subset R^2$ stands for the spatial locations and $t \in [1,T]$ for the monthly time steps. 197 S(x,t) is in the same unit as the data (here kg/hr fished as data are CPUE for survey trawls 198 or LPUE for commercial landing declarations). The term $\alpha_{S}(t)$ is a time varying intercept 199 modeled as a fixed effect and $\delta(x,t)$ is a GRF spatio-temporal process which represents 200 the spatio-temporal correlation structure of the biomass field. As commercial data may not 201 always cover all time steps, the temporal correlation is a critical component that allows to 202 interpolate between time-steps. Here, the spatio-temporal term has a classical stationary 203 first-order autoregressive form (eq.2) following (Cameletti et al. 2013):

204
$$\delta(x,t) = \varphi \cdot \delta(x,t-1) + \omega(x,t) \quad \text{for } t = 2,...T \quad (2)$$

The autocorrelation coefficient φ is a scalar with $\varphi \in -1,1$, $\omega(x,t)$ are spatial random effects that represents the spatial innovation, modeled as a 0 mean GRF (with no temporal correlation). Spatial random effects $\omega(x,t)$ are parameterized through a range parameter ρ that corresponds to the distance at which spatial autocorrelation falls below 0.1.

Note that no covariate is included in the latent field to keep the model as simple as possible. If any, the covariates effects are captured through the spatio-temporal term $\delta(x,t)$. Similarly, the intercept $\alpha_S(t)$ was modeled through a simple fixed effect but more complex specifications including some seasonal, yearly and interaction effects could be adopted such as in Thorson et al. (2020).

214 **2.3.2** Sampling process for the commercial fishing points

215 For most scientific surveys targeting a variety of species or covering wide areas, even if 216 the sampling plan is a stratified-random sampling and sampling takes into account the 217 most important commercial species in the area, the final distribution of the sampling 218 location can be considered independent from the biomass field distribution (see S3 for the 219 scientific survey used in this paper), and scientific sampling locations do not need to be 220 modeled explicitly (Diggle et al. 2010). By contrasts, the dependence between the 221 commercial fishing locations and the biomass field has to be modeled and included in the 222 likelihood function to capture preferential sampling. We extended the model proposed by 223 Alglave et al. (2022) to account for temporal variations in PS. Because the fishing behavior is potentially different among fishing fleet, PS is modeled specifically for each fleet 224 225 denoted *j* (in the next section, *j* is also used to denote the scientific suvey). Observed 226 fishing locations of any fishing fleet *i* are integrated in the likelihood through an Page 11 of 47

inhomogeneous point process (X_{comj} in the Figure 2) whose intensity $\lambda_j(x,t)$ (eq.3) controls

the expected number of fishing points within a given area:

229
$$\log(\lambda_j(x,t)) = \alpha_{Xj}(t) + b_j(t) \cdot \log(S(x,t)) + \eta_j(x,t)$$
(3)

230 where:

- the time varying intercept $\alpha_{Xj}(t)$ quantifies the average fishing intensity on the whole area; it is modeled as a fixed effect;

- the time varying $b_j(t)$ quantifies the strength of PS towards biomass S(x,t); it is modeled as a fixed effect too. If $b_j(t) = 0$, then PS is null. If $b_j(t) > 0$, then PS occurs and the greater, the stronger PS. Alternatively, $b_j(t) < 0$ would indicate that fisher have a repulsive behavior towards the resource.

- the pure spatial GRF $\eta_j(x,t)$ captures the remaining spatial variability in the fishing point pattern not captured by the PS term (for instance, dependence of the fishing locations towards management regulations, distribution of other targeted species, habits/tradition).

241 2.3.3 Observation process

All observations for both scientific and commercial data of any fleet *j* are assumed all mutually independent conditionally on the latent field of biomass and the sampling locations. As data (both scientific and commercial) eventually present a high proportion of zero values, we model the observations through a Poisson-link zero-inflated model introduced by Thorson (2018) and already used in Alglave et al. (2022). The observation model explicitly considers that each fleet can have its own catchability and its own zero inflation parameter. The probability to observe a catch data y_i conditionally on the location x_i (with *i* the observation index), the time-step t_i , the biomass field value $S(x_i, t_i)$ and the fleet *j* is expressed as follow:

252
$$P(Y_i = y_i | x_i, t_i, S(x_i, t_i), j) = \begin{cases} p_i & \text{if } y_i = 0\\ (1 - p_i) \cdot L\left(y_i, \frac{\mu_j(x_i, t_i)}{(1 - p_i)}, \sigma_j^2\right) & \text{if } y_i > 0 \end{cases}$$
(4)

253
$$p_i = \exp\left(-e^{\xi_j} \cdot \mu_j(x_i, t_i)\right)$$
(5)

 $\mu_j(x_i, t_i) = q_j S(x_i, t_i)$ is the expected catch of fleet *j* at location x_i and time step t_i . It is the product of the latent field value $S(x_i, t_i)$ and of the relative catchability coefficient specific for fleet *j* denoted q_j . Specifically for each fleet *j*, ξ_j is a zero-inflation parameter controlling the proportion of zero in the data, σ_j^2 is the observation variance when the catch is positive. Equation (4) shows the two components that compose the probability to observe a catch y_i :

• the probability to obtain a zero catch $(y_i = 0)$. It is modeled as a Bernoulli variable with probability $p_i = \exp(-e^{\xi_j} \cdot \mu_j(x_i,t_i))$. p_i is equivalent to the probability to obtain a 0 value with a Poisson distribution of intensity $e^{\xi_j} \cdot \mu_j(x_i,t_i)$. The value of ξ_j controls the intensity of the zero inflation, when ξ_j increases, the amount of zero in the data decreases. Then the probability to obtain a positive catch is given by $1 - p_i$. the value of the positive catch is modeled through a lognormal distribution L with

expected value
$$\frac{\mu_j(x_i,t_i)}{(1-p_i)}$$
 and observation error σ_j^2 . The standardization by $(1-p_i)$
allows to keep the expectancy of the observation model to $\mu_j(x)$.

