

Package: kwb.geosalz (via r-universe)

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Title R Package for Documenting Workflow Used in Project ``geosalz"

Version 0.7.0

Description R Package for Documenting Workflow Used in Project ``geosalz".

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

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

Depends R (>= 2.10)

Imports archive, cowplot, cellranger (>= 1.1.0), crayon (>= 1.3.4), data.table (>= 1.12.0), dplyr (>= 0.7.8), forcats, fs (>= 1.2.6), geosalz.phreeqc, ggplot2 (>= 3.1.0), janitor (>= 1.1.1), jsonlite, kwb.base, kwb.file, kwb.nextcloud, kwb.utils, magrittr (>= 1.5), parallel, readODS, readr (>= 1.4.0), readxl (>= 1.2.0), RColorBrewer, rlang (>= 0.3.1), rmarkdown (>= 1.11), sf, sftp, stringr (>= 1.4.0), tibble (>= 2.0.1), tidyr (>= 0.8.2), tidyselect (>= 1.1.2), withr, zoo

Suggests covr (>= 3.2.1), DT, ggforce, htmltools, htmlwidgets, knitr (>= 1.21), kwb.pkgbuild, leaflet, openxlsx, phreeqc, sessioninfo (>= 1.1.1), wasserportal

VignetteBuilder knitr

Remotes github::KWB-R/geosalz.phreeqc, github::KWB-R/kwb.base, github::KWB-R/kwb.nextcloud@dev, github::KWB-R/kwb.pkgbuild, github::KWB-R/kwb.utils, github::KWB-R/wasserportal, github::stenevang/sftp

ByteCompile true

Encoding UTF-8

LazyData true

RoxygenNote 7.3.1

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

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

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<code>add_para_metadata</code>	<i>add_para_metadata</i>
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Description

`add_para_metadata`

Usage

`add_para_metadata(df, lookup_para_path, parameters_path)`

Arguments

<code>df</code>	<code>df</code>
<code>lookup_para_path</code>	<code>lookup_para_path</code>
<code>parameters_path</code>	<code>parameters_path</code>

Value

return "df" with added parameter metadata

add_site_metadata	<i>add_site_metadata</i>
-------------------	--------------------------

Description

add_site_metadata

Usage

```
add_site_metadata(df, site_path)
```

Arguments

df	df
site_path	site_path

Value

data frame with added site metadata

all_defined	<i>Check if all strings are not empty</i>
-------------	---

Description

Check if all strings are not empty

Usage

```
all_defined(x)
```

Arguments

x	vector of character
---	---------------------

Value

TRUE or FALSE

cat_green_bold_0 *Helper function: cat_green_bold_0*

Description

Helper function: cat_green_bold_0

Usage

cat_green_bold_0(...)

Arguments

... text passed to crayon::green()

Value

formatted text output

cat_red_bold_0 *Helper function: cat_red_bold_0*

Description

Helper function: cat_red_bold_0

Usage

cat_red_bold_0(...)

Arguments

... text passed to crayon::red

Value

formatted text output

column_pattern_gather_ignore

Helper function: column_pattern_gather_ignore

Description

Helper function: column_pattern_gather_ignore

Usage

```
column_pattern_gather_ignore(
  fields = c("Datum", "KN", "[iI]nterne Nr.", "Name der", "Ort", "Probe", "Prü",
            "Untersuchung", "Labor", "Jahr", "Galer", "Detail", "Meß", "Zeit", "Bezei", "Monat")
)
```

Arguments

fields column names to be ignored for gathering (default: c("Datum", "KN", "[iI]nterne Nr.", "Name der", "Ort", "Probe", "Prü", "Untersuchung", "Labor", "Jahr", "Galer", "Detail", "Me\u00DF", "Zeit", "Bezei", "Monat"))

Value

vector with ignored columns for gathering

column_pattern_gather_ignore_clean

Helper function: column_pattern_gather_ignore_clean

Description

Helper function: column_pattern_gather_ignore_clean

Usage

```
column_pattern_gather_ignore_clean(
  fields = c("LabSampleCode", "Date", "Time", "Waterbody", "ExSiteCode", "Site")
)
```

Arguments

fields column names to be ignored for gathering (default: c("LabSampleCode", "Date", "Time", "Waterbody", "ExSiteCode", "Site"))

