



## End-to-end model of Icelandic waters using the Atlantis framework: Exploring system dynamics and model reliability

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### ABSTRACT

Icelandic waters are very productive and the fisheries are economically important for the Icelandic nation. The importance of the fisheries has led to progressive fisheries management and extensive monitoring of the ecosystem. However, fisheries management is mainly built on single species stock assessment models, and multi-species or ecological models are essential for building capacity around ecosystem-based fisheries management. This paper describes the first end-to-end model for the Icelandic waters using the Atlantis modeling framework. The modeled area is 1,600,000 km<sup>2</sup>, and covers the area from Greenland through Icelandic waters to the Faroe Islands. The ocean area was divided into 51 spatial boxes, each with multiple vertical layers. There were 52 functional groups in the model: 20 fish groups (8 at a species level), 5 groups of mammals, 1 seabird group, 16 invertebrates, 5 primary producers, 2 bacteria and 3 detritus groups. The reliability of the model was evaluated using a skill assessment and a sensitivity analysis was conducted to understand the dynamics of the system. The sensitivity study revealed that saithe, redbfish and tooth whales had the greatest effect on other groups in the system. The skill assessment showed that the model was able to replicate time-series of biomass and landings for the most important commercial groups and that modeling of the recruitment processes was important for some of the groups. This model now provides a solid basis for evaluating alternative ecosystem and fisheries management scenarios, and should produce reliable results for the most important commercial groups.

### 1. Introduction

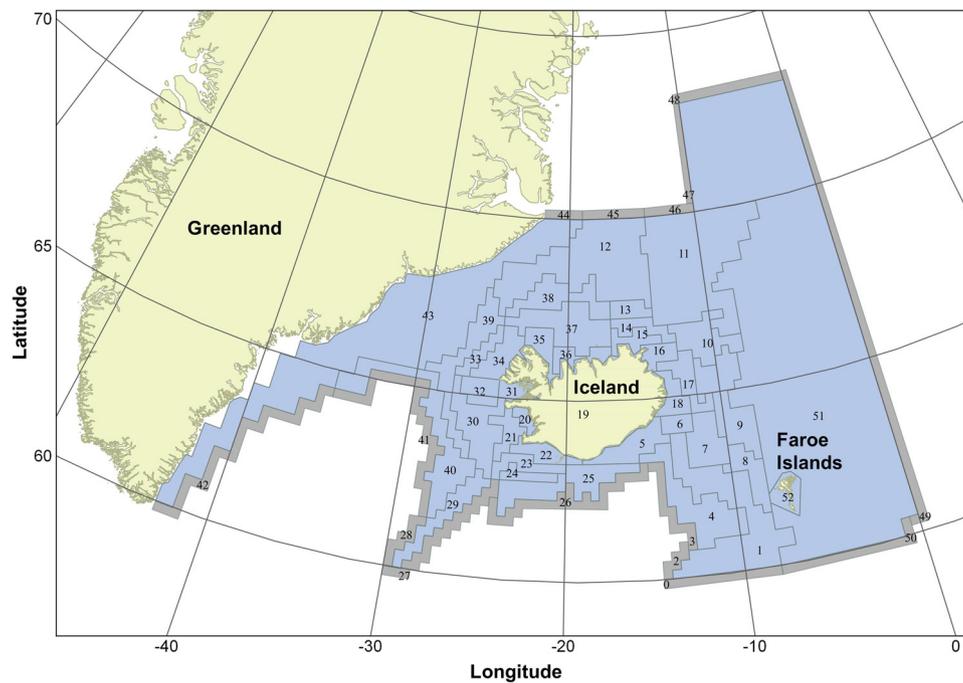
Icelandic waters, where the relatively warm Atlantic water and the cold Arctic water meet, are very productive (Astthorsson et al., 2007). The annual harvest from these waters is around 1.3 million tones, which is 1.4% of the world's harvest (Statistics Iceland, 2017). The fisheries are economically important for the Icelandic nation and they have, along with fish processing, accounted for 6–11% of the GDP and 37–63% of the exports since 2002 (Statistics Iceland, 2017). The highest catches are of capelin (*Mallotus villosus*), but cod (*Gadus morhua*) has the highest commercial value.

The importance of the fisheries has led to progressive fisheries management, and Iceland was one of the first nations to implement a quota system (Hilborn, 2007; Matthíasson, 2003). The ecosystem monitoring program is extensive and a bottom trawl survey is carried out twice annually (Anon., 2010) while acoustic surveys are conducted

for pelagic species (Anon., 2016; Vilhjálmsson and Carscadden, 2002). The environmental conditions around Iceland are also monitored annually where nutrients, temperature, salinity and plankton is measured (Anon., 2015). In spite of extended datasets, including data on stomach contents, fisheries management advice is mainly built on single species stock assessment models for the most important commercial species (Anon., 2016). Nevertheless, there has been increased demand for ecosystem-based fisheries management (EBFM) in recent years. Single species models do not consider species interactions, which are an important factor in EBFM (Link, 2002). Multi-species and ecosystem models, where species interactions, and in some cases environmental factors, are considered are tools that can be used to support EBFM (Plagányi, 2007). Two preliminary food web models have been built for Icelandic waters (Buchary, 2001; Mendy, 1998), but have not been tested or used for fisheries management. A dynamic ecosystem model could support an EBFM and allow fisheries scenarios concerning the

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**Fig. 1.** The modeled area and the locations of the 53 spatial boxes. Active boxes are in blue and boundary boxes in grey. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article).

most important commercial groups, e.g. the effects of stop fishing capelin, an important prey species in the system, to be evaluated.

Modeling of marine ecosystems has increased in recent years, along with developments in computational power, along with a growing understanding of ecosystem functioning and increased data sampling (Fulton, 2010). End-to-end models have become possible, where ecosystem and human components are integrated. They are not appropriate for tactical management advice (e.g., quota setting), unlike the single species models, but are useful to evaluate system-level trade-offs of alternative management strategies. Ecopath with Ecosim (EwE), a trophically-focused ecosystem model, has become widely used (Christensen and Walters, 2004; Fulton, 2010), but more complex models such as Atlantis are becoming more widely used (Fulton, 2010; Fulton et al., 2011; Nyamweya et al., 2016; Ortega-Cisneros et al., 2017).

Atlantis (Audzijonyte et al., 2017a, 2017b; Fulton et al., 2011) is a spatially resolved deterministic end-to-end model designed for exploited marine ecosystems. The modeling framework consists of four sub-models: biophysical, fisheries, management and socio-economic. It has been used to explore major processes and responses in systems (Kaplan et al., 2014; Nyamweya et al., 2016) and it has been used for management strategy evaluations (MSE, Fulton et al., 2007).

Different ecosystem models (e.g. Atlantis vs. EwE) for the same areas are not always consistent and have shown contradicting predictions (Forrest et al., 2015; Pope et al., 2018). Such an ensemble modeling approach can provide major insights into uncertainty around system structure and function. This is important as the modeling process for these models is subjective, as formal parameter estimation is prohibited by the complexity of the models. Instead, they are currently typically manually calibrated to historical data. This source of potential uncertainty means that even when not being used in an ensemble, a model skill assessment is an important means of determining how reliable models are, i.e. how well they fit to existing data and how well they predict (Olsen et al., 2016). The prediction ability of models is however usually not assessed because all existing data are used to calibrate the model (with subsequent use focused on relative projections rather than focusing on absolute predictions). Olsen et al. (2016) however performed a skill assessment on the predictive capacity of the Atlantis model for the northeast US, ten years after the calibration,

when new data had been acquired. They recommend using a several metrics to assess the different aspects of the skill of the model, e.g. one that measures correlation and another that measures scale mismatch. They concluded that the forecasting skill of the model for the northeast US was comparable with the hindcasting skill, and did not degenerate for a medium-term forecasting.

Finding means of assessing uncertainties and performance for large ecosystem models is important, as they have both inherent structural and parametric uncertainty. Unfortunately, their size has meant traditional approaches to assessing parametric uncertainty (let alone structural uncertainty) have been impractical due to the curse of dimensionality, rapid growth of complexity in multi-parametric analyses and sensitivity to experimental design due to the feedback influences on time dependence of parametric sensitivity results (Fulton, 2010; Fulton et al., 2011). A sensitivity analysis can give insight into which parameters contribute the most towards uncertainty in the output (Pantus, 2007; Saltelli et al., 2006). However, a complete sensitivity analysis is not feasible for Atlantis because it has thousands of parameters and numerous possible interactions. Therefore, sensitivity analysis of Atlantis models have been carried out for each parameter one-at-time (Murray and Parslow, 1997) or for interactions between a selection of parameters, which are already known to have a strong influence on model performance or are particularly pertinent to that system type (Ortega-Cisneros et al., 2017).

This paper describes the first end-to-end model for Icelandic waters using the Atlantis modeling framework. The aim with this work is to describe the model, compare its output to available data and evaluate its reliability using a skill assessment. The aim is also to investigate how sensitive the output is to changes in parameters and to use a partial sensitivity analysis to understand the dynamics of the system.

## 2. Material and methods

### 2.1. Study area

The study area, the Icelandic waters, extends from 60° to 73°N and from 43° to 0°W (Fig. 1). Two water masses meet in this area, the relatively warm and saline Atlantic water and cold Arctic water with low

salinity (Astthorsson et al., 2007). The mixing of these two water masses causes rather unstable environmental conditions, which has a substantial impact on the production of the lower trophic levels (Gislason et al., 2016). Primary production is higher in the warm Atlantic water south of Iceland than in the cold Arctic water to the north and east of the island (Astthorsson et al., 2007). This has an impact on both the productivity and distribution of the fish stocks. The main spawning grounds of the most important commercial species are in the warm water to the south while the nursery grounds are in the colder water to the north (Astthorsson et al., 2007). Around 30 fish species and seven invertebrate species are commercially harvested in Icelandic waters, along with whales and seals (Anon., 2016). The most important commercial species is cod because of its high catches and commercial value. Another important fish stock is the capelin, being the most abundant pelagic stock in Icelandic waters and an important prey for many demersal species such as cod, saithe (*Pollachius virens*) and Greenland halibut (*Reinhardtius hippoglossoides*). Capelin feed in the northern most part of the area and transfer large amount of energy to the more southern grounds with their feeding and spawning migrations (Vilhjálmsón, 1994). Over 20 species of mammals inhabit Icelandic waters where they have great influence on the ecosystem; it has been estimated that marine mammals consume over 6 million tons of fish, cephalopods and zooplankton annually (Sigurjonsson and Vikingsson, 1997). Icelandic waters also support large populations of seabirds (Lilliendahl and Solmundsson, 1997).

