

Annex IV - STREAM: Auxiliary scripts for the conversion from RCG Sampling and Landing tables to COST format

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Tools

R, Rstudio and packages.

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

#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(reshape2)
library(reshape)
library(dplyr)
library(knitr)
library(pander)
library(data.table)
library(COSTcore)
library(COSTdbe)
library(COSTeda)

#pander options
panderOptions('table.split.table', 60)
panderOptions('table.style', 'grid')
panderOptions('table.split.cells', 10)
panderOptions('table.alignment.default', 'left')
panderOptions('table.alignment.rownames', 'right')
panderOptions('decimal.mark', ',')
panderOptions('graph.fontsize', '10')
```

script 01: RCG_to_CS_COST_object

This script allow to convert the RCG Data call format into the SDEF format as a CS COST object.

The RCG Data call format allows to provide the data according the following two modalities: 1) aggregated data related to sex, maturity and age by length class (leaving empty fish ID and

individual weight) 2) individual data related to length, sex, maturity, age and weight (necessarily providing the Fish ID to avoid duplicated records)

Settings

set the working directory

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

```
setwd(myWD)
```

```
log_varFilePath <- paste(getwd(), "/log_CS.csv", sep="")  
path.data <- getwd()
```

```
log_var <- TRUE  
error <- FALSE
```

```
fpKey <- function(tab, colIndex, sep = ":-:") {  
  key <- tab[, colIndex]  
  key <- apply(key, 1, paste, collapse = sep)  
  key <- gsub("[[:space:]]", "", key)  
  return(key)  
}
```

CS can be in both above-mentioned modalities

```
CS <- read.csv(file="DPS_GSA99.csv",  
              stringsAsFactors=FALSE, sep=";")
```

```
# CS <- read.csv(file="SDEF_STREAM_final_HKE_examples_NOid.csv",  
# stringsAsFactors=FALSE, sep=";")
```

```
dataset_proj <- "STREAM project"
```

Input Data

RCG Datacall format

Table continues below

Sampling.type	Flag.country	Year	Trip.code
S	COUNTRY1	1900	01_18_2017
S	COUNTRY1	1900	01_18_2017
S	COUNTRY1	1900	01_18_2017
S	COUNTRY1	1900	01_18_2017

S	COUNTRY1	1900	01_18_2017
S	COUNTRY1	1900	01_18_2017

Table continues below

Harbour	Number.of.sets...hauls.on.trip	Days.at.sea
ITBCE	5	1
ITBCE	5	1
ITBCE	5	1
ITBCE	5	1
ITBCE	5	1
ITBCE	5	1

Table continues below

Sampling.method	Aggregation.level	Station.number
SelfSampling	TRUE	999
SelfSampling	TRUE	999
SelfSampling	TRUE	999
SelfSampling	TRUE	999
SelfSampling	TRUE	999
SelfSampling	TRUE	999

Table continues below

Duration.of.fishing.operation	Initial.latitude
1200	NA
1200	NA
1200	NA
1200	NA
1200	NA
1200	NA

Table continues below

Initial.longitude	Final.latitude	Final.longitude
NA	NA	NA

NA	NA	NA
NA	NA	NA
NA	NA	NA
NA	NA	NA
NA	NA	NA

Table continues below

Depth.of.fishing.operation	Water.depth
65	NA
65	NA
65	NA
65	NA
65	NA
65	NA

Table continues below

Catch.registration	Species.registration	Date
Lan	All	08/01/1900
Lan	All	08/01/1900
Lan	All	08/01/1900
Lan	All	08/01/1900
Lan	All	08/01/1900
Lan	All	08/01/1900

Table continues below

Area	Fishing.activity.category.National
GSA99	OTB_shelf
GSA99	OTB_shelf
GSA99	OTB_shelf
GSA99	OTB_shelf
GSA99	OTB_shelf

GSA99 OTB_shelf

Table continues below

Fishing.activity.category.European.lvl.6	Species
OTB_DEF_>=40_0_0	Parapenaeus longirostris
OTB_DEF_>=40_0_0	Parapenaeus longirostris
OTB_DEF_>=40_0_0	Parapenaeus longirostris
OTB_DEF_>=40_0_0	Parapenaeus longirostris
OTB_DEF_>=40_0_0	Parapenaeus longirostris
OTB_DEF_>=40_0_0	Parapenaeus longirostris

Table continues below

Catch.category	Weight	Subsample.weight	Sex
Lan	6100	3050	M
Lan	6100	3050	F
Lan	6100	3050	F
Lan	6100	3050	M
Lan	6100	3050	F
Lan	6100	3050	F

Table continues below

Maturity.method	Maturity.Scale	Maturity.Stage
Macr	Medit's scale	2d
Macr	Medit's scale	2e
Macr	Medit's scale	2e
Macr	Medit's scale	2d
Macr	Medit's scale	2e
Macr	Medit's scale	2b

Table continues below

Ageing.method	Age	Length.code	Length.class
NA	NA	mm	18
NA	NA	mm	19

NA	NA	mm	20
NA	NA	mm	21
NA	NA	mm	21
NA	NA	mm	22

Table continues below

Number.at.length	Commercial.size.category.scale	
1	COUNTRY1	
1	COUNTRY1	
2	COUNTRY1	
1	COUNTRY1	
1	COUNTRY1	
1	COUNTRY1	
Commercial.size.category	fish.ID	Individual.weight
1	NA	NA
1	NA	NA
1	NA	NA
1	NA	NA
1	NA	NA
1	NA	NA

Processing tables

Using some data in the DG MARE Med&BS format :

