

1 **EwE-F 1.0: An implementation of Ecopath with Ecosim in**  
2 **Fortran 95/2003 for coupling and integration with other**  
3 **models**

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5 **Ekin Akoglu<sup>1,2</sup>, Simone Libralato<sup>1</sup>, Baris Salihoglu<sup>2</sup>, Temel Oguz<sup>2</sup>, Cosimo**  
6 **Solidoro<sup>2</sup>**

7 [1]{OGS (Istituto Nazionale di Oceanografia e di Geofisica Sperimentale), Via Beirut 2/4  
8 (Ex-SISSA building), 34151, Trieste, Italy}

9 [2]{Middle East Technical University, Institute of Marine Sciences, P.O. Box 28, 33731,  
10 Erdemli, Mersin, Turkey}

11 Correspondence to: Ekin Akoglu (eakoglu@ogs.trieste.it)

12

13 **Abstract**

14 Societal and scientific challenges foster the implementation of the ecosystem approach to  
15 marine ecosystem analysis and management, which is a comprehensive means of integrating  
16 the direct and indirect effects of multiple stressors on the different components of ecosystems,  
17 from physical to chemical and biological and from viruses to fishes and marine mammals.  
18 Ecopath with Ecosim (EwE) is a widely used software package, which offers capability for a  
19 dynamic description of the multiple interactions occurring within a food web, and potentially,  
20 a crucial component of an integrated platform supporting the ecosystem approach. However,  
21 being written for the Microsoft .NET framework, seamless integration of this code with  
22 Fortran-based physical oceanographic and/or biogeochemical models is technically not  
23 straightforward. In this work we release a re-coding of EwE in Fortran (EwE-F). We believe  
24 that the availability of a Fortran version of EwE is an important step towards setting-up  
25 coupled/integrated modelling schemes utilising this widely adopted software because it i)  
26 increases portability of the EwE models, ii) provides additional flexibility towards integrating  
27 EwE with Fortran-based modelling schemes. Furthermore, EwE-F might help modellers using  
28 Fortran programming language to get close to the EwE approach. In the present work, first  
29 fundamentals of EwE-F are introduced, followed by validation of EwE-F against standard

1 EwE utilising sample models. Afterwards, an end-to-end (E2E) ecological representation of  
2 the Trieste Gulf (Northern Adriatic Sea) ecosystem is presented as an example of online two-  
3 way coupling between an EwE-F food web model and a biogeochemical model. Finally, the  
4 possibilities that having EwE-F opens up are discussed.

5

6 **1 Introduction**

7

8 Oceanographic models, particularly computationally intensive hydrodynamic and  
9 biogeochemical models, have mostly been written in Fortran (e.g. hydrodynamic models:  
10 NEMO (Madec 2008), ROMS (Shchepetkin and McWilliams, 2005), POM (Blumberg and  
11 Mellor, 1978), MITGCM (Adcroft et al., 2004), MOM (Stock et al., 2014); and  
12 biogeochemical models: ERSEM (Blackford et al., 2004), BFM (Vichi et al 2015), ERGOM  
13 (Neumann, 2000)). In fact, Fortran was the first programming language specifically designed  
14 for solving engineering and scientific computing problems (Backus et al., 1957) and proved to  
15 be one of the most efficient for performing complicated mathematical tasks with its collection  
16 of predefined high-level mathematical functions. Over the years, frequent revision of the  
17 Fortran standard and the addition of new capabilities to the language to meet changing  
18 demands enabled it to remain as *de facto* standard for writing computationally intensive  
19 scientific and engineering applications.

20 Ecopath with Ecosim (hereinafter EwE) (Christensen and Walters, 2004; Christensen et al.,  
21 2005) is the most widely adopted tool for building models of marine and freshwater  
22 ecosystems, and possibly the first choice for analysis of food web dynamics. Freely available  
23 at [www.ecopath.org](http://www.ecopath.org), EwE has long been used for scientific studies related to fisheries, and  
24 also including some aspects of aquaculture, marine ecology, climate and pollution. There are  
25 thousands of users of the software worldwide (last record in 2008, reported 5649 users;  
26 [www.ecopath.org](http://www.ecopath.org)) and more than 400 scientific publications utilising EwE as a modelling  
27 tool have been issued only in the last two decades (search on Web of Science on 29/09/2014  
28 for “Ecopath with Ecosim” or “Ecospace” or “Ecopath” resulted in 469 items published  
29 between 1997 and 2014). Because many EwE models for a variety of aquatic ecosystems are  
30 available, it makes sense to capitalise on such experience when developing coupled/integrated  
31 modelling applications. This would require only minimal modifications in these models and  
32 remove the burden of starting from scratch. However, being written for the Microsoft .NET

1 framework constrains EwE's ability to integrate with models written in Fortran and the  
2 Fortran recoding of EwE presented in this paper will facilitate this.

3 EwE is designed for interoperability with other models, which is crucial considering that  
4 ecological modelling is facing an important challenge to set a basis for the comprehensive  
5 description of marine ecosystems through integrated modelling schemes that incorporate  
6 multiple models (e.g. hydrodynamic, biogeochemical, ecological and socioeconomic)  
7 interactively with one another (e.g. end-to-end (hereinafter E2E) models (Fulton, 2010)). This  
8 interoperability leads to insightful linking of these models into EwE (e.g. Christensen et al.,  
9 2014) and EwE's flexibility already permits to link physical/biogeochemical oceanographic  
10 models with EwE (e.g. Libralato and Solidoro, 2009). This one-way linking permits  
11 exchanges of information between models that are run separately and is valid, robust and  
12 usually faster to implement than a two-way coupling. In spite of the interesting results  
13 obtained, however, one-way linking lacks a complete representation of feedbacks that  
14 propagate two-ways between the coupled models. These feedbacks were proven to be  
15 important and reveal important ecological mechanisms (Kearney et al., 2012) that need to be  
16 accounted explicitly for a full representation of ecosystem effects due to climatic changes,  
17 aquaculture, socioeconomic changes and other important drivers (Fulton, 2010). The  
18 scientific requirements for such modelling approaches, therefore, mandate two-way coupling  
19 with existing oceanographic models which are mostly written in Fortran. Because these  
20 models and EwE use different programming languages, the technical differences complicate  
21 the coupling task more than anticipated (e.g. Beecham et al., 2010). One possible solution is  
22 the offline coupling of EwE and Fortran-coded models via two-way data transfer between the  
23 models at predefined time intervals while pausing the other model (i.e. turn-based run).  
24 Another solution could be utilising inter-process communications such as pipes and/or sockets  
25 between EwE and the model to be coupled while simultaneously running the models.  
26 However, coupled model construction will benefit from a Fortran version of EwE that will  
27 permit direct integration of the EwE modelling approach with mainly, but not limited to,  
28 physical and biogeochemical models in Fortran, and will allow a straightforward and two-way  
29 propagating feedback between high trophic level (HTL) and low trophic level (LTL) models.  
30 Hence, the development of a Fortran version of EwE will be useful for integration of HTL  
31 food web models with potentially any other model written in Fortran which simulates, for  
32 example, socioeconomic, bioenergetic dynamics.

1 In this work, we present (Section 3) the first version of EwE re-coded in Fortran 95/2003  
2 language standard (EwE-F, version 1.0). In Section 3.3, we provide evidence of the full  
3 reliability of the code by comparing EwE-F with standard EwE (version 6.5) utilising sample  
4 food web models. In Section 4, we present how EwE-F allows for easy coupling with other  
5 models, by providing an example of integration with a biogeochemical model of the Gulf of  
6 Trieste in the Northern Adriatic Sea. Finally, in the same section, we discuss the possibilities  
7 opened up by the availability of EwE-F. We believe that EwE-F will appeal also to the  
8 scientific community previously sceptical to the EwE approach (usually more confident with  
9 Fortran programming) and provide the possibility of both easy modification of the EwE-F  
10 structure and parameterisation for specific cases and easy integration with other  
11 biogeochemical, population dynamics, individual-based and/or any type of ecological model  
12 written in Fortran.

