# Numerical Models for Monitoring and Forecasting Ocean Ecosystems: a short description of present status

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Abstract. Understanding and managing marine ecosystems under potential stress from human activities or climate change
 requires the development of models with different degree of sophistication in order to be capable of predicting changes in
 living components and in relation to human pressures and environmental variables. Recent advances in ecosystem modelling
 are the focus of this paper, which reviews numerical approaches to analyse the characteristics of marine conditions in terms of
 typical units, i.e., individuals, populations, communities and ecosystems. In particular, lit specifically examines the current
 classification of numerical models of increasing complexity – from individuals and population and stock assessment models
 to models representing the whole ecosystem by covering all trophic levels – and presents examples and their operational
 maturity and readiness, finally demonstrating their use for supporting marine resource management, conservation, planning
 and mitigation actions.

#### **1** Introduction

In recent decades, a Understanding and managing marine ecosystems under potential stress from human activities and climate
 change requires the development of modelling tools able to monitor and forecast ocean ecosystem dynamics, from physics to fish (De Young et al., 2004). The challenge is to relate processes occurring at individual, population or community levelies with environmental variables, i.e., to connect the dynamics of marine ecosystem with the quite well established physical and biogeochemical products that exists for the ocean (Fennel et al., 2022). A large variety of numerical ecosystem models have been developed to predict the growth and dynamics of individuals and populations of marine resources. According to the scope

25 the approaches are very diverse ranging from single- to multi-species and might include the effects of various environmental changes and human impacts on marine biological resources (Hollowed et al., 2013; Nielsen et al., 2018). To illustrate In order to shed light on approaches that have the potential to become the next generation operational tools for ocean ecosystem forecast, this paper provides a structured synthesis of models applied to marine higher trophic levels (i.e., from zooplankton to fish and top predators) that can be connected with lower trophic level models (physics and physics).

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30 biogeochemistry).

A comprehensive analysis is challenging, <u>although but</u> models can be mapped in terms of their main scope and distinguishing approaches that can incorporatinge age structure, environmental factors, represent trophic interactions, and spatial structure (Hollowed et al., 2000; <u>Plaganyi, 2007</u>). Based on the above characteristics, numerical models <u>for marine ecosystems</u> can be divided into six broad classes:

Bioenergetic models representing the processes related to growth, respiration, excretion of an individual;

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- population and stock assessment-fisheries models (typically for single species without trophic interactions and possibly age-structured);
  - connectivity models (considering propagules dispersal, larval cycle, spatial structures, and environmental factors);
  - species distribution models (statistical models based on representation of spatial environmental variables and biota);
- minimal realistic models (typically age-structured, with representing a few species -with trophic interactions);
- whole ecosystem models (typically covering all trophic levels and based on trophic interactions, which may include size structure and spatial variation).

These <u>six five</u> classes of models are reviewed in the following sections <u>below</u>, considering <u>also</u> available syntheses and reviews (e.g., Plaganyi, 2007; <u>Cowen et al., 2009</u>; Stock et al., 2011; <u>Hilborn and Walters, 2013</u>; Itoh et al., 2018; <u>Nielsen et al., 2018</u>;

- 45 Rose et al., 2024).-The work does not pretend to be exhaustive and readers are referred to original reviews that are providing in depth analyses of each class of models. It aims to provide a synthetic integration across different classes, with examples provided to illustrate their application in operational coupling with lower trophic level models. The work intends to synthetically bridge across classes and examples are provided to shed light on their usage for operational coupling with lower trophic level models. For this purpose, readiness and maturity of each model was subjectively elaborated on the basis ofbased
- 50 on its current application. The mMaturity of each example werewas assessed on the basis of based on the availability of code, documentation, test cases, routines for assessing model performances, diagnostics, and isare used by a community of developers that can provide support, updates and advancement. Stock assessment models routinely applied for fisheries management, for example, were considered more mature because the code is publicly available and documented and input and output test cases are developed and accessible. Readiness for operational purposes was defined on the basis of based on existing
- 55 knowledge abouton possible connection of the model example to physical and biogeochemical spatio-temporal models. Existence of such applications, even if scarce, might shed light on theshow the difficulties in connecting (one-way or twoway) with low trophic level models. Operational readiness may be regarded as more tentative and less precise, owing to the challenges in establishing a clearly objective definition, particularly in light of its potentially limited application. Readiness for operational purposes might be considered more tentative and less precise, because it is more difficult to define in a very 60 objective way also for the potentially very sparse application.

For each class of models some examples are reported shown in Table 1., includingand their characteristics in terms of typical units, elemental structure, number of species typically represented, representation of eventual trophic interactions are reported in Table 1. The table 1 also contains synthetic information on primary model focus and main output, as well as if each model is maturity and readiness for operational or not purposes.