```
bad.rm <- FALSE

if (log_var) {
  log_varCs <- CS
  log_varCs$duplicated_TR <- FALSE
  log_varCs$duplicated_HH <- FALSE
  log_varCs$duplicated_SL <- FALSE
  log_varCs$duplicated_HL <- FALSE
  log_varMsg <- ""
} else {
  log_varMsg <- " Consider using the log_var=TRUE parameter."
}
```

```

names(CS)[which(tolower(names(CS)) == "sampling.type")] <- "sampType"
names(CS)[which(tolower(names(CS)) == "sampling_type")] <- "sampType"
names(CS)[which(tolower(names(CS)) == "samplingtype")] <- "sampType"
names(CS)[which(tolower(names(CS)) == "flag.country")] <- "vslFlgCtry"
names(CS)[which(tolower(names(CS)) == "flag_country")] <- "vslFlgCtry"
names(CS)[which(tolower(names(CS)) == "flagcountry")] <- "vslFlgCtry"
names(CS)[which(tolower(names(CS)) == "year")] <- "year"
names(CS)[which(tolower(names(CS)) == "trip.code")] <- "trpCode"
names(CS)[which(tolower(names(CS)) == "trip_code")] <- "trpCode"
names(CS)[which(tolower(names(CS)) == "tripcode")] <- "trpCode"
names(CS)[which(tolower(names(CS)) == "number.of.sets...hauls.on.trip")] <- "foNum"
"

names(CS)[which(tolower(names(CS)) == "number_of_setshauls")] <- "foNum"
names(CS)[which(tolower(names(CS)) == "nsets")] <- "foNum"
names(CS)[which(tolower(names(CS)) == "days.at.sea")] <- "daysAtSea"
names(CS)[which(tolower(names(CS)) == "days_at_sea")] <- "daysAtSea"
names(CS)[which(tolower(names(CS)) == "daysatsea")] <- "daysAtSea"
names(CS)[which(tolower(names(CS)) == "sampling.method")] <- "sampMeth"
names(CS)[which(tolower(names(CS)) == "sampling_method")] <- "sampMeth"
names(CS)[which(tolower(names(CS)) == "aggregation.level")] <- "aggLev"
names(CS)[which(tolower(names(CS)) == "aggregation_level")] <- "aggLev"
names(CS)[which(tolower(names(CS)) == "aggregationlevel")] <- "aggLev"
names(CS)[which(tolower(names(CS)) == "station.number")] <- "staNum"
names(CS)[which(tolower(names(CS)) == "station_number")] <- "staNum"
names(CS)[which(tolower(names(CS)) == "stationnumber")] <- "staNum"
names(CS)[which(tolower(names(CS)) == "catch.registration")] <- "catReg"
names(CS)[which(tolower(names(CS)) == "catch_registration")] <- "catReg"
names(CS)[which(tolower(names(CS)) == "catchregistration")] <- "catReg"
names(CS)[which(tolower(names(CS)) == "species.registration")] <- "sppReg"
names(CS)[which(tolower(names(CS)) == "species_registration")] <- "sppReg"
names(CS)[which(tolower(names(CS)) == "speciesregistration")] <- "sppReg"
names(CS)[which(tolower(names(CS)) == "date")] <- "date"
names(CS)[which(tolower(names(CS)) == "area")] <- "area"
names(CS)[which(tolower(names(CS)) ==
    "fishing.activity.category.national")] <- "foCatNat"
names(CS)[which(tolower(names(CS)) ==
    "fishingactivitycategorynational")] <- "foCatNat"
names(CS)[which(tolower(names(CS)) == "fac_national")] <- "foCatNat"
names(CS)[which(tolower(names(CS)) ==
    "fishing.activity.category.european.lvl.6")] <- "foCatEu6"
names(CS)[which(tolower(names(CS)) == "metier")] <- "foCatEu6"
names(CS)[which(tolower(names(CS)) == "fac_ec_lvl6")] <- "foCatEu6"
names(CS)[which(tolower(names(CS)) == "species")] <- "spp"
names(CS)[which(tolower(names(CS)) == "catch.category")] <- "catchCat"
names(CS)[which(tolower(names(CS)) == "catch_category")] <- "catchCat"
names(CS)[which(tolower(names(CS)) == "catchcategory")] <- "catchCat"
names(CS)[which(tolower(names(CS)) == "commercial.size.category")] <- "commCat"
names(CS)[which(tolower(names(CS)) == "commercial_size_category")] <- "commCat"

```

```

names(CS)[which(tolower(names(CS)) == "weight")] <- "wt"
names(CS)[which(tolower(names(CS)) == "subsample.weight")] <- "subSampWt"
names(CS)[which(tolower(names(CS)) == "subsample_weight")] <- "subSampWt"
names(CS)[which(tolower(names(CS)) == "subsampleweight")] <- "subSampWt"
names(CS)[which(tolower(names(CS)) == "length.code")] <- "lenCode"
names(CS)[which(tolower(names(CS)) == "length_code")] <- "lenCode"
names(CS)[which(tolower(names(CS)) == "lengthcode")] <- "lenCode"
names(CS)[which(tolower(names(CS)) == "length.class")] <- "lenCls"
names(CS)[which(tolower(names(CS)) == "length_class")] <- "lenCls"
names(CS)[which(tolower(names(CS)) == "lengthclass")] <- "lenCls"
names(CS)[which(tolower(names(CS)) == "number.at.length")] <- "lenNum"
names(CS)[which(tolower(names(CS)) == "number_at_length")] <- "lenNum"
names(CS)[which(tolower(names(CS)) == "numberatlength")] <- "lenNum"

```

new Fields RCG 2018 -----