13

## 14 **2 A brief description of the EwE Model**

15

16 EwE modelling software includes a suite of modules that enables the building and analysis of  
17 food web models. EwE includes three main modules; i) Ecopath; the mass-balance  
18 representation, ii) Ecosim; the time-dynamic simulation, and iii) Ecospace; the 2D spatial-  
19 temporal dynamics, plus other complementary routines: Network Analysis (Ulanowicz,  
20 1986), Monte-Carlo Simulation and Time Series Fitting. EwE-F comprises only Ecopath and  
21 Ecosim modules thus only these two are briefly summarised here.

22 The Ecopath module comprises a series of linear equations that defines a mass-balance  
23 stationary state of the food web. The functional groups are regulated by gains (consumption,  
24 production, and immigration) and losses (mortality and emigration), and are linked to each  
25 other by predatory relationships. Fisheries extract biomass from the targeted and by-catch  
26 groups. In Ecopath, a set of linear equations describes flows of mass into and out of discrete  
27 biomass pools of the form

$$B_i * \left(\frac{P}{B}\right)_i - \sum_{j=1}^n B_j * \left(\frac{Q}{B}\right)_j * DC_{ji} - B_i * \left(\frac{P}{B}\right)_i * (1 - EE_i) - Y_i - E_i - BA_i = 0 \quad (1)$$

28 where, for each functional group  $i$ ,  $B$  stands for biomass,  $(P/B)$  stands for the production rate  
29 per unit of biomass,  $(Q/B)$  stands for the consumption rate per unit of biomass of predator  $j$ ,

1  $DC_{ji}$  is the fraction of prey  $i$  in the average diet of predator  $j$ ,  $Y$  is the landings,  $E$  is net  
2 emigration rate, and  $BA$  is the biomass accumulation rate (Christensen et al, 2005).  $EE$  is the  
3 ecotrophic efficiency representing the proportion of mortality of a group that is not  
4 attributable to predators or fishing activities. As it can be seen, Equation (1) is quite simple as  
5 a result of the fact that it represents the budget of biomass fluxes in a given time window  
6 within an ecosystem. Ecopath is also characterized by a top-down solution of the system of  
7 equations, i.e. consumption on a group is a function of predator biomass, which differs from  
8 bottom-up approaches used in other inverse modelling methods (Steele, 2009).

9 In the time-dynamic module of EwE (Ecosim), dynamics of a state variable are defined with a  
10 differential equation composed of sources and sinks terms. Each state variable represents the  
11 biomass of a functional group representing species and/or groups of species or populations  
12 split into age-size categories (multi-stanza). The definition of such differential equation in  
13 Ecosim is as follows:

$$\frac{dB_i}{dt} = \gamma_i * \sum_{j=1}^n Q_{ji} - \sum_{j=1}^n Q_{ij} + I_i - (M_i + F_i + e_i) * B_i \quad (2)$$

14 where  $dB_i/dt$  is the rate of change of biomass ( $B$ ) of group  $i$  over time  $t$ ,  $\gamma$  is the growth  
15 efficiency of group  $i$ ,  $\sum Q_{ji}$  is the sum of the consumptions of group  $i$  over all of its preys,  $\sum$   
16  $Q_{ij}$  is the sum of the predation on group  $i$  by all of its predators,  $I$  is the immigration,  $M$  is the  
17 non-predation mortality,  $F$  is the fisheries mortality and  $e$  is the emigration rate of group  $i$   
18 (Walters et al., 1997).  $Q_{ij}$  is defined on the basis of biomasses of predator and prey in a form  
19 that represents a slightly modified version of Holling Type II functional response in order to  
20 consider only the part of the biomass of the prey  $i$  that is accessible to the predator  $j$  (foraging  
21 arena theory; Ahrens et al., 2012). For each trophic interaction, the accessible biomass is  
22 dynamically defined on the basis of a parameter called “*vulnerability*” (for details refer to  
23 Walters et al., 1997; Walters et al., 2000; Ahrens et al., 2012). This system of differential  
24 equations is numerically integrated over time under the influence of forcing functions  
25 (typically fishing mortalities or efforts, changes in primary productivity) starting from the  
26 initial condition settings defined by the Ecopath module.

27

### 28 **3 The EwE-F Software**

29

1 The EwE software was translated to Fortran 95/2003 language in its core architecture and  
2 kept limited to; i) the Ecopath mass-balance routine including multi-stanza calculations, and  
3 ii) the Ecosim time-dynamic simulation including multi-stanza calculations. Due to  
4 modularity considerations, EwE-F was implemented under two separate components; i)  
5 Ecopath-F: the Ecopath mass-balance algorithm, and ii) Ecosim-F: the Ecosim time-dynamic  
6 simulation algorithm. EwE-F v1.0 includes only core routines of Ecopath and Ecosim:  
7 complementary routines for calculation of indicators for Network Analysis, as well as  
8 routines for Monte-Carlo Simulation, Time Series Fitting and Ecospace are not included. Also  
9 the capability to define mediation functions is not yet implemented in EwE-F v1.0, although  
10 is planned to be addressed in future versions. A schematic view of the EwE-F components  
11 and the input/output (I/O) files necessary for information exchange are given in Figure 1. In  
12 the following two sections (3.1 and 3.2), the structure and functioning of the components in  
13 Figure 1 are described in detail.

14

### 15 **3.1 Ecopath-F**

16

17 Ecopath-F is the component of EwE-F that carries out mass-balance calculations given in  
18 Equation (1). Similar to stock Ecopath, it requires the same fundamental input parameters to  
19 be entered via four tab-delimited ASCII (American Standard Code for Information  
20 Interchange) encoded text input files; i) a scenario file containing the basic input and multi-  
21 stanza parameters and catches, ii) a file comprising the diet composition matrix of the state  
22 variables, iii) a file comprising the detritus fate of the state variables and, iv) if applicable, a  
23 file including the growth parameters of the multi-stanza groups. Furthermore, Ecopath-F  
24 requires a Fortran “namelist” file that includes the full paths and names of the above-  
25 mentioned four input files and, in addition, the path and name of the output HDF5  
26 (Hierarchical Data Format version 5, [www.hdfgroup.org/HDF5](http://www.hdfgroup.org/HDF5)) file which the mass-balance  
27 calculation results will be output to and be used to initialise and run Ecosim-F (Figure 1).

28 An Ecopath-F run produces two output files; i) an ASCII file which includes the summary of  
29 estimated parameters and basic statistical information, and ii) an HDF5 file specifically  
30 formatted to define the initial conditions for the Ecosim-F simulation (Figure 1). The output  
31 HDF5 file includes all the parametric details about the state variables of Ecopath run and

1 further comprises the diet composition matrix, detritus fate matrix and multi-stanza group  
2 parameters.

3 Ecopath-F is independent of the Ecosim-F implementation; however, Ecosim-F requires  
4 output data from Ecopath-F plus additional parameter settings. The data transfer from  
5 Ecopath-F to Ecosim-F is carried out via the intermediary HDF5 data file.

6

7 **3.2 Ecosim-F**

8

9 Ecosim-F is the component of EwE-F that carries out time-dynamic simulation calculations  
10 given in Equation (2). Ecosim-F requires the HDF5 output file from the Ecopath-F run and,  
11 depending on the compile time options, at least three additional tab-delimited ASCII encoded  
12 text input files; i) a scenario file containing group information of state variables, ii) a file  
13 comprising the vulnerability matrix between predator-prey pairs, and iii) a file comprising the  
14 monthly fishing mortality/effort time series forcing functions for all state variables (Figure 1).  
15 Similar to Ecopath-F, Ecosim-F also requires a namelist file that includes the full paths and  
16 names of the input files as well as the values of some particular variables; i.e. number of time  
17 steps per month, base proportion of free nutrients, relaxation parameter and simulation time in  
18 years, to prepare the Ecosim simulation (for details see Christensen et al., 2005, p. 78; Akoglu  
19 et al., 2015).

