Annex 3 - R script to construct Table 1A by country

Joel Vigneau, with the collaboration of Matt Elliott, Jon Elson, Kirsten Birch Hakansson, Marie Storr-Paulsen, Nuno Prista, Katja Ringhdal, Lies Vansteenbrugge, Sieto Verver

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Filling of the EU-MAP table 1A requires to report on country shares of landings and shares of EU TAC when relevant, for all the stocks listed in table 1A of the EU-MAP Regulation (EU Decision 1254/2016). This process necessitates to gather information on landings and EU TAC from an official database, namely EUROSTAT for EU landings and MARE/FIDES for EU TAC.

Two datasets were added to complete the references, (1) the Nephrops FU landings provided by ICES and (2) the Mediterranean and Black Sea landings fisgures put together by 2016 RCM Mediterranean and Black Sea.

First of all, the datasets listed above contain information from all EU Member States, which means that the script has the potential to be used by all Member States, and by STECF for control of the NWP submitted for 2017.

Setting the parameters for the analysis

The variables needed for the work are the working directory, the country code (2-letter code) and the reference years

```
library(tidyr)
library(stringr)

CTRY <- 'DK'
refYears <- 2016:2018</pre>
```

Importing the datasets

The list of datasets are the following:

1. Landings and TAC shares files:

DF <- rbind.data.frame(DF, DF47)

- EUROSTAT landings files: http://ec.europa.eu/eurostat/web/fisheries/data/database
- MARE/FIDES TAC file: https://webgate.ec.europa.eu/fides/index.cfm
- ICES Nephrops fishery units landings per country for 2015
- RCM Mediterranean and Black Sea 2016 landings compilation

```
DF27 <- read.table(file.path(input_path_common, "fish_ca_atl27.tsv"),header=TRUE, sep='\t', as.is=TRUE
DF21 <- read.table(file.path(input_path_common, "fish_ca_atl21.tsv"),header=TRUE, sep='\t', as.is=TRUE
DF34 <- read.table(file.path(input_path_common, "fish_ca_atl34.tsv"),header=TRUE, sep='\t', as.is=TRUE
DF41 <- read.table(file.path(input_path_common, "fish_ca_atl41.tsv"),header=TRUE, sep='\t', as.is=TRUE
DF47 <- read.table(file.path(input_path_common, "fish_ca_atl47.tsv"),header=TRUE, sep='\t', as.is=TRUE
DF51 <- read.table(file.path(input_path_common, "fish_ca_ind51.tsv"),header=TRUE, sep='\t', as.is=TRUE
DF37 <- read.table(file.path(input_path_common, "fish_ca_atl37.tsv"),header=TRUE, sep='\t', as.is=TRUE
DF <- rbind.data.frame(DF27, DF21)
DF <- rbind.data.frame(DF, DF34)
DF <- rbind.data.frame(DF, DF37)
DF <- rbind.data.frame(DF, DF41)
```

```
DF <- rbind.data.frame(DF, DF51)

TAC <- read.csv(file.path(input_path_common, 'EU opening quota 2018.csv'), header=TRUE, sep=';',as.is=TR

NEP <- read.csv(file.path(input_path_common, 'Nephrops landings 2015.csv'), header=TRUE, sep=';', as.is='

MED <- read.csv(file.path(input_path_common, 'RCM MED landings.csv'), header=TRUE, sep=';', as.is=TRUE)
```

- 2. Reference tables:
- EuroStat Geo.def: full names of countries
- ASFIS file: FAO species naming and coding
- Linkage table mirroring EU-MAP Table 1A naming of species and stock area, and lining to EUROSTAT and MARE/FIDES species and area naming

```
GEO <- read.table(file.path(input_path_common,'geo.def'),header=TRUE,sep=";", as.is=TRUE)

ASFIS <- read.table(file.path(input_path_common,'ASFIS_sp_Feb_2016.txt'), header=TRUE, sep="\t", as.i

table1A <- read.csv(file.path(input_path_common,'EUMAP_Table1A_Linkage_EUROSTAT and EC_TAC.csv'), sep='
```

data.frame preparation

The country names are matching between GEO and TAC data.frame, except for UK, so the following lines enables the full match.

```
TAC$StockID <- paste(TAC$Species.Code,TAC$Area.Code,sep="")
TAC$Level.Description[substring(TAC$Level.Description,1,3) %in% 'U.K'] <- 'United Kingdom'
```