```

names(CS)[which(tolower(names(CS)) == "commercial.size.category.scale")] <- "commC
atSc1"
names(CS)[which(tolower(names(CS)) == "commercial_size_category_scale")] <- "commC
atSc1"
names(CS)[which(tolower(names(CS)) == "commercialsizecategoryscale")] <- "commCatS
cl"

names(CS)[which(tolower(names(CS)) == "fish.id")] <- "fishId"
names(CS)[which(tolower(names(CS)) == "fish_id")] <- "fishId"
names(CS)[which(tolower(names(CS)) == "fishid")] <- "fishId"

names(CS)[which(tolower(names(CS)) == "individual.weight")] <- "indWt"
names(CS)[which(tolower(names(CS)) == "individual_weight")] <- "indWt"
names(CS)[which(tolower(names(CS)) == "individualweight")] <- "indWt"

names(CS)[which(tolower(names(CS)) == "harbour")] <- "harbour"

names(CS)[which(tolower(names(CS)) == "duration.of.fishing.operation")] <- "foDur"
names(CS)[which(tolower(names(CS)) == "duration_of_fishing_operation")] <- "foDur"
names(CS)[which(tolower(names(CS)) == "durationoffishing_operation")] <- "foDur"

names(CS)[which(tolower(names(CS)) == "initial.latitude")] <- "latIni"
names(CS)[which(tolower(names(CS)) == "initial_latitude")] <- "latIni"
names(CS)[which(tolower(names(CS)) == "initiallatitude")] <- "latIni"

names(CS)[which(tolower(names(CS)) == "initial.longitude")] <- "lonIni"
names(CS)[which(tolower(names(CS)) == "initial_longitude")] <- "lonIni"

names(CS)[which(tolower(names(CS)) == "final.latitude")] <- "latFin"
names(CS)[which(tolower(names(CS)) == "final_latitude")] <- "latFin"
names(CS)[which(tolower(names(CS)) == "finallatitude")] <- "latFin"

```



```

names(CS)[which(tolower(names(CS)) == "final.longitude")] <- "lonFin"
names(CS)[which(tolower(names(CS)) == "final_longitude")] <- "lonFin"
names(CS)[which(tolower(names(CS)) == "finallongitude")] <- "lonFin"

names(CS)[which(tolower(names(CS)) == "depth.of.fishing.operation")] <- "foDep"
names(CS)[which(tolower(names(CS)) == "depth_of_fishing_operation")] <- "foDep"
names(CS)[which(tolower(names(CS)) == "depthoffishingoperation")] <- "foDep"

names(CS)[which(tolower(names(CS)) == "water.depth")] <- "waterDep"
names(CS)[which(tolower(names(CS)) == "water_depth")] <- "waterDep"
names(CS)[which(tolower(names(CS)) == "waterdepth")] <- "waterDep"

names(CS)[which(tolower(names(CS)) == "sex")] <- "sex"

names(CS)[which(tolower(names(CS)) == "maturity.method")] <- "matMeth"
names(CS)[which(tolower(names(CS)) == "maturity_method")] <- "matMeth"
names(CS)[which(tolower(names(CS)) == "maturitymethod")] <- "matMeth"

names(CS)[which(tolower(names(CS)) == "maturity.scale")] <- "matScale"
names(CS)[which(tolower(names(CS)) == "maturity_scale")] <- "matScale"
names(CS)[which(tolower(names(CS)) == "maturityscale")] <- "matScale"

names(CS)[which(tolower(names(CS)) == "maturity.stage")] <- "matStage"
names(CS)[which(tolower(names(CS)) == "maturity_stage")] <- "matStage"
names(CS)[which(tolower(names(CS)) == "maturitystage")] <- "matStage"

names(CS)[which(tolower(names(CS)) == "ageing.method")] <- "ageMeth"
names(CS)[which(tolower(names(CS)) == "ageing_method")] <- "ageMeth"
names(CS)[which(tolower(names(CS)) == "ageingmethod")] <- "ageMeth"

names(CS)[which(tolower(names(CS)) == "age")] <- "age"

CS[is.na(CS[,])] <- -1

CS$aggLev <- as.character(CS$aggLev)
CS$aggLev[ CS$aggLev %in% c("t","TRUE",TRUE)] <- "T" ###

CS$spp <- unlist(lapply(CS$spp, function(x) paste(toupper(substring(x,1, 1)),
                                                  tolower(substring(x, 2)), sep = "")))

CS <- CS %>% mutate(proj=dataset_proj,landCtry=vs1FlgCtry) ###

trPk <- c("sampType", "vs1FlgCtry", "year", "trpCode","proj","landCtry") ###
trOther <- c("foNum", "daysAtSea", "sampMeth",
            "harbour" ) # new : harbour

```

```

hhPk <- c(trPk, "staNum")
hhOther <- c("aggLev", "catReg", "sppReg", "date", "area",
            "foCatNat", "foCatEu6",
            "foDur", "latIni", "lonIni", "latFin", "lonFin", "foDep", "waterDep")
# new

# modified respect to the previous version
slPk <- c(hhPk, "spp", "catchCat", "commCat", "commCatScl", "sex") # new
slOther <- c("wt", "subSampWt", "lenCode")

#slPk <- c(hhPk, "spp", "catchCat", "commCat", "commCatScl", "sex", "wt", "subSampW
t") # new
#slOther <- c("lenCode")

hlPk <- c(slPk, "lenCls")
hlOther <- c("lenNum")

