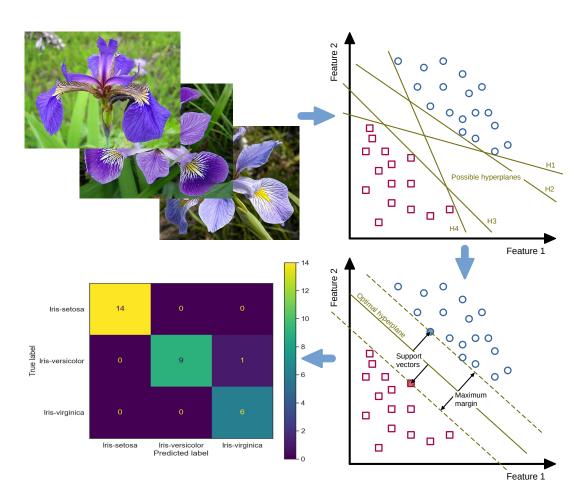
# Analysis of survey data from SoSci Survey in the Edible Cities project using the Analytic Hierarchy Process (AHP) and the R-package ahpsurvey

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This is a placeholder for the abstract that needs to be added later.

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### 1 Introduction

Title drafting:

- Performing the Analytic Hierarchy Process with survey data from \emph{SoSci Survey} and R-package \emph{ahpsurvey}
- Auswertung der Umfragedaten aus \emph{SoSci Survey} zum Thema \emph{Essbare Städte} mit Hilfe des Analytic Hierarchy Process (AHP) und dem R-Paket \emph{ahpsurvey}
- Analysis of survey data from \emph{SoSci Survey} in the \emph{Edible Cities} project using the Analytic Hierarchy Process (AHP) and the R-package \emph{ahpsurvey}

Why we use a Jupyter notebook to to publish the R program examples:

Jupyter is a new **open source** alternative to the proprietary numerical software Mathematica from **Wolfram Research** that is well on the way to become a **standard for exchanging research results** (Somers 2018; Romer 2018).

Originally Jupyter was intended as an IDE for the programming languages **Julia** and **Python**. Besides that it is also possible to install other interpreter kernels, such as the **IRkernel** for R. This can be interesting if the IDE **RStudio Desktop** is not available on the target platform used. For example, it is very difficult to install RStudio on the ARM-based embedded computer **Raspberry Pi** due to many technical dependencies. In contrast, using the R kernel in JupyterLab on the Raspberry Pi works very well and performant.

# 2 Global loading of the used R packages and definition of globally used functions

### 2.1 Install missing packages if not present yet

**Attention:** For some R packages several dependencies have to be installed first with apt install package name>.

Dependencies for package ahpsurvey:

- R package randomNames (it depends on R 4.0, refer to https://cran.r-project.org/web/packages/randomNames/index.html)

Drawback for Raspbian buster: the dependency randomNames is not available for R v3.5.2 as it depends on R (4.0). Upgrading R in Raspbian following the instruction on https://cran.rstudio.com/bin/linux/debian/#debian-buster-stable does not work so far ...

[1] "All required packages are installed."

### 2.2 Load package data.table

The package data.table is used to read and manipulate tables (data.table inherits from data.frame).

[2]: library(data.table)

### 2.3 Load package ggplot2

The package ggplot2 is used to plot diagrams.

[3]: library(ggplot2)

### 2.4 Load packages knitr and IRdisplay

The kable() function from the package knitr is used to output dataframes as a markdown tables.

The display\_markdown() function from the package IRdisplay renders the markdown table in the notebook as well as in the PDF version.

[4]: library(knitr) library(IRdisplay)

### 2.5 Load package tidyr

The package tidyr is used to **reshape** the dataframes and provides functions like **gather()** or **spread()**. Some examples for the application can be found here: Reshaping Your Data with tidyr.

[5]: library(tidyr)

### 2.6 Load package dplyr

The package dplyr is necessary to manipulate dataframes using functions like select(), mutate() and left\_join().

Hint: Setting the parameter warn.conflicts=FALSE when calling the library() function silences annoying messages about masked functions.

[6]: library(dplyr, warn.conflicts=FALSE)

### 2.7 Use pipes for better coding

The package magrittr provides the pipe functionality and can be used to create more effective code for processing large datasets. What pipes of the form like %>% are and how to use them is described here: https://statistik-dresden.de/archives/15679.

**HINT:** The pipe functionality is already available by loading the library tidyr - so you don't have to load it explicitly.

Before pipes can be used in R, the package magrittr has to be loaded first:

[7]: library(magrittr, warn.conflicts=FALSE)

### 2.8 Load package forcats

The fct\_inorder() function from the package forcats is used to reorder the discrete levels of diagram axes according to the intended order of attributes.

[8]: library(forcats)

### 2.9 Load package appsurvey

The package ahpsurvey contains all the necessary mathematical and statistical methods to run the analytical hierarchy process (AHP).

```
[9]: library(ahpsurvey)
```

### 2.10 Function to format dataframes as markdown tables

Following function formats given dataframes as markdown tables using the kable() function from the knitr package.

The display\_markdown() function from the package IRdisplay renders the markdown table in the Jupyter notebook as well as in the compiled PDF version.

```
[10]: func_render_md_tables <- function(df_table, str_table_header) {
    # format the dataframe as a markdown table using the 'kable()' function from
    • the 'knitr' package
    table_out <- kable(
        df_table,
        format = "markdown",
        # digits = 2,
        caption = str_table_header)

    display_markdown(as.character(table_out))
}</pre>
```

# 3 Prepare raw CSV input data from ScoSciSurvey for analytical hierarchy process (AHP)

### 3.1 Set globally used input and output folders for preparing raw CSV data

```
[11]: str_input_path_prep = "./input_data_from_survey"
   str_output_path_prep = "./output_data_manipulated"
```

### 3.2 Define functions to prepare the survey data for further analysis

The following functions are used to read the survey data from the CSV files and prepare the data structure for further analysis with the R package ahpsurvey.

