

A Review of Inputs and Outputs of Fisheries Tools

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1. Ecology and fisheries simulators

1.1 Atlantis

- Use Atlantis as an example. For more details on using Atlantis see this link.
- Atlantis model structure
 - Geometry submodel
 - * Description of model box shapes
 - Physics submodel
 - * Physical processes parameters
 - * Initial concentrations and distributions of model variables
 - * Hydrodynamic forcing of the transport model
 - * Environmental drivers such as temperature, salinity, nutrient inputs, river flows, and contaminants
 - Biology submodel
 - * Initial concentrations and distributions of model variables
 - * List of functional groups and their key biological characteristics
 - * Biological interaction parameters
 - * Biological drivers such as recruitment, extra mortality, and dispersal
 - Harvest submodel
 - * List of fisheries and their key characteristics
 - * Human impacts parameters
 - * Catches and discards
 - Economics submodel
 - * Socio-economic interaction parameters
 - * prices

- The structure of input files (* means essential files)

##	levelName	description
## 1	Inputs	
## 2	--catchts	
## 3	--boundary.ts	Blank catch file for use with boundary boxes
## 4	--catch1.ts	Historical catch time series file for box 1
## 5	--catch2.ts	Historical catch time series file for box 2
## 6	--catch3.ts	Historical catch time series file for box 3
## 7	--catch4.ts	Historical catch time series file for box 4
## 8	--catch5.ts	Historical catch time series file for box 5
## 9	--essential_optional_files	
## 10	--Biology.prm	*Set up ecological parameters, etc
## 11	--Fisheries.csv	*Fisheries name, recreation info, etc
## 12	--Forcings.prm	*Lays out the forcing file pathways
## 13	--Functional_groups.csv	*Biological, fishing, and management info
## 14	--Geography.bgm	*Geometry of boxes
## 15	--Harvest.prm	Fisheries and management parameters
## 16	--Initial_condition.cdf	Convert .nc to readable and editable file
## 17	--Initial_condition.nc	*Initial biomass and age structure
## 18	--Physics.prm	*Physics parameters
## 19	--Run_settings.prm	*Time setup, run and stop time, etc
## 20	--forcisets	
## 21	--pCO2RCP85.ts	pCO2 data time series
## 22	--SETAS_pH.nc	pH data for physics submodel
## 23	--SETAS_VMPAArsenic.nc	Contaminants tracer of functional groups
## 24	--SETAS_VMPAhydroA.nc	*Hydrodynamics forcing input file
## 25	--SETAS_VMPAhydroB.nc	*Hydrodynamics forcing input file
## 26	--SETAS_VMPAhydroC.nc	*Hydrodynamics forcing input file
## 27	--SETAS_VMPAhydroD.nc	*Hydrodynamics forcing input file
## 28	--SETAS_VMPAhydroE.nc	*Hydrodynamics forcing input file
## 29	--SETAS_VMPAOxygen.cdf	Convert .nc to readable and editable file
## 30	--SETAS_VMPAOxygen.nc	Oxygen data
## 31	--SETAS_VMPAsalt.nc	Salinity data
## 32	--SETAS_VMPAtemp.cdf	Convert .nc to readable and editable file
## 33	--SETAS_VMPAtemp.nc	Temperature data
## 34	--solar9195.ts	Solar radiation data
## 35	--meteor	
## 36	--84-95.nc	Evaporation data
## 37	--times1.6.nc	Precipitation data