The TAC dataset is well structured and thus ready for the analysis

```
head(TAC,3)
```

```
##
     Load_ind Definition.Year Species.Code
## 1
         INIV
                          2018
                                     AGO_CSQ
## 2
         INIV
                          2018
                                     AGO CSQ
## 3
         INIV
                          2018
                                     AGO_CSQ
                                                          Species.Name Area.Code
## 1 Angola direct agreement fishing category (Coastal State quota)
                                                                           DIR AG
## 2 Angola direct agreement fishing category (Coastal State quota)
                                                                           DIR_AG
## 3 Angola direct agreement fishing category (Coastal State quota)
                                                                           DIR AG
##
      Area.Description Level.Code Level.Description Initial.Quantity
## 1 Direct Agreements
                               BEL
                                              Belgium
## 2 Direct Agreements
                               BGR.
                                             Bulgaria
                                                                     NA
## 3 Direct Agreements
                               CYP
                                               Cyprus
                                                                     NA
     Adapted.Quota Eurlex.Ref OJ.Ref Publication.Date Page.Number
##
## 1
                NA 32017R2403
                                 L347
                                             2017-12-28
## 2
                NA 32017R2403
                                 L347
                                             2017-12-28
                                                                  81
## 3
                NA 32017R2403
                                 L347
                                             2017-12-28
                                                                  81
##
     In.regulation Compute.uptake
                                          StockID
                                 Y AGO_CSQDIR_AG
## 1
                 Y
## 2
                 Y
                                 Y AGO_CSQDIR_AG
## 3
                 Y
                                 Y AGO_CSQDIR_AG
names(GEO)[2] <- "Country"</pre>
GEO$geo <- toupper(GEO$geo) #2-letter code should be in capitals
SRG <- strsplit(as.character(DF$species.fishreg.unit.geo.time),split=",")</pre>
SRG.m <- matrix(unlist(SRG), ncol=4, byrow=TRUE)</pre>
coln <- sapply(refYears, function(x) which(grepl(x,names(DF))))</pre>
```

```
DFT <- data.frame(X3A_CODE = toupper(SRG.m[,1]), area = toupper(SRG.m[,2]), geo = SRG.m[,4],
Y1 = DF[,coln[1]], Y2 = DF[,coln[2]], Y3 = DF[,coln[3]])
DFM <- merge(DFT, GEO, all.x=TRUE)
DFM$Y1 <- as.numeric(as.character(DFM$Y1))
DFM$Y2 <- as.numeric(as.character(DFM$Y2))
DFM$Y3 <- as.numeric(as.character(DFM$Y3))
DFM <- DFM[!is.na(DFM$Country),]
DFM <- merge(DFM, ASFIS[,c(3:6)], all.x=TRUE)</pre>
```

Let's have a look at the workable structure of EuroStat dataset. Note that Y1, Y2 and Y3 are the 3-year period demanded, and the presence of NA's. The assumption made here (further in the Construction of the table section) is to exclude NA from the average, i.e. like if MS had omitted to report, instead of a NA which would mean 0. The confusion comes because lots of 0 are reported in EuroStat (implicitly meaning that NA is not a 0). This point may be subject of a STECF agreement or suggestion for modification.

head(DFM,3)

```
Y1 Y2 Y3
##
     X3A_CODE geo
                                                         Country
                         area
                                NA NA NA European union (28 MS)
## 1
          AAS EU28 27 3 C 22
## 2
          AAS EU28
                      27_4_B 0.03 NA NA European union (28 MS)
## 3
          AAS
                DK
                         27_4 0.03 NA NA
                                                         Denmark
                       English_name
##
     Scientific_name
                                                  French_name
## 1 Astacus astacus Noble crayfish Écrevisse à pieds rouges
## 2 Astacus astacus Noble crayfish Écrevisse à pieds rouges
## 3 Astacus astacus Noble crayfish Écrevisse à pieds rouges
NEP <- merge(NEP, GEO, all.x=TRUE)
NEP$geo[is.na(NEP$geo)] <- 'UK'</pre>
NEP2 <- data.frame(X3A_CODE='NEP', geo=NEP$geo, area=NEP$Stock, Y1=round(NEP$TotalLanding.in.kg/1000,0)
```