Value

vector with ignored columns for gathering

```
convert_phreeqc_input_to_wide
  Convert Phreeqc input to "wide" format
```

Description

Convert Phreeqc input to "wide" format

Usage

```
convert_phreeqc_input_to_wide(phreeqc_input)
```

Arguments

phreeqc_input Phreeqc input as retrieved by [get_phreeqc_data](#)

Value

Phreeqc input in "wide" format

```
convert_to_sf          Convert to SF
```

Description

Convert to SF

Usage

```
convert_to_sf(
  df,
  crs_source = 25833,
  crs_target = 4326,
  col_coord_x = "Rechtswert_UTM_33_N",
  col_cood_y = "Hochwert_UTM_33_N"
)
```

Arguments

df	data frame or tibble with spatial data
crs_source	original CRS (default: 25833)
crs_target	target CRS (default: 4326)
col_coord_x	column name of latitude (default: "Rechtswert_UTM_33_N")
col_cood_y	column name of longitude (default: "Hochwert_UTM_33_N")

Value

data frame or tibble converted to sf

Examples

```
gwl_master <- jsonlite::fromJSON("https://kwb-r.github.io/wasserportal/stations_gwl_master.json")
convert_to_sf(gwl_master)
```

convert_xls_as_xlsx *Convert xls to xlsx*

Description

Convert xls to xlsx

Usage

```
convert_xls_as_xlsx(  
  input_dir,  
  export_dir = tempdir(),  
  office_folder = safe_office_folder(),  
  dbg = TRUE  
)
```

Arguments

input_dir	input directory containing .xls files
export_dir	export directory (default: tempdir())
office_folder	office folder path (default: safe_office_folder)
dbg	debug (default: TRUE)

convert_xls_to_xlsx *Helper function: convert_xls_to_xlsx*

Description

Helper function: convert_xls_to_xlsx

Usage

```
convert_xls_to_xlsx(exe, xls, xlsx, i, n_files, dbg = TRUE)
```


Arguments

exe	exe
xls	xls
xlsx	xlsx
i	i
n_files	n_files
dbg	debug (default: TRUE)

copy_lookup_para_file *Helper function: copy_lookup_para_file*

Description

Helper function: copy_lookup_para_file

Usage

```
copy_lookup_para_file(
  from_dir,
  to_dir,
  overwrite = FALSE,
  recursive = TRUE,
  file_pattern = "^lookup_para\\.csv$"
)
```

Arguments

from_dir	input directory with xlsx files
to_dir	target directory where to copy the xlsx files
overwrite	should existing files be overwritten (TRUE) otherwise (FALSE) ? (default: FALSE)
recursive	if TRUE recursively find all xlsx files in the directory specified in parameter "from_dir" (default: TRUE)
file_pattern	pattern for identifying lookup_para file (default: "^lookup_para\\.csv\$")

copy_xlsx_files *Helper function: copy_xlsx_files*

Description

Helper function: copy_xlsx_files

Usage

```
copy_xlsx_files(
  from_dir,
  to_dir,
  overwrite = FALSE,
  recursive = TRUE,
  file_pattern = "[xX][lL][sS][xX]"
)
```

Arguments

from_dir	input directory with xlsx files
to_dir	target directory where to copy the xlsx files
overwrite	should existing files be overwritten (TRUE) otherwise (FALSE) ? (default: FALSE)
recursive	if TRUE recursively find all xlsx files in the directory specified in parameter "from_dir" (default: TRUE)
file_pattern	pattern for identifying xlsx files (default: "[xX][lL][sS][xX]")

create_emshoff91_import
Create EMSHOFF91 Import Data Frame

Description

Create EMSHOFF91 Import Data Frame

Usage

```
create_emshoff91_import(
  ods_dir,
  files_to_ignore = c("cl25", "c11iste", "rupelauf", "salzlast")
)
```

Arguments

`ods_dir` directory to ".ods" files created manually by importing original ".wq1" files into LibreOffice 7.0 on Ubuntu with encoding (Western Europe (DOS/OS2-437/US)) and exporting to ".ods" format

`files_to_ignore` tidied names of files to ignore due to complex data input structure not yet covered by importer (default: c("cl25", "clliste", "gwnguete", "rupelauf", "salzlast"))

Value

data frame with columns "ods_paths" (full paths to ".ods" files), "ods_files" (their "basenames") and "ods_names_clean" (tidied names used as identifier)