#### 65 2 Bioenergetic models

Traditional bioenergetic models describe energy intake from feeding and its allocation to maintenance, activity, growth, reproduction, and excretion (for a review see: Rose et al., 2024), A bioenergetic model is any mechanicistic model describing how individuals take energy from the environment and allocate it to different processes (Kooijman, 2010; Sibly et al., 2013). Bioenergetic models are typically used for representing the growth of the individual, while accounting for respirations,

70 exerctions and other losses. Energy intake can be a model input or output, depending on whether it's modeled dynamically or derived from energy requirements (Pirotta et al., 2022). Acquisition and allocation can vary based on the individual's state and environmental conditions (e.g., Libralato and Solidoro, 2009; Nisbet et al., 2012)

Indeed, bioenergetic models and can account of for external oceanographic conditions influencing uptakes, such as light, nutrients and temperature for autotrophs (e.g., Bocci et al., 1997) or food availability and temperature for heterotrophs (e.g.,

- 75 Libralato and Solidoro, 2009), while losses are usually related to temperature and internal conditions (Koojiman, 2010). Bioenergetic models can also consider explicitly the gonadic development and egg release (Pastres et al., 2000). Because of these characteristics, bioenergetic models, other than providing realistic individual-level response to environmental conditions, permits to project responses at the population and food web levels and can support other classes of approaches (Rose et al., 2024).
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Traditional bioenergetic models describe energy intake from feeding and its allocation to maintenance, activity, growth, reproduction, and exerction. These models are advantageous due to their clear empirical interpretation and measurable units but tend to be parameter heavy and difficult to generalize across species. A widely used bioenergetic approach for fish and invertebrates is represented by the Dynamic Energy Budget (DEB, Koojiman, 2010) which is characterized by an explicit

- 85 representation of <u>energy dynamics into</u> somatic, gonadic and storage tissues <u>(Koojiman, 2010)</u>. Dynamic Energy Budget (DEB) theory offers a more general approach by using mass-energy balance principles to link sub-organismal processes with overall organismal performance (Koojiman, 2010; Nisbet et al., 2012). However, this generality leads to abstract concepts that are more challenging to measure empirically (Pirotta et al., 2022). <u>Although</u>, Tthe presence of the storage in DEBis challenging to be measured empirically (Pirotta et al., 2022), it allows <u>-allow</u>-representation of delayed use of energy in the individual
- 90 development resulting in improved generality of the approach (Koojiman, 2010; Nisbet et al., 2012). Thus DEB has been developed into a theory for scaling the parameters for all life cycles of the individual (from eggs to larvae to juveniles and adults)—and, provides setting parameters for a large number of marine species (see also https://www.bio.vu.nl/thb/deb/deblab/add\_my\_pet/) and is well documented (Nisbet et al., 2012; Koojiman, 2020). Thus DEB is considered of high maturity for being used routinely and adapted to operational applications, and because it is seldom
   95 connected to spatiotemporal physical and biogeochemical models the readiness is considered of intermediate level (Table 1).5
- thus it has a maturity for being used routinely and adapted to operational applications.



#### 3 Population and stock assessment fisheries models

Various types of numerical models of single populations are used worldwide to support <u>fisheries</u> management by determining
 the population at sea and the current status of exploited marine populations, thus providing insight for management <u>in a process</u> <u>called stock assessment (for a review see Hilborn and Walters, 2013)</u>. Such sStock assessment models typically represent the biomass or abundance of <u>one individual species (Table 1)</u>, are routinely used by management agencies, and include probability models to incorporate various sources of observational data (Maunder and Punt, 2013). In cases where stock assessments are based on limited observations, i.e., no catch by age or size, surplus production models are used. The general underlying assumption of these models is a theta-logistic function for the evolution of biomass (B; but it can also be applied to the number of individuals) over time:

$$\frac{dB}{dt} = \frac{r}{n-1} B_{\varepsilon} \left( 1 - \left[ \frac{B_{\varepsilon}}{K} \right]^{\theta-1} \right) - F_{\varepsilon} B_{\varepsilon} \quad (Eq.1)$$

where F is the fishing mortality, so F<sub>t</sub>B<sub>t</sub> corresponds to the catches Ct, while r is the population growth rate, and K is the capabilities of the system to support the population (through living space, habitat, food, etc.), and generally called as carrying capacity. The parameter O allows generalization of the equation (in the case of O=2, the classical logistic curve is obtained). Several packages use the surplus production model as principal approach, are used routinely to perform stock assessment and are including several useful diagnostics. Among them the most used are the CMSY (Froese et al., 2023) and the SPiCT (Pedersen and Berg, 2017) models. CMSY uses a time series of catch data and fishing effort to estimate parameters, reconstruct
biomass and establish reference points using a Bayesian approach (Froese et al., 2023). The Stochastic surplus Production