# modified respect to the previous version
#caPk <- c(hlPk, "age", "area", "fishId")
#caOther <- c("matMeth", "matScale", "matStage", "ageMeth", "indWt", "lenCode")

caPk <- c(hlPk, "age", "area", "fishId", "matStage")
caOther <- c("matMeth", "matScale", "ageMeth", "indWt", "lenCode")

allFields <- c(caPk, trOther, hhOther, slOther, hlOther, caOther)
missingFields <- allFields[!allFields %in% names(CS)]

# check if all fields are used

# names(CS)[!names(CS) %in% allFields]

if (length(missingFields) > 0) {
  stop("Missing fields : ", paste(missingFields, collapse = ", ",
                                   sep = ""))
}

csTr <- unique(CS[, c(trPk, trOther)])

trPkV <- fpKey(csTr, trPk)

trPkVDup <- trPkV %in% trPkV[duplicated(trPkV)]
if (any(trPkVDup)) {
  print(trPkVDup)
  if (log_var) {
    log_varCs$duplicated_TR <- fpKey(CS, trPk) %in% trPkV[trPkVDup]
  }
  if (bad.rm) {
    message("Integrity problem for CS/TR, ", sum(trPkVDup),
            " row(s) removed.", log_varMsg)
  }
}

```

```

    print("Removed following row(s):", quote = F)
    print(csTr[trPkVDup, ])
    CS <- merge(CS, csTr[!trPkVDup, ])
  } else {
    error <- TRUE
    message("Integrity problem for CS/TR, ", sum(trPkVDup),
            " row(s) concerned.", log_varMsg)
  }
}
print(paste("No rows TR =", nrow(csTr)), quote=F)

## [1] No rows TR = 69

csHh <- unique(CS[, c(hhPk, hhOther)])
hhPkv <- fpKey(csHh, hhPk)
hhPkVDup <- hhPkv %in% hhPkv[duplicated(hhPkv)]
if (any(hhPkVDup)) {
  if (log_var) {
    log_varCs$duplicated_HH <- fpKey(CS, hhPk) %in% hhPkv[hhPkVDup]
  }
  if (bad.rm) {
    message("Integrity problem for CS/HH, ", sum(hhPkVDup),
            " row(s) removed.", log_varMsg)
    print("Removed following row(s):", quote = F)
    print(csHh[hhPkVDup, ])
    CS <- merge(CS, csHh[!hhPkVDup, ])
  }
  else {
    error <- TRUE
    message("Integrity problem for CS/HH, ", sum(hhPkVDup),
            " row(s) concerned.", log_varMsg)
  }
}
print(paste("No rows HH =", nrow(csHh)), quote=F)

## [1] No rows HH = 69

csSl <- unique(CS[, c(slPk, slOther)])
slPkv <- fpKey(csSl, slPk)
slPkVDup <- slPkv %in% slPkv[duplicated(slPkv)]
if (any(slPkVDup)) {
  if (log_var) {
    log_varCs$duplicated_SL <- fpKey(CS, slPk) %in% slPkv[slPkVDup]
  }
  if (bad.rm) {
    message("Integrity problem for CS/SL, ", sum(slPkVDup),
            " row(s) removed.", log_varMsg)
    print("Removed following row(s):", quote = F)
    print(csSl[slPkVDup, ])
    CS <- merge(CS, csSl[!slPkVDup, ])
  }
  else {

```

```

    error <- TRUE
    message("Integrity problem for CS/SL, ", sum(slPkVDup),
           " row(s) concerned.", log_varMsg)
  }
}
print(paste("No rows SL =", nrow(csSl)), quote=F)
## [1] No rows SL = 413

CS_aggregated_by_length <- aggregate(CS$lenNum, by=list(CS$sampType , CS$vslFlgC
try ,
               CS$year, CS$trpCode, CS$proj, CS$landCtry, CS$harbou
r,
               CS$foNum, CS$daysAtSea , CS$sampMeth, CS$aggLev, CS$
staNum,
               CS$foDur, CS$foDep, CS$catReg, CS$sppReg, CS$date, CS
$area,
               CS$foCatNat, CS$foCatEu6, CS$spp, CS$catchCat, CS$wt
,
               CS$subSampWt, CS$sex, CS$lenCode, CS$lenCls,
               CS$commCatSc1, CS$commCat), FUN="sum")

colnames(CS_aggregated_by_length) <- c("sampType" , "vslFlgCtry" , "year" , "trpC
ode" ,
               "proj" , "landCtry","harbour" , "foNum" , "daysAtSea"
,
               "sampMeth" , "aggLev" , "staNum" , "foDur" , "foDep"
,
               "catReg", "sppReg" , "date" , "area" , "foCatNat" ,
               "foCatEu6", "spp" , "catchCat" , "wt" , "subSampWt" ,
               "sex" , "lenCode" , "lenCls" , "commCatSc1" , "commCat
",
               "lenNum")

# sum(CS$LenNum)
# sum(CS_aggregated_by_length$LenNum)

csH1 <- unique(CS_aggregated_by_length[, c(h1Pk, h1Other)])
h1PkV <- fpKey(csH1, h1Pk)
h1PkVDup <- h1PkV %in% h1PkV[duplicated(h1PkV)]
if (any(h1PkVDup)) {
  if (log_var) {
    log_varCs$duplicated_HL <- fpKey(CS, h1Pk) %in% h1PkV[h1PkVDup]
  }
  else {
    error <- TRUE
    message("Integrity problem for CS/HL, ", sum(h1PkVDup),
           " row(s) concerned.", log_varMsg)
    print("Check the following row(s):", quote = F)
    csH1[h1PkVDup, ]
  }
}

```

