

Annex III - STREAM: Auxiliary scripts for the conversion from the SDEF format to DG MARE Med&BS Data Call

E. Mantzouni

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```
#R general option:
options(stringsAsFactors = FALSE)

options(warn=0)

options(scipen = 999) # disable scientific notation

#chunk option
knitr::opts_chunk$set(cache=TRUE,echo=TRUE, warning=FALSE,
  message=FALSE, fig.height=6,progress=FALSE,verbose=FALSE,
  include=TRUE,dev='png',autodep=FALSE)

#Load packages

library(COSTcore)
library(COSTeda)
library(COSTdbe)

library(tidyr)
library(dplyr)
library(data.table)
library(knitr)
```

Example of use of the script B_LANDINGS

This script implements the Calculation of the raised numbers by length required for the DG MARE Med&BS Data Call, using as input file the SDEF format (CS table) and COST as the raising procedure

Settings

```
path_in <- paste("C:\\Users\\Bitetto Isabella\\OneDrive - Coispa Tecnologia & Rice
rca S.C.A.R.L\\MARE22\\STREAM\\FINAL REVISION OF DELIVERABLES\\SDEF_to_DG_MARE_Med
BS", sep="")

setwd(path_in)
```

Input Data

Validated csData and clData of the SDEF format and stratification object

```
load("fri-TEST.Rdata")
```

```
fri_strD1 <- strIni(timeStrata="quarter", techStrata = "foCatEu5",  
  spaceStrata = "area")
```

```
fri_strD1
```

```
## An object of class "strIni"
```

```
## Slot "timeStrata":
```

```
## [1] "quarter"
```

```
##
```

```
## Slot "spaceStrata":
```

```
## [1] "area"
```

```
##
```

```
## Slot "techStrata":
```

```
## [1] "foCatEu5"
```

```
##
```

```
## Slot "tpRec":
```

```
## [[1]]
```

```
## [1] NA
```

```
##
```

```
##
```

```
## Slot "spRec":
```

```
## [[1]]
```

```
## [1] NA
```

```
##
```

```
##
```

```
## Slot "tcRec":
```

```
## [[1]]
```

```
## [1] NA
```

```
  fri_csc <- csDataCons(fri_cs1, fri_strD1)
```

```
fri_clc <- clDataCons(fri_cl1, fri_strD1)
```

```
head(fri_csc )
```

```
## An object of class "csDataCons"
```

```
## Slot "desc":
```

```
## [1] "Unknown stock"
```

```
##
```

```
## Slot "tr":
```

##	PSUid	time	space	technical	sampType	landCtry	vslFlgCtry	proj
## 1	1 9999	- 1	GSA99 OTB_-1_VL2440_-1		S	COUNTRY1	COUNTRY1	DCF
## 2	2 9999	- 1	GSA99 OTB_-1_VL2440_-1		S	COUNTRY1	COUNTRY1	DCF
## 3	3 9999	- 1	GSA99 OTB_-1_VL2440_-1		S	COUNTRY1	COUNTRY1	DCF
## 4	4 9999	- 1	GSA99 OTB_-1_VL2440_-1		S	COUNTRY1	COUNTRY1	DCF
## 5	5 9999	- 1	GSA99 OTB_-1_VL2440_-1		S	COUNTRY1	COUNTRY1	DCF
## 6	6 9999	- 1	GSA99 OTB_-1_VL2440_-1		S	COUNTRY1	COUNTRY1	DCF

```

##   trpCode foNum daysAtSea vsId sampCtry sampMeth
## 1      2      3          1    NA COUNTRY1 Observer
## 2      1      2          1    NA COUNTRY1 Observer
## 3      3      3          1    NA COUNTRY1 Observer
## 4      4      3          1    NA COUNTRY1 Observer
## 5      5      3          1    NA COUNTRY1 Observer
## 6      8      3          1    NA COUNTRY1 Observer
##
## Slot "hh":
##   PSuid SSuid      time space      technical sampType landCtry vsIdFlgCtry
## 1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1
## 2      1      2 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1
## 3      1      3 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1
## 4      2      1 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1
## 5      2      2 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1
## 6      3      1 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1
##   proj trpCode staNum foVal aggLev catReg sppReg      date      foDur
## 1 DCF      2      1      V      H      All      All 2017-02-20 3.500000
## 2 DCF      2      2      V      H      All      All 2017-02-20 1.750000
## 3 DCF      2      3      V      H      All      All 2017-02-20 3.750000
## 4 DCF      1      1      V      H      All      All 2017-02-21 6.083333
## 5 DCF      1      2      V      H      All      All 2017-02-21 5.916667
## 6 DCF      3      1      V      H      All      All 2017-02-23 5.083333
##   latIni lonIni latFin lonFin foDep
## 1      NA      NA      NA      NA      NA
## 2      NA      NA      NA      NA      NA
## 3      NA      NA      NA      NA      NA
## 4      NA      NA      NA      NA      NA
## 5      NA      NA      NA      NA      NA
## 6      NA      NA      NA      NA      NA
##
## Slot "sl":
##   PSuid SSuid TSuid      time space      technical      sort sampType
## 1      1      1      3 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 2      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 DIS-HUC-NA-NA      S
## 3      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 DIS-HUC-NA-NA      S
## 4      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 5      1      1      4 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 6      1      2      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
##   landCtry vsIdFlgCtry proj trpCode staNum      spp sex  wt
## 1 COUNTRY1  COUNTRY1 DCF      2      1 Merluccius merluccius <NA> 2400
## 2 COUNTRY1  COUNTRY1 DCF      2      1 Merluccius merluccius <NA> 90
## 3 COUNTRY1  COUNTRY1 DCF      2      1 Mullus barbatus <NA> 60
## 4 COUNTRY1  COUNTRY1 DCF      2      1 Mullus barbatus <NA> 2020
## 5 COUNTRY1  COUNTRY1 DCF      2      1 Merluccius merluccius <NA> 110
## 6 COUNTRY1  COUNTRY1 DCF      2      2 Mullus barbatus <NA> 350
##   subSampWt lenCode
## 1      2400      mm
## 2      45      mm
## 3      30      mm
## 4      2020      mm

```

```

## 5      110      mm
## 6      350      mm
##
## Slot "hl":
##   PSUid SSUid TSUid      time space      technical      sort sampType
## 1      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 2      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 3      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 4      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 5      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 6      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
##   landCtry vsIflgCtry proj trpCode staNum      spp sex lenCls
## 1 COUNTRY1 COUNTRY1 DCF      2      1 Mullus barbatus <NA> 170
## 2 COUNTRY1 COUNTRY1 DCF      2      1 Mullus barbatus <NA> 180
## 3 COUNTRY1 COUNTRY1 DCF      2      1 Mullus barbatus <NA> 150
## 4 COUNTRY1 COUNTRY1 DCF      2      1 Mullus barbatus <NA> 160
## 5 COUNTRY1 COUNTRY1 DCF      2      1 Mullus barbatus <NA> 140
## 6 COUNTRY1 COUNTRY1 DCF      2      1 Mullus barbatus <NA> 200
##   lenNum
## 1      7
## 2      7
## 3      7
## 4      3
## 5      7
## 6      3
##
## Slot "ca":
##   PSUid SSUid      time space      technical      sort sampType landCtry
## 1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA      S COUNTRY1
## 2      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA      S COUNTRY1
## 3      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA      S COUNTRY1
## 4      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA      S COUNTRY1
## 5      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA      S COUNTRY1
## 6      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA      S COUNTRY1
##   vsIflgCtry proj trpCode staNum      spp sex
## 1 COUNTRY1 DCF      2      1 Merluccius merluccius U
## 2 COUNTRY1 DCF      2      1 Mullus barbatus U
## 3 COUNTRY1 DCF      2      1 Mullus barbatus U
## 4 COUNTRY1 DCF      2      1 Merluccius merluccius U
## 5 COUNTRY1 DCF      2      1 Mullus barbatus U
## 6 COUNTRY1 DCF      2      1 Mullus barbatus U
##
##   stock lenCls age fishId lenCode ageMeth plusGrp otoWt
## 1 Merluccius merluccius_22 170 NA 20 mm OWR <NA> NA
## 2 Mullus barbatus_22 130 NA 37 mm OWR <NA> NA
## 3 Mullus barbatus_22 170 NA 7 mm OWR <NA> NA
## 4 Merluccius merluccius_22 160 NA 19 mm OWR <NA> NA
## 5 Mullus barbatus_22 160 NA 36 mm OWR <NA> NA
## 6 Mullus barbatus_22 110 NA 22 mm OWR <NA> NA
##   otoSide indWt matMeth matScale matStage
## 1 <NA> NA <NA> <NA> <NA>
## 2 <NA> NA <NA> <NA> <NA>

```

```
## 3      <NA>      NA      <NA>      <NA>      <NA>
## 4      <NA>      NA      <NA>      <NA>      <NA>
## 5      <NA>      NA      <NA>      <NA>      <NA>
## 6      <NA>      NA      <NA>      <NA>      <NA>
```

```
head(fri_clc)
```

```
## An object of class "clDataCons"
```

```
## Slot "desc":
```

```
## [1] "Unknown stock"
```

```
##
```

```
## Slot "cl":
```

```
##   landCtry vs1FlgCtry      time space      technical
```

```
## 1 COUNTRY1  COUNTRY1 9999 - 1 GSA99 GNS_-1_VL1218_-1
```

```
## 2 COUNTRY1  COUNTRY1 9999 - 1 GSA99 LLS_-1_VL1218_-1
```

```
## 3 COUNTRY1  COUNTRY1 9999 - 1 GSA99 OTB_-1_VL1824_-1
```

```
## 4 COUNTRY1  COUNTRY1 9999 - 1 GSA99 OTB_-1_VL1824_-1
```

```
## 5 COUNTRY1  COUNTRY1 9999 - 1 GSA99 OTB_-1_VL2440_-1
```

```
## 6 COUNTRY1  COUNTRY1 9999 - 1 GSA99 OTB_-1_VL2440_-1
```

```
##           taxon landCat commCatScl commCat unallocCatchWt
```

```
## 1 Merluccius merluccius      HUC      <NA>      <NA>          NA
```

```
## 2 Merluccius merluccius      HUC      <NA>      <NA>          NA
```

```
## 3 Merluccius merluccius      HUC      <NA>      <NA>          NA
```

```
## 4      Mullus barbatus      HUC      <NA>      <NA>          NA
```

```
## 5 Merluccius merluccius      HUC      <NA>      <NA>          NA
```

```
## 6      Mullus barbatus      HUC      <NA>      <NA>          NA
```

```
##   misRepCatchWt landWt landMult landValue
```

```
## 1              NA     46         1         NA
```

```
## 2              NA    140         1         NA
```

```
## 3              NA   2151         1         NA
```

```
## 4              NA   2285         1         NA
```

```
## 5              NA  32979         1         NA
```

```
## 6              NA   9144         1         NA
```

```
# extract COUNTRY and YEAR
```

```
COUNTRY<-unique(fri_cl1@cl$landCtry)
```

```
YEAR=unique(fri_cl1@cl$year)
```

DG MARE Med&BS LANDINGS Table template

```
lan.temp2<- read.table("xxx_LANDINGS.csv",sep="," ,header=T)
```

```
head(lan.temp2)
```

```
##   [1] ID                COUNTRY                YEAR
```

```
##   [4] QUARTER              VESSEL_LENGTH         GEAR
```

```
##   [7] MESH_SIZE_RANGE      FISHERY                AREA
```

```
##  [10] SPECON              SPECIES                LANDINGS
```

```
##  [13] UNIT                LENGTHCLASS0          LENGTHCLASS1
```

```
##  [16] LENGTHCLASS2        LENGTHCLASS3          LENGTHCLASS4
```

```
##  [19] LENGTHCLASS5        LENGTHCLASS6          LENGTHCLASS7
```

```
##  [22] LENGTHCLASS8        LENGTHCLASS9          LENGTHCLASS10
```

```
## [25] LENGTHCLASS11      LENGTHCLASS12      LENGTHCLASS13
## [28] LENGTHCLASS14      LENGTHCLASS15      LENGTHCLASS16
## [31] LENGTHCLASS17      LENGTHCLASS18      LENGTHCLASS19
## [34] LENGTHCLASS20      LENGTHCLASS21      LENGTHCLASS22
## [37] LENGTHCLASS23      LENGTHCLASS24      LENGTHCLASS25
## [40] LENGTHCLASS26      LENGTHCLASS27      LENGTHCLASS28
## [43] LENGTHCLASS29      LENGTHCLASS30      LENGTHCLASS31
## [46] LENGTHCLASS32      LENGTHCLASS33      LENGTHCLASS34
## [49] LENGTHCLASS35      LENGTHCLASS36      LENGTHCLASS37
## [52] LENGTHCLASS38      LENGTHCLASS39      LENGTHCLASS40
## [55] LENGTHCLASS41      LENGTHCLASS42      LENGTHCLASS43
## [58] LENGTHCLASS44      LENGTHCLASS45      LENGTHCLASS46
## [61] LENGTHCLASS47      LENGTHCLASS48      LENGTHCLASS49
## [64] LENGTHCLASS50      LENGTHCLASS51      LENGTHCLASS52
## [67] LENGTHCLASS53      LENGTHCLASS54      LENGTHCLASS55
## [70] LENGTHCLASS56      LENGTHCLASS57      LENGTHCLASS58
## [73] LENGTHCLASS59      LENGTHCLASS60      LENGTHCLASS61
## [76] LENGTHCLASS62      LENGTHCLASS63      LENGTHCLASS64
## [79] LENGTHCLASS65      LENGTHCLASS66      LENGTHCLASS67
## [82] LENGTHCLASS68      LENGTHCLASS69      LENGTHCLASS70
## [85] LENGTHCLASS71      LENGTHCLASS72      LENGTHCLASS73
## [88] LENGTHCLASS74      LENGTHCLASS75      LENGTHCLASS76
## [91] LENGTHCLASS77      LENGTHCLASS78      LENGTHCLASS79
## [94] LENGTHCLASS80      LENGTHCLASS81      LENGTHCLASS82
## [97] LENGTHCLASS83      LENGTHCLASS84      LENGTHCLASS85
## [100] LENGTHCLASS86      LENGTHCLASS87      LENGTHCLASS88
## [103] LENGTHCLASS89      LENGTHCLASS90      LENGTHCLASS91
## [106] LENGTHCLASS92      LENGTHCLASS93      LENGTHCLASS94
## [109] LENGTHCLASS95      LENGTHCLASS96      LENGTHCLASS97
## [112] LENGTHCLASS98      LENGTHCLASS99      LENGTHCLASS100_PLUS
## <0 rows> (or 0-length row.names)
```

Communication Table for FISHERY

```
fishery<- read.table("communicationTable_for_fishery.csv",sep=";",header=T)
```

```
head(fishery)
```

```
##   SDEF_codification DGMARE_Med_BS_codification
## 1                MOL                      MOL
## 2                DES                      DEMSP
## 3                DWS                      DWS
## 4                MDD                      MDD
## 5                SPF                      SPF
## 6                FIF                      FIF
```

Auxiliary table: species_LAN.csv

```
# species file : selected species with FAO three alpha code
```

```
sel_spe <-read.table("species_LANDINGS.csv",sep=";",header=T)
```

```
head(sel_spe)
```

```
##           SPECIES SPE   GSA LC_RANGE lanEstim_methodDesc SPECON
## 1 Merluccius merluccius HKE GSA99      10      analytical      .
## 2      Mullus barbatus MUT GSA99      10      analytical      .
```

Data analysis- raising

Analysis by stock

```
for (i in 1:dim(sel_spe)[1]) {

  STK<- sel_spe$SPECIES[i]

  AREA <- sel_spe$GSA[i]

  fri_csc1<- subset(fri_csc, space==sel_spe$GSA[i],table="ca",link=T)
  fri_clc1<- subset(fri_clc, space==sel_spe$GSA[i],table="cl")

  # The first step is to create the empty object, that will be given
# the appropriate values for the descriptor fields.

  lanEstim <-
    dbeObject(
      desc = paste(STK, AREA,"Landings", sep="_"),
      species = STK,
      catchCat = "LAN",
      strataDesc = fri_strD1,
      methodDesc = sel_spe$lanEstim_methodDesc[i]
    )

  # the only arguments to pass to the function are the dbe object,
# the consolidated cs and cl datasets.

  if ( sel_spe$lanEstim_methodDesc[i]=="analytical"){
lanEstim <- RaiseLgth(lanEstim, fri_csc1, fri_clc1,incl.precision =F)
  } else {
lanEstim <- RaiseLgthBoot(lanEstim, fri_csc1, fri_clc1,
                        incl.precision =F,B=15)
  }

  # totalW\$estim : total weight,
  aa <-lanEstim$totalW$estim

  aa$value<- aa$value/1000 # tons

  aa<- rename(aa, "totalW"=value)
```

```

# LenStruc\$estim : numbers-at-length estimates

bb<- lanEstim@lenStruc$estim

# define LCs and UNIT Len
UNIT <- as.character( unique(fri_csc@ca$lenCode[fri_csc@ca$spp==STK]) )

if (UNIT %in% c("mm", "MM") & sel_spe$LC_RANGE[i]==10) {
bb$length<-as.numeric(bb$length)/10
UNIT1<- "CM"
}

if (UNIT %in% c("mm", "MM") & sel_spe$LC_RANGE[i]==1) {
bb$length<-as.numeric(bb$length)
UNIT1<- "MM"
}

if (UNIT %in% c("mm", "MM") & sel_spe$LC_RANGE[i]==5) {
bb$length<-as.numeric(bb$length)/10
UNIT1<- "CM"
}

if (UNIT %in% c("cm", "CM") ) {
bb$length<-as.numeric(bb$length)
UNIT1<- "CM"
}

bb$length<- plyr::round_any( bb$length,1,floor)

bb$value<- bb$value/1000 # '000 ind

ab=left_join(bb,aa ,by = c("time", "space", "technical"))

ab<- ab %>% separate(technical, c("gear","FISHERY", "VL","MESH_SIZE_RANGE"),
                    sep = "_",remove=T)

ab$length<- as.numeric(as.character(ab$length))

ab<- ab%>% group_by(time, space , gear ,FISHERY, VL,MESH_SIZE_RANGE ) %>%
mutate(minlc=min(length,na.rm=T),maxlc=max(length,na.rm=T))

# matrix with all combinations of "time" "space" "gear" "VL"
# "length" , "MESH_SIZE_RANGE"

dt <- as.data.table(ab)

dt[,c(1:7)][is.na(dt[,c(1:7)])]<- -1

seq_1 <- seq(0, max(dt$length,na.rm = T), by = 1) #

```



```

dt$id<- paste(dt$time,dt$space,dt$gear,dt$FISHERY,dt$VL,
             dt$MESH_SIZE_RANGE,sep=":")

dt1<- dt[, list(length = seq_1), by = id]

dt1<- dt1 %>% separate(id, c("time", "space", "gear", "FISHERY","VL",
                             "MESH_SIZE_RANGE"), sep = ":")

ab[is.na(ab)]<- -1

dt2<- left_join(dt1,ab)
dt2$stock<- STK

##

dt3 <- data.table::dcast(dt2,as.formula(paste(paste(names(dt2)[! names(dt2) %in%
c("length","value")], collapse='+'), "length", sep="~")),
value.var = "value")

dt3=dt3[complete.cases(dt3[,c(7:9)]), ]

dt3 <- dt3 %>% separate(time, c("Year","Quarter")," - ")

dt3$MESH_SIZE_RANGE<-as.character(dt3$MESH_SIZE_RANGE)

# numbers at LC : NA-->0
dt3<- dt3 %>% mutate_at(vars( -(Year:stock) ),
  funs( if_else( is.na(.), 0, .) ) )

LANDINGS <- data.frame(

  ID = NA ,
  COUNTRY = COUNTRY ,
  YEAR = YEAR ,
  QUARTER =dt3$Quarter,
  VESSEL_LENGTH = dt3$VL,
  GEAR = dt3$gear,
  MESH_SIZE_RANGE = dt3$MESH_SIZE_RANGE,
  FISHERY = dt3$FISHERY ,
  AREA = sel_spe$GSA[i],
  SPECON = -1 ,
  SPECIES = STK ,
  LANDINGS = dt3$totalw ,
  UNIT = UNIT1
)

```

```

LANDINGS<-left_join(LANDINGS,dt3[, -c(1,3,8:11)],by=c( "QUARTER" ="Quarter" ,
"GEAR"="gear" , "VESSEL_LENGTH" = "VL" ,
"MESH_SIZE_RANGE", "FISHERY" ))

# take care of number of Length classes (max is 100 acc. to DG MARE Med&BS template)
zz<-dim(LANDINGS[-c(1:13))][2]
names(LANDINGS)[-c(1:13)]<- paste("LENGTHCLASS",seq(0,zz-1,1),sep="")

if(zz>=100){

  LANDINGS$LENGTHCLASS100_PLUS<- rowSums(LANDINGS[, !1:113])

  LANDINGS<-LANDINGS %>% select(ID:LENGTHCLASS99, LENGTHCLASS100_PLUS)
}

# FISHERY to DG MARE Med&BS codification
LANDINGS$FISHERY <- fishery$SDEF_codification[match(LANDINGS$FISHERY ,
fishery$DGMARE_Med_BS_codification)]