### 3.2.1 Function to read the survey data from CSV files to dataframe objects

Define a function for reading in a CSV file to 4 different dateframes by selecting different columns.

```
# environmental sub-criteria
  df_mySurvey_2 <- fread(</pre>
    file = str_CSVfilename, encoding = "UTF-8".
    header = TRUE, sep = "\t", quote = "\"",
    # dec = ".", row.names = "CASE",
    select = c("CASE", "AU01", "AU02", "AU03",
               "RU01_01", "RU02_01", "RU03_01", "RU04_01", "RU05_01", "RU06_01")
    )
  # social sub-criteria
  df_mySurvey_3 <- fread(</pre>
    file = str_CSVfilename, encoding = "UTF-8",
    header = TRUE, sep = "\t", quote = "\"",
    # dec = ".", row.names = "CASE",
    select = c("CASE", "ASO1", "ASO2", "ASO3",
               "RS01_01", "RS02_01", "RS03_01", "RS04_01", "RS05_01", "RS06_01")
    )
  # economic sub-criteria
  df_mySurvey_4 <- fread(</pre>
    file = str_CSVfilename, encoding = "UTF-8",
    header = TRUE, sep = "\t", quote = "\"",
    # dec = ".", row.names = "CASE",
    select = c("CASE", "AW01", "AW02", "AW03",
               "RW01_01", "RW02_01", "RW03_01", "RW04_01", "RW05_01", "RW06_01")
    )
  output <- list(df_mySurvey_1, df_mySurvey_2, df_mySurvey_3, df_mySurvey_4)</pre>
 return(output)
}
```

### 3.2.2 Function to prepare the data and store it in new dataframes

```
[13]: func_scrambleData <- function(df_inputData, vec_colnames_search_1,_
       svec_colnames_search_2, vec_colnames_out) {
        # Generate new dataframe ...
        df_outputData <- data.frame(matrix(ncol = 3, nrow = 0))</pre>
        # ... and name the columns
        colnames(df_outputData) <- vec_colnames_out</pre>
        # Generate 1. column
        for ( row_idx in 1:nrow(df_inputData) ) {
          # filter column names by vector element
          if (df_inputData[row_idx, colnames(df_inputData) %in% vec_colnames_search_1[1],_
       ⇔with=FALSE] == 1) {
            int_tmp_val <- as.integer(df_inputData[row_idx, colnames(df_inputData) %in%u
       ⇔vec_colnames_search_2[1], with=FALSE])
            int_tmp_val <- int_tmp_val * -1 - 1</pre>
            df_outputData[row_idx, vec_colnames_out[1]] <- int_tmp_val</pre>
          else if (df_inputData[row_idx, colnames(df_inputData) %in%_

    vec_colnames_search_1[1], with=FALSE] == -1) {
            df_outputData[row_idx, vec_colnames_out[1]] <- 1</pre>
```

```
else if (df_inputData[row_idx, colnames(df_inputData) %in%_

    vec_colnames_search_1[1], with=FALSE] == 2) {
    int_tmp_val <- as.integer(df_inputData[row_idx, colnames(df_inputData) %in%u
⇔vec_colnames_search_2[2], with=FALSE])
    int_tmp_val <- int_tmp_val + 1</pre>
    df_outputData[row_idx, vec_colnames_out[1]] <- int_tmp_val</pre>
  }
# Generate 2. column
for ( row_idx in 1:nrow(df_inputData) ) {
  # filter column names by vector element
  if (df_inputData[row_idx, colnames(df_inputData) %in% vec_colnames_search_1[2],_
⇔with=FALSE] == 1) {
    int_tmp_val <- as.integer(df_inputData[row_idx, colnames(df_inputData) %in%u
⇔vec_colnames_search_2[3], with=FALSE])
    int_tmp_val <- int_tmp_val * -1 - 1</pre>
    df_outputData[row_idx, vec_colnames_out[2]] <- int_tmp_val</pre>
  else if (df_inputData[row_idx, colnames(df_inputData) %in%_

    vec_colnames_search_1[2], with=FALSE] == -1) {
    df_outputData[row_idx, vec_colnames_out[2]] <- 1</pre>
  }
  else if (df_inputData[row_idx, colnames(df_inputData) %in%_
ovec_colnames_search_1[2], with=FALSE] == 2) {
    int_tmp_val <- as.integer(df_inputData[row_idx, colnames(df_inputData) %in%_

→vec_colnames_search_2[4], with=FALSE])
    int_tmp_val <- int_tmp_val + 1</pre>
    df_outputData[row_idx, vec_colnames_out[2]] <- int_tmp_val</pre>
  }
}
# Generate 3. column
for ( row_idx in 1:nrow(df_inputData) ) {
  # filter column names by vector element
  if (df_inputData[row_idx, colnames(df_inputData) %in% vec_colnames_search_1[3],_
⇔with=FALSE] == 1) {
    int_tmp_val <- as.integer(df_inputData[row_idx, colnames(df_inputData) %in%_

→vec_colnames_search_2[5], with=FALSE])
    int_tmp_val <- int_tmp_val * -1 - 1</pre>
    df_outputData[row_idx, vec_colnames_out[3]] <- int_tmp_val</pre>
  }
  else if (df_inputData[row_idx, colnames(df_inputData) %in%_
→vec_colnames_search_1[3], with=FALSE] == -1) {
    df_outputData[row_idx, vec_colnames_out[3]] <- 1</pre>
  else if (df_inputData[row_idx, colnames(df_inputData) %in%_
ovec_colnames_search_1[3], with=FALSE] == 2) {
    int_tmp_val <- as.integer(df_inputData[row_idx, colnames(df_inputData) %in%u
→vec_colnames_search_2[6], with=FALSE])
    int_tmp_val <- int_tmp_val + 1</pre>
```

```
df_outputData[row_idx, vec_colnames_out[3]] <- int_tmp_val
}

# return scrambled dataframe
return(df_outputData)
}</pre>
```

### 3.2.3 Function to write resulting dataframes to CSV files

```
[14]: func_writeDataframe_to_CSVfile <- function(str_path, str_CSVfilename, df_dataframe, u
       str_filenameExtension) {
        # Split file name on second underscore, found here:
        # https://stackoverflow.com/questions/32398427/
       r-split-a-character-string-on-the-second-underscore/32398489#32398489
       list_str_split <- strsplit(sub('(^[^_]+_[^_]+)_(.*)$', '\\1 \\2',_
       str_CSVfilename), ' ')
        # extend the file name prefix and glue together with old suffix
        str_CSVfilename_extended <- paste(list_str_split[[1]][1], str_filenameExtension,_
       Glist_str_split[[1]][2], sep="_")
        # extend file name by path
        str_CSVfilename_extended <- paste(str_path, str_CSVfilename_extended, sep="/")
        write.table(df_dataframe, file = str_CSVfilename_extended,
                    fileEncoding = "UTF-8", row.names = FALSE,
                    col.names = TRUE, sep = "\t", quote = TRUE)
      }
```

# 3.3 Create dataframe (table) handling the file names of input CSV data (raw data from survey)

Table 1: File table for handling the file names of input CSV data (raw data from survey)

file_idx keys	filenames	descriptions
1 all	$\begin{array}{c} {\rm rdata\_all\_AHP\_edible\_Cities\_2022\text{-}03\text{-}18\_09\text{-}} \\ {\rm 53.csv} \end{array}$	all target groups together

file_idx	keys	filenames	descriptions
2	CA	rdata_CA_AHP_edible_Cities_2022-03-18_10-28.csv	City Administrations
3	NGO	rdata_NGO_AHP_edible_Cities_2022-03-18 10-40.csv	Non-Governmental Organisations
4	PE	$\begin{array}{c}$	Practitioners and Experts

