

Package ‘R4MFCL’

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Title R functions for MFCL

Version 0.1

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Description miscellaneous funtions for dealing with input to and output from MFCL

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add.catch.frq	<i>add.catch.frq.Rd</i>
---------------	-------------------------

Usage

```
add.catch.frq(frq, filepath, fshries)
```

Arguments

```
frq
filepath
fshries
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

add.cpue.frq	<i>add.cpue.frq.Rd</i>
--------------	------------------------

Usage

```
add.cpue.frq(CPUE.file = "P:/yft/2009/Data Preparation/CPUE/indices/yft_JPstd_R1
```

Arguments

```
CPUE.file
data
fishery
add.cv
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

add.flag	<i>add.flag.Rd</i>
----------	--------------------

Usage

```
add.flag(doitall, flagtype, flagnum, newval, phase)
```

Arguments

```
doitall  
flagtype  
flagnum  
newval  
phase
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

alb.clean.lfdata	<i>alb.clean.lfdata.Rd</i>
------------------	----------------------------

Usage

```
alb.clean.lfdata(infrq)
```

Arguments

```
infrq
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

```
alb.initial_clean_2008
      alb.initial_clean_2008.Rd
```

Usage

```
alb.initial_clean_2008(frq.obj)
```

Arguments

```
frq.obj
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
alb.initial_clean_2009
      alb.initial_clean_2009.Rd
```

Usage

```
alb.initial_clean_2009(frq.obj)
```

Arguments

```
frq.obj
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
alb.initial_clean_2011  
    alb.initial_clean_2011.Rd
```

Usage

```
alb.initial_clean_2011(frq.obj)
```

Arguments

```
freq.obj
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

```
alb.pagocatch_2009 alb.pagocatch_2009.Rd
```

Usage

```
alb.pagocatch_2009(frq.obj)
```

Arguments

```
freq.obj
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

```
carry.effort.frq    carry.effort.frq.Rd
```

Usage

```
carry.effort.frq(data = out.data, fishery = 1, last = 2008)
```

Arguments

```
data
fishery
last
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

```
change.fishflag    change.fishflag.Rd
```

Usage

```
change.fishflag(a, fisheries, flagnum, newvals)
```

Arguments

```
a
fisheries
flagnum
newvals
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

change.flag	<i>change.flag.Rd</i>
-------------	-----------------------

Usage

```
change.flag(doitall, flagtype, flagnum, newval)
```

Arguments

```
doitall
flagtype
flagnum
newval
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

change.negflag	<i>change.negflag.Rd</i>
----------------	--------------------------

Usage

```
change.negflag(doitall, flagtype, flagnum, newval)
```

Arguments

```
doitall
flagtype
flagnum
newval
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

change.size.frq	<i>change.size.frq.Rd</i>
-----------------	---------------------------

Usage

```
change.size.frq(ver = 6, data = data, FISH = 1, LF.FILE = "P:/yft/2009/Data Prep
```

Arguments

```
ver
data
FISH
LF.FILE
WT.FILE
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

change_data	<i>change_data.Rd</i>
-------------	-----------------------

Usage

```
change_data(obj, searchtext, xlines, newline)
```

Arguments

```
obj
searchtext
xlines
newline
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

check.eff.devs	<i>check.eff.devs.Rd</i>
----------------	--------------------------

Usage

```
check.eff.devs(parfile, repfile, frqfile, parlim = 5.9)
```

Arguments

```
parfile  
repfile  
frqfile  
parlim
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

check_flag_value	<i>check_flag_value.Rd</i>
------------------	----------------------------

Usage

```
check_flag_value(parname, flagtype, flagnums, fishery = NA, flaglist = T)
```

Arguments

```
parname  
flagtype  
flagnums  
fishery  
flaglist
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

clean.lfdata	<i>clean.lfdata.Rd</i>
--------------	------------------------

Usage

```
clean.lfdata(infrq)
```

Arguments

```
infrq
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

compare.ce.frq	<i>compare.ce.frq.Rd</i>
----------------	--------------------------

Usage

```
compare.ce.frq(file1, file2, fm, plotname, fdesc = "")
```

Arguments

```
file1
```

```
file2
```

```
fm
```

```
plotname
```

```
fdesc
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

compare.frq	<i>compare.frq.Rd</i>
-------------	-----------------------

Usage

```
compare.frq(file1, file2, fm = "all", plotname, fdesc = "")
```

Arguments

```
file1
file2
fm
plotname
fdesc
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

compare.size.frq	<i>compare.size.frq.Rd</i>
------------------	----------------------------

Usage

