

EU Farm Fish CEA - 1. Consumption Prep File

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This file brings together estimates of per country EU consumption of Sea bream, Sea bass, Carp, Salmon and Rainbow Trout into a single dataset in long format.

Preparation

Clear environment

```
rm(list=ls())
```

Load packages

```
library(readxl)
library(tidyverse)
```

```
-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
v dplyr      1.1.2      v readr      2.1.4
v forcats    1.0.0      v stringr    1.5.0
v ggplot2    3.4.2      v tibble     3.2.1
v lubridate  1.9.2      v tidyr      1.3.0
v purrr      1.0.1
```

```
-- Conflicts ----- tidyverse_conflicts() --
```

```
x dplyr::filter() masks stats::filter()
```

```
x dplyr::lag()     masks stats::lag()
```

```
i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become
```

```
library(scales)
```

Attaching package: 'scales'

The following object is masked from 'package:purrr':

discard

The following object is masked from 'package:readr':

col_factor

```
library(DT)
```

Session Info

```
sessionInfo()
```

R version 4.3.0 (2023-04-21)

Platform: aarch64-apple-darwin20 (64-bit)

Running under: macOS 14.3

Matrix products: default

BLAS: /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/lib/libRblas.0.dylib

LAPACK: /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/lib/libRlapack.dylib;

locale:

[1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8

time zone: Europe/London

tzcode source: internal

attached base packages:

[1] stats graphics grDevices utils datasets methods base

other attached packages:

[1] DT_0.28 scales_1.2.1 lubridate_1.9.2 forcats_1.0.0

[5] stringr_1.5.0 dplyr_1.1.2 purrr_1.0.1 readr_2.1.4

```
[9] tidyr_1.3.0      tibble_3.2.1      ggplot2_3.4.2      tidyverse_2.0.0
[13] readxl_1.4.2
```

loaded via a namespace (and not attached):

```
[1] gtable_0.3.3      jsonlite_1.8.4      compiler_4.3.0      tidyselect_1.2.0
[5] yaml_2.3.7        fastmap_1.1.1       R6_2.5.1            generics_0.1.3
[9] knitr_1.43        htmlwidgets_1.6.2   munsell_0.5.0       pillar_1.9.0
[13] tzdb_0.4.0        rlang_1.1.1         utf8_1.2.3          stringi_1.7.12
[17] xfun_0.39         timechange_0.2.0    cli_3.6.1           withr_2.5.0
[21] magrittr_2.0.3    digest_0.6.31       grid_4.3.0          rstudioapi_0.14
[25] hms_1.1.3         lifecycle_1.0.3     vctrs_0.6.2         evaluate_0.21
[29] glue_1.6.2        cellranger_1.1.0    fansi_1.0.4         colorspace_2.1-0
[33] rmarkdown_2.22    tools_4.3.0         pkgconfig_2.0.3     htmltools_0.5.5
```

Open data files

This file contains data on:

- EU country codes (from [Eurostat](#))
- EUFOMA apparent consumption estimates (live weight equivalent) in various EU countries of [Seabass \(2016\)](#), [Seabream \(2019\)](#) and [Carp\(2018\)](#)
- Portion trout and large trout estimates come from the EUFOMA [large trout in the EU \(2020\)](#) study, using apparent consumption estimates for large trout and all trout respectively
- Salmon consumption estimates are extracted from the chart from page 42 of the [Mowi \(2023\) industry report](#) for France, Germany, Italy, Spain and Sweden only.
- Population data for EU27 countries to estimate salmon consumption (from Eurostat)

```
xl_data <- "../1_input_data/EUFOMA_consumption_data.xlsx"
country_codes <- read_excel(xl_data,sheet = "country_codes")
bass <- read_excel(xl_data,sheet = "seabass_2016")
breame <- read_excel(xl_data,sheet = "seabream_2019")
carp <- read_excel(xl_data,sheet = "carp_2018")
trout <- read_excel(xl_data,sheet = "trout_2020")
salmon <- read_excel(xl_data,sheet = "salmon_2022_raw")
population <- read_excel("../1_input_data/country populations.xlsx",sheet = "EU27population")
```