- model in Continuous Time (SPiCT)<u>, for example</u>, provides estimates of exploitable biomass and fishing mortality at any point in time from <u>catch and survey</u> data collected at arbitrary and possibly irregular intervals (Pedersen and Berg, 2017). <del>The model</del> allows the inclusion of prior distributions for parameters that are difficult to estimate such as growth rate and carrying capacity. SPiCT is available as an R package (R Core Team 2015) in the online GitHub repository: <u>https://github.com/mawp/spict</u>.
- 120 Surplus production models are simplistic representation of the population that is lumped with no size and/or age details. More sophisticated approaches (such as SS3, a4a, XSA, etc.) are used when use catch data by age or size classes are available for the exploited population (catch-at-age or catch-at-length models; Maunder and Punt, 2013). These stock assessment models reconstruct the number of individuals in to reconstruct the cohorts based on catch and assuming natural mortality for each class by age class, as well as and considering information on species growth, fecundity, and fisheries selectivity (Methot and Wetzel, 2013). The basic dynamics are described by the number of individuals N at time t and age a, as in the following:

$$N = R_{\varepsilon} + \left(N_{\varepsilon-1,\alpha=1}e^{-\frac{M}{2}} - C_{\varepsilon-1,\alpha=1}\right)e^{-\frac{M}{2}} + \left(N_{\varepsilon-1,\alpha=2}e^{-\frac{M}{2}} - C_{\varepsilon-1,\alpha=2}\right)e^{-\frac{M}{2}} + \cdots \left(N_{\varepsilon-1,\chi-1}e^{-\frac{M}{2}} - C_{\varepsilon-1,\chi-1}\right)e^{-\frac{M}{2}} + \left(N_{\varepsilon-1,\chi}e^{-\frac{M}{2}} - C_{\varepsilon-1,\chi}\right)e^{-\frac{M}{2}} - C_{\varepsilon-1,\chi}e^{-\frac{M}{2}} - C_{\varepsilon-1,\chi-1}e^{-\frac{M}{2}} - C_{\varepsilon-1,\chi-1}e^{-\frac{M$$

where each year the population comprises all age classes from the new juvenile individuals entering the population as recruits
 the same year ( R, age 0 ), all age classes a, from 1 up to the oldest age modelled (age x) surviving from the year before. The number of individuals are decreasing through time on the basis of catches C at age and time, and assuming instantaneous natural mortality M.

Typically, these models report juvenile recruitment R as a function of a combination of fecundity by age class estimated from data (Stock et al., 2011). This class of models includes, for example, the a4a tool (assessment for all, Jadim et al., 2014), a modeling framework for fitting statistical age-structured fishery models using nonlinear statistical submodels. The submodels

- can include linear functions of age and year, smoothing splines with fixed degrees of freedom that vary with age and/or year and environmental indicators as covariates. The tool a4a is implemented in R Fishery Library and includes the optimization procedure. Stock synthesis (SS3; Anderson et al., 2014) is the most an example widely used of catch-at-age stock assessment model that can incorporate age or length composition information from surveys, abundance indices, multi-gear effort,
- 140 selectivity, and spatial data in the most recent and advanced applications (e.g., Punt, 2019; Privitera-Johnson et al., 2022). In all eases, pProjections of from stock assessment models are generally made for annual to decadal time periods and .-Catch or effort limitation scenarios can be used to SS3 provides estimates for biological reference points for management decisions (indicators based on maximum sustainable yield; Hilborn and Walters, 2013). As for many stock assessment fisheries models, Although in most cases, SS3 stock assessment models are is routinely used in formal assessments, well documented and easily
- 145 accessible (https://github.com/nmfs-ost/ss3-source-code ), thus it has a very high degree of maturity. -Nevertheless, it is not spatially explicit and <u>it\_does</u> not consider explicitly oceanographic forcings they are routinely used in formal assessments for management and might be considered <u>of intermediate readiness</u> as ready for operational oceanographic applications (<u>Table 1</u>).

#### 4 Connectivity models

The distribution and survival of small eggs and larvae of marine fishes and invertebrates, as well as propagules of algae and seagrass' seeds are advected and thus are strongly influenced by currents, which can disperse individuals both near spawning sites and in distant areas (Cowen et al., 2007). Therefore, biophysical dispersal (advection, diffusion, and migratory behaviorbehaviour of organisms) is fundamental to explaining marine population dynamics and connectivity (for a review see Cowen et al., 2009). NumericalConnectivity models are used to quantitatively integrate the large spatial and temporal variability of oceanographic processes (physical connectivity) with processes inherent in the biology of marine organisms (life history traits) to investigate connectivity between and within populations and alsoand across larval stages (Gawarkiewicz et al., 2007; Melaku Canu et al., 2021). Connectivity models <u>such as Larval TRANSport Lagrangian model (LTRANS, North et al., 2008)</u> typically uses offline physical parameters (velocity, density, temperature, and salinity) obtained from hydrodynamic models and estimate the distribution of organisms. <u>The advection-diffusion-reaction equation is typically used for biomass</u> distribution (e.g., Sibert et al., 1999; Faugeras and Maury, 2005), while Lagrangian approaches are used to track particles and