Examples

```
## Not run:
ods_dir <- "<replace-with-path-to-files>/emshoff91/converted_ods"
emshoff91_import <- create_emshoff91_import(ods_dir)

## End(Not run)
```

create_sftp_connection

Measurement Chains: Create an SFTP Connection

Description

Measurement Chains: Create an SFTP Connection

Usage

```
create_sftp_connection()
```

Value

sftp connection

delete_registry	<i>Helper function: delete_registry</i>
-----------------	---

Description

Helper function: delete_registry

Usage

```
delete_registry(office_folder = safe_office_folder(), dbg = TRUE)
```

Arguments

office_folder	office folder path (default: safe_office_folder)
dbg	debug (default: TRUE)

download_measurementchains_data	<i>Measurement Chains: download data</i>
---------------------------------	--

Description

Measurement Chains: download data

Usage

```
download_measurementchains_data(
  sftp_paths,
  target_directory = temp_dir(),
  sftp_connection = create_sftp_connection(),
  run_parallel = TRUE,
  debug = FALSE
)
```

Arguments

sftp_paths	character vector with paths to files to be downloaded. As retrieved by get_measurementchains_files column "sftp_path"
target_directory	target directory
sftp_connection	an SFTP connection as retrieved by create_sftp_connection
run_parallel	default: TRUE
debug	show debug messages (default: FALSE)

Value

tibble with columns file_id, sftp_path and local_path of csv files

Examples

```
## Not run:  
mc_files <- kwb.geosalz::get_measurementchains_files()  
target_directory <- tempdir()  
local_paths <- kwb.geosalz::download_measurementchains_data(  
  sftp_paths = mc_files$sftp_path,  
  target_directory)  
  
## End(Not run)
```

emshoff91_list_to_df *Emshoff 91: list to data frame*

Description

Emshoff 91: list to data frame

Usage

```
emshoff91_list_to_df(emshoff91_list)
```

Arguments

emshoff91_list list as retrieved by [read_multiple_emshoff91_ods](#)

Value

tibble

emshoff91_remap_values
Emshoff 91: remap values from imported tibble

Description

Emshoff 91: remap values from imported tibble

Usage

```
emshoff91_remap_values(
  emshoff91_df,
  remap_list = list(fi_mi = "fi_mi_m_nn", ku_sto = "kupp_st", lf = "el_lf", progr =
    "beprob_progr", strat = "stratigr", uv254 = "uv_ext"),
  delete_cols = TRUE
)
```

Arguments

emshoff91_df	tibble as retrieved by <code>emshoff91_list_to_df</code>
remap_list	list with values to be remapped. Names of the list are columns values contained in list values should be mapped to (default: <code>list(fi_mi = "fi_mi_m_nn", ku_sto = "kupp_st", lf = "el_lf", progr = "beprob_progr", strat = "stratigr", uv254 = "uv_ext")</code>)
delete_cols	should unneeded columns be deleted, i.e. the ones where data were mapped from (default: TRUE)

Value

data frame with remapped values and deleted columns were this values were copied from (default: TRUE)

<code>gather_and_join_1</code>	<i>Helper function: <code>gather_and_join_1</code></i>
--------------------------------	--

Description

Helper function: `gather_and_join_1`

Usage

```
gather_and_join_1(tmp_data, columns_keep, metadata, dbg = FALSE)
```

Arguments

tmp_data	tmp_data
columns_keep	columns_keep
metadata	metadata
dbg	dbg (default: FALSE)

Value

gathered and joined data frame

gather_and_join_2 *Helper function: gather_and_join_2*

Description

Helper function: gather_and_join_2

Usage

```
gather_and_join_2(tmp_content, columns_keep, header)
```

Arguments

tmp_content	tmp_content
columns_keep	columns_keep
header	header

Value

gathered and joined data frame

get_excelcnv_exe *Helper function: get_excelcnv_exe*

Description

Helper function: get_excelcnv_exe

Usage

```
get_excelcnv_exe(office_folder = safe_office_folder())
```

Arguments

office_folder	office folder path (default: safe_office_folder)
---------------	--

Value

path containing 'excelcnv.exe'

get_foerdermengen *get_foerdermengen*

Description

get_foerdermengen

Usage

```
get_foerdermengen(  
  xlsx_path,  
  sheet_name = "WW Q Rhow ",  
  sheet_range = "A4:S127"  
)
```

