

# Package: kwb.flusshygiene (via r-universe)

August 24, 2024

**Title** Functions used within FLUSSHYGIENE project (BMBF)

**Version** 0.3.0

**Description** Easy and transferable functions for creating and managing models for hygiene data in rivers. This package is developed within the FLUSSHYGIENE project. See <https://bmbf.nawam-rewam.de/en/projekt/flusshygiene/> for details.

**License** MIT + file LICENSE

**URL** <https://github.com/KWB-R/kwb.flusshygiene>

**BugReports** <https://github.com/KWB-R/kwb.flusshygiene/issues>

**Depends** R (>= 3.4), stats, utils, graphics, grDevices

**Imports** MASS (>= 7.3-47), dplyr (>= 0.7.4), tidyR (>= 0.7.2), tibble (>= 1.3.4), readr (>= 1.1.1), rlang (>= 0.4.0), lubridate (>= 1.7.1), ggplot2 (>= 2.2.1), rstanarm (>= 2.15.3), kwb.utils (>= 0.4.3), purrr (>= 0.3.2)

**Suggests** knitr (>= 1.23), readxl (>= 1.3.1), rmarkdown (>= 1.13), testthat (>= 2.2.1)

**VignetteBuilder** knitr

**Remotes** `github::kwb-r/kwb.utils`

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 6.1.1

**Repository** <https://kwb-r.r-universe.dev>

**RemoteUrl** <https://github.com/KWB-R/kwb.flusshygiene>

**RemoteRef** HEAD

**RemoteSha** b9fee176e392f7932673e8efb092114cfdcbd9d0

## Contents

assess_bathing_quality_eu . . . . .	2
build_model . . . . .	3
build_new_model . . . . .	4
choose_model . . . . .	4
correlation_scatterplot . . . . .	5
get_mpn_ci . . . . .	6
get_paths . . . . .	6
import_riverdata . . . . .	7
kwb.flusshygience . . . . .	8
plot_data_overview . . . . .	9
plot_hygiene_overview . . . . .	9
plot_predicted_quality . . . . .	10
plot_q_overview . . . . .	10
plot_rain_overview . . . . .	11
plot_stan_model . . . . .	11
predict_quality . . . . .	12
readTableData . . . . .	13
river_model_prediction . . . . .	13
unroll_physical_data . . . . .	14

## Index

15

---

assess\_bathing\_quality\_eu  
*Bathing quality assessment (EU)*

---

### Description

Computes the quality assessment according to european bathing directive 2006/7/EC from E.Coli values. The four possible quality levels are: excellent, good, sufficient and poor

### Usage

```
assess_bathing_quality_eu(e.coli, log = TRUE)
```

### Arguments

e.coli	A numeric vector with e.coli values
log	logical. Are the values log-values?

### Value

Returns a single factor with all quality levels

---

build_model	<i>Build a Model for E.Coli</i>
-------------	---------------------------------

---

**Description**

Functions for modelbuilding \n build\_model: takes the riverdata, handles the other functions and invokes [stan\\_lm](#)

**Usage**

```
build_model(riverdata, variables = ask_for_variables(riverdata),
            with_interaction = TRUE)

ask_for_variables(riverdata)

process_model_riverdata(riverdata, variables)

create_formula(variables, with_interaction = FALSE)
```

**Arguments**

riverdata	a list with riverdata (hygiene + physical data)
variables	character. Selected variables for the model
with_interaction	logical. Formula with interactions? Default set to TRUE

**Details**

Build the model from hygiene data and physical data like flow, rain, wwtp. Asks for user input to select variables. Computes the data.frame with data for hygiene and chosen variables and creates a formula of the form:  $Q*(K + R)$  while multiple Qs will be multiplied, multiple Ks and Rs will be added.

**Value**

- Returns a model of the riverdata.
- Returns a character-vector with the chosen model variables
- Returns a data.frame with data for hygiene and chosen variables
- Returns parsed model-formula. (Like model\$formula)

**Functions**

- `ask_for_variables`: Internal function. Quite time consuming
- `process_model_riverdata`: Internal usage
- `create_formula`: Internal usage

## Examples

```
## Not run: variables <- c("e.coli","q_havel",...)
lm(formula = eval(create_formula(variables)),
  data = process_model_riverdata(riverdata, variables))
## End(Not run)

create_formula(c("log_e.coli","q_havel","ka_ruhleben","r_berlin"))
create_formula(c("e.coli","r_mitte","r_charlottenburg","r_spandau"))
```

build\_new\_model

*Create a new Model*

## Description

Main function for creation, object handling and saving of new models. The models will be saved in a own subdirectory as r-objects as a side-effect. The models will not be returned. They have to be loaded by other functions.