## 2.2. Model structure

### 2.2.1. The oceanography model

The modeled area is 1,600,000 km<sup>2</sup> and covers the area from Greenland through the Icelandic waters and to the Faroe Islands (Fig. 1). The area has been divided into 51 ocean boxes and two land boxes based on work described in Stefánsson and Pálsson (1997) and in Taylor (2005) where the division was mainly based on hydrography, bathymetry and species distribution. Areas outside of the survey coverage are divided into larger boxes because of less information on species distribution. Active boxes (where the biology was modeled) numbered 36, with an additional 15 boundary boxes to buffer water fluxes to and from waters beyond the model domain. Each box was further divided into vertical layers depending on the depth of the box. The boxes each have one sediment layer and can have a maximum of six water column layers (0–50 m, 50–150 m, 150–300 m, 300–600 m, 600–1000 m and 1000 m+). The oceanographic data were taken from a hydrodynamic model (Logemann et al., 2013) and water fluxes, temperature and salinity were calculated for each box and layer each day to create a forcing time series running from 1948 to 2012. A full model run is therefore 65 years and the time step of the model is 12 h.

### 2.2.2. Biological model

There were 52 functional groups in the model: 20 fish groups (8 represented at a species level), 5 groups of mammals, 1 seabird group, 16 invertebrates, 5 primary producers, 2 bacteria and 3 detritus groups (Tables 1 and 2). The vertebrate groups could have up to ten age classes, each of which could contain multiple annual cohorts. The model tracks numbers and weight (mg N) per age class and the weight was divided into reserve and structural weight, where reserve weight was soft tissues and structural weight the bone structure. Cephalopods and shrimp have two age classes, juveniles and adults. Other groups were represented as aggregate biomass pools with no explicit age structure. The initial conditions of most of the vertebrate groups, i.e. their biomass and weight per individual were acquired from data sampled by the Marine and Freshwater Research Institute (MFRI) or from reports from the Institute (Anon., 2016). Initial condition for zooplankton and primary producers were acquired from Astthorsson et al. (2007).

The consumption rate of prey *i* by predator *j* ( $CR_{ij}$ ) was modeled

**Table 1**

The vertebrate groups: the group code, group name and species, their maximum age, reproduction function (BH = Beverton-Holt, BH-f = Beverton-Holt with recruitment scalars, C = constant per adult), if the group is being harvested and if a group is migratory.

Code	Group	Max age	Reprod.	Harvest.	Mig.
FCD	Cod ( <i>Gadus morhua</i> )	20	BH-f	Yes	No
FHA	Haddock ( <i>Melanogrammus aeglefinus</i> )	20	BH-f	Yes	No
FSA	Saithe ( <i>Pollachius virens</i> )	20	BH-f	Yes	No
FRF	Redfish ( <i>Sebastes</i> sp)	50	BH	Yes	No
FGH	Greenland halibut ( <i>Reinhardtius hippoglossoides</i> )	20	BH	Yes	No
FFF	Flatfish	20	BH	Yes	No
FHE	Herring ( <i>Clupea harengus</i> )	20	BH-f	Yes	No
FCA	Capelin ( <i>Mallotus villosus</i> )	6	BH-f	Yes	No
FMI	Blue whiting ( <i>Micromesistius poutassou</i> )	20	BH	Yes	Yes
FMA	Mackerel ( <i>Scomber scombrus</i> )	20	BH	Yes	Yes
FOC	Other codfish	20	BH	Yes	No
FDC	Demersal commercial	20	BH	Yes	No
FDL	Other demersal fish	10	BH	No	No
FSD	Sandeel fish	10	BH	No	No
FDL	Long lived demersal	30	BH	No	No
FMP	Large pelagic fish	30	BH	No	No
FBP	Small pelagic fish	10	BH	No	No
SSR	Skates	30	BH	Yes	No
SSD	Small sharks	50	C	No	No
SSH	Large sharks	100	C	No	No
SB	Seabird	40	C	No	Yes
PIN	Pinniped	40	C	No	No
WMW	Minke whale ( <i>Balaenoptera acutorostrata</i> )	50	C	No	Yes
WHB	Baleen whale	100	C	No	Yes
WHT	Tooth whale	70	C	No	No
WTO	Other tooth whale	30	C	No	No

**Table 2**

Invertebrates, primary producers and detritus groups in the model – indicating their major habitat type (benthic vs pelagic), whether the group is explicitly age structured and whether the group is harvested.

Code	Group	Age-structure	Benthic/pelagic	Harvested
CEP	Cephalopod	Yes	Pelagic	No
PWN	Shrimp	Yes	Pelagic	No
ZS	Microzooplankton	No	Pelagic	No
ZM	Mesozooplankton	No	Pelagic	No
ZL	Macrozooplankton	No	Pelagic	No
ZG	Gelatinous zooplankton	No	Pelagic	No
LOB	Norway lobster	No	Benthic	No
BML	Other megazoobenthos	No	Benthic	No
SCA	Iceland scallop	No	Benthic	No
QUA	Ocean quahog	No	Benthic	No
CUC	Cucumbers	No	Benthic	No
BD	Deposit feeder	No	Benthic	No
BFF	Other benthic filter feeders	No	Benthic	No
BG	Benthic grazer	No	Benthic	No
BC	Benthic carnivore	No	Benthic	No
BO	Meiobenthos	No	Benthic	No
PL	Diatom	No	Pelagic	No
PS	Pico-phytoplankton	No	Pelagic	No
MA	Macroalgae	No	Benthic	No
SG	Seagrass	No	Benthic	No
DF	Dinoflagellates	No	Pelagic	No
PB	Pelagic bacteria	No	Pelagic	No
BB	Sediment bacteria	No	Benthic	No
DL	Labile detritus	No		No
DR	Refractory detritus	No		No
DC	Carrion	No		No

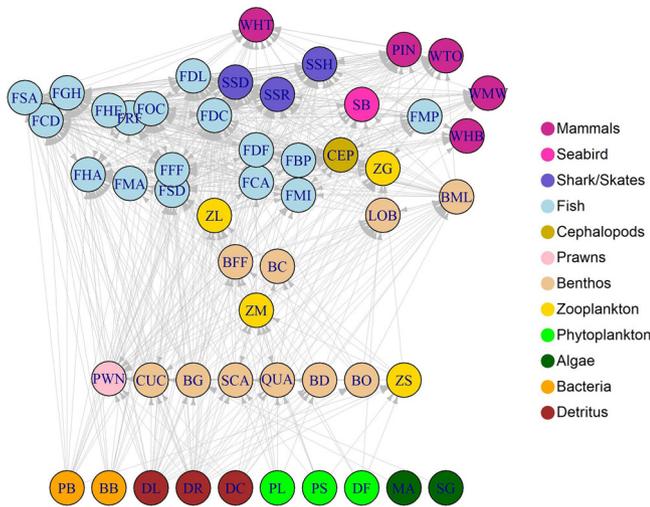


Fig. 2. Food web connections between the modeled functional groups (see Tables 1 and 2 for group codes).

using an adjusted Holling type II:

$$CR_{ij} = \frac{C_j \cdot a_{ij} \cdot B_i}{1 + \frac{C_j}{mum_j} [\sum_{k=1}^n a_{kj} \cdot B_k \cdot E_{kj}]} \quad (1)$$

where  $mum_j$  is the maximum growth rate and  $C_j$  is the clearance rate of predator  $j$ ,  $B_i$  is the biomass of prey  $i$ , and  $a_{ij}$  is the availability of prey  $i$  to predator  $j$ . The ratio between  $C$  and  $mum$  determines the steepness of the consumption curve and  $E_{kj}$  is the assimilation rate of prey  $k$  for predator  $j$ . The diet composition for each predator was adjusted by tuning the availability of each prey. Data from the MFRI on stomach content and information from the literature (Gunnarsson et al., 1998; Jónsson and Pálsson, 2013) was used as a guideline when tuning the availability of each prey. The modeled food web is quite complex (Fig. 2, graphed using the cheddar and igraph packages in R, Csardi and Nepusz, 2006; Hudson et al., 2016).

Recruitment of the fish groups was modeled with the Beverton-Holt function that describes the relationship between the spawning stock biomass and number of recruits as follows:

$$R = \frac{\alpha * SSB}{\beta + SSB} \quad (2)$$

where  $R$  is the number of recruits,  $\alpha$  is the maximum number of recruits,  $\beta$  represents the size of the spawning stock which gives half of the maximum recruitment, and  $SSB$  is spawning stock biomass, which depends on individual weight and on the proportion spawning in each age-class across the model domain (Audzijonyte et al., 2017a). Data from MFRI were available for the most important commercial groups to parameterize the recruitment curve. For other groups, the assumed natural mortality and initial numbers were used to set the maximum number of recruits. The recruitment of the mammals and the seabird groups was modeled as a constant per adult (Table 1). It is possible to induce recruitment spikes in Atlantis by scaling the recruitment from the Beverton-Holt curve and this was done for the cod, haddock (*Melanogrammus aeglefinus*), saithe, herring (*Clupea harengus*) and capelin. Numbers per age-class as estimated by MFRI were used to calculate time-series of recruitment scales. The recruitment of the mackerel (*Scomber scombrus*) was scaled down before the year 2000 to imitate the invasion that took place after 2000.

The functional groups had different spatial distributions that were allowed to vary by season and for juveniles and adults. The distributions of the groups, which were kept fixed, were based on survey data from the MFRI or from the literature (Jónsson and Pálsson, 2013). Groups could also migrate into and out of the model area. The model

includes five migratory groups: blue whiting (*Micromesistius poutassou*), mackerel, seabirds, minke whale and baleen whales.

### 2.2.3. Fisheries model

Harvests are modeled for the most important commercial species (Table 1 and 2). Each group is harvested by fishing gear represented using a length-specific logistic selectivity curve. Data on the size distribution of the catch and stock, as estimated by the MFRI, were used to parameterize the selectivity curves for cod, haddock and saithe. The current model did not have a dynamic fisheries model connected to economics. Instead, time-series of harvest mortality were used to drive the fisheries. The harvest mortality for each day is multiplied by selectivity for each age class of each species. The harvest mortality was allowed to change with time, but selectivity was assumed temporally invariant. Harvest mortality was the same in all boxes and layers. Discards were included in the model for two functional groups, cod and haddock, and were based on Pálsson et al. (2012).

