Measure 1: Savings potentials, Normalization, Proportionalization and Creation of Plots and Tables

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Code for the calculation of the savings potential using the results of the LCA-studies. Normalization and Proportionalization is performed as well as the creation of plots and tables.

To execute, change:

- Login data for Nextcloud
- Adjust download location
- Adjust upload location
- Adjust URLs for uploading

Load Packages:

```
# install.packages("httr")
# install.packages("readxl")
# install.packages("dplyr")
# install.packages("writexl")
# install.packages("gaplot2")
# install.packages("tidyr")
# install.packages("openxlsx")
# install.packages("flextable")
# install.packages("officer")
library(readx1)
library(httr)
library(dplyr)
library(writexl)
library(ggplot2)
library(tidyr)
library(openxlsx)
library(flextable)
library(officer)
```

Parameters

Parameters to be adjusted:

```
measure <- "01" # "01", "02", "03" or "04" sheet <- 2 # 2,3,4 or 5
```

```
# name for columns
name_with_TP <- "digital TP"</pre>
name_with_BT <- "digital BT"</pre>
name_with_MT <- "digital MT"</pre>
name without TP <- "analogue TP"
name_without_BT <- "analogue BT"</pre>
name without MT <- "analogue MT"</pre>
name_SP_TP <- "SP TP"</pre>
name SP BT <- "SP BT"
name SP MT <- "SP MT"
name_SP_total <- "SP total"</pre>
name with TP norm <- "norm. digital TP"
name_with_BT_norm <- "norm. digital BT"</pre>
name_with_MT_norm <- "norm. digital MT"</pre>
name_without_TP_norm <- "norm. analogue TP"</pre>
name_without_BT_norm <- "norm. analogue BT"</pre>
name without MT norm <- "norm. analogue MT"
name_SP_TP_norm <- "norm. SP TP"</pre>
name SP BT norm <- "norm. SP BT"
name_SP_MT_norm <- "norm. SP MT"</pre>
name_SP_total_norm <- "norm. SP total"</pre>
name_with_TP_prop <- "prop. digital TP"</pre>
name_with_BT_prop <- "prop. digital BT"</pre>
name_with_MT_prop <- "prop. digital MT"</pre>
name_without_TP_prop <- "prop. analogue TP"</pre>
name_without_BT_prop <- "prop. analogue BT"</pre>
name without MT prop <- "prop. analogue MT"
name_SP_TP_prop <- "prop. SP TP"</pre>
name_SP_BT_prop <- "prop. SP BT"</pre>
name_SP_MT_prop <- "prop. SP MT"</pre>
name SP total prop <- "prop. SP total"</pre>
# vector with excluded categories due to missing normalization factor
excluded_categories_nf <- c("Climate change: biogenic",
                               "Climate change: fossil",
                               "Climate change: land use and land use change",
                               "Ecotoxicity: freshwater, inorganics",
                               "Ecotoxicity: freshwater, organics",
                               "Human toxicity: carcinogenic, inorganics",
                               "Human toxicity: carcinogenic, organics",
                               "Human toxicity: non-carcinogenic, inorganics",
                               "Human toxicity: non-carcinogenic, organics")
# vector with excluded toxicity categories for Pareto rule
excluded_categories_pr <- c("Human toxicity: carcinogenic",
                               "Human toxicity: non-carcinogenic",
                               "Ecotoxicity: freshwater")
# storage location
location_download <- "C:/Users/Klene/Documents/Uni_Bremen/WS24_25/Masterarbei</pre>
```

```
t/R/Daten/"
location_upload <- "C:/Users/Klene/Documents/Uni_Bremen/WS24_25/Masterarbeit/
R/Results/"
# Login data for NextCloud
username <- "mkindler@uni-bremen.de" # username
password <- "xxxxxxxx" # password
# WebDAV-URL to file
nextcloud_url <- "https://nc.uni-bremen.de/remote.php/dav/files/mkindler%40un
i-bremen.de/Masterarbeit/Masterarbeit_II/R/Daten/Ergebnisse_LCA_basic.xlsx"
nextcloud_url_2 <- "https://nc.uni-bremen.de/remote.php/dav/files/mkindler%40
uni-bremen.de/Masterarbeit/Masterarbeit_II/R/Results/"</pre>
```

