

DLMtool CHEAT SHEET 1



Getting Started

Install package: `install.packages('DLMtool')`

User Guide: `userguide()`

Website: <http://datalimitedtoolkit.org>

Report Issues: <https://github.com/DLMtool/DLMtool/issues>

Acronyms

DLMtool	Data-Limited Methods Toolkit
MP	Management Procedure
MSE	Management Strategy Evaluation
OM	Operating Model
PM	Performance Metrics

Main Object Classes

Class	Contents
Stock	Biological Properties
Fleet	Exploitation Properties
Obs	Observation Error
Imp	Implementation Error
OM	Operating Model
MSE	Management Strategy Evaluation Results
MP	Management Procedure

Find Available Objects: `avail('Object Class')`

e.g. `avail('Stock')`
 `avail('Fleet')`
 `avail('Obs')` **More Objects:** `DLMextra()`
 ...
 `avail('MP')`

Slot Names: `slotNames('Object Class')`

e.g. `slotNames('Stock')`

Create New OM

Blank OM: `OM <- new('OM')`

New OM from available objects: `OM <- new('OM', 'Stock', 'Fleet', 'Obs', 'Imp')`

e.g. `OM <- new('OM', Albacore, Generic_Fleet, Generic_Obs, Perfect_Imp)`

Initialize Excel OM and OM Report: `OMinit('myOM')`

Import OM from Excel: `OM <- XL2OM('myOM')`

Generate OM Report: `OMdoc()`

Customize OM

Sketch Historical Fishing: `Fleet/OM <- ChooseEffort(Fleet/OM)`

Sketch Selectivity: `Fleet <- ChooseSelect(Fleet, FstYr = ...)`

Sketch Age Specific *M*: `OM <- ChooseM(OM)`

Sketch Length Specific *M*: `OM <- ChooseM(OM, 'Length')`

Predicting Life-History Parameters:

e.g. `OM <- new('OM')`
 `OM@Species <- 'Scomber japonicus'`
 `OM <- LH2OM(OM)`

Custom Parameters:

e.g. `OM <- new('OM', Albacore, Generic_Fleet, Generic_Obs, Perfect_Imp)`
 `OM@cpars$M <- rlnorm(OM@nsim, log(0.2), 0.05)`

Remove Process and Observation Error: `OM <- tinyErr(OM)`

Replace OM Component: `OM <- Replace(OM, Blue_shark)`

Examine OM

Plot OM Components:

e.g. `plot(Albacore)`
 `plot(Generic_Fleet)`

Plot OM: `plot(OM)`

Plot Existing MPA: `plotMPA(OM)`

Plot *M*: `plotM(OM)`

Plot Selectivity: `plotSelect(OM)`

OM Excel: myOM.xlsx

	A	B	C	D
1	Slot			
2	Name			
3	Common_Name			
4	Species			
5	maxage			
6	R0			
7	M			
8	M2			
9	Mexp			
10	Msd			
11	Mgrad			
12	h			
13	SRrel			
14	Perr			
15	AC			
16	Period			
17	Amplitude			
18	Linf			
19	K			
20	t0			
21	LenCV			
22	Ksd			
23	Kgrad			
24	Linfsd			
25	Linfgrad			
26	L50			
27	L50_95			
28	D			
29	a			
30	b			
31	Size_area_1			
32	Frac_area_1			
33	Prob_staying			
34	Fdisc			
35	Source			
36				

OM Report: myOM.rmd

1	# Title
2	Include name and Location of the Fishery. One line only.
3	
4	# Subtitle
5	Subtitle, one line only.
6	
7	# Author(s)
8	Name and contact details (e.g email, affiliation) for each author.
9	One line per author.
10	
11	# Date
12	Optional, date that the operating model was created. If none provided, today's date will be used.
13	
14	# Introduction
15	
16	
17	## Completing the OM Documentation
18	This document is used to generate a HTML OM report document.
19	
20	The document is separated into 7 sections:
21	1. Introduction (this section)
22	2. Custom Parameters (optional)
23	3. Stock Parameters
24	4. Fleet Parameters
25	5. Obs (Observation) Parameters
26	6. Imp (Implementation) Parameters
27	7. References
28	
29	The Introduction section is used to briefly describe the fishery and the details of the operating Model.
30	It should include an explanation for the OM parameters:
31	* nsim: the number of simulations.
32	* proyears: the number of projectio years.
33	* interval: the management interval.
34	* pstar: the percentile of the sample of the management recommendation for each method.
35	* mmaxf: the maximum instantaneous fishing mortality rate that may be simulated for any given age class.
36	* reps: the number of samples of the management recommendation for each method.
37	

Management Procedures

MP Type	Returns
Output	TAC (total allowable catch)
Input	TAE, SL, Spatial (total allowable effort, size limit, spatial closure)
Mixed	Combination of Output and Input
Reference	TAC (assuming perfect data)

Find MP type: `MPtype()`

e.g. `MPtype(c('AvC', 'curE', 'matlenlim', 'FMSYref'))`

Run MSE

Run MSE: `runMSE()`

e.g. `MSE <- runMSE(OM, MP=c('AvC', 'curE', 'matlenlim', 'FMSYref'))`

Run MSE in parallel: `runMSE(parallel=TRUE)`

e.g. `MSE <- runMSE(OM, MP=c('AvC', 'curE', 'matlenlim', 'FMSYref'), parallel=TRUE)`

Check Convergence: `Converge(MSE)`

Run Historical Simulations:

`Hist <- runMSE(OM, Hist=TRUE)`