# species to FAO three alpha code and set ID (COUNTRY, AREA, GEAR, VESSEL_LENGTH
,
# MESH_SIZE_RANGE, QUARTER, SPECIES)

land.tab <-LANDINGS %>% mutate(SPECIES=spe_spe$SPE[match(SPECIES,spe_spe$SPECIES
)],
ID = paste(COUNTRY, AREA, GEAR,FISHERY, VESSEL_L
ENGTH,
MESH_SIZE_RANGE, YEAR, QUARTER, SPECIES, sep = "_"))

lan.temp2<-bind_rows(lan.temp2,land.tab)

lan.temp2[, -c(1:13)][is.na(lan.temp2[, -c(1:13)])] <- 0
}

```

Output

```

lan.temp2 <- lan.temp2[, 2:ncol(lan.temp2)]
write.table(format(lan.temp2,digits=3,scientific=F), file = "LANDINGS.csv",
row.names=FALSE,sep=";",na="-1")

head(lan.temp2)

```

##	COUNTRY	YEAR	QUARTER	VESSEL_LENGTH	GEAR	MESH_SIZE_RANGE	FISHERY	AREA
## 1	COUNTRY1	9999	1	VL2440	OTB	-1	NONE	GSA99
## 2	COUNTRY1	9999	3	VL0612	GNS	-1	NONE	GSA99
## 3	COUNTRY1	9999	3	VL2440	OTB	-1	NONE	GSA99
## 4	COUNTRY1	9999	4	VL0006	GNS	-1	NONE	GSA99
## 5	COUNTRY1	9999	4	VL0612	GNS	-1	NONE	GSA99
## 6	COUNTRY1	9999	4	VL0612	GTR	-1	NONE	GSA99
##	SPECON	SPECIES	LANDINGS	UNIT	LENGTHCLASS0	LENGTHCLASS1	LENGTHCLASS2	
## 1	-1	HKE	241.996000	CM	0	0	0	
## 2	-1	HKE	140.230571	CM	0	0	0	
## 3	-1	HKE	196.598000	CM	0	0	0	
## 4	-1	HKE	10.177558	CM	0	0	0	
## 5	-1	HKE	87.679445	CM	0	0	0	
## 6	-1	HKE	2.382747	CM	0	0	0	
##	LENGTHCLASS3	LENGTHCLASS4	LENGTHCLASS5	LENGTHCLASS6	LENGTHCLASS7			
## 1	0	0	0	0	0			
## 2	0	0	0	0	0			
## 3	0	0	0	0	0			
## 4	0	0	0	0	0			
## 5	0	0	0	0	0			
## 6	0	0	0	0	0			
##	LENGTHCLASS8	LENGTHCLASS9	LENGTHCLASS10	LENGTHCLASS11	LENGTHCLASS12			
## 1	0	0	0	0	0			
## 2	0	0	0	0	0			
## 3	0	0	0	0	0			
## 4	0	0	0	0	0			
## 5	0	0	0	0	0			
## 6	0	0	0	0	0			
##	LENGTHCLASS13	LENGTHCLASS14	LENGTHCLASS15	LENGTHCLASS16	LENGTHCLASS17			
## 1	0	0.000000	19.793306	24.827583	73.370333			
## 2	0	0.000000	0.000000	0.000000	0.000000			
## 3	0	4.138302	8.188831	17.894727	20.699818			
## 4	0	0.000000	0.000000	0.000000	0.000000			
## 5	0	0.000000	2.836237	1.134495	3.403485			
## 6	0	0.000000	0.000000	0.000000	0.000000			
##	LENGTHCLASS18	LENGTHCLASS19	LENGTHCLASS20	LENGTHCLASS21	LENGTHCLASS22			
## 1	144.63101	170.341556	152.4106110	188.78637	115.6763957			
## 2	0.000000	0.000000	0.0000000	0.000000	19.2228335			
## 3	24.95705	34.569465	52.1754254	65.54129	51.2356341			
## 4	0.000000	0.000000	0.0000000	0.000000	1.7191821			
## 5	6.64339	7.962294	5.9055523	11.74862	15.5695889			
## 6	0.000000	0.000000	0.1604868	0.000000	0.1604868			
##	LENGTHCLASS23	LENGTHCLASS24	LENGTHCLASS25	LENGTHCLASS26	LENGTHCLASS27			
## 1	144.5208194	86.5762343	125.5708742	129.9463867	110.9232050			
## 2	0.0000000	0.0000000	0.0000000	0.0000000	0.0000000			
## 3	53.7350790	64.4316644	63.9193107	43.7347026	38.3000654			
## 4	0.0000000	1.7191821	3.4383642	3.4383642	1.7191821			
## 5	17.7096472	24.4611394	32.0892633	32.0962731	37.2363491			
## 6	0.1604868	0.1604868	0.1604868	0.1604868	0.3209735			
##	LENGTHCLASS28	LENGTHCLASS29	LENGTHCLASS30	LENGTHCLASS31	LENGTHCLASS32			
## 1	109.01789	52.6093754	50.0200945	42.1897758	22.8384296			

## 2	0.00000	19.2228335	19.2228335	38.4456671	38.4456671
## 3	29.49994	42.5713246	27.4307910	32.6078231	23.4171370
## 4	0.00000	1.7191821	3.4383642	0.0000000	0.0000000
## 5	26.28628	18.0104432	23.4942145	14.2945541	13.5494027
## 6	0.00000	0.3209735	0.4814603	0.1604868	0.6419471
##	LENGTHCLASS33	LENGTHCLASS34	LENGTHCLASS35	LENGTHCLASS36	LENGTHCLASS37
## 1	25.8835535	25.8835535	16.7481817	7.6128099	12.180496
## 2	38.4456671	38.4456671	38.4456671	38.4456671	19.222834
## 3	31.0538825	25.3616402	26.5250182	22.6443216	13.586593
## 4	1.7191821	1.7191821	0.0000000	1.7191821	0.0000000
## 5	14.8379526	11.9465417	8.1090350	15.3843710	7.432411
## 6	0.1604868	0.3209735	0.1604868	0.1604868	0.0000000
##	LENGTHCLASS38	LENGTHCLASS39	LENGTHCLASS40	LENGTHCLASS41	LENGTHCLASS42
## 1	13.703058	6.0902479	6.0902479	4.5676859	1.522562
## 2	0.0000000	0.0000000	0.0000000	19.2228335	0.0000000
## 3	12.680820	9.9635015	6.5980152	11.1268795	9.963501
## 4	1.719182	0.0000000	0.0000000	0.0000000	0.0000000
## 5	6.974540	5.5392491	2.2689900	2.1357642	4.729399
## 6	0.0000000	0.8024339	0.4814603	0.4814603	0.0000000
##	LENGTHCLASS43	LENGTHCLASS44	LENGTHCLASS45	LENGTHCLASS46	LENGTHCLASS47
## 1	3.0451239	7.6128099	0.0000000	0.0000000	0.0000000
## 2	0.0000000	19.2228335	19.222834	0.0000000	0.0000000
## 3	6.3404100	3.6230915	2.717319	2.717319	1.8115457
## 4	0.0000000	1.7191821	0.0000000	0.0000000	0.0000000
## 5	2.4604090	1.8931615	0.0000000	1.325914	1.1344950
## 6	0.1604868	0.1604868	0.0000000	0.0000000	0.1604868
##	LENGTHCLASS48	LENGTHCLASS49	LENGTHCLASS50	LENGTHCLASS51	LENGTHCLASS52
## 1	4.5676859	0.0000000	0.0000000	0.0000000	0.0000000
## 2	19.2228335	0.0000000	0.0000000	0.0000000	0.0000000
## 3	0.9057729	3.6230915	1.811546	0.9057729	0.9057729
## 4	3.4383642	0.0000000	0.0000000	0.0000000	0.0000000
## 5	0.5672475	0.5672475	0.0000000	0.0000000	0.0000000
## 6	0.0000000	0.0000000	0.0000000	0.0000000	0.0000000
##	LENGTHCLASS53	LENGTHCLASS54	LENGTHCLASS55	LENGTHCLASS56	LENGTHCLASS57
## 1	0.0000000	0.0000000	0.0000000	0.0000000	0.0000000
## 2	0.0000000	0.0000000	0.0000000	0.0000000	0.0000000
## 3	0.0000000	0.9057729	0.0000000	1.811546	0.9057729
## 4	0.0000000	0.0000000	0.0000000	0.0000000	0.0000000
## 5	0.5672475	0.0000000	0.0000000	0.0000000	0.0000000
## 6	0.0000000	0.0000000	0.1604868	0.0000000	0.0000000
##	LENGTHCLASS58	LENGTHCLASS59	LENGTHCLASS60	LENGTHCLASS61	LENGTHCLASS62
## 1	0.0000000	0.0000000	0.0000000	0.0000000	0
## 2	0.0000000	0.0000000	0.0000000	0.0000000	0
## 3	1.811546	2.717319	1.811546	0.9057729	0
## 4	0.0000000	0.0000000	0.0000000	0.0000000	0
## 5	0.0000000	0.0000000	0.0000000	0.0000000	0
## 6	0.0000000	0.0000000	0.0000000	0.0000000	0
##	LENGTHCLASS63	LENGTHCLASS64	LENGTHCLASS65	LENGTHCLASS66	LENGTHCLASS67
## 1	1.5225620	0.0000000	0	0	0.0000000
## 2	0.0000000	0.0000000	0	0	0.0000000
## 3	0.9057729	0.9057729	0	0	0.9057729

## 4	0.0000000	0.0000000	0	0	0.0000000
## 5	0.0000000	0.0000000	0	0	0.0000000
## 6	0.0000000	0.0000000	0	0	0.0000000
##	LENGTHCLASS68	LENGTHCLASS69	LENGTHCLASS70	LENGTHCLASS71	LENGTHCLASS72
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS73	LENGTHCLASS74	LENGTHCLASS75	LENGTHCLASS76	LENGTHCLASS77
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS78	LENGTHCLASS79	LENGTHCLASS80	LENGTHCLASS81	LENGTHCLASS82
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS83	LENGTHCLASS84	LENGTHCLASS85	LENGTHCLASS86	LENGTHCLASS87
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS88	LENGTHCLASS89	LENGTHCLASS90	LENGTHCLASS91	LENGTHCLASS92
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS93	LENGTHCLASS94	LENGTHCLASS95	LENGTHCLASS96	LENGTHCLASS97
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS98	LENGTHCLASS99	LENGTHCLASS100_PLUS		
## 1	0	0	0		
## 2	0	0	0		
## 3	0	0	0		
## 4	0	0	0		

## 5	0	0	0
## 6	0	0	0

Raising of the length distributions by ID of the discards

E. Mantzouni

Wed Jul 17 09:33:29 2019

```
#R general option:
options(stringsAsFactors = FALSE)

options(warn=0)

options(scipen = 999) # disable scientific notation

#chunk option
knitr::opts_chunk$set(cache=TRUE,echo=TRUE, warning=FALSE,
  message=FALSE, fig.height=6,progress=FALSE,verbose=FALSE,
  include=TRUE,dev= 'png',autodep=FALSE)

#Load packages

library(COSTcore)
library(COSTeda)
library(COSTdbe)
library(tidyr)
library(dplyr)
library(data.table)
library(knitr)
```

Example of use of the script C_DISCARDS

Settings

```
path_in <- paste("C:\\Users\\Bitetto Isabella\\OneDrive - Coispa Tecnologia & Rice
rca S.C.A.R.L\\MARE22\\STREAM\\FINAL REVISION OF DELIVERABLES\\SDEF_to_DG_MARE_Med
BS", sep="")
setwd(path_in)
```

Input Data

Validated csData , ceData, and clData

```
load("fri-TEST.Rdata")

head(fri_cs1@hh$foCatEu5)
```

```

## [1] "OTB_-1_VL2440_-1" "OTB_-1_VL2440_-1" "OTB_-1_VL2440_-1"
## [4] "OTB_-1_VL2440_-1" "OTB_-1_VL2440_-1" "OTB_-1_VL2440_-1"

head(fri_cl1@cl$foCatEu5)

## [1] "GNS_-1_VL1218_-1" "LLS_-1_VL1218_-1" "OTB_-1_VL1824_-1"
## [4] "OTB_-1_VL1824_-1" "OTB_-1_VL2440_-1" "OTB_-1_VL2440_-1"

head(fri_cev@ce$foCatEu5)

## [1] "FPN_-1_VL0006_-1" "FPN_-1_VL0006_-1" "FPN_-1_VL0006_-1"
## [4] "FPN_-1_VL0006_-1" "FPN_-1_VL0006_-1" "FPN_-1_VL0006_-1"

fri_strD1 <- strIni(timeStrata="quarter", techStrata = "foCatEu5",
                    spaceStrata = "area")

fri_strD1

## An object of class "strIni"
## Slot "timeStrata":
## [1] "quarter"
##
## Slot "spaceStrata":
## [1] "area"
##
## Slot "techStrata":
## [1] "foCatEu5"
##
## Slot "tpRec":
## [[1]]
## [1] NA
##
##
## Slot "spRec":
## [[1]]
## [1] NA
##
##
## Slot "tcRec":
## [[1]]
## [1] NA

fri_csc <- csDataCons(fri_cs1, fri_strD1)
fri_clc <- clDataCons(fri_cl1, fri_strD1)
fri_cec <- ceDataCons(fri_cev, fri_strD1)

head(fri_csc )

## An object of class "csDataCons"
## Slot "desc":
## [1] "Unknown stock"
##

```

```

## Slot "tr":
##   PSUid      time space      technical sampType landCtry vs1FlgCtry proj
## 1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1 COUNTRY1 DCF
## 2      2 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1 COUNTRY1 DCF
## 3      3 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1 COUNTRY1 DCF
## 4      4 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1 COUNTRY1 DCF
## 5      5 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1 COUNTRY1 DCF
## 6      6 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1 COUNTRY1 DCF
##   trpCode foNum daysAtSea vs1Id sampCtry sampMeth
## 1      2      3          1      NA COUNTRY1 Observer
## 2      1      2          1      NA COUNTRY1 Observer
## 3      3      3          1      NA COUNTRY1 Observer
## 4      4      3          1      NA COUNTRY1 Observer
## 5      5      3          1      NA COUNTRY1 Observer
## 6      8      3          1      NA COUNTRY1 Observer
##
## Slot "hh":
##   PSUid SSUid      time space      technical sampType landCtry vs1FlgCtry
## 1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1 COUNTRY1
## 2      1      2 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1 COUNTRY1
## 3      1      3 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1 COUNTRY1
## 4      2      1 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1 COUNTRY1
## 5      2      2 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1 COUNTRY1
## 6      3      1 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1 COUNTRY1
##   proj trpCode staNum foVal aggLev catReg sppReg      date      foDur
## 1 DCF      2      1      V      H      All      All 2017-02-20 3.500000
## 2 DCF      2      2      V      H      All      All 2017-02-20 1.750000
## 3 DCF      2      3      V      H      All      All 2017-02-20 3.750000
## 4 DCF      1      1      V      H      All      All 2017-02-21 6.083333
## 5 DCF      1      2      V      H      All      All 2017-02-21 5.916667
## 6 DCF      3      1      V      H      All      All 2017-02-23 5.083333
##   latIni lonIni latFin lonFin foDep
## 1      NA      NA      NA      NA      NA
## 2      NA      NA      NA      NA      NA
## 3      NA      NA      NA      NA      NA
## 4      NA      NA      NA      NA      NA
## 5      NA      NA      NA      NA      NA
## 6      NA      NA      NA      NA      NA
##
## Slot "sl":
##   PSUid SSUid TSuid      time space      technical      sort sampType
## 1      1      1      3 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 2      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 DIS-HUC-NA-NA      S
## 3      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 DIS-HUC-NA-NA      S
## 4      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 5      1      1      4 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 6      1      2      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
##   landCtry vs1FlgCtry proj trpCode staNum      spp sex wt
## 1 COUNTRY1 COUNTRY1 DCF      2      1 Merluccius merluccius <NA> 2400
## 2 COUNTRY1 COUNTRY1 DCF      2      1 Merluccius merluccius <NA> 90
## 3 COUNTRY1 COUNTRY1 DCF      2      1      Mullus barbatus <NA> 60

```



```

## 4 COUNTRY1 COUNTRY1 DCF 2 1 Mullus barbatus <NA> 2020
## 5 COUNTRY1 COUNTRY1 DCF 2 1 Merluccius merluccius <NA> 110
## 6 COUNTRY1 COUNTRY1 DCF 2 2 Mullus barbatus <NA> 350
## subSampWt lenCode
## 1 2400 mm
## 2 45 mm
## 3 30 mm
## 4 2020 mm
## 5 110 mm
## 6 350 mm
##
## Slot "h1":
## PSUid SSUid TSUid time space technical sort sampType
## 1 1 1 1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA S
## 2 1 1 1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA S
## 3 1 1 1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA S
## 4 1 1 1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA S
## 5 1 1 1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA S
## 6 1 1 1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA S
## landCtry vs1FlgCtry proj trpCode staNum spp sex lenCls
## 1 COUNTRY1 COUNTRY1 DCF 2 1 Mullus barbatus <NA> 170
## 2 COUNTRY1 COUNTRY1 DCF 2 1 Mullus barbatus <NA> 180
## 3 COUNTRY1 COUNTRY1 DCF 2 1 Mullus barbatus <NA> 150
## 4 COUNTRY1 COUNTRY1 DCF 2 1 Mullus barbatus <NA> 160
## 5 COUNTRY1 COUNTRY1 DCF 2 1 Mullus barbatus <NA> 140
## 6 COUNTRY1 COUNTRY1 DCF 2 1 Mullus barbatus <NA> 200
## lenNum
## 1 7
## 2 7
## 3 7
## 4 3
## 5 7
## 6 3
##
## Slot "ca":
## PSUid SSUid time space technical sort sampType landCtry
## 1 1 1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA S COUNTRY1
## 2 1 1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA S COUNTRY1
## 3 1 1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA S COUNTRY1
## 4 1 1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA S COUNTRY1
## 5 1 1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA S COUNTRY1
## 6 1 1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA S COUNTRY1
## vs1FlgCtry proj trpCode staNum spp sex
## 1 COUNTRY1 DCF 2 1 Merluccius merluccius U
## 2 COUNTRY1 DCF 2 1 Mullus barbatus U
## 3 COUNTRY1 DCF 2 1 Mullus barbatus U
## 4 COUNTRY1 DCF 2 1 Merluccius merluccius U
## 5 COUNTRY1 DCF 2 1 Mullus barbatus U
## 6 COUNTRY1 DCF 2 1 Mullus barbatus U
##
## stock lenCls age fishId lenCode ageMeth plusGrp otoWt
## 1 Merluccius merluccius_22 170 NA 20 mm OWR <NA> NA

```

```
## 2      Mullus barbatus_22      130 NA      37      mm      OWR      <NA>      NA
## 3      Mullus barbatus_22      170 NA       7      mm      OWR      <NA>      NA
## 4 Merluccius merluccius_22      160 NA      19      mm      OWR      <NA>      NA
## 5      Mullus barbatus_22      160 NA      36      mm      OWR      <NA>      NA
## 6      Mullus barbatus_22      110 NA      22      mm      OWR      <NA>      NA
##   otoSide indWt matMeth matScale matStage
## 1      <NA>    NA      <NA>      <NA>      <NA>
## 2      <NA>    NA      <NA>      <NA>      <NA>
## 3      <NA>    NA      <NA>      <NA>      <NA>
## 4      <NA>    NA      <NA>      <NA>      <NA>
## 5      <NA>    NA      <NA>      <NA>      <NA>
## 6      <NA>    NA      <NA>      <NA>      <NA>
```

`head(fri_clc)`

```
## An object of class "clDataCons"
## Slot "desc":
## [1] "Unknown stock"
##
## Slot "cl":
##   landCtry vs1FlgCtry      time space      technical
## 1 COUNTRY1  COUNTRY1 9999 - 1 GSA99 GNS_-1_VL1218_-1
## 2 COUNTRY1  COUNTRY1 9999 - 1 GSA99 LLS_-1_VL1218_-1
## 3 COUNTRY1  COUNTRY1 9999 - 1 GSA99 OTB_-1_VL1824_-1
## 4 COUNTRY1  COUNTRY1 9999 - 1 GSA99 OTB_-1_VL1824_-1
## 5 COUNTRY1  COUNTRY1 9999 - 1 GSA99 OTB_-1_VL2440_-1
## 6 COUNTRY1  COUNTRY1 9999 - 1 GSA99 OTB_-1_VL2440_-1
##           taxon landCat commCatSc1 commCat unallocCatchWt
## 1 Merluccius merluccius      HUC      <NA>      <NA>      NA
## 2 Merluccius merluccius      HUC      <NA>      <NA>      NA
## 3 Merluccius merluccius      HUC      <NA>      <NA>      NA
## 4      Mullus barbatus      HUC      <NA>      <NA>      NA
## 5 Merluccius merluccius      HUC      <NA>      <NA>      NA
## 6      Mullus barbatus      HUC      <NA>      <NA>      NA
##   misRepCatchWt landWt landMult landValue
## 1             NA     46         1         NA
## 2             NA    140         1         NA
## 3             NA   2151         1         NA
## 4             NA   2285         1         NA
## 5             NA  32979         1         NA
## 6             NA   9144         1         NA
```

`head(fri_cec)`

```
## An object of class "ceDataCons"
## Slot "desc":
## [1] "Unknown stock"
##
## Slot "ce":
##   vs1FlgCtry      time space      technical      trpNum foNum foDur
## 1 COUNTRY1 9999 - 2 GSA99 FPN_-1_VL0006_-1 417.8571    NA    NA
## 2 COUNTRY1 9999 - 2 GSA99 FPN_-1_VL0006_-1 167.1429    NA    NA
```

```
## 3  COUNTRY1 9999 - 3 GSA99 FPN_-1_VL0006_-1 501.4286 NA NA
## 4  COUNTRY1 9999 - 3 GSA99 FPN_-1_VL0006_-1 417.8571 NA NA
## 5  COUNTRY1 9999 - 3 GSA99 FPN_-1_VL0006_-1 417.8571 NA NA
## 6  COUNTRY1 9999 - 4 GSA99 FPN_-1_VL0006_-1 128.1429 NA NA
##   effKwDays effGtDays daysAtSea
## 1    3071.25 133.71429      NA
## 2    1228.50  53.48571      NA
## 3    3685.50 160.45714      NA
## 4    3071.25 133.71429      NA
## 5    3071.25 133.71429      NA
## 6     941.85  41.00571      NA
```

extract COUNTRY and YEAR

```
COUNTRY<-unique(fri_cl1@cl$landCtry)
```

```
YEAR=unique(fri_cl1@cl$year)
```

DG MARE Med&BS DISCARDS Table template

```
dis.temp2<- read.table("xxx_DISCARDS.csv",sep="," ,header=T)
names(dis.temp2)
```

```
## [1] "ID" "COUNTRY" "YEAR"
## [4] "QUARTER" "VESSEL_LENGTH" "GEAR"
## [7] "MESH_SIZE_RANGE" "FISHERY" "AREA"
## [10] "SPECON" "SPECIES" "DISCARDS"
## [13] "UNIT" "LENGTHCLASS0" "LENGTHCLASS1"
## [16] "LENGTHCLASS2" "LENGTHCLASS3" "LENGTHCLASS4"
## [19] "LENGTHCLASS5" "LENGTHCLASS6" "LENGTHCLASS7"
## [22] "LENGTHCLASS8" "LENGTHCLASS9" "LENGTHCLASS10"
## [25] "LENGTHCLASS11" "LENGTHCLASS12" "LENGTHCLASS13"
## [28] "LENGTHCLASS14" "LENGTHCLASS15" "LENGTHCLASS16"
## [31] "LENGTHCLASS17" "LENGTHCLASS18" "LENGTHCLASS19"
## [34] "LENGTHCLASS20" "LENGTHCLASS21" "LENGTHCLASS22"
## [37] "LENGTHCLASS23" "LENGTHCLASS24" "LENGTHCLASS25"
## [40] "LENGTHCLASS26" "LENGTHCLASS27" "LENGTHCLASS28"
## [43] "LENGTHCLASS29" "LENGTHCLASS30" "LENGTHCLASS31"
## [46] "LENGTHCLASS32" "LENGTHCLASS33" "LENGTHCLASS34"
## [49] "LENGTHCLASS35" "LENGTHCLASS36" "LENGTHCLASS37"
## [52] "LENGTHCLASS38" "LENGTHCLASS39" "LENGTHCLASS40"
## [55] "LENGTHCLASS41" "LENGTHCLASS42" "LENGTHCLASS43"
## [58] "LENGTHCLASS44" "LENGTHCLASS45" "LENGTHCLASS46"
## [61] "LENGTHCLASS47" "LENGTHCLASS48" "LENGTHCLASS49"
## [64] "LENGTHCLASS50" "LENGTHCLASS51" "LENGTHCLASS52"
## [67] "LENGTHCLASS53" "LENGTHCLASS54" "LENGTHCLASS55"
## [70] "LENGTHCLASS56" "LENGTHCLASS57" "LENGTHCLASS58"
## [73] "LENGTHCLASS59" "LENGTHCLASS60" "LENGTHCLASS61"
## [76] "LENGTHCLASS62" "LENGTHCLASS63" "LENGTHCLASS64"
## [79] "LENGTHCLASS65" "LENGTHCLASS66" "LENGTHCLASS67"
## [82] "LENGTHCLASS68" "LENGTHCLASS69" "LENGTHCLASS70"
## [85] "LENGTHCLASS71" "LENGTHCLASS72" "LENGTHCLASS73"
## [88] "LENGTHCLASS74" "LENGTHCLASS75" "LENGTHCLASS76"
## [91] "LENGTHCLASS77" "LENGTHCLASS78" "LENGTHCLASS79"
```