### 2.3. Skill assessment

A skill assessment was conducted to measure how well the model fits to available data with and without recruitment spikes. Numerous metrics exist to compare the model output with data (Bennett et al., 2013; Stow et al., 2009) and while some of them are redundant (Olsen et al., 2016), multiple metrics are necessary to evaluate model skill (Olsen et al., 2016). In the present study three metrics were chosen, demonstrating in three different ways how the model fits to data: 1) model efficiency (MEF), 2) Pearson’s correlation ( $r$ ) and 3) the reliability index (RI). They are defined as follows:

$$MEF = \frac{\sum_{i=1}^n (O_i - \bar{O})^2 - \sum_{i=1}^n (P_i - O_i)^2}{\sum_{i=1}^n (O_i - \bar{O})^2}, \quad (3)$$

$$r = \frac{\sum_{i=1}^n (O_i - \bar{O})(P_i - \bar{P})}{\sqrt{\sum_{i=1}^n (O_i - \bar{O})^2 \sum_{i=1}^n (P_i - \bar{P})^2}}, \quad (4)$$

$$RI = \exp \sqrt{\frac{1}{n} \sum_{i=1}^n \left( \log \frac{O_i}{P_i} \right)^2}, \quad (5)$$

where  $O_i$  and  $P_i$  are the  $i$ th of  $n$  observations and predictions, respectively and the  $\bar{O}$  and  $\bar{P}$  are the corresponding averages. These metrics capture different aspects of model performance. MEF measures how well the model fits to the data compared to the average. A perfect fit is 1, 0 is no better than using the average of the data points and negative values correspond to a model that is worse than simply using the average of the data in terms of providing direct biomass estimates (although a model with negative MEF may still be useful if it has the same trend as the data). The correlation measures how the observations and the model prediction vary together, i.e. if the model has a similar trend to the data. The correlation is from  $-1$  to  $1$ , where  $1$  is a perfect positive, linear association,  $0$  is no linear association and,  $-1$  is a perfect negative, linear association. The closer this metric is to  $1$ , the better the model. However, this metric can be  $1$  even if the model is far from the observations, i.e., when the predictions differ from the observations by a constant factor. The third metric, the reliability index, measures how far on average the predictions and the observations are from each other. Ideally, this measure should be close to  $1$ .

The model biomass was compared to observed estimates of biomass from the MFRI, which were available for five groups: cod, haddock, saithe, herring and capelin. The predicted landings were compared to landings data for 12 functional groups: cod, haddock, saithe, herring, capelin, redfish (*Sebastes* sp), Greenland halibut, flatfish, blue whiting, mackerel, other codfish and demersal commercial.

**Table 3**

Growth rate of pico-phytoplankton (PS), diatoms (PL) and macrozooplankton (ZL) in the base run (b) and where these parameters were decreased by 20% (d) or increased by 20% (i).

Model run	PS	PL	ZL
PSb_PLb_ZLb	0.7	0.4	0.8
PSb_PLb_ZLd	0.7	0.4	0.64
PSb_PLb_ZLi	0.7	0.4	0.96
PSb_PLd_ZLb	0.7	0.32	0.8
PSb_PLd_ZLd	0.7	0.32	0.64
PSb_PLd_ZLi	0.7	0.32	0.96
PSb_PLi_ZLb	0.7	0.48	0.8
PSb_PLi_ZLd	0.7	0.48	0.64
PSb_PLi_ZLi	0.7	0.48	0.96
PSd_PLb_ZLb	0.56	0.4	0.8
PSd_PLb_ZLi	0.56	0.4	0.64
PSd_PLb_ZLd	0.56	0.4	0.96
PSd_PLd_ZLb	0.56	0.32	0.8
PSd_PLd_ZLd	0.56	0.32	0.64
PSd_PLd_ZLi	0.56	0.32	0.96
PSd_PLi_ZLb	0.56	0.48	0.8
PSd_PLi_ZLd	0.56	0.48	0.64
PSd_PLi_ZLi	0.56	0.48	0.96
PSi_PLb_ZLb	0.84	0.4	0.8
PSi_PLb_ZLi	0.84	0.4	0.64
PSi_PLb_ZLd	0.84	0.4	0.96
PSi_PLd_ZLb	0.84	0.32	0.8
PSi_PLd_ZLd	0.84	0.32	0.64
PSi_PLd_ZLi	0.84	0.32	0.96
PSi_PLi_ZLb	0.84	0.48	0.8
PSi_PLi_ZLd	0.84	0.48	0.64
PSi_PLi_ZLi	0.84	0.48	0.96

#### 2.4. Sensitivity analysis

A sensitivity analysis was conducted to determine how sensitive the model output was to some of the model inputs, i.e., parameters and oceanographic data. Some effort has been put into developing methods for sensitivity analysis for complex and large models in recent years (Oakley and O'Hagan, 2004; Pantus, 2007; Saltelli et al., 2010). These methods require restricting consideration to a subset of parameters from the Atlantis model, as there are thousands of parameters in the model and that would overwhelm the analytical approaches.

Instead of conducting a rigorous and computationally expensive sensitive analysis, a preliminary and simpler sensitivity analysis was carried out. To see how sensitive the model was to the production of the vertebrate groups, parameters that control recruitment ( $\alpha$  in the Beverton-Holt function, Eq. (2), and a parameter controlling the constant recruitment per adult) were increased and decreased one at a time by 20%. A sensitivity analysis was also conducted to assess how sensitive the model was to production of the low trophic levels groups (phytoplankton and zooplankton) by altering their production by 20% and considering interactions between the selected parameters (Table 3), as was done for the Atlantis model of the Benguela system (Ortega-Cisneros et al., 2017). How sensitive the model was to the oceanographic data (temperature, salinity and water fluxes) was also explored. It is common to repeat years of oceanographic data in Atlantis models for the hindcast (Nyamweya et al., 2016; Ortega-Cisneros et al., 2017) and forecast (Kaplan et al., 2012). In the present model there are time-series of oceanographic data of 65 years which cover the whole simulated period. To explore the effects of oceanographic data, five of the warmest years (2003–2007) were repeated 13 times to cover the 65 year simulation run (Fig. 3) and effects on biomass of the functional groups in the model were assessed.

When only one parameter was perturbed at a time, as was done with the recruitment parameters, a measure of model sensitivity was calculated as described in Murray and Parslow (1997):

$$S_{ij} = \frac{V_i(1.2\alpha_j) - V_i(0.8\alpha_j)}{0.4V_i(\alpha_j)}, \quad (6)$$

where  $S_{ij}$  is the sensitivity measure for the biomass of group  $i$  when recruitment parameter ( $\alpha$ ) in the Beverton-Holt function (see Eq. (2)) is perturbed for group  $j$ ,  $V_i(-)$  is the average biomass of group  $i$  for the whole simulated period, i.e.  $V_i(1.2\alpha_j)$  is the average biomass of group  $i$  when the recruitment of group  $j$  is increased by 20%. If  $S = 1$  then the biomass changes by 20% when the recruitment is changed by 20%. If  $S > 1$  then the change in biomass is higher and if  $S < 1$  then the change in biomass is lower.

The measure  $S$  is no longer applicable when sensitivity to interactions between parameters is explored. Instead percentage change in biomass is used to measure sensitivity. The same was also done when examining the effects of the oceanographic data. In that case, the effects the oceanographic data had on the trend of the biomass was also considered. This was achieved by calculating the correlation of biomass from the base run and biomass under the modified oceanographic run.

### 3. Results and discussion

#### 3.1. Species interactions

An important part of ecosystem modeling is modeling the species (or functional groups) interactions. This was done using the Holling type II function (Eq. (1)). The diet composition of the predators resembled what was observed in the stomach content data for most groups (Figs. 4 and 5). The predators were feeding on the correct groups, but they were relying too much on zooplankton and benthic invertebrates in the model than what the stomach data indicated. The zooplankton could however be under-represented in the stomach content data because of differences in digestion rates (Hyslop, 1980). Bias towards the invertebrates will result in weaker species interactions between the vertebrate groups. Also, sandeel were not as large a component of the diet of its predators as they should have been. However that group collapsed over time in the model, which it also did in the ecosystem (Lilliendahl et al., 2013).

How much each group needs to feed to maintain their individual weight was modeled with the assimilation parameter ( $E$ ) in the Holling type II function (Eq. (1)) and with the respiration function (Audzijonyte et al., 2017a). The weight each group loses due to spawning also affects how much they need to consume to maintain their weight. This, along with the diet composition, controls how strong the species interactions are. The Atlantis model does not report detailed consumption statistics and therefore results on this are not shown. The strength of the species interactions can be investigated using a sensitivity analysis as was the case in this study (see Section 3.3).

#### 3.2. Simulated biomass and landings

There were few groups with biomass that decreased substantially towards the end of the model run. These groups were: sandeel (6% of initial biomass), gelatinous zooplankton (10% of initial biomass), pico-phytoplankton (10% of initial biomass), macroalgae (7% of initial biomass) and dinoflagellates (0.1% of initial biomass). Sandeel was also observed to decline in the ecosystem (Lilliendahl et al., 2013). The biomass of pico-phytoplankton has very large seasonal variations and is reported for January in the model when it is lowest, but the initial biomass represents the situation during the summer months. The biomass of gelatinous zooplankton and macroalgae decreased to 7–10% of its initial biomass, but because they still exist in the system and because they are not important part of the diet of other groups (except benthic grazers) this should not cause problems for the behavior of the system. The dinoflagellates were a group that went close to extinction. The sensitivity study showed that this group is outcompeted by the pico-phytoplankton (see Section 3.3.2). The abundance of the other two