20 Once completed, Ecosim-F simulation produces five tab-delimited ASCII coded text files  
21 comprising the annual and monthly absolute and relative biomass values of the state variables  
22 and a file comprising monthly catches of the fished state variables throughout the simulation  
23 in the model directory (Figure 1).

24

25 **3.3 The skill assessment of EwE-F**

26

27 In order to assess the skill of EwE-F with respect to EwE, two test case simulations, “Generic  
28 37” and “Tampa Bay”, which are distributed with the installation of the EwE software were  
29 used. The test case simulations were run both with EwE version 6.5 and EwE-F version 1.0

1 and the residuals between simulated absolute biomasses of state variables were used to  
2 evaluate the performance of EwE-F. It is worth noting that other EwE versions may produce  
3 slightly different results compared to EwE-F v1.0. The residuals for each state variable in the  
4 respective simulations were visualised with box-whisker plots showing the minimum value,  
5 25th percentile, median, 75th percentile and maximum values respectively (Figure 2 and  
6 Figure 3).

7 The residuals between the simulated biomass values of EwE-F and EwE ranged from  $10^{-8}$  to  
8  $10^{-5}$ , with the maximum difference found to be on the order of  $10^{-5}$ . The residuals calculated  
9 from the comparison of the simulations justified that EwE-F possessed the necessary skill to  
10 reproduce the results of EwE for the Generic 37 and Tampa Bay simulations. The magnitude  
11 of the misfits concluded that EwE-F was capable of being used in conjunction with other  
12 models without introducing significant sources of error to the resulting modelling scheme.

13

#### 14 **4 Exploring EwE-F flexibilities: example from a complex coupling exercise**

15

16 The Fortran recoding of EwE creates great flexibility for customisation, modification or  
17 coupling to different models written in Fortran. An example, which illustrated the potential of  
18 such flexibility, came from the integration of EwE-F to a biogeochemical Fortran model. In  
19 fact, the direct integration of these two models required to address, and subsequently solve a  
20 number of problems. These included defining the links between the two models and  
21 modifying them accordingly, exchanging information between the two models, dealing with  
22 different model time steps, and accounting for different model currencies.

23 The HTL model is an updated version of the EwE model of the Northern Adriatic Sea  
24 originally developed by Coll et al. (2007). The original model which is composed of 40  
25 functional groups (FG) has been updated by i) removing discards and by-catch FGs, ii)  
26 splitting phytoplankton and zooplankton in two FGs each to represent small and large taxa;  
27 iii) adding bacteria to explicitly represent the microbial loop; iv) adjusting diet of plankton  
28 feeders to split the diet into the new plankton FGs. The updated model has 44 FGs and  
29 parameters for the plankton groups were updated considering literature information (see  
30 Cossarini and Solidoro, 2008 and references therein). The model currency is wet weight. The  
31 time step of the model is one month, the default time step of the EwE software.

1 The biogeochemical model is a Fasham-like (Fasham et al., 1990) 0D box model of the  
2 Northern Adriatic Sea (Cossarini and Solidoro, 2008) and consists of phytoplankton,  
3 zooplankton, and heterotrophic bacteria groups, one pool of inorganic phosphorus ( $\text{PO}_4^{3-}$ );  
4 one dissolved organic matter compartment in terms of phosphorus (DOP) and carbon (DOC),  
5 and one particulate organic matter compartment in terms of phosphorus (POP) and carbon  
6 (POC) (Figure 4). The model is a multi-currency model calculating the biomasses of its  
7 particular state variables (sediment, dissolved organic matter, particulate organic matter) both  
8 in terms of carbon and phosphorus. The time step of the model is one hour. Full description of  
9 the biogeochemical model is reported in Cossarini and Solidoro (2008).

10 For the harmonisation of both models in an E2E coupled scheme, first, the state variables that  
11 were already present in the LTL model were removed from the HTL model as well as their  
12 links (grey-shaded area and links in Figure 4). Then the linkages between the state variables  
13 of the HTL model and the state variables of the LTL model were set up in accordance with  
14 the removed state variables as shown in Figure 4 (links in dashed and continuous black lines).  
15 In this way, a coupled model scheme that consisted of 44 functional groups was set up: 9 FG  
16 represented the state variables of the biogeochemical model, i.e., plankton groups plus  
17 inorganic and organic nutrient forms (Figure 4). For simplicity, the HTL and LTL groups are  
18 not given in detail in the figure, however, sources and sinks of the whole HTL compartment  
19 and the linkages between the HTL and LTL domains and state variables are shown.

20 The second step in the harmonisation of models consisted of accounting for the different  
21 currencies used. Considering the multiple currency utilisation of the biogeochemical model  
22 for some of its state variables and the fact that the application of a similar principle in the  
23 HTL model would require the modification of the various calculations in the state equation of  
24 the original EwE software, the state variables of the HTL model, which were in wet weight  
25 (tons), were converted to phosphorus ( $\mu\text{mol P}$ ) weight utilising C:N:P ratios taken from  
26 literature.

27 The third step in the harmonisation procedure was to reconcile the differences in the  
28 integration time step between the two models. Considering that the biogeochemical model  
29 consisted of state variables with faster dynamics compared to HTL model, it was convenient  
30 to make the HTL model comply with the integration step of the biogeochemical model. For  
31 this purpose, the rates of the HTL model, which were “per year ( $\text{yr}^{-1}$ )”, were converted to “per

1 hour ( $\text{h}^{-1}$ )” by simply dividing the rates by 8760 (365  $\text{d}^{-1}$  x 24  $\text{h}^{-1}$ ) so that the HTL variables  
2 could be integrated with the same time step of the biogeochemical model.

3 The final step in the harmonisation process would be to adjust the closure terms of the  
4 biogeochemical model (mortality rates of zooplankton and phytoplankton groups) so as to  
5 compensate the additional losses through explicit predation of these groups by the HTL state  
6 variables. However, for our specific application, we decided to keep these values identical to  
7 the standalone biogeochemical model as the coupled model produced similar seasonal cycles  
8 observed in the standalone biogeochemical model except the missing second cycle in  
9 mesozooplankton (Figure 6) and as our aim was indeed to have plankton dynamics  
10 qualitatively comparable to the biogeochemical model.

11 The technical overview of the coupling scheme is given in Figure 5. As shown in the figure,  
12 the coupled simulation was carried out in four consecutive stages. In the first stage, a static  
13 mass-balance model of the whole system, which comprised all the HTL and LTL state  
14 variables in the ecosystem, was set up utilising Ecopath-F. In this stage, the LTL state  
15 variables were ordered in advance of the HTL state variables so that the LTL state variables  
16 were numbered from 1-9 and the HTL state variables from 10-35 in the resulting scheme.  
17 Following this procedure, Ecopath-F was run to calculate the basic parameters and exchange  
18 rates between the state variables of the HTL and LTL compartments which were necessary to  
19 perform a dynamic simulation after completing all of the harmonisation steps. In the second  
20 stage, utilising the calculations from the previous stage, the HTL and LTL models were  
21 initialised by calculating initial conditions for each of their respective state variables utilising  
22 their specific internal routines. In the third stage, the sources and sinks of HTL and LTL state  
23 variables were computed via utilising their respective derivative functions during the whole  
24 simulation period. The selection of the derivative function to be used to calculate the  
25 differentials of the state variables depended on the rank of the state variables determined  
26 during the Ecopath-F set-up in the first stage. This stage continued iteratively until the end of  
27 the simulation and, at the end of each time step, stage four was executed so that the results  
28 calculated at each time step were, if required, post-processed and then written to the results  
29 files. Post-processing of LTL results might not be necessary in all cases but only if the LTL  
30 model is a multi-currency model and calculates its variables in more than one currency. In our  
31 example, because the LTL model represented some of its state variables both in carbon and  
32 phosphorus but the coupled HTL model only in phosphorus, a post-processing step was

1 necessary to compute the corresponding phosphorus values of variables that were in carbon  
2 units while interchanging information between the HTL and LTL derivative functions as well  
3 as before writing the results into the output files. The coupled simulation was run ten years,  
4 two of which were for spin-off. In the simulations, we used default values for vulnerabilities  
5 ( $v_{ij} = 2$ ) that represent a mixed control (Christensen et al., 2005).