### 3.4 Prepare the data and store it in new CSV files for each criterion

### 3.4.1 Criteria (main criteria)

Walk over all input CSV files, select necessary columns, filter cells by given algorithm, and write the results to output CSV files:

### 3.4.2 Environmental sub-criteria

Walk over all input CSV files, select necessary columns, filter cells by given algorithm, and write the results to output CSV files:

### 3.4.3 Social sub-criteria

Walk over all input CSV files, select necessary columns, filter cells by given algorithm, and write the results to output CSV files:

### 3.4.4 Economic sub-criteria

Walk over all input CSV files, select necessary columns, filter cells by given algorithm, and write the results to output CSV files:

# 4 Application of the processed survey data in the analytical hierarchy process (AHP)

### 4.1 Set globally used input and output folders for processing AHP

```
[20]: str_input_path_ahp = "./output_data_manipulated"
str_output_path_ahp = "./output_data_AHP"
```

### 4.2 Define functions for processing AHP

### 4.2.1 Function to read in the processed survey data from CSV files to dataframes

Define a function for reading in a CSV file to a date frame.

```
[21]: func_readCSVdata_to_dataframe <- function(str_CSVfilename) {

    df_CSVdata <- fread(
        file = str_CSVfilename, encoding = "UTF-8",
        header = TRUE, sep = "\t", quote = "\""
    )

    return(df_CSVdata)
}</pre>
```

### 4.2.2 Function to generate a dataframe with eigentrue values (weights)

```
[22]: func_genEigentrue_to_dataframe <- function(df_surveyData, vec_attributes) {
    list_mat_judgement <- df_surveyData %>%
        ahp.mat(vec_attributes, negconvert = TRUE)

    df_eigentrue <- ahp.indpref(list_mat_judgement, vec_attributes, method = "eigen")
    return(df_eigentrue)
}</pre>
```

### 4.2.3 Function to generate an array with consistency ratios

```
[23]: func_genCR_to_arr <- function(df_surveyData, vec_attributes) {
    arr_cr <- df_surveyData %>%
        ahp.mat(vec_attributes, negconvert = TRUE) %>%
        ahp.cr(vec_attributes, ri=0.58)

    return(arr_cr)
}
```

### 4.2.4 Function to generate a dataframe with consistency ratios

```
mutate(rowid = 1:length(arr_cr), arr_cr.dum = as.factor(ifelse(arr_cr <=
u
consistency_thres, 1, 0)))

# rename column with consistency ratios
colnames(df_cr)[1] <- str_CRlabel

return(df_cr)
}</pre>
```

### 4.2.5 Function to visualize individual priorities and consistency ratios

```
[25]: func_visuPriosCRs <- function(df_surveyData, df_cr, arr_cr, consistency_thres = 0.
       41, vec_attributes, df_eigentrue, vec_labels, str_image_filename, str_title) {
        # Select columns 'arr cr.dum' and 'rowid' from input dataframe 'df cr'
        # 'arr_cr.dum': Binary representation of the consistency ratio (0: inconsistent;
       →1: consistent)
        df_cr_sel <- df_cr %>%
         select(arr_cr.dum, rowid)
        # Generate AHP pairwise matrices from survey data
        mat_ahp <- ahp.mat(df_surveyData, atts = vec_attributes, negconvert = TRUE)</pre>
        # Compute priority weights of individual decision-makers
        df_prio_weights <- ahp.indpref(mat_ahp, vec_attributes, method = "eigen")</pre>
        # Add column 'rowid' from dataframe 'df_eigentrue'
        df_prio_weights <- mutate(df_prio_weights, rowid = 1:nrow(df_eigentrue))</pre>
        # Left join dataframes 'df_prio_weights' and 'df_cr_sel' by column 'rowid'
        df_prio_weights_binCR <- left_join(df_prio_weights, df_cr_sel, by = "rowid")</pre>
        # Gather columns of 'df_prio_weights_binCR' into key-value pairs
        # The function 'all_of(vec_attributes)' selects data-variables listed in the
       ⇔character vector 'vec_attributes'
        li_binCR_attr_weights <- gather(df_prio_weights_binCR, all_of(vec_attributes),_</pre>
       # Create the violin plots with overlaid box plots.
        # Important: The function "fct_inorder()" is necessary to reorder the discrete_
       \hookrightarrow levels
        # of the diagram axes according to the intended order of the attributes.
        # Otherwise, the order will be automatically set alphanumerically and will not
        # the attribute labels later.
        # refer: https://stackoverflow.com/a/41417136
        plt <- ggplot(li_binCR_attr_weights, aes(x = fct_inorder(var), y = pref)) +</pre>
          # Add a violin plot
          geom_violin(alpha = 0.6, width = 0.8, color = "transparent", fill = "gray") +
          # 'geom_jitter()' is a shortcut for 'geom_point(position = "jitter")'
          # Adds a small amount of random variation to the location of each point
          # to handle overplotting caused by discreteness in smaller datasets
          geom_jitter(alpha = 0.6, height = 0, width = 0.1, aes(color = arr_cr.dum)) +
          # Add a box plot
          geom_boxplot(alpha = 0, width = 0.3, color = "#808080") +
          \# Set discrete levels of the diagram X-axis according to the corresponding \sqcup
       ⇔attribute labels
```

```
scale_x_discrete("Attribute", label = vec_labels) +
    # Configure the diagram Y-axis to display continuos data with
    # scale in percent and choose where the ticks appear by setting 'breaks'
    scale_y_continuous("Weight (dominant eigenvalue)",
                         labels = scales::percent,
                         breaks = c(seq(0,0.7,0.1))) +
    # Hide the title of the legend
    guides(color=guide_legend(title=NULL)) +
    # Set the discrete color scale according to the binarized consistency ratio
    # and use the Unicode character '\u2264' for '<='
    scale_color_discrete(breaks = c(0,1),
                         labels = c(paste("CR >", consistency_thres),
                                    paste("CR \u2264", consistency_thres))) +
    # Set caption text to be displayed in the bottom-right of the plot
    # with number of rows and mean value of the consistency ratio
    labs(NULL, caption = paste("n =", nrow(df_surveyData), ",", "Mean CR =",
                               round(mean(arr_cr), 3))) +
    # Set theme of the plot to 'theme_light()'
    theme_light() +
    # Set the title of the diagram
    ggtitle(str_title)
  print(plt)
  # Save generated ggplot graphics to PNG image files
  ggsave(filename = str_image_filename, width = 7, height = 7, dpi = 300)
}
```