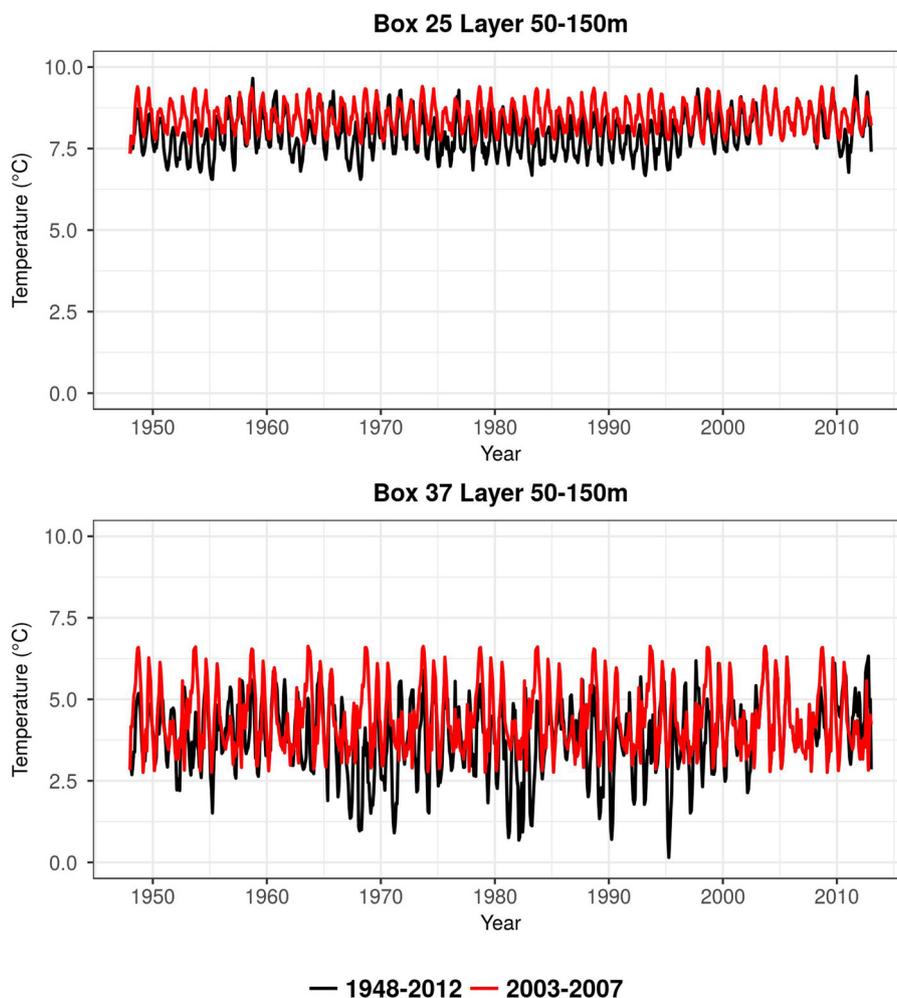


Fig. 3. Temperature in the 50–150 m layer in box 25 (south of Iceland, see Fig. 1) and in box 37 (north of Iceland, see Fig. 1) from 1948 to 2012 when the full time-series was used (1948–2012, base run) and when five years (2003–2007) were repeated.

phytoplankton groups seems to be sufficient to make the system dynamics work, but the model would be more stable if it was possible to achieve balance between the phytoplankton groups.

The simulated biomass changes either because individual weight changes (Fig. 6), the numbers changes, or both. The simulated biomass of some of the commercial fish groups (Fig. 7) increased in the start of the model run, but decreased rapidly when fishing pressure was increased. The exception was for the blue whiting, which dropped in biomass at the start of the model run, when the individual weight dropped (Fig. 6k), and did not increase until around 1980, but then showed a decreasing trend when harvesting of that group began around 2000 (Fig. 7k). Mackerel was modeled as an invasive species and its biomass increased rapidly after 2000 (Fig. 7l). The burn in period was included in the model run, which can explain the rapid changes at the start of the simulation when the model has not stabilized.

The simulated biomass trajectories of five groups were compared to estimated biomass with and without recruitment spikes (Fig. 7). The skill assessment showed that the simulated biomass of cod fitted best to the estimated biomass (Fig. 7a), both MEF and the correlation were the highest and RI the lowest (Table 4). The simulated biomass of the pelagic species, herring and capelin (Fig. 7d and e), did not fit the observations as well as the biomass of the demersal species and had a MEF lower than zero and a weak correlation. When the recruitment spikes were taken out of the model, the fit became worse for all these five groups. The cod and the saithe had positive MEF and high, positive correlation without the recruitment spikes but haddock, capelin and herring in the model were not able to get the trend in biomass without

having the recruitment spikes included.

The model cannot be expected to achieve the correct trend when forecasting for the groups that required recruitment spikes to achieve positive correlation in the skill assessment. However, it is possible to achieve uncertainty in the forecast by running the model under different recruitment processes. It also must be kept in mind that this is an ecosystem model and not intended for tactical advice, but rather for strategic advice where the accuracy of specific annual recruitment estimates is not as important.

The simulated landings, which were forced in the model with a time-series of harvest rates had reasonable fit to the data for most of the 12 groups (Fig. 8). Saithe (Fig. 8c), other codfish (Fig. 8i) and the demersal commercial group (Fig. 8j) had a good fit in terms of all three metrics (Table 4). The landings of the flatfish group (Fig. 8h) had the poorest fit, with the MEF just above zero and a correlation of 0.58. The MEF was positive for all groups, except for haddock when it was modeled without recruitment spikes (Fig. 8b and Table 4). The magnitude of the simulated landings was not far off for most groups; seven of the groups were within 50% of their observed values. Note that the RI index is very sensitive if there are a few years where the magnitude is incorrect. This was the case for blue whiting and mackerel where the landings were very low in some years. In these cases, the total difference in tons was not high, but because the landings were low, the difference in magnitude could be large. All groups had positive correlation to the landings data except haddock, which had a negative correlation when the model had no recruitment spikes. The correlation decreased for the cod when the recruitment spikes were taken out of the model, but

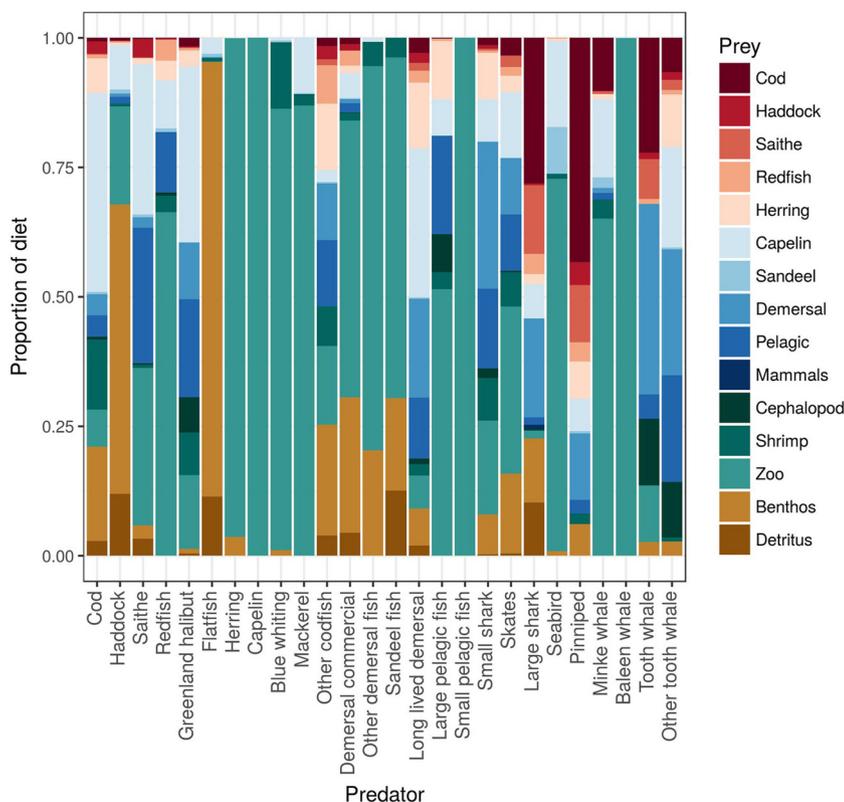


Fig. 4. The average simulated diet composition for the vertebrate groups (age-class 4) over the period 1948–2012. Some of the prey groups have been aggregated into the following groups: demersal (all other demersal fish), pelagic (all pelagic fish except herring and capelin), mammals (all mammals and seabirds), zoo (all zooplankton groups), benthos (all benthic invertebrates) and detritus (detritus and discards).

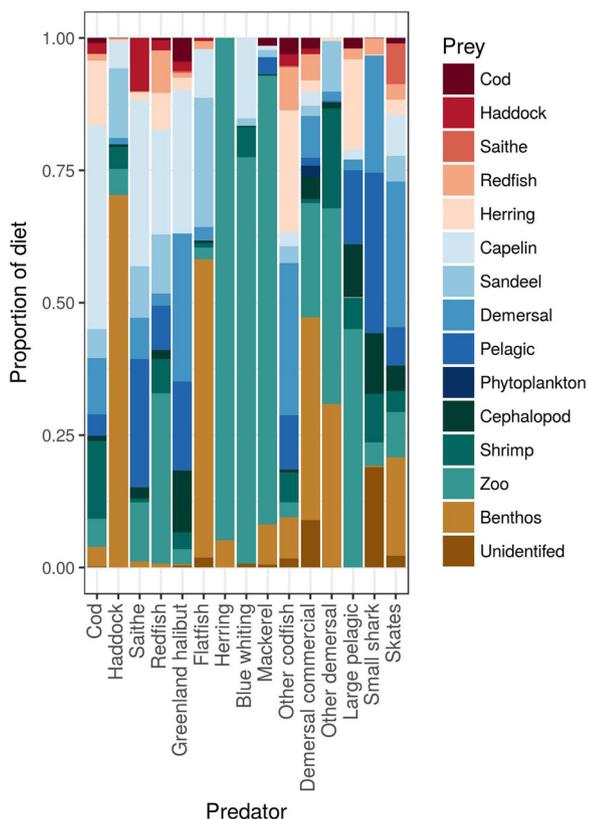


Fig. 5. Average diet composition from stomach content data that was available for 15 of the 20 fish groups.

increased for the saithe. Modeling without the recruitment spikes did not have as much of a negative effect on the landings as it did on the biomass for the pelagic groups (i.e. capelin and herring). It was possible

to get a negative correlation for the biomass but positive correlation for the landings (Table 4).

Skill assessment has now been conducted for most recent Atlantis models, e.g. for Lake Victoria (Nyamweya et al., 2016) and the Benguela and Agulhas currents (Ortega-Cisneros et al., 2017). The present model has similar skill as these two models where most groups had correlation higher than 0.5 and positive MEF. Skill assessment has been performed for Atlantis models using biomass estimates (Ortega-Cisneros et al., 2017) or catch per unit effort where biomass estimates were not available and landings data (Nyamweya et al., 2016). Olsen et al. (2016) also used ecosystem indicators to conduct skill assessment for the Atlantis model for the northeast US. This was not attempted for the present model as the data needed to calculate these indicators were not available.

Discards were simulated for cod and haddock and compared to estimated discards (Fig. 8). The discard rate was on average 3.8% for cod and 7.0% for haddock over the simulated period when recruitment was modeled with recruitment spikes. This is consistent with what has been estimated for haddock, where the discard rate has been estimated to be from 2 to 22% in the last three decades (Pálsson, 2002; Pálsson et al., 2012). The simulated discard rate for cod was higher than what has been estimated, about 1% in recent years (Pálsson et al., 2012), but no estimates exist from the last century. Note that the method used to estimate the discard rate estimated the minimum rate and that the actual discard rate can be assumed to be higher.

### 3.3. Sensitivity analysis

#### 3.3.1. Recruitment parameters

For each group, altering the recruitment parameter for the group had an effect on that group, but the size of that effect varied greatly among groups (Fig. 9). The groups which that were the most sensitive to changes in the recruitment parameters were herring, mackerel, sandeel, seabirds and pinnipeds. Redfish, capelin, long lived demersal, small sharks and tooth whales were the least sensitive.

