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Probabilistic estimates of Daily Intake

Using LitStream data

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This R code is only for BCEP

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Latest modification: 2/27/2024

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Updated the FUE values

#####

rm(list=ls(all=TRUE))

library(dplyr)

library(tidyr)

library(openxlsx)

library("writexl")

library("readxl")

library(basictabler)

library(psych)

library(goft)

```
library(ggplot2)

#install.packages("goft")

#install.packages("EnvStats")

library(EnvStats);

#####

## Location for the results

path_results = "D:\\UC-OneDrive\\OneDrive - University of Cincinnati\\Desktop\\F\\Dr. Lynne -
biomonitoring task\\Results";

#path_results = "C:\\Users\\hsuwe\\OneDrive - University of Cincinnati\\Desktop\\F\\Dr. Lynne -
biomonitoring task\\Results";

## As the results in the file named:

exceldata_name = "\\4. LitStream_Distributional_data_results.xlsx";

#####

#### FUE parameters ####

#####

# Old values

# c(best estimate, high, low)

#FUE_BCEP = c(0.2, 0.41, 0.04)

#FUE_BCIPP = c(0.39, 0.78, 0.08)

#FUE_BDCIPP = c(0.42, 0.5, 0.08)

#FUE_BCIPP_BCIPHIPP = c(0.78, 0.94, 0.16)

## New values as of 2/24/2024

FUE_BCEP = c(0.13, 0.27, 0.03)
```

```
FUE_BCIPP = c(0.23, 0.47, 0.05)
FUE_BDCIPP = c(0.23, 0.28, 0.05)
FUE_BCIPP_BCIPHIPP = c(0.58, 0.90, 0.12)
```

```
#####
```

```
#### UFRBW parameters ####
```

```
#####
```

```
# From NHANES 2017-2018 data
```

```
# NOTE: the position of the value is important
```

```
groups = c("3-5 yrs", "6-11 yrs", "12-17 yrs", "18+ yrs")
```

```
UFRBW_GM = c(0.063917438, 0.024195805, 0.016092702, 0.01293934)
```

```
GSD = c(2.639638606, 2.241928384, 2.129355164, 2.072937724)
```

```
parameters_UFRBW = as.data.frame(cbind(groups, UFRBW_GM, GSD) )
```

```
parameters_UFRBW$UFRBW_GM = as.numeric(parameters_UFRBW$UFRBW_GM)
```

```
parameters_UFRBW$GSD = as.numeric(parameters_UFRBW$GSD)
```

```
#####
```

```
## Estimated Chemicals from LitStream ##
```

```
## By age groups          ##
```

```
#####
```

```
# NOTE: the position of value should be in this order : c("6-11 years", "20-39 years")
```

```
#####
```

```
#### Use the following concentration levels from the studies for each age group ####
```

```
## Study: "379-Percy et al-2022", "379-Percy et al-2022" and "382-Yang et al-2023"
```

```
age_groups =c("3 years", "5 years", "20-39 years");
```

```
GM_conc_BCEP = c(1 , 0.74, 0.51)
```

```
GSD_conc_BCEP = c(4.6, 3.64, 4.33)
```

```
#####  
####
```

```
#install.packages("EnvStats")
```

```
library(EnvStats); # a package for generating random values from a triangular distribution
```

```
#####
```

```
### By Chemicals    ###
```

```
#####
```

```
# a random sample of 1,000
```

```
n = 1000
```

```
## BCEP
```

```
BCEP_matrix=matrix(,1000,3); ## for 3 age-groups
```

```
yr=0;
```

```
for (agrp in c(1,1,4)){ ## select the first (3-5), 1st (3-5) and 4th (18+) age groups from NHANES 2017-18  
cycle
```

```
yr=yr+1;
```

```
FUE_BCEP_rand = rtri(n, min=FUE_BCEP[[3]], max=FUE_BCEP[[2]], mode=FUE_BCEP[[1]])
```

```
UFRBW_BCEP_rand = rlnorm(n, meanlog = log(parameters_UFRBW$UFRBW_GM[[agrp]]), sdlog =  
log(parameters_UFRBW$GSD[[agrp]]))
```

```
GM_conc_BCEP_rand = rlnorm(n, meanlog = log(GM_conc_BCEP[[yr]]), sdlog =  
log(GSD_conc_BCEP[[yr]]))
```

```
DI_BCEP = (GM_conc_BCEP_rand * UFRBW_BCEP_rand) / FUE_BCEP_rand; ## pay attention to the unit
```

```
BCEP_matrix[,yr]=DI_BCEP
```

```
}
```

```
BCEP_matrix = as.data.frame(BCEP_matrix)
```

```
names(BCEP_matrix)= age_groups
```

```
BCEP_summary = BCEP_matrix %>% summarise(
```

```
  mean_grp1 = (mean(BCEP_matrix[[1]], na.rm = TRUE)),
```

```
  mean_grp2 = (mean(BCEP_matrix[[2]], na.rm = TRUE)),
```

```
  mean_grp3 = (mean(BCEP_matrix[[3]], na.rm = TRUE)),
```

```
  SD_grp1 = (sd(BCEP_matrix[[1]], na.rm = TRUE)),
```

```
  SD_grp2 = (sd(BCEP_matrix[[2]], na.rm = TRUE)),
```

```
  SD_grp3 = (sd(BCEP_matrix[[3]], na.rm = TRUE)),
```

```
  Gmean_grp1 = exp(mean(log(BCEP_matrix[[1]]), na.rm = TRUE)),
```