```
compare.size.frq(frq1, frq2, fishery = 5)
```

Arguments

```
frq1
frq2
fishery
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
compare.size.frq2  compare.size.frq2.Rd
```

Usage

```
compare.size.frq2(frq1, frq2, fishery = 5, wt = T)
```

Arguments

```
frq1
frq2
fishery
wt
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

```
compare_par_flags  compare_par_flags.Rd
```

Usage

```
compare_par_flags(par1, par2, flaglist = T)
```

Arguments

```
par1
par2
flaglist
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

`condor.go`*condor.go.Rd*

Usage

```
condor.go(run.dir, frq.obj, tag.obj, doitall.obj, ini.obj, sub.obj = suball, spe
```

Arguments

```
run.dir
frq.obj
tag.obj
doitall.obj
ini.obj
sub.obj
species
condor_f
par.obj
run_now
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

`condor.go2`*condor.go2.Rd*

Usage

```
condor.go2(run.dir, frq.obj, tag.obj, doitall.obj, ini.obj, sub.obj = suball, sp
```

Arguments

```
run.dir
frq.obj
tag.obj
doitall.obj
ini.obj
sub.obj
species
condor_f
par.obj
run_now
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
Convert.frq.ver6    Convert.frq.ver6.Rd
```

Usage

```
Convert.frq.ver6(a)
```

Arguments

```
a
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
create.missing.ce    create.missing.ce.Rd
```

Usage

```
create.missing.ce(data = data, year = 2008)
```

Arguments

```
data
year
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

create.ver6.frq	<i>create.ver6.frq.Rd</i>
-----------------	---------------------------

Usage

```
create.ver6.frq(frq = base.frq)
```

Arguments

```
frq
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

crit.fishery.summary	<i>crit.fishery.summary.Rd</i>
----------------------	--------------------------------

Usage

```
crit.fishery.summary(crit)
```

Arguments

```
crit
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

crit.summary	<i>crit.summary.Rd</i>
--------------	------------------------

Usage

```
crit.summary(crit, years)
```

Arguments

```
crit  
years
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

datfromstr	<i>datfromstr.Rd</i>
------------	----------------------

Usage

```
datfromstr(datstring)
```

Arguments

```
datstring
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

do.critical.calcs	<i>do.critical.calcs.Rd</i>
-------------------	-----------------------------

Usage

```
do.critical.calcs(repfile = "P:/yft/2007/BaseYFT/yftfinal2007.rep", ests = "P:/y
```

Arguments

```
repfile
ests
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

doit.rm_flag	<i>doit.rm_flag.Rd</i>
--------------	------------------------

Usage

```
doit.rm_flag(a, flagtype, flag, value)
```

Arguments

```
a
flagtype
flag
value
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

effortcreep	<i>effortcreep.Rd</i>
-------------	-----------------------

Usage

```
effortcreep(frq.obj, fisheries, creep)
```

Arguments

```
freq.obj
fisheries
creep
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

fix_growth	<i>fix_growth.Rd</i>
------------	----------------------

Usage

```
fix_growth(a)
```

Arguments

```
a
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

frq.change.nint *frq.change.nint.Rd*

Usage

```
frq.change.nint(in.frq, add.lfint, add.wfint)
```

Arguments

```
in.frq
add.lfint
add.wfint
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

frq.remove.size.or.weight.data
frq.remove.size.or.weight.data.Rd

Usage

```
frq.remove.size.or.weight.data(data = test.data, exclude = exclude, size = T, we
```

Arguments

```
data
exclude
size
weight
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
get.critical.age    get.critical.age.Rd
```

Usage

```
get.critical.age(data = Base.rep)
```

Arguments

data

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

```
get.length.output    get.length.output.Rd
```

Usage

```
get.length.output(REGION = 1, DIR = "P:/yft/2009/Data Preparation/size data/")
```

Arguments

REGION

DIR

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

get.outcomes	<i>get.outcomes.Rd</i>
--------------	------------------------

Usage

```
get.outcomes(file.rep, file.par, nofish = T)
```

Arguments

```
file.rep  
file.par  
nofish
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

get.outcomes.alb	<i>get.outcomes.alb.Rd</i>
------------------	----------------------------

Usage