```

}
print(paste("No rows HL =", nrow(csH1)), quote=F)

## [1] No rows HL = 4383

##

# # check CA -----

csCa <- unique(CS[, c(caPk, caOther)])
caPkV <- fpKey(csCa, caPk)
caPkVDup <- caPkV %in% caPkV[duplicated(caPkV)]
if (any(caPkVDup)) {
  if (log_var) {
    log_varCs$duplicated_CA <- fpKey(CS, caPk) %in% caPkV[caPkVDup]
  }
  if (bad.rm) {
    message("Integrity problem for CS/CA, ", sum(caPkVDup),
            " row(s) removed.", log_varMsg)
    print("Removed following row(s):", quote = F)
    print(csCa[caPkVDup, ])
    CS <- merge(CS, csCa[!caPkVDup, ])
  } else {
    error <- TRUE
    message("Integrity problem for CS/CA, ", sum(caPkVDup),
            " row(s) concerned.", log_varMsg)
    print("Check the following row(s):", quote = F)
    csCa[caPkVDup, ]
  }
}

##

if (log_var) {
  if (missing(log_varFilePath)) {
    log_varFilePath <- tempfile(fileext = ".csv")
  }
  write.table(log_varCs, file = log_varFilePath, row.names = FALSE,
              sep = ";")
}
if (error) {
  print("See errors on log_var file!", quote = F)
  message("log_var file: ", log_varFilePath)
}

# ALL CS tables names -----

### Simpler way to define names for the CS tables
obj <- new("csData")

```

```

tr0 <- obj@tr
hh0 <- obj@hh
sl0 <- obj@sl
hl0 <- obj@hl
ca0 <- obj@ca

TR.col <- names(tr0)
HH.col <- names(hh0)
SL.col <- names(sl0)
HL.col <- names(hl0)
CA.col <- names(ca0)

## end names ###

if (!error) {

  missTR<-TR.col[!TR.col %in% names(csTr)]
  costCS.TR <- csTr
  costCS.TR[ ,missTR] <- NA
  costCS.TR<- costCS.TR[,TR.col]

  write.table(costCS.TR, file.path(path.data, "SDEF CS-TR data.csv"),
              sep = ";", row.names = FALSE)

  missHH<-HH.col[!HH.col %in% names(csHh)]
  costCS.HH <- csHh
  costCS.HH[ , missHH] <- NA

  costCS.HH<- costCS.HH[,HH.col]

  write.table(costCS.HH, file.path(path.data, "SDEF CS-HH data.csv"),
              sep = ";", row.names = FALSE)

  missSL<-SL.col[!SL.col %in% names(csSl)]
  costCS.SL <- csSl
  costCS.SL[ , missSL] <- NA

  costCS.SL<- costCS.SL[,SL.col]

  write.table(costCS.SL, file.path(path.data, "SDEF CS-SL data.csv"),
              sep = ";", row.names = FALSE)

  missHL<-HL.col[!HL.col %in% names(csHl)]
  costCS.HL <- csHl
  costCS.HL[ , missHL] <- NA

  costCS.HL <- costCS.HL[,HL.col]

```

```

write.table(costCS.HL, file.path(path.data, "SDEF CS-HL data.csv"),
            sep = ";", row.names = FALSE)

if (all(CS$fishId == -1)) {

  ## add subSampWt (SL) and lenNum (HL) to CA

  ca.sl.hl.COL<-c(caPk, caOther,"subSampWt","lenNum")
  ca.sl.hl <- unique(CS[, ca.sl.hl.COL])

  # indWt
  ca.sl.hl<-ca.sl.hl %>% mutate(indWt=subSampWt/lenNum)
  ca.sl.hl1 <- ca.sl.hl[rep(row.names(ca.sl.hl), ca.sl.hl$lenNum), 1:ncol(ca.sl.
hl)]
  ca.sl.hl1$fishId <- 1:nrow(ca.sl.hl1)
  costCS.CA <- ca.sl.hl1[, names( ca.sl.hl1) %in% CA.col]

  missCA<-CA.col[!CA.col %in% names(costCS.CA)]
  costCS.CA[ , missCA] <- NA
  costCS.CA<- costCS.CA[,CA.col]
  costCS.CA$indWt <- -1

} else {

  missCA <- CA.col[!CA.col %in% names(csCa)]
  costCS.CA <- csCa
  costCS.CA[ , missCA] <- NA

  costCS.CA$proj<- dataset_proj
  costCS.CA$landCtry<-costCS.CA$vs1FlgCtry
  costCS.CA<- costCS.CA[,CA.col]

  costCS.CA$fishId=seq(1,dim(costCS.CA)[1],by=1)

}

print(paste("No rows CA =", nrow(costCS.CA)), quote=F)

write.table(costCS.CA, file.path(path.data, "SDEF CS-CA data.csv"),
            sep = ";", row.names = FALSE)

costCS = csData(tr =costCS.TR, hh = costCS.HH, sl = costCS.SL,
               hl = costCS.HL, ca=costCS.CA)

saveRDS(costCS, "costCS.rds")