6 Comparison of uncoupled and coupled model results (Figure 6) demonstrated that the  
7 coupling scheme worked successfully and highlighted the effects of integration of LTL and  
8 HTL models. Because the aim of this exercise was only to demonstrate the capability of EwE-  
9 F to be used in integration with other models, the ecological interpretations of these results are  
10 not the focus of this work and thus are only briefly discussed here. Comparing the seasonal  
11 dynamics of LTL state variables before and after coupling showed that explicit addition of  
12 HTL dynamics influenced the seasonality of the LTL state variables (grey-shaded plots in  
13 Figure 6). It is worth noting that presence of several detrital and predatory links between HTL  
14 and LTL models (as shown in Figure 4) resulted in clear top-down impacts on the LTL  
15 variables, particularly in non-living and bacteria. Furthermore, the comparison between the  
16 simulation results of HTL model forced with primary productivity changes (green lines in  
17 Figure 6) in stock EwE and the fully coupled HTL/LTL models (black lines) showed that  
18 changes in the biogeochemical dynamics, namely nutrient recycling, not only impacted the  
19 LTL groups but also propagated up through the food web (bottom-up) to impact the  
20 biomasses of HTL organisms. While most of the bottom-associated state variables decreased  
21 by the incorporation of the biogeochemical model in the coupled scheme, pelagic-associated  
22 state variables increased due to the explicit representation of resuspension of detritus and  
23 remineralisation that favoured plankton. Thus as evidenced in Figure 6, the consequences of  
24 two-way coupling were not only one directional. These proved that the proper exchange of  
25 information and the establishment of successful interaction between the two models were  
26 realised in the final coupled scheme.

27

28 **5 Discussions**

29 **5.1 Potential and flexibility of the application**

30

1 In this work, the reliability of EwE-F was proven by utilising two sample models as test cases  
2 and comparing the absolute biomass values simulated by EwE-F against the simulated  
3 absolute biomass values by stock EwE version 6.5. Further, the applicability of EwE-F in an  
4 E2E modelling framework was exemplified with a test case for the Gulf of Trieste ecosystem.  
5 This example proved the adaptability of EwE-F for coupled modelling frameworks,  
6 facilitating its integration with other hydrodynamic and biogeochemical Fortran models for  
7 aquatic ecosystems in ecosystem research. The scheme used in this work successfully  
8 conveyed two-way dynamics of HTL and LTL domains along the whole food web. As a step  
9 forward, this opened up the opportunity for using EwE, by utilising EwE-F implementation,  
10 as an HTL component of holistic ecosystem representations in various ecosystems.

11 According to Rose et al. (2010), the main difficulty encountered in coupling models of  
12 different realms lies in the reconciliation of the differences in time and spatial resolutions.  
13 However, difficulties may extend beyond these two areas, e.g. differences in model  
14 currencies. The coupling scheme used in this work is able to provide solutions to overcome  
15 such constraints highlighted by Rose et al. (2010) and others (Fulton, 2010; Kearney et al.,  
16 2012; Salihoglu et al., 2013) via its simplistic but ecologically capable approach to form E2E  
17 representations of aquatic ecosystems through the incorporation of EwE-F. In addition, the  
18 EwE-F enables significant opportunities for integrating it with any kind of Fortran models as  
19 depicted in Figure 7. The figure represents a typical EwE food web model in the middle  
20 rectangular box and elaborates the possibilities of modifying EwE-F in different ways by  
21 replacing different components with sophisticated model representations for selected state  
22 variables or incorporating additional Fortran models to enhance the applicability of the  
23 original EwE approach. These solutions and possibilities are explored in detail in the  
24 following sections; i) reconciling different integration steps (Section 5.1.1), ii) dealing with  
25 models that use multiple currencies (Section 5.1.2), iii) other possibilities: incorporation of  
26 population demographic structure, physiological processes, socioeconomical frames (Section  
27 5.1.4).

28

## 29 5.1.1 Reconciling different integration steps

30

1 There are two possibilities when combining two models with different integration (time)  
2 steps; i) keeping the integrator function of the two models intact and averaging the outputs of  
3 the model with faster dynamics (high turnover rate) over the time frame of the model with  
4 slower dynamics (low turnover rate) and vice versa when exchanging information (time-  
5 averaged coupling), and ii) utilising a common integrator for both models and adjusting the  
6 rates of the model with slower dynamics to comply with the time window of the model with  
7 faster dynamics (real-time coupling). Although Ecosim, by default, works with monthly time  
8 steps it is capable of simulating high frequency dynamics using shorter time steps. In the  
9 present work, we opted for the latter to showcase the possibility of harmonisation in terms of  
10 integration step size when using EwE-F in coupled modelling schemes. The difference in the  
11 time resolution of both models was remedied by adjusting the HTL model's time step (one  
12 month) to conform to the time step of the biogeochemical model (one hour) in order to render  
13 the use of one common ordinary differential equation (ODE) solver (the Runge-Kutta 4th  
14 order) possible. Furthermore, due to this change in the time step of the HTL model, the annual  
15 rates of the HTL groups were converted to hourly rates by simple arithmetic calculations.

16

### 17 5.1.2 Dealing with models that use multiple currencies

18

19 Some biogeochemical models may carry out their computations in more than one currency for  
20 explicit representation of the ratios of fundamental nutrients in the system and their rate  
21 limiting conditions on nutrient uptake and primary productivity that can vary in space and  
22 time. The multiple currency approach, however, is usually not applied in HTL models,  
23 although implicit nutrient-based limitations can be represented in EwE (Araujo et al., 2006;  
24 Christensen et al., 2005). Hence, the coupling exercise presented here provided a simple  
25 solution for such situations. In order to reconcile the currency differences, one may opt to  
26 pick one of the currencies utilised in the biogeochemical model as the one considered to be  
27 the limiting nutrient, use it for the final coupled scheme incorporating the EwE-F model and  
28 post-process the derivative function outputs of the two models when exchanging information.  
29 In the coupling example given in this work, the difference in the currencies of the models was  
30 adjusted by converting the currency of the HTL model from wet weight to phosphorus (P)  
31 utilising the conversion rates and equations available in the literature for HTL groups (stage 1

1 of the coupling scheme in Figure 5). In addition, the simulated results of the biogeochemical  
2 model (which were in dual currency, phosphorus and carbon) were post-processed prior to  
3 output and transferred to EwE-F so as to comply with the currency of the HTL compartment  
4 (stage 4 in Figure 5). The approach used in this work proved to be a practical solution for the  
5 issue in cases where there is no particular consideration to have simultaneously tracking  
6 multiple currencies in the HTL food web. However, with the availability of EwE-F, HTL  
7 models with computations of multiple model currencies can even be set up if desired,  
8 although this will require significant modification of various calculations in the EwE state  
9 equations.

10

### 11 5.1.3 Spatial simulations

12

13 Given the current experience with biogeochemical models coupled with hydrodynamic  
14 models (e.g. Lazzari et al., 2012), explicit accounting for spatial variability is important for  
15 any assessment of marine ecosystem dynamics. Future efforts are required to add spatial  
16 simulation capabilities to EwE-F, either by implementing Ecospace in Fortran or by direct  
17 integration of Ecosim-F in a spatially explicit coupled hydrodynamic-biogeochemical model.  
18 This planned future work could lead EwE-F to play a substantial role in spatial simulations.