```
## [94] "LENGTHCLASS80"      "LENGTHCLASS81"      "LENGTHCLASS82"
## [97] "LENGTHCLASS83"      "LENGTHCLASS84"      "LENGTHCLASS85"
## [100] "LENGTHCLASS86"      "LENGTHCLASS87"      "LENGTHCLASS88"
## [103] "LENGTHCLASS89"      "LENGTHCLASS90"      "LENGTHCLASS91"
## [106] "LENGTHCLASS92"      "LENGTHCLASS93"      "LENGTHCLASS94"
## [109] "LENGTHCLASS95"      "LENGTHCLASS96"      "LENGTHCLASS97"
## [112] "LENGTHCLASS98"      "LENGTHCLASS99"      "LENGTHCLASS100_PLUS"
```

```
fishery<- read.table("communicationTable_for_fishery.csv",sep=";",
                     header=T)
```

```
head(fishery)
```

```
## SDEF_codification DGMARE_Med_BS_codification
## 1                MOL                      MOL
## 2                DES                      DEMSP
## 3                DWS                      DWS
## 4                MDD                      MDD
## 5                SPF                      SPF
## 6                FIF                      FIF
```

Auxiliary table: species_DIS.csv

```
# species file : selected species with FAO three alpha code
```

```
sel_spe <-read.table("species_DISCARDS.csv",sep=";",header=T)
head(sel_spe)
```

```
##          SPECIES SPE   GSA LC_RANGE FISHERY type landSp
## 1 Merluccius merluccius HKE GSA99      10     -1 trip    NA
## 2 Mullus barbatus MUT GSA99      10     -1 trip    NA
```

Data analysis- raising

Analysis by stock

```
for (i in 1:dim(sel_spe)[1]) {

  STK<- sel_spe$SPECIES[i]

  AREA <- sel_spe$GSA[i]

  fri_csc1<- subset(fri_csc, space==sel_spe$GSA[i],table="ca",link=T)
  fri_clc1<- subset(fri_clc, space==sel_spe$GSA[i],table="cl")

  fri_cec1<- subset(fri_cec, space==sel_spe$GSA[i],table="ce")

# The first step is to create the empty object, that will be given
# the appropriate values for the descriptor fields.

  DIS_dbe <- dbEObject(desc= paste(STK, AREA,"Discards", sep="_"),
                       species=STK,
                       catchCat="DIS",
```

```

        strataDesc=fri_strD1,
        methodDesc="analytical"
    )

    if (sel_spe$type[i]=="landings" ) {

DIS_dbe <- totVolume(DIS_dbe,fri_csc1,fri_cec1, fri_clc1,
    type=sel_spe$type[i],val="nAtLength",landSpp=sel_spe$landSpp[i])
    } else {
        DIS_dbe <- totVolume(DIS_dbe,fri_csc1,fri_cec1, type=sel_spe$type[i],
            val="nAtLength")
    }

    # totalW$estim : total weight,
aa <-DIS_dbe@totalW$estim

aa$value<- aa$value/1000 # tons

aa<- rename(aa, "totalW"=value)
    # LenStruc$estim : numbers-at-length estimates,

bb<- DIS_dbe@lenStruc$estim

bb$length=as.numeric(bb$length)

    # define LCs and UNIT Len
UNIT <- as.character( unique(fri_csc@ca$lenCode[fri_csc@ca$spp==STK]) )

if (UNIT %in% c("mm", "MM")& sel_spe$LC_RANGE[i]==10) {
bb$length<-as.numeric(bb$length)/10
UNIT1<- "cm"
}

if (UNIT %in% c("mm", "MM") & sel_spe$LC_RANGE[i]==1) {
bb$length<-as.numeric(bb$length)
UNIT1<- "mm"
}

if (UNIT %in% c("mm", "MM")& sel_spe$LC_RANGE[i]==5) {
bb$length<-as.numeric(bb$length)/10
UNIT1<- "cm"
}

if (UNIT %in% c("cm", "CM") ) {
bb$length<-as.numeric(bb$length)
UNIT1<- "cm"
}
bb$length<- plyr::round_any( bb$length,1,floor)

```

```

bb$value<- bb$value/1000 # '000 ind

ab=left_join(bb,aa ,by = c("time", "space", "technical"))

ab<- ab %>% separate(technical, c("gear","FISHERY", "VL","MESH_SIZE_RANGE"),
                      sep = "_",remove=T)

ab$length<- as.numeric(as.character(ab$length))

ab<- ab%>% group_by(time, space , gear ,FISHERY, VL,MESH_SIZE_RANGE ) %>%
  mutate(minlc=min(length,na.rm=T),maxlc=max(length,na.rm=T))

# matrix with all combinations of "time" "space" "gear" "VL"
# "length" , "MESH_SIZE_RANGE"

dt <- as.data.table(ab)

dt[,c(1:6)][is.na(dt[,c(1:6)])]<- -1

seq_1 <- seq(0, max(dt$length,na.rm = T), by = 1) #

dt$id<- paste(dt$time,dt$space,dt$gear,dt$FISHERY,dt$VL,
             dt$MESH_SIZE_RANGE,sep=":")

dt1<- dt[, list(length = seq_1), by = id]

dt1<- dt1 %>% separate(id, c("time", "space", "gear", "FISHERY","VL",
                           "MESH_SIZE_RANGE"), sep = ":")

# ab[,`1:6`]: NA-->-1
ab<- ab %>%ungroup()%>% mutate_at(vars(c(time:MESH_SIZE_RANGE) ),
  funs( ifelse( is.na(.), -1, .) ) )

dt2<- left_join(dt1,ab)
dt2$stock<- STK

##

dt3 <- data.table::dcast(dt2,as.formula(paste(paste(names(dt2)[! names(dt2) %in%
c("length","value")], collapse='+'), "length", sep=~")),
value.var = "value")

dt3=dt3[complete.cases(dt3[,c(7:9)]), ]

dt3 <- dt3 %>% separate(time, c("Year","Quarter")," - ")

dt3$MESH_SIZE_RANGE<-as.character(dt3$MESH_SIZE_RANGE)

```

```

# numbers at LC : NA-->0
dt3<- dt3 %>% mutate_at(vars( -(Year:stock) ),
  funs( if_else( is.na(.), 0, .) ) )

DISCARDS <- data.frame(

  ID = NA ,
  COUNTRY = COUNTRY ,
  YEAR = YEAR ,
  QUARTER =dt3$Quarter,
  VESSEL_LENGTH = dt3$VL,
  GEAR = dt3$gear,
  MESH_SIZE_RANGE = dt3$MESH_SIZE_RANGE,
  FISHERY= dt3$FISHERY ,
  AREA = sel_spe$GSA[i],
  SPECON = "",
  SPECIES = STK ,
  DISCARDS = dt3$totalW ,
  UNIT = UNIT1
)

DISCARDS<-left_join(DISCARDS,dt3[, -c(1,3,8:11)],by=c( "QUARTER" ="Quarter" ,
  "GEAR"="gear" , "VESSEL_LENGTH" = "VL" , "MESH_SIZE_RANGE", "FISHERY" ))

# take care of number of Length classes (max is 100 acc. to JRC template)
zz<-dim(DISCARDS[-c(1:13)])[2]
names(DISCARDS)[-c(1:13)]<- paste("LENGTHCLASS",seq(0,zz-1,1),sep="")

if(zz>=100){
  DISCARDS$LENGTHCLASS100_PLUS<- rowSums(DISCARDS[,!1:113],na.rm = T)
  DISCARDS<-DISCARDS %>% select(ID:LENGTHCLASS99,LENGTHCLASS100_PLUS)
}

# FISHERY to DG MARE Med&BS specification
DISCARDS$FISHERY <- fishery$SDEF_codification[match(DISCARDS$FISHERY ,
  fishery$DGMARE_Med_BS_codification)]

# species to FAO three alpha code and set ID (COUNTRY, AREA, GEAR,
# VESSEL_LENGTH, MESH_SIZE_RANGE,QUARTER, SPECIES)

dis.tab <-DISCARDS %>% mutate(SPECIES=sel_spe$SPE[match(SPECIES,sel_spe$SPECIES)],
  ID = paste(COUNTRY, AREA, GEAR,FISHERY, VESSEL_LENGTH,
    MESH_SIZE_RANGE,YEAR, QUARTER, SPECIES, sep = "_"))

dis.temp2<-bind_rows(dis.temp2,dis.tab)

# col after 13: set -1 or NA to 0
dis.temp2[, -c(1:13)][is.na(dis.temp2[, -c(1:13)])] <- 0

```

```

dis.temp2<-setDT(dis.temp2)
for (jj in c(14:114)) set(dis.temp2, i = which(dis.temp2[[jj]]== -1), j = jj, v = 0
)

dis.temp2<-setDF(dis.temp2)

}

```

Output

```

write.table(format(dis.temp2,digits=3,scientific=F),
            file = "DISCARDS.csv",row.names=FALSE,sep=";",na="-1")

```

```
head(dis.temp2)
```

```

##                                ID  COUNTRY YEAR  QUARTER
## 1 COUNTRY1_GSA99_OTB_NONE_VL2440_-1_9999_1_HKE COUNTRY1 9999      1
## 2 COUNTRY1_GSA99_GNS_NONE_VL0612_-1_9999_3_HKE COUNTRY1 9999      3
## 3 COUNTRY1_GSA99_OTB_NONE_VL2440_-1_9999_3_HKE COUNTRY1 9999      3
## 4 COUNTRY1_GSA99_FPO_NONE_VL0612_-1_9999_4_HKE COUNTRY1 9999      4
## 5 COUNTRY1_GSA99_GNS_NONE_VL0006_-1_9999_4_HKE COUNTRY1 9999      4
## 6 COUNTRY1_GSA99_GNS_NONE_VL0612_-1_9999_4_HKE COUNTRY1 9999      4
##  VESSEL_LENGTH GEAR MESH_SIZE_RANGE FISHERY  AREA SPECON SPECIES DISCARDS
## 1      VL2440  OTB              -1    NONE GSA99              HKE 5.738667
## 2      VL0612  GNS              -1    NONE GSA99              HKE 0.000000
## 3      VL2440  OTB              -1    NONE GSA99              HKE 8.398101
## 4      VL0612  FPO              -1    NONE GSA99              HKE 0.000000
## 5      VL0006  GNS              -1    NONE GSA99              HKE 1.388400
## 6      VL0612  GNS              -1    NONE GSA99              HKE 2.693500
##  UNIT LENGTHCLASS0 LENGTHCLASS1 LENGTHCLASS2 LENGTHCLASS3 LENGTHCLASS4
## 1    cm              0              0              0              0              0
## 2    cm              0              0              0              0              0
## 3    cm              0              0              0              0              0
## 4    cm              0              0              0              0              0
## 5    cm              0              0              0              0              0
## 6    cm              0              0              0              0              0
##  LENGTHCLASS5 LENGTHCLASS6 LENGTHCLASS7 LENGTHCLASS8 LENGTHCLASS9
## 1              0              0              0      0.00000      0.00000
## 2              0              0              0      0.00000      0.00000
## 3              0              0              0     16.53608     73.01889
## 4              0              0              0      0.00000      0.00000
## 5              0              0              0      0.00000      0.00000
## 6              0              0              0      0.00000      0.00000
##  LENGTHCLASS10 LENGTHCLASS11 LENGTHCLASS12 LENGTHCLASS13 LENGTHCLASS14
## 1          0.0000          3.2000     11.2000000      37.33333      77.8666667
## 2          0.0000          0.0000      0.0000000      0.00000      0.0000000
## 3       164.6176       209.2093     152.5407338     112.59401      41.0615130
## 4          0.0000          0.0000      0.0000000      0.00000      0.0000000
## 5          0.0000          0.0000      0.0000000      0.00000      0.0000000
## 6          0.0000          0.0000      0.5271038       1.054208      0.5271038
##  LENGTHCLASS15 LENGTHCLASS16 LENGTHCLASS17 LENGTHCLASS18 LENGTHCLASS19

```


## 1	67.733333	61.866667	20.800000	0.0000000	0.0000000
## 2	0.000000	0.000000	0.000000	0.0000000	0.0000000
## 3	15.235494	0.000000	3.158578	0.0000000	0.0000000
## 4	0.000000	0.000000	0.000000	0.0000000	0.0000000
## 5	0.000000	0.000000	0.000000	0.0000000	0.0000000
## 6	2.635519	3.162623	1.581311	0.5271038	0.5271038
##	LENGTHCLASS20	LENGTHCLASS21	LENGTHCLASS22	LENGTHCLASS23	LENGTHCLASS24
## 1	0.0000000	0.000000	0.000000	0.0000000	0
## 2	0.0000000	0.000000	0.000000	0.0000000	0
## 3	0.0000000	0.000000	0.000000	0.0000000	0
## 4	0.0000000	0.000000	0.000000	0.0000000	0
## 5	0.0000000	0.000000	0.000000	3.6536842	0
## 6	0.5271038	1.581311	1.054208	0.5271038	0
##	LENGTHCLASS25	LENGTHCLASS26	LENGTHCLASS27	LENGTHCLASS28	LENGTHCLASS29
## 1	0.000000	0.000000	0.000000	0.000000	0.000000
## 2	0.000000	0.000000	0.000000	0.000000	0.000000
## 3	0.000000	0.000000	0.000000	0.000000	0.000000
## 4	0.000000	0.000000	0.000000	0.000000	0.000000
## 5	0.000000	0.000000	0.000000	3.653684	0.000000
## 6	3.162623	1.581311	2.108415	1.581311	2.635519
##	LENGTHCLASS30	LENGTHCLASS31	LENGTHCLASS32	LENGTHCLASS33	LENGTHCLASS34
## 1	0.000000	0.0000000	0.0000000	0.000000	0
## 2	0.000000	0.0000000	0.0000000	0.000000	0
## 3	0.000000	0.0000000	0.0000000	0.000000	0
## 4	0.000000	0.0000000	0.0000000	0.000000	0
## 5	0.000000	0.0000000	0.0000000	3.653684	0
## 6	1.581311	0.5271038	0.5271038	0.000000	0
##	LENGTHCLASS35	LENGTHCLASS36	LENGTHCLASS37	LENGTHCLASS38	LENGTHCLASS39
## 1	0	0	0	0.0000000	0
## 2	0	0	0	0.0000000	0
## 3	0	0	0	0.0000000	0
## 4	0	0	0	0.0000000	0
## 5	0	0	0	0.0000000	0
## 6	0	0	0	0.5271038	0
##	LENGTHCLASS40	LENGTHCLASS41	LENGTHCLASS42	LENGTHCLASS43	LENGTHCLASS44
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS45	LENGTHCLASS46	LENGTHCLASS47	LENGTHCLASS48	LENGTHCLASS49
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS50	LENGTHCLASS51	LENGTHCLASS52	LENGTHCLASS53	LENGTHCLASS54
## 1	0	0	0	0	0
## 2	0	0	0	0	0

## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS55	LENGTHCLASS56	LENGTHCLASS57	LENGTHCLASS58	LENGTHCLASS59
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS60	LENGTHCLASS61	LENGTHCLASS62	LENGTHCLASS63	LENGTHCLASS64
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS65	LENGTHCLASS66	LENGTHCLASS67	LENGTHCLASS68	LENGTHCLASS69
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS70	LENGTHCLASS71	LENGTHCLASS72	LENGTHCLASS73	LENGTHCLASS74
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS75	LENGTHCLASS76	LENGTHCLASS77	LENGTHCLASS78	LENGTHCLASS79
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS80	LENGTHCLASS81	LENGTHCLASS82	LENGTHCLASS83	LENGTHCLASS84
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS85	LENGTHCLASS86	LENGTHCLASS87	LENGTHCLASS88	LENGTHCLASS89
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0

```
## 5      0      0      0      0      0
## 6      0      0      0      0      0
##  LENGTHCLASS90 LENGTHCLASS91 LENGTHCLASS92 LENGTHCLASS93 LENGTHCLASS94
## 1      0      0      0      0      0
## 2      0      0      0      0      0
## 3      0      0      0      0      0
## 4      0      0      0      0      0
## 5      0      0      0      0      0
## 6      0      0      0      0      0
##  LENGTHCLASS95 LENGTHCLASS96 LENGTHCLASS97 LENGTHCLASS98 LENGTHCLASS99
## 1      0      0      0      0      0
## 2      0      0      0      0      0
## 3      0      0      0      0      0
## 4      0      0      0      0      0
## 5      0      0      0      0      0
## 6      0      0      0      0      0
##  LENGTHCLASS100_PLUS
## 1      0
## 2      0
## 3      0
## 4      0
## 5      0
## 6      0
```

Example of use of the script A_CATCH

Settings

```
path_in <- paste("C:\\Users\\Bitetto Isabella\\OneDrive - Coispa Tecnologia & Rice
rca S.C.A.R.L\\MARE22\\STREAM\\FINAL REVISION OF DELIVERABLES\\SDEF_to_DG_MARE_Med
BS", sep="")
```

```
setwd(path_in)
```

Input Data

Validated csData , ceData, and clData

```
load("fri-TEST.Rdata")
```

```
head(fri_cs1@hh$foCatEu5)
```

```
## [1] "OTB_-1_VL2440_-1" "OTB_-1_VL2440_-1" "OTB_-1_VL2440_-1"
## [4] "OTB_-1_VL2440_-1" "OTB_-1_VL2440_-1" "OTB_-1_VL2440_-1"
```

```
head(fri_cl1@cl$foCatEu5)
```

```

## [1] "GNS_-1_VL1218_-1" "LLS_-1_VL1218_-1" "OTB_-1_VL1824_-1"
## [4] "OTB_-1_VL1824_-1" "OTB_-1_VL2440_-1" "OTB_-1_VL2440_-1"

head(fri_cev@ce$foCatEu5)

## [1] "FPN_-1_VL0006_-1" "FPN_-1_VL0006_-1" "FPN_-1_VL0006_-1"
## [4] "FPN_-1_VL0006_-1" "FPN_-1_VL0006_-1" "FPN_-1_VL0006_-1"

fri_strD1 <- strIni(timeStrata="quarter", techStrata = "foCatEu5", spaceStrata = "area")

fri_strD1

## An object of class "strIni"
## Slot "timeStrata":
## [1] "quarter"
##
## Slot "spaceStrata":
## [1] "area"
##
## Slot "techStrata":
## [1] "foCatEu5"
##
## Slot "tpRec":
## [[1]]
## [1] NA
##
##
## Slot "spRec":
## [[1]]
## [1] NA
##
##
## Slot "tcRec":
## [[1]]
## [1] NA

fri_csc <- csDataCons(fri_cs1, fri_strD1)

fri_clc <- clDataCons(fri_cl1, fri_strD1)

fri_cec <- ceDataCons(fri_cev, fri_strD1)

head(fri_csc )

## An object of class "csDataCons"
## Slot "desc":
## [1] "Unknown stock"
##
## Slot "tr":
##   PSUid      time space      technical sampType landCtry vsFlgCtry proj
## 1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1          S COUNTRY1  COUNTRY1 DCF

```

```

## 2      2 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1  DCF
## 3      3 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1  DCF
## 4      4 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1  DCF
## 5      5 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1  DCF
## 6      6 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1  DCF
##      trpCode foNum daysAtSea vslId sampCtry sampMeth
## 1      2      3          1      NA COUNTRY1 Observer
## 2      1      2          1      NA COUNTRY1 Observer
## 3      3      3          1      NA COUNTRY1 Observer
## 4      4      3          1      NA COUNTRY1 Observer
## 5      5      3          1      NA COUNTRY1 Observer
## 6      8      3          1      NA COUNTRY1 Observer
##
## Slot "hh":
##      PSUid SSUid      time space      technical sampType landCtry vslFlgCtry
## 1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1
## 2      1      2 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1
## 3      1      3 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1
## 4      2      1 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1
## 5      2      2 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1
## 6      3      1 9999 - 1 GSA99 OTB_-1_VL2440_-1      S COUNTRY1  COUNTRY1
##      proj trpCode staNum foVal aggLev catReg sppReg      date      foDur
## 1 DCF      2      1      V      H      All      All 2017-02-20 3.500000
## 2 DCF      2      2      V      H      All      All 2017-02-20 1.750000
## 3 DCF      2      3      V      H      All      All 2017-02-20 3.750000
## 4 DCF      1      1      V      H      All      All 2017-02-21 6.083333
## 5 DCF      1      2      V      H      All      All 2017-02-21 5.916667
## 6 DCF      3      1      V      H      All      All 2017-02-23 5.083333
##      latIni lonIni latFin lonFin foDep
## 1      NA      NA      NA      NA      NA
## 2      NA      NA      NA      NA      NA
## 3      NA      NA      NA      NA      NA
## 4      NA      NA      NA      NA      NA
## 5      NA      NA      NA      NA      NA
## 6      NA      NA      NA      NA      NA
##
## Slot "sl":
##      PSUid SSUid TSUid      time space      technical      sort sampType
## 1      1      1      3 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 2      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 DIS-HUC-NA-NA      S
## 3      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 DIS-HUC-NA-NA      S
## 4      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 5      1      1      4 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 6      1      2      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
##      landCtry vslFlgCtry proj trpCode staNum      spp sex wt
## 1 COUNTRY1  COUNTRY1 DCF      2      1 Merluccius merluccius <NA> 2400
## 2 COUNTRY1  COUNTRY1 DCF      2      1 Merluccius merluccius <NA> 90
## 3 COUNTRY1  COUNTRY1 DCF      2      1 Mullus barbatus <NA> 60
## 4 COUNTRY1  COUNTRY1 DCF      2      1 Mullus barbatus <NA> 2020
## 5 COUNTRY1  COUNTRY1 DCF      2      1 Merluccius merluccius <NA> 110
## 6 COUNTRY1  COUNTRY1 DCF      2      2 Mullus barbatus <NA> 350

```