The catchabilities q_j are not identifiable per se and some additional constraints need to be set to estimate the relative catchability of each fleet (Alglave et al., (2022)). To ensure identifiability, one fleet catchability is set as reference level (e.g. $q_{ref} = 1$, here OTB_DEF was used as the reference fleet) and the other fleets' catchabilities are estimated relatively to the reference fleet through the equation:

 $q_j = k_j.q_{ref} \qquad (6)$

274 2.3.4 Maximum likelihood estimation

The estimation of the spatio-temporal model is achieved through maximum likelihood 275 276 estimation. We used the Stochastic Partial Differential Equation (SPDE) approach that 277 allows to benefit from the nice computational properties of Gaussian Markov Random 278 Fields while working on a continuous domain (Lindgren et al. 2011). In practice our model 279 is coded with Template Model Builder (TMB - Kristensen et al., 2016) which benefits from 280 the Laplace approximation to integrate over random effects, automatic differentiation and 281 sparse matrix computation technics for a fast estimation of the model through maximum 282 likelihood estimation. Details on estimation are provided in SM 4, 5 and 6.

283 **2.4** Evaluating the interest of integrating multiple fleets

Integrating several fleets in inference allows to cover the whole area (Figure 1) and is expected to improve inferences. To illustrate the value of integrating the data from multiple fleets within a single integrated model, we compared the spatial predictions obtained by fitting the model to all available data with those obtained by integrating only one fleet. In addition, we investigated if integrating all the fleets in inference increased the correlation between scientific data and model predictions. We also compared the coefficient of variation of the prediction between each model on a single time step (here November 2018).

Note that scientific data was systematically integrated into inference (either in the integrated model or in the single-fleet models). However, due to the low sample size compared with intensive 'VMS x logbooks' data (about 80 scientific samples each year in November compared with 17000 samples per month on average), they have very low contribution to inference. This was extensively discussed in Alglave et al. (2022). Here they mainly provide some standardized and reference data to assess the performance of the framework.

299 2.5 Evaluating the value of modeling PS

300 **2.5.1 Comparing the inferences with and without PS**

We first assessed the impact of PS on the distribution of biomass by comparing estimations obtained from integrated models (i.e. models fitted to all data sources) accounting for PS with those obtained when ignoring PS. We computed the log-likelihood related to each data source (commercial and scientific data) to assess if there is an improvement in model goodness-of-fit when accounting or not for PS. Note that fitting a model without PS is straightforward as it only requires to remove the sampling process component from the likelihood function.

308 2.5.2 Interpreting the intensity of preferential sampling

The estimates of PS parameters $b_j(t)$ in eq. (3) may bring valuable information on the dynamics of the fishery as they inform on the strength of the relationship between commercial sampling distribution and species distribution. We investigate the variability of the PS parameters among the three species and the different fleets. Then, focusing on the sole case study, we highlight the insights brought by the model on the temporalevolution of PS and its seasonal variations.

315 2.6 Investigating spatio-temporal distribution and identifying reproduction 316 grounds

The spatio-temporal model provides some insight on the temporal dynamics of species distribution both at inter- and intra-annual levels. We applied a method to identify recurrent aggregation areas from the maps of abundance inferred at each time step.

320 2.6.1 Aggregation index

We used the Getis and Ord index $G_d(x,t)$ (Getis and Ord 1992; Ord and Getis 1995) to 321 322 determine persistent aggregation areas (see for instance Milisenda et al., (2021)). The 323 generalized version of the Getis and Ord index is given in Bivand and Wong (2018) and 324 Ord and Getis (1995). Basically, $G_d(x,t)$ is a normalized version of the ratio between the 325 sum of the log-biomass (denoted s(x,t)) within a fixed neighborhood d and the sum of s 326 (x,t) on the entire area (for a fixed time step) (Getis and Ord 1992). We computed these indices on $s(x,t) = \log(S(x,t))$ so that the s(x,t) are Gaussian, which makes G_d Gaussian 327 328 too. In the application, we used a neighborhood distance d=7.5 km which defines a small 329 neighborhood of 8 cells (the direct neighbors of each cell grid) and allows to identify very localized aggregation areas. Positive values for the aggregation index G_d indicates that s 330 (x,t) fall within a local patch of high values while negative G_d indicates that s(x,t) fall within 331 332 a local patch of low values. Near 0 values G_d , indicates that s(x,t) does not fall in some 333 local aggregation patch. As G_d follows a standardized Gaussian distribution, the 334 comparison between the value of the index and the quantiles of a standard Gaussian 335 distribution can be used to evaluate whether or not the latent field of biomass fall within a 336 statistically significant high or low aggregation patch. We used the quantile 99% (2.58) as a threshold to ensure a high level of significance for patch detection (only local patch of
 positive values are considered) and applied the Bonferroni correction to account for the
 multiple statistical tests that are conducted.