### 4.2.6 Function to generate geometric mean values from individual judgement matrices

```
[26]: func_aggpref_gmean <- function(df_surveyData, vec_attributes, arr_cr,_u
       ⇔consistency_thres=0.1, str_CRlabel) {
        df_cr <- df_surveyData %>%
          ahp.mat(vec_attributes, negconvert = TRUE) %>%
          ahp.cr(vec_attributes, ri=0.58) %>%
          data.frame() %>%
          mutate(rowid = 1:length(arr_cr), arr_cr.dum = as.factor(ifelse(arr_cr <=_</pre>
       ⇔consistency_thres, 1, 0)))
        # rename column with consistency ratios
        colnames(df_cr)[1] <- str_CRlabel</pre>
        # combine dataframe 'df_cr' with raw survey data ('df_surveyData')
        df_cr_wRaw <- cbind(df_cr, df_surveyData)</pre>
        # remove rows, where 'arr_cr.dum' == 0 (inconsistent data)
        df_cr_wRaw_cons <- df_cr_wRaw[df_cr_wRaw$arr_cr.dum != 0, ]</pre>
        # get individual judgement matrices from last 3 columns
        list_mat_judgement <- df_cr_wRaw_cons[tail(names(df_cr_wRaw_cons), 3)] %>%
```

```
ahp.mat(vec_attributes, negconvert = TRUE)

# get geometric mean values from judgement matrices
list_gmean_1 <- ahp.aggpref(list_mat_judgement, vec_attributes, method = "eigen",
aggmethod = "geometric")

return(list_gmean_1)
}</pre>
```

### 4.2.7 Function to normalize the geometric mean values

```
[27]: func_norm_gmean <- function(list_gmeans) {
        # normalization so that the sum of the geometric mean values is 1 (corresponds to \Box
        →100%)
        df_gmean_l <- data.frame(list_gmeans)</pre>
        # rename column with geometric mean values (raw)
        colnames(df_gmean_1)[1] <- "gmean.raw"</pre>
        gmean_sum <- 0</pre>
        for ( val in list_gmeans ) {
          gmean_sum <- gmean_sum + val</pre>
        df_gmean_l["Sum", 1] <- gmean_sum</pre>
        for (idx in 1:length(list_gmeans)) {
           gmean_norm <- list_gmeans[[idx]] / gmean_sum</pre>
           df_gmean_l[idx, "gmean.norm"] <- gmean_norm</pre>
        }
        gmean_sum_norm <- 0</pre>
        # iterate over all rows except the last, because this is the sum itself
        for ( row in 1:(nrow(df_gmean_1)-1) ) {
           gmean_sum_norm <- gmean_sum_norm + df_gmean_l[row, 2]</pre>
        df_gmean_1["Sum", 2] <- gmean_sum_norm</pre>
        return(df_gmean_1)
      }
```

### 4.2.8 Function to process the ahpsurvey and create violin plots with overlaid box plots

```
func_process_ahpsurvey <- function(df_csvInputFiles, str_participants_group,__
str_input_path_ahp, str_output_path_ahp, df_attributes_labels) {
   row_start = 1
   row_end = 3

   df_outputTable <- data.table()

   for ( file_idx in 1:nrow(df_csvInputFiles) ) {
        # create dataframe from current input CSV file
        str_filename <- paste(str_input_path_ahp, df_csvInputFiles[file_idx,__
filenames], sep="/")
        df_processed_survey_data <- func_readCSVdata_to_dataframe(str_filename)</pre>
```

```
# create vectors for attributes and labels from a subset of dataframe_
→ 'df_attributes_labels_XXX'
      vec_atts <- df_attributes_labels[c(row_start:row_end), attr]</pre>
      vec_labels <- df_attributes_labels[c(row_start:row_end), labels]</pre>
      # shift row interval for next iteration
      row_start = row_start + 3
      row_end = row_end + 3
      # create title string for violin plots
      str\_violin\_title = "Violins display priorities and consistency ratios for_{\sqcup}
⇔attributes:"
      str_violin_title <- paste(str_violin_title, vec_atts[1], sep = " ")</pre>
      # extend title string by looping through attributes, starting with 2nd_{\sqcup}
      for ( idx in 2:length(vec_atts) ) {
           str_violin_title <- paste(str_violin_title, vec_atts[idx], sep = ", ")</pre>
      # print(str_violin_title)
      # generate dataframe with eigentrue values (weights)
      df_eigentrue_weights <-
□func_genEigentrue_to_dataframe(df_processed_survey_data, vec_atts)
      # generate an array with consistency ratios
      arr_CRs <- func_genCR_to_arr(df_processed_survey_data, vec_atts)</pre>
      # generate an extended dataframe with consistency ratios
      consistency_thres = 0.1
      str_CRlabel <- paste("CR", df_csvInputFiles[file_idx, keys], sep="_")</pre>
      df_CRs <- func_genCR_to_dataframe(df_processed_survey_data, vec_atts,_
⇔arr_CRs, consistency_thres, str_CRlabel)
      str image filename <- paste("ahp violin", str participants group, ...
df_csvInputFiles[file_idx, keys], sep="_")
      str_image_filename <- paste(str_image_filename, ".png", sep="")</pre>
      str_image_filename <- paste(str_output_path_ahp, str_image_filename, sep="/
ç")
      func_visuPriosCRs(df_processed_survey_data, df_CRs, arr_CRs,__
Gonsistency_thres, vec_atts, df_eigentrue_weights, vec_labels,
str_image_filename, str_violin_title)
       # combine dataframes of eigentrue values (weights) with consistency ratios
      df_outputTable <- cbind(df_outputTable, df_eigentrue_weights)</pre>
       # add only specific columns of 'df_CRs' (omit column 'row_id')
      df_outputTable <- cbind(df_outputTable, df_CRs[c(1, 3)])</pre>
  }
  # extend file name by path
  str_CSVfilename_output <- paste("rdata", str_participants_group,_

¬"eigentrue_CRs", sep="_")

  str_CSVfilename_output <- paste(str_CSVfilename_output, ".csv", sep="")
  str_CSVfilename_output <- paste(str_output_path_ahp, str_CSVfilename_output,_
⇔sep="/")
  \# write dataframe 'df_outputTable' to CSV file for further statistical analysis
  write.table(df_outputTable, file = str_CSVfilename_output,
```

```
fileEncoding = "UTF-8", row.names = FALSE,
col.names = TRUE, sep = "\t", quote = TRUE)
}
```