```
get.outcomes.alb(file.rep, file.par, catch.rep, nofish = T, SBcurr_1yr = F)
```

Arguments

```
file.rep  
file.par  
catch.rep  
nofish  
SBcurr_1yr
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
get.outcomes.test  get.outcomes.test.Rd
```

Usage

```
get.outcomes.test(file.rep, file.par, catch.rep, nofish = T)
```

Arguments

```
file.rep
file.par
catch.rep
nofish
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==>  Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

```
get.weight.output  get.weight.output.Rd
```

Usage

```
get.weight.output(REGION = 1, DIR = "P:/yft/2009/Data Preparation/size data/")
```

Arguments

```
REGION
DIR
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==>  Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

initial_clean_2008 *initial_clean_2008.Rd*

Usage

```
initial_clean_2008(frq.obj)
```

Arguments

```
freq.obj
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

labels_store *labels_store.Rd*

Usage

```
labels_store(labelfile = basecase.labels)
```

Arguments

```
labelfile
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

load.LFdata *load.LFdata.Rd*

Usage

```
load.LFdata(species = "ALB", gear = "L")
```

Arguments

```
species  
gear
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
make.projection.betyft.frq
      make.projection.betyft.frq.Rd
```

Usage

```
make.projection.betyft.frq(frq.in = base.frq, fish = 1:24, years = 10)
```

Arguments

```
frq.in
fish
years
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
map_all_pacific      map_all_pacific.Rd
```

Usage

```
map_all_pacific(plot_title = "", lims = c(100, 300, -45, 45), eezfile = "L:/alb/
```

Arguments

```
plot_title
lims
eezfile
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

`merge.frq`*merge.frq.Rd*

Usage

```
merge.frq(frq.obj, oldf, newf, mergelf = FALSE)
```

Arguments

```
frq.obj
```

```
oldf
```

```
newf
```

```
mergelf
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

`merge.tag`*merge.tag.Rd*

Usage

```
merge.tag(tag.obj, oldf, newf)
```

Arguments

```
tag.obj
```

```
oldf
```

```
newf
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

merge_tag_objs	<i>merge_tag_objs.Rd</i>
----------------	--------------------------

Usage

```
merge_tag_objs(obj1, obj2, relgrps)
```

Arguments

```
obj1  
obj2  
relgrps
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

NZtrollglm	<i>NZtrollglm.Rd</i>
------------	----------------------

Usage

```
NZtrollglm(frq.obj, fishery, coef.file)
```

Arguments

```
frq.obj  
fishery  
coef.file
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

oldsetup	<i>oldsetup.Rd</i>
----------	--------------------

Usage

```
oldsetup(olddbasedir)
```

Arguments

```
olddbasedir
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

pack.fisheries.frq	<i>pack.fisheries.frq.Rd</i>
--------------------	------------------------------

Usage

```
pack.fisheries.frq(frq.obj)
```

Arguments

```
frq.obj
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

pagocatch	<i>pagocatch.Rd</i>
-----------	---------------------

Usage

```
pagocatch(frq.obj)
```

Arguments

```
frq.obj
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
plot.base.comparison
      plot.base.comparison.Rd
```

Usage

```
plot.base.comparison(baseres, labs)
```

Arguments

```
baseres
labs
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
plot.biomass      plot.biomass.Rd
```

Usage

```
plot.biomass(plotdir = "H:/rmfcl/test/figs/", plotrep = test, varfile = NULL, ty
```

Arguments

```
plotdir
plotrep
varfile
type
plotname
plottype
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
plot.biomass.combined  
plot.biomass.combined.Rd
```

Usage

```
plot.biomass.combined(plotdir = "H:/rmfcl/test/figs/", plotrep = test, varfile =
```

Arguments

```
plotdir  
plotrep  
varfile  
type  
plotname  
plottype
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
plot.F.time plot.F.time.Rd
```

Usage

```
plot.F.time(plotdir = "H:/rmfcl/test/figs/", plotrep = "C:/assessments/alb/2008/
```

Arguments

```
plotdir  
plotrep  
inifile  
plotname  
plottype  
COL
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
plot.fishery.impact.r
      plot.fishery.impact.r.Rd
```

Usage

```
plot.fishery.impact.r(plotdir = "H:/rmfcl/test/figs/", type = "Total", plotrep =
```