```

##      subSampWt lenCode
## 1      2400      mm
## 2       45      mm
## 3       30      mm
## 4     2020      mm
## 5      110      mm
## 6      350      mm
##
## Slot "hl":
##      PSuid SSuid TSuid      time space      technical      sort sampType
## 1      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 2      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 3      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 4      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 5      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
## 6      1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA-NA      S
##      landCtry vs1FlgCtry proj trpCode staNum      spp sex lenCls
## 1 COUNTRY1 COUNTRY1 DCF      2      1 Mullus barbatus <NA> 170
## 2 COUNTRY1 COUNTRY1 DCF      2      1 Mullus barbatus <NA> 180
## 3 COUNTRY1 COUNTRY1 DCF      2      1 Mullus barbatus <NA> 150
## 4 COUNTRY1 COUNTRY1 DCF      2      1 Mullus barbatus <NA> 160
## 5 COUNTRY1 COUNTRY1 DCF      2      1 Mullus barbatus <NA> 140
## 6 COUNTRY1 COUNTRY1 DCF      2      1 Mullus barbatus <NA> 200
##      lenNum
## 1      7
## 2      7
## 3      7
## 4      3
## 5      7
## 6      3
##
## Slot "ca":
##      PSuid SSuid      time space      technical      sort sampType landCtry
## 1      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA      S COUNTRY1
## 2      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA      S COUNTRY1
## 3      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA      S COUNTRY1
## 4      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA      S COUNTRY1
## 5      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA      S COUNTRY1
## 6      1      1 9999 - 1 GSA99 OTB_-1_VL2440_-1 LAN-HUC-NA      S COUNTRY1
##      vs1FlgCtry proj trpCode staNum      spp sex
## 1 COUNTRY1 DCF      2      1 Merluccius merluccius U
## 2 COUNTRY1 DCF      2      1 Mullus barbatus U
## 3 COUNTRY1 DCF      2      1 Mullus barbatus U
## 4 COUNTRY1 DCF      2      1 Merluccius merluccius U
## 5 COUNTRY1 DCF      2      1 Mullus barbatus U
## 6 COUNTRY1 DCF      2      1 Mullus barbatus U
##
##      stock lenCls age fishId lenCode ageMeth plusGrp otoWt
## 1 Merluccius merluccius_22 170 NA 20 mm OWR <NA> NA
## 2 Mullus barbatus_22 130 NA 37 mm OWR <NA> NA
## 3 Mullus barbatus_22 170 NA 7 mm OWR <NA> NA
## 4 Merluccius merluccius_22 160 NA 19 mm OWR <NA> NA

```

```
## 5 Mullus barbatus_22 160 NA 36 mm OWR <NA> NA
## 6 Mullus barbatus_22 110 NA 22 mm OWR <NA> NA
## otoSide indWt matMeth matScale matStage
## 1 <NA> NA <NA> <NA> <NA>
## 2 <NA> NA <NA> <NA> <NA>
## 3 <NA> NA <NA> <NA> <NA>
## 4 <NA> NA <NA> <NA> <NA>
## 5 <NA> NA <NA> <NA> <NA>
## 6 <NA> NA <NA> <NA> <NA>
```

`head(fri_clc)`

```
## An object of class "clDataCons"
## Slot "desc":
## [1] "Unknown stock"
##
## Slot "cl":
## landCtry vs1FlgCtry time space technical
## 1 COUNTRY1 COUNTRY1 9999 - 1 GSA99 GNS_-1_VL1218_-1
## 2 COUNTRY1 COUNTRY1 9999 - 1 GSA99 LLS_-1_VL1218_-1
## 3 COUNTRY1 COUNTRY1 9999 - 1 GSA99 OTB_-1_VL1824_-1
## 4 COUNTRY1 COUNTRY1 9999 - 1 GSA99 OTB_-1_VL1824_-1
## 5 COUNTRY1 COUNTRY1 9999 - 1 GSA99 OTB_-1_VL2440_-1
## 6 COUNTRY1 COUNTRY1 9999 - 1 GSA99 OTB_-1_VL2440_-1
##
## taxon landCat commCatSc1 commCat unallocCatchWt
## 1 Merluccius merluccius HUC <NA> <NA> NA
## 2 Merluccius merluccius HUC <NA> <NA> NA
## 3 Merluccius merluccius HUC <NA> <NA> NA
## 4 Mullus barbatus HUC <NA> <NA> NA
## 5 Merluccius merluccius HUC <NA> <NA> NA
## 6 Mullus barbatus HUC <NA> <NA> NA
## misRepCatchWt landWt landMult landValue
## 1 NA 46 1 NA
## 2 NA 140 1 NA
## 3 NA 2151 1 NA
## 4 NA 2285 1 NA
## 5 NA 32979 1 NA
## 6 NA 9144 1 NA
```

`head(fri_cec)`

```
## An object of class "ceDataCons"
## Slot "desc":
## [1] "Unknown stock"
##
## Slot "ce":
## vs1FlgCtry time space technical trpNum foNum foDur
## 1 COUNTRY1 9999 - 2 GSA99 FPN_-1_VL0006_-1 417.8571 NA NA
## 2 COUNTRY1 9999 - 2 GSA99 FPN_-1_VL0006_-1 167.1429 NA NA
## 3 COUNTRY1 9999 - 3 GSA99 FPN_-1_VL0006_-1 501.4286 NA NA
## 4 COUNTRY1 9999 - 3 GSA99 FPN_-1_VL0006_-1 417.8571 NA NA
## 5 COUNTRY1 9999 - 3 GSA99 FPN_-1_VL0006_-1 417.8571 NA NA
```

```
## 6    COUNTRY1 9999 - 4 GSA99 FPN_-1_VL0006_-1 128.1429    NA    NA
##    effKwDays effGtDays daysAtSea
## 1    3071.25 133.71429    NA
## 2    1228.50  53.48571    NA
## 3    3685.50 160.45714    NA
## 4    3071.25 133.71429    NA
## 5    3071.25 133.71429    NA
## 6    941.85  41.00571    NA
```

extract COUNTRY and YEAR

```
COUNTRY<-unique(fri_cl1@cl$landCtry)
```

```
YEAR=unique(fri_cl1@cl$year)
```

DG MARE Med&BS DISCARDS Table template

```
catch.temp2<- read.table("xxx_CATCH.csv",sep="," ,header=T)
```

```
names(catch.temp2)
```

```
##    [1] "ID"
##    [3] "YEAR"
##    [5] "VESSEL_LENGTH"
##    [7] "MESH_SIZE_RANGE"
##    [9] "AREA"
##   [11] "SPECIES"
##   [13] "DISCARDS"
##   [15] "NO_LENGTH_MEASUREMENTS_LANDINGS"
##   [17] "NO_SAMPLES_DISCARDS"
##   [19] "NO_AGE_MEASUREMENTS_DISCARDS"
##   [21] "NO_LENGTH_MEASUREMENTS_CATCH"
##   [23] "MIN_AGE"
##   [25] "AGE_0"
##   [27] "AGE_0_MEAN_WEIGHT_LANDED"
##   [29] "AGE_0_NO_DISCARD"
##   [31] "AGE_0_MEAN_LENGTH_DISCARD"
##   [33] "AGE_1_NO_LANDED"
##   [35] "AGE_1_MEAN_WEIGHT_LANDED"
##   [37] "AGE_1_MEAN_LENGTH_DISCARD"
##   [39] "AGE_2"
##   [41] "AGE_2_MEAN_WEIGHT_LANDED"
##   [43] "AGE_2_NO_DISCARD"
##   [45] "AGE_2_MEAN_LENGTH_DISCARD"
##   [47] "AGE_3_NO_LANDED"
##   [49] "AGE_3_MEAN_WEIGHT_LANDED"
##   [51] "AGE_3_MEAN_LENGTH_DISCARD"
##   [53] "AGE_4"
##   [55] "AGE_4_MEAN_WEIGHT_LANDED"
##   [57] "AGE_4_NO_DISCARD"
##   [59] "AGE_4_MEAN_LENGTH_DISCARD"
##   [61] "AGE_5_NO_LANDED"
##   [63] "AGE_5_MEAN_WEIGHT_LANDED"
##   [65] "AGE_5_MEAN_LENGTH_DISCARD"
```


## [67]	"AGE_6"	"AGE_6_NO_LANDED"
## [69]	"AGE_6_MEAN_WEIGHT_LANDED"	"AGE_6_MEAN_LENGTH_LANDED"
## [71]	"AGE_6_NO_DISCARD"	"AGE_6_MEAN_WEIGHT_DISCARD"
## [73]	"AGE_6_MEAN_LENGTH_DISCARD"	"AGE_7"
## [75]	"AGE_7_NO_LANDED"	"AGE_7_MEAN_WEIGHT_LANDED"
## [77]	"AGE_7_MEAN_LENGTH_LANDED"	"AGE_7_NO_DISCARD"
## [79]	"AGE_7_MEAN_WEIGHT_DISCARD"	"AGE_7_MEAN_LENGTH_DISCARD"
## [81]	"AGE_8"	"AGE_8_NO_LANDED"
## [83]	"AGE_8_MEAN_WEIGHT_LANDED"	"AGE_8_MEAN_LENGTH_LANDED"
## [85]	"AGE_8_NO_DISCARD"	"AGE_8_MEAN_WEIGHT_DISCARD"
## [87]	"AGE_8_MEAN_LENGTH_DISCARD"	"AGE_9"
## [89]	"AGE_9_NO_LANDED"	"AGE_9_MEAN_WEIGHT_LANDED"
## [91]	"AGE_9_MEAN_LENGTH_LANDED"	"AGE_9_NO_DISCARD"
## [93]	"AGE_9_MEAN_WEIGHT_DISCARD"	"AGE_9_MEAN_LENGTH_DISCARD"
## [95]	"AGE_10"	"AGE_10_NO_LANDED"
## [97]	"AGE_10_MEAN_WEIGHT_LANDED"	"AGE_10_MEAN_LENGTH_LANDED"
## [99]	"AGE_10_NO_DISCARD"	"AGE_10_MEAN_WEIGHT_DISCARD"
## [101]	"AGE_10_MEAN_LENGTH_DISCARD"	"AGE_11"
## [103]	"AGE_11_NO_LANDED"	"AGE_11_MEAN_WEIGHT_LANDED"
## [105]	"AGE_11_MEAN_LENGTH_LANDED"	"AGE_11_NO_DISCARD"
## [107]	"AGE_11_MEAN_WEIGHT_DISCARD"	"AGE_11_MEAN_LENGTH_DISCARD"
## [109]	"AGE_12"	"AGE_12_NO_LANDED"
## [111]	"AGE_12_MEAN_WEIGHT_LANDED"	"AGE_12_MEAN_LENGTH_LANDED"
## [113]	"AGE_12_NO_DISCARD"	"AGE_12_MEAN_WEIGHT_DISCARD"
## [115]	"AGE_12_MEAN_LENGTH_DISCARD"	"AGE_13"
## [117]	"AGE_13_NO_LANDED"	"AGE_13_MEAN_WEIGHT_LANDED"
## [119]	"AGE_13_MEAN_LENGTH_LANDED"	"AGE_13_NO_DISCARD"
## [121]	"AGE_13_MEAN_WEIGHT_DISCARD"	"AGE_13_MEAN_LENGTH_DISCARD"
## [123]	"AGE_14"	"AGE_14_NO_LANDED"
## [125]	"AGE_14_MEAN_WEIGHT_LANDED"	"AGE_14_MEAN_LENGTH_LANDED"
## [127]	"AGE_14_NO_DISCARD"	"AGE_14_MEAN_WEIGHT_DISCARD"
## [129]	"AGE_14_MEAN_LENGTH_DISCARD"	"AGE_15"
## [131]	"AGE_15_NO_LANDED"	"AGE_15_MEAN_WEIGHT_LANDED"
## [133]	"AGE_15_MEAN_LENGTH_LANDED"	"AGE_15_NO_DISCARD"
## [135]	"AGE_15_MEAN_WEIGHT_DISCARD"	"AGE_15_MEAN_LENGTH_DISCARD"
## [137]	"AGE_16"	"AGE_16_NO_LANDED"
## [139]	"AGE_16_MEAN_WEIGHT_LANDED"	"AGE_16_MEAN_LENGTH_LANDED"
## [141]	"AGE_16_NO_DISCARD"	"AGE_16_MEAN_WEIGHT_DISCARD"
## [143]	"AGE_16_MEAN_LENGTH_DISCARD"	"AGE_17"
## [145]	"AGE_17_NO_LANDED"	"AGE_17_MEAN_WEIGHT_LANDED"
## [147]	"AGE_17_MEAN_LENGTH_LANDED"	"AGE_17_NO_DISCARD"
## [149]	"AGE_17_MEAN_WEIGHT_DISCARD"	"AGE_17_MEAN_LENGTH_DISCARD"
## [151]	"AGE_18"	"AGE_18_NO_LANDED"
## [153]	"AGE_18_MEAN_WEIGHT_LANDED"	"AGE_18_MEAN_LENGTH_LANDED"
## [155]	"AGE_18_NO_DISCARD"	"AGE_18_MEAN_WEIGHT_DISCARD"
## [157]	"AGE_18_MEAN_LENGTH_DISCARD"	"AGE_19"
## [159]	"AGE_19_NO_LANDED"	"AGE_19_MEAN_WEIGHT_LANDED"
## [161]	"AGE_19_MEAN_LENGTH_LANDED"	"AGE_19_NO_DISCARD"
## [163]	"AGE_19_MEAN_WEIGHT_DISCARD"	"AGE_19_MEAN_LENGTH_DISCARD"
## [165]	"AGE_20_PLUS"	"AGE_20_PLUS_NO_LANDED"
## [167]	"AGE_20_PLUS_MEAN_WEIGHT_LANDED"	"AGE_20_PLUS_MEAN_LENGTH_LANDED"

```
## [169] "AGE_20_PLUS_NO_DISCARD" "AGE_20_PLUS_MEAN_WEIGHT_DISCARD"
## [171] "AGE_20_PLUS_MEAN_LENGTH_DISCARD"
```

Auxiliary table: species_DIS.csv

species file : selected species with FAO three alpha code

```
sel_spe <-read.table("species_CATCH.csv",sep=";",header=T, na.strings = c("", "-1",
, " ", ".", "NA"))
head(sel_spe)
```

```
##          SPECIES SPE LC_RANGE   GSA typeALK mcrs specon_catch type
## 1 Merluccius merluccius HKE      10 GSA99  fixed   NA          NA trip
## 2 Mullus barbatus MUT      10 GSA99  fixed   70          NA trip
## 3 Merluccius merluccius HKE      10 GSA99  fixed  110          NA trip
## landSpp lanEstim_methodDesc methodDesc_LAN.age.wght
## 1      NA      analytical      analytical
## 2      NA      analytical      analytical
## 3      NA      analytical      analytical
## methodDesc_LAN.len.age methodDesc_DIS.age.wght methodDesc_DIS.len.age
## 1      analytical      analytical      analytical
## 2      analytical      analytical      analytical
## 3      analytical      analytical      analytical
## adjust_L.w.a adjust_D.w.a adjust_L.len.a adjust_D.len.a
## 1      TRUE      FALSE      TRUE      FALSE
## 2      FALSE      TRUE      FALSE      TRUE
## 3      TRUE      FALSE      TRUE      FALSE
```

```
fishery<- read.table("communicationTable_for_fishery.csv",sep=";",header=T)
```

```
head(fishery)
```

```
## SDEF_codification DGMARE_Med_BS_codification
## 1      MOL      MOL
## 2      DES      DEMSP
## 3      DWS      DWS
## 4      MDD      MDD
## 5      SPF      SPF
## 6      FIF      FIF
```

Checking data

Check if there are stocks with no age data and impute "-1", otherwise the bpEstim() method for length at age gives error :

```
# get col w. all=NA by spp
aa=fri_csc@ca %>% group_by(spp) %>% summarize_all(~all(is.na(.)))
```

```
# to make it run for Len if age=NA, impute age == "-1"
fri_csc@ca$age[fri_csc@ca$spp %in% aa$spp[aa$age==TRUE]]=-1
```

```
### error if sp-quarter-gear-VL not found in both CS and CL
```

Analysis by stock

```
for (i in 1:dim(sel_spe)[1]) {  
  
  STK<- sel_spe$SPECIES[i]  
  
  AREA <- sel_spe$GSA[i]  
  
  fri_csc1<- subset(fri_csc, space==sel_spe$GSA[i],table="ca",link=T)  
  fri_clc1<- subset(fri_clc, space==sel_spe$GSA[i],table="cl")  
  
  fri_cec1<- subset(fri_cec, space==sel_spe$GSA[i],table="ce")  
  
# Estimating age structure LAN - -----  
  
lanEstim <-  
  dbeObject(  
    desc = paste(STK, "Landings.str", sep="_"),  
    species = STK,  
    catchCat = "LAN",  
    strataDesc = fri_strD1,  
    methodDesc = sel_spe$lanEstim_methodDesc[i]  
  )  
  
  # Length str LAN  
  
if ( sel_spe$lanEstim_methodDesc[i]=="analytical"){  
  lanEstim <- RaiseLgth(lanEstim, fri_csc1, fri_clc1,incl.precision =F)  
} else {  
  lanEstim <- RaiseLgthBoot(lanEstim, fri_csc1, fri_clc1,incl.precision =F,B=15)  
}  
  
# subset Lenstruc for MCRS  
  
if (!is.na(sel_spe$mcrs[i]) & sel_spe$mcrs[i]!=".") {  
  lanEstim@lenStruc$estim =  
  lanEstim@lenStruc$estim[as.numeric(lanEstim@lenStruc$estim$length) <= sel_spe$mcrs[i],]  
  lanEstim@lenVar = lanEstim@lenVar[as.numeric(lanEstim@lenVar$length) <= sel_spe$mcrs[i],]  
}  
  
#####>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>  
Estimation of total numbers-at-age from market sampling  
  
lanEstim <- RaiseAge(lanEstim, fri_csc1, fri_clc1, type = sel_spe$typeALK[i],  
                    strataDesc = fri_strD1)
```

```

# Estimating age structure DIS - -----

DIS_dbe <- dbeObject(desc= paste(STK, AREA,"Discards", sep="_"),
                    species=STK,
                    catchCat="DIS",
                    strataDesc=fri_strD1,
                    methodDesc="analytical")

# discards raising

if (sel_spe$type[i]=="landings" ) {
  DIS_dbe <- totVolume(DIS_dbe,fri_csc1,fri_cec1, fri_clc1,
                      type=sel_spe$type[i],landSpp=sel_spe$landSpp[i])
} else {
  DIS_dbe <- totVolume(DIS_dbe,fri_csc1,fri_cec1, type=sel_spe$type[i])
}

DIS_dbe <- RaiseAge(DIS_dbe, fri_csc1, fri_clc1,type = sel_spe$typeALK[i],
                   strataDesc = fri_strD1)

# CATCH1 : cols ID: MAX_AGE -----
---

# NO SAMPLES -----
-

### No Samples == No trips (see ANNEX2- DG MARE Med&BS data call spec.).
# Note: dbe estimates n.samples as trips*fo , thus the estimation must be based on
HL

newhl<-mergecsData(fri_cs1)@hl %>%
  rename("space"=area, "technical"=foCatEu5) %>%
  mutate(time=paste(year, quarter, sep=" - ")) %>% filter( spp==STK)

if (!is.na(sel_spe$mcrs[i]) & sel_spe$mcrs[i]!=".") { # MCRS

no.samples<- data.frame(newhl) %>% filter(lenCls<=sel_spe$mcrs[i]) %>%
  dplyr::group_by(time,space,technical,catchCat)%>%
  summarize(value=n_distinct(trpCode))

no.samples<- no.samples%>% spread(catchCat,value)

L.no.samples<- no.samples%>% select(-DIS)%>% rename("value"=LAN)
D.no.samples<- no.samples%>% select(-LAN)%>% rename("value"=DIS)

} else {

no.samples<- data.frame(newhl) %>%
  dplyr::group_by(time,space,technical,catchCat)%>%
  summarize(value=n_distinct(trpCode))

```

```

no.samples<- no.samples%>% spread(catchCat,value)

      L.no.samples<- no.samples%>% select(-DIS)%>% rename("value"=LAN)
      D.no.samples<- no.samples%>% select(-LAN)%>% rename("value"=DIS)
}

# MCRS : no age and Len measur. for LAN < MCRS-----

if (!is.na(sel_spe$mcrs[i]) & sel_spe$mcrs[i]!="."){

# No age measurements LAN

newca <- mergecsData(fri_cs1)@ca %>%
  rename("space"=area, "technical"=foCatEu5) %>%
  mutate(time=paste(year, quarter, sep=" - ")) %>% filter( spp==STK)

no.age.meas.lan.mcrs<- data.frame(newca) %>%
  filter(lenCls<=sel_spe$mcrs[i]&!is.na(age)&catchCat=="LAN" )%>%
  dplyr::group_by(time,space,technical)%>% summarize(value=n())

# remove technical if all ==NA
no.age.meas.lan.mcrs<- no.age.meas.lan.mcrs[,!apply(is.na(no.age.meas.lan.mcrs),
2, all)]

# No Len measurements LAN

no.len.meas.lan.mcrs<- data.frame(newca) %>%
  filter(lenCls<=sel_spe$mcrs[i]&!is.na(lenCls)&catchCat=="LAN")%>%
  dplyr::group_by(time,space,technical)%>% summarize(value=n())

}

### end

# separate merge for age samples: may not have technical strata (use : space, time
)

if (!is.na(sel_spe$mcrs[i]) & sel_spe$mcrs[i]!=".") {
  # if mcrs use no.age.meas.lan.mcrs
  list.age.smp<-list(no.age.meas.lan.mcrs,DIS_dbe@nMeas$age)
} else{
  list.age.smp<-list(lanEstim@nMeas$age,DIS_dbe@nMeas$age)
}

names(list.age.smp)<-c("L.nmeas.age", "D.nmeas.age")
list.age.smp.merge = data.table::rbindlist(list.age.smp,id=T)
age.smp= spread(list.age.smp.merge,key=.id,value=value)

# List all remaining output tables (excl. age meas & no samples)

```

```

if (!is.na(sel_spe$mcrcs[i]) & sel_spe$mcrcs[i]!=".") { # if mcrcs use no.len.meas.La
n.mcrcs

list2<- list(no.len.meas.lan.mcrcs,DIS_dbe@nMeas$len,lanEstim@totalN$estim,
            lanEstim@totalW$estim ,DIS_dbe@totalW$estim ,DIS_dbe@totalN$estim,
            L.no.samples,D.no.samples)

} else {

    list2<- list(lanEstim@nMeas$len,DIS_dbe@nMeas$len,
                lanEstim@totalN$estim,lanEstim@totalW$estim,
                DIS_dbe@totalW$estim ,DIS_dbe@totalN$estim,
                L.no.samples,D.no.samples)

}

names(list2)<-c("LnMeas.len", "DnMeas.len", "LtotalN", "LtotalW", "totalWDIS",
               "totalNDIS", "L.no.samples", "D.no.samples")

list3=append(list.age.smp,list2)
list..merge = data.table::rbindlist(list3,id=T,fill=T)

all.merge= spread(list..merge,key=.id,value=value)

all.merge <- all.merge%>% mutate( "stock"=STK) %>% select(stock,everything())

# delete rows (age...) with no "technical": a number should be given only
# if it relates to this fishery only
# (see DG MARE Med&BS Data Call specifiation - Annex 2 - Catch)

all.merge<- all.merge[complete.cases(all.merge$technical), ]

aa.len <- all.merge
aa.len$totalN=all.merge$LtotalN/1000 # '000 ind
aa.len$totalW=all.merge$LtotalW/1000 # tons

aa.len$totalNDIS=all.merge$totalNDIS/1000 # '000 ind
aa.len$totalWDIS=all.merge$totalWDIS/1000 # tons