### 4.2.9 Function to calculate aggregated preference weights for consistent datasets

```
[29]: func_calc_pref_weights <- function(df_csvInputFiles, str_input_path_ahp,_u

¬df_attributes_labels) {
          row_start = 1
          row_end = 3
          for ( file_idx in 1:nrow(df_csvInputFiles) ) {
              # create dataframe from current input CSV file
              str_filename <- paste(str_input_path_ahp, df_csvInputFiles[file_idx,_u
        ⇔filenames], sep="/")
              str_file_description <- df_csvInputFiles[file_idx, descriptions]</pre>
              df_processed_survey_data <- func_readCSVdata_to_dataframe(str_filename)</pre>
              # create vectors for attributes and labels from a subset of dataframe_{\sqcup}
        → 'df_attributes_labels_XXX'
              vec_atts <- df_attributes_labels[c(row_start:row_end), attr]</pre>
              vec_labels <- df_attributes_labels[c(row_start:row_end), labels]</pre>
              # shift row interval for next iteration
              row_start = row_start + 3
              row_end = row_end + 3
              # generate an array with consistency ratios
              arr_CRs <- func_genCR_to_arr(df_processed_survey_data, vec_atts)</pre>
              str_CRlabel <- paste("CR", df_csvInputFiles[file_idx, keys], sep="_")</pre>
              list_gmean <- func_aggpref_gmean(df_processed_survey_data, vec_atts,_</pre>
        Garr_CRs, consistency_thres=0.1, str_CRlabel)
              df_gmean <- func_norm_gmean(list_gmean)</pre>
              # store dataframe of main criteria to calculate the total weights of \Box
        \hookrightarrow sub-criteria
              if ( str_file_description == "criteria (main criteria)" ) {
                  df_gmean_main_criteria <- df_gmean</pre>
              } else if ( str_file_description == "environmental sub-criteria" ) {
                   # multiply column 'gmean.norm' of sub-criteria dataframe by 'Envi'u
        ⇔value from main criteria dataframe
                   # and store the result in the new column 'gmean.total'
                   df_gmean$gmean.total <- df_gmean$gmean.norm *_

df_gmean_main_criteria["Envi", "gmean.norm"]

                   # sum up the values of the new column 'qmean.total' and write to row_
       → 'Sum' of the same column
                  df_gmean["Sum", "gmean.total"] <- sum(df_gmean[c(1:3), "gmean.total"])</pre>
              } else if ( str_file_description == "social sub-criteria" ) {
                   df_gmeangmean.total <- df_gmeangmean.norm *_

→df_gmean_main_criteria["Soci", "gmean.norm"]
```

```
df_gmean["Sum", "gmean.total"] <- sum(df_gmean[c(1:3), "gmean.total"])
} else if ( str_file_description == "economic sub-criteria" ) {
    df_gmean$gmean.total <- df_gmean$gmean.norm *_U

df_gmean_main_criteria["Econ", "gmean.norm"]

df_gmean["Sum", "gmean.total"] <- sum(df_gmean[c(1:3), "gmean.total"])
}

func_render_md_tables(df_gmean, paste("Aggregated weights for ",_U

str_file_description))
}
</pre>
```

### 4.3 Create dataframes (tables) handling the file names of processed survey data

### 4.3.1 File table for all participants

Table 2: File table for all participants

$file\_idx$	keys	filenames	descriptions
1	crit	rdata_all_crit_AHP_edible_Cities_2022-03-18_09-	criteria (main criteria)
2	env	53.csv rdata_all_env_AHP_edible_Cities_2022-03-18_09- 53.csv	environmental sub-criteria
3	soc	rdata_all_soc_AHP_edible_Cities_2022-03-18_09-	social sub-criteria
4	eco	53.csv rdata_all_eco_AHP_edible_Cities_2022-03-18_09- 53.csv	economic sub-criteria

### 4.3.2 File table for city administrations

Table 3: File table for city administrations

file_idx	keys	filenames	descriptions
1	crit	rdata_CA_crit_AHP_edible_Cities_2022-03-	criteria (main criteria)
2	env	18_10-28.csv rdata_CA_env_AHP_edible_Cities_2022-03- 18_10-28.csv	environmental sub-criteria
3	soc	rdata_CA_soc_AHP_edible_Cities_2022-03-	social sub-criteria
4	eco	18_10-28.csv rdata_CA_eco_AHP_edible_Cities_2022-03- 18_10-28.csv	economic sub-criteria

### 4.3.3 File table for non-governmental organizations

Table 4: File table for non-governmental organizations

file_idx	keys	filenames	descriptions
1	$\operatorname{crit}$	rdata_NGO_crit_AHP_edible_Cities_2022-03-	criteria (main criteria)
2	env	18_10-40.csv rdata_NGO_env_AHP_edible_Cities_2022-03- 18_10-40.csv	environmental sub-criteria
3	soc	rdata_NGO_soc_AHP_edible_Cities_2022-03-	social sub-criteria
4	eco	18_10-40.csv rdata_NGO_eco_AHP_edible_Cities_2022-03- 18_10-40.csv	economic sub-criteria

### 4.3.4 File table for practitioners and experts

```
[33]: df_csvInputFiles_PE <- data.table(
    file_idx = 1:4,
    keys = c("crit", "env", "soc", "eco"),
    filenames = c("rdata_PE_crit_AHP_edible_Cities_2022-03-18_10-41.csv",</pre>
```

Table 5: File table for practitioners and experts

file_idx	keys	filenames	descriptions
1	crit	rdata_PE_crit_AHP_edible_Cities_2022-03-	criteria (main criteria)
2	env	18_10-41.csv rdata_PE_env_AHP_edible_Cities_2022-03- 18_10-41.csv	environmental sub-criteria
3	soc	$rdata\_PE\_soc\_AHP\_edible\_Cities\_2022-03-$	social sub-criteria
4	eco	18_10-41.csv rdata_PE_eco_AHP_edible_Cities_2022-03- 18_10-41.csv	economic sub-criteria

### 4.3.5 Table with attributes and labels

```
[34]: df_attributes_labels <- data.table(
    # idx = 1:12,
    attr = c("Envi", "Soci", "Econ",
        "Clim", "BDiv", "CiEc",
        "KEdu", "Comm", "Part",
        "Qual", "LVCs", "Affo"),
    labels = c("Environment", "Society", "Economy",
        "Microclimate and Hydrology", "Biodiversity", "Circular economy",
        "Knowledge and Education", "Community Building", "Participation",
        "Food Quality and Safety", "Local Value Chains", "Food Affordability")
)

# print.data.frame(df_attributes_labels, right=FALSE)
func_render_md_tables(df_attributes_labels, "Table with attributes and labels")</pre>
```

Table 6: Table with attributes and labels

attr	labels
Envi	Environment
Soci	Society
Econ	Economy
Clim	Microclimate and Hydrology
BDiv	Biodiversity
CiEc	Circular economy
KEdu	Knowledge and Education
Comm	Community Building
Part	Participation
Qual	Food Quality and Safety
LVCs	Local Value Chains

attr	labels
Affo	Food Affordability

# 4.4 Visualize datasets of survey with package ahpsurvey for each group of participants

### 4.4.1 All participants

```
[35]: func_process_ahpsurvey(df_csvInputFiles_all, "all", str_input_path_ahp,_u

str_output_path_ahp, df_attributes_labels)
```