} else {

  print("An error occurred in the trasformation.

```

```

    Impossible to save the CS COST object!")
}

## [1] No rows CA = 80632

```

Output

CS COST object

```

## An object of class "csData"
## Slot "desc":
## [1] "Unknown stock"
##
## Slot "tr":
##      sampType landCtry vslFlgCtry year      proj      trpCode vslLen
## 1          S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017    NA
## 83         S COUNTRY1  COUNTRY1 1900 STREAM project 02_18_2017    NA
## 204        S COUNTRY1  COUNTRY1 1900 STREAM project 03_18_2017    NA
## 357        S COUNTRY1  COUNTRY1 1900 STREAM project 09_18_2017    NA
## 391        S COUNTRY1  COUNTRY1 1900 STREAM project 101_18_2017   NA
## 530        S COUNTRY1  COUNTRY1 1900 STREAM project 110_18_2017   NA
##      vslPwr vslSize vslType harbour foNum daysAtSea vslId sampCtry
## 1          NA      NA      <NA>  ITBCE      5          1      NA      <NA>
## 83          NA      NA      <NA>  ITBDS      2          1      NA      <NA>
## 204         NA      NA      <NA>  ITBDS      4          1      NA      <NA>
## 357         NA      NA      <NA>  ITMFR     11          2      NA      <NA>
## 391         NA      NA      <NA>  ITMNP      4          1      NA      <NA>
## 530         NA      NA      <NA>  ITMFR     11          2      NA      <NA>
##      sampMeth
## 1  SelfSampling
## 83   Observer
## 204  Observer
## 357 SelfSampling
## 391   Observer
## 530 SelfSampling
##
## Slot "hh":
##      sampType landCtry vslFlgCtry year      proj      trpCode staNum
## 1          S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017    999
## 83         S COUNTRY1  COUNTRY1 1900 STREAM project 02_18_2017    999
## 204        S COUNTRY1  COUNTRY1 1900 STREAM project 03_18_2017    999
## 357        S COUNTRY1  COUNTRY1 1900 STREAM project 09_18_2017    999
## 391        S COUNTRY1  COUNTRY1 1900 STREAM project 101_18_2017   999
## 530        S COUNTRY1  COUNTRY1 1900 STREAM project 110_18_2017   999
##      foVal aggLev catReg sppReg      date time foDur latIni lonIni latFin
## 1      <NA>      T   Lan   All 08/01/1900 <NA>  1200     -1     -1     -1
## 83      <NA>      T   All   All 21/01/1900 <NA>   510     -1     -1     -1
## 204      <NA>      T   All   All 31/01/1900 <NA>  1155     -1     -1     -1
## 357      <NA>      T   Lan   All 29/01/1900 <NA>  2880     -1     -1     -1
## 391      <NA>      T   All   All 17/01/1900 <NA>   960     -1     -1     -1
## 530      <NA>      T   Lan   All 22/01/1900 <NA>  1980     -1     -1     -1
##      lonFin area rect subRect foDep waterDep foCatNat foCatEu5

```



```

## 1      -1 GSA99 <NA>      <NA>      65      -1 OTB_shelf      <NA>
## 83      -1 GSA99 <NA>      <NA>      188     -1 OTB_shelf      <NA>
## 204     -1 GSA99 <NA>      <NA>      115     -1 OTB_shelf      <NA>
## 357     -1 GSA99 <NA>      <NA>      148     -1 OTB_shelf      <NA>
## 391     -1 GSA99 <NA>      <NA>       81     -1 OTB_shelf      <NA>
## 530     -1 GSA99 <NA>      <NA>      104     -1 OTB_shelf      <NA>
##          foCatEu6 meshSize selDev meshSizeSelDev
## 1  OTB_DEF_>=40_0_0      NA  <NA>      NA
## 83  OTB_DEF_>=40_0_0      NA  <NA>      NA
## 204 OTB_DEF_>=40_0_0      NA  <NA>      NA
## 357 OTB_DEF_>=40_0_0      NA  <NA>      NA
## 391 OTB_DEF_>=40_0_0      NA  <NA>      NA
## 530 OTB_DEF_>=40_0_0      NA  <NA>      NA
##
## Slot "s1":
##      sampType landCtry vs1FlgCtry year      proj      trpCode staNum
## 1      S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017    999
## 2      S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017    999
## 33     S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017    999
## 36     S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017    999
## 60     S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017    999
## 61     S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017    999
##          spp catchCat landCat commCatScl commCat subSampCat
## 1 Parapenaeus longirostris      Lan  <NA>  COUNTRY1      1      <NA>
## 2 Parapenaeus longirostris      Lan  <NA>  COUNTRY1      1      <NA>
## 33 Parapenaeus longirostris      Lan  <NA>  COUNTRY1      2      <NA>
## 36 Parapenaeus longirostris      Lan  <NA>  COUNTRY1      2      <NA>
## 60 Parapenaeus longirostris      Lan  <NA>  COUNTRY1      3      <NA>
## 61 Parapenaeus longirostris      Lan  <NA>  COUNTRY1      3      <NA>
##      sex      wt subSampWt lenCode
## 1      M  6100      3050      mm
## 2      F  6100      3050      mm
## 33     M 38080      2720      mm
## 36     F 38080      2720      mm
## 60     F  5600      1400      mm
## 61     M  5600      1400      mm
##
## Slot "h1":
##      sampType landCtry vs1FlgCtry year      proj      trpCode staNum
## 1      S COUNTRY1  COUNTRY1 1900 STREAM project 14_19_2017    999
## 2      S COUNTRY1  COUNTRY1 1900 STREAM project 02_19_2017    999
## 3      S COUNTRY1  COUNTRY1 1900 STREAM project 38_19_2017    999
## 4      S COUNTRY1  COUNTRY1 1900 STREAM project 02_19_2017    999
## 5      S COUNTRY1  COUNTRY1 1900 STREAM project 125_18_2017    999
## 6      S COUNTRY1  COUNTRY1 1900 STREAM project 14_19_2017    999
##          spp catchCat landCat commCatScl commCat subSampCat
## 1 Parapenaeus longirostris      Lan  <NA>  COUNTRY1      1      <NA>
## 2 Parapenaeus longirostris      Lan  <NA>  COUNTRY1      1      <NA>
## 3 Parapenaeus longirostris      Lan  <NA>  COUNTRY1      1      <NA>
## 4 Parapenaeus longirostris      Lan  <NA>  COUNTRY1      1      <NA>
## 5 Parapenaeus longirostris      Lan  <NA>  COUNTRY1      1      <NA>

```