A look at the Nephrops dataset on the same format as EuroStat dataset, so they can be merged

head(NEP2)

```
area Y1 Y2 Y3 Country
                                                Scientific name
##
     X3A_CODE geo
                                                                   English name
## 1
          NEP
              BE nep-22
                            5 NA NA Belgium Nephrops norvegicus Norway lobster
## 2
               BE nep-15
                            O NA NA Belgium Nephrops norvegicus Norway lobster
          NEP
               BE nep-33 299 NA NA Belgium Nephrops norvegicus Norway lobster
## 3
          NEP
               BE nep-5 146 NA NA Belgium Nephrops norvegicus Norway lobster
## 4
          NEP
                            O NA NA Belgium Nephrops norvegicus Norway lobster
## 5
          NEP
               BE nep-14
                            O NA NA Belgium Nephrops norvegicus Norway lobster
## 6
          NEP
               BE nep-6
     French name
##
## 1 Langoustine
## 2 Langoustine
## 3 Langoustine
## 4 Langoustine
## 5 Langoustine
## 6 Langoustine
DFM <- rbind.data.frame(DFM, NEP2)</pre>
MEDA <- merge(MED, ASFIS[,c(3,4,5,6)], by.x='Species', by.y='Scientific_name', all.x=TRUE)
MEDA <- tidyr::gather(MEDA, "Country", "n", 4:13)</pre>
MEDAG <- merge(MEDA, GEO, all.x=TRUE)</pre>
```

and a look at the Mediterranean dataset

```
head(MEDAG,3)
      Country
                        Species
                                                 Area RefYears
## 1 Bulgaria Alopias vulpinus All areas in the Med 2013-2015
## 2 Bulgaria Anguilla anguilla all areas in the Med 2013-2015
## 3 Bulgaria
                   Aphia minuta
                                 GSA 9,10,16 and 19 2013-2015
    Total.average.landings..t. X3A_CODE
                                             English name
                                                                 French name n
## 1
                            9.0
                                                  Thresher
                                     ALV
                                                                      Renard 0
## 2
                          308.0
                                     ELE
                                             European eel Anguille d'Europe 0
## 3
                           50.7
                                     FIM Transparent goby
                                                                      Nonnat 0
    geo
##
## 1 BG
## 2 BG
## 3 BG
MED <- data.frame(X3A_CODE=MEDAG$X3A_CODE, geo=MEDAG$geo, area=MEDAG$Area, Y1=round(MEDAG$n,0),
                   Y2=NA, Y3=NA, Country=MEDAG$Country, Scientific_name=MEDAG$Species, English_name=MED
                   French_name=NA)
DFM <- rbind.data.frame(DFM, MED)</pre>
```