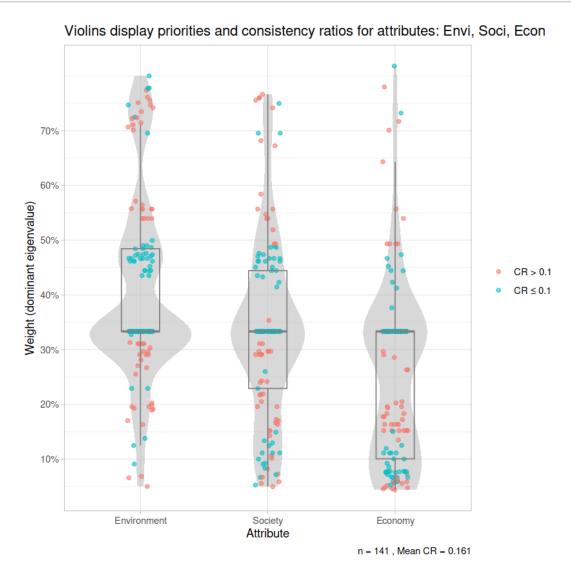


Figure 1: Series of violin plots that display the priorities and consistency ratios of all participants

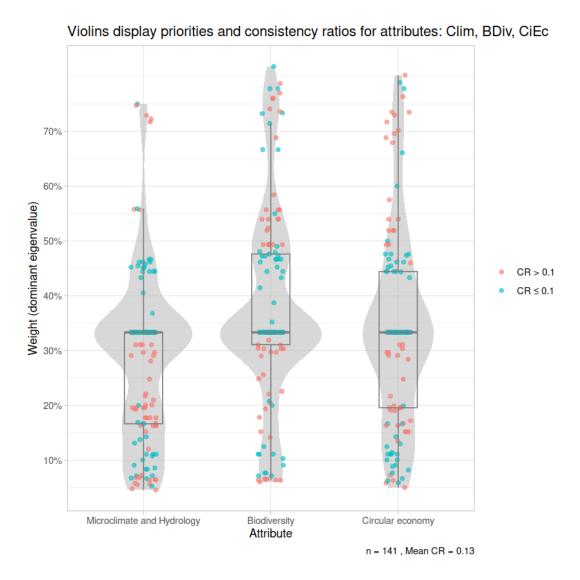


Figure 2: Series of violin plots that display the priorities and consistency ratios of all participants

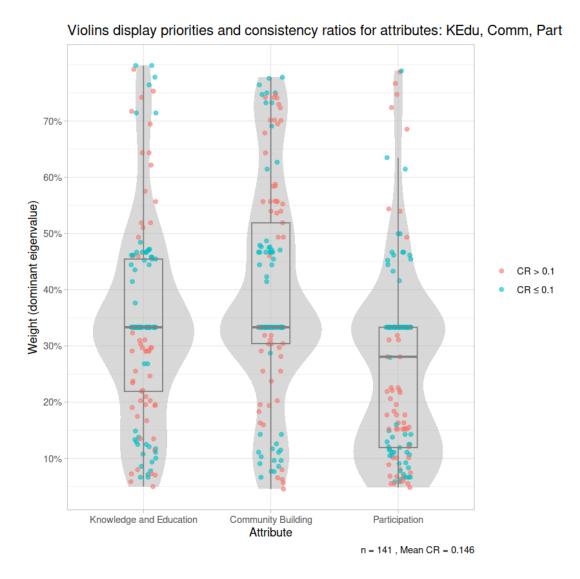


Figure 3: Series of violin plots that display the priorities and consistency ratios of all participants

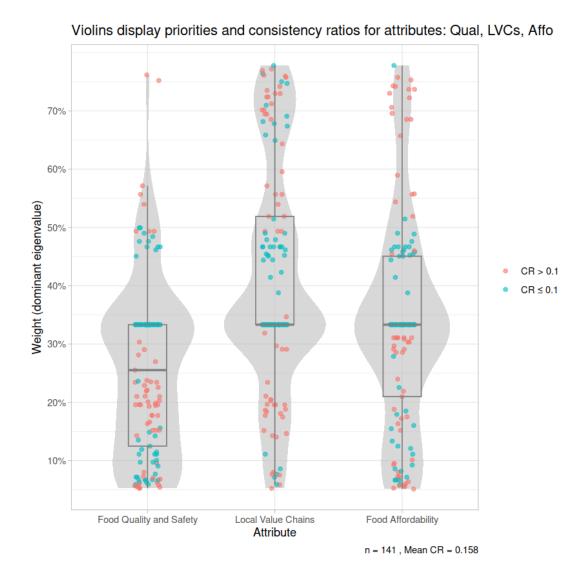


Figure 4: Series of violin plots that display the priorities and consistency ratios of all participants

### 4.4.2 Participants of city administrations

```
[36]: func_process_ahpsurvey(df_csvInputFiles_CA, "CA", str_input_path_ahp, ustr_output_path_ahp, df_attributes_labels)
```

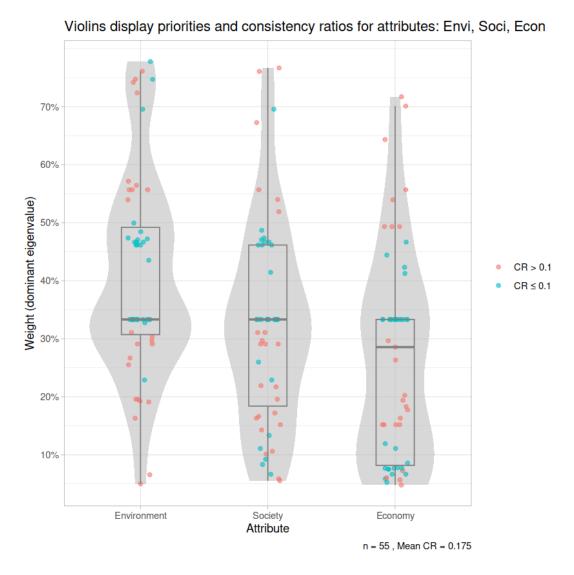


Figure 5: Series of violin plots that display the priorities and consistency ratios of city administrations

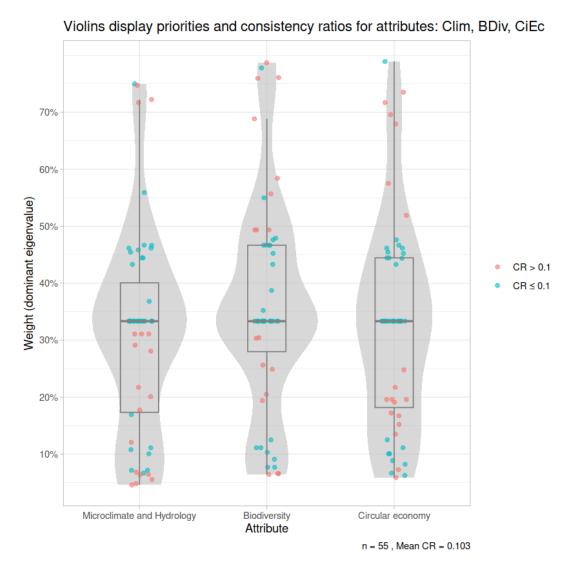


Figure 6: Series of violin plots that display the priorities and consistency ratios of city administrations