Gmean_grp2 = exp(mean(log(BCEP_matrix[[2]]), na.rm = TRUE)),

Gmean_grp3 = exp(mean(log(BCEP_matrix[[3]]), na.rm = TRUE)),

GSD_grp1 = exp(sd(log(BCEP_matrix[[1]]), na.rm = TRUE)),

GSD_grp2 = exp(sd(log(BCEP_matrix[[2]]), na.rm = TRUE)),

GSD_grp3 = exp(sd(log(BCEP_matrix[[3]]), na.rm = TRUE)),

p25_grp1 = quantile(BCEP_matrix[[1]],0.25),

p25_grp2 = quantile(BCEP_matrix[[2]],0.25),

p25_grp3 = quantile(BCEP_matrix[[3]],0.25),

p50_grp1 = quantile(BCEP_matrix[[1]],0.5),

p50_grp2 = quantile(BCEP_matrix[[2]],0.5),

p50_grp3 = quantile(BCEP_matrix[[3]],0.5),

p75_grp1 = quantile(BCEP_matrix[[1]],0.75),

p75_grp2 = quantile(BCEP_matrix[[2]],0.75),

p75_grp3 = quantile(BCEP_matrix[[3]],0.75),

p90_grp1 = quantile(BCEP_matrix[[1]],0.9),

p90_grp2 = quantile(BCEP_matrix[[2]],0.9),

p90_grp3 = quantile(BCEP_matrix[[3]],0.9),

p95_grp1 = quantile(BCEP_matrix[[1]],0.95),

p95_grp2 = quantile(BCEP_matrix[[2]],0.95),

p95_grp3 = quantile(BCEP_matrix[[3]],0.95)