Data preparation

Load data from NextCloud:

```
# download file
response <- GET(nextcloud_url, authenticate(username, password, type = "basic")</pre>
"))
# Ckeck successful downloading
if (status_code(response) == 200) {
  # Save file
  writeBin(content(response, "raw"), paste0(location download, "Ergebnisse LC
A basic.xlsx"))
  cat("The file was successfully downloaded and saved locally.\n")
} else {
  cat("Error downloading the file. Status code: ", status code(response), "\n
")
}
## The file was successfully downloaded and saved locally.
# Load results data from Excel file
results <- read_excel(pasteO(location_download, "Ergebnisse_LCA_basic.xlsx"),
sheet = sheet)
## New names:
## • `Wirkungskategorien` -> `Wirkungskategorien...1`
## • `Einheit` -> `Einheit...2`
## • `Wirkungskategorien` -> `Wirkungskategorien...5`
## • `Einheit` -> `Einheit...6`
## • `Wirkungskategorien` -> `Wirkungskategorien...9`
## • `Einheit` -> `Einheit...10`
# Check data type: DataFrame
if (is.data.frame(results)) {
  results <- results
} else {
  results <- as.data.frame(results)
# Rename columns
results <- results %>%
```

```
rename(
    `impact category` = `Wirkungskategorien...1`,
    unit = `Einheit...2`
    ) %>%
  select(-"Wirkungskategorien...5", -"Einheit...6", -"Wirkungskategorien...9"
, -"Einheit...10")
# Load normalization factors from Excel file
NF <- read excel(paste0(location download, "Ergebnisse LCA basic.xlsx"), shee
t = 1
## New names:
## • `` -> `...2`
## • `` -> `...3`
# Check data type: DataFrame
if (is.data.frame(NF)) {
  NF <- NF
} else {
  NF <- as.data.frame(NF)</pre>
# Set column names right
NF <- NF[-1, ]
colnames(NF) <- as.character(NF[1, ])</pre>
NF <- NF[-1, ]
# Select relevant columns
NF <- NF %>%
  select(`impact category`,
         NF = `normalization factor\r\n(Matuštík 2024)`)
```

Calculations

Savings potential

Calculation of Savings Potentials:

Normalization

Calculation of normalized values (to the planetary boudaries (Matuštík 2024)):

```
# Join of results and NF according to the impact categories
results <- left_join(results, NF, by = c("impact category" = "impact category"))
# Convert all columns except the first two into numerical values
results <- results %>%
    mutate_at(vars(-1, -2), as.numeric)
```

```
# Remove categories without normalization factors
results_norm <- results %>%
  filter(!`impact category` %in% excluded_categories_nf)

# Calculation of the normalised values by dividing the column values by the n
ormalization factors
results_norm <- results_norm %>%
  mutate(
    across(!c(`impact category`, unit, NF), ~ .x / NF, .names = "norm. {.col}")
    )
    )
```