Arguments

```
plotdir
type
plotrep
impnames
plotname
plottype
COL
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
plot.Kobe      plot.Kobe.Rd
```

Usage

```
plot.Kobe(plotdir = "S:/OFP Publications/Tuna Fishery Assessment Report/2007/Fig
```

Arguments

```
plotdir
plotrep
type
plotname
plottype
COL
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
plot.Kobe.template.bw  
plot.Kobe.template.bw.Rd
```

Usage

```
plot.Kobe.template.bw (Type)
```

Arguments

Type

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
plot.Kobe.template.col  
plot.Kobe.template.col.Rd
```

Usage

```
plot.Kobe.template.col (Type)
```

Arguments

Type

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
plot.mfcl.betyft09 plot.mfcl.betyft09.Rd
```

Usage

```
plot.mfcl.betyft09(lims = c(100, 260, -45, 45))
```

Arguments

```
lims
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
plot.nofishing plot.nofishing.Rd
```

Usage

```
plot.nofishing(plotdir = "H:/rmfcl/test/figs/", plotrep = testq0, type = "SSB",
```

Arguments

```
plotdir
plotrep
type
plotname
plottype
COL
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
plot.nofishing.combined  
    plot.nofishing.combined.Rd
```

Usage

```
plot.nofishing.combined(plotdir = "H:/rmfcl/test/figs/", plotrep = testq0, type
```

Arguments

```
plotdir  
plotrep  
type  
plotname  
plotype  
COL
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

```
plot.pacific.alb    plot.pacific.alb.Rd
```

Usage

```
plot.pacific.alb(plot_title = "", eez_dir = "I:/assessments/Pop dy modeling/MFCL
```

Arguments

```
plot_title  
eez_dir  
plot_eez
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

```
plot.pacific.skj    plot.pacific.skj.Rd
```

Usage

```
plot.pacific.skj(plot_title = "")
```

Arguments

```
plot_title
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

```
plot.pacific.species  
    plot.pacific.species.Rd
```

Usage

```
plot.pacific.species(plot_title = "", uselims = NA, add.WCPFC = F, add.EPO = F,
```

Arguments

```
plot_title  
uselims  
add.WCPFC  
add.EPO  
sp  
add.EEZ  
eez_file
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

`plot.pacific.WCPFC` *plot.pacific.WCPFC.Rd*

Usage

```
plot.pacific.WCPFC(plot_title = "", lims = c(100, 260, -45, 45))
```

Arguments

```
plot_title  
lims
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

`plot.pacific.yft` *plot.pacific.yft.Rd*

Usage

```
plot.pacific.yft(plot_title = "", lims = c(100, 260, -45, 45), add.WCPFC = F)
```

Arguments

```
plot_title  
lims  
add.WCPFC
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

plot.recruitment *plot.recruitment.Rd*

Usage

```
plot.recruitment(plotdir = "H:/rmfcl/test/figs/", plotrep = test, varfile = NULL)
```

Arguments

```
plotdir
plotrep
varfile
plotname
plottype
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

plot.recruitment.combined
 plot.recruitment.combined.Rd

Usage

```
plot.recruitment.combined(plotdir = "H:/rmfcl/test/figs/", plotrep = test, varfile = NULL)
```

Arguments

```
plotdir
plotrep
varfile
plotname
plottype
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

plot_cpue_cv_frq	<i>plot_cpue_cv_frq.Rd</i>
------------------	----------------------------

Usage

```
plot_cpue_cv_frq(frq, parf, fisheries)
```

Arguments

```
frq  
parf  
fisheries
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

read.catchrep	<i>read.catchrep.Rd</i>
---------------	-------------------------

Usage

```
read.catchrep(catchrep.file)
```

Arguments

```
catchrep.file
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

read.ests	<i>read.ests.Rd</i>
-----------	---------------------

Usage

```
read.ests(rep.obj, ests = "C:/assessments/alb/2008/6_area/28.splitgr3/ests.rep",
```

Arguments

```
rep.obj
```

```
ests
```

```
x
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

read.fit	<i>read.fit.Rd</i>
----------	--------------------

Usage

```
read.fit(fit.file)
```

Arguments

```
fit.file
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

read.frq	<i>read.frq.Rd</i>
----------	--------------------

Usage

```
read.frq(frq.file, frq.title = "", ntop = 0, fishdefs = NA)
```

Arguments

```
frq.file
frq.title
ntop
fishdefs
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

read.impact	<i>read.impact.Rd</i>
-------------	-----------------------

Usage