<sup>160</sup> thus distribute individuals (e.g., Laurent et al., 2020). These approaches take into accountconsider life history traits such as



growth, mortality and the behavior of target organisms in terms of seasonal variability, spawning sites, vertical movement and settlement preferences (Melaku Canu et al., 2021; Paris et al., 2013; Lett et al., 2008). :- since in most of the cases living organisms have negligible influences on physical oceanographic processes parameters, modeling the biophysical LTRANS is frequently applied and well documented, applied and the code is available at (https://github.com/LTRANS/LTRANS/L2b)
makingdesignating it asof intermediate level of maturity. It is coupled dispersion offline from thewith hydrodynamic models is considered a reliable strategy also considering time evolutions, and can include incorporate several biological features (North et al., 2008) that makesplacing its operational readiness for operational purposes of at an intermediate level (Table 1). The advection diffusion reaction equation is typically used for biomass distribution (e.g., Sibert et al., 1999; Faugeras and Maury, 2005), while Lagrangian approaches are used to track particles and thus distribute individuals (e.g., Laurent et al., 2020). These approaches take into account life history traits such as growth, mortality and the behavior of target organisms in terms of seasonal variability, spawning sites, vertical movement and settlement preferences [e.g., Melaku Canu et al., 2021; Paris et al., 2013; Lett et al., 2008]. Connected with oceanographic variables and spatially explicit these models however, appear less mature for operational applications.

### **5** Species distribution models

- 175 Species distribution models (SDM, also called habitat suitability models) are statistical models that predict the occurrence, abundance, or biomass of organisms using geoposition, biotic and environmental data (<u>for a review see: Elith and Leathwick</u>, <u>2009Brodie et al., 2020</u>). Particularly useful when applied to <u>standardized</u>, <u>spatio-temporal</u> scientific <u>monitoring and</u> surveys of <u>-biotie dataspecies abundance</u>, these approaches can also exploit <u>publicly available opportunistic biological</u> datasets (e.g., <u>OBIS, www.obis.org</u>; <u>GBIF, www.gbif.org</u>). SDMs are implemented using various <u>statistical approaches</u> approaches.
- 180 including linear models (LM), generalized linear models (GLM), generalized additive models (GAM) (Melo-Merino et al., 2020; Maravelias et al., 2003; Melo-Merino et al., 2020; Brodie et al., 2020), machine learning,- artificial neural networks methods such as random forest (RF, Breiman et al., 2018) or artificial neural networks (ANN),- (Catucci et al., 2025) and maximum entropy (Jones et al., 2012; Pittman and Brown, 2011; Reiss et al., 2011). The inclusion of physical and biogeochemical oceanographic covariates, which can have direct and indirect effects on species distributions, can improve the
- capabilities of SDMs to explain observed biotic data compared to using only geopositional variables (Panzeri et al., 2021; Thorson et al., 2015). Recent advances include combining the approaches into an ensemble (Jones et al., 2012; Panzeri et al., 2024) and including multiple species as covariates into the so called Joint-Species Distribution Models (JSDM, Pollock et al., 2015; Thorson et al., 2016). These classes of SDMs are increasingly being used to describe current and future distributions of exploited and endangered species, identify hotspots, map essential fish habitat, support conservation development, and feed
  other ecosystem models (Jones et al., 2012; Colloca et al., -2015; Grüss et al., 2014; Dolder et al., 2018).

The Dynamic Bioclimate Envelope Model (DBEM) estimates species distributions based on environmental preferences and considers population dynamics and dispersal (Cheung et al., 2009). The DBEM makes predictions of future envelopes using

physical and biogeochemical data from oceanographic models and <u>also</u> considers the response of organisms to natural/anthropogenic environmental changes such as growth, mortality, larval dispersal, and migration (Cheung et al., 2013).
In general SDMs are widely applied, <u>well documented and available (see for example: https://github.com/helixen/sdm\_r\_packages)</u> thus have an intermediate level of maturity but giving their direct integration with physical-biogeochemical models they have a good readiness level for operational use (Table 1)and although at the moment they are not used operationally, they can be easily implemented within an operational chain.

#### 6 Minimal realistic models

- 200 Dynamic multispecies models or Minimal Realistic Models (MRM), Punt and Butterworth, 1995) are approaches that represent a limited number of species (usually less than 10 species) that have important interactions with a target species (for a review see Plaganyi, 2007). The MRMs often represent an evolution of single species stock assessment models: for example, Multispecies Virtual Population Analysis (MSVPA) is an extension of virtual population analysis (Gislason, 1999), while GADGET (Globally applicable Area-Disaggregated General Ecosystem Toolbox) is an extension of stock synthesis in the
- 205 multispecies framework, where populations can be partitioned by species, size classes, age groups, areas, and time steps (e.g., Andonegi et al., 2011). In particular, GADGET is flexible, allowing easy addition/replacement of alternative model components for biological processes such as growth, maturation, and predator-prey interactions representing some species in age classes. GADGET provides estimates of population dynamics under technical fisheries and biological interactions with the ability to use different growth functions and fitness functions (Plaganyi, 2007). <u>Although well documented (see</u>
- 210 https://gadget-framework.github.io/gadget2/userguide/) its fitting is quite complex and thus hasve few applications: for these reasons maturity is considered intermediate and readiness for operational purposes is low because of lack of interactions with physical and biogeochemical models (Table 1).