19

### 20 5.1.4 Other possibilities: population demographic structure, physiological 21 processes, socioeconomical frames

22

23 Similar to the flexibility of EwE provided by its plugin system, EwE-F gives broad  
24 possibilities for interconnecting HTL models with other Fortran models sophisticating and/or  
25 incorporating HTL processes. Examples span from fish population to socioeconomic dynamic  
26 models.

27 For instance, EwE-F permits incorporating sophisticated population dynamic models written  
28 in Fortran within the EwE-F scheme (Figure 7, C). These population models can be of any  
29 kind, including population's demographic structure (age/size classes) used for stock

1 assessment and account for differences in fecundity by ages or size (Hilborn and Walters,  
2 1992).

3 Moreover, EwE-F allows for parameterising various rates for HTL groups (e.g. assimilation  
4 efficiency, respiration) under the influence of various environmental factors (e.g. temperature,  
5 pH, light) that is not always straightforward otherwise (Figure 7, D). In addition, EwE-F  
6 allows for replacing the growth of certain state variables in the food web with sophisticated  
7 bioenergetics models coded in Fortran. In this way, various physiological processes of the  
8 selected HTL organisms can be related directly and explicitly to the ambient physical factors  
9 such as light, temperature and nutrient availability (Figure 7, B). With EwE-F, in fact, as  
10 demonstrated in this work, the dynamics of any desired additional state variable in the final  
11 coupled scheme could be resolved using derivative functions defined in other models during  
12 run-time. This allows for a two-way coupling of, potentially, any number of models  
13 (including earth system ones) in one coupling scheme.

14 Given the calls for ecosystem-based management for marine ecosystems, one can also  
15 incorporate socioeconomical dynamics on holistic ecosystem representations that deal with  
16 fisheries on top of EwE-F. Considering its modular structure and ease of integration with  
17 other models as demonstrated in this work, such holistic representations of ecological and  
18 socioeconomical systems have been significantly improved also including frameworks that  
19 involve integration of multiple models written in Fortran (Figure 7, A).

20

## 21 **5.2 Other practical considerations and future development**

22

23 In contrast to the EwE, the introduction of namelist and HDF5 files to be used for the  
24 operation of EwE-F may create a hindrance to its users. However, it is not necessarily more  
25 complicated than the current EwE database files (MS Access). EwE-F requires an HDF5  
26 database file only when transferring information from Ecopath-F to Ecosim-F, and output to  
27 and input from this file does not require any user intervention. In addition, the results of both  
28 Ecopath-F and Ecosim-F models are output into TAB-delimited ASCII files, which are quite  
29 similar to the EwE's output files, i.e. comma-separated value (CSV) ASCII files. These files  
30 can easily be opened with spreadsheet programs. The only hindrance for the user could be the

1 preparation of the TAB-delimited ASCII input files for Ecopath-F and Ecosim-F, which  
2 however is explained in the User's Manual in detail. On the other hand, through this simple  
3 input/output scheme utilising ASCII encoded text files, the availability of EwE-F provides a  
4 further opportunity by giving Fortran modellers the possibility to perform detailed sensitivity  
5 and uncertainty analyses using hundreds of ensemble scenarios that can easily be prepared  
6 also by using modern high-level languages (e.g. Perl, Python, NCL) in addition to Fortran.  
7 For their convenience, users of EwE-F are advised to set up, test and fit their models to time  
8 series data using EwE also benefiting from the several routines included in EwE and,  
9 thereafter, transfer their models to EwE-F.

10 Ecospace (Walters et al., 1999) and other complementary routines aforementioned (see  
11 section 3) were not implemented considering that EwE-F was not designed to be an EwE  
12 replacement but a bare-bones incarnation that can be used easily for purposes summarised in  
13 Section 5.1.4. Therefore, analyses requiring the aforementioned specific routines (e.g. Monte-  
14 Carlo analysis, Network Analysis etc.) in uncoupled or coupled EwE-F simulations can be  
15 done by coding the required specific routines or alternatively EwE could be employed for  
16 such purposes. The current lack of such useful tools that are present in EwE 6.5 is considered  
17 as a drawback for the EwE-F v1.0, which may represent an obstacle for some users. However,  
18 these technical shortcomings and the lack of these tools including mediation function and  
19 time series fitting via vulnerability parameter search are planned to be addressed in the future  
20 by incorporating these routines in EwE-F and developing a Visual Basic plug-in for stock  
21 EwE which will prepare input files required by EwE-F through EwE's graphical user  
22 interface in a straightforward way. Furthermore, considering advancements for coupling in  
23 the spatial scale, future efforts of developing EwE-F may also focus on incorporating 2D  
24 spatial dynamics by implementing the Ecospace module of EwE to facilitate the use of EwE-F  
25 in schemes that require spatial-temporal dynamics to be resolved.

26 Another important consideration to be discussed is to keep up EwE-F on par with EwE. With  
27 every new release of EwE software, many things are prone to change. However, the majority  
28 of these changes are related to the ancillary functionalities (graphical user interface, network  
29 analysis routines etc. but not the core state equations and its related calculations) that are not  
30 included in EwE-F. Furthermore, the changes to the basic model structure and dynamics have  
31 remained almost unchanged since EwE version 5. Hence, it is believed that the core structure  
32 of EwE-F (state equations and other related calculations) can be kept on par with the original

1 EwE with little effort, considering that the development of EwE-F is a joint effort of two  
2 prominent marine science institutes and is not strictly bound to any individual.

3

4 **6 Code availability**

5

6 The source code of EwE-F version 1.0 detailed in the present work and the corresponding  
7 User's Manual can be obtained as supplementary material to this article. In the User's  
8 Manual, detailed instructions to obtain the current and future versions of EwE-F along with  
9 building and running EwE-F on different platforms are described. Further versions of EwE-F  
10 model and their respective documentations can be obtained on Bitbucket.org  
11 (<https://bitbucket.org/ewe-f>). The system requirements, license and other basic information  
12 regarding EwE-F version 1.0 are given in Table 1.

13

14 **7 Conclusions**

15

16 It has been shown that a Fortran version of EwE software could open up various possibilities  
17 in terms of coupling and integration with other Fortran-coded biogeochemical and  
18 hydrodynamic models where an HTL compartment is required. In order to exemplify the  
19 applicability of the approach, a coupled biogeochemical-EwE-F E2E modelling example was  
20 demonstrated (Section 4). However, this was done to demonstrate the feasibility of the  
21 approach and it does not mean that EwE-F can be applied only in E2E modelling frameworks.  
22 As discussed in section 5.1.4, many other exploitations of EwE-F are possible.

23 EwE-F is still in its infancy and future development efforts will focus on maturing the  
24 software and implementing missing useful features like times series fitting via vulnerability  
25 search, capability to define multiple fishing fleets and explicit spatial simulation. We believe  
26 that the development pace of EwE-F will accelerate with the adoption and utilisation of the  
27 software in the scientific community.