```
read.impact(impdir = "H:/rmfcl/test/", impnames = c("ll", "psassoc", "psunassoc"))
```

Arguments

```
impdir
impnames
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

read.ini	<i>read.ini.Rd</i>
----------	--------------------

Usage

```
read.ini(ini.file)
```

Arguments

```
ini.file
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

read.par	<i>read.par.Rd</i>
----------	--------------------

Usage

```
read.par(par.file)
```

Arguments

```
par.file
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

read.rep	<i>read.rep.Rd</i>
----------	--------------------

Usage

```
read.rep(rep.file)
```

Arguments

```
rep.file
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

read.tag	<i>read.tag.Rd</i>
----------	--------------------

Usage

```
read.tag(tagfile)
```

Arguments

```
tagfile
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

read.tags.JP	<i>read.tags.JP.Rd</i>
--------------	------------------------

Usage

```
read.tags.JP(tagfile, reg_areas, fisheries)
```

Arguments

```
tagfile  
reg_areas  
fisheries
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

read.var	<i>read.var.Rd</i>
----------	--------------------

Usage

```
read.var(var.file)
```

Arguments

```
var.file
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

read_nmd.frq	<i>read_nmd.frq.Rd</i>
--------------	------------------------

Usage

```
read_nmd.frq(frq.file, frq.title = "", ntop = 0, fishdefs = NA)
```

Arguments

```
frq.file
frq.title
ntop
fishdefs
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

read_nmd.par	<i>read_nmd.par.Rd</i>
--------------	------------------------

Usage

```
read_nmd.par(par.file)
```

Arguments

```
par.file
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
reconstruct.frq.ce reconstruct.frq.ce.Rd
```

Usage

```
reconstruct.frq.ce(CPUE.file = "X:/yft/2009/Data Preparation/CPUE/indices/yft_JP
```

Arguments

```
CPUE.file
```

```
data
```

```
fishery
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
reconstruct.frq.ce2008  
reconstruct.frq.ce2008.Rd
```

Usage

```
reconstruct.frq.ce2008(data = out.data, fishery = 1)
```

Arguments

```
data  
fishery
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
reconstruct.frq.size  
reconstruct.frq.size.Rd
```

Usage

```
reconstruct.frq.size(data = data, FISH = 1, LF.FILE = "P:/yft/2009/Data Preparat
```

Arguments

```
data  
FISH  
LF.FILE  
WT.FILE
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
region_single_frq  region_single_frq.Rd
```

Usage

```
region_single_frq(frq, region)
```

Arguments

```
frq  
region
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

```
region_single_ini  region_single_ini.Rd
```

Usage

```
region_single_ini(ini)
```

Arguments

```
ini
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

```
region_single_tag  region_single_tag.Rd
```

Usage

```
region_single_tag(tag, region, keepfish)
```

Arguments

```
tag
region
keepfish
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==>  Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

```
regroup_fishery_grps.doitall
      regroup_fishery_grps.doitall.Rd
```

Usage

```
regroup_fishery_grps.doitall(doitall, f, flag, newgrps)
```

Arguments

```
doitall
f
flag
newgrps
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==>  Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

```
rename.fisheries.doitall  
      rename.fisheries.doitall.Rd
```

Usage

```
rename.fisheries.doitall(doitall, oldfs, newfs)
```

Arguments

```
doitall  
oldfs  
newfs
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

```
rename.fisheries.frq  
      rename.fisheries.frq.Rd
```

Usage

```
rename.fisheries.frq(frq.obj, oldfish, newfish)
```

Arguments

```
frq.obj  
oldfish  
newfish
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

```
rename.fisheries.tag  
      rename.fisheries.tag.Rd
```

Usage

```
rename.fisheries.tag(tag.obj, oldfish, newfish)
```

Arguments

```
tag.obj  
oldfish  
newfish
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
rename.fishery.grps.doitall  
      rename.fishery.grps.doitall.Rd
```

Usage

```
rename.fishery.grps.doitall(doitall, oldfs, newfs, flag, keep = T, newgrps = c(0
```

Arguments