# AgeStruc : n.at.age LAN / DIS -----

# Landings
bb<-lanEstim@ageStruc$estim

bb$value=bb$value/1000 # '000 ind

bb<-rename(bb, "n.at.age"=value)

# discards
bbd<-DIS_dbe@ageStruc$estim

```

```

bbd$value=bbd$value/1000 # '000 ind

bbd<-rename(bbd, "DIS.n.at.age"=value)

ab<- left_join(bb,bbd) %>% left_join(aa.len)

ab<- ab %>% separate(technical, c("gear","FISHERY", "VL","MESH_SIZE_RANGE"),
                      sep = "_")

# min age/ max age-----

ab$age <- as.numeric(as.character(ab$age))

ab <- ab%>% group_by(time, space , gear,FISHERY , VL ,MESH_SIZE_RANGE ) %>%
  mutate(minage=min(age,na.rm=T),maxage=max(age,na.rm=T))

ab <- ab %>% separate(time, c("Year","Quarter")," - ",remove=F)

# ### >>>>>>>>. catch1: info by row -----

catch1= data.frame(
  ID = NA ,
  COUNTRY =COUNTRY ,
  YEAR = YEAR ,
  QUARTER = ab$Quarter ,
  VESSEL_LENGTH = ab$VL ,
  GEAR = ab$gear ,
  MESH_SIZE_RANGE = ab$MESH_SIZE_RANGE ,
  FISHERY = ab$FISHERY[i] ,
  AREA = AREA ,
  SPECON = sel_spe$specon_catch[i] ,
  SPECIES = STK ,
  LANDINGS = ab$totalW , # MCRS: meanW.at.age * n.at.age
  DISCARDS = ab$totalWDIS,
  NO_SAMPLES_LANDINGS =ab$L.no.samples, # = TRIPS
  NO_LENGTH_MEASUREMENTS_LANDINGS = ab$LnMeas.len ,
  NO_AGE_MEASUREMENTS_LANDINGS = ab$L.nmeas.age ,
  NO_SAMPLES_DISCARDS = ab$D.no.samples , # = TRIPS
  NO_LENGTH_MEASUREMENTS_DISCARDS = ab$DnMeas.len ,
  NO_AGE_MEASUREMENTS_DISCARDS = ab$D.nmeas.age,
  NO_SAMPLES_CATCH = 0,
  NO_LENGTH_MEASUREMENTS_CATCH = 0,
  NO_AGE_MEASUREMENTS_CATCH = 0 ,
  MIN_AGE = ab$minage ,
  MAX_AGE = ab$maxage
)

```

```
# if mcrs delete LANDINGS: estimated below
if (!is.na(sel_spe$mcrs[i]) & sel_spe$mcrs[i]!="."){
  catch1<-catch1 %>% select(-LANDINGS)
}

# ### >>>>>>. catch1: no samples & no meas. -----

# NO_SAMPLES_CATCH - NO_LENGTH_MEASUREMENTS_CATCH -NO_AGE_MEASUREMENTS_CATCH

catch1 <- catch1 %>% mutate(
  NO_SAMPLES_CATCH = rowSums( cbind (NO_SAMPLES_LANDINGS , NO_SAMPLES_DISCARDS),
                                na.rm=TRUE),
  NO_LENGTH_MEASUREMENTS_CATCH = rowSums( cbind (NO_LENGTH_MEASUREMENTS_DISCARDS ,
                                                  NO_LENGTH_MEASUREMENTS_LANDINGS),
                                            na.rm=TRUE),

  NO_AGE_MEASUREMENTS_CATCH=rowSums( cbind (NO_AGE_MEASUREMENTS_DISCARDS,
                                             NO_AGE_MEASUREMENTS_LANDINGS),
                                       na.rm=TRUE)) %>% distinct()

# ##### >>>>> LAN- DIS n.at.age -----

# matrix with all combinations of "time" "space" "gear" "VL"
# "MESH_SIZE_RANGE" "length"

ab[,c(1:8)][is.na(ab[,c(1:8)])]<-1

dt <- as.data.table(ab )

seq_1 <- try(seq(0, 19, by = 1),silent=T)
if(class(seq_1=="try-error")){seq_1=-1}

dt$id<- paste(dt$time,dt$space,dt$gear,dt$FISHERY,dt$VL,dt$MESH_SIZE_RANGE,
              sep=":")
dt1<- dt[, list(age = seq_1), by = id]
dt1<- dt1 %>% separate(id, c("time", "space", "gear", "FISHERY",
                           "VL","MESH_SIZE_RANGE"), sep = ":")

dt2=left_join(dt1,ab %>% select("time" , "space", "gear" , "FISHERY",
                              "VL" , "MESH_SIZE_RANGE", "age", "n.at.age",
                              "DIS.n.at.age", "stock" ))

dt2$stock=STK

# MEAN LENGTH LAND, MEAN WEIGHT LAND -----

#####>>>>>>>>>> LAN weight at age!!
```



```

wtEstim_An <-
  dbeObject(
    desc = "Weights at age",
    species = STK,
    catchCat = "LAN",
    param = "weight",
    strataDesc = fri_strD1, # strBP,
    methodDesc = sel_spe$methodDesc_LAN.age.wght[i]
  )

if(sel_spe$methodDesc_LAN.age.wght[i]=="analytical") {
  wtEstim_An <- bpEstim(wtEstim_An, fri_csc1, adjust = sel_spe$adjust_L.w.a[i])
} else{
  wtEstim_An <- bpBoot(wtEstim_An, fri_csc1, adjust = sel_spe$adjust_L.w.a[i])
}

# # LAN mean weight at age -----

cc=wtEstim_An@ageStruc$estim

cc$value<- cc$value/1000 # g ->kg

cc$age=as.numeric(as.character(cc$age))
cc=rename(cc, "meanW.at.age"=value)
cc=cc %>% separate(technical, c("gear", "FISHERY", "VL", "MESH_SIZE_RANGE"),
                  sep = "_")

# LAN Length at age -----

LEstim_An <-
  dbeObject(
    desc = "Length at age",
    species = STK,
    catchCat = "LAN",
    param = "length",
    strataDesc = fri_strD1, # strBP,
    methodDesc = sel_spe$methodDesc_LAN.len.age[i]
  )

if(sel_spe$methodDesc_LAN.len.age[i]=="analytical") {
  LEstim_An <- bpEstim(LEstim_An, fri_csc1, adjust = sel_spe$adjust_L.len.a[i])
} else{
  LEstim_An <- bpBoot(LEstim_An, fri_csc1, adjust = sel_spe$adjust_L.len.a[i])
}

# LAN mean Length at age -----

```

```

ff=LEstim_An@ageStruc$estim

UNIT <- unique(fri_csc1@ca$lenCode[fri_csc1@ca$spp==STK])
if (UNIT=="mm" || "MM"){
  ff$value <- ff$value/10 # mm-> cm
}

ff$age=as.numeric(as.character(ff$age))
ff=rename(ff, "meanL.at.age"=value)
ff=ff %>% separate(technical, c("gear", "FISHERY", "VL", "MESH_SIZE_RANGE"),
                  sep = "_")

# MEAN LENGTH DISCARD, MEAN WEIGHT DISCARD -----

# DIS weight at age -----

DwtEstim_An <-
  dbeObject(
    desc = "Weights at age",
    species = STK,
    catchCat = "DIS",
    param = "weight",
    strataDesc = fri_strD1, # strBP,
    methodDesc = sel_spe$methodDesc_DIS.age.wght[i]
  )

if(sel_spe$methodDesc_DIS.age.wght[i]=="analytical") {
  DwtEstim_An <- bpEstim(DwtEstim_An, fri_csc1, adjust = sel_spe$adjust_D.w.a[i])
} else{
  DwtEstim_An <- bpBoot(DwtEstim_An, fri_csc1, adjust = sel_spe$adjust_D.w.a[i])
}

# DIS mean weight at age
ccD=DwtEstim_An@ageStruc$estim

ccD$value <- ccD$value/1000 # g -> kg

ccD$age=as.numeric(as.character(ccD$age))
ccD=rename(ccD, "DmeanW.at.age"=value)
ccD=ccD %>% separate(technical, c("gear", "FISHERY", "VL", "MESH_SIZE_RANGE"),
                  sep = "_")

# DIS Length at age -----

DLEstim_An <-

```

```

dbeObject(
  desc = " Length at age",
  species = STK,
  catchCat = "DIS",
  param = "length",
  strataDesc = fri_strD1, # strBP,
  methodDesc = sel_spe$methodDesc_DIS.len.age[i]
)

if(sel_spe$methodDesc_DIS.len.age[i]=="analytical") {
  DLEstim_An <- bpEstim(DLEstim_An, fri_csc1, adjust = sel_spe$adjust_D.len.a[i])
} else{
  DLEstim_An <- bpBoot(DLEstim_An, fri_csc1, adjust = sel_spe$adjust_D.len.a[i])
}

##### >> DIS mean length at age
ffD=DLEstim_An@ageStruc$estim

UNIT <- unique(fri_csc1@ca$lenCode[fri_csc1@ca$spp==STK])
if (UNIT=="mm" || "MM"){
ffD$value <- ffD$value/10 # mm-> cm
}

ffD$age=as.numeric(as.character(ffD$age))
ffD=rename(ffD, "DmeanL.at.age"=value)
ffD=ffD %>% separate(technical, c("gear", "FISHERY", "VL", "MESH_SIZE_RANGE"),
                     sep = "_")

# ## combine: w.age, L.age, no.age (LAN- DIS)... to get CATCH cols -----

l3=list(dt2,cc,ff,ccD,ffD)

l3=lapply(l3, function(x){ x[,c(1:6)][is.na(x[,c(1:6)])]<- -1;return(x)})

cfdt2=Reduce(function(x, y) merge(x, y, by = c("time", "space", "gear" ,
        "FISHERY", "VL", "MESH_SIZE_RANGE" ,"age" ),all.x=T), l3)

dt3 = try(data.table::dcast(setDT(distinct(cfdt2)),
                           time + space + gear +FISHERY+ VL+MESH_SIZE_RANGE ~ age
,
                           value.var = c("n.at.age", "DIS.n.at.age", "meanW.at.age"
,
                           "meanL.at.age", "DmeanW.at.age", "DmeanL.at.age",
                           "age" )),silent=T)

# if mcrs
# Landings= meanW.at.age*n.at.age
if (!is.na(sel_spe$mcrs[i]) & sel_spe$mcrs[i]!="."){

```

```

# tonnes
landings <- cfdt2 %>% mutate(LANDINGS= meanW.at.age*n.at.age) %>%
  select(time, space, gear, FISHERY, VL, MESH_SIZE_RANGE, LANDINGS)

landings <- landings[!is.na(landings$LANDINGS ), ]

dt3<-left_join(dt3,landings)

}

## rename col to match CATCH

names(dt3)[grep("DIS.n.at.age", names(dt3)) ]<-paste("AGE",
  0:(length(grep("DIS.n.at.age", names(dt3)))-1),"NO_DISCARD",sep="_")

names(dt3)[grep("n.at.age", names(dt3)) ]<-paste("AGE",
  0:(length(grep("n.at.age", names(dt3)))-1),"NO_LANDED",sep="_")

names(dt3)[grep("DmeanW.at.age", names(dt3)) ]<-paste("AGE",
  0:(length(grep("DmeanW.at.age", names(dt3)))-1),"MEAN_WEIGHT_DISCARD",
  ",
  sep="_")

names(dt3)[grep("DmeanL.at.age", names(dt3)) ]<-paste("AGE",
  0:(length(grep("DmeanL.at.age", names(dt3)))-1),"MEAN_LENGTH_DISCARD",
  ",
  sep="_")

names(dt3)[grep("meanW.at.age", names(dt3)) ]<-paste("AGE",
  0:(length(grep("meanW.at.age", names(dt3)))-1),"MEAN_WEIGHT_LANDED",
  sep="_")

names(dt3)[grep("meanL.at.age", names(dt3)) ]<-paste("AGE",
  0:(length(grep("meanL.at.age", names(dt3)))-1),"MEAN_LENGTH_LANDED",
  sep="_")

names(dt3)[grep("age.1", names(dt3))]<-paste("AGE",
  0:(length(grep("age.1", names(dt3)))-1),sep="_")

dt3 <- dt3 %>% separate(time, c("Year","Quarter")," - ",remove=T)

#####\\

# FINAL CATCH TAB -----

catch1<- catch1 %>% mutate_at(vars( c(ID:SPECIES) ),
  funs( ifelse( is.na(.), -1, .) ) )

```

```

dt3<- dt3 %>% mutate_at(vars( c(Year:MESH_SIZE_RANGE) ),
  funs( ifelse( is.na(.), -1, .) ) )

catch1$YEAR=as.character(catch1$YEAR)
catch1$FISHERY=as.character(catch1$FISHERY)
catch.tab <- left_join(catch1,dt3,by=c( "QUARTER" = "Quarter" , "YEAR"="Year",
  "AREA"="space", "GEAR"="gear" , "VESSEL_LENGTH" = "VL" ,
  "MESH_SIZE_RANGE","FISHERY" ))

# FISHERY to DG MARE Med&BS codification
catch.tab$FISHERY <- fishery$SDEF_codification[match(catch.tab$FISHERY ,
  fishery$DGMARE_Med_BS_codification)]

# species to FAO three alpha code and set ID (COUNTRY, AREA, GEAR, VESSEL_LENGTH
,
# MESH_SIZE_RANGE,QUARTER, SPECIES)
catch.tab <- catch.tab %>%
  mutate(SPECIES=sel_spe$SPE[match(SPECIES,sel_spe$SPECIES)],
  ID = paste(COUNTRY, AREA, GEAR,FISHERY, VESSEL_LENGTH, MESH_SIZE_RANGE,
  YEAR, QUARTER, SPECIES, sep = "_"))

catch.tab$YEAR=as.numeric(catch.tab$YEAR)
catch.temp2<-bind_rows(catch.temp2,catch.tab)

}

```

Output

```

# col after 12: set -1 or NA to 0
catch.temp2[, -c(1:11)][is.na(catch.temp2[, -c(1:11)])] <- 0

catch.temp2<-setDT(catch.temp2)
for (jj in c(12:171)) set(catch.temp2, i = which(catch.temp2[[jj]]==-1),
  j = jj, v = 0)

catch.temp2<-setDF(catch.temp2)

write.table(catch.temp2, file = "CATCH.csv",row.names=FALSE,sep=";",
  na="-1")

head(catch.temp2)