## Usage

```
build_new_model(river)
```

## Arguments

river	character. river you want to build a model on
-------	---

## Value

This function returns merely a message what happend.

choose\_model

*Set User Input for a List of Models*

## Description

Choose model asks the user inside the console for input. Options are Exit, New Model, or one of a list of existing models. If no integer number was presented by the user an ERROR message will be created but no ERROR will be thrown. This way this can be inserted inside a loop.

## Usage

```
choose_model(rivermodels)
```

## Arguments

rivermodels	A list of named models
-------------	------------------------

**Value**

Returns the user input as character vector, or an ERROR message.

**Examples**

```
choose_model(list())
choose_model(list(fake_river_model = 1))
```

---

**correlation\_scatterplot**

*Scatterplotmatrix of similar Variables to E.Coli*

---

**Description**

Takes similar named variables and produces a matrix of scatterplots and their correlation coefficients to E.Coli.

**Usage**

```
correlation_scatterplot(df, ...)
correlation_values(df, ...)
```

**Arguments**

df	data.frame with data for e.coli and chosen variables in lagdays
...	Arguments passed to stats::cor

**Value**

Plotting function. Returns a plot.

Returns correlation values.

**Functions**

- **correlation\_values**: Internal function

**Examples**

```
correlation_values(data.frame(datum = rep("egal",10), e.coli = 1:10, var = 1:10), variable = "var")
```

`get_mpn_ci`*Get MPN Confidence Intervals for E.Coli***Description**

Lookup laboratory tables for MPN values for E.Coli to get upper and lower 0.95 confidence interval for the given values. If value is not directly found in table it will be generated by interpolating nearest neighbors.

**Usage**

```
get_mpn_ci(e.coli)
```

**Arguments**

<code>e.coli</code>	numeric. A vector for e.coli values
---------------------	-------------------------------------

**Value**

A data.frame with 3 columns: e.coli, lo, up

**Examples**

```
## Not run:
print(get_mpn_ci(c(15,30,35,60,61,71,120,1959,25000,369990)))
## End(Not run)
```

`get_paths`*Get List of Paths used in the Flusshygiene Project***Description**

Get List of Paths used in the Flusshygiene Project

**Usage**

```
get_paths(resolve = TRUE, ...)
```

**Arguments**

<code>resolve</code>	if TRUE (default) path placeholders are resolved
<code>...</code>	arguments passed to <code>resolve</code> if <code>resolve</code> is TRUE

`import_riverdata`

7

## Examples

```
## Not run:  
paths <- get_paths()  
  
# Paths to the different work package folders  
paths$ap2  
paths$ap3  
paths$ap4  
  
# What tables are contained in the ODM database?  
kwb.db::hsTables(paths$odm)  
  
# Get all Flusshygiene data into one data frame  
data <- kwb.ogre.model::get_lab_values(paths$odm)  
  
## End(Not run)
```

---

`import_riverdata`      *Read existing River Data*

---

## Description

Read existing, preprocessed csv files with first column datetime and other columns variable information.

## Usage

```
import_riverdata(path)
```

## Arguments

`path`      character-string to a DATA\_preprocessed\_csv directory

## Value

Returns a list of data.frames containing the river data

## Description

The kwb.flusshygiene package provides functions in three major categories: model handling, model creation and model prediction.

### Model handling

`river_model_prediction` is the main function in this package. It uses all of the following functions from within.

`get_paths` reads a serverpath library for easy directory accessing

`search_existing_models` searches saved R-objects in the river directories.

### Model creation

`build_new_model` again a overhead function for model creation. Asks the user whether or not the new model shall be saved as R-object.

`import_riverdata` reads all river data from a directory.

`build_model` is a small function handling model creation and invokes `stan_lm` for model building.

`ask_for_variables` asks the user which variables shall be included in the model. Creates plots as a side effect.