Salmon consumption estimates

Mowi industry report only provides salmon consumption estimates for 5 EU countries. For other countries, we need estimate salmon consumption. I do this by assuming:

- per capita consumption in the Denmark and Finland is equal to per capita consumption in Sweden
- per capita consumption in Netherlands, Belgium, Austria, Luxembourg and Ireland is equal to per capita consumption in Germany
- per capita consumption in Portugal, Greece, Cyprus and Malta is equal to per capita consumption in Italy
- the remainder is equally distributed across the rest of the EU

I also assume that the Mowi industry reports estimate for EU+UK market size actually excludes the UK, based on comparisons with EUFOMA balance sheet data (<https://www.eumofa.eu/supply-balance>).

I first extract aggregate EU consumption of salmon...

```
EU_salmon <- salmon %>% filter(Country=="EU27") %>% pull(Tons) %>% mean()  
EU_salmon
```

```
[1] 1141000
```

I then estimate per capita consumption of salmon (in Tons) for the three reference countries, Sweden, Italy and Germany.

```
per_capita_salmon <-  
  salmon %>%  
  filter(Country %in% c("Sweden", "Germany", "Italy")) %>%  
  left_join(population, by="Country") %>%  
  mutate(per_capita=Tons/pop) %>%  
  select(Country, per_capita) %>%  
  rename(ref_country=Country)  
  
per_capita_salmon
```

```
# A tibble: 3 x 2  
  ref_country per_capita  
  <chr>         <dbl>
```

1 Germany	0.00285
2 Italy	0.00235
3 Sweden	0.00555

I then assign these per capita consumption values to selected other countries, and calculate aggregate consumption in those countries.

```
#Assign country groups
nordic <- c("Denmark","Finland")
germanic <- c("Austria","Netherlands","Belgium","Luxembourg","Ireland")
italian <- c("Portugal","Greece","Malta","Cyprus")

#Estimate salmon consumption in these countries
salmon_ref_countries <- population %>%
  filter(Country %in% c(nordic,germanic,italian)) %>%
  mutate(
    ref_country=
      case_when(
        Country %in% nordic ~ "Sweden",
        Country %in% germanic ~ "Germany",
        Country %in% italian ~ "Italy",
        TRUE ~ "Error"
      )
  ) %>%
  left_join(per_capita_salmon,by="ref_country") %>%
  mutate(
    Tons=pop*per_capita,
    Estimate=str_c("ref country per capita - ",str_to_lower(ref_country))) %>%
  select(Country,Tons,Estimate)

salmon_ref_countries
```

A tibble: 11 x 3

	Country	Tons	Estimate
	<chr>	<dbl>	<chr>
1	Belgium	33079.	ref country per capita - germany
2	Denmark	32592.	ref country per capita - sweden
3	Ireland	14407.	ref country per capita - germany
4	Greece	24630.	ref country per capita - italy
5	Cyprus	2130.	ref country per capita - italy
6	Luxembourg	1838.	ref country per capita - germany
7	Malta	1227.	ref country per capita - italy
8	Netherlands	50086.	ref country per capita - germany

```

9 Austria      25566. ref country per capita - germany
10 Portugal    24376. ref country per capita - italy
11 Finland     30787. ref country per capita - sweden

```

I then estimate consumption in the remainder of EU countries, by assuming per capita consumption in these countries is equal.

```

#Estimate residual aggregate salmon consumption
salmon_residual_aggregate<- EU_salmon*2-sum(salmon$Tons)-sum(salmon_ref_countries$Tons)

#Calculate consumption in each country by multiplying aggregate by their population share
salmon_residual_countries <- population %>%
  filter(!(Country %in% c(salmon$Country,germanic,italian,nordic))) %>%
  mutate(
    Tons=pop/sum(pop)*salmon_residual_aggregate,
    Estimate="residual EU per capita"
  ) %>%
  select(-pop)

salmon_residual_countries