MICE ("Models of intermediate complexity for ecosystem assessment"; Plagányi et al., 2014) are developed considering the specific problem and data availability. MICE represents a limited number of populations (usually 10) exposed to fisheries or

- 215 anthropogenic interactions and includes ecological processes (Angelini et al., 2016). These models have different levels of detail for the species represented: MICE can simultaneously represent focal populations in age-structured classes, while others take a surplus production approach (Morello et al., 2014). MICE can be a fairly complex but flexible tool that overcomes the many complexities of whole ecosystem models and is useful for providing tactical advice for focal species management (Plagányi et al., 2014). An example of minimum realistic model is the Spatial Environmental POpulation Dynamics Model
- 220 (SEAPODYM), which is a two-dimensional coupled physical-biological model originally developed for tropical tunas in the Pacific (Lehodey et al., 2003). SEAPODYM includes an age-structured population model for tuna species top predators and a movement model based on a diffusion-advection equation such that swimming behavior is modeledmodelled as a function of habitat quality (sea surface temperature (SST), ocean currents, and primary production) predicted obtained from oceanographic models and satellites (Lehodey et al., 2015). This model describes spatial structures that are essential to account for the

225 distribution of fishing effort, swimming behavior, and environmental variations typically determined by ocean circulation models or derived from satellites (Lehodey et al., 2015; Senina et al., 2020). <u>SEAPODYM is well documented and already</u> used for operational global projections (https://github.com/PacificCommunity/seapodym-codebase) thus can be considered to have a high degree of maturity and readiness for operational purposes (Table 1).

#### 7 Whole ecosystem models

230 Whole ecosystem models (WEM) are designed to represent all trophic levels in an ecosystem, from primary producers to top predators and take advantage of .- Thus, WEMs typically use a very large set of data collected infrom different disciplines a variety of disciplines, including results from oceanographic models and stock assessments (e.g., Agnetta et al., 2022).

The main distinguishing feature between the different WEM is the way in which the ecosystem is described: i) <u>through</u> <u>compartments representing in flexible compartments representing species of groups of species (Christensen and Walters, 2004;</u>

- 235 <u>Fulton et al., 2011</u>), ecologically meaningful groups of species, or size- and age-structured populations, such as Ecopath with Ecosim (hereafter EwE, Christensen and Walters, 2004) and ATLANTIS (Fulton et al., 2005); ii) <u>through compartments that</u> <u>represent in</u>-size-structured communities, typically benthic and pelagic communities (<u>Shin and Cury, 2004</u>; <u>Travers et al., 2010</u>), such as Osmose (Shin and Cury, 2001), Feisty (Petrik et al., 2019), and DBEM (Blanchard et al., 2009), for example; iii) in a mixture of size-structured <u>and trophic</u> communities (typically pelagic, mesopelagic migratory, and non-migratory) and
- 240 age structured species as in Apecosm (Maury, 2010); iv) using the ecosystem is described by dynamic spectra of trophic levels (e.g., Gashe and Gascuel, 2013) as in Ecotroph (Gasche and Gascuel, 2013). All these models are based on biomass and consider rules such as biomass conservation (Table 1; for a review see Plaganyi, 2007).

Ecopath with Ecosim (EwE; Christensen and Walters, 2004), undoubtedly is the most widely used WEM, it is a freely available (www.ecopath.org) and has a flexible structure, general, Hit representsis a suite of models developed in more than 30 years

- 245 for the whole ecosystem descriptions that have been developed over 35 years (Christensen and Walters, 2004) and <u>. EwE</u> has been used to analyze past and future impacts of fisheries, nutrient inputs, invasive species, and climate <u>change</u> (e.g., Heymans et al., 2014; Libralato et al., 2015; Serpetti et al., 2017; Piroddi et al., 2021). It consists of three different interconnected main modules, i) a static mass-balanced ecosystem network (Ecopath, Christensen and Pauly, 1992), ii) a temporally dynamic simulation module (Ecosim, Walters et al., 2000), and iii) a spatially and temporally dynamic module (Ecospace, Walters et al., 2000).
- 250 al., 1999). EwE contains a large number ofmany additional modules for calibration, uncertainty analysis, calculation of indicators, and simulation of pollutant dynamics (Steenbeek et al., 2016). Recent advances allow the direct embedding of two-dimensional monthly results from oceanographic physical-biogeochemical models (Steenbeek et al., 2013). EwE can be considered an approach of high maturity and intermediate degree of readiness for operational applications (Table 1). ATLANTIS spatially resolves the full trophic spectrum of ecosystem types, including age structured formulations for high
- 255 trophic levels, potentially in multiple vertical layers (Fulton et al., 2011). ATLANTIS includes a nutrient pool formulation that



can be used to test effects such as nutrient inputs (Audzijonyte et al., 2019). The model has been used for site-specific analyses and to examine general aspects of fishery's impacts on fish communities (Link et al., 2010).

OSMOSE (Objected oriented Simulator of Marine ecoSystems Exploitation; Shin and Cury, 2004) is an individual-based ecosystem model that simulates size-based communities on a 2-D spatial cell grid and can be coupled with a planktonic
 ecosystem model (Travers et al., 2010). The model has been used to study the effects of various aspects of fisheries on the food web (e.g., Shin and Cury, 20019.