`process_model_riverdata` processes a data.frame with the necessary data for the data argument in `stan_lm`

`create_formula` creates a hygiene formula out of the variables with the form `e.coli ~ Q * (R + Ka)`

### Model prediction

`predict_quality` is the overhead function for the prediction. It also invokes `posterior_predict.stanreg`

`get_newdata` gathers the latest data for prediction.

`print_latest` prints the prediction of the latest day.

`plot_predicted_quality` plots a whole season with quality assessment.

### Utility functions

`unroll_physical_data` unrolls a list with data with lagday combinations to 5 days (default)

`correlation_scatterplot` plots a scatterplot matrix of the unrolled physical data together with correlation values.

### Plotting functions

`plot_stan_model` plots a stan\_lm posterior prediction with quality assessment.  
`plot_data_overview` plot data overview  
`plot_hygiene_overview` a statistical hygiene data overview  
`plot_q_overview` plot a overview of all q values  
`plot_rain_overview` plot a monthly overview of all rain gauges

---

`plot_data_overview` *Plot Data Timeline Overview*

---

#### Description

Creates a plot with segments or points of the data availability.

#### Usage

```
plot_data_overview(riverdata, type = "segment")
```

#### Arguments

<code>riverdata</code>	A list of hygiene and physical data of the river
<code>type</code>	Either "segment" or "point" for more precise information

#### Value

Returns a plot

---

`plot_hygiene_overview` *Plot Hygiene Overview*

---

#### Description

Creates a plot with three graphs: Histogramm of all e.coli values, a density curve of the last 16 values, and a boxplot of all values again

#### Usage

```
plot_hygiene_overview(hygiene_df)
```

#### Arguments

<code>hygiene_df</code>	A data.frame with the hygiene data of a given river
-------------------------	---

#### Value

Returns a plot

**plot\_predicted\_quality**  
*Plot Quality*

### Description

Window function for [plot\\_stan\\_model](#)

### Usage

```
plot_predicted_quality(model, prediction, ...)
```

### Arguments

model	stan.lm model for the river
prediction	list of season, ppd of predcit and ppd of means
...	Further parameter passed to plot.default

### Value

Plotting function. Returns a plot.

**plot\_q\_overview**      *Plot Flowing Conditions*

### Description

Creates a plot with the standard flowing conditions over the year. The data of all years will be taken into account.

### Usage

```
plot_q_overview(q_df)
```

### Arguments

q_df	The data.frame with 2 columns: datum and q
------	--

### Value

Returns a plot

---

plot\_rain\_overview      *Plot Monthly Rain Summary*

---

**Description**

Creates a plot with a monthly summary overview over the different rain sites

**Usage**

```
plot_rain_overview(df)
```

**Arguments**

df                  A data frame with different rain gauges.

**Value**

Returns a plot

---

plot\_stan\_model      *Plot Model Prediction with Quality Assessment*

---

**Description**

Plots a sample of posterior predictions and means. Furthermore colours an hygiene quality assessment as background (see EU Bathing Water Directive) Dark blue means excellent quality. Steelblue means good quality. Yellow means sufficient quality. Red means insufficient quality.

**Usage**

```
plot_stan_model(timestamp, predict, linpred, log = FALSE, q90, q95,
                 nlines = 250, nlinesCenter = 100, ...)
```

**Arguments**

timestamp	POSIX. The x-axis timestamp
predict	ppd. The posterior prediction of the model
linpred	ppd. The linpred (predicted means) of the model
log	logical. Is E.Coli log01-transformed?
q90	numeric. The 90. percentile of predict.
q95	numeric. The 95. percentile of predict.
nlines	numeric. How many lines for posterior predictions?
nlinesCenter	numeric. How many lines for predicted means?
...	Further parameters for plot.default

**Value**

Plotting function. Returns a plot.

predict_quality	<i>Predict Hygiene Quality</i>
-----------------	--------------------------------

**Description**

Main function for invoking and object handling. E.Coli hygiene models will be used to predict hygiene quality on differnt scopes.