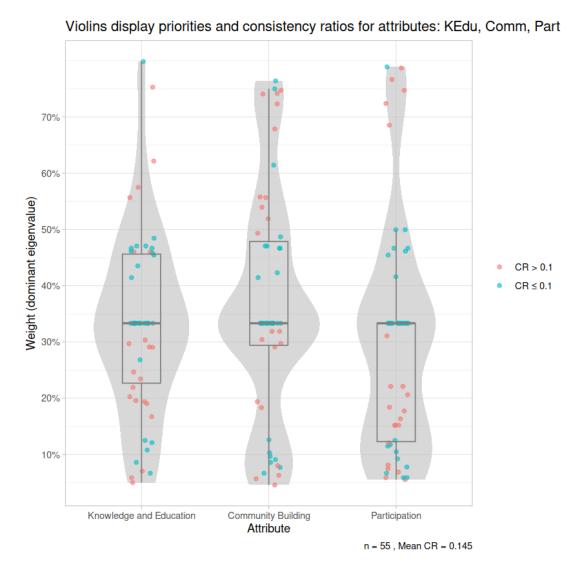


Figure 7: Series of violin plots that display the priorities and consistency ratios of city administrations

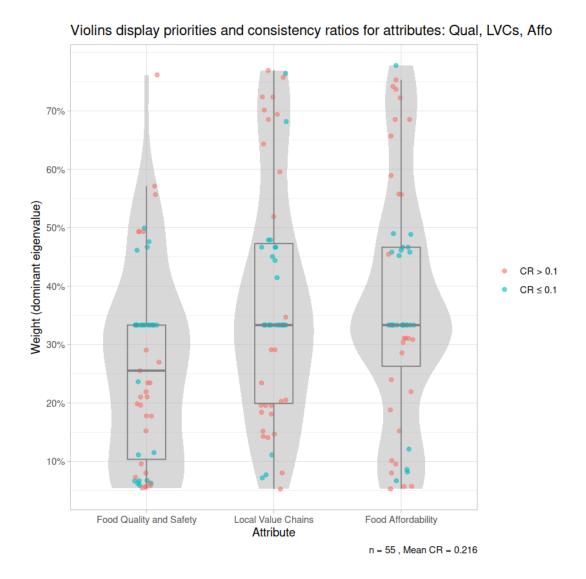


Figure 8: Series of violin plots that display the priorities and consistency ratios of city administrations

### 4.4.3 Participants of non-governmental organizations

```
[37]: func_process_ahpsurvey(df_csvInputFiles_NGO, "NGO", str_input_path_ahp,_

str_output_path_ahp, df_attributes_labels)
```

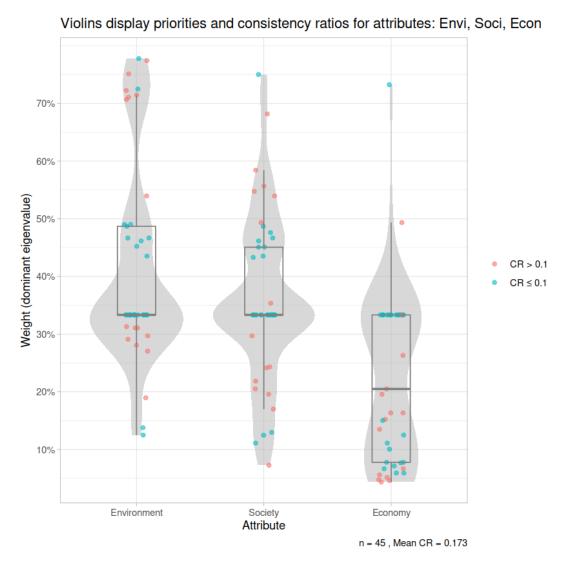


Figure 9: Series of violin plots that display the priorities and consistency ratios of non-governmental organizations

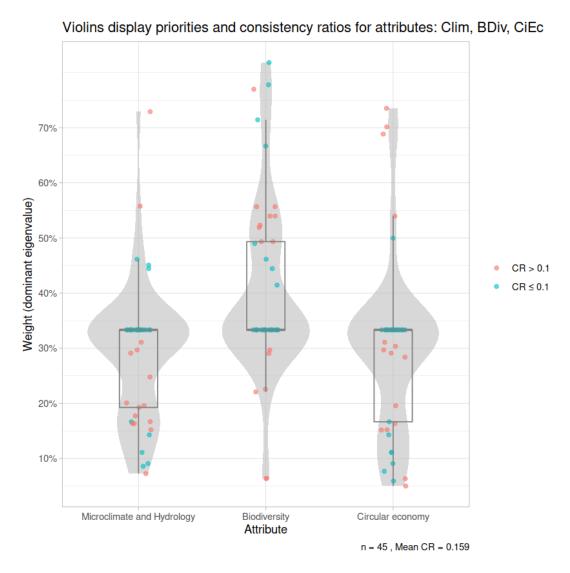


Figure 10: Series of violin plots that display the priorities and consistency ratios of non-governmental organizations

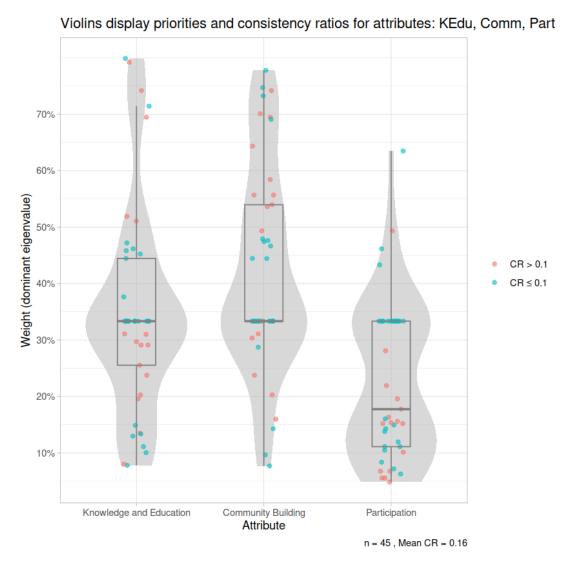


Figure 11: Series of violin plots that display the priorities and consistency ratios of non-governmental organizations