Determine relevant impact categories

Calculation of proportion of total savings potential of each measure:

```
# Removing the toxicity categories
results_prop <- results_norm %>%
  filter(!`impact category` %in% excluded_categories_pr)
# Calculate the sum of each normalized column and divide the values by this s
results prop <- results prop %>%
  mutate(
    !!sym(name_without_TP_prop) := !!sym(name_without_TP_norm) / sum(abs(!!sy
m(name_without_TP_norm)), na.rm = TRUE),
    !!sym(name_with_TP_prop) := !!sym(name_with_TP_norm) / sum(abs(!!sym(name_
with TP norm)), na.rm = TRUE),
    !!sym(name_SP_TP_prop) := !!sym(name_SP_TP_norm) / sum(abs(!!sym(name_SP_
TP_norm)), na.rm = TRUE),
    !!sym(name_without_BT_prop) := !!sym(name_without_BT_norm) / sum(abs(!!sy
m(name_without_BT_norm)), na.rm = TRUE),
    !!sym(name with BT prop) := !!sym(name with BT norm) / sum(abs(!!sym(name
with BT norm)), na.rm = TRUE),
    !!sym(name_SP_BT_prop) := !!sym(name_SP_BT_norm) / sum(abs(!!sym(name_SP_
BT_norm)), na.rm = TRUE),
    !!sym(name_without_MT_prop) := !!sym(name_without_MT_norm) / sum(abs(!!sy
m(name_without_MT_norm)), na.rm = TRUE),
    !!sym(name with MT prop) := !!sym(name with MT norm) / sum(abs(!!sym(name
with MT norm)), na.rm = TRUE),
    !!sym(name_SP_MT_prop) := !!sym(name_SP_MT_norm) / sum(abs(!!sym(name_SP_
MT norm)), na.rm = TRUE),
    !!sym(name_SP_total_prop) := !!sym(name_SP_total_norm) / sum(abs(!!sym(na
me_SP_total_norm)), na.rm = TRUE)
  )
norm sum <- sum(abs(results prop[[name SP total norm]]))</pre>
```

Determine relevant impact categories:

```
# Sort categories by proportion of total savings potential
results_prop <- results_prop %>%
    arrange(desc(abs(!!sym(name_SP_total_prop))))
# Calculation of the cumulative total and mark categories that contribute to
the sum of 0.8
results_prop <- results_prop %>%
    mutate(
        `cum. sum` = cumsum(abs(!!sym(name_SP_total_prop))),
        relevant = (`cum. sum` < 0.8) | lag(`cum. sum` < 0.8, default = FALSE)
)
# Mark at least the highest three impact categories
results_prop$relevant[1:3] <- TRUE</pre>
```

Save as csv-file

```
write.csv(results_prop, paste0(location_upload, measure, "_data.csv"), row.na
mes = FALSE)
```