Arguments

xlsx_path path to xlsx file with pumping rates
sheet_name sheet_name (default: "WW Q Rhow ")
sheet_range sheet_range (default: "A4:S127")

Value

data frame with annual pumping rates per waterworks

get_foerdermengen_gal_fri
Get Abtraction of Friedrichshagen Well Galleries

Description

Get Abtraction of Friedrichshagen Well Galleries

Usage

```
get_foerdermengen_gal_fri(path)
```

Arguments

path path to "2018-04-27 Rohwasser Bericht - Galeriefördermengen.xlsx"

Value

tidy data frame with abstraction rates for waterworks Friedrichshagen

`get_measurementchains_files`*Measurement Chains: Get Tidied Files Metadata*

Description

Measurement Chains: Get Tidied Files Metadata

Usage

```
get_measurementchains_files(  
  sftp_connection = create_sftp_connection(),  
  debug = FALSE  
)
```

Arguments

`sftp_connection` an SFTP connection as retrieved by [create_sftp_connection](#)

`debug` show debug messages (default: FALSE)

Value

tibble with information on available files and tidied meta-information based on file naming

Examples

```
## Not run:  
mc_files <- kwb.geosalz::get_measurementchains_files()  
str(mc_files)  
  
## End(Not run)
```

`get_measurementchains_metadata`*Measurement Chains: Get Metadata*

Description

Measurement Chains: Get Metadata

Usage

```
get_measurementchains_metadata(file = extdata_file("metadata_messketten.csv"))
```

Arguments

`file` path to measurement chains metadata file. Default: `kwb.geosalz:::extdata_file("metadata_messketten.csv")`

Value

tibble with measurement chains metadata

Examples

```
mc_metadata <- kwb.geosalz::get_measurementchains_metadata()
str(mc_metadata)
mc_metadata
```

```
get_measurement_chain_data_on_cloud
```

Get Measurement Chain Data on KWB Cloud

Description

Get Measurement Chain Data on KWB Cloud

Usage

```
get_measurement_chain_data_on_cloud(dbg = TRUE)
```

Arguments

`dbg` logical indicating whether or not to show debug messages

Value

data frame with the content of "mc_data.zip" in the GeoSalz project folder on the Nextcloud server. The SFTP paths to the files from which the data in "mc_data.zip" originate are returned in attribute "sftp_paths". If either of the files "mc_data.zip" or "mc_files.csv" does not exist, NULL is returned.

get_measurmentchains_data_stats

Measurement Chains: get statistics for data

Description

Measurement Chains: get statistics for data

Usage

get_measurmentchains_data_stats(mc_data)

Arguments

mc_data tibble with measurement chains data as retrieved by [read_measurementchains_data](#)

Value

tibble with columns datetime min/max, q10 (10 median, q90 (90

get_meta_sheet_or_stop

Helper function: get_meta_sheet_or_stop

Description

Helper function: get_meta_sheet_or_stop

Usage

get_meta_sheet_or_stop(sheets, pattern, file)

Arguments

sheets	sheets
pattern	pattern
file	file

Value

meta sheet name

get_pandoc_info	<i>Get Information on Pandoc</i>
-----------------	----------------------------------

Description

Get Information on Pandoc

Usage

```
get_pandoc_info()
```

Value

data frame with columns `pandoc_directory`, `pandoc_version` if Pandoc is installed, otherwise a message is printed that pandoc is not installed.

get_parameters_meta	<i>get_parameters_meta</i>
---------------------	----------------------------

Description

get_parameters_meta

Usage

```
get_parameters_meta(xlsx_path, sheet_name = "nur Parameterliste")
```

Arguments

<code>xlsx_path</code>	path to EXCEL spreadsheet with parameter metadata
<code>sheet_name</code>	name of sheet containing metadata table (default: "nur Parameterliste")

Value

imported parameter metadata with cleaned columns names

get_phreeqc_data	<i>Get PhreeqC data</i>
------------------	-------------------------

Description

Get PhreeqC data

Usage

```
get_phreeqc_data(lab_bwb)
```

Arguments

lab_bwb imported BWB lab data as retrieved by [read_lab_bwb](#)