```
doitall  
oldfs  
newfs  
flag  
keep  
newgrps
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

retro.frq	<i>retro.frq.Rd</i>
-----------	---------------------

Usage

```
retro.frq(frq.obj, retro.tag.obj = NA)
```

Arguments

```
frq.obj  
retro.tag.obj
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

retro.tag	<i>retro.tag.Rd</i>
-----------	---------------------

Usage

```
retro.tag(tag.obj, yr)
```

Arguments

```
tag.obj  
yr
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
rm_fisheries.doitall  
rm_fisheries.doitall.Rd
```

Usage

```
rm_fisheries.doitall(a, rmfisheries)
```

Arguments

```
a  
rmfisheries
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
rm_fisheries.frq rm_fisheries.frq.Rd
```

Usage

```
rm_fisheries.frq(frq.obj, fishery)
```

Arguments

```
frq.obj  
fishery
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

rm_fisheries.tag	<i>rm_fisheries.tag.Rd</i>
------------------	----------------------------

Usage

```
rm_fisheries.tag(tag.obj, fisheries)
```

Arguments

```
tag.obj  
fisheries
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

rm_fishflag	<i>rm_fishflag.Rd</i>
-------------	-----------------------

Usage

```
rm_fishflag(doitall, flag)
```

Arguments

```
doitall  
flag
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

rm_flag.doitall	<i>rm_flag.doitall.Rd</i>
-----------------	---------------------------

Usage

```
rm_flag.doitall(a, flagtype, flag, value)
```

Arguments

```
a  
flagtype  
flag  
value
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

run.profile	<i>run.profile.Rd</i>
-------------	-----------------------

Usage

```
run.profile(rundir, rungrp, startpar = NA, ptype = "Fmult", target, nsteps = 300)
```

Arguments

```
rundir  
rungrp  
startpar  
ptype  
target  
nsteps  
penalty
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

seas.flag	<i>seas.flag.Rd</i>
-----------	---------------------

Usage

```
seas.flag(a, fishery, flagnum, seasf.list)
```

Arguments

```
a  
fishery  
flagnum  
seasf.list
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

seas.frq	<i>seas.frq.Rd</i>
----------	--------------------

Usage

```
seas.frq(frq.obj, seas.fish)
```

Arguments

```
frq.obj  
seas.fish
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

seas.tag	<i>seas.tag.Rd</i>
----------	--------------------

Usage

```
seas.tag(tag.obj, fishlist)
```

Arguments

```
tag.obj  
fishlist
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

setup.cpue	<i>setup.cpue.Rd</i>
------------	----------------------

Usage

```
setup.cpue(rungrp, sourcedir, cpue, spp)
```

Arguments

```
rungrp  
sourcedir  
cpue  
spp
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

setup.ffmpeg	<i>setup.ffmpeg.Rd</i>
--------------	------------------------

Usage

```
setup.ffmpeg(rungrp, creeprate)
```

Arguments

```
rungrp
```

```
creeprate
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

setup.growth	<i>setup.growth.Rd</i>
--------------	------------------------

Usage

```
setup.growth(rungrp, VBopt)
```

Arguments

```
rungrp
```

```
VBopt
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
setup.growth.offsets  
      setup.growth.offsets.Rd
```

Usage

```
setup.growth.offsets(rungrp, ageclasses, penwt, phase, tog)
```

Arguments

```
rungrp  
ageclasses  
penwt  
phase  
tog
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
setup.idphcatch      setup.idphcatch.Rd
```

Usage

```
setup.idphcatch(rungrp, sourcedir, idph, spp)
```

Arguments

```
rungrp  
sourcedir  
idph  
spp
```

Examples

```
##---- Should be DIRECTLY executable !! ----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

setup.lensel	<i>setup.lensel.Rd</i>
--------------	------------------------

Usage

```
setup.lensel(rungrp, fisheries, tog)
```

Arguments

```
rungrp
fisheries
tog
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

setup.LFwt	<i>setup.LFwt.Rd</i>
------------	----------------------

Usage

```
setup.LFwt(rungrp, newLFwt)
```

Arguments

```
rungrp
newLFwt
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

setup.M	<i>setup.M.Rd</i>
---------	-------------------

Usage

```
setup.M(rungrp, newM)
```

Arguments

```
rungrp
```

```
newM
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

setup.pscatch	<i>setup.pscatch.Rd</i>
---------------	-------------------------

Usage