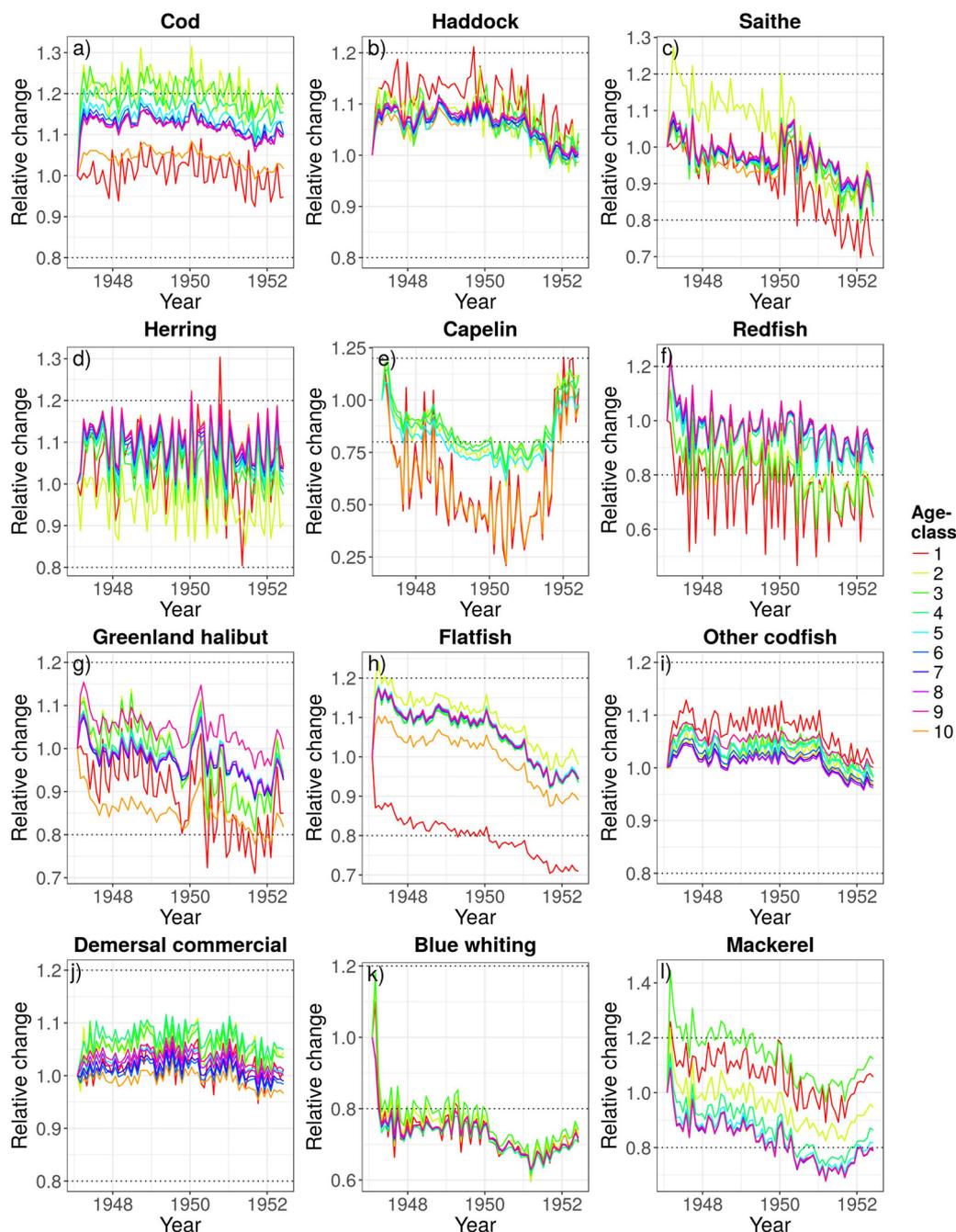


Fig. 6. Relative change in weight (reserve + structural weight) of each age-class of 12 commercial fish groups.

The change in the recruitment parameter was 20%, but the change in actual recruitment could be much greater over time because of the shape of the Beverton-Holt curve. The asymptote of the curve was increased, which leads to higher recruitment. With higher recruitment the stock can recover over time and produce higher spawning stock that shifts the population to the right of the curve which will consequently lead to even more recruitment, if the recruitment was not at the maximum in the base run. This was the case for herring, mackerel and sandeel, which had  $S > 1$  (Fig. 9). The seabird, large shark and the mammal groups were modeled with constant recruitment per adult with no explicit asymptote. Changing the recruitment parameter for these groups by 20% also led to a greater change in actual recruitment by the end of the model run.

Most of the groups had  $S < 1$  which means that changing the recruitment parameters by 20% resulted in less than a 20% change in the

biomass. This was because other parameters affected how much groups could increase in biomass. The quadratic mortality parameter controls the density dependent non-predation mortality; as numbers in an age-class increase so does the mortality. Cannibalism also restrains population growth when recruitment is increased and this may be the reason why the stock size for the redfish group changed little with increased recruitment. Also, redfish is a long-lived group (5 years within an age-class), which means that it takes a long time for the change in recruitment to have an effect on the numbers in the older age-classes. Changing the maximum recruitment of capelin by 20% had little effect on its biomass ( $S = 0.11$ ). The capelin numbers increased when maximum recruitment was increased by 20%, but the individual weight dropped, resulting in very similar biomass as in the base run. The inverse happened when maximum recruitment was decreased, their numbers decreased but individual weight increased and resulted in

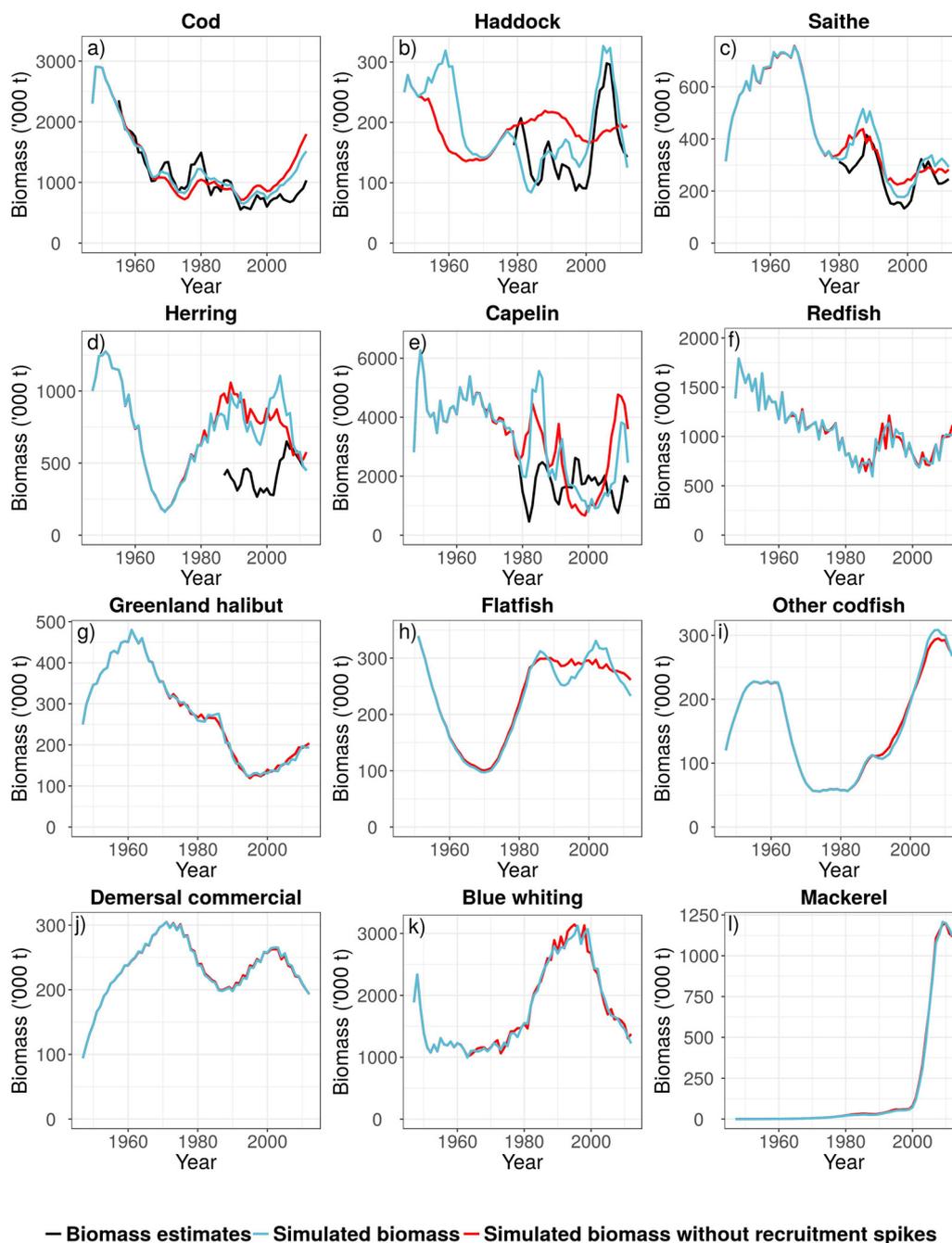


Fig. 7. Simulated biomass with and without recruitment spikes for 12 commercial fish groups and compared to biomass estimates for 5 commercial fish groups.

similar biomass.

Saithe, redfish and tooth whales had the largest effect on other vertebrate groups which were mostly negative effect on their prey groups. They also had positive effects on few groups by affecting the biomass of their predators. Vertebrate groups that were the most sensitive to changes in the recruitment of other groups were mackerel, flatfish, sandeel, small pelagic fish, seabirds and minke whale. It was necessary to analyze what the effects were on individual weight (reserve and structural weight) and numbers rather than simply considering their biomass to understand how the groups affected each other. It is also helpful to look at the diet composition of the groups to understand their interactions.

Groups can have an effect on other groups if they feed on that group or if they feed on its predators. For example, redfish had an effect on saithe ( $S = -0.28$ ) because it feeds on it, even though saithe only

makes up a very small portion of the redfish diet. Redfish had an opposite effects on the flatfish group ( $S = 0.35$ ) by consuming the predators of flatfish, such as saithe.

There can also be indirect effects on a group if they compete for the same prey groups. For example, altering the recruitment of sandeel affected the flatfish group ( $S = -0.33$ ). The individual weight of flatfish increased when the recruitment of sandeel was decreased and vice versa when the recruitment was increased. That affected the spawning biomass of the flatfish, which consequently affected the recruitment leading to changes in their numbers and biomass.