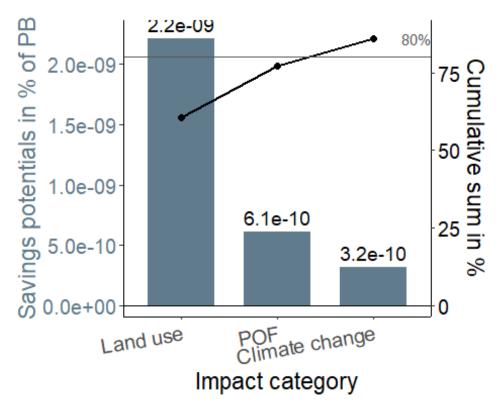
Plot

Prepare data

```
data_plot <- results_prop %>%
  # Select only relevant categories
  filter(relevant == TRUE) %>%
  # Shorten Long category names
  mutate(`impact category` = case when(
    `impact category` == "Photochemical oxidant formation: human health" ~ "P
OF",
    TRUE ~ `impact category`
  )) %>%
  select(`impact category`, name_SP_total_norm) %>%
  # Arrange descending
  arrange(desc(abs(!!sym(name_SP_total_norm))))
## Warning: Using an external vector in selections was deprecated in tidysele
ct 1.1.0.
## i Please use `all_of()` or `any_of()` instead.
##
     # Was:
##
     data %>% select(name SP total norm)
##
##
     # Now:
##
     data %>% select(all of(name SP total norm))
##
## See <https://tidyselect.r-lib.org/reference/faq-external-vector.html>.
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
## generated.
# create cummulative sum
data plot <- data plot %>%
mutate(
```

```
cumsum = cumsum(abs(data_plot$`norm. SP total`/norm_sum))*100
  )
# Define impact categories as factor to ensure order
data_plot$`impact category` <- factor(data_plot$`impact category`, levels = d</pre>
ata_plot$`impact category`)
# max. values for scaling
max norm <- max(data plot[[name SP total norm]])</pre>
max cumsum <- max(data plot$cumsum)</pre>
# Conversion factor
factor <- max_norm / max_cumsum</pre>
Create plot:
ggplot(data plot, aes(x = `impact category`)) +
  geom_bar(aes(y = !!sym(name_SP_total_norm)), stat = "identity",
           width = 0.7, fill = "lightskyblue4", show.legend = FALSE) +
  geom_line(aes(y = cumsum * factor, group = 1), color = "black", size = 1) +
  geom_point(aes(y = cumsum * factor), color = "black", size = 2) +
  geom_hline(yintercept = 0, color = "black", linetype = "solid") +
  geom_hline(yintercept = 80 * factor, color = "grey35", linetype = "solid")
  annotate("text", x = 3.3, y = 80 * factor, label = "80%", color = "grey35",
size = 4, hjust = 0, vjust = -1) +
  geom_text(aes(
    y = !!sym(name SP total norm),
    label = format(!!sym(name SP total norm), scientific = TRUE, digits = 2))
    vjust = -0.5, size = 5) +
  scale_y_continuous(
    name = "Savings potentials in % of PB",
    sec.axis = sec_axis(~ . / factor, name = "Cumulative sum in %"),
  expand = expansion(mult = c(0.04, 0.07))
  labs(x = "Impact category") +
  theme classic() +
  theme(
    axis.text.x = element text(angle = 10, hjust = 1, size = 15),
    axis.text.y.left = element_text(color = "lightskyblue4", size = 15),
    axis.text.y.right = element_text(color = "black", size = 15),
    axis.title.x = element text(size = 17),
    axis.title.y.left = element text(size = 17, color = "lightskyblue4"),
    axis.title.y.right = element_text(size = 17, color = "black")
  )
## Warning: Using `size` aesthetic for lines was deprecated in ggplot2 3.4.0.
## i Please use `linewidth` instead.
## This warning is displayed once every 8 hours.
```

Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
generated.



```
# Save plot
ggsave(paste0(location_upload, measure, "_relevant_SP_norm_allin1.png"), dpi
= 600)
## Saving 5 x 4 in image
```

Tables

Prepare data:

```
# Select relevant columns
results_prop <- results_prop %>%
  select(`impact category`, unit,
         !!sym(name_without_TP_prop), !!sym(name_without_BT_prop), !!sym(name_
_without_MT_prop),
         !!sym(name_with_TP_prop), !!sym(name_with_BT_prop), !!sym(name_with_
MT prop),
         !!sym(name SP TP prop), !!sym(name SP BT prop), !!sym(name SP MT pro
p), !!sym(name_SP_total_prop))
results_norm <- results_norm %>%
  select(`impact category`, unit,
         !!sym(name_without_TP_norm), !!sym(name_without_BT_norm), !!sym(name
without MT norm),
         !!sym(name_with_TP_norm), !!sym(name_with_BT_norm), !!sym(name_with_
MT_norm),
         !!sym(name_SP_TP_norm), !!sym(name_SP_BT_norm), !!sym(name_SP_MT_nor
```

```
m), !!sym(name_SP_total_norm))
# Join of 'results' and 'results_norm' according to the impact categories
results <- left_join(results, results_norm, by = c("impact category" = "impac
t category"))
results <- results %>%
  select(-unit.y) %>%
  rename(unit = unit.x)
# Join of 'results' and 'results_prop' according to the impact categories
results <- left join(results, results prop, by = c("impact category" = "impac
t category"))
results <- results %>%
  select(-unit.y) %>%
  rename(unit = unit.x)
# Sort descending according to normalized savings potential
results <- results %>%
  arrange(desc(abs(!!sym(name SP total norm))))
```