Then, we define the persistence indices IP(x,m) as the proportion of times point x falls significantly within an aggregation area for a specific month/season m (can be either a month or several months) among several years. Areas marked with high values of IP are persistent aggregation areas throughout the time series.

344 **2.6.2** Comparing the results with the available literature

Persistent aggregation areas derived from our model during the reproduction period (as
defined from the literature) were interpreted as potential recurrent reproduction grounds.
We compare those inferences with the information of reproduction ground available from
the literature (for sole and whiting).

Arbault et al. (1986) investigated the reproduction of sole along the Bay of Biscay based on several egg surveys occurring in 1982. Five surveys were conducted between January and May. Egg density was sampled in different locations from Hendaye to Pointe du Raz (43°30N-48°N) and allowed to map the distribution of egg production on the full study domain. The peak of reproduction occurred in February; thus we compare the maps obtained from the February survey with the persistence index obtained from our model in February.

For whiting, only two EVHOE trawl surveys occurred during spring (considered as the reproduction period of whiting) between 1987 and 1992 in the Bay of Biscay (Houise and Forest 1993). For each haul, the individuals were counted and aged. Individual of two years and older were considered mature. We compare the distribution of mature

individuals obtained with these surveys and the index of persistence from our modelduring spring (March to May).

No available information exists regarding the reproduction grounds of squids in this area, however the study from Moreno et al. (2002) investigated the reproduction period for *Loligo vulgaris* in the Eastern Atlantic and highlighted that their reproduction falls in winter and spring with a peak from January to April. We compute the persistence index for this period to identify the spatial aggregation patterns that emerge from the model outputs and that could be considered as spawning grounds.

368 To assess whether the aggregation patterns within the reproduction period are stable over

the time period, we iteratively computed the persistence index over a 5-year mobile time-

370 span while pushing forward one year each time.

371 **3 Results**

372 **3.1** Assessing the contribution of each data sources to inference

Results highlight how combining several commercial fleets in the framework brings a better picture of the spatial distribution on the whole domain. For instance, when comparing model predictions with survey data (for the month of the survey), integrating several fleets into the analysis improves correlation with scientific data (Figure 3). It also reduces the standard deviation of the predictions on the full domain (SM 7, Figure S7).

378 When looking at the predictions within the spatial range of the fleets, single-fleet models 379 logically provide similar spatial predictions compared with the integrated model (Figure 4; 380 red dots). However, when using single fleet data, predictions realized outside the spatial 381 range of the fleet largely depart from the ones realized through the integrated models 382 (black dots, Figure 4), emphasizing the contribution of the other fleets to improve 383 inferences on areas poorly covered by single fleet. This is particularly evidenced with the 384 OTB CEP and OTT DEF fleets that partially cover the study area compared with 385 OTB DEF that better cover the whole study area (Figure 1).

386 3.2 Interpreting estimates PS intensity

Estimates of the PS intensity (b parameters in eq. (3)) for the different species, the 387 different fleets and the different time steps provide information on the targeting behavior 388 389 that are consistent with expertise. Estimates of *b* are positive for each species and each 390 fleet (Figure 5, left column). For squids, PS is the strongest for OTB CEP followed by 391 OTB DEF and OTT DEF. This is consistent with the expert knowledge of the targeting behavior of these fleets: OTB CEP target cephalopods and catch on average 15% of 392 393 squids while OTB DEF and OTT DEF catch respectively 5% and 1% of squid). A similar pattern can be identified for whiting $(b_{OTB_{CFP}} > b_{OTB} > b_{OTT})$; this is consistent with species 394

spatial distribution as whiting (like squids) are found in coastal areas where the OTB_CEP fleet is preferentially operating (Figure 1). For sole, the strength of PS is on average higher for OTB_CEP and OTT_DEF than for OTB_DEF but with less contrast between the three commercial fleets which is also consistent with expertise as those three fleet target this high commercial value species.

Interestingly, some of the *b* parameters time series emphasize seasonal patterns (Figure 6, top). For instance, in the sole case study for the OTB_CEP fleet, the *b* parameters are higher in summer and autumn emphasizing relatively stronger PS, while being lower in winter and early spring (but see section 3.4 below for a more detailed interpretation of this seasonality pattern).

405 **3.3** Evaluating the influence of PS on spatial inferences

406 Because estimates of b are positive, spatial density of fishing points is positively correlated 407 with biomass density. Then logically, ignoring PS leads to a positive bias (overestimate) 408 in biomass estimates in areas not sampled by the commercial fleets compared to 409 estimates obtained while considering (Figure 5, right column, black points), but does not 410 strongly affect predictions in locations within the range of the fleets (blue points). 411 Considering PS only slightly improves the fit of the model to the data. For the Sole case 412 study, some improvement of the likelihood occurred for both the likelihood associated with 413 the commercial and the scientific data (Table 1). For whiting and squids, there are no 414 strong modifications in both scientific and commercial likelihoods. Overall, accounting or 415 not for PS does not strongly modify the overall pattern of species abundance (SM 8).