##              ID  COUNTRY YEAR QUARTER
## 1 COUNTRY1_GSA99_OTB_NONE_VL2440_-1_9999_1_HKE COUNTRY1 9999      1
## 2 COUNTRY1_GSA99_GNS_NONE_VL0612_-1_9999_3_HKE COUNTRY1 9999      3
## 3 COUNTRY1_GSA99_OTB_NONE_VL2440_-1_9999_3_HKE COUNTRY1 9999      3
## 4 COUNTRY1_GSA99_GNS_NONE_VL0006_-1_9999_4_HKE COUNTRY1 9999      4
## 5 COUNTRY1_GSA99_GNS_NONE_VL0612_-1_9999_4_HKE COUNTRY1 9999      4
## 6 COUNTRY1_GSA99_GTR_NONE_VL0612_-1_9999_4_HKE COUNTRY1 9999      4
##  VESSEL_LENGTH GEAR MESH_SIZE_RANGE FISHERY  AREA SPECON SPECIES

```

## 1	VL2440	OTB	-1	NONE	GSA99	-1	HKE
## 2	VL0612	GNS	-1	NONE	GSA99	-1	HKE
## 3	VL2440	OTB	-1	NONE	GSA99	-1	HKE
## 4	VL0006	GNS	-1	NONE	GSA99	-1	HKE
## 5	VL0612	GNS	-1	NONE	GSA99	-1	HKE
## 6	VL0612	GTR	-1	NONE	GSA99	-1	HKE
##	LANDINGS	DISCARDS	NO_SAMPLES_LANDINGS	NO_LENGTH_MEASUREMENTS_LANDINGS			
## 1	241.996000	5.7386667	6	990			
## 2	140.230571	0.0000000	2	20			
## 3	196.598000	8.3981013	11	1247			
## 4	10.177558	1.3884000	4	18			
## 5	87.679445	2.6935003	34	657			
## 6	2.382747	0.5006894	30	54			
##	NO_AGE_MEASUREMENTS_LANDINGS	NO_SAMPLES_DISCARDS					
## 1		0	6				
## 2		0	0				
## 3		0	11				
## 4		0	1				
## 5		0	9				
## 6		0	6				
##	NO_LENGTH_MEASUREMENTS_DISCARDS	NO_AGE_MEASUREMENTS_DISCARDS					
## 1		158	0				
## 2		0	0				
## 3		566	0				
## 4		3	0				
## 5		55	0				
## 6		8	0				
##	NO_SAMPLES_CATCH	NO_LENGTH_MEASUREMENTS_CATCH	NO_AGE_MEASUREMENTS_CATCH				
## 1	12	1148	0				
## 2	2	20	0				
## 3	22	1813	0				
## 4	5	21	0				
## 5	43	712	0				
## 6	36	62	0				
##	MIN_AGE	MAX_AGE	AGE_0	AGE_0_NO_LANDED	AGE_0_MEAN_WEIGHT_LANDED		
## 1	0	5	0	843.1723427	0.06205808		
## 2	0	3	0	12.0142710	0.07634600		
## 3	0	5	0	259.8784047	0.02516775		
## 4	0	4	0	4.1846456	0.10845659		
## 5	0	4	0	80.1428431	0.09104605		
## 6	0	3	0	0.6587253	0.10588056		
##	AGE_0_MEAN_LENGTH_LANDED	AGE_0_NO_DISCARD	AGE_0_MEAN_WEIGHT_DISCARD				
## 1	20.65193	67.733333	0.06217778				
## 2	22.00000	0.000000	0.02465982				
## 3	14.17963	787.972292	0.02465982				
## 4	24.17651	3.653684	0.03872180				
## 5	22.30149	12.243183	0.03872180				
## 6	24.08543	0.000000	0.03872180				
##	AGE_0_MEAN_LENGTH_DISCARD	AGE_1	AGE_1_NO_LANDED	AGE_1_MEAN_WEIGHT_LANDED			
## 1	20.66667	1	674.988360	0.1611525			
## 2	14.02727	1	154.873488	0.2537027			

## 3	14.02727	1	460.251467	0.1669250
## 4	15.74194	1	15.502594	0.1976597
## 5	15.74194	1	217.383326	0.1889448
## 6	15.74194	1	2.608883	0.2201388
##	AGE_1_MEAN_LENGTH_LANDED	AGE_1_NO_DISCARD	AGE_1_MEAN_WEIGHT_DISCARD	
## 1	28.12088	0.000000	0.1650500	
## 2	31.59069	0.000000	0.1828965	
## 3	27.29795	0.000000	0.1828965	
## 4	29.55358	6.576632	0.1914079	
## 5	28.97150	11.859835	0.1914079	
## 6	30.46054	1.112643	0.1914079	
##	AGE_1_MEAN_LENGTH_DISCARD	AGE_2	AGE_2_NO_LANDED	AGE_2_MEAN_WEIGHT_LANDED
## 1	28.35000	2	130.922358	0.3723365
## 2	28.11538	2	150.288995	0.3443452
## 3	28.11538	2	132.648716	0.3820961
## 4	29.01471	2	4.381310	0.3785201
## 5	29.01471	2	47.558245	0.3911659
## 6	29.01471	2	1.707482	0.4578178
##	AGE_2_MEAN_LENGTH_LANDED	AGE_2_NO_DISCARD	AGE_2_MEAN_WEIGHT_DISCARD	
## 1	36.66217	0.0000000	0.3918095	
## 2	35.13610	0.0000000	0.3789413	
## 3	36.34099	0.0000000	0.3789413	
## 4	36.51664	0.7307368	0.4064423	
## 5	36.88426	0.6708593	0.4064423	
## 6	38.93109	0.1788176	0.4064423	
##	AGE_2_MEAN_LENGTH_DISCARD	AGE_3	AGE_3_NO_LANDED	AGE_3_MEAN_WEIGHT_LANDED
## 1	37.19048	3	7.1052892	0.5340000
## 2	36.27500	3	48.0570838	0.7162028
## 3	36.27500	3	35.2480868	0.7249392
## 4	37.36667	3	2.5787732	1.0823878
## 5	37.36667	3	12.6647254	0.7583605
## 6	37.36667	3	0.8024339	0.8602206
##	AGE_3_MEAN_LENGTH_LANDED	AGE_3_NO_DISCARD	AGE_3_MEAN_WEIGHT_DISCARD	
## 1	42.32727	0.000000	0.6838182	
## 2	45.92308	0.000000	0.7623153	
## 3	44.93343	0.000000	0.7623153	
## 4	49.59184	0.000000	0.8041833	
## 5	44.95651	0.000000	0.8041833	
## 6	46.96346	0.960317	0.8041833	
##	AGE_3_MEAN_LENGTH_DISCARD	AGE_4	AGE_4_NO_LANDED	AGE_4_MEAN_WEIGHT_LANDED
## 1	45.72727	4	0.0000000	0.000000
## 2	45.57895	4	0.0000000	0.000000
## 3	45.57895	4	13.4507270	1.530725
## 4	45.70833	4	0.8595911	1.050696
## 5	45.70833	4	0.8674070	1.260696
## 6	45.70833	4	0.0000000	1.228488
##	AGE_4_MEAN_LENGTH_LANDED	AGE_4_NO_DISCARD	AGE_4_MEAN_WEIGHT_DISCARD	
## 1	0.00000	0	1.131250	
## 2	0.00000	0	1.620727	
## 3	56.02192	0	1.620727	
## 4	51.12000	0	1.460415	

## 5	52.69554	0	1.460415
## 6	53.06459	0	1.460415
##	AGE_4_MEAN_LENGTH_DISCARD	AGE_5	AGE_5_NO_LANDED
## 1	53.50000	5	4.567686
## 2	57.09091	5	0.000000
## 3	57.09091	5	2.400298
## 4	55.46154	5	0.000000
## 5	55.46154	5	0.000000
## 6	55.46154	5	0.000000
##	AGE_5_MEAN_LENGTH_LANDED	AGE_5_NO_DISCARD	AGE_5_MEAN_WEIGHT_DISCARD
## 1	48.00000	0	1.37450
## 2	0.00000	0	2.04452
## 3	61.15508	0	2.04452
## 4	54.00000	0	2.18600
## 5	0.00000	0	2.18600
## 6	0.00000	0	2.18600
##	AGE_5_MEAN_LENGTH_DISCARD	AGE_6	AGE_6_NO_LANDED
## 1	55.66667	6	0
## 2	63.10000	6	0
## 3	63.10000	6	0
## 4	62.62500	6	0
## 5	62.62500	6	0
## 6	62.62500	6	0
##	AGE_6_MEAN_LENGTH_LANDED	AGE_6_NO_DISCARD	AGE_6_MEAN_WEIGHT_DISCARD
## 1	0	0	1.517000
## 2	0	0	2.804000
## 3	0	0	2.804000
## 4	0	0	3.177754
## 5	0	0	3.177754
## 6	0	0	3.177754
##	AGE_6_MEAN_LENGTH_DISCARD	AGE_7	AGE_7_NO_LANDED
## 1	59.00000	7	0
## 2	69.50000	7	0
## 3	69.50000	7	0
## 4	71.07692	7	0
## 5	71.07692	7	0
## 6	71.07692	7	0
##	AGE_7_MEAN_LENGTH_LANDED	AGE_7_NO_DISCARD	AGE_7_MEAN_WEIGHT_DISCARD
## 1	0	0	0.0000
## 2	0	0	0.0000
## 3	0	0	0.0000
## 4	0	0	3.4315
## 5	0	0	3.4315
## 6	0	0	3.4315
##	AGE_7_MEAN_LENGTH_DISCARD	AGE_8	AGE_8_NO_LANDED
## 1	0.00000	8	0
## 2	0.00000	8	0
## 3	0.00000	8	0
## 4	74.07143	8	0
## 5	74.07143	8	0
## 6	74.07143	8	0

##	AGE_8_MEAN_LENGTH_LANDED	AGE_8_NO_DISCARD	AGE_8_MEAN_WEIGHT_DISCARD	
## 1	0	0	0.0000	
## 2	0	0	3.6825	
## 3	0	0	3.6825	
## 4	0	0	5.2066	
## 5	0	0	5.2066	
## 6	0	0	5.2066	
##	AGE_8_MEAN_LENGTH_DISCARD	AGE_9	AGE_9_NO_LANDED	AGE_9_MEAN_WEIGHT_LANDED
## 1	0.0	9	0	0
## 2	76.5	9	0	0
## 3	76.5	9	0	0
## 4	82.8	9	0	0
## 5	82.8	9	0	0
## 6	82.8	9	0	0
##	AGE_9_MEAN_LENGTH_LANDED	AGE_9_NO_DISCARD	AGE_9_MEAN_WEIGHT_DISCARD	
## 1	0	0	0.000	
## 2	0	0	0.000	
## 3	0	0	0.000	
## 4	0	0	6.195	
## 5	0	0	6.195	
## 6	0	0	6.195	
##	AGE_9_MEAN_LENGTH_DISCARD	AGE_10	AGE_10_NO_LANDED	
## 1	0	10	0	
## 2	0	10	0	
## 3	0	10	0	
## 4	87	10	0	
## 5	87	10	0	
## 6	87	10	0	
##	AGE_10_MEAN_WEIGHT_LANDED	AGE_10_MEAN_LENGTH_LANDED	AGE_10_NO_DISCARD	
## 1	0	0	0	
## 2	0	0	0	
## 3	0	0	0	
## 4	0	0	0	
## 5	0	0	0	
## 6	0	0	0	
##	AGE_10_MEAN_WEIGHT_DISCARD	AGE_10_MEAN_LENGTH_DISCARD	AGE_11	
## 1	0	0	11	
## 2	0	0	11	
## 3	0	0	11	
## 4	0	0	11	
## 5	0	0	11	
## 6	0	0	11	
##	AGE_11_NO_LANDED	AGE_11_MEAN_WEIGHT_LANDED	AGE_11_MEAN_LENGTH_LANDED	
## 1	0	0	0	
## 2	0	0	0	
## 3	0	0	0	
## 4	0	0	0	
## 5	0	0	0	
## 6	0	0	0	
##	AGE_11_NO_DISCARD	AGE_11_MEAN_WEIGHT_DISCARD	AGE_11_MEAN_LENGTH_DISCARD	
## 1	0	0	0	

##	2	0	0	0
##	3	0	0	0
##	4	0	0	0
##	5	0	0	0
##	6	0	0	0
##	AGE_12	AGE_12_NO_LANDED	AGE_12_MEAN_WEIGHT_LANDED	
##	1	12	0	0
##	2	12	0	0
##	3	12	0	0
##	4	12	0	0
##	5	12	0	0
##	6	12	0	0
##	AGE_12_MEAN_LENGTH_LANDED	AGE_12_NO_DISCARD	AGE_12_MEAN_WEIGHT_DISCARD	
##	1	0	0	0
##	2	0	0	0
##	3	0	0	0
##	4	0	0	0
##	5	0	0	0
##	6	0	0	0
##	AGE_12_MEAN_LENGTH_DISCARD	AGE_13	AGE_13_NO_LANDED	
##	1	0	13	0
##	2	0	13	0
##	3	0	13	0
##	4	0	13	0
##	5	0	13	0
##	6	0	13	0
##	AGE_13_MEAN_WEIGHT_LANDED	AGE_13_MEAN_LENGTH_LANDED	AGE_13_NO_DISCARD	
##	1	0	0	0
##	2	0	0	0
##	3	0	0	0
##	4	0	0	0
##	5	0	0	0
##	6	0	0	0
##	AGE_13_MEAN_WEIGHT_DISCARD	AGE_13_MEAN_LENGTH_DISCARD	AGE_14	
##	1	0	0	14
##	2	0	0	14
##	3	0	0	14
##	4	0	0	14
##	5	0	0	14
##	6	0	0	14
##	AGE_14_NO_LANDED	AGE_14_MEAN_WEIGHT_LANDED	AGE_14_MEAN_LENGTH_LANDED	
##	1	0	0	0
##	2	0	0	0
##	3	0	0	0
##	4	0	0	0
##	5	0	0	0
##	6	0	0	0
##	AGE_14_NO_DISCARD	AGE_14_MEAN_WEIGHT_DISCARD	AGE_14_MEAN_LENGTH_DISCARD	
##	1	0	0	0
##	2	0	0	0
##	3	0	0	0

## 4		0		0		0
## 5		0		0		0
## 6		0		0		0
##	AGE_15	AGE_15_NO_LANDED	AGE_15_MEAN_WEIGHT_LANDED			
## 1	15		0		0	
## 2	15		0		0	
## 3	15		0		0	
## 4	15		0		0	
## 5	15		0		0	
## 6	15		0		0	
##	AGE_15_MEAN_LENGTH_LANDED	AGE_15_NO_DISCARD	AGE_15_MEAN_WEIGHT_DISCARD			
## 1		0		0		0
## 2		0		0		0
## 3		0		0		0
## 4		0		0		0
## 5		0		0		0
## 6		0		0		0
##	AGE_15_MEAN_LENGTH_DISCARD	AGE_16	AGE_16_NO_LANDED			
## 1		0	16		0	
## 2		0	16		0	
## 3		0	16		0	
## 4		0	16		0	
## 5		0	16		0	
## 6		0	16		0	
##	AGE_16_MEAN_WEIGHT_LANDED	AGE_16_MEAN_LENGTH_LANDED	AGE_16_NO_DISCARD			
## 1		0		0		0
## 2		0		0		0
## 3		0		0		0
## 4		0		0		0
## 5		0		0		0
## 6		0		0		0
##	AGE_16_MEAN_WEIGHT_DISCARD	AGE_16_MEAN_LENGTH_DISCARD	AGE_17			
## 1		0		0	17	
## 2		0		0	17	
## 3		0		0	17	
## 4		0		0	17	
## 5		0		0	17	
## 6		0		0	17	
##	AGE_17_NO_LANDED	AGE_17_MEAN_WEIGHT_LANDED	AGE_17_MEAN_LENGTH_LANDED			
## 1		0		0		0
## 2		0		0		0
## 3		0		0		0
## 4		0		0		0
## 5		0		0		0
## 6		0		0		0
##	AGE_17_NO_DISCARD	AGE_17_MEAN_WEIGHT_DISCARD	AGE_17_MEAN_LENGTH_DISCARD			
## 1		0		0		0
## 2		0		0		0
## 3		0		0		0
## 4		0		0		0
## 5		0		0		0

## 6	0	0	0
##	AGE_18	AGE_18_NO_LANDED	AGE_18_MEAN_WEIGHT_LANDED
## 1	18	0	0
## 2	18	0	0
## 3	18	0	0
## 4	18	0	0
## 5	18	0	0
## 6	18	0	0
##	AGE_18_MEAN_LENGTH_LANDED	AGE_18_NO_DISCARD	AGE_18_MEAN_WEIGHT_DISCARD
## 1	0	0	0
## 2	0	0	0
## 3	0	0	0
## 4	0	0	0
## 5	0	0	0
## 6	0	0	0
##	AGE_18_MEAN_LENGTH_DISCARD	AGE_19	AGE_19_NO_LANDED
## 1	0	19	0
## 2	0	19	0
## 3	0	19	0
## 4	0	19	0
## 5	0	19	0
## 6	0	19	0
##	AGE_19_MEAN_WEIGHT_LANDED	AGE_19_MEAN_LENGTH_LANDED	AGE_19_NO_DISCARD
## 1	0	0	0
## 2	0	0	0
## 3	0	0	0
## 4	0	0	0
## 5	0	0	0
## 6	0	0	0
##	AGE_19_MEAN_WEIGHT_DISCARD	AGE_19_MEAN_LENGTH_DISCARD	AGE_20_PLUS
## 1	0	0	0
## 2	0	0	0
## 3	0	0	0
## 4	0	0	0
## 5	0	0	0
## 6	0	0	0
##	AGE_20_PLUS_NO_LANDED	AGE_20_PLUS_MEAN_WEIGHT_LANDED	
## 1	0	0	
## 2	0	0	
## 3	0	0	
## 4	0	0	
## 5	0	0	
## 6	0	0	
##	AGE_20_PLUS_MEAN_LENGTH_LANDED	AGE_20_PLUS_NO_DISCARD	
## 1	0	0	
## 2	0	0	
## 3	0	0	
## 4	0	0	
## 5	0	0	
## 6	0	0	
##	AGE_20_PLUS_MEAN_WEIGHT_DISCARD	AGE_20_PLUS_MEAN_LENGTH_DISCARD	

```
## 1          0          0
## 2          0          0
## 3          0          0
## 4          0          0
## 5          0          0
## 6          0          0
```

Example of use of the script ML

This script implements the Calculation of maturity at length, required for the DG MARE Med&BS Data Call, using as input file the SDEF format (CS table) and COST as the raising procedure

Settings

```
path <- paste("C:\\Users\\Bitetto Isabella\\OneDrive - Coispa Tecnologia & Ricerca
S.C.A.R.L\\MARE22\\STREAM\\FINAL REVISION OF DELIVERABLES\\SDEF_to_DG_MARE_MedBS",
sep="")
```

```
setwd(path)
```

```
knitr::opts_knit$set(root.dir = path)
```

Input Data

csData of the SDEF format

```
load("fri_cs_test.Rdata")
head( fri_cs@ca)
```

```
##      sampType landCtry vsIflgCtry year   proj trpCode staNum quarter month
## 58          V COUNTRY1  COUNTRY1 9999 PROJECT      3    999         1      2
## 105         V COUNTRY1  COUNTRY1 9999 PROJECT      7    999         1      2
## 106         V COUNTRY1  COUNTRY1 9999 PROJECT      2    999         1      2
## 107         V COUNTRY1  COUNTRY1 9999 PROJECT      6    999         1      2
## 122         V COUNTRY1  COUNTRY1 9999 PROJECT     10    999         1      2
## 127         V COUNTRY1  COUNTRY1 9999 PROJECT      5    999         3      7
##
##              spp sex catchCat landCat commCatScl commCat stock  area
## 58 Mullus barbatus M      LAN      HUC      <NA>    <NA> <NA> GSA99
## 105 Mullus barbatus M      LAN      HUC      <NA>    <NA> <NA> GSA99
## 106 Mullus barbatus M      LAN      HUC      <NA>    <NA> <NA> GSA99
## 107 Mullus barbatus F      LAN      HUC      <NA>    <NA> <NA> GSA99
## 122 Mullus barbatus M      LAN      HUC      <NA>    <NA> <NA> GSA99
## 127 Mullus barbatus F      LAN      HUC      <NA>    <NA> <NA> GSA99
##
##      rect subRect lenCls age fishId lenCode ageMeth plusGrp otoWt otoSide
## 58 <NA> THR-LIM    110  0      1      mm      <NA>    <NA>    NA    <NA>
## 105 <NA> THR-LIM     90  0      1      mm      <NA>    <NA>    NA    <NA>
## 106 <NA> THR-LIM    110  0      1      mm      <NA>    <NA>    NA    <NA>
## 107 <NA> THR-LIM    110  0      1      mm      <NA>    <NA>    NA    <NA>
## 122 <NA> THR-LIM    110  0      1      mm      <NA>    <NA>    NA    <NA>
## 127 <NA> THR-LIM    180  3      1      mm      <NA>    <NA>    NA    <NA>
##
##      indWt matMeth matScale matStage
```

```
## 58 18.00 <NA> MEDITS 2
## 105 11.10 <NA> MEDITS 1
## 106 13.80 <NA> MEDITS 1
## 107 17.30 <NA> MEDITS 2
## 122 17.76 <NA> MEDITS 2
## 127 68.12 <NA> MEDITS 4
```

```
# extract COUNTRY
```

```
COUNTRY<-unique(fri_cs@ca$landCtry)
```

stratification: spatial only

```
fri_strD <- strIni(spaceStrata="area")
```

DG MARE Med&BS ML Table template

```
m1.temp2<- read.table("xxx_ML.csv",sep="," ,header=T)
names(m1.temp2)
```

```
## [1] "COUNTRY"      "AREA"          "START_YEAR"    "END_YEAR"      "SPECIES"
## [6] "SEX"          "LENGTHCLASS"  "UNIT"          "SAMPLE_SIZE"   "PRM"
## [11] "METHOD_USED"
```

Auxiliary table: species_BIO_ML.csv

```
# auxiliary Table
```

```
sel_spe<-read.table( "species_BIO_ML.csv", header=TRUE, sep=";",
                     row.names=NULL)
```

```
head(sel_spe)
```

```
##           SPECIES SPE   GSA SEX START_YEAR END_YEAR IMMATURE
## 1 Merluccius merluccius HKE GSA99  M      9999      9999    0_1_2a
## 2 Mullus barbatus MUT GSA99 M_F      9999      9999      1
##           MAT_METHOD methodDesc nboot
## 1 macroscopically analytical      .
## 2 macroscopically bootstrap      10
```

```
sel_spe<- sel_spe %>%
  mutate(SEX = strsplit(as.character(SEX), "_")) %>%
  unnest(SEX)
```

```
imm=sel_spe%>%separate(IMMATURE,c("a", "b","c","d"),
                       extra = "drop", fill = "right",sep="_")
```

Checking data

Check if there are stocks with no age data and impute "-1", otherwise the bpEstim() method for maturity at length gives error :

```
# get col w. all=NA by spp
```

```
aa<- fri_cs@ca %>% group_by(spp) %>% summarize_all(~all(is.na(.)))
```

```
# aa$spp[aa$age==FALSE] # spp with age data
```

```
# impute "-1" if a stock has no age data
```

```
fri_cs@ca$age[fri_cs@ca$spp %in% aa$spp[aa$age==TRUE]]<- -1
```

Set immature stages based on Input Auxiliary Table

```
# get col w. all=NA by spp
```

```
aa<- fri_cs@ca %>% group_by(spp) %>% summarize_all(~all(is.na(.)))
```

```
# if no matScale is spcified, impute "-1"
```

```
fri_cs@ca$matScale[fri_cs@ca$spp %in% aa$spp[aa$matScale==TRUE]]<- "-1"
```

Data analysis

Analysis by stock

```
for (i in 1:dim(sel_spe)[1]) {
```

```
  STK<- sel_spe$SPECIES[i]
```

```
  fri_cs1<- subset(fri_cs, year%in% seq(sel_spe$START_YEAR[i],  
    sel_spe$END_YEAR[i],by=1),table="ca",link=T)
```

```
    # set immature stages as "I"
```

```
if (na.omit(unique(fri_cs1@ca$matScale[fri_cs1@ca$spp==STK &  
  fri_cs1@ca$area==sel_spe$GSA[i]]))!="1-7") {  
  fri_cs1@ca$matStage[fri_cs1@ca$spp==STK & fri_cs1@ca$area==sel_spe$GSA[i] &  
    fri_cs1@ca$matStage %in% imm[i,6:9][!is.na(imm[i, 6:9])]]="I"
```

```
} else{ #if "1-7" matScale is used, immature (I)==1  
  fri_cs1@ca$matStage[fri_cs1@ca$spp==STK & fri_cs1@ca$area==sel_spe$GSA[i] &  
    fri_cs1@ca$matStage ==1]="I"
```

```
}
```

```
# sample size
```

```
if (sel_spe$SEX[i]=="C"){
```

```
  fri_csv <- csDataVal(fri_cs1)
```

```
  # get sample size
```

```
  nml<- data.frame(fri_cs1@ca) %>% filter(!is.na(matStage))%>%  
    dplyr::group_by(area,spp,lenCls)%>% summarize(SAMPLE_SIZE=n())
```

```
} else {
```

```

    fri_cs1=subset(fri_cs1,sex==sel_spe$SEX[i],table="ca",link=T)
    fri_csv <- csDataVal(fri_cs1)

    # get sample size
    nml<- data.frame(fri_cs1@ca) %>% filter(!is.na(matStage))%>%
      dplyr::group_by(area,spp,lenCls,sex)%>% summarize(SAMPLE_SIZE=n())
  }

  fri_csv1<- subSetSpp(fri_csv, STK)
  fri_csv1<- subset(fri_csv1, area%in% sel_spe$GSA[i],table="ca")

  fri_csc1 <- csDataCons(fri_csv1, fri_strD)

  fri_csc1@ca$matStage=as.character(fri_csc1@ca$matStage)

  MS_MAT_An<-dbeObject(desc="Maturity-at-length",species=STK,
    param="maturity",strataDesc=fri_strD,
    methodDesc=sel_spe$methodDesc[i],nboot=sel_spe$nboot[i])

  if (sel_spe$methodDesc[i]=="bootstrap"){
    MS_MAT_An<-bpBoot(MS_MAT_An,fri_csc1,adjust=F,immature.scale="I")
  } else { # analytical
    MS_MAT_An<-bpEstim(MS_MAT_An,fri_csc1,adjust=F,immature.scale="I")
  }

  pp=dbPlot(MS_MAT_An,elmt="lenStruc$estim",ylab="Maturity ratio",
    main=paste("Maturity ratio-at-length estimates
      for",STK,sel_spe$SEX[i],sel_spe$GSA[i],sel_spe$START_YEAR[i]
    ,
      sel_spe$END_YEAR[i],sep=" "))

  # pdf(file="p.pdf")
  # print(pp)

  dfML <-
    data.frame(
      COUNTRY =COUNTRY ,
      AREA = sel_spe$GSA[i],
      START_YEAR = sel_spe$START_YEAR[i] ,
      END_YEAR = sel_spe$END_YEAR[i] ,
      SPECIES = STK ,
      SEX = sel_spe$SEX[i],
      UNIT = unique(fri_csc1@ca$lenCode) ,
      LENGTHCLASS = as.numeric(as.character(MS_MAT_An@lenStruc[["estim"]][["length"
    ]]) ),

```



```

    PRM = MS_MAT_An@lenStruc[["estim"]][["value"]] ,
    METHOD_USED = sel_spe$MAT_METHOD[i]
  )

# get sample size

if (sel_spe$SEX[i]=="C"){

dfML <- dfML %>%
  left_join(nml, by = c("AREA" = 'area', 'SPECIES' = 'spp',
                        "LENGTHCLASS"="lenCls")) %>%
  mutate(SPECIES = sel_spe$SPE[i])
# FAO three alpha code species code

} else{

dfML <- dfML %>%
  left_join(nml, by = c("AREA" = 'area', 'SPECIES' = 'spp',
                        "SEX"="sex", "LENGTHCLASS"="lenCls")) %>%
  mutate(SPECIES = sel_spe$SPE[i])
# FAO three alpha code species code

}

ml.temp2<- bind_rows(ml.temp2,na.omit(dfML))

}

## csData subset by species European hake Merluccius merluccius
## New data set consists of 21 trip records
## 74 length records
## and 368 age or maturity records
## csData subset by species Red mullet Mullus barbatus
## New data set consists of 19 trip records
## 28 length records
## and 329 age or maturity records
## csData subset by species Red mullet Mullus barbatus
## New data set consists of 20 trip records
## 33 length records
## and 759 age or maturity records

```

Output

```

# export DG MARE Med&BS ML table
write.table(format(ml.temp2,digits=3, scientific=F),
            file=paste("ML.csv",sep=""),dec=".",sep="," ,
            col.names=TRUE,row.names=FALSE,na="-1")

head(ml.temp2)

```

```
##      COUNTRY AREA START_YEAR END_YEAR SPECIES SEX LENGTHCLASS UNIT
## 1 COUNTRY1 GSA99      9999      9999      HKE  M          90    mm
## 2 COUNTRY1 GSA99      9999      9999      HKE  M         100    mm
## 3 COUNTRY1 GSA99      9999      9999      HKE  M         110    mm
## 4 COUNTRY1 GSA99      9999      9999      HKE  M         120    mm
## 5 COUNTRY1 GSA99      9999      9999      HKE  M         130    mm
## 6 COUNTRY1 GSA99      9999      9999      HKE  M         140    mm
##  SAMPLE_SIZE PRM      METHOD_USED
## 1           4  0 macroscopically
## 2           5  0 macroscopically
## 3          11  0 macroscopically
## 4          21  0 macroscopically
## 5          12  0 macroscopically
## 6           4  0 macroscopically
```

Example of use of the script MA

This script implements the Calculation of maturity at age, required for the JRC MED &BS Data Call, using as input file the SDEF format (CS table) and COST as the raising procedure

Settings

```
path <- paste("C:\\Users\\Bitetto Isabella\\OneDrive - Coispa Tecnologia & Ricerca
S.C.A.R.L\\MARE22\\STREAM\\FINAL REVISION OF DELIVERABLES\\SDEF_to_DG_MARE_MedBS",
sep="")
setwd(path)
```

Input Data

csData of the SDEF format and estimate sample size

```
load("fri_cs_test.Rdata")
head( fri_cs@ca)

##      sampType landCtry vsFlgCtry year      proj trpCode staNum quarter month
## 58          V COUNTRY1  COUNTRY1 9999 PROJECT          3    999          1     2
## 105         V COUNTRY1  COUNTRY1 9999 PROJECT          7    999          1     2
## 106         V COUNTRY1  COUNTRY1 9999 PROJECT          2    999          1     2
## 107         V COUNTRY1  COUNTRY1 9999 PROJECT          6    999          1     2
## 122         V COUNTRY1  COUNTRY1 9999 PROJECT         10    999          1     2
## 127         V COUNTRY1  COUNTRY1 9999 PROJECT          5    999          3     7
##
##      spp sex catchCat landCat commCatScl commCat stock area
## 58 Mullus barbatus M      LAN      HUC      <NA>      <NA>      <NA> GSA99
## 105 Mullus barbatus M      LAN      HUC      <NA>      <NA>      <NA> GSA99
## 106 Mullus barbatus M      LAN      HUC      <NA>      <NA>      <NA> GSA99
## 107 Mullus barbatus F      LAN      HUC      <NA>      <NA>      <NA> GSA99
## 122 Mullus barbatus M      LAN      HUC      <NA>      <NA>      <NA> GSA99
## 127 Mullus barbatus F      LAN      HUC      <NA>      <NA>      <NA> GSA99
##      rect subRect lenCls age fishId lenCode ageMeth plusGrp otoWt otoSide
## 58 <NA> THR-LIM      110  0      1      mm      <NA>      <NA>      NA      <NA>
## 105 <NA> THR-LIM       90  0      1      mm      <NA>      <NA>      NA      <NA>
```

```
## 106 <NA> THR-LIM      110    0    1      mm      <NA>      <NA>      NA      <NA>
## 107 <NA> THR-LIM      110    0    1      mm      <NA>      <NA>      NA      <NA>
## 122 <NA> THR-LIM      110    0    1      mm      <NA>      <NA>      NA      <NA>
## 127 <NA> THR-LIM      180    3    1      mm      <NA>      <NA>      NA      <NA>
##      indWt matMeth matScale matStage
## 58  18.00      <NA>    MEDITS         2
## 105  11.10      <NA>    MEDITS         1
## 106  13.80      <NA>    MEDITS         1
## 107  17.30      <NA>    MEDITS         2
## 122  17.76      <NA>    MEDITS         2
## 127  68.12      <NA>    MEDITS         4
```

```
# extract COUNTRY
COUNTRY<-unique(fri_cs@ca$landCtry)
```

stratification: spatial only

```
fri_strD <- strIni(spaceStrata="area")
```

DG MARE Med&BS MA Table template

```
ma.temp2<- read.table("xxx_MA.csv",sep=";",header=T)
names(ma.temp2)

## [1] "COUNTRY"      "AREA"          "START_YEAR"    "END_YEAR"      "SPECIES"
## [6] "SEX"          "AGECLASS"      "SAMPLE_SIZE"   "PRM"            "METHOD_USED"
```

Auxiliary table: species_BIO_MA.csv

```
# species file : selected species with FAO three alpha code

sel_spe<-read.table( "species_BIO_MA.csv", header=TRUE, sep=";",row.names=NULL)

head(sel_spe)

##              SPECIES SPE    GSA SEX START_YEAR END_YEAR IMMATURE
## 1 Merluccius merluccius HKE GSA99  M      9999      9999    0_1_2a
## 2 Mullus barbatus MUT GSA99 M_F      9999      9999         1
##      MAT_METHOD methodDesc nboot adjust
## 1 macroscopically analytical .    TRUE
## 2 macroscopically bootstrap   10  FALSE

sel_spe<- sel_spe %>%
  mutate(SEX = strsplit(as.character(SEX), "_")) %>%
  unnest(SEX)

imm=sel_spe%>%separate(IMMATURE,c("a", "b","c","d"), extra = "drop",
  fill = "right",sep="_")
```

Checking data

Set immature stages based on Auxiliary Table

```
# get col w. all=NA by spp
aa<- fri_cs@ca %>% group_by(spp) %>% summarize_all(~all(is.na(.)))