```
## 6 Parapenaeus longirostris      Lan      <NA>      COUNTRY1      1      <NA>
##      sex lenCls lenNum
## 1      F      11      6
## 2      F      11      3
## 3      M      11      1
## 4      F      12      8
## 5      F      12      1
## 6      M      12      2
##
## Slot "ca":
##      sampType landCtry vsFlgCtry year      proj      trpCode staNum
## 1              S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017 999
## 2              S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017 999
## 3              S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017 999
## 3.1            S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017 999
## 4              S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017 999
## 5              S COUNTRY1  COUNTRY1 1900 STREAM project 01_18_2017 999
##      quarter month      spp sex catchCat landCat commCatSc1
## 1          NA      NA Parapenaeus longirostris      M      Lan      <NA>      COUNTRY1
## 2          NA      NA Parapenaeus longirostris      F      Lan      <NA>      COUNTRY1
## 3          NA      NA Parapenaeus longirostris      F      Lan      <NA>      COUNTRY1
## 3.1        NA      NA Parapenaeus longirostris      F      Lan      <NA>      COUNTRY1
## 4          NA      NA Parapenaeus longirostris      M      Lan      <NA>      COUNTRY1
## 5          NA      NA Parapenaeus longirostris      F      Lan      <NA>      COUNTRY1
##      commCat stock  area rect subRect lenCls age fishId lenCode ageMeth
## 1          1 <NA> GSA99 <NA>      <NA>      18 -1      1      mm      -1
## 2          1 <NA> GSA99 <NA>      <NA>      19 -1      2      mm      -1
## 3          1 <NA> GSA99 <NA>      <NA>      20 -1      3      mm      -1
## 3.1        1 <NA> GSA99 <NA>      <NA>      20 -1      4      mm      -1
## 4          1 <NA> GSA99 <NA>      <NA>      21 -1      5      mm      -1
## 5          1 <NA> GSA99 <NA>      <NA>      21 -1      6      mm      -1
##      plusGrp otoWt otoSide indWt matMeth      matScale matStage
## 1          <NA>      NA      <NA>      -1      Macr Medits scale      2d
## 2          <NA>      NA      <NA>      -1      Macr Medits scale      2e
## 3          <NA>      NA      <NA>      -1      Macr Medits scale      2e
## 3.1        <NA>      NA      <NA>      -1      Macr Medits scale      2e
## 4          <NA>      NA      <NA>      -1      Macr Medits scale      2d
## 5          <NA>      NA      <NA>      -1      Macr Medits scale      2e
```

script 02: RCG_CL_to_COST_CL

This script allow to convert the RCG Landing table into CL COST object

Settings

```
# set the working directory
myWD <- paste("C:\\Users\\Bitetto Isabella\\OneDrive - Coispa Tecnologia & Ricerca
S.C.A.R.L\\MARE22\\STREAM\\FINAL REVISION OF DELIVERABLES\\RCG_to_COST", sep="")

logFilePath <- paste(getwd(), "/log_CL.csv", sep="")
path.data <- getwd()
```


Table continues below

commCatScl	commCat	foCatNat	foCatEu5
NA	NA	OTB_shelf	NA
NA	NA	OTB_shelf	NA
NA	NA	OTB_shelf	NA
NA	NA	OTB_shelf	NA
NA	NA	OTB_shelf	NA
NA	NA	OTB_shelf	NA

Table continues below

foCatEu6	harbour	vslLenCat
OTB_DEF_>=40_0_0	Port	NA
OTB_DEF_>=40_0_0	Port	NA
OTB_DEF_>=40_0_0	Port	NA
OTB_DEF_>=40_0_0	Port	NA
OTB_DEF_>=40_0_0	Port	NA
OTB_DEF_>=40_0_0	Port	NA

Table continues below

unallocCatchWt	misRepCatchWt	landWt	landMult
NA	NA	73453	NA
NA	NA	78742	NA
NA	NA	82021	NA
NA	NA	89023	NA
NA	NA	103912	NA
NA	NA	102987	NA

landValue

372659
392665
460281
433524
494103

498299

Processing tables

Using some data in the DG MARE Med&BS format :

```
if (log) {
logCl <- CL
logCl$duplicated <- FALSE
logMsg <- ""
} else {
logMsg <- " Consider using the log=TRUE parameter."
}
error <- FALSE

names(CL)[which(tolower(names(CL)) == "flag.country")] <- "vslFlgCtry"
names(CL)[which(tolower(names(CL)) == "flag_country")] <- "vslFlgCtry"

names(CL)[which(tolower(names(CL)) == "year")] <- "year"
names(CL)[which(tolower(names(CL)) == "quarter")] <- "quarter"
names(CL)[which(tolower(names(CL)) == "month")] <- "month"
names(CL)[which(tolower(names(CL)) == "area")] <- "area"
names(CL)[which(tolower(names(CL)) == "species")] <- "taxon"

names(CL)[which(tolower(names(CL)) ==
                 "fishing.activity.category.national")] <- "foCatNat"
names(CL)[which(tolower(names(CL)) == "fac_national")] <- "foCatNat"

names(CL)[which(tolower(names(CL)) ==
                 "fishing.activity.category.european.lvl.6")] <- "foCatEu6"
names(CL)[which(tolower(names(CL)) == "fac_ec_lvl6")] <- "foCatEu6"

names(CL)[which(tolower(names(CL)) == "harbour")] <- "harbour"

names(CL)[which(tolower(names(CL)) == "official.landings.weight")] <- "landWt"
names(CL)[which(tolower(names(CL)) == "official_landings_weight")] <- "landWt"

names(CL)[which(tolower(names(CL)) == "official.landings.value")] <- "landValue"
names(CL)[which(tolower(names(CL)) == "official_landings_value")] <- "landValue"