28

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17

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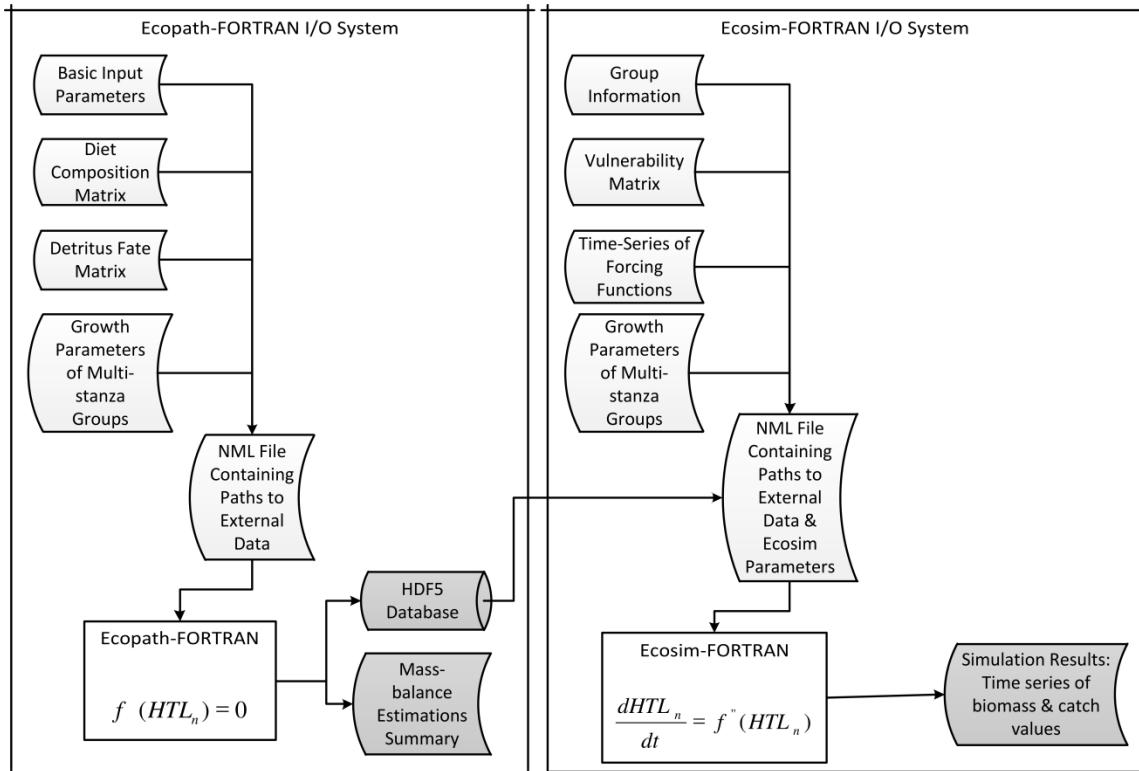
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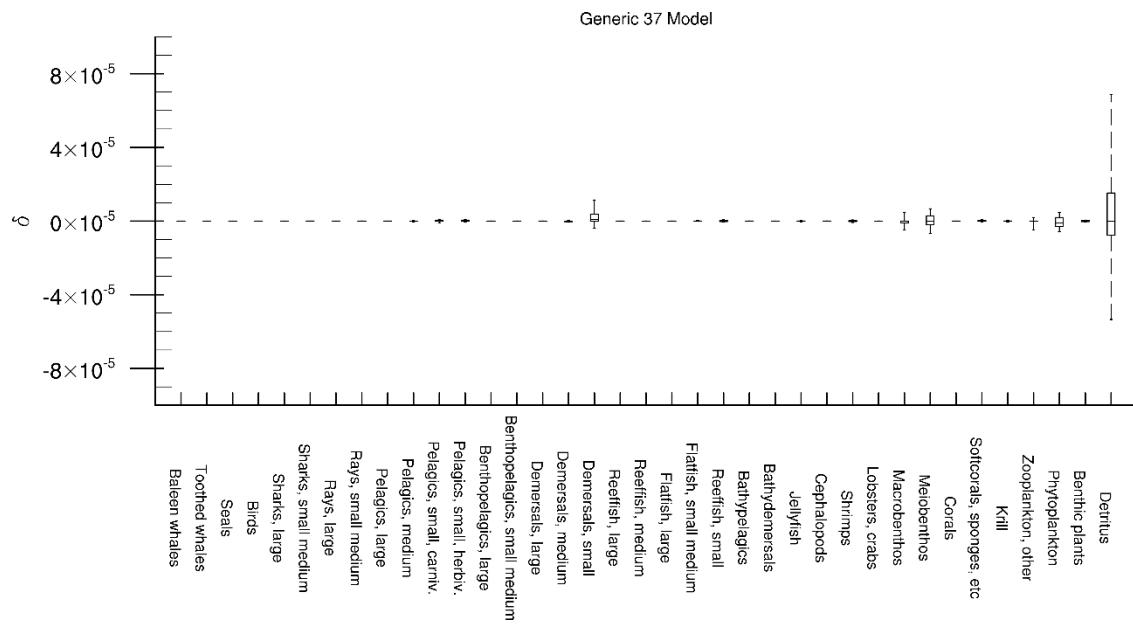
1 Table 1. General system and software related requirements of EwE-F v1.0.

<b>Name</b>	EwE-F (Ecopath with Ecosim in Fortran)
<b>Operating Systems</b>	Unix-like operating systems (Linux, *BSD, Mac OS X) and Microsoft Windows
<b>Processor</b>	Intel or AMD x86 processor
<b>Disk Space</b>	30 MB
<b>Compiler</b>	Fortran 95/2003 standards compliant compiler (e.g. GNU Fortran, Intel® Fortran Compiler, PGI® Fortran, Oracle® Solaris Studio, Absoft® Pro Fortran Compiler)
<b>Version Control System</b>	GIT (optional, for version controlled development)
<b>Building</b>	GNU Make (only required for building on Unix-like systems)
<b>Required External Libraries</b>	HDF5 version 1.8.11 or above
<b>License</b>	GNU Public License (GPL) version 2
<b>Homepage</b>	<a href="https://bitbucket.org/ewe-f">https://bitbucket.org/ewe-f</a>
<b>Obtaining and Documentation</b>	supporting information (SI) “EwE-F User’s Manual”

2

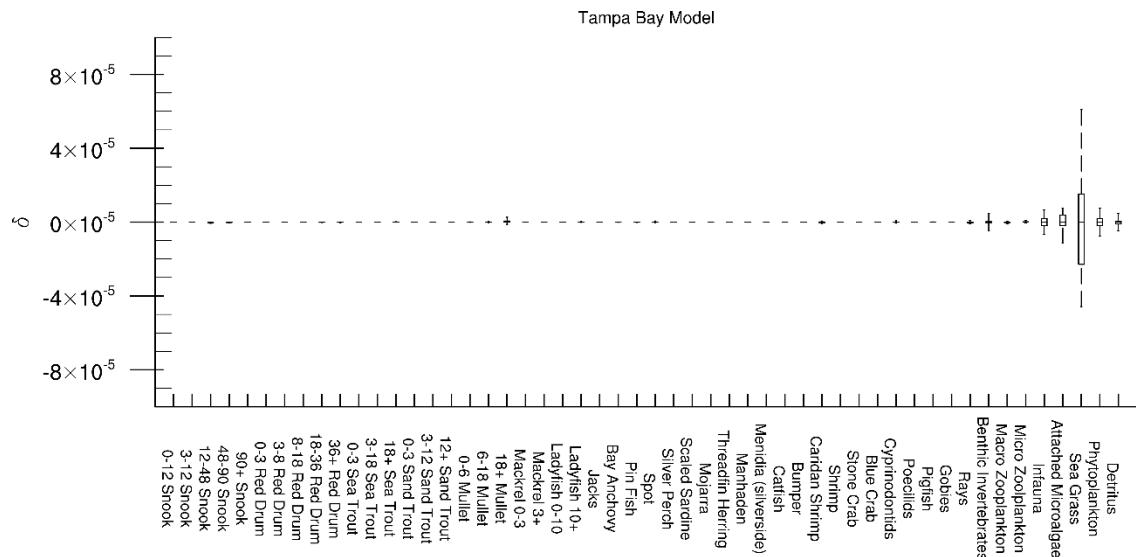


1      Figure 1. The EwE-F data input/output scheme. Curved white rectangular boxes denote tab-  
2      delimited ASCII files providing external data input to the EwE-F models (rectangles). Curved  
3      grey-shaded rectangles and the cylindrical box denote the model output via tab-delimited  
4      ASCII and HDF5 files respectively. For details see sections 3.1 and 3.2.  
5  
6