Construction of the table

```
T1A <- data.frame()
for (i in 1:nrow(table1A)) {
    ctry2 <- GEO$Country[GEO$geo %in% CTRY]</pre>
   reg <- strsplit(as.character(table1A$areaBis[i]), split=',')</pre>
    if (table1A$region[i] %in% 'Mediterranean and Black Sea') {
       DT <- DFM[DFM$Scientific name %in% table1A$latinName[i] & tolower(DFM$area) %in% tolower(paste(
      } else {
        DT <- DFM[DFM$Scientific_name %in% table1A$latinName[i] & tolower(DFM$area) %in% tolower(reg[[1]
   DT$MOY <- apply(DT[,4:6],1,mean,na.rm=TRUE)
    #RFMO <- 'ICES'
    if (substring(table1A$region[i],1,3) %in% 'Med') RFMO <- 'GFCM'
   T1 <- data.frame(MS=CTRY, refYears=paste(min(refYears), '-', max(refYears), sep=""), spp=table1A$latinN
        RFMO=table1A$RFMO[i], area = table1A$area[i],select=NA, landings=NA, TAC=NA,shareLanding=NA,Thr
    ind <- which(DT$geo %in% CTRY)</pre>
    if (length(ind)>0) {
        T1$landings <- sum(DT$MOY[DT$geo %in% CTRY],na.rm=TRUE)
        T1$shareLanding <- T1$landings/sum(DT$MOY, na.rm=TRUE)
        } else {
        T1$landings <- 0
        T1$shareLanding <- 0
        }
    ## TAC
    ind.ct<-NULL
    if (!(table1A$FIDES_stockID[i] %in% 'No TAC')) {
        aa<-strsplit(as.character(table1A$FIDES_stockID[i]),split=',')[[1]]
        TACi <- TAC[TAC$StockID %in% aa,]
        if (length(aa)>1)
            TACi <- aggregate(list(Initial.Quantity = TACi$Initial.Quantity),
                by=list(Level.Code=TACi$Level.Code, Level.Description=TACi$Level.Description), sum)
```

```
ind.ct <- TACi$Initial.Quantity[which(TACi$Level.Description %in% ctry2)]
        ind.eu <- TACi$Initial.Quantity[which(TACi$Level.Code %in% 'EEC')]</pre>
        if (length(ind.ct) == 1) T1$TAC <- ind.ct/ind.eu</pre>
        T1$Comments<-NA
        TT <- tapply(TACi$Initial.Quantity, TACi$Level.Description,sum,na.rm=TRUE)/TACi$Initial.Quantit
        TT <- TT[names(TT) %in% GEO$Country] #Keep only the EU countries to calculate the 25% rule
        if (!(is.na(T1$TAC)) & T1$TAC <0.1 & T1$TAC>0) T1$Comments <- sum(TT[which(TT<0.1)])
        if (!(is.na(T1$Comments)) & T1$Comments >=.25) {
            print(T1)
            print(TT[TT<.1])</pre>
            cat('\n')
        }
   }
    #Add-on Sept 2019 Joel
    if (length(ind.ct)>0) {
      if (!is.na(ind.ct)) {
      T1$Comments2 <- paste('FIDES Initial.Quantity =',ind.ct)
   }}
    ##
   T1A <- rbind.data.frame(T1A, T1)
    T1A$Thresh <- as.character(T1A$Thresh)
    #Threshold ruling
    # T1A$Thresh[T1A$TAC >=.1 & T1A$landings >=200] <- 'M' #rule (a) & (c)
    # T1A$Thresh[is.na(T1A$TAC) & T1A$shareLanding >=.1 & T1A$landings >=200] <- 'M' #rule (b) & (c)
    # T1A$Thresh[T1A$TAC <.1 & T1A$Comments >=.25] <- 'C' # 25% rule, sampling to be coordinated betw
    #Threshold ruling specified like the EU Reg
    T1A$Thresh[T1A$TAC <.1] <- 'Y' #rule (a)
   T1A$Thresh[is.na(T1A$TAC) & T1A$shareLanding <.1] <- 'Y' #rule (b)
   T1A$Thresh[T1A$landings < 200] <- 'Y' #rule (c)
    T1A$Thresh[T1A$TAC <.1 & T1A$Comments >=.25] <- 'N'
                                                         # 25% rule, sampling to be coordinated betwee
}
    MS refYears
                                spp
                                        region RFMO area select landings
## 1 DK 2016-2018 Sprattus sprattus Baltic Sea ICES 22-32
##
            TAC shareLanding Thresh Comments Comments2
                                  N 0.2628188
## 1 0.09864283
                  0.08089579
                 Finland
      Denmark
                            Germany Lithuania
## 0.09864283 0.05163738 0.06249476 0.05004384
##
    MS refYears
                                                          region RFMO area
                                spp
## 1 DK 2016-2018 Macrourus berglax North Sea and Eastern Arctic ICES
     select landings TAC shareLanding Thresh Comments Comments2
                   0 0.1
## 1
                                    0
                                           N
##
          Denmark
                         Germany United Kingdom
##
              0.1
                             0.1
                                            0.1