## primary keys & fields

clPk <- c("vslFlgCtry", "year", "quarter", "month", "area", "taxon",
         "foCatNat", "foCatEu6")
clOther <- c("landWt", "landValue")

# check fields
allFields <- c(clPk, clOther)
```

```

missingFields <- allFields[! allFields %in% names(CL)]
if (length(missingFields) > 0) {
stop("Missing fields : ", paste(missingFields, collapse = ", ", sep=""))
}

c1PkV <- fpKey(CL, c1Pk)
c1PkVDup <- c1PkV %in% c1PkV[duplicated(c1PkV)]

# test integrity
if (any(c1PkVDup)) {
if (log) {
logCl$duplicated <- fpKey(CL, c1Pk)
}

if (bad.rm) {
message("Integrity problem for CL, ", sum(c1PkVDup),
" row(s) removed.", logMsg)
CL <- CL[! c1PkVDup,]
} else {
error <- TRUE
message("Integrity problem for CL, ", sum(c1PkVDup),
" row(s) concerned.", logMsg)
}
}

if (log) {
if (missing(logFilePath)) {
logFilePath <- tempfile(fileext = ".csv")
}
write.table(logCl, file=logFilePath, row.names = FALSE, sep=";")
message("Log file: ", logFilePath)
}
if (error) {
stop("Stop on reported errors.")
}

# forming

CL$taxon <- unlist(lapply(CL$taxon, function(x)
paste(toupper(substring(x, 1, 1)), tolower(substring(x, 2)), sep="")))

## df
c1Df <- data.frame(
landCtry=NA,
vs1FlgCtry=CL$vs1FlgCtry,

```

```

year=CL$year,
quarter=CL$quarter,
month=CL$month,
area=CL$area,
rect=NA,
subRect=NA,
taxon=CL$taxon,
landCat=NA,
commCatSc1=NA,
commCat=NA,
foCatNat=CL$foCatNat,
foCatEu5=NA,
foCatEu6=CL$foCatEu6,
harbour=CL$harbour,
vs1LenCat=NA,
unallocCatchWt=NA,
misRepCatchWt=NA,
landWt=CL$landWt,
landMult=NA,
landValue=CL$landValue,
stringsAsFactors=FALSE)

```

```

write.table(c1Df, file.path(path.data, "SDEF CL data.csv"), sep=";", row.names =
FALSE)

```

```

costCL = c1Data(cl=c1Df)
saveRDS(costCL, "costCL.rds")

```

Output

CL COST object

```

## An object of class "c1Data"
## Slot "desc":
## [1] "Unknown stock"
##
## Slot "cl":
##   landCtry vs1FlgCtry year quarter month   area rect subRect
## 1    <NA>   COUNTRY1 1900         1     1 GSA99 <NA>   <NA>
## 2    <NA>   COUNTRY1 1900         1     2 GSA99 <NA>   <NA>
## 3    <NA>   COUNTRY1 1900         1     3 GSA99 <NA>   <NA>
## 4    <NA>   COUNTRY1 1900         2     4 GSA99 <NA>   <NA>
## 5    <NA>   COUNTRY1 1900         2     5 GSA99 <NA>   <NA>
## 6    <NA>   COUNTRY1 1900         2     6 GSA99 <NA>   <NA>
##
##           taxon landCat commCatSc1 commCat  foCatNat foCatEu5
## 1 Parapenaeus longirostris    <NA>      <NA>      <NA> OTB_shelf    <NA>
## 2 Parapenaeus longirostris    <NA>      <NA>      <NA> OTB_shelf    <NA>
## 3 Parapenaeus longirostris    <NA>      <NA>      <NA> OTB_shelf    <NA>
## 4 Parapenaeus longirostris    <NA>      <NA>      <NA> OTB_shelf    <NA>
## 5 Parapenaeus longirostris    <NA>      <NA>      <NA> OTB_shelf    <NA>
## 6 Parapenaeus longirostris    <NA>      <NA>      <NA> OTB_shelf    <NA>

```

##	foCatEu6	harbour	vs1LenCat	unallocCatchWt	misRepCatchWt
## 1	OTB_DEF_>=40_0_0	Port	<NA>	NA	NA
## 2	OTB_DEF_>=40_0_0	Port	<NA>	NA	NA
## 3	OTB_DEF_>=40_0_0	Port	<NA>	NA	NA
## 4	OTB_DEF_>=40_0_0	Port	<NA>	NA	NA
## 5	OTB_DEF_>=40_0_0	Port	<NA>	NA	NA
## 6	OTB_DEF_>=40_0_0	Port	<NA>	NA	NA
##	landWt	landMult	landValue		
## 1	73452.53	NA	372658.8		
## 2	78741.52	NA	392665.2		
## 3	82021.10	NA	460280.8		
## 4	89022.59	NA	433524.2		
## 5	103911.92	NA	494103.4		
## 6	102987.30	NA	498298.8		