416 **3.4** Investigating spatio-temporal dynamics of fish biomass

Results provide biomass density maps on a monthly time step that emphasize seasonaldistribution patterns and from which aggregation index were calculated. The temporal

419 correlation parameter (φ) is estimated around 0.8 for all the species emphasizing strong 420 between months temporal correlations in the biomass field values. The range parameters 421 are estimated to 55 km for sole and squids while being estimated to 67 km for whiting 422 emphasizing wider spatial autocorrelation for this species.

423 Concerning the sole case study, model predictions highlight the relatively offshore 424 distribution from November to April and a more coastal distribution from June to October 425 suggesting some offshore-coastal migrations between these 2 periods (Figure 6, bottom). 426 In particular, the migration in June/July is associated with a contraction of the sole 427 distribution around the Vendée coast, the Gironde Estuary and the Landes coast (45.5°N-428 46°N) while the migration in November leads to an expansion of the species distribution 429 towards the offshore areas all along the Bay of Biscay. Interestingly, such seasonality 430 coincides with the seasonality of PS intensity for the OTB CEP (Top of Figure 6 and SM 431 9). Higher PS parameter values are associated with a coastal distribution of sole while lower values correspond to offshore distribution of sole. 432

433 Similar maps can be computed for the other species and are presented in SM10.

434 **3.5** Aggregation index and reproduction grounds

For both sole and squids, areas of aggregation persistent over the years identified during 435 436 the spawning period exhibit strong aggregation patterns that match the available 437 knowledge of their reproduction grounds. For sole, the aggregation areas globally match 438 with the observed area of maximum egg concentration (Figure 7), although the spawning grounds identified by egg maps are slightly further East of those identified by our method. 439 440 This slight discrepancy could be interpreted as an effect of the larval drift as the maps 441 provided by Arbault et al. (1986) are concentration of eggs and not reproduction grounds 442 per se. This is consistent with the simulation analysis of Ramzi et al. (2001) showing that the eggs and larval drift is this area of the Biscay Bay is oriented to the East. Overall,these aggregation areas are stable over time (Figure 8).

445 For whiting, similar patterns can be identified during the reproduction period (Figure 7); they match with previous studies investigating the spatial distribution of mature whitings 446 447 (Houise and Forest 1993). In particular, the Northern (3°W-47°N) and the Southern (2°W-45.5°N) aggregation patches are almost systematically significantly considered as 448 449 aggregation areas (aggregation index equals 1) while the other middle one (2.5°W-46.5°N) 450 is classified as an aggregation area that appears less frequently over the years. An 451 additional persistent aggregation area can be identified in the North of the Bay of Biscay 452 (4.5°W-48°N) suggesting that reproduction may also occur in this area which was not 453 identified in the report of Houise and Forest (1993). Interestingly, the Northern aggregation 454 area (3°W-47°N) is more intense at the end of the period (Figure 8).

For squids, no information related to any reproduction ground exists in the literature, only the time period of the reproduction is known (the peak fall between January to April). On this time period, some persistent aggregation areas can be evidenced in coastal areas (Figure 7) along the Vendée coast (2.5°W-46.5°N), the Landes coasts (1.5°W-44°N to 45°N) and around Belle-Île-en-Mer (3°W-47.25°N). Interestingly, the two Northern aggregation areas are more intense at the end of the time series compared to the beginning of the time series (Figure 8).

Maps of persistent aggregation areas are available for all month and evidence some other
aggregation areas outside of the reproduction period (SM11). For instance, for sole a
persistent patch can be identified offshore the Gironde Estuary (1.5°W – 45.5°N) from
August to December.

466 4 DISCUSSION

467 Main findings

In this paper, we develop a framework to infer fish spatio-temporal distribution on a 468 monthly time step while combining scientific survey data and commercial catch 469 470 declarations from several fleets. Commercial catch data constitute a valuable data source that complements scientific survey or onboard sampling programs by providing much 471 higher spatio-temporal sampling density. Those complementary sources of data were 472 473 integrated through a spatio-temporal hierarchical model taking into account spatio-474 temporal variation within the biomass field and PS on a monthly time step. We fitted the 475 model to VMS-logbooks data filtered and processed over the period 2010-2018 for 3 476 demersal species (sole, squids and whiting) in the Bay of Biscay.

477 We emphasize the benefit of integrating several spatially complementary fleets to infer 478 fish distribution throughout the year. We demonstrate how the within-year dynamic of the 479 PS parameters can be interpreted with regards to the joint dynamics of species distribution and fishing distribution and to the overall targeting behavior of the fleets (e.g. OTB CEP 480 for the squids case study). Even though PS parameters are not fishing intention per se 481 482 (Bourdaud et al. 2019), these could advantageously complement information provided by 483 landing profiles to estimate the targeting behavior of any group of vessels (either 484 métier/fleet or any group that would seem appropriate).

Interestingly, although interpretation of the PS parameters provide insight into the spatiotemporal fleet dynamics, accounting for PS in the inferences does not significantly improve model fitting even when some fleets emphasize strong PS (e.g. squids, OTB_CEP). These results contrast with Alglave et al. (2022), and could result from the integration of several

fleets in the analysis that allow a full coverage of the area. Indeed, in Alglave et al. (2022), the fleet emphasizing strong PS only covered a restricted (and coastal) part of the area. As introducing PS mainly affects inferences on poorly sampled areas, predictions in the offshore areas where mostly affected. Here, as the fleets are all estimated to have a positive PS and cover the whole area, PS only downscale the predictions in the few areas that remain unsampled.