# if no matScale is spcified, impute "-1"
fri_cs@ca$matScale[fri_cs@ca$spp %in% aa$spp[aa$matScale==TRUE]]<- "-1"
```

Data analysis- raising

Analysis by stock

```
for (i in 1:dim(sel_spe)[1]) {

  STK<- sel_spe$SPECIES[i]

  fri_cs1<- subset(fri_cs, year%in% seq(sel_spe$START_YEAR[i],
    sel_spe$END_YEAR[i],by=1),table="ca",link=T)

  # set immature stages as "I"
  if (na.omit(unique(fri_cs1@ca$matScale[fri_cs1@ca$spp==STK &
    fri_cs1@ca$area==sel_spe$GSA[i]]))!="1-7") {
    fri_cs1@ca$matStage[fri_cs1@ca$spp==STK & fri_cs1@ca$area==sel_spe$GSA[i] &
      fri_cs1@ca$matStage %in% imm[i,6:9][!is.na(imm[i, 6:9])]]="I"
  } else{ #if "1-7" matScale is used, immature (I)==1
    fri_cs1@ca$matStage[fri_cs1@ca$spp==STK & fri_cs1@ca$area==sel_spe$GSA[i] &
      fri_cs1@ca$matStage ==1]="I"
  }

  # sample size
  if (sel_spe$SEX[i]=="C"){

    fri_csv <- csDataVal(fri_cs1)

    # get sample size

    nml<- data.frame(fri_cs1@ca) %>% filter(!is.na(matStage)& !is.na(age))%>%
      dplyr::group_by(area,spp,age)%>% summarize(SAMPLE_SIZE=n())

  } else {

    fri_cs1=subset(fri_cs1,sex==sel_spe$SEX[i],table="ca",link=T)
    fri_csv <- csDataVal(fri_cs1)
```

```

# get sample size
nm1<- data.frame(fri_csv1@ca) %>% filter(!is.na(matStage)& !is.na(age))%>%
  dplyr::group_by(area,spp,age,sex)%>% summarize(SAMPLE_SIZE=n())
}

fri_csv1<- subSetSpp(fri_csv, STK)
fri_csv1<- subset(fri_csv1, area%in% sel_spe$GSA[i],table="ca")

fri_csc1 <- csDataCons(fri_csv1, fri_strD)

fri_csc1@ca$matStage=as.character(fri_csc1@ca$matStage)

MS_MAT_An<-dbeObject(desc=" Maturity-at-age",species=STK,param="maturity",
  strataDesc=fri_strD,methodDesc=sel_spe$methodDesc[i],
  nboot=sel_spe$nboot[i])

if (sel_spe$methodDesc[i]=="bootstrap"){
MS_MAT_An<-bpBoot(MS_MAT_An,fri_csc1,adjust=sel_spe$adjust[i],immature.scale="I")
} else { # analytical
MS_MAT_An<-bpEstim(MS_MAT_An,fri_csc1,adjust=sel_spe$adjust[i],immature.scale="I")
}

dfMA <-
  data.frame(

    COUNTRY =COUNTRY ,
    AREA = unique(MS_MAT_An@ageStruc[["estim"]][["space"]]) ,
    START_YEAR = sel_spe$START_YEAR[i] ,
    END_YEAR = sel_spe$END_YEAR[i] ,
    SPECIES = STK ,
    SEX = sel_spe$SEX[i],

    AGECLASS = as.numeric(as.character(MS_MAT_An@ageStruc[["estim"]][["age"]]) ),

    PRM = MS_MAT_An@ageStruc[["estim"]][["value"]] ,
    METHOD_USED = sel_spe$MAT_METHOD[i]

  )

if (sel_spe$SEX[i]=="C"){

dfMA <- dfMA %>%

```

```

    left_join(nml, by = c("AREA" = 'area', 'SPECIES' = 'spp',
                          "AGECLASS"="age")) %>% mutate(SPECIES = sel_spe$SPE[
i])
# FAO three alpha code
} else{

  dfMA <- dfMA %>%
    left_join(nml, by = c("AREA" = 'area', 'SPECIES' = 'spp',
                          "SEX"="sex", "AGECLASS"="age")) %>%
    mutate(SPECIES = sel_spe$SPE[i])
# FAO three alpha code
}

ma.temp2<- bind_rows(ma.temp2,na.omit(dfMA))

}

## csData subset by species European hake Merluccius merluccius
## New data set consists of 21 trip records
## 74 length records
## and 368 age or maturity records
## csData subset by species Red mullet Mullus barbatus
## New data set consists of 19 trip records
## 28 length records
## and 329 age or maturity records
## csData subset by species Red mullet Mullus barbatus
## New data set consists of 20 trip records
## 33 length records
## and 759 age or maturity records

```

Output

```

# export DG MARE Med&BS MA table
write.table(format(ma.temp2,digits=3, scientific=F), file=paste("MA.csv",sep=""),
            dec=".",sep=";",col.names=TRUE, row.names=FALSE,na="-1")

```

```
head(ma.temp2)
```

```

##   COUNTRY AREA START_YEAR END_YEAR SPECIES SEX AGECLASS SAMPLE_SIZE
## 1 COUNTRY1 GSA99      9999      9999   HKE  M         0          67
## 2 COUNTRY1 GSA99      9999      9999   HKE  M         1         107
## 3 COUNTRY1 GSA99      9999      9999   HKE  M         2          51
## 4 COUNTRY1 GSA99      9999      9999   HKE  M         3          17
## 5 COUNTRY1 GSA99      9999      9999   MUT  M         0          14
## 6 COUNTRY1 GSA99      9999      9999   MUT  M         1         133
##   PRM      METHOD_USED
## 1 0.4072577 macroscopically
## 2 0.9801666 macroscopically
## 3 1.0000000 macroscopically
## 4 1.0000000 macroscopically

```

```
## 5 1.0000000 macroscopically
## 6 0.9789111 macroscopically
```

Example of use of the script SRL

This script implements the Calculation of sex ratio at length, required for the DG MARE Med&BS Data Call, using as input file the SDEF format (CS table) and COST as the raising procedure

Settings

```
path <- paste("C:\\Users\\Bitetto Isabella\\OneDrive - Coispa Tecnologia & Ricerca
S.C.A.R.L\\MARE22\\STREAM\\FINAL REVISION OF DELIVERABLES\\SDEF_to_DG_MARE_MedBS",
sep="")
```

```
setwd(path)
```

Input Data

csData of the SDEF format

```
load("fri_cs_test.Rdata")
head( fri_cs@ca)
```

```
##      sampType landCtry vs1FlgCtry year   proj trpCode staNum quarter month
## 58          V COUNTRY1  COUNTRY1 9999 PROJECT      3    999        1      2
## 105         V COUNTRY1  COUNTRY1 9999 PROJECT      7    999        1      2
## 106         V COUNTRY1  COUNTRY1 9999 PROJECT      2    999        1      2
## 107         V COUNTRY1  COUNTRY1 9999 PROJECT      6    999        1      2
## 122         V COUNTRY1  COUNTRY1 9999 PROJECT     10    999        1      2
## 127         V COUNTRY1  COUNTRY1 9999 PROJECT      5    999        3      7
##
##              spp sex catchCat landCat commCatScl commCat stock  area
## 58  Mullus barbatus  M      LAN      HUC      <NA>    <NA>    <NA> GSA99
## 105 Mullus barbatus  M      LAN      HUC      <NA>    <NA>    <NA> GSA99
## 106 Mullus barbatus  M      LAN      HUC      <NA>    <NA>    <NA> GSA99
## 107 Mullus barbatus  F      LAN      HUC      <NA>    <NA>    <NA> GSA99
## 122 Mullus barbatus  M      LAN      HUC      <NA>    <NA>    <NA> GSA99
## 127 Mullus barbatus  F      LAN      HUC      <NA>    <NA>    <NA> GSA99
##
##      rect subRect lenCls age fishId lenCode ageMeth plusGrp otoWt otoSide
## 58  <NA> THR-LIM    110  0      1      mm      <NA>    <NA>    NA    <NA>
## 105  <NA> THR-LIM     90  0      1      mm      <NA>    <NA>    NA    <NA>
## 106  <NA> THR-LIM    110  0      1      mm      <NA>    <NA>    NA    <NA>
## 107  <NA> THR-LIM    110  0      1      mm      <NA>    <NA>    NA    <NA>
## 122  <NA> THR-LIM    110  0      1      mm      <NA>    <NA>    NA    <NA>
## 127  <NA> THR-LIM    180  3      1      mm      <NA>    <NA>    NA    <NA>
##
##      indWt matMeth matScale matStage
## 58  18.00    <NA>    MEDITS      2
## 105  11.10    <NA>    MEDITS      1
## 106  13.80    <NA>    MEDITS      1
## 107  17.30    <NA>    MEDITS      2
## 122  17.76    <NA>    MEDITS      2
## 127  68.12    <NA>    MEDITS      4
```

```
# extract COUNTRY
COUNTRY<-unique(fri_cs@ca$landCtry)
```

stratification: spatial only

```
fri_strD <- strIni(spaceStrata="area")
```

DG MARE Med&BS SRL Table template

```
srl.temp2<- read.table("xxx_SRL.csv",sep=",",header=T)
```

```
names(srl.temp2)
```

```
## [1] "COUNTRY"      "AREA"          "START_YEAR"    "END_YEAR"      "SPECIES"
## [6] "LENGTHCLASS" "UNIT"          "SEX_RATIO"     "COMMENTS"
```

Auxilliary table: species_BIO_SRL.csv

```
# species file : selected species with FAO three alpha code
```

```
sel_spe<-read.table( "species_BIO_SRL.csv", header=TRUE, sep=";",row.names=NULL)
```

```
head(sel_spe)
```

```
##           SPECIES SPE   GSA START_YEAR END_YEAR methodDesc nboot
## 1 Merluccius merluccius HKE GSA99      9999      9999 analytical   .
## 2 Mullus barbatus MUT GSA99      9999      9999 bootstrap    10
## COMMENTS
## 1      NA
## 2      NA
```

Checking data

Check if there are stocks with no age data and impute "-1", otherwise the bpEstim() method for sex ratio at length gives error :

```
# get col w. all=NA by spp
aa<- fri_cs@ca %>% group_by(spp) %>% summarize_all(~all(is.na(.)))
```

```
aa$spp[aa$age==FALSE] # spp with age data
```

```
## [1] "Merluccius merluccius" "Mullus barbatus"
```

```
# impute "-1" if a stock has no age data
```

```
fri_cs@ca$age[fri_cs@ca$spp %in% aa$spp[aa$age==TRUE]]<- -1
```

Data analysis- raising

Analysis by stock

```
for (i in 1:dim(sel_spe)[1]) {
```

```
  STK<- sel_spe$SPECIES[i]
```



```

fri_cs1<- subset(fri_cs, year%in% seq(sel_spe$START_YEAR[i],
                                     sel_spe$END_YEAR[i],by=1),table="ca",link=T)

# Validate- consolidate
fri_csv <- csDataVal(fri_cs1)
fri_csc<- csDataCons(fri_csv, fri_strD)

fri_csc1<- subset(fri_csc, space==sel_spe$GSA[i],table="ca")

MS_SEX_An<-dbeObject(desc="SEX-ratio",species=STK,param="sex",
                     strataDesc=fri_strD,methodDesc=sel_spe$methodDesc[i],
                     nboot=sel_spe$nboot[i])

if (sel_spe$methodDesc[i]=="bootstrap"){
MS_SEX_An<-bpBoot(MS_SEX_An,fri_csc1,adjust=F)
} else {
MS_SEX_An<-bpEstim(MS_SEX_An,fri_csc1,adjust=F)
}

dfSRL <-
  data.frame(

    COUNTRY = COUNTRY ,
    AREA = MS_SEX_An@lenStruc[["estim"]][["space"]] ,
    START_YEAR = sel_spe$START_YEAR[i] ,
    END_YEAR = sel_spe$END_YEAR[i] ,
    SPECIES = STK ,

    UNIT = unique(fri_cs@ca$lenCode[fri_cs@ca$spp==STK]) ,
    LENGTHCLASS = as.character(MS_SEX_An@lenStruc[["estim"]][["length"]]) ,

    SEX_RATIO = MS_SEX_An@lenStruc[["estim"]][["value"]],
    COMMENTS = sel_spe$COMMENTS[i]
  )

dfSRL<-dfSRL %>% mutate(SPECIES=sel_spe$SPE[match(STK,sel_spe$SPECIES)])

dfSRL <- dfSRL[complete.cases(dfSRL[, -which(names(dfSRL) %in% c("COMMENTS"))]),]

sr1.temp2=bind_rows(sr1.temp2,dfSRL)

}

```

Output

```

# export DG MARE Med&BS SRL table
write.table(format(sr1.temp2,digits=3, scientific=F),

```

```
file="SRL.csv",dec=".",sep=","col.names=TRUE,row.names=FALSE,na="-1")
```

```
head(sr1.temp2)
```

```
##      COUNTRY  AREA START_YEAR END_YEAR SPECIES LENGTHCLASS UNIT  SEX_RATIO
## 1 COUNTRY1  GSA99      9999      9999      HKE           80   mm 1.00000000
## 2 COUNTRY1  GSA99      9999      9999      HKE           90   mm 0.00000000
## 3 COUNTRY1  GSA99      9999      9999      HKE          100   mm 0.00000000
## 4 COUNTRY1  GSA99      9999      9999      HKE          110   mm 0.21428571
## 5 COUNTRY1  GSA99      9999      9999      HKE          120   mm 0.08695652
## 6 COUNTRY1  GSA99      9999      9999      HKE          130   mm 0.40000000
##      COMMENTS
## 1          NA
## 2          NA
## 3          NA
## 4          NA
## 5          NA
## 6          NA
```

Example of use of the script SRA

This script implements the Calculation of sex ratio at age, required for the DG MARE Med&BS Data Call, using as input file the SDEF format (CS table) and COST as the raising procedure

Settings

```
path <- paste("C:\\Users\\Bitetto Isabella\\OneDrive - Coispa Tecnologia & Ricerca
S.C.A.R.L\\MARE22\\STREAM\\FINAL REVISION OF DELIVERABLES\\SDEF_to_DG_MARE_MedBS",
sep="")
```

```
setwd(path)
```

Input Data

csData of the SDEF format

```
load("fri_cs_test.Rdata")
```

```
head( fri_cs@ca)
```

```
##      sampType landCtry vs1FlgCtry year   proj trpCode staNum quarter month
## 58          V COUNTRY1  COUNTRY1 9999 PROJECT      3    999        1      2
## 105         V COUNTRY1  COUNTRY1 9999 PROJECT      7    999        1      2
## 106         V COUNTRY1  COUNTRY1 9999 PROJECT      2    999        1      2
## 107         V COUNTRY1  COUNTRY1 9999 PROJECT      6    999        1      2
## 122         V COUNTRY1  COUNTRY1 9999 PROJECT     10    999        1      2
## 127         V COUNTRY1  COUNTRY1 9999 PROJECT      5    999        3      7
##
##      spp sex catchCat landCat commCatSc1 commCat stock  area
## 58 Mullus barbatus  M      LAN      HUC      <NA>      <NA>      <NA> GSA99
## 105 Mullus barbatus  M      LAN      HUC      <NA>      <NA>      <NA> GSA99
## 106 Mullus barbatus  M      LAN      HUC      <NA>      <NA>      <NA> GSA99
```

```
## 107 Mullus barbatus F LAN HUC <NA> <NA> <NA> GSA99
## 122 Mullus barbatus M LAN HUC <NA> <NA> <NA> GSA99
## 127 Mullus barbatus F LAN HUC <NA> <NA> <NA> GSA99
## rect subRect lenCls age fishId lenCode ageMeth plusGrp otoWt otoSide
## 58 <NA> THR-LIM 110 0 1 mm <NA> <NA> NA <NA>
## 105 <NA> THR-LIM 90 0 1 mm <NA> <NA> NA <NA>
## 106 <NA> THR-LIM 110 0 1 mm <NA> <NA> NA <NA>
## 107 <NA> THR-LIM 110 0 1 mm <NA> <NA> NA <NA>
## 122 <NA> THR-LIM 110 0 1 mm <NA> <NA> NA <NA>
## 127 <NA> THR-LIM 180 3 1 mm <NA> <NA> NA <NA>
## indWt matMeth matScale matStage
## 58 18.00 <NA> MEDITS 2
## 105 11.10 <NA> MEDITS 1
## 106 13.80 <NA> MEDITS 1
## 107 17.30 <NA> MEDITS 2
## 122 17.76 <NA> MEDITS 2
## 127 68.12 <NA> MEDITS 4
```

extract COUNTRY

```
COUNTRY<-unique(fri_cs@ca$landCtry)
```

stratification: spatial only

```
fri_strD <- strIni(spaceStrata="area")
```

DG MARE Med&BS SRA Table template

```
sra.temp2<- read.table("xxx_SRA.csv",sep=";",header=T)
```

```
names(sra.temp2)
```

```
## [1] "COUNTRY" "AREA" "START_YEAR" "END_YEAR" "SPECIES"
## [6] "AGECLASS" "SEX_RATIO" "COMMENTS"
```

Auxilliary table: species_BIO_SRA.csv

species file : selected species with FAO three alpha code

```
sel_spe<-read.table("species_BIO_SRA.csv", header=TRUE, sep=";",row.names=NULL)
```

```
head(sel_spe)
```

```
## SPECIES SPE GSA START_YEAR END_YEAR COMMENTS methodDesc
## 1 Merluccius merluccius HKE GSA99 9999 9999 NA analytical
## 2 Mullus barbatus MUT GSA99 9999 9999 NA bootstrap
## nboot adjust
## 1 . FALSE
## 2 10 FALSE
```

Data analysis- raising

Analysis by stock

```

for (i in 1:dim(sel_spe)[1]) {

  STK<- sel_spe$SPECIES[i]

  fri_cs1<- subset(fri_cs, year%in% seq(sel_spe$START_YEAR[i],
                                         sel_spe$END_YEAR[i],by=1),table="ca",lin
k=T)

# Validate- consolidate
fri_csv <- csDataVal(fri_cs1)
fri_csc<- csDataCons(fri_csv, fri_strD)

  fri_csc1<- subset(fri_csc, space==sel_spe$GSA[i],table="ca")

MS_SEX_An<-dbeObject(desc="SEX-ratio",species=STK,param="sex",
                      strataDesc=fri_strD,methodDesc=sel_spe$methodDesc[i],
                      nboot=sel_spe$nboot[i])

if (sel_spe$methodDesc[i]=="bootstrap"){
MS_SEX_An<-bpBoot(MS_SEX_An,fri_csc1,adjust=sel_spe$adjust[i])
} else {
MS_SEX_An<-bpEstim(MS_SEX_An,fri_csc1,adjust=sel_spe$adjust[i])
}

dfSRA <-
  data.frame(

    COUNTRY = COUNTRY ,
    AREA = MS_SEX_An@ageStruc[["estim"]][["space"]] ,
    START_YEAR = sel_spe$START_YEAR[i] ,
    END_YEAR = sel_spe$END_YEAR[i] ,
    SPECIES = STK ,
    AGECLASS = MS_SEX_An@ageStruc[["estim"]][["age"]] ,
    SEX_RATIO = MS_SEX_An@ageStruc[["estim"]][["value"]],
    COMMENTS = sel_spe$COMMENTS[i]
  )

dfSRA<-dfSRA %>% mutate(SPECIES=sel_spe$SPE[match(STK,sel_spe$SPECIES)])

dfSRA <- dfSRA[complete.cases(dfSRA[, -which(names(dfSRA) %in% c("COMMENTS"))]),]

dfSRA$AGECLASS <- as.numeric(as.character(dfSRA$AGECLASS))

sra.temp2=bind_rows(sra.temp2, dfSRA)

}

```

Output

```
# export DG MARE Med&BS SRA table
write.table(format(sra.temp2,digits=3, scientific=F),
            file="SRA.csv",dec=".",sep="," ,col.names=TRUE,
            row.names=FALSE,na="-1")
```

```
head(sra.temp2)
```

##	COUNTRY	AREA	START_YEAR	END_YEAR	SPECIES	AGECLASS	SEX_RATIO	COMMENTS
## 1	COUNTRY1	GSA99	9999	9999	HKE	0	0.4765625	NA
## 2	COUNTRY1	GSA99	9999	9999	HKE	1	0.6310345	NA
## 3	COUNTRY1	GSA99	9999	9999	HKE	2	0.6871166	NA
## 4	COUNTRY1	GSA99	9999	9999	HKE	3	0.8172043	NA
## 5	COUNTRY1	GSA99	9999	9999	HKE	4	0.9493671	NA
## 6	COUNTRY1	GSA99	9999	9999	HKE	5	0.9454545	NA

Example of use of the script ALK

This script implements the Calculation of the age length key, required for the DG MARE Med&BS Data Call, using as input file the SDEF format (CS table) and COST as the raising procedure. Additional methods exist for filling in the potential gaps for length classes.

Settings