Create tables:

```
# Only results
table results <- results %>%
  select("impact category", "unit",
         !!sym(name without TP), !!sym(name without BT), !!sym(name without M
T),
         !!sym(name with TP), !!sym(name with BT), !!sym(name with MT),
         !!sym(name_SP_TP), !!sym(name_SP_BT), !!sym(name_SP_MT), !!sym(name_
SP total)) %>%
  arrange(desc(abs(!!sym(name_SP_total)))) %>%
  mutate(
    across(-all_of(c("impact category", "unit")), ~ formatC(.x, format = "e",
digits = 2)))
# Create flextable
ft <- flextable(table results) %>%
  theme vanilla() %>%
  autofit() %>%
  bold(part = "header") %>%
  border outer() %>%
  border_inner_h() %>%
  border inner v()
# Save as PNG
tf <- tempfile(fileext = ".png")</pre>
save as image(x = ft, path = paste0(location upload, measure, " results", ".p
ng"))
## [1] "C:/Users/Klene/Documents/Uni_Bremen/WS24_25/Masterarbeit/R/Results/01
results.png"
# Saving potentials
table results <- results %>%
  select(`impact category`, unit, !!sym(name_SP_total), !!sym(name_SP_total_n
orm), !!sym(name SP total prop)) %>%
  arrange(desc(abs(!!sym(name_SP_total_norm)))) %>%
```

```
mutate(
    across(-all_of(c("impact category", "unit")), ~ formatC(.x, format = "e",
digits = 2)))
# Create flextable
ft <- flextable(table results) %>%
  theme vanilla() %>%
  autofit() %>%
  bold(part = "header") %>%
  border outer() %>%
  border inner h() %>%
  border_inner_v()
# Save as PNG
tf <- tempfile(fileext = ".png")</pre>
save as image(x = ft, path = paste0(location upload, measure, " results SPs",
".png"))
## [1] "C:/Users/Klene/Documents/Uni_Bremen/WS24_25/Masterarbeit/R/Results/01
_results_SPs.png"
# Normalized results
table results <- results %>%
  select("impact category", "unit",
         !!sym(name_without_TP_norm), !!sym(name_without_BT_norm), !!sym(name
_without_MT_norm),
         !!sym(name with TP norm), !!sym(name with BT norm), !!sym(name with
MT_norm),
         !!sym(name_SP_TP_norm), !!sym(name_SP_BT_norm), !!sym(name_SP_MT_nor
m), !!sym(name SP total norm)) %>%
  arrange(desc(abs(!!sym(name_SP_total_norm)))) %>%
  mutate(
    across(-all of(c("impact category", "unit")), ~ formatC(.x, format = "e",
digits = 2)))
# Removing the categories without normalization factors
table_results <- table_results %>%
  filter(!`impact category` %in% excluded_categories_nf)
# Rename columns
table results <- table results %>%
  rename(!!sym(name_without_TP) := !!sym(name_without_TP_norm),
         !!sym(name_without_BT) := !!sym(name_without_BT_norm),
         !!sym(name_without_MT) := !!sym(name_without_MT_norm),
          !!sym(name_with_TP) := !!sym(name_with_TP_norm),
         !!sym(name with BT) := !!sym(name with BT norm),
         !!sym(name_with_MT) := !!sym(name_with_MT_norm),
           !!sym(name_SP_TP) := !!sym(name_SP_TP_norm),
         !!sym(name_SP_BT) := !!sym(name_SP_BT_norm),
         !!sym(name_SP_MT) := !!sym(name_SP MT norm),
         !!sym(name_SP_total) := !!sym(name_SP_total_norm)
           )
# Create flextable
ft <- flextable(table_results) %>%
  theme vanilla() %>%
  autofit() %>%
```

```
bold(part = "header") %>%
  border outer() %>%
  border inner h() %>%
  border_inner_v()
# Save as PNG
tf <- tempfile(fileext = ".png")</pre>
save as image(x = ft, path = paste0(location upload, measure, " results norm"
, ".png"))
## [1] "C:/Users/Klene/Documents/Uni Bremen/WS24 25/Masterarbeit/R/Results/01
_results_norm.png"
# Proportion data
table results <- results %>%
  select("impact category", "unit",
         !!sym(name without TP prop), !!sym(name without BT prop), !!sym(name
_without_MT_prop),
         !!sym(name_with_TP_prop), !!sym(name_with_BT_prop), !!sym(name_with_
MT_prop),
         !!sym(name SP TP prop), !!sym(name SP BT prop), !!sym(name SP MT pro
p), !!sym(name SP total prop)) %>%
  arrange(desc(abs(!!sym(name SP total prop)))) %>%
  mutate(
    across(-all_of(c("impact category", "unit")), ~ formatC(.x, format = "e",
digits = 2)))
# Removing the categories without normalization factor
table results <- table results %>%
  filter(!`impact category` %in% excluded categories nf)
# Removing the toxicity categories
table results <- table results %>%
  filter(!`impact category` %in% excluded categories pr)
# Rename columns
table_results <- table_results %>%
  rename(!!sym(name_without_TP) := !!sym(name_without_TP_prop),
         !!sym(name without BT) := !!sym(name without BT prop),
         !!sym(name_without_MT) := !!sym(name_without_MT_prop),
          !!sym(name with TP) := !!sym(name with TP prop),
         !!sym(name_with_BT) := !!sym(name_with_BT_prop),
         !!sym(name_with_MT) := !!sym(name_with_MT_prop),
           !!sym(name_SP_TP) := !!sym(name_SP_TP_prop),
         !!sym(name SP BT) := !!sym(name SP BT prop),
         !!sym(name SP MT) := !!sym(name SP MT prop),
         !!sym(name_SP_total) := !!sym(name_SP_total_prop)
# Create flextable
ft <- flextable(table results) %>%
  theme vanilla() %>%
  autofit() %>%
  bold(part = "header") %>%
  border_outer() %>%
  border inner h() %>%
  border_inner_v()
```

```
# Save as PNG
tf <- tempfile(fileext = ".png")
save_as_image(x = ft, path = paste0(location_upload, measure, "_results_prop"
, ".png"))
## [1] "C:/Users/Klene/Documents/Uni_Bremen/WS24_25/Masterarbeit/R/Results/01
_results_prop.png"</pre>
```