495 Filtering the mature fraction of the population in both the scientific and the commercial 496 data make possible to infer the spatio-temporal distribution of the fraction of the biomass 497 that could be potentially mature through the year on a monthly time step. We developed 498 an index to infer aggregation areas of the potentially mature fraction of the biomass that 499 are persistent across years. When calculated on a temporal window predefined following 500 the available information on the reproduction period for each species, the aggregation 501 index enables to identify the main recurrent spatial aggregation areas within the 502 reproduction period. Results demonstrate that the recurrent aggregation areas identified 503 from our method for Sole and Whiting were highly consistent with those already identified 504 in the literature. Our results demonstrate how the aggregation index can provide new 505 insights on the spawning grounds for species like squids for which no information on the 506 spawning grounds is available on the literature. Areas of high aggregation persistent 507 across years were identified during the expected period of reproduction and could be 508 interpreted as spawning grounds. This opens perspectives for applying more 509 systematically the approach for species where no information of reproduction grounds is 510 available to fill the gaps in our knowledge with minimum cost (Delage and Le Pape 2016; 511 Regimbart et al. 2018).

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514 Combining our results with other data sources to refine the identification of 515 spawning grounds

516 Persistent aggregation areas should be considered as potential spawning areas rather than actual spawning areas. Indeed, although the mature fraction of the biomass was 517 518 filtered in the data, our maps do not directly inform whether individuals are actually 519 reproducing or not. The outputs of the model provide maps of the mature fraction of the 520 population (i.e. the individuals that can reproduce) and not the spawning fraction of the 521 population (i.e. the individuals that are actually spawning). However, by focusing on the 522 temporal window identified as reproduction period in the literature, we limit the risk of 523 misinterpreting the aggregation areas as reproduction areas.

524 Our results can also be used to help gathering additional data to identify reproduction 525 grounds. Typically, our maps could be of great help to design surveys to record eggs, 526 larvae and spawning individuals that would provide direct information of species 527 reproduction (Fox et al. 2008). Because developing such additional surveys would be 528 highly expensive, our maps could provide valuable a priori information to optimize the 529 survey design and potentially find a compromise between the cost, the spatial extent, the 530 temporal coverage of the survey and the accuracy of the expected estimates/predictions. 531 Similar ideas were already applied to the sole case study to investigate more precisely 532 the space-time variation of sole reproduction. Arbault et al. (1986) work provided a priori 533 information of reproduction grounds that allowed designing more localized surveys to 534 study inter- and intra-annual variability of one specific sole spawning area (Petitgas 1997). 535 Several statistical methods have been developed since and are suitable to optimize such 536 adaptive sampling design; see for instance the recent work of Leach et al. (2021).

537 Our results could also be combined with fisher expert knowledge (Yochum et al. 2011) to 538 complement our knowledge of fish reproduction (Delage and Le Pape 2016). For instance, 539 Bezerra et al. (2021) and Silvano et al. (2006) proved the usefulness of fishers knowledge 540 to determine the temporality of fish spawning and to identify some spawning grounds by 541 crossing the information of aggregation areas provided by several fisher. These were 542 proved complementary with scientific data as they can be available at low cost and provide 543 local knowledge of fish ecology.

544 Limits and perspectives for the approach

545 Our framework has several limitations that are all material for future research avenues.

546 Our model remains relatively simple with regards to all the temporal processes that 547 actually occur within a fishery. It is both a strength and a weakness: the model remains 548 relatively generic, but one might want to extend it further to account for other temporal and 549 spatio-temporal processes affecting fisheries dynamics. For instance, we opted for a non-550 seasonal representation of the model. One could make it seasonal by decomposing the 551 intercepts $\alpha_{S}(t)$ and $\alpha_{Xi}(t)$ as well as the random effects $\delta(x,t)$ and $\eta(x,t)$ into yearly and 552 seasonal terms in addition to some 'season x year' interaction terms as performed in 553 Thorson et al. (2020). In their work, such specification mainly allowed to provide 554 information over the time-steps where data was lacking. In the configuration of our case 555 studies, data is available for all time steps and have a relatively good coverage of the 556 study domain. Consequently, even though it provides a nice conceptual view of 557 seasonality, complexifying our model in that direction should not deeply modify our 558 inference of the biomass field. Alternatively, our framework could integrate orthogonal 559 spatio-temporal terms in the latent field to capture the main mode of variability of the

560 biomass field (Thorson et al. 2020b). Such orthogonal terms would allow to capture the 561 main spatial patterns that structure the latent field as well as their variation in time. These 562 could prove very useful to identify the structuring processes that affect species distribution 563 and could give a valuable insight in the space-time dynamics of the species. Another 564 exciting research avenue would consist in integrating population dynamics in the latent 565 field of biomass (Cao et al. 2020). This would require to refine further the demographic 566 resolution of the VMS-logbooks data (see for instance Azevedo and Silva 2020), but once 567 done, it would give access to huge data for inferring the space-time dynamics of fish 568 populations. Finally, our model considers fisher preferentially sample areas where the 569 biomass is higher (preferential sampling), but does not consider any other drivers and specifically the temporal and spatio-temporal relations that can affect fishers behavior. 570 571 These can be highly complex and may depend on the distribution of the resource, 572 tradition/habits, management regulations (Abbott et al., 2015; Girardin et al., 2017; Salas 573 and Gaertner, 2004; Hintzen, 2021). These drivers are rarely studied in both space and 574 time (although see Tidd et al., 2015). Our framework could allow to jointly model the 575 dynamics of the species, the distribution of the effort, the link that relates species 576 distribution and effort in space/time and all the other spatial and/or temporal drivers that 577 affect the distribution of fishing effort. For instance, we could relate the fishing intensity to 578 the biomass field from the previous time steps, or alternatively consider that the locational 579 choice depends on the catches of the previous time steps. Adding such covariates and 580 spatio-temporal dependencies in the sampling equation (eq.3) will probably not modify the 581 overall pattern of biomass distribution, but it would make possible to quantify the drivers 582 of fisher behavior and give valuable insight to the fishery dynamics.