```
path <- paste("C:\\Users\\Bitetto Isabella\\OneDrive - Coispa Tecnologia & Ricerca  
S.C.A.R.L\\MARE22\\STREAM\\FINAL REVISION OF DELIVERABLES\\SDEF_to_DG_MARE_MedBS",  
sep="")
```

```
setwd(path)
```

Input Data

csData of the SDEF format

```
load("fri_cs_test.Rdata")
head(fri_cs@ca)
```

##	sampType	landCtry	vs1FlgCtry	year	proj	trpCode	staNum	quarter	month
## 58	V	COUNTRY1	COUNTRY1	9999	PROJECT	3	999	1	2
## 105	V	COUNTRY1	COUNTRY1	9999	PROJECT	7	999	1	2
## 106	V	COUNTRY1	COUNTRY1	9999	PROJECT	2	999	1	2
## 107	V	COUNTRY1	COUNTRY1	9999	PROJECT	6	999	1	2
## 122	V	COUNTRY1	COUNTRY1	9999	PROJECT	10	999	1	2
## 127	V	COUNTRY1	COUNTRY1	9999	PROJECT	5	999	3	7
##	spp	sex	catchCat	landCat	commCatSc1	commCat	stock	area	
## 58	Mullus barbatus	M	LAN	HUC	<NA>	<NA>	<NA>	GSA99	
## 105	Mullus barbatus	M	LAN	HUC	<NA>	<NA>	<NA>	GSA99	
## 106	Mullus barbatus	M	LAN	HUC	<NA>	<NA>	<NA>	GSA99	
## 107	Mullus barbatus	F	LAN	HUC	<NA>	<NA>	<NA>	GSA99	
## 122	Mullus barbatus	M	LAN	HUC	<NA>	<NA>	<NA>	GSA99	

```
## 127 Mullus barbatus F LAN HUC <NA> <NA> <NA> GSA99
## rect subRect lenCls age fishId lenCode ageMeth plusGrp otoWt otoSide
## 58 <NA> THR-LIM 110 0 1 mm <NA> <NA> NA <NA>
## 105 <NA> THR-LIM 90 0 1 mm <NA> <NA> NA <NA>
## 106 <NA> THR-LIM 110 0 1 mm <NA> <NA> NA <NA>
## 107 <NA> THR-LIM 110 0 1 mm <NA> <NA> NA <NA>
## 122 <NA> THR-LIM 110 0 1 mm <NA> <NA> NA <NA>
## 127 <NA> THR-LIM 180 3 1 mm <NA> <NA> NA <NA>
## indWt matMeth matScale matStage
## 58 18.00 <NA> MEDITS 2
## 105 11.10 <NA> MEDITS 1
## 106 13.80 <NA> MEDITS 1
## 107 17.30 <NA> MEDITS 2
## 122 17.76 <NA> MEDITS 2
## 127 68.12 <NA> MEDITS 4
```

```
# extract COUNTRY
```

```
COUNTRY<-unique(fri_cs@ca$landCtry)
```

stratification: spatial only

```
fri_strD <- strIni(spaceStrata="area")
```

DG MARE Med&BS ALK Table template

```
alk.temp2<- read.table("xxx_ALK.csv",sep="," ,header=T)
head(alk.temp2)
```

```
## [1] COUNTRY
## [2] AREA
## [3] START_YEAR
## [4] END_YEAR
## [5] SPECON
## [6] SPECIES
## [7] SEX
## [8] APPLY_TO_CATCHES_FILE
## [9] TOTAL_NUMBER_OF_HARD_STRUCTURE_READ_BY_AGE
## [10] CV
## [11] UNIT
## [12] AGE
## [13] LENGTHCLASS0
## [14] LENGTHCLASS1
## [15] LENGTHCLASS2
## [16] LENGTHCLASS3
## [17] LENGTHCLASS4
## [18] LENGTHCLASS5
## [19] LENGTHCLASS6
## [20] LENGTHCLASS7
## [21] LENGTHCLASS8
## [22] LENGTHCLASS9
## [23] LENGTHCLASS10
## [24] LENGTHCLASS11
```

##	[25]	LENGTHCLASS12
##	[26]	LENGTHCLASS13
##	[27]	LENGTHCLASS14
##	[28]	LENGTHCLASS15
##	[29]	LENGTHCLASS16
##	[30]	LENGTHCLASS17
##	[31]	LENGTHCLASS18
##	[32]	LENGTHCLASS19
##	[33]	LENGTHCLASS20
##	[34]	LENGTHCLASS21
##	[35]	LENGTHCLASS22
##	[36]	LENGTHCLASS23
##	[37]	LENGTHCLASS24
##	[38]	LENGTHCLASS25
##	[39]	LENGTHCLASS26
##	[40]	LENGTHCLASS27
##	[41]	LENGTHCLASS28
##	[42]	LENGTHCLASS29
##	[43]	LENGTHCLASS30
##	[44]	LENGTHCLASS31
##	[45]	LENGTHCLASS32
##	[46]	LENGTHCLASS33
##	[47]	LENGTHCLASS34
##	[48]	LENGTHCLASS35
##	[49]	LENGTHCLASS36
##	[50]	LENGTHCLASS37
##	[51]	LENGTHCLASS38
##	[52]	LENGTHCLASS39
##	[53]	LENGTHCLASS40
##	[54]	LENGTHCLASS41
##	[55]	LENGTHCLASS42
##	[56]	LENGTHCLASS43
##	[57]	LENGTHCLASS44
##	[58]	LENGTHCLASS45
##	[59]	LENGTHCLASS46
##	[60]	LENGTHCLASS47
##	[61]	LENGTHCLASS48
##	[62]	LENGTHCLASS49
##	[63]	LENGTHCLASS50
##	[64]	LENGTHCLASS51
##	[65]	LENGTHCLASS52
##	[66]	LENGTHCLASS53
##	[67]	LENGTHCLASS54
##	[68]	LENGTHCLASS55
##	[69]	LENGTHCLASS56
##	[70]	LENGTHCLASS57
##	[71]	LENGTHCLASS58
##	[72]	LENGTHCLASS59
##	[73]	LENGTHCLASS60
##	[74]	LENGTHCLASS61
##	[75]	LENGTHCLASS62

```
## [76] LENGTHCLASS63
## [77] LENGTHCLASS64
## [78] LENGTHCLASS65
## [79] LENGTHCLASS66
## [80] LENGTHCLASS67
## [81] LENGTHCLASS68
## [82] LENGTHCLASS69
## [83] LENGTHCLASS70
## [84] LENGTHCLASS71
## [85] LENGTHCLASS72
## [86] LENGTHCLASS73
## [87] LENGTHCLASS74
## [88] LENGTHCLASS75
## [89] LENGTHCLASS76
## [90] LENGTHCLASS77
## [91] LENGTHCLASS78
## [92] LENGTHCLASS79
## [93] LENGTHCLASS80
## [94] LENGTHCLASS81
## [95] LENGTHCLASS82
## [96] LENGTHCLASS83
## [97] LENGTHCLASS84
## [98] LENGTHCLASS85
## [99] LENGTHCLASS86
## [100] LENGTHCLASS87
## [101] LENGTHCLASS88
## [102] LENGTHCLASS89
## [103] LENGTHCLASS90
## [104] LENGTHCLASS91
## [105] LENGTHCLASS92
## [106] LENGTHCLASS93
## [107] LENGTHCLASS94
## [108] LENGTHCLASS95
## [109] LENGTHCLASS96
## [110] LENGTHCLASS97
## [111] LENGTHCLASS98
## [112] LENGTHCLASS99
## [113] LENGTHCLASS100_PLUS
## [114] COMMENTS
## <0 rows> (or 0-length row.names)
```

Auxiliary table: species_BIO_ALK.csv

species file : selected species with FAO three alpha code

```
sel_spe<-read.table( "species_BIO_ALK.csv", header=TRUE, sep=";",row.names=NULL)
head(sel_spe)
```

```
##           SPECIES SPE   GSA SEX START_YEAR END_YEAR LC_RANGE
## 1 Merluccius merluccius HKE GSA99  C      9999      9999      10
## 2      Mullus barbatus MUT  GSA99 F_M      9999      9999      10
```



```
## APPLY_TO_CATCHES_FILE typeALK valueALK methodDesc_LAN.len.age adjust
## 1 Y stepIncr 10 analytical TRUE
## 2 N stepIncr 10 analytical FALSE
## SPECON COMMENTS
## 1 NA NA
## 2 NA NA

sel_spe<- sel_spe %>%
  mutate(SEX = strsplit(as.character(SEX), "_")) %>%
  unnest(SEX)
```

Data analysis- raising

Analysis by stock

```
for (i in 1:dim(sel_spe)[1]) {

  STK<- sel_spe$SPECIES[i]

  fri_cs1<- subset(fri_cs, year%in% seq(sel_spe$START_YEAR[i],
    sel_spe$END_YEAR[i],by=1),table="ca",link=T)

  # estimate sample size (number of otoliths per stock, sex and age)
  if (sel_spe$SEX[i]=="C"){

    fri_csv <- csDataVal(fri_cs1)

    nml<- data.frame(fri_cs1@ca) %>% filter(!is.na(age))%>%
      dplyr::group_by(area,spp,age)%>%
      summarize(TOTAL_NUMBER_OF_HARD_STRUCTURE_READ_BY_AGE=n())

  } else { # ALK for selected sex

    fri_cs1=subset(fri_cs1,sex==sel_spe$SEX[i],table="ca",link=T)
    fri_csv <- csDataVal(fri_cs1)

    # get sample size: number of otoliths
    nml<- data.frame(fri_cs1@ca) %>% filter(!is.na(age))%>%
      dplyr::group_by(area,spp,age,sex)%>%
      summarize(TOTAL_NUMBER_OF_HARD_STRUCTURE_READ_BY_AGE=n())

  }

  fri_csv1<- subSetSpp(fri_csv, STK)
  fri_csv1<- subset(fri_csv1, area%in% sel_spe$GSA[i],table="ca",link=T)
  fri_csc1 <- csDataCons(fri_csv1, fri_strD)

  ## ### CV from individual length-at-age
  LEstim_An <-
```

```

dbeObject(
desc = "Length at age",
species = STK,
catchCat = "LAN",
param = "length",
strataDesc = fri_strD, # ,
methodDesc = "analytical" #sel_spe$methodDesc_LAN.Len.age[i]
)

if(sel_spe$methodDesc_LAN.Len.age[i]=="analytical") {
LEstim_An <- bpEstim(LEstim_An, fri_csc1, adjust = sel_spe$adjust[i])
} else{
LEstim_An <- bpBoot(LEstim_An, fri_csc1, adjust = sel_spe$adjust[i])
}

## ALK

res1 <- alkLgthRec(fri_csc1,type=sel_spe$typeALK[i],value=sel_spe$valueALK[i],
                  update=F, preview=F,postview = F)

if (sel_spe$typeALK[i]=="fillALKmult"){
  fri_csc2 <- fillALKmult(fri_csc1,STK,p=10,trace=T)
  res1 <- alkLgthRec(fri_csc2,update=F, preview=F,postview = F,
                    value=sel_spe$valueALK[i])
}

dfALK <-
  data.frame(
    COUNTRY =COUNTRY ,
    AREA =sel_spe$GSA[i],
    START_YEAR = unique(sel_spe$START_YEAR[i]) ,
    END_YEAR = unique( sel_spe$END_YEAR[i]) ,
    SPECIES = STK ,
    SEX = sel_spe$SEX[i],
    UNIT = unique(fri_cs1@ca$lenCode[fri_cs1@ca$spp==STK]) ,
    SPECON= -1,
    APPLY_TO_CATCHES_FILE= unique( sel_spe$APPLY_TO_CATCHES_FILE[i]),
    CV=NA,
    AGE = as.numeric(colnames(res1$alk)),
    COMMENTS= sel_spe$COMMENTS[i],
    LENGTHCLASS100_PLUS=0
  )

# get sample size

if (sel_spe$SEX[i]=="C"){

```

```

dfALK <- dfALK %>%
  left_join(nml, by = c("AREA" = 'area', 'SPECIES' = 'spp', "AGE"="age")) %>%
  mutate(SPECIES = sel_spe$SPE[i])
# FAO Three alpha code
} else{

dfALK <- dfALK %>%
  left_join(nml, by = c("AREA" = 'area', 'SPECIES' = 'spp', "AGE"="age",
                        "SEX"="sex")) %>%
  mutate(SPECIES = sel_spe$SPE[i])
# FAO Three alpha code

}

aa=data.frame(res1$alk)
names(aa)=colnames(res1$alk)

aa=aa%>% mutate(LC=rownames(res1$alk))

## fix LC
UNIT <- as.character( unique(fri_csc1@ca$lenCode[fri_csc1@ca$spp==STK]) )

if (UNIT %in% c("mm", "MM") & sel_spe$LC_RANGE[i]==10) {
aa$LC<-as.numeric(aa$LC)/10
UNIT1<- "cm"
}

if (UNIT %in% c("mm", "MM") & sel_spe$LC_RANGE[i]==1) {
aa$LC<-as.numeric(aa$LC)
UNIT1<- "mm"
}

if (UNIT %in% c("mm", "MM") & sel_spe$LC_RANGE[i]==5) {
aa$LC<-as.numeric(aa$LC)/10
UNIT1<- "cm"
}

if (UNIT %in% c("cm", "CM") ) {
aa$LC<-as.numeric(aa$LC)
UNIT1<- "cm"
}

aa$LC<- plyr::round_any( aa$LC,1,floor)
###

dfALK$UNIT<-UNIT1

aa1=aa %>% gather(age, n.at.len, -LC)

```

```

aa1 <- as.data.table(aa1)

seq_1 <- seq(0, max(aa$LC, na.rm = T), by = 1) #
dt1<- aa1[, list(LC = seq_1), by = age]

dt2<- left_join(dt1,aa1)

dt3 <- data.table::dcast(dt2,as.formula(paste(paste(names(dt2)[! names(dt2)
      %in% c("LC","n.at.len")], collapse='+'), "LC", sep="~")),
      value.var = "n.at.len")

dt3$age=as.numeric(dt3$age)

dt3<- dt3 %>% mutate_at(vars( -(age) ),
      funs( if_else( is.na(.), 0, .) ) )

dfALK<-left_join(dfALK,dt3,by=c("AGE"="age"))

## CV
# LEstim_An @ageNum$ cv

LEstim_An@ageNum[["cv"]]$age=as.numeric(LEstim_An@ageNum[["cv"]]$age)

dfALK<- dfALK%>% left_join(LEstim_An@ageNum[["cv"]]%>% select(age,value),
      by=c( "AGE"="age"))%>% mutate(CV=value)%>% select(-c(value))

# take care of number of Length classes (max is 100 acc. to DG MARE Med&BS template)
zz<-dim(dfALK[-c(1:14)])[2]
names(dfALK)[-c(1:14)]<- paste("LENGTHCLASS",seq(0,zz-1,1),sep="")

if(zz>=100){
  dfALK$LENGTHCLASS100_PLUS<- rowSums(dfALK[, -c(1:114)])
}

dfALK<-dfALK%>% select(one_of(as.vector(names(alk.temp2))))

alk.temp2<- bind_rows(alk.temp2,(dfALK))

alk.temp2[, -c(1:13,114)][is.na(alk.temp2[, -c(1:13,114)])] <- 0

# export updated CsDataCons

if (sel_spe$APPLY_TO_CATCHES_FILE[i] == "Y") {

```

```

if (sel_spe$typeALK[i] == "fillALKmult") {
  fri_csc2 <- fillALKmult(fri_csc1, STK, p = 10, trace = T)

  save(fri_csc2,
       file = paste("upd", STK, sel_spe$SEX[i], sel_spe$GSA[i], ".Rdata", sep
=" _"))

} else{
  res1 <-
    alkLgthRec(
      fri_csc1,
      type = sel_spe$typeALK[i],
      value = sel_spe$valueALK[i],
      update = T,
      preview = F,
      postview = F
    )

  save(res1,
       file = paste("upd", STK, sel_spe$SEX[i], sel_spe$GSA[i], ".Rdata", sep
=
       "_"))
}

}

}

## csData subset by species European hake Merluccius merluccius
## New data set consists of 197 trip records
## 6573 length records
## and 23481 age or maturity records
## csData subset by species Red mullet Mullus barbatus
## New data set consists of 20 trip records
## 33 length records
## and 759 age or maturity records
## csData subset by species Red mullet Mullus barbatus
## New data set consists of 19 trip records
## 28 length records
## and 329 age or maturity records

```

Output

```

# export DG MARE Med&BS ALK table
write.table(format(alk.temp2,digits=3, scientific=F),
  file="ALK.csv",dec=".",sep="," ,col.names=TRUE,
  row.names=FALSE,na="-1")

head(alk.temp2)

##   COUNTRY  AREA START_YEAR END_YEAR SPECON SPECIES SEX
## 1 COUNTRY1 GSA99      9999      9999     -1     HKE   C

```

## 2	COUNTRY1	GSA99		9999	9999	-1	HKE	C	
## 3	COUNTRY1	GSA99		9999	9999	-1	HKE	C	
## 4	COUNTRY1	GSA99		9999	9999	-1	HKE	C	
## 5	COUNTRY1	GSA99		9999	9999	-1	HKE	C	
## 6	COUNTRY1	GSA99		9999	9999	-1	HKE	C	
##	APPLY_TO_CATCHES_FILE TOTAL_NUMBER_OF_HARD_STRUCTURE_READ_BY_AGE								
## 1			Y						337
## 2			Y						293
## 3			Y						165
## 4			Y						94
## 5			Y						79
## 6			Y						55
##		CV	UNIT	AGE	LENGTHCLASS0	LENGTHCLASS1	LENGTHCLASS2	LENGTHCLASS3	
## 1		NA	cm	0	0	0	39		40
## 2	0.018500742		cm	1	0	0	0		0
## 3	0.007583819		cm	2	0	0	0		0
## 4	0.007301511		cm	3	0	0	0		0
## 5	0.008633827		cm	4	0	0	0		0
## 6	0.011211429		cm	5	0	0	0		0
##	LENGTHCLASS4	LENGTHCLASS5	LENGTHCLASS6	LENGTHCLASS7	LENGTHCLASS8				
## 1	22	23		21	17		18		
## 2	0	0		0	0		0		
## 3	0	0		0	0		0		
## 4	0	0		0	0		0		
## 5	0	0		0	0		0		
## 6	0	0		0	0		0		
##	LENGTHCLASS9	LENGTHCLASS10	LENGTHCLASS11	LENGTHCLASS12	LENGTHCLASS13				
## 1	12	6		10	0		12		
## 2	0	1		1	0		0		
## 3	0	0		0	0		0		
## 4	0	0		0	0		0		
## 5	0	0		0	0		0		
## 6	0	0		0	0		0		
##	LENGTHCLASS14	LENGTHCLASS15	LENGTHCLASS16	LENGTHCLASS17	LENGTHCLASS18				
## 1	8	12		10	7		6		
## 2	9	7		4	9		23		
## 3	0	0		0	0		0		
## 4	0	0		0	0		0		
## 5	0	0		0	0		0		
## 6	0	0		0	0		0		
##	LENGTHCLASS19	LENGTHCLASS20	LENGTHCLASS21	LENGTHCLASS22	LENGTHCLASS23				
## 1	0	1		0	0		0		
## 2	25	38		27	21		0		
## 3	0	0		0	1		0		
## 4	0	0		0	0		0		
## 5	0	0		0	0		0		
## 6	0	0		0	0		0		
##	LENGTHCLASS24	LENGTHCLASS25	LENGTHCLASS26	LENGTHCLASS27	LENGTHCLASS28				
## 1	0	0		0	0		0		
## 2	25	28		28	20		12		
## 3	7	3		9	13		10		

## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS29	LENGTHCLASS30	LENGTHCLASS31	LENGTHCLASS32	LENGTHCLASS33
## 1	0	0	0	0	0
## 2	5	6	1	3	0
## 3	17	19	17	25	13
## 4	0	0	1	3	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS34	LENGTHCLASS35	LENGTHCLASS36	LENGTHCLASS37	LENGTHCLASS38
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	13	5	2	6
## 4	0	4	7	17	5
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS39	LENGTHCLASS40	LENGTHCLASS41	LENGTHCLASS42	LENGTHCLASS43
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	3	1	0	1	0
## 4	9	7	10	6	6
## 5	1	1	5	2	2
## 6	0	0	0	1	2
##	LENGTHCLASS44	LENGTHCLASS45	LENGTHCLASS46	LENGTHCLASS47	LENGTHCLASS48
## 1	0	1	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	6	0	3	5	0
## 5	5	0	6	3	7
## 6	0	0	0	1	2
##	LENGTHCLASS49	LENGTHCLASS50	LENGTHCLASS51	LENGTHCLASS52	LENGTHCLASS53
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	1	3	1	0	0
## 5	3	6	5	4	7
## 6	2	1	6	1	3
##	LENGTHCLASS54	LENGTHCLASS55	LENGTHCLASS56	LENGTHCLASS57	LENGTHCLASS58
## 1	0	0	10	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	4	3	0	5	4
## 6	2	5	0	4	1
##	LENGTHCLASS59	LENGTHCLASS60	LENGTHCLASS61	LENGTHCLASS62	LENGTHCLASS63
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	1	2	1	1	1

## 6	7	3	5	1	4
##	LENGTHCLASS64	LENGTHCLASS65	LENGTHCLASS66	LENGTHCLASS67	LENGTHCLASS68
## 1	0	0	0	18	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	1	2	0	0	0
##	LENGTHCLASS69	LENGTHCLASS70	LENGTHCLASS71	LENGTHCLASS72	LENGTHCLASS73
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	1	0	0	0
##	LENGTHCLASS74	LENGTHCLASS75	LENGTHCLASS76	LENGTHCLASS77	LENGTHCLASS78
## 1	0	0	0	0	17
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS79	LENGTHCLASS80	LENGTHCLASS81	LENGTHCLASS82	LENGTHCLASS83
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS84	LENGTHCLASS85	LENGTHCLASS86	LENGTHCLASS87	LENGTHCLASS88
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS89	LENGTHCLASS90	LENGTHCLASS91	LENGTHCLASS92	LENGTHCLASS93
## 1	27	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS94	LENGTHCLASS95	LENGTHCLASS96	LENGTHCLASS97	LENGTHCLASS98
## 1	0	0	0	0	0
## 2	0	0	0	0	0
## 3	0	0	0	0	0
## 4	0	0	0	0	0
## 5	0	0	0	0	0
## 6	0	0	0	0	0
##	LENGTHCLASS99	LENGTHCLASS100_PLUS	COMMENTS		

## 1	0	0	NA
## 2	0	0	NA
## 3	0	0	NA
## 4	0	0	NA
## 5	0	0	NA
## 6	0	0	NA