Altering the recruitment of the vertebrate groups also had an effect on the invertebrates, primary producers and the detritus groups (Fig. 10). The most sensitive groups were pico-phytoplankton and microzooplankton. The seagrass and benthic invertebrate groups were not sensitive to a change in the recruitment of the vertebrates except the

**Table 4**

Skill assessment with and without recruitment spikes: The three metrics model efficiency (MEF), reliability index (RI) and correlation (r) for biomass and landings (see Eqs. (3)–(5) for the metrics).

Group	Metrics with recruitment spikes						Metrics without recruitment spikes					
	Landings			Biomass			Landings			Biomass		
	MEF	RI	r	MEF	RI	r	MEF	RI	r	MEF	RI	r
Cod	0.41	1.24	0.74	0.73	1.23	0.88	0.23	1.32	0.64	0.55	1.32	0.78
Haddock	0.38	1.31	0.75	0.64	1.27	0.84	-0.75	1.53	-0.28	-0.29	1.50	-0.31
Saithe	0.55	1.23	0.86	0.40	1.24	0.83	0.72	1.20	0.89	0.58	1.24	0.81
Herring	0.65	1.70	0.87	-0.59	1.38	0.28	0.76	1.67	0.90	-2.22	1.56	-0.45
Capelin	0.73	1.98	0.87	-6.21	1.90	0.32	0.65	2.29	0.82	-9.51	2.33	-0.22
Redfish	0.19	1.36	0.77				0.20	1.35	0.78			
Greenland halibut	0.54	1.69	0.75				0.54	1.68	0.75			
Flatfish	-0.05	1.44	0.50				0.13	1.41	0.59			
Other codfish	0.53	1.24	0.79				0.48	1.26	0.75			
Demersal commercial	0.52	1.28	0.77				0.52	1.28	0.77			
Blue whiting	0.93	2.29	0.97				0.93	2.32	0.97			
Mackerel	0.82	2.33	0.98				0.82	2.44	0.98			

benthic grazer group, which was sensitive towards a change in sandeel and haddock recruitment. Capelin and sandeel seemed to have the most effect on the plankton groups but had opposite effects on the pico-phytoplankton. Both these groups had high biomass (at least at the start of the model run) and have zooplankton as the most important component of their diet, but sandeel also relies on benthic invertebrates. Dinoflagellates were a group that went close to extinction (0.1% of its initial biomass) throughout the base model run, but altered recruitment of cod and sandeel changed its biomass. The low biomass of the dinoflagellates should not be a problem as the other two phytoplankton groups compensated for the low biomass of them.

It has been observed that removal of top predators can have cascading effects down the food web, where the removal led to 45% decrease in large zooplankton and a slight increase in phytoplankton (Frank et al., 2005). The changes in recruitment of vertebrate groups led to large changes in phytoplankton biomass that may not be realistic and this should be studied further before the model is used for management strategy evaluation.

### 3.3.2. Growth parameters of plankton groups

The effects of changing the growth parameters of pico-phytoplankton, diatoms and macrozooplankton on the biomass of the functional groups in the model are given in Fig. 11. Changing the growth rate of the macrozooplankton had very little effect except when the growth rate of diatoms had also been increased. The results of altering the growth rate of the macrozooplankton are therefore not discussed further here. Dinoflagellates appeared to suffer competitive exclusion, being outcompeted by the pico-phytoplankton (for example, dinoflagellates multiply when the growth rate of the pico-phytoplankton is decreased; Fig. 11). This increased biomass of dinoflagellates led to an increased biomass of microzooplankton.

The change in the growth of the phytoplankton also had a considerable effect on the vertebrate groups. The vertebrate groups that were the most sensitive were: baleen whales, minke whale, seabirds, small and large pelagic fish, sandeel, mackerel and blue whiting, capelin and redfish (Fig. 11). Zooplankton is a large portion of the diet of these groups (Fig. 4) and as the zooplankton responded to the phytoplankton, these planktivorous feeders did as well. The redfish biomass increased by 84% when the growth rate of pico-phytoplankton was decreased and the rate of diatoms was increased. All vertebrate groups increased in biomass when the growth rate of both the phytoplankton groups was increased and all vertebrate groups except mackerel decreased in biomass when just the diatom growth rate was reduced (Fig. 11). A sensitivity study on the Atlantis model for the Benguela system also showed strong reactions to changes in the growth rate of the plankton groups, where some of the fish groups showed more than

100% increase in biomass (Ortega-Cisneros et al., 2017).

### 3.3.3. Oceanographic data

Repeating the five warm years instead of having the full oceanographic time series affected some of the functional groups. Temperature affects the respiration of the vertebrate groups and consequently their growth. The water fluxes were also different between the two model runs, influencing advection of nutrients and plankton groups; the shift in nutrients also affected the growth of the primary producers. This methodology can be used when testing the effect of climate change that could have larger effects on the groups, especially the vertebrate groups if density-dependent movement is turned on in the model, which was not the case in the present model.

The oceanographic data had a large effect on the biomass of the phytoplankton groups: pico-phytoplankton and diatoms (Fig. 12a). The biomass of the pico-phytoplankton increased, while the biomass of the diatoms decreased. Diatoms need higher nutrient concentrations than pico-phytoplankton to achieve their maximum growth rates. The warm conditions and associated water fluxes were therefore more favorable for the pico-phytoplankton. The warm conditions were also favorable for cephalopods, capelin and mackerel, leading to increased biomasses compared to the base run. In contrast, these conditions led to decreased biomass of herring. Capelin and herring feed mostly on zooplankton, but capelin feeds both on mesozooplankton and macrozooplankton, while herring feeds mainly on macrozooplankton, which increased in biomass while the mesozooplankton decreased in biomass. It was necessary to look at the output from the model for each spatial box to better understand why the capelin biomass was higher in the case when the five warm years were repeated. At the individual spatial box level, it was observed that the biomasses of both mesozooplankton and macrozooplankton were much higher in one box in the north, where the capelin feed during the winter (Box 12, Fig. 1), with the modified oceanography than in the base run. The model output had to be explored at a high temporal resolution to better understand why the herring biomass was lower when the warm years were repeated. The reserve to structural weight ratio of the herring is lower in the early years of the simulation with modified oceanography, which affected the recruitment and ultimately led to a lower number of adults later in the simulation (perpetuating lower recruitment).

Using the five warm years did not have a substantial effect on the trend of the biomass of the vertebrate groups (Fig. 12b), not even herring and capelin, which had changes in total biomass. It did have some effect on the large pelagic fish, baleen whales and the other toothed whales. The correlation between the zooplankton and phytoplankton groups was low, indicating that the trend was different from the base run. The other benthic filter feeders group had a negative

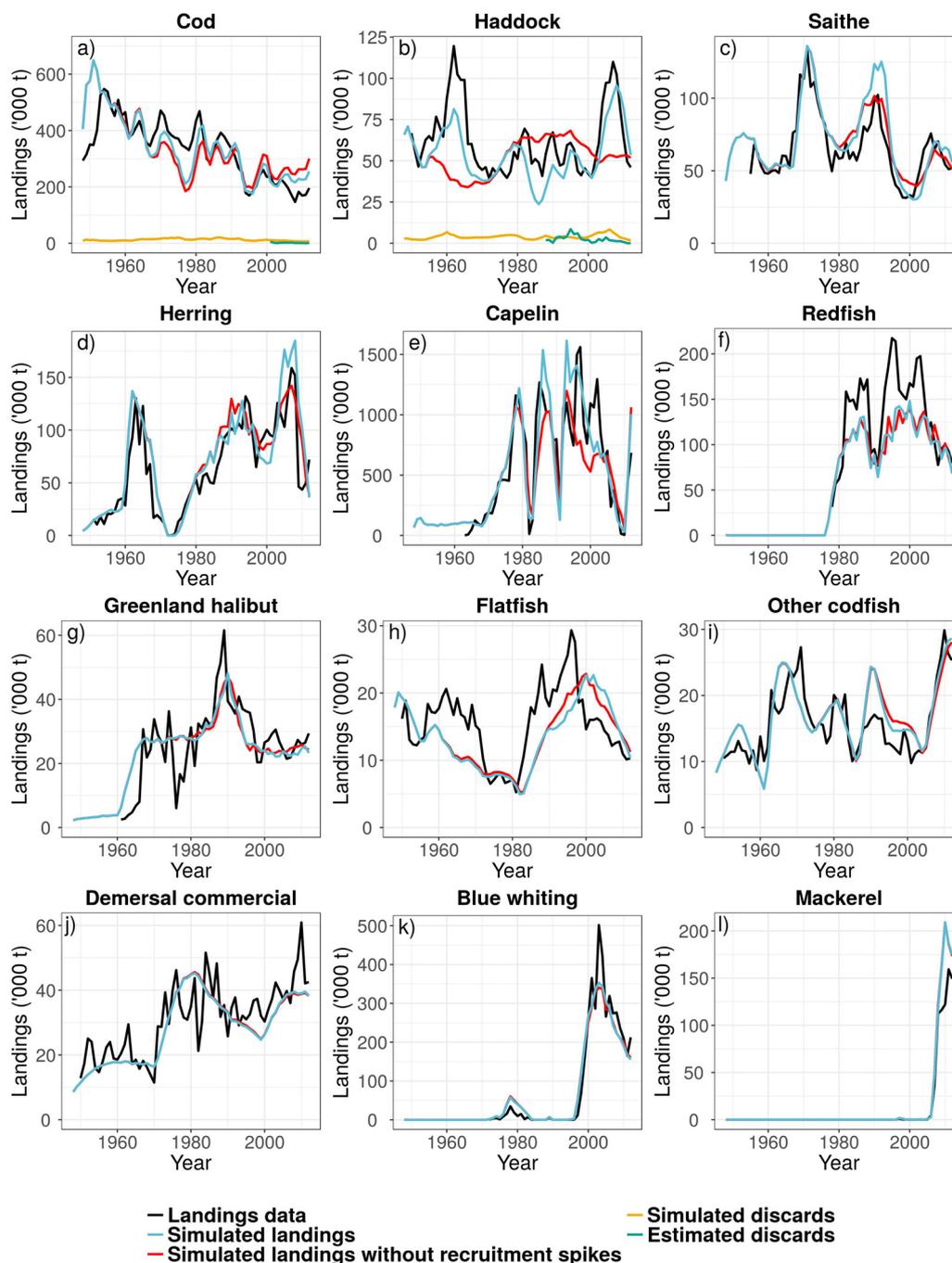


Fig. 8. Simulated landings with and without recruitment spikes and discards from the Atlantis model compared to landings data for 12 commercial fish groups.

correlation, which means that the trend of the biomass under the modified oceanography was in the opposite direction compared to its biomass in the base run. It is therefore important to have correct oceanographic data if trends of low tropic levels are the focus of a study.