583 Including discards would potentially improve our approach. Indeed, logbooks data are 584 landings declarations data which means they inform on the landings and not on the true 585 catch. Thus, by assuming the landings per unit effort are proportional to the biomass, we 586 make the hypothesis that the discard rate is constant in space and time and does not 587 affect model predictions. This should not be a problem for sole and squids as the discards 588 are low and TAC have not been really binding during the studied period. However, the 589 issue might be more stringent for whiting and/or other species with a high and non-590 stationary level of discards. Integrating discards data in the analysis could help solving 591 this issue. Stock et al. (2019) and Yan et al. (2022) used observer data to model bycatch 592 in both space and time and Breivik et al. (2017) used bycatch data from onboard surveys 593 to predict the temporal evolution of bycatch realized in the full commercial data. Similarly, 594 we could integrate into the same analysis the logbooks and the observer data by assuming 595 that the catch of observer represents the sum of landings (which is also observed in the 596 logbooks data) and discards (which is unobserved in the logbooks data). This way, the 597 discards information available from observer data would be shared with the logbooks data 598 and would allow correcting for the missing portion of catch declarations data while possibly 599 accounting for possible space or time variation in the discard rate.

Our analysis rely on the hypothesis that the spawning season is known a priori. Extending the approach to infer the spawning season based on the temporal dynamics of the aggregation patterns could improve our knowledge of species spatio-temporal distribution. In particular, identifying the main species phenomenological phases and their consistency (or shift) in time is crucial in the context of global change (Thorson et al. 2020a). In our study, we computed the aggregation index on a predefined temporal window based on literature assumed to be the reproduction period (Arbault et al. 1986; Houise and Forest

607 1993; Moreno et al. 2002). Several methods exist and could be adapted to extract the 608 spatial patterns that shape model outputs, their related temporal variation and identify the 609 main phenological phases that characterize species distribution (e.g. reproduction, 610 feeding - see for instance Empirical Orthogonal Functions or Principal Oscillation Patterns 611 - Cressie and Wikle 2015; Wikle et al. 2019). While the approach we adopted in the 612 manuscript requires to know the reproduction period of the species and would be 613 inappropriate in a context of a changing reproduction time-span, those alternative 614 methods would not require any a priori. Hence, these methods would be more appropriate 615 to identify phenological modifications in species life cycle in response to climate change. 616 Applying those kind of methods to the huge amount of data available from mandatory declarations, would make possible to track the effect of climate change on fish phenology 617 618 at a monthly/seasonal scale, while it is generally only possible at a yearly time step 619 through scientific survey data (Maureaud et al. 2020).

Last, confidentiality remains a major limitation to the massive use of VMS data (Hintzen 2021). Indeed, there are strong confidentiality constraints on these data due to the huge information available on fisher fishing grounds. Few countries are now giving free access to their data (e.g. Norway), but in most cases administrative procedures to get access to the data remain a burden and still constitute a limitation for the use of 'VMS x logbooks' data for routine operational use.

626 Future use for Marine Spatial Planning

Our model has potential application for Marine Spatial Planning (MSP). Janßen et al. (2018) highlighted that one of the main requirements for implementing MSP is the availability of fine scale information on species distribution and of their essential habitats. Here we propose a method which can provide such information for the fraction of the 631 population available through catch declarations (i.e. mainly the adult fraction and in some 632 cases part of the juvenile fraction). This knowledge is required to design Marine Protected 633 Areas (MPA – see for instance Lambert et al. (2017) or Loiselle et al. (2003)), Fishery 634 Conservation Zones (Delage et Le Pape, 2016; Regimbart et al., 2018), or alternatively 635 identify areas that should be kept for fishing in a context where many other human 636 activities are competing in space and time with fishing (Campbell et al. 2014; Bastardie et 637 al. 2015). This would require to integrate our results into bio-economic models in order to 638 evaluate alternative management regulations and assess their tradeoffs in regards to all 639 the sets of ecosystem services provided through activities such as fishing, aquaculture, 640 energy, shipping, recreation and conservation (Nielsen et al. 2018).

642 **ACKNOWLEDGMENT**

- 643 The authors acknowledge the Pôle de Calcul et de Données Marines (PCDM;
- 644 https://wwz.ifremer.fr/en/Research-Technology/Research-Infrastructures/Digital-
- 645 infrastructures/Computation-Centre) for providing DATARMOR supercomputer on which
- 646 the model has been fitted.
- 647 The authors are grateful to the Direction des pêches maritimes et de l'aquaculture (DPMA)
- and Ifremer (Système d'Informations Halieutiques SIH) who provided the aggregated
- 649 VMS and logbooks data. The findings and conclusions of the present paper are those of
- 650 the authors.