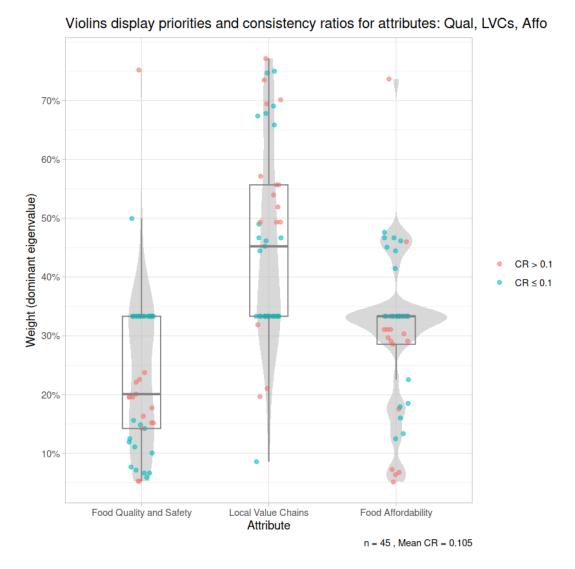


Figure 12: Series of violin plots that display the priorities and consistency ratios of non-governmental organizations

### 4.4.4 Participants of practitioners and experts

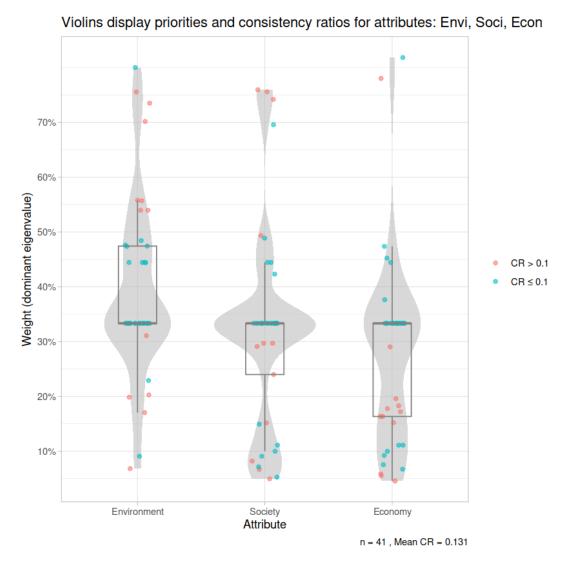


Figure 13: Series of violin plots that display the priorities and consistency ratios of practitioners and experts

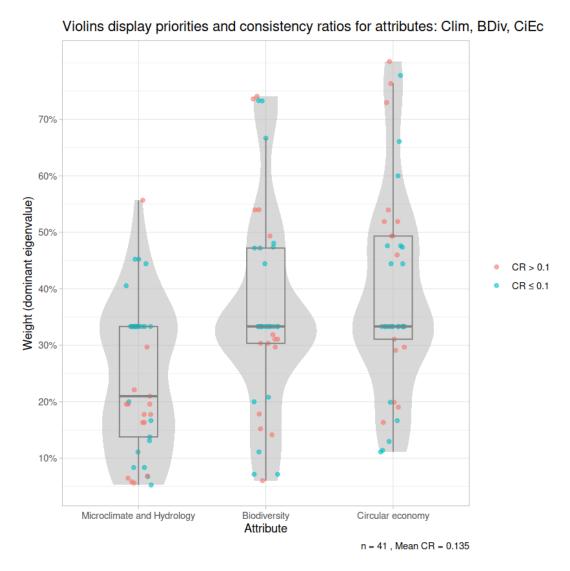


Figure 14: Series of violin plots that display the priorities and consistency ratios of practitioners and experts

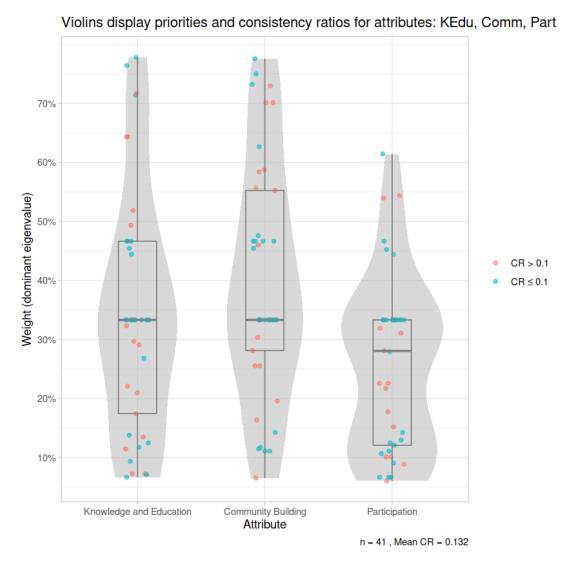


Figure 15: Series of violin plots that display the priorities and consistency ratios of practitioners and experts

# Violins display priorities and consistency ratios for attributes: Qual, LVCs, Affo 60% 60% 60% CR > 0.1 CR ≤ 0.1 10% Food Quality and Safety Local Value Chains Attribute n = 41, Mean CR = 0.137

## Figure 16: Series of violin plots that display the priorities and consistency ratios of practitioners and experts

### 4.5 Calculate aggregated preference weights for consistent datasets

[39]: func\_calc\_pref\_weights(df\_csvInputFiles\_all, str\_input\_path\_ahp,\_u

df\_attributes\_labels)

Table 7: Aggregated weights for criteria (main criteria)

	gmean.raw	gmean.norm
Envi	0.3739039	0.4179807
Soci	0.3023657	0.3380094
Econ	0.2182787	0.2440099
Sum	0.8945482	1.0000000

Table 8: Aggregated weights for environmental sub-criteria

	gmean.raw	gmean.norm	gmean.total
Clim	0.2741300	0.3089871	0.1291506
BDiv	0.3291718	0.3710277	0.1550824
CiEc	0.2838874	0.3199852	0.1337476
Sum	0.8871892	1.0000000	0.4179807

Table 9: Aggregated weights for social sub-criteria

	gmean.raw	gmean.norm	gmean.total
KEdu	0.3018698	0.3512801	0.1187360
Comm	0.3212634	0.3738480	0.1263641
Part	0.2362090	0.2748719	0.0929093
Sum	0.8593422	1.0000000	0.3380094

Table 10: Aggregated weights for economic sub-criteria

	gmean.raw	gmean.norm	gmean.total
Qual	0.2329901	0.2625724	0.0640703
LVCs	0.3666029	0.4131498	0.1008127
Affo	0.2877435	0.3242777	0.0791270
$\operatorname{Sum}$	0.8873365	1.0000000	0.2440099

### 5 Summary and outlook

### 6 References

### Online references

Romer, Paul (Apr. 13, 2018). Jupyter, Mathematica, and the Future of the Research Paper. English. URL: https://paulromer.net/jupyter-mathematica-and-the-future-of-the-research-paper/ (visited on 09/08/2022) (cit. on p. 3).

Somers, James (Apr. 5, 2018). The Scientific Paper Is Obsolete. English. The Atlantic. URL: https://www.theatlantic.com/science/archive/2018/04/the-scientific-paper-is-obsolete/556676/ (visited on 09/08/2022) (cit. on p. 3).