1  
2 Figure 2. The residuals between absolute biomasses simulated by EwE 6.5 and EwE-F 1.0 for  
3 the Generic 37 model. X-axis denotes all state variables in the model.  
4

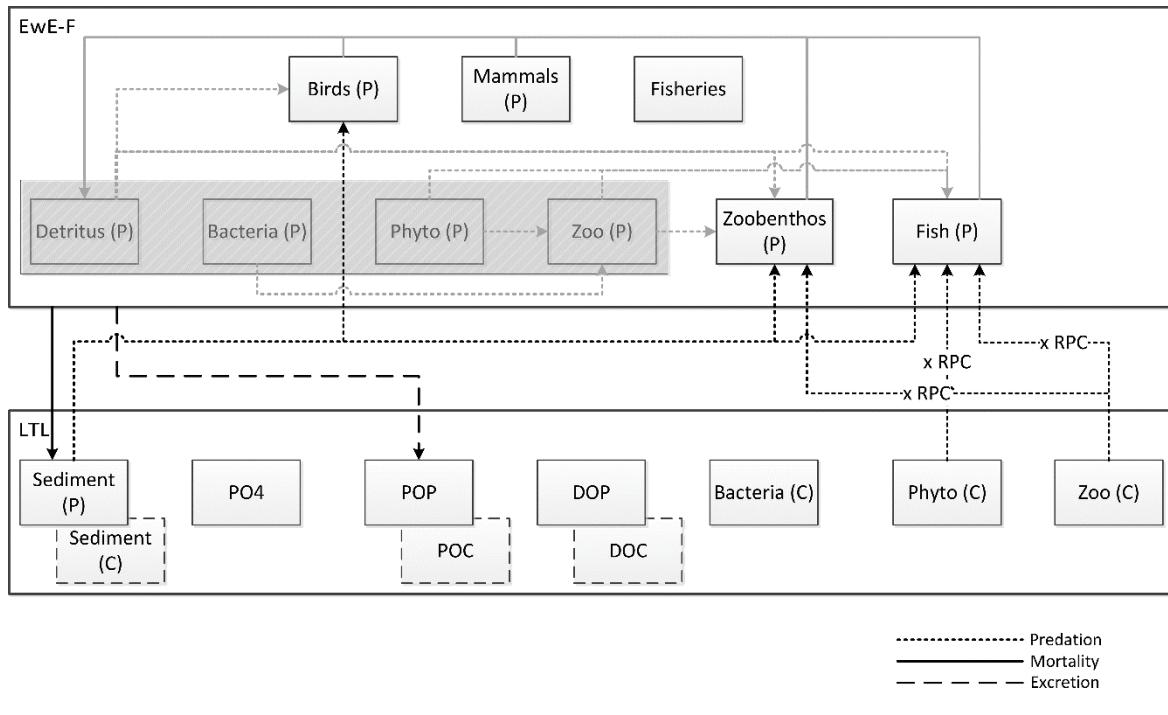
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3 Figure 3. The residuals between absolute biomasses simulated by EwE 6.5 and EwE-F 1.0 for  
4 the Tampa Bay model. X-axis denotes all state variables in the model.