Upload files

```
# # List of Excel files to be uploaded
# files_to_upload <- list.files(path = location_upload, pattern = "\\.xlsx$",
full.names = TRUE)
# # Loop over all files and upload
# for (file_path in files_to_upload) {
  # Creating the file name
# file_name <- basename(file_path)</pre>
# # Upload file
#
  response <- PUT(
#
      url = paste0(nextcloud_url_2, file_name),
      authenticate(username, password),
#
      body = upload file(file path)
  # Check upload status
   if (status_code(response) == 201) {
#
      print(paste(file_name, "was successfully uploaded."))
#
    } else if (status code(response) == 204) {
#
      print(paste(file_name, "was successfully replaced (no content returned)
      print(paste("Error uploading", file_name, ". Status-Code:", status_code
(response)))
# }
# # List of PNG files to be uploaded
# files to upload <- list.files(path = location upload, pattern = "\\.png$",
full.names = TRUE)
# # Loop over all files and upload
# for (file_path in files_to_upload) {
  # Create the file name
# file_name <- basename(file_path)</pre>
# # Upload file
#
  response <- PUT(
#
      url = paste0(nextcloud_url_2, file_name),
#
      authenticate(username, password),
#
      body = upload_file(file_path)
#
   # Check upload status
# if (status_code(response) == 201) {
      print(paste(file_name, "was successfully uploaded."))
# } else if (status_code(response) == 204) {
```

```
# print(paste(file_name, "was successfully replaced (no content returned)
."))
# } else {
# print(paste("Error uploading", file_name, ". Status-Code:", status_code
(response)))
# }
# }
```