##
    MS refYears
                                  spp
                                                             region RFMO area
## 1 DK 2016-2018 Trachurus trachurus North Sea and Eastern Arctic ICES IIa
                            TAC shareLanding Thresh Comments Comments2
##
     select landings
                   0 0.09913542
                                                  N 0.3265306
## 1
##
                         Denmark
                                         France
                                                                        Latvia
          Belgium
                                                        Germany
##
     0.00000000
                     0.099135418
                                    0.039810998
                                                    0.077349955
                                                                   0.00000000
##
       Lithuania
                        Portugal
                                         Sweden United Kingdom
##
      0.00000000
                     0.010163869
                                                   0.093284407
                                    0.006785966
```

```
##
                                               region RFMO
##
    MS refYears
                                  spp
## 1 DK 2016-2018 Trachurus trachurus North Atlantic ICES
                                   area select landings
                                                                TAC
## 1 IIa, IVa, Vb, VIa, VIIa-c, e-k, VIIIabde
                                             NA 6839.608 0.09913542
     shareLanding Thresh Comments Comments2
##
                       N 0.3265306
## 1
       0.05597712
##
          Belgium
                         Denmark
                                          France
                                                        Germany
                                                                        Latvia
##
      0.000000000
                     0.099135418
                                    0.039810998
                                                    0.077349955
                                                                   0.00000000
##
        Lithuania
                        Portugal
                                          Sweden United Kingdom
      0.00000000
                     0.010163869
                                    0.006785966
                                                    0.093284407
##Formatting
T1B <- T1A
T1B$landings <- round(T1B$landings,0)
T1B$landings[T1B$landings == 0] <- '-'
T1B$TAC <- paste(round(100*T1B$TAC,0),'%',sep='')
T1B$TAC[T1B$TAC %in% c('NA%','NaN%','Inf%')] <- '-'
T1B$shareLanding <- paste(round(100*T1B$shareLanding,0),'%',sep='')
T1B$shareLanding[T1B$shareLanding %in% c('NA%','NaN%','Inf%')] <- '-'
T1B$Thresh[T1B$landings %in% '-' & T1B$TAC %in% '-'] <- T1B$shareLanding[T1B$landings %in% '-' & T1B$TA
ind <- which(T1B$Comments>.25)
T1B$Comments <- paste(round(100*T1B$Comments,0),'%',sep='')
T1B$Comments[T1B$Comments<.25] <- '-'
T1B$Comments[T1B$Comments %in% c('NA%','NaN%','Inf%')] <- '-'
T1B$Comments[ind] <- paste('Sum of MS shares <10% = ',T1B$Comments[ind],'%',sep='')
T1B[!ind]<-'-'
T1B$select <- '-'
T1B$select[T1B$Thresh %in% c('N')] <- 'Y'
T1B$select[T1B$Thresh %in% 'Y'] <- 'N'
T1B[T1B$RFMO %in% c('ICCAT', 'IOTC', 'WCPFC') & T1B$landings>0,c('select', 'Thresh')] <- c('Y', 'N')
T1B[T1B$spp %in% 'Anguilla anguilla' & T1B$landings>0,c('select','Thresh')] <- c('Yes','No')
T1B[T1B$spp %in% 'Nephrops norvegicus' & !(grepl('TAC', T1B$area)),'TAC'] <- '-'
levels(T1B$refYears) <- c(levels(T1B$refYears), '2015')</pre>
T1B[T1B$spp %in% 'Nephrops norvegicus' & !(grepl('TAC', T1B$area)), 'refYears'] <- '2015'
T1B[T1B$RFMO %in% 'GFCM', 'refYears'] <- '2015'
```

Export of Table 1A

the rule sum of quotas for coutries <10% (less or more than 25%) is noted in the comments column

```
ind <- order(as.character(T1B$region), as.character(T1B$RFMO), as.character(T1B$spp), as.character(T1B$
write.table(T1B[ind,], file=paste(output_path, CTRY,'_table1A_filled_common.csv',sep=';',row.n</pre>
```