<u>A large set of These models WEM models (Table 1)</u> are <u>used</u> increasingly being used to address the need for holistic ecosystem approaches, and their framework is often <u>used\_applied</u> to answer strategic medium-term questions related to management strategies, fisheries issues, and climate or <u>environmeentalenvironmental</u> change (e.g., Tittensor et al., 2021). Notably, WEM

265 can be coupled with other classes of models (population dynamic, SDM, connectivity models), as well <u>as with biogeochemical models, which is why therefore most of the approaches in this class have a high to intermediate level of maturity and readiness (Table 1).</u>

#### Conclusions

- 270 A large setwide range of models are usedexist that were developed tofor representing ocean ecosystems at different level of organisations, fromincluding individuals, populations, communities and whole entire ecosystems. Although dividedcategorised into in 6 classes for the sake of clarity, some modelling approaches are not esclusiveconfined to a single of a class. For instance, the DEB modelling approach is also-used also-to also represent the growth of individuals in Connectivity models and MRM classes (see for example Maury 2010). Conversely MICE ("Models of intermediate complexity for ecosystem assessment"; Plagányi et al., 2014) of the MRM class, are developed using different levels of detail for the species
- represented by combining for example age-structured and surplus production approaches (Morello et al., 2014). These models have been developed for specific objectives that embrace many issues important for society societal issues, i.e., from effects of climate change, pollution, nutrient enrichment and, fisheries-ete.
- The numerical approaches analysed here have characteristic spatio-temporal resolutions (Table 1) that are generally decreasing when moving from individual species models to whole ecosystem models (Table 1). Moreover, while increased representeding complexity with MRM and WEM classes result in there is a general improvement of realism but also lower at the cost of accuracy (generally declining from individual models to WEM)ies. Overall, the first set of approaches (bioenergetic and population models) are more adapted for tactical analyses while especially the WEM are at the moment currently considered useful especially in strategic analyses (see Table 1). Although very few of the reviewed approaches are currently
- 285 used-in operationally (i.e., SEAPODYM), many -approaches are routinely usedapplied for supporting management (e.g., fisheries stock assessment models). -Most of the approaches reviewed have repository for documentation, code and testing cases, thus have high degree of maturity (Table 1). Conversely, approaches under the MRM class are not widely applied, are

often quite complex to fit and therefore were considered-were categorised to be at aof poor level of readiness for operational purposes (Table 1). Nevertheless, all the tools have some degree of coupling (mainly off-line) with physical and biogeochemical variables, thus have athey have great potentials for becoming operational, by and to be used for analysinge ecosystem dynamics and make useful scenarios, which can be useful for, on a very wide range of issues and management actions that might-could be eventually prioritized.

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#### 540 Competing interests

The contact author has declared that none of the authors has any competing interests.

# Data and/or code availability

Not applicable.

### Authors contribution

545 Not Applicable



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#### **Bioenergetic models Readiness for** Elemental Time Spatial Trophic Primary model Physical/biogeochem Number **Repository** <u>ModelName</u> **Model Units Model**Acronym **Maturity** <u>o</u>Operational units if open access) structure structure of species interactions focus, output ical processes use Yes used as forcings ttps://www.bio.v PotentiallyGo Individual weight (gww, DEB Dynamic Energy Budget Individual day No No growth High (temperature, light, 1.nl/thb/deb/debla 1 gC or others) or length <u>od</u> b/add\_my\_pet/ food, nutrients) Population and stock assessment fisheries models Number **Physical/biogeochem Trophic** Primary model **Readiness for** <u>Time</u> **Elemental Spatial** <u>of</u> ical **Repository** <u>nteractions</u>T focus, <u>operational</u> <u>Acronym</u>Model <u>unitsTim</u> processesPhysical/bi ModelName structure Eleme Model Units Model Units structureSpa <u>species</u>Nu **Maturity** rophic outputPrimary <u>use</u>Operation if open access) ogeochemical ntal structure e units tial structure mber of interactions model focus, output al species <del>processes</del> Sstochastic surplus https://github.co Biological reference Surplus SPICTSpict pProduction model Iin Biomass Year No No <u>High</u> Pooryes No 1 prodution points for fisheriesm/mawp/spict. Ceontinuous Ttime **Biological reference** ttps://github.com/ SISTA16/cmsy Catches at Maximum Surplus points for CMSY (tonn) Year No 1 No Intermediate <u>YesPoor</u> No prodution fisheriesBiological sustainable Yield reference points **Biological reference** https://github.com/ **Yes**Intermedi points for A4a All for all Catch-at-age Year No No Biomass (tonn) No 1 Intermediate fisheries Biological <u>a4a</u> ate reference points **Biological reference** tps://github.com Number of Individuals; Potentially points for Intermediate¥ SS3 Catch-at-age Year No Potentially yes Stock Synthesis <u>High</u> 1 nmfs-ost/ss3biomass (ton) fisheriesBiological yes es source-code reference points https://noaa-Biological reference fisheries-Number of Individuals; points for VPA Virtual population analysis Catch-at-age Year No No no Poor Poorno integrated-1 fisheries Biological biomass (ton) lbox.github.i reference points <u>VPA</u> **Connectivity models** Readiness for Physical/biogeochem Elemental Time Spatial Number Trophic Primary model <u>operational</u> **Repository** <u>Acronym</u>Model <u>ModelName</u> **Model Units Maturity** structure units structure of species interactions focus, output use Operation ical processes (if open access) al Distribution of Typically ttps://github.com species and Agents (super individuals) Intermediate<sub>N</sub> Yes (physical LTRANS Lagrangian Transport Number of individuals days Yes No Intermediate TRANS/LTRAN North et al one connectivity among processes) θ species Sv.2b sites Study effects of Typically Individuals physical and Lagrangian tool for Yes (physical ttps://ichthyop.or Ichthyop (early life Number of individuals Intermediate Intermediate simulating ichthyoplankton <u>days</u> Yes No biological factors on one processes) <u>g/</u> ichthyoplankton dynamics stages) species