651 **Fundings**

652 The authors declare no specific funding for this work.

653 **COMPETING INTERESTS**

654 The authors declare there are no competing interests.

655 **DATA AVAILABILITY STATEMENT**

556 Survey data are available through the DATRAS portal (https://www.ices.dk/data/data-557 portals/Pages/DATRAS.aspx) with the package 'icesDatras' (https://cran.r-558 project.org/web/packages/icesDatras/index.html). Logbooks and VMS data are 559 confidential data and they are available on specific request to DPMA.

660 **CODES**

661 Toy example codes of the model are available on the github link:

662 https://github.com/balglave/sdm_vms_logbooks

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TABLES 888

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Table 1. Ratio between the negative log-likelihood values (either commercial or scient

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ntific) from the integrated	l model	accounting	for preferential	sampling and
integrated m	odel ign	noring prefer	rential sampling	j.

	Scientific data	Commercial data
Sole	0.97	0.92
Squids	1.00	1.0
Whiting	0.99	1.0

the

896 **FIGURES CAPTION**

Figure 1. Spatial distribution of each fleet on the whole period (2010-2018). Unit: fishing effort in fishing hour. Coordinate system: WGS84.

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900 Figure 2. Diagram of the integrated spatio-temporal model.

901

902 Figure 3. Sole case study. Comparison between the observed scientific CPUE (y-axis) 903 and the corresponding model predictions (x-axis) on the month of the survey, based on 904 model integrating data from one commercial fleet only (either OTB CEP, OTB DEF, OTT DEF) or from all commercial fleets (Integrated model). x-axis: model predictions. y-905 axis: scientific data observations (CPUE in kg/hour). Black line: linear regression 906 907 'log(scientific observations) ~ log(model predictions)'. r: Spearman correlation coefficient. Scientific data are integrated to inference for all models. *** stands for the level of 908 909 significance. Each point is a grid cell in which fall a scientific data point. We compare the 910 scientific observation values to the related prediction values.

911

912 Figure 4. Sole case study. Comparison between predictions (each point is a combination 913 'grid cell x time step'.) from the integrated model (using all fishing fleets) and the model 914 integrating only one commercial fleet for the 12 months of year 2018. Left: OTB CEP 915 fleet, middle: OTB DEF fleet, right: OTT DEF fleet. x-axis: integrated model predictions. y-axis: single-fleet model predictions. The prediction values are log-scaled. Red points: 916 predictions within the sampling area of the related fleets (i.e. the cells sampled by the 917 918 fleet). Black points: predictions outside the sampling area of the related fleets. Black line: 919 x = y axis. Note that the intercept of the x-y line has been scaled to account for differences 920 in the intercept values between models. Scientific data are integrated to inference for all 921 models.

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924 Figure 5. Estimates of PS parameters for each commercial fleet (left) and effect of PS 925 on model outputs (right). Left: boxplot represent the variability of maximum likelihood 926 estimates of parameters b across all monthly time steps. Right: log-predictions of the 927 integrated model accounting for PS (y-axis) versus log-predictions of the integrated model 928 ignoring PS (x-axis) for the 12 months of year 2018. Blue points: predictions within the 929 sampling area of the commercial fleets (i.e. the cells sampled by commercial fleets). Black 930 point: predictions outside the sampling area of the commercial fleets. Black line: x = y931 axis.

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Figure 6. Sole case study. (Top) Temporal evolution of the *b* parameters for the three commercial fleets fitted to the integrated model. Blue vertical lines: January. (Bottom) Monthly biomass distribution averaged on the full period. Only quantile values are represented. Model predictions come from the integrated model accounting for PS.Coordinate system: WGS84.

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939 Figure 7. Left: index of persistence (average over 2010-2018) during the reproduction 940 period of sole (February), whiting (March-May) and squids (January-April). Reproduction 941 period defined from ecological expertise. Right: literature information on reproduction 942 grounds when available. For sole, the map represents egg concentration from an egg and larvae survey conducted in 1982 (Arbault et al., 1986). For whiting, the map represents 943 944 records of age-2+ whiting (i.e. mature individuals), from two spring trawl surveys that 945 occurred between 1987 and 1992 (Houise and Forest, 1993). Model predictions come 946 from the integrated model accounting for PS. Coordinate system: WGS84.

947

- 948 Figure 8. Persistence indices within the reproduction period computed on a 5-years
- mobile time-span for each 3 species (5-years time span indicated on the top of each map).
- 950 Model predictions come from the integrated model accounting for PS. Coordinate system:
- 951 WGS84.