)

```
names(BCEP_summary) =
c(paste0("Mean_",age_groups),paste0("SD_",age_groups),paste0("GM_",age_groups),paste0("GSD_",age_
e_groups),
      paste0("p25_",age_groups),
paste0("p50_",age_groups),paste0("p75_",age_groups),paste0("p90_",age_groups),paste0("p95_",age_
groups))
```

```
#####
#### Save the summary results #####
#####
```

```
study = c("379-Percy et al-2022", "379-Percy et al-2022", "382-Yang et al-2023")
Age_grp = c("3 years", "5 years", "20-39 years")
```

```
Mean = c(BCEP_summary[1:3])
SD = c(BCEP_summary[4:6])
GM = c(BCEP_summary[7:9])
GSD = c(BCEP_summary[10:12])
p25 = c(BCEP_summary[13:15])
p50 = c(BCEP_summary[16:18])
p75 = c(BCEP_summary[19:21])
p90 = c(BCEP_summary[22:24])
p95 = c(BCEP_summary[25:27])
```

```
NH_result_matrix = as.data.frame(cbind(study ,Age_grp,Mean, SD, GM, GSD, p25, p50, p75, p90, p95))
```

```
names(NH_result_matrix) = c("Study", "Age Group", "Arithmetic mean", "Arithmetic SD", "Geometric Mean", "Geometric SD", "25th percentile", "50th percentile", "75th percentile", "90th percentile", "95th percentile");
```

```
NH_result_matrix
```

```
#####
```

```
### Save results as a worksheet in the Excel file ###
```

```
#####
```

```
Readme="Readme";
```

```
write.xlsx(Readme, file=paste0(path_results,exceldata_name),
```

```
  sheetName = "Readme", colNames = TRUE, rowNames = F, append = T)
```

```
##### BCEP #####
```

```
wb <- loadWorkbook(paste0(path_results,exceldata_name));
```

```
addWorksheet(wb,"Simulated_BCEP")
```

```
writeData(wb, sheet = "Simulated_BCEP", x =BCEP_matrix, borders = "columns", startCol = 1, startRow = 1)
```

```
### graphs
```

```
BCEP_data_long <- gather(BCEP_matrix, age_groups, BCEP, age_groups, factor_key=TRUE)
```

```
g2.s = ggplot(BCEP_data_long, aes(x=BCEP, color=age_groups, fill=age_groups)) +
```

```
geom_histogram(alpha=0.6, show.legend = FALSE) +
```

```
  scale_x_continuous(name = "BCEP level") +
```

```
  scale_y_continuous(name = "Frequency")+
```

```
facet_wrap(~ age_groups, scale="free", ncol=2) +
```



```

theme(
  axis.text.y = element_text(size=10),
  axis.title.y = element_text(size=10),
  axis.text.x = element_text(size=10),
  axis.title = element_text( size = 10, face = "bold" ),
  strip.text = element_text(size = 10),
  axis.title.x = element_text(size=10)
)
#g2.s

plot(g2.s)
insertPlot(wb, "Simulated_BCEP", startCol = 6, startRow = 2)

### log-normal distribution test
BCEP_Inorm_test_results1 = Inorm_test(BCEP_matrix[["3 years"]])
BCEP_Inorm_test_results2 = Inorm_test(BCEP_matrix[["5 years"]])
BCEP_Inorm_test_results3 = Inorm_test(BCEP_matrix[["20-39 years"]])

BCEP_Inorm = rbind(

cbind(BCEP_Inorm_test_results1$method,BCEP_Inorm_test_results1$data.name,BCEP_Inorm_test_results1$p.value),

cbind(BCEP_Inorm_test_results2$method,BCEP_Inorm_test_results2$data.name,BCEP_Inorm_test_results2$p.value),

cbind(BCEP_Inorm_test_results3$method,BCEP_Inorm_test_results3$data.name,BCEP_Inorm_test_results3$p.value)
)

colnames(BCEP_Inorm) = c("Method:", "Data:", "p-value=")

```

```
BCEP_Inorm=transform(BCEP_Inorm)
```

```
BCEP_Inorm[[3]]=as.numeric(BCEP_Inorm[[3]])
```

```
summary_stat_BCEP= rbind(describe(BCEP_matrix[["3 years"]]),describe(BCEP_matrix[["5  
years"]]),describe(BCEP_matrix[["20-39 years"]]))
```

```
summary_stat_BCEP= summary_stat_BCEP[,-1]
```

```
rownames(summary_stat_BCEP) = age_groups;
```

```
writeData(wb, sheet = "Simulated_BCEP", x =summary_stat_BCEP, rowNames = T, startCol = 14, startRow  
= 2)
```

```
writeData(wb, sheet = "Simulated_BCEP", x =BCEP_Inorm , startCol = 6, startRow = 22)
```

```
sheet_name = "BCEP_Estimates"
```

```
addWorksheet(wb,paste0(sheet_name))
```

```
writeData(wb, sheet = paste0(sheet_name), x ="Table. Daily intake - Estimated parameters based on  
probabilistic models using LitStream data", borders = "columns", startCol = 1, startRow = 1)
```

```
## Table Caption
```

```
writeData(wb, sheet = sheet_name, x = "Unit: ug/kg-day", borders = "columns", startCol = 1, startRow =  
2)
```

```
row_headers = paste0(rownames(NH_result_matrix))
```

```
tbl <- BasicTable$new()
```

```
tbl$addData(NH_result_matrix, firstColumnAsRowHeaders=T,
```

```
explicitColumnHeaders=c("Study", "Age Group", "Arithmetic mean", "Arithmetic SD", "Geometric Mean", "Geometric SD", "25th percentile", "50th percentile", "75th percentile", "90th percentile", "95th percentile");
```

```
# the arguments are (rFrom, cFrom, rTo, cTo, declarations)
```

```
tbl$setStyling(2, 3, 16, 11, declarations=list("xl-value-format"="##0.0000"))
```

```
tbl$writeToExcelWorksheet(wb=wb, wsName=sheet_name,
```

```
topRowNumber=4, leftMostColumnNumber=1,
```

```
applyStyles=T, mapStylesFromCSS=F, outputValuesAs="rawValue")
```

```
saveWorkbook(wb, file=paste0(path_results,exceldata_name), overwrite = TRUE);
```

```
#####  
#####
```

```
#####
```

```
##### THE END #####
```

```
#####
```