# Table 1: Main characteristics of some widely used numerical models for of marine biological resources divided into the six classes

dynamics

Doso at al 2024		4		
Rose et al., 2024		1	Tobolla formattata	
	-			
Reference	<b>.</b>		Eormattato, Allinoato a cinistra	
<u>Koojiman, 2020</u>	<b>*</b>		Formattato: Allineato a sinistra	
Hilborn and Walters,	<b>20</b> 1	3-	Formattato: Allineato a sinistra	
			Formattato: Allineato a sinistra	
Pedersen and Berg, 2017	<b>*</b>		Formattato: Allineato a sinistra	
	_			
Froese et al., 2023	<b>*</b>		Formattato: Allineato a sinistra	
Jardim et al., 2014	<b>.</b>		Formattato: Allineato a sinistra	
			Tormatator Annicato a Sinistra	
. 1				
Anderson et al., 2014	•		Formattato: Allineato a sinistra	
Gislason, 1999	<b>.</b>		Formattato: Allineato a sinistra	
<u>Cowen et al., 2009</u>	*		Formattato: Allineato a sinistra	
	<b>*</b>		Formattato: Allineato a sinistra	
North et al., 2008	<b>.</b>		Formattato: Allineato a sinistra	
Lett et al., 2008				

IBM/ABM	Individual-based and Agent Based Models	Individual	Biomass	days	Yes	Typically a few species	Efficient predator	Ecosystem effects on target population and connectivity	Poor	<u>PoorNo</u> (computationa lly complex)	Yes	NA	Rose et al., 2015	<b>*</b>	- Formattato: Allineato a sinistra	
	Species Distribution Models												Elith and Leathwick.	2009	Formattato: Allineato a sinistra	
<u>Acronym</u> Model	<u>Model</u> Name	Elemental structure	Model Units	Time units	Spatial structure	Number of species	<b>Trophic</b> interactions	Primary model focus, output	<u>Maturity</u>	Readiness for operational useOperation	Physical/biogeochem ical processes	<u>Repository</u> (if open access)		<b>*</b>	Formattato: Allineato a sinistra	
										al	•					
Ens <u>e</u> amble of SDM	Ensamble of Species Distribution models	Species adundance, presence or	Number of individuals or weight per unit surface or presence/absence	Month, year, climatolo	Yes	1	No	Species distribution; essential fish habitats	Good	<u>GoodPotential</u> <del>ly</del>	Environmental factors can be included	https://github.com/ helixcn/sdm_r_pac kages	Panzeri et al., 2024	*	ha formattato: Italiano (Italia)	
		Species		gy						Poor		https://github.com/		Ň	Formattato: Italiano (Italia)	
Joint-SDM	Joint Species Distribution models	adundance, presence or biomass	Number of individuals or weight per unit surface	Month, year	Yes	A few species	implicit	Distribution of target species	Intermediate	(computationa <u>lly</u> intensive)No	Environmental factors can be included	<u>James-</u> <u>Thorson/spatial_D</u> <u>FA</u>	<u>Thorson et al., 2016</u>	<b>*</b>	Formattato: Allineato a sinistra	
DEBM	Dynamic Bioclimate Envelope Model	Species biomass	biomass	year	Yes	Several species	No	Distribution of multiple species	Intermediate	<u>Good</u> No	Yes included for developing the bicenvelope	NA	Cheung et al., 2013	<b>*</b>	- Formattato: Allineato a sinistra	
	bioenvelope													_		
					Minimal	Realistic m	odels						<u>Plagányi 2007</u>	***	ha formattato: Tipo di carattere: 11 pt, carattere: Nero	Grassetto, Colore
										Readiness for					Formattato: Allineato a sinistra	
<u>Acronym</u> Model	<u>Model</u> Name	Elemental structure	Model Units	Time units	Spatial structure	Number of species	<b>Trophic</b> interactions	Primary model focus, output	<u>Maturity</u>	operational useOperation	Physical/biogeochem ical processes	<u>Repository</u> (if open access)		<b>*</b>	Formattato: Allineato a sinistra	
CADCET	Globally applicable Area Disaggregated General	Population in	Biomass derived from	Voor	Yes, can be	Typically	Yes, suitability-	Ecosystem effects on	Intermediate	notontiallylayy	Can be coupled with physical-	https://gadget- framework.github.	Andonazi et el 2011			
GADGET	(derived from BORMICON)	age structure	population size structure	I cai	included	species	based, flexible	yearly biomass	mtermediate	potentiany <u>iow</u>	biogeochemical model	io/gadget2/usergui <u>de/</u>	Andonegi et al., 2011		Formattato: Allineato a sinistra	