- The structure of output files

	levelName	description
##		
## 1	Outputs	
## 2	--Biology.xml	Readable converted input file
## 3	--BiomIndx.txt	Biomass in tonnes of each species
## 4	--delete_to_halt_run	Write outputs to date
## 5	--export.ts	Net mass changes
## 6	--Fisheries.xml	Readable converted input file
## 7	--Functional_groups.xml	Readable converted input file
## 8	--Geography.bgm	Box model geometry
## 9	--Harvest.xml	Readable converted input file
## 10	--implementation.xml	Readable converted input file
## 11	--inputs.ts	Total tracer inputs
## 12	--log.txt	Messages generated during the run
## 13	--outputs.nc	3D snapshots of the tracer values
## 14	--outputs_BrokenStick.txt	Broken stick harvest strategy indices
## 15	--outputs_management.xml	Readable converted input file
## 16	--outputsAgeBiomIndx.txt	Biomass of all species per age group
## 17	--outputsAnnualAgeBiomIndx.txt	Annual biomass index per age
## 18	--outputsBiomIndx.txt	Biomasses of all active species
## 19	--outputsBiomReg.txt	Similar to BiomIndx outputs?
## 20	--outputsBoxBiomass.txt	Biomass of all species per box
## 21	--outputsCATCH.nc	Landing per species per fishery
## 22	--outputsCatch.txt	Landing per species over all fisheries
## 23	--outputsCatchPerFishery.txt	Landing per species per fishery
## 24	--outputsDietCheck.txt	Indication of diet pressure
## 25	--outputsDiscard.txt	Discards per species over all fisheries
## 26	--outputsDiscardPerFishery.txt	Discards per species per fishery
## 27	--outputsEffort.txt	Number of days per fishery for fishing
## 28	--outputsHarvestIndx.txt	Fisheries and management indices
## 29	--outputsMigration.txt	Migrating biomass per adults & juveniles
## 30	--outputsMort.txt	Natural and fishing mortality per species
## 31	--outputsMortPerPred.txt	Mortality of species by each predator?
## 32	--outputsPredPropCheck.txt	Indication of predation?
## 33	--outputsPROD.nc	2D snapshots of diagnostic tracers
## 34	--outputsRecCatch.txt	Recreational catch per species
## 35	--outputsSpecificMort.txt	3 values of instantaneous mortality
## 36	--outputsSpecificPredMort.txt	Mortality of species by each predator
## 37	--outputsSSB.txt	Mature biomass at spawning time
## 38	--outputsTAC.txt	TAC per species per year
## 39	--outputsTOT.nc	2D snapshots of tracers per box
## 40	--outputsTOTCATCH.nc	Total catch per species in tonnes
## 41	--outputsVertSize.txt	Sizes of migrating age structured group
## 42	--outputsYOY.txt	Biomass of recruits at spawning time
## 43	Run_settings.xml	Readable converted input file

2. Stock assessment operating tools

2.1 Atlantisom

- Use Atlantisom as an example. For more details on using Atlantisom see this link.
- Atlantisom model structure
 - Import simulated truth form Atlantis
 - * Get biomass, abundance, age composition, length composition, weight-at-age for each species
 - * Get fishery catch, catch-at-age, fishery length composition, and fishery weight-at-age for each species
 - Specify uncertainty in observed data
 - * Survey specifications (e.g., timing and spatial coverage) and uncertainty (e.g., observation error and effective sample size)
 - * Fishery specifications (e.g., sampling areas) and uncertainty (e.g., observation error and effective sample size)
 - Get other life history parameters from Atlantis Biology.prm file
 - * Use weight-length relationship parameters from Atlantis to calculate h, R0
 - * Use natural mortality from Atlantis
 - * Use survey length-at-age from Atlantis to estimate growth curve
 - Format data to run stock assessment models
- The structure of input files (use ANNAGEBIO.nc and ANNAGECATCH.nc if available)

```
##                                         levelName          description
## 1  Inputs
## 2  |--Biology.prm                      Set up ecological parameters, etc
## 3  |--BiomIndx.txt                     Biomass in tonnes of each species
## 4  |--Fisheries.csv                    Fisheries name, recreation info, etc
## 5  |--Functional_groups.csv           Biological, fishing, and management info
## 6  |--Geography.bgm                   Geometry of boxes
## 7  |--Initial_condition.nc            Initial biomass and age structure
## 8  |--outputs.nc                      3D snapshots of the tracer values
## 9  |--outputsCATCH.nc                 Landing per species per fishery
## 10 |--outputsCatch.txt                Landing per species over all fisheries
## 11 |--outputsCatchPerFishery.txt      Landing per species per fishery
## 12 |--outputsDietCheck.txt            Indication of diet pressure
## 13 |--outputsPROD.nc                  2D snapshots of diagnostic tracers
## 14 |--outputsYOY.txt                 Biomass of recruits at spawning time
## 15 |--Run_settings.xml               Time setup, run and stop time, etc
```

- The structure of output files

	levelName	description
##		
## 1 Outputs		
## 2 --outputscatchage_census.rds		Catch at age per species per time
## 3 --outputscaughtlengthwt_census.rds		Len and wt at age, num at length
## 4 --outputscatchN_census.rds		Catch number per species per time
## 5 --outputslengthwt_census.rds		Len and wt at age, num at length
## 6 --outputsnatage_census.rds		Survey num at age per species per time
## 7 --outputsrun_truth.RData		Atlantis true outputs (biomass etc)
## 8 --outputssurveyBcensus.rds		Survey biomass per species per time
## 9 °--outputssurveyNcensus.rds		Survey abundance per species per time

3. Stock assessment tools

3.1 Stock Synthesis (SS)

- Use SS as an single-species assessment example. For more details on using SS see this link.
- SS structure