```
setup.pscatch(rungrp, sourcedir, PScatch, spp)
```

Arguments

```
rungrp
```

```
sourcedir
```

```
PScatch
```

```
spp
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

setup.startyr	<i>setup.startyr.Rd</i>
---------------	-------------------------

Usage

```
setup.startyr(rungrp, newstartyr)
```

Arguments

```
rungrp
```

```
newstartyr
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

setup.steepness	<i>setup.steepness.Rd</i>
-----------------	---------------------------

Usage

```
setup.steepness(rungrp, newsteep)
```

Arguments

```
rungrp
```

```
newsteep
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

setup.timesplit	<i>setup.timesplit.Rd</i>
-----------------	---------------------------

Usage

```
setup.timesplit(rungrp, splitx, storefish)
```

Arguments

```
rungrp  
splitx  
storefish
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

sort.frq	<i>sort.frq.Rd</i>
----------	--------------------

Usage

```
sort.frq(frq.obj)
```

Arguments

```
frq.obj
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do  help(data=index)  for the standard data sets.
```

start_year.frq	<i>start_year.frq.Rd</i>
----------------	--------------------------

Usage

```
start_year.frq(frq.obj, start_yr, halfyr = F)
```

Arguments

```
frq.obj  
start_yr  
halfyr
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

steepness.doit	<i>steepness.doit.Rd</i>
----------------	--------------------------

Usage

```
steepness.doit(doitall, new.steepness, add_header = T, gap = 2)
```

Arguments

```
doitall  
new.steepness  
  
add_header  
gap
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

```
summarise.size.frq.bet
      summarise.size.frq.bet.Rd
```

Usage

```
summarise.size.frq.bet(frql, fishery = 5)
```

Arguments

```
frql
fishery
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

```
tag_grps_rm      tag_grps_rm.Rd
```

Usage

```
tag_grps_rm(tag.obj, keep)
```

Arguments

```
tag.obj
keep
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

timesplit.doitall *timesplit.doitall.Rd*

Usage

```
timesplit.doitall(doitall, fishsplit, qsplit = T)
```

Arguments

```
doitall
fishsplit
qsplit
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

timesplit.frq *timesplit.frq.Rd*

Usage

```
timesplit.frq(frq.obj, divyrs, div.fish)
```

Arguments

```
frq.obj
divyrs
div.fish
```

Examples

```
##---- Should be DIRECTLY executable !! ----
##-- ==> Define data, use random,
##--or do  help(data=index)  for the standard data sets.
```

timesplit.tag	<i>timesplit.tag.Rd</i>
---------------	-------------------------

Usage

```
timesplit.tag(tag.obj, fishsplit)
```

Arguments

```
tag.obj
fishsplit
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

varfromstr	<i>varfromstr.Rd</i>
------------	----------------------

Usage

```
varfromstr(datstring, cols = c(2:3))
```

Arguments

```
datstring
cols
```

Examples

```
##----- Should be DIRECTLY executable !! -----
##-- ==> Define data, use random,
##--or do help(data=index) for the standard data sets.
```

write.frq	<i>write.frq.Rd</i>
-----------	---------------------

Usage

```
write.frq(new.frq, frq.obj)
```

Arguments

```
new.frq  
frq.obj
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

write.ini	<i>write.ini.Rd</i>
-----------	---------------------

Usage

```
write.ini(ini.file, ini.obj)
```

Arguments

```
ini.file  
ini.obj
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

write.par	<i>write.par.Rd</i>
-----------	---------------------

Usage

```
write.par(par.file, par.obj)
```

Arguments

```
par.file
```

```
par.obj
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

write.tag	<i>write.tag.Rd</i>
-----------	---------------------

Usage

```
write.tag(tagfile, tag.obj)
```

Arguments

```
tagfile
```

```
tag.obj
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

write_nmd.frq	<i>write_nmd.frq.Rd</i>
---------------	-------------------------

Usage

```
write_nmd.frq(new.frq, frq.obj)
```

Arguments

```
new.frq  
frq.obj
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

write_nmd.par	<i>write_nmd.par.Rd</i>
---------------	-------------------------

Usage

```
write_nmd.par(par.file, par.obj)
```

Arguments

```
par.file  
par.obj
```

Examples

```
##----- Should be DIRECTLY executable !! -----  
##-- ==> Define data, use random,  
##--or do help(data=index) for the standard data sets.
```

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