MSVDA and	Multi-species Virtual	Populations in				Typically	Yes; Suitability-	Ecosystem effects on		<u>Poor</u> No		https://noaa- fisheries-				
MSFOR	multi-species Forecasting	age structure	Numbers at age; Biomass	Year	No	3-4 species	based; Efficient	target population; yearly biomass	<u>Poor</u>	(seldom applied)	Not usually included	integrated- toolbox.github.io/	<u>Gislason, 1999</u>	<b>+</b>	<b>Formattato:</b> Allineato a sinistra	Coloro carattoro: Noro
	Widder						predator					MSVPA_X2				
MICE	Model of Intermediate Complexity for Ecosystem assessments	Populations in surplus production and age structure	Numbers at age, Biomass	Year	No	Typically 6-7 species	Efficient predator	Dynamics of focal species and their predators or preys	<u>Difficult to</u> establish: programmed on purpose	Poor (only <u>few</u> <u>applications)</u> P otentially	Environmental effects can be included	<u>NA</u>	<u>Plagányi et al., 2014</u>	<b>*</b>	- Formattato: Allineato a sinistra	
						т : н				HighYes	Can be coupled with	https://github.com/		<b>*</b>	- Formattato: Giustificato	
SEAPODYM	Spatial Ecosystem, and population Dynamics Model	Populations in age structure	Biomass	Year	Yes	3-4	Efficient predator	Ecosystem effects on target population	High	( <u>already</u> <u>applied for top</u> predators i e	physical- biogeochemical	PacificCommunity /seapodym-	<u>Lehodey et al., 2015</u>	<b>-</b>	Formattato: Allineato a sinistra	
	Widder					species				tuna <u>s</u> )	model	<u>codebase</u>				
ERSEM II	Commission for the Conservation of Antarctic	Functional group approach	Nutrient	month	Yes	Limited number of HTL	Type II	Effects in both directions	Intermediate	<u>Too</u> <u>complex</u> No	Yes, detailed	https://github.com/ pmlmodelling/erse	Butenschön et al., 2018	<b>*</b>	Formattato: Allineato a sinistra	
	warme Living Resources	**				groups						m				
<u>Apecosm</u>	<u>Apex Predators</u> ECOSystem Model	Size spectra approach	<u>Biomass</u>	month	Yes	<u>Few</u> species	<u>Few top</u> predators	Top predator group dynamics	Poor (few applications)	Poor (model complexity)	<u>Tes, included</u>	https://github.com/ apecosm/python- apecosm	<u>Maury, 2010</u>			

Whole Ecosystem Models													<u>Plagán</u>
<u>Acronym</u> Model	<u>Model</u> Name	Elemental structure	Model Units	Time units	Spatial structure	Number of species	Trophic interactions	Primary model focus, output	<u>Maturity</u>	Readiness for operational useOperation al	Physical/biogeochem ical processes	<u>Repository</u> (if open access)	
ATLANTIS	Atlantis	Functional group approach; Populations in age structure	Nutrient	month	Yes	Can be a very large number; typically order 40	Flexible, Type II, type III or other	Effects of ecosystem and fisheries in both directions; yearly outputs	<u>High</u>	<u>poorNo</u> (model complexity)	Yes, detailed	https://github.com/ runatlantis/atlantis	Fulton e
EwE	Ecopath with Ecosim	Functional group approach; Populations also in age structure	Biomass, Nutrient	month	Yes (ECOSPACE)	Can be a very large number; typically order 40	Foraging arena, flexible approach	Effects of ecosystem and fisheries in both directions; yearly outputs	High	<u>PooNor</u> (model complexity)	Included as off-line coupling	https://ecopath.org	<sup>I</sup> Christens
OSMOSE	Object-oriented Simulator of marine ecosystem Exploitation	Size spectra approach	Biomass at different levels of aggregation	year	Yes	Large number of species	Efficient predator but can starve	Multispecies dynamics	Intermediate	Intermediate (model complexity) <del>N</del> <del>o</del>	Included as off-line coupling	https://osmose- model.org/	<u>Shin and</u>
FEISTY	<u>FishErIes Size and</u> <u>functional TYpe model</u>	Size spectra approach	Biomass at different levels of aggregation	year	Yes	Large number of species	Flexible approach	Multispecies dynamics	Intermediate	IntermediateN o	Included as off-line coupling	https://github.com/ Kenhasteandersen/ FEISTY	Blancha
Apecosm	<b>A</b>												