### 3.4. Model reliability

Kaplan and Marshall (2016) have set some standards that end-to-end models such as Atlantis should reach before they are used in management strategy evaluations. Models should fulfill the following:

- 1) All biological functional groups should persist throughout the model run.
- 2) The model should achieve equilibrium, i.e., under fixed

environmental forcing the unfished model should have stable biomass over the final 20 years for most vertebrate groups.

- 3) The biomass trends should be compared to survey time-series (hindcast).
- 4) Qualitative model comparisons to survey data.
- 5) The model should capture the dynamics of abundant species.
- 6) Most functional groups should qualitatively match expected productivity.
- 7) Natural mortality should be realistic.
- 8) Age and length structure of vertebrate groups should match data.
- 9) Diet composition of the functional groups should match diet data.

In relation to these standards, all groups except the dinoflagellates (which had only 0.1% of its initial biomass) persisted throughout the model run. Few groups (sandeel, gelatinous zooplankton, pico-

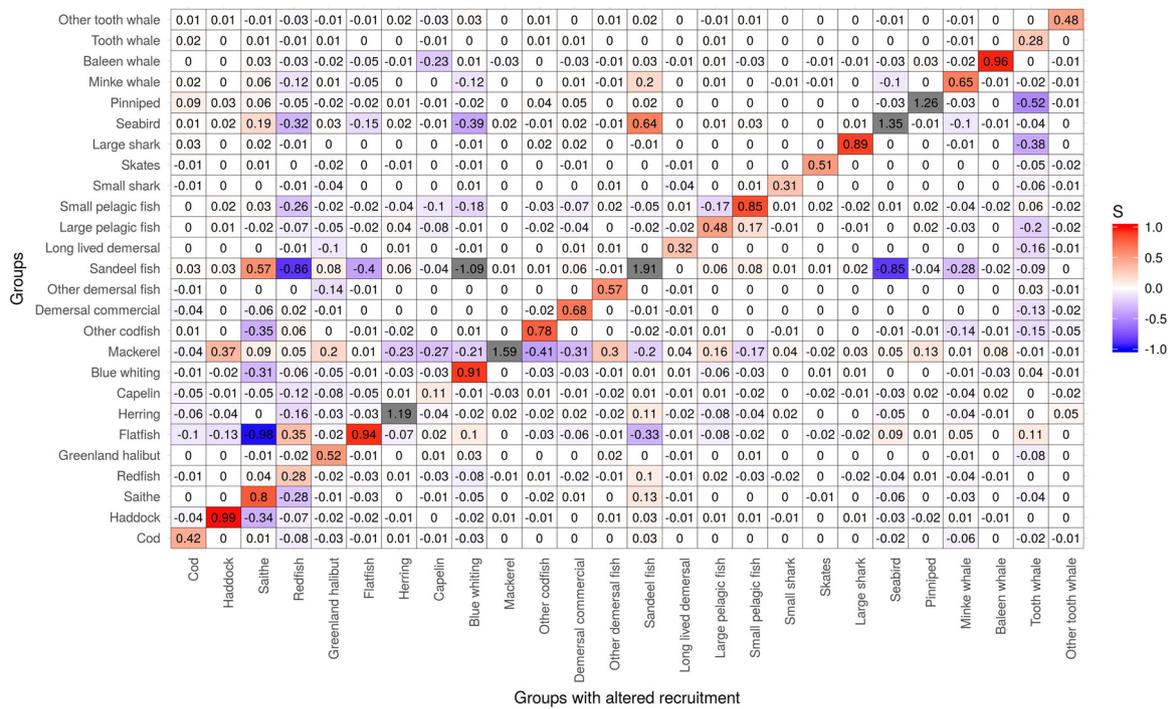
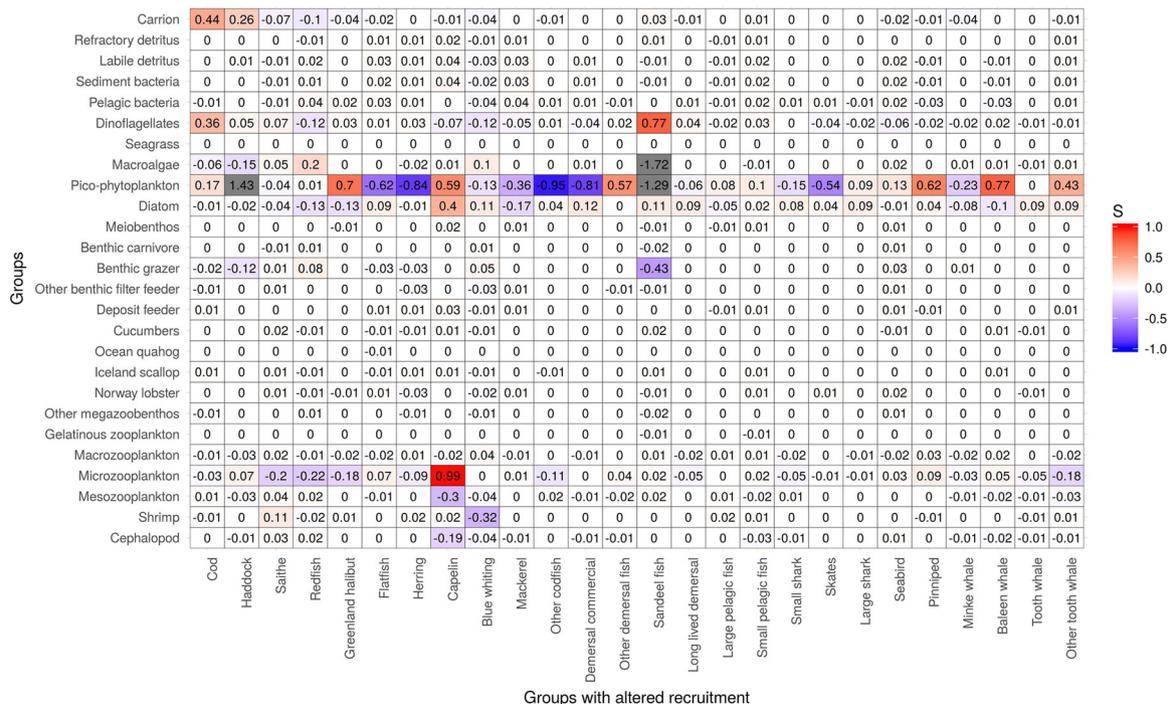


Fig. 9. Sensitivity analysis: the metric S (see Eq. (6)) for the change in biomass of the vertebrate groups when recruitment was altered for the vertebrate groups. The grey color represent  $S < -1$  or  $S > 1$ .

phytoplankton and macroalgae) decreased considerably in biomass but still persisted at the end of the model when they had 7–10% of its initial biomass.

Kaplan and Marshall (2016) did not define what is considered stable biomass. In this study stable biomass will be considered if the biomass changes by less than 10% over the last 20 years of the simulated period. The model was run 100 years further, with the same fishing pressure that was used in the final year of the base model and with

oceanography rerunning from the start of the model run. After 100 years of forecast 23 of the 26 vertebrate and 14 of the 16 invertebrate groups had stable biomass in the last 20 years of the model run. If fishing was excluded from the model and the model run was 165 years only 15 of the 26 vertebrate groups had stable biomass in the last 20 years of the model run. Herring showed fluctuations and did not have any trend if more than 20 years were considered. It is not realistic here for all groups to reach a perfectly constant biomass as the underlying



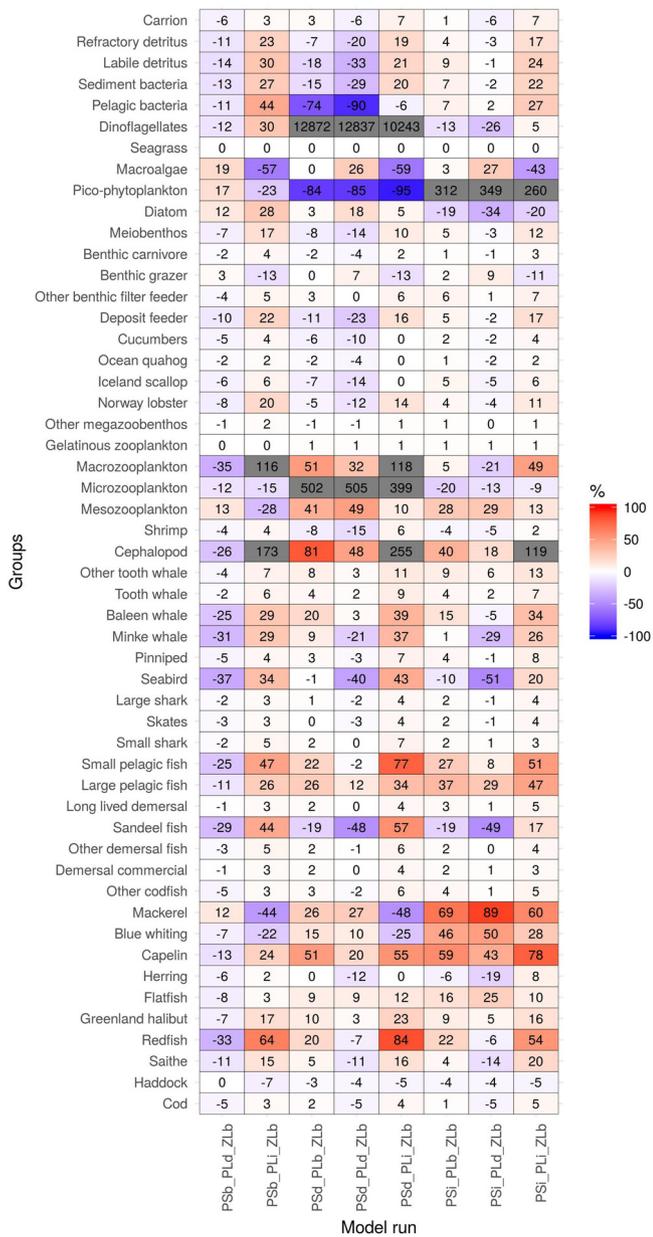


Fig. 11. Sensitivity analysis: Percentage change in biomass of all functional groups when the growth rate of pico-phytoplankton (PS), diatoms (PL) and macrozooplankton (ZL) were decreased by 20% (d), increased by 20% (i) or kept unchanged (b). See Table 3 for parameter values in the model runs. Grey color represents more than 100% change.

physical model always has some fluctuations. The physical forcing (oceanographic data) contains seasonal and annual variations in temperature, salinity and water fluxes, which consequently leads to fluctuations in phytoplankton biomass, which escalates up the food chain. It is difficult to assess what groups compose the same trend as the phytoplankton because of the large variation in phytoplankton biomass, both in space and time.