```

```

# A tibble: 11 x 3
  Country      Tons Estimate
  <chr>      <dbl> <chr>
1 Bulgaria   4549. residual EU per capita
2 Czechia    6995. residual EU per capita
3 Estonia     886. residual EU per capita
4 Croatia    2569. residual EU per capita
5 Latvia     1248. residual EU per capita
6 Lithuania  1866. residual EU per capita
7 Hungary    6444. residual EU per capita
8 Poland     25045. residual EU per capita
9 Romania    12666. residual EU per capita
10 Slovenia   1402. residual EU per capita
11 Slovakia   3615. residual EU per capita

```

Finally I combine the data into final table

```

#Combined table
salmon<-salmon %>%
  filter(Country!="EU27") %>%

```

```
mutate(Estimate="mowi industry report") %>%
rbind(salmon_ref_countries,salmon_residual_countries)

salmon
```

```
# A tibble: 27 x 3
  Country      Tons Estimate
  <chr>      <dbl> <chr>
1 France  269000 mowi industry report
2 Germany 237000 mowi industry report
3 Italy   139000 mowi industry report
4 Spain   130000 mowi industry report
5 Sweden   58000 mowi industry report
6 Belgium 33079. ref country per capita - germany
7 Denmark 32592. ref country per capita - sweden
8 Ireland 14407. ref country per capita - germany
9 Greece  24630. ref country per capita - italy
10 Cyprus  2130. ref country per capita - italy
# i 17 more rows
```

And check that the values look reasonable (TRUE means working ok).

```
#Check
near(EU_salmon,sum(salmon$Tons))
```

```
[1] TRUE
```

Finally I plot some charts for reference.

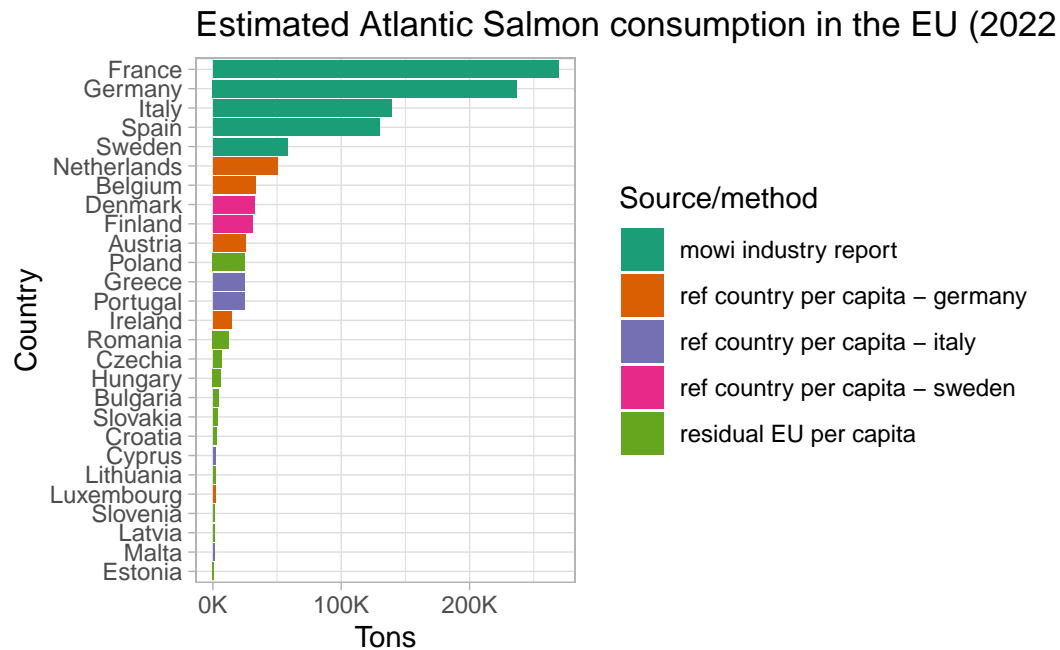
Estimated salmon consumption in each EU country

```
salmon %>%
  ggplot(
    aes(
      x=Tons,
      y=reorder(Country,Tons),
      fill=Estimate
    ) +
    geom_col()+
    labs(
      title = "Estimated Atlantic Salmon consumption in the EU (2022)",
```

```

y="Country",
fill="Source/method") +
scale_fill_brewer(palette="Dark2") +
theme_light() +
scale_x_continuous(labels = label_number(suffix = "K", scale = 1e-3))