**Usage**

```
predict_quality(model, river_dir, output = "season")

get_newdata(variables, river_dir)

print_latest(model, newdata)

get_latest_season(newdata)
```

**Arguments**

model	stan_lm. A model of e.coli concentration in given river
river_dir	character. Path to river-data for up-to-date predictions.
output	character. "season" will return a list with prediction, "latest" will return console output
variables	character. A vector with all variables used in the model
newdata	data.frame with physical data used in the model

**Value**

Returns a list of physical data and prediction and linpred from model

Returns a data.frame with the merged data found

**Functions**

- `get_newdata`: Internal Usage
- `print_latest`: Internal Usage
- `get_latest_season`: Internal Usage

---

readTableData      *Read Data for ODM Tables*

---

### Description

Read data for ODM tables from CSV files stored in the package

### Usage

```
readTableData(sourcedir = system.file("extdata", "ODM", package =
  "kwb.flusshygiene"))
```

### Arguments

sourcedir      path to input directory

### Value

list of data frames

---

river\_model\_prediction      *Programm for Model Handling and Prediction*

---

### Description

This function is the front-end for model search on the server, model building with existing data, or prediction with new data. It invokes all other functions and handles their objects. It is a main function.

### Usage

```
river_model_prediction(river)

search_existing_models(river_dir)
```

### Arguments

river      character. The desired river, like "isar".  
river\_dir      character. Path to server and river directory

### Value

(invisible) The data.frame returned by the prediction plus a date column for easy plotting.  
Returns a list with the existing models for that river (empty if no model was found).

## Functions

- `search_existing_models`: directory searching

## Examples

```
river_model_prediction(river = "isar")
```

```
serverpath <- "//poseidon/projekte$/SUW_Department/Projects/FLUSSHYGIENE/Data-Work packages/Daten"
river_dir <- search_existing_river_dir(river = "isar", server = serverpath)
search_existing_models(river_dir = river_dir)
```

`unroll_physical_data`    *Unroll Lagdays of Data*

## Description

Unrolls the lagdays of data.frames.

## Usage

```
unroll_physical_data(physical_data)

unroll_lagdays(df, n = 5)
```

## Arguments

<code>physical_data</code>	list of river data (without hygiene)
<code>df</code>	data.frame of 2 columns: datum and var
<code>n</code>	numeric. unto to which day shall be lagged behind?

## Value

Returns a list of data.frames for each variable. The data.frames contain the unrolled lagdays (with `maxday = 5`, `length(df) == 17`)

## Functions

- `unroll_lagdays`: Internal usage mostly

## Examples

```
df1 <- data.frame(datum = rep("egal", 25), var = 1:25)
df2 <- data.frame(datum = rep("egal", 25), var2 = 51:75, var3 = 101:125)
unroll_lagdays(df1)
summary(unroll_physical_data(list(var1 = df1, var2 = df2)))
```

# Index

ask\_for\_variables, 8  
ask\_for\_variables (build\_model), 3  
assess\_bathing\_quality\_eu, 2  
  
build\_model, 3, 8  
build\_new\_model, 4, 8  
  
choose\_model, 4  
correlation\_scatterplot, 5, 8  
correlation\_values  
    (correlation\_scatterplot), 5  
create\_formula, 8  
create\_formula (build\_model), 3  
  
get\_latest\_season (predict\_quality), 12  
get\_mpn\_ci, 6  
get\_newdata, 8  
get\_newdata (predict\_quality), 12  
get\_paths, 6, 8  
  
import\_riverdata, 7, 8  
  
kwb.flusshygiene, 8  
kwb.flusshygiene-package  
    (kwb.flusshygiene), 8  
  
plot\_data\_overview, 9, 9  
plot\_hygiene\_overview, 9, 9  
plot\_predicted\_quality, 8, 10  
plot\_q\_overview, 9, 10  
plot\_rain\_overview, 9, 11  
plot\_stan\_model, 9, 10, 11  
posterior\_predict.stanreg, 8  
predict\_quality, 8, 12  
print\_latest, 8  
print\_latest (predict\_quality), 12  
process\_model\_riverdata, 8  
process\_model\_riverdata (build\_model), 3  
  
readTableData, 13  
resolve, 6  
  
river\_model\_prediction, 8, 13  
  
search\_existing\_models, 8  
search\_existing\_models  
    (river\_model\_prediction), 13  
stan\_lm, 3, 8  
  
unroll\_lagdays (unroll\_physical\_data),  
    14  
unroll\_physical\_data, 8, 14