5

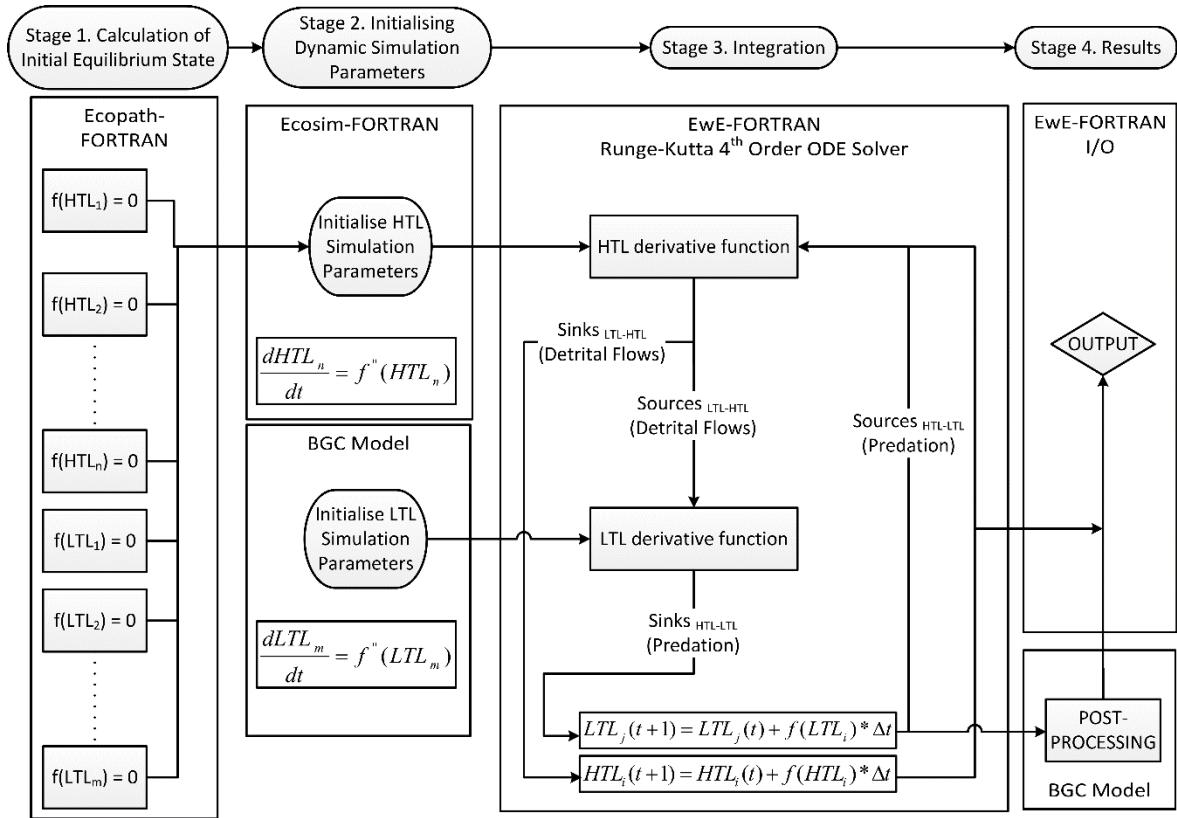


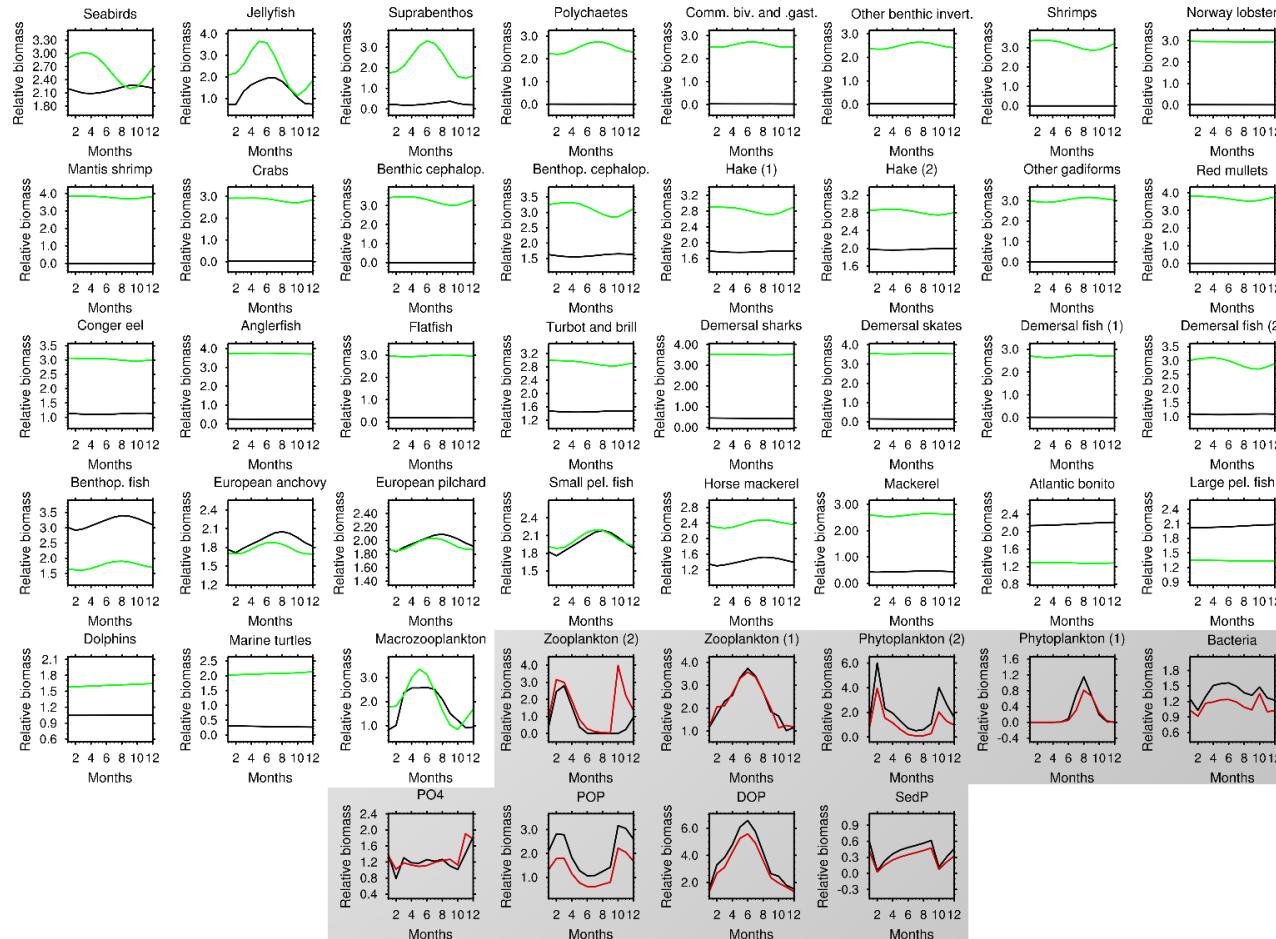
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2 Figure 4. Coupled trophodynamic model scheme of the Gulf of Trieste (Northern Adriatic  
 3 Sea) showing the linkages between the HTL and LTL models. Phosphorus (denoted with P)  
 4 was used as the currency for all of the HTL state variables and flows linking the two models.  
 5 Flows originating from the state variables of the LTL model, which were expressed in carbon  
 6 (denoted with C); i.e. phytoplankton and zooplankton, to the HTL model were converted to  
 7 phosphorus (by multiplying variable-specific phosphorus to carbon (RPC) ratios) before being  
 8 transferred. Grey-shaded state variables and flows in the HTL model were replaced by the  
 9 LTL model's corresponding state variables and the new linked flows are shown in black  
 10 dashed and continuous lines. Abbreviations: Zoo (small and large zooplankton groups), Phyto  
 11 (small and large phytoplankton groups), PO4 (phosphate), POP (particulate organic  
 12 phosphorus), DOP (dissolved organic phosphorus).

13

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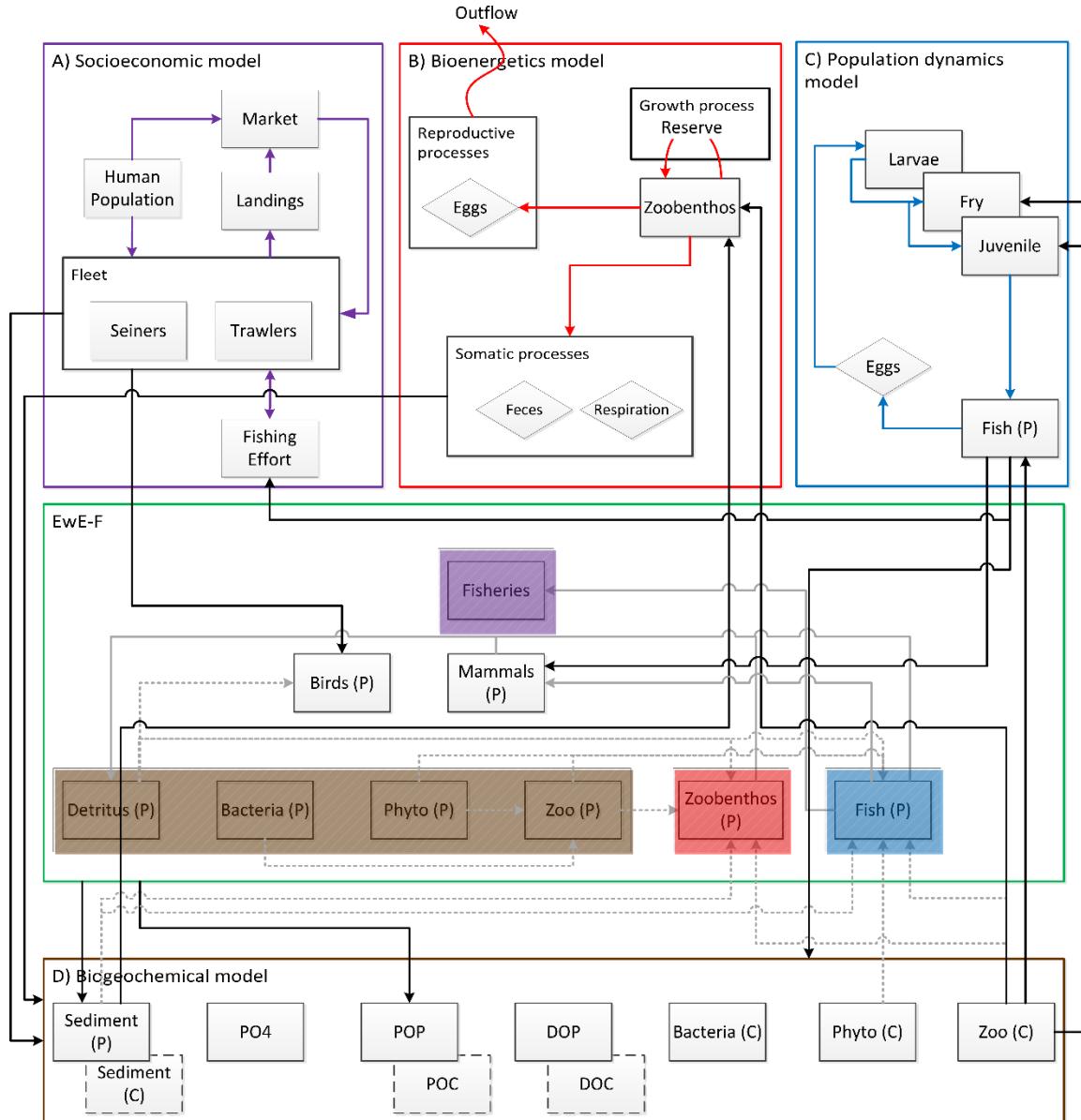




1

2 Figure 6. Monthly results of the final year in a 10-year simulation of the coupled (black lines) model versus simulations of uncoupled EwE 6.5  
 3 (green lines for HTL variables – unshaded boxes) and uncoupled biogeochemical (red lines for LTL variables – grey shaded boxes) models.

1



2

3 Figure 7. Potentialities provided by the EwE-F approach. Coloured arrows denote flows  
 4 specific to the integrating Fortran models. Black arrows denote linking flows and grey-shaded  
 5 arrows denote flows replaced/augmented by the linking flows. Boxes denoted by the letters A,  
 6 B, C and D and bordered by coloured lines replace the respective colour-shaded regions in the  
 7 EwE-F box (bordered green) under the coupling/integration scheme.

8