```



Per capita Atlantic salmon production

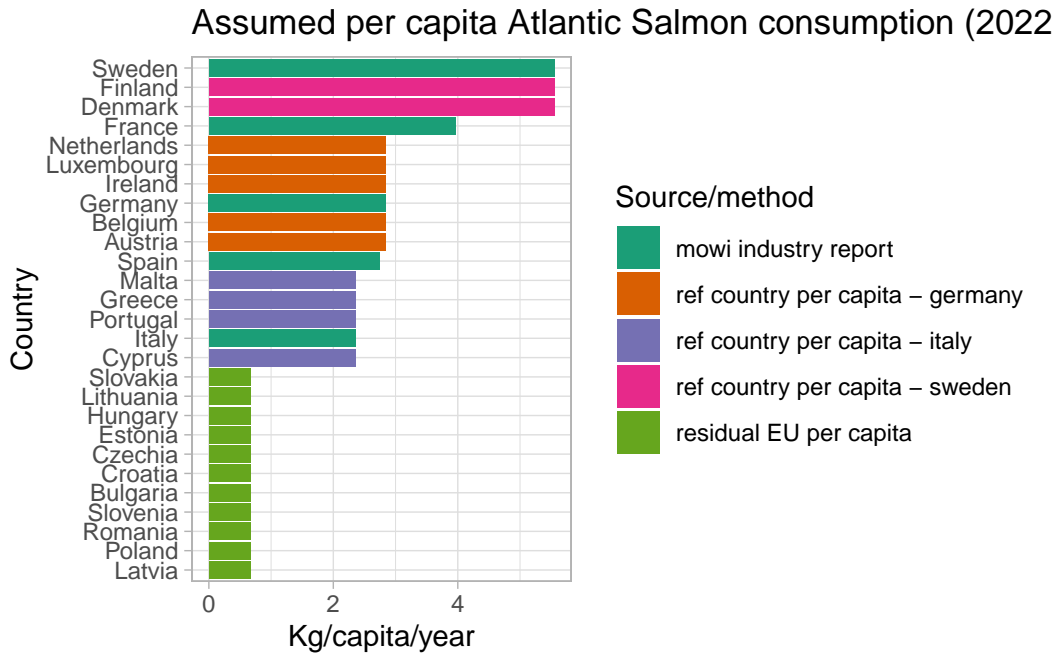
```

salmon %>%
left_join(population,by="Country") %>%
mutate(per_capita=Tons*1000/pop) %>%
ggplot(
aes(
x=per_capita,
y=reorder(Country,per_capita),
fill=Estimate
)) +
geom_col()+
labs(
title = "Assumed per capita Atlantic Salmon consumption (2022)",
y="Country",
x="Kg/capita/year",

```



```
fill="Source/method") +
scale_fill_brewer(palette="Dark2") +
theme_light()
```



Combine data into single file

We now prepare/clean the country species consumption data and assemble to into a “long” dataframe. Steps involved are:

1. linking country code and country code data
2. adding species names
3. apportioning consumption by portion trout and large trout share (iv) combining the species data into a single data frame
4. making column names lower case for ease of matching

```
# Add country codes and species name
breame <- left_join(breame,country_codes,by="Code") %>% mutate(Species="Sea Bream")
bass <- left_join(bass,country_codes,by="Country") %>% mutate(Species="Sea Bass")
carp <- left_join(carp,country_codes,by="Code") %>% mutate(Species="Carp")
```

```

salmon<- left_join(salmon,country_codes,by="Country") %>% mutate(Species="Atlantic Salmon")

# Add country codes and fish size and separate into two observations
trout<- trout %>%
  left_join(country_codes,by="Code") %>%
  pivot_longer(
    cols=c("Rainbow Trout (Small)","Rainbow Trout (Large)"),
    names_to="Species",values_to="Tons"
  ) %>%
  select(-Notes,-Total)

# combine into single dataframe
cons_data <- bind_rows(bream,bass,carp,trout) %>%
  mutate(Estimate="EUFOMA") %>%
  bind_rows(salmon)

# make column names lower case and reorder
names(cons_data) <- str_to_lower(names(cons_data))
cons_data <- relocate(cons_data, species, country, code, tons)

```

We do a simple plot of the data to make sure it looks reasonable.

```

cons_data <- cons_data %>%
  group_by(country) %>%
  mutate(total_cons=sum(tons)) %>%
  ungroup()

cons_data %>%
  ggplot(
    aes(
      x=tons,
      y=reorder(country,total_cons),
      fill=species
    ) +
  geom_col() +
  labs(
    title = "Annual consumption of salmon, carp, trout and seabream/bass in EU",
    x = "Metric tons (live weight equivalent)",
    y = "",
    fill = "Species"
  )

```

```

) +
scale_x_continuous(labels = label_number(suffix = "K", scale = 1e-3)) +
scale_fill_brewer(palette = "Dark2") +
theme_light() +
theme(legend.position = "top")

```

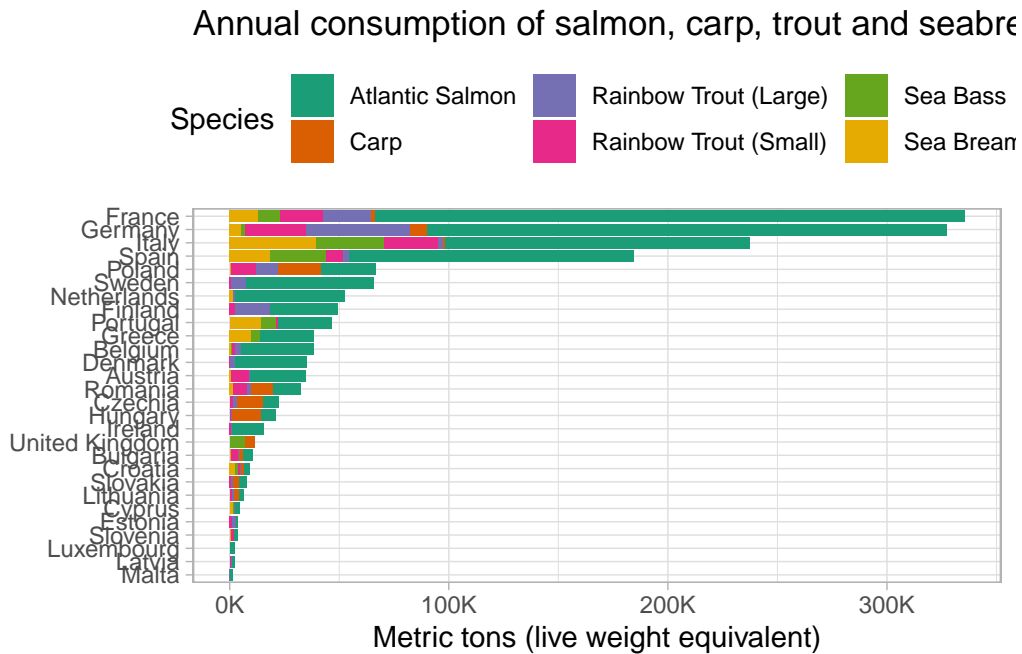


Figure 1: Annual consumption of selected species in the EU (kg)

Save output file

```

saveRDS(cons_data,file= "../3_intermediate_data/cons_data.rds")

```