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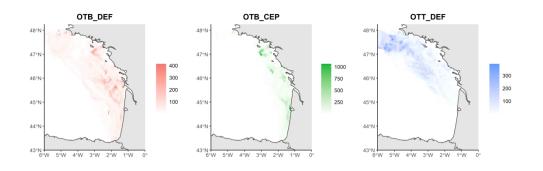
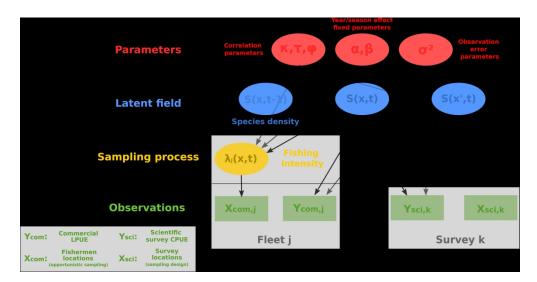
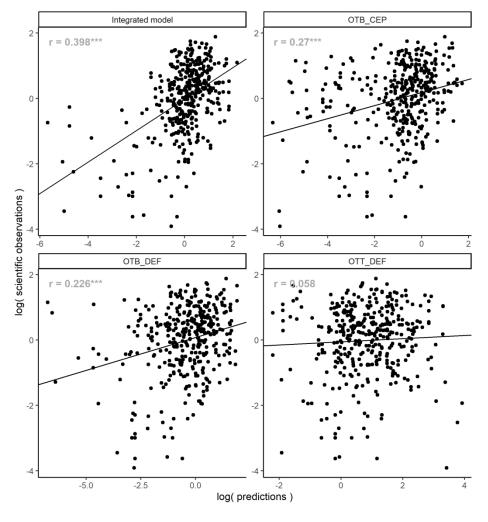


Figure 1 774x258mm (118 x 118 DPI)



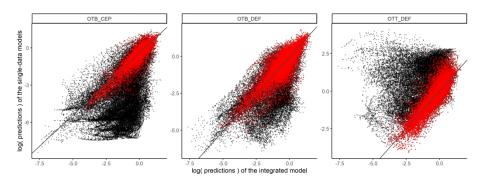


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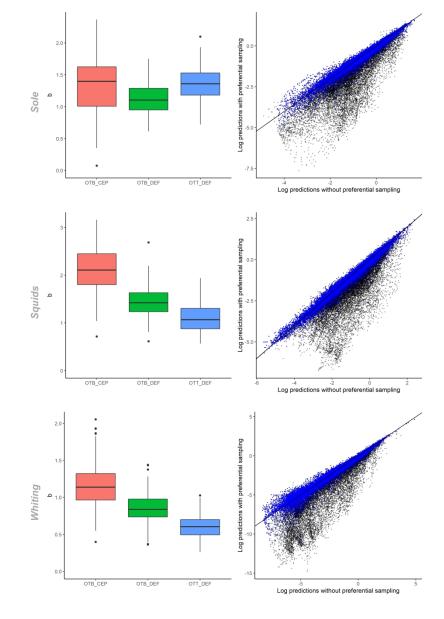


Figure 5 645x968mm (118 x 118 DPI)

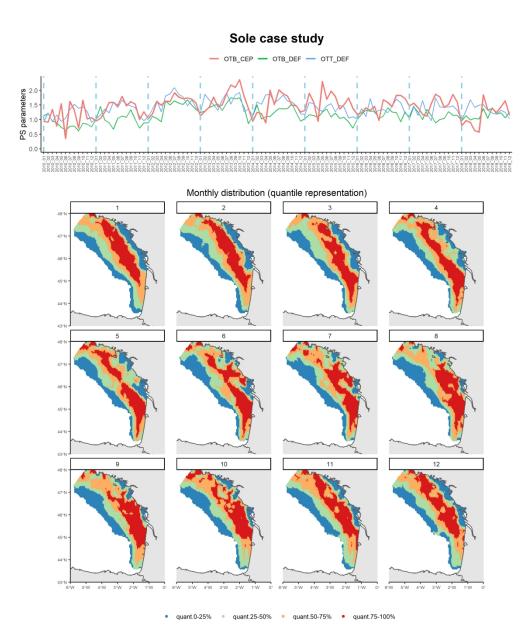
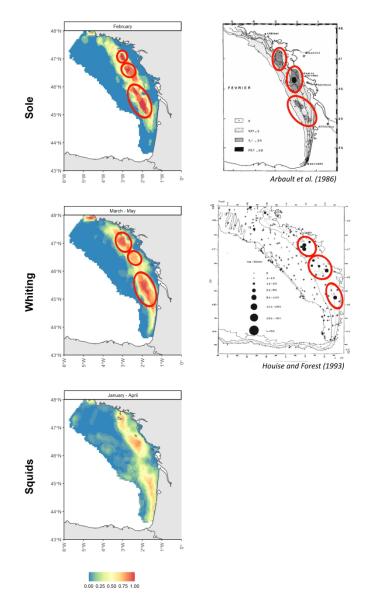
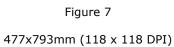


Figure 6

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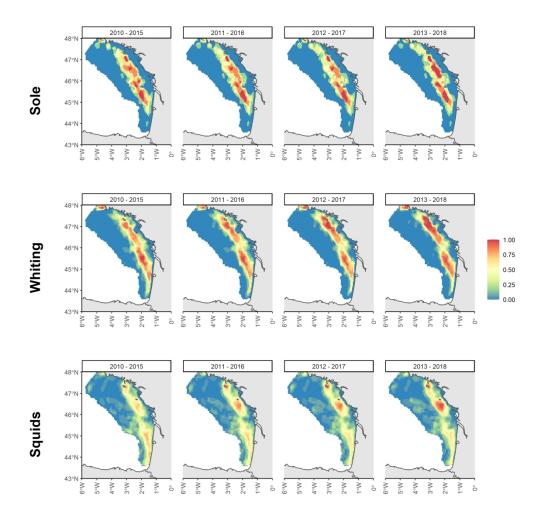


Figure 8 645x645mm (118 x 118 DPI)