Most of the fish groups had realistic natural mortality that decreased with age. Capelin had higher mortality for adults than juveniles, but that group is set as semelparous, i.e., all fish die after spawning. The whale groups had lower mortality for the juveniles. This is a consequence of the quadratic mortality parameter settings used for this group (which is different between juveniles and adults). This should be fixed by increasing the mortality for the juveniles and the recruitment rate in order to maintain suitable biomass levels and trajectories before

the model is used to predict effects on the whale groups or scenarios regarding whaling.

The model was calibrated so that the weight (or length) of each age-class within a group would be relatively constant over time (< 20% change from initial conditions). This was achieved for most groups, but some age-classes went over (or under) the limit (Fig. 6).

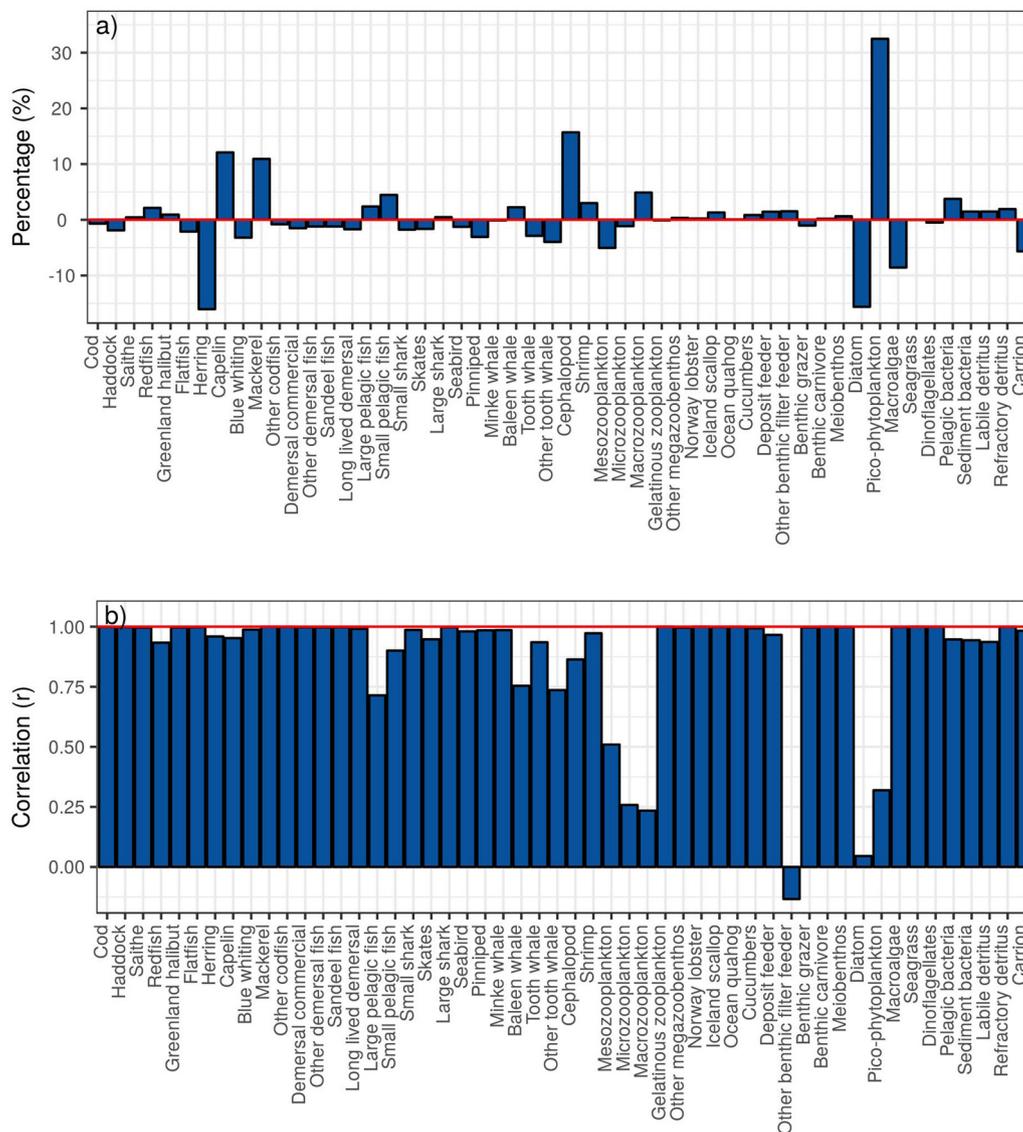
The diet composition matched the data (Section 3.1). It is a challenging task to calibrate an ecosystem model of this complexity such that both diet composition and changes in diet over time are represented correctly. The most important is to get the proportions similar in order for the species interactions to be realistic and that has been accomplished for the current model. Minor adjustments to the availability of prey or to spatial distribution of the functional groups may improve the model and that can be determined by conducting another skill assessment of an adjusted model.

The simulated biomass was compared to biomass estimates and simulated landings to landings data and a skill assessment conducted (Section 3.2). Kaplan and Marshall (2016) did not state how well the model needs to fit to the data before it is considered reliable and neither did Olsen et al. (2016). However, it can be concluded that models that cannot replicate historical time-series will not be able to produce reliable predictions about the future. Modeled groups that have negative correlation in the skill assessment do not replicate historical time-series. It is more difficult to define what values the metrics MEF and RI need to reach to be able to conclude that the model is reliable. When  $MEF < 0$  then a straight line through the average would give a better fit, but if the correlation is positive the model can still be useful and the same can be said about high RI. The model could however be used with more confidence when the correlation, MEF and RI are close to 1 but the minimum requirement is that the correlation is positive. The skill assessment showed that the model had a good fit to the data, when modeled with recruitment spikes, for the most important commercial groups, i.e. the correlation was positive,  $MEF > 0$  and  $RI < 1.5$  for most groups (Table 4). However, most of the groups in the model had no biomass estimates and are not targeted by the fisheries and therefore had no landings data for calibration and skill assessment purposes. It was therefore not possible to determine the skill of the model for those groups. Some of the non-commercial groups are caught in the survey and it is possible to calculate a survey index for those groups and calculate correlation between the simulated biomass and the survey index. However, such analysis was not conducted for this paper because these indices may not be reliable as the survey is not designed to capture population trends for these species. Skill assessment could also be carried out for plankton groups. However, this is challenging as these groups show large variation, both in space and time. It would also be of interest to assess how well the model captures the ocean currents. It can then be evaluated if the model would benefit from an improvement of the spatial structure of the model. This is a work for further research.

Ideally, one should not use the same data for model calibration and for skill assessment of that model (Bennett et al., 2013). This was not possible in this case as all available data were needed for calibration and therefore the model reliability could be overestimated.

### 3.5. How can this model be used for ecosystem based fisheries management?

The goals of the Icelandic fisheries management act have largely been met. Since its inception, harvest rates of commercially exploited fish stocks have gradually been decreased, with the aim of achieving single species maximum sustainable yield. The need for alternative management strategies that take ecosystem and socio-economic considerations into account has, however, increased in recent years. The model could be used to evaluate fisheries scenarios for the most important commercial species taking interactions into account. It provides a solid basis for the testing of alternative ecosystem and fisheries management scenarios and will be of further use when the socio-ecological model component has been integrated with the biophysical and fisheries



**Fig. 12.** The effects of the oceanographic data, when five years of oceanographic data were repeated from 1948 to 2012 a) Change in biomass from the base run where the full oceanographic data were used. b) Correlation between the biomass from the model with the full oceanographic data and from the model with five years repeated. The correlation measure the change of the trend of the biomass of the functional groups.

model components.

This model can be used for examining scenarios regarding discards and by-catch. It has already been used to explore the economic and ecological effects of discarding in the cod and haddock fisheries where five scenarios with different selectivity and discard rates were compared (Sturludottir, 2018). The small scale lump sucker fishery, which is an important part of a local economy, has been under scrutiny for considerable by-catch of marine mammals and seabirds (Anon., 2018; Pálsson et al., 2015) and the model could be used to evaluate trade-offs of alternative management actions, such as area closures and/or effort reduction, which could have substantial consequences for the coastal communities.

The spatial component of the Atlantis model allows diverse scenarios to be explored. For example the effects of changes in spatial movement of species because of climate change can be explored, but such changes have already been observed in Icelandic waters (Carscadden et al., 2013). The effects of seasonal or permanent closures already implemented in Icelandic waters have been studied (Schopka, 2007; Woods et al., 2017), but the effects of potential closures could be explored with the current model.

The model is being used as an operating model to evaluate the

performance of other simpler ecosystem models such as EwE and Gadget. It is very difficult to evaluate the reliability of ecosystem models, but skill assessment has been used in that purpose. Atlantis models can be used to test if other models can imitate the “Atlantis ecosystem”. Data are simulated from the known ecosystem and imported into the other models and then compared to the true values from the Atlantis model. Using this method it can be studied how well the models need to fit to historical data to be deemed reliable when used for forecasting. This work is now ongoing using the present model.

#### 4. Conclusion

An end-to-end model has been constructed that resembles the ecosystem of the Icelandic waters. This is the first dynamic end-to-end model for this area. Preliminary EwE models (Buchary, 2001; Mendy, 1998) and multi-species models (Elvarsson, 2015) have been constructed but no dynamic models where the entire ecosystem is simulated. Fisheries management in Iceland aims to be ecosystem based and this model can be used to support EBFM. It can be used to evaluate different fisheries and climate scenarios and to conduct management strategy evaluations. The focus of the model is on the most important

commercial groups, to explore the effects of various harvesting strategies, such as area closures and selectivity changes, on both commercial groups and on other parts of the ecosystem. The model needs to be judged to be reliable to have confidence in the model as a basis for assessments and projections. It is very difficult to estimate the reliability of a model as complex as Atlantis. Nevertheless, it was possible to carry out a skill assessment, which showed that the model was able to simulate the biomass trends for the most important commercial groups. There were limited data available on the non-commercial groups and therefore the reliability of the model for those groups could not be assessed. Therefore, predicted effects on those groups need to be taken with caution.

The sensitivity analysis revealed the influence of key parameters and inputs, e.g., recruitment parameters and oceanographic data, on model projections. This showed that there was uncertainty due to sensitivity to the form of the recruitment relationships used and due to the effect of environmental conditions (something the model may not capture well). Uncertainty also exists due to sensitivity of the model to oceanographic forcing, meaning that care must be taken around assumptions regarding prevailing oceanographic conditions used when running simulations, such as warm vs. cold periods.

Ecosystem models such as Atlantis are in a constant process of further improvement and this is the case with the current model. This work, the skill assessment and the sensitivity study will facilitate that process by increasing the understanding of the dynamics of the system.

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