3.2 Globally applicable Area Disaggregated General Ecosystem Toolbox (Gadget)

- Use Gadget as an multi-species assessment example. For more details on using Gadget see these links:
 - gadget2
 - rgadget
 - gadget courses
- Gadget structure
 - Specify model years, time steps, areas, temperature of areas
 - Provide information of stocks
 - * Biological information such as minage and minlength
 - * Key components such as growth model, natural mortality for each age group, predator-prey determination, initial condition, migration, maturity, movement, renew, spawn, and stray information
 - * Predation is modelled in terms of size and it includes consumption and cannibalistic relationships
 - * Recruitment can be modeled as result of spawning and independent renewal
 - Add human interactions as removals of stock
 - * Provide fleet type,selectivity, and catch data
- The structure of input files

```
##                                         levelName           description
## 1  Inputs
## 2  |--Aggfiles
## 3  |  |--lingimm.stock.len.agg          Bounds of length class
## 4  |  `--lingmat.stock.len.agg          Bounds of length class
## 5  |--Data
## 6  |  |--fleet.bmt.data                Catch weight per time per area
## 7  |  |--fleet.foreign.data             Catch weight per time per area
## 8  |  |--fleet.gil.data                Catch weight per time per area
## 9  |  |--fleet.lln.data                Catch weight per time per area
## 10 |  `--fleet.surv.data               Survey number per time per area
## 11 |--lingimm                         Biological information of stock
## 12 |--lingmat                          Biological information of stock
## 13 |--main                            File paths
## 14 |--Modelfiles
## 15 |  |--area                           Area and temperature per time step
## 16 |  |--fleet                          Fleet selectivity information etc
## 17 |  |--lingimm.init.normalparam      Initial condition equations
## 18 |  |--lingimm.maturity              Maturity parameter values
## 19 |  |--lingimm.refwgt              Weight at length class
## 20 |  |--lingmat.init.normalparam      Initial condition equations
## 21 |  |--lingmat.refwgt              Weight at length class
## 22 |  `--time                           Year, time step and its duration
## 23 `--params.forsim                  Parameter values for simulation
```

- The structure of output files

	levelName	description
##		
## 1	Outputs	
## 2	--FIT	
## 3	--main.print	File paths
## 4	--out.fit	
## 5	--lingimm.full	Num and wt of stock by length class
## 6	--lingimm.prey	Num and wt caught by all fleets
## 7	--lingimm.prey.bmt	Num and wt caught by one fleet
## 8	--lingimm.prey.foreign	Num and wt caught by one fleet
## 9	--lingimm.prey.gil	Num and wt caught by one fleet
## 10	--lingimm.prey.lln	Num and wt caught by one fleet
## 11	--lingimm.prey.surv	Num and wt caught by one fleet
## 12	--lingimm.recruitment	Num and wt of recruits by length and step
## 13	--lingimm.std	Biological info by age, step, and area
## 14	--lingmat.full	Num and wt of stock by length class
## 15	--lingmat.prey	Num and wt caught by all fleets
## 16	--lingmat.prey.bmt	Num and wt caught by one fleet
## 17	--lingmat.prey.foreign	Num and wt caught by one fleet
## 18	--lingmat.prey.gil	Num and wt caught by one fleet
## 19	--lingmat.prey.lln	Num and wt caught by one fleet
## 20	--lingmat.prey.surv	Num and wt caught by one fleet
## 21	--lingmat.recruitment	Num and wt of recruits by length and step
## 22	°--lingmat.std	Biological info by age, step, and area
## 23	--print.aggfiles	
## 24	--lingimm.age.agg	Min to max age
## 25	--lingimm.allages.agg	Min to max age
## 26	--lingimm.alllen.agg	Min and max length classes
## 27	--lingimm.area.agg	Area
## 28	--lingimm.len.agg	Bounds of length classes
## 29	--lingimm.rec.age.agg	Recruitment age
## 30	--lingmat.age.agg	Min to max age
## 31	--lingmat.allages.agg	Min to max age
## 32	--lingmat.alllen.agg	Min and max length classes
## 33	--lingmat.area.agg	Area
## 34	--lingmat.len.agg	Bounds of length classes
## 35	°--lingmat.rec.age.agg	Recruitment age
## 36	--printfile.fit	File paths
## 37	--SS.print	List of parameters used in the run
## 38	°--WGTS.Rdata	Outputs in Rdata format
## 39	--ling_log	Log of model run
## 40	°--params.out	Estimation values of parameters

4. Forecasting tools

- Required inputs and forecasting outputs of potential tools
 - Gadget projections
 - * Need to set up stock biological parameterizations
 - * Need to keep original parameter settings or: adding a recruitment series; modifying reference points used to control fishing behavior, modifying harvest levels and errors
 - * Can produce projections of immature and mature portions of stocks and projections of each species in multispecies model
 - * Can project fleet and its response to a harvest control rule/modifications of fishing behavior (e.g., inputs or outputs regulation, split effort/quota among fleets)
 - * Can produce yield and spawning stock biomass over F
 - * Can produce biomass, abundance, and catch over years under different harvest rates
- Capabilities of incorporating climate, ecosystem, and economic considerations
- Scenarios to explore and comparison criteria

5. Diagnostic tools