Value

tibble with columns solution, par_name_phreeqc (which are not empty or NA) and numeric_value

get_site_id	<i>Helper function: get_site_id</i>
-------------	-------------------------------------

Description

Helper function: get_site_id

Usage

```
get_site_id(string, pattern = "^[0-9]{1,4}")
```

Arguments

string vector with character strings
pattern pattern used for identifying site_id (default: "^[0-9]1,4")

Value

extracted site_id's from input string

import_labor	<i>import_labor</i>
--------------	---------------------

Description

import_labor

Usage

```
import_labor(files, export_dir, func = read_bwb_header2)
```

Arguments

files	vector with full paths of xlsx input files
export_dir	export directory
func	function to be used (default: read_bwb_header2)

Value

list with length equal to number of input files

order_measurement_chain_data	<i>Order Measurement Chain Data</i>
------------------------------	-------------------------------------

Description

Order Measurement Chain Data

Usage

```
order_measurement_chain_data(data)
```

Arguments

data	data frame as retrieved by read_measurementchains_data
------	--

Value

data, ordered by "parameter", "sensor_id", "datum_uhrzeit"

`plot_measurementchains`*Measurement Chains: plott*

Description

Measurement Chains: plott

Usage

```
plot_measurementchains(mc_data, para = "Leitfaehigkeit")
```

Arguments

<code>mc_data</code>	as retrieved by read_measurementchains_data
<code>para</code>	parameter to plot "Leitfaehigkeit" or "Temperatur" (default: Leitfaehigkeit)

Value

plot of selected measurement chain parameter

`plot_measurementchain_and_well_operation`*Plot measurementchain and well operation in combined plot*

Description

Plot measurementchain and well operation in combined plot

Usage

```
plot_measurementchain_and_well_operation(  
  mc_dat,  
  well_op_data_meta,  
  brunnen_nr = 9,  
  para = "Leitfaehigkeit",  
  y_label = "elektr. Leitfähigkeit (µS/cm)",  
  date_min = as.Date("2023-05-10"),  
  date_max = Sys.Date()  
)
```

Arguments

mc_dat	mc_dat
well_op_data_meta	well_op_data_meta
brunnen_nr	well id (default: 9)
para	parameter (either: "Leitfaehigkeit" or "Temperatur")
y_label	y label (default: "elektr. Leitfaehigkeit ($\mu\text{S}/\text{cm}$)")
date_min	minimum date for plotting (default: <code>as.Date("2023-05-10")</code>)
date_max	maximum date for plotting (default: <code>Sys.Date()</code>)

Value

combined plot

prepare_phreeqc_input *Prepare PhreeQC input*

Description

Prepare PhreeQC input

Usage

```
prepare_phreeqc_input(lab_bwb_phreeqc, title = "")
```

Arguments

lab_bwb_phreeqc	selected BWB lab data as retrieved by get_phreeqc_data
title	user defined title (default: "")

Value

data frame with input structure for `kwb.phreeqc`

```
print_datatype_info_if
```

Helper function: print_datatype_info_if

Description

Helper function: print_datatype_info_if

Usage

```
print_datatype_info_if(dbg, tbl_datatypes, columns_keep)
```

Arguments

dbg	dbg
tbl_datatypes	tbl_datatypes
columns_keep	columns_keep

```
read_bwb_data
```

Import: read_bwb_data

Description

wrapper around read_bwb_header2 and read_bwb_header1_meta

Usage

```
read_bwb_data(
  files,
  meta_pattern = "META",
  keep_pattern = column_pattern_gather_ignore(),
  site_id_pattern = "^[0-9]{1,4}",
  dbg = TRUE
)
```

Arguments

files	file path(s) to EXCEL spreadsheet
meta_pattern	(default: "META")
keep_pattern	(default: column_pattern_gather_ignore)
site_id_pattern	(default: "^[0-9]1,4")
dbg	debug (default: TRUE)

Value

data.table with imported xls(x) files

read_bwb_header1_meta *Import: read_bwb_header1_meta*

Description

Import: read_bwb_header1_meta

Usage

```
read_bwb_header1_meta(
  file,
  meta_pattern = "META",
  keep_pattern = column_pattern_gather_ignore(),
  dbg = FALSE
)
```

Arguments

file	path(s) to EXCEL spreadsheet
meta_pattern	meta_pattern default("META")
keep_pattern	keep_pattern (default: column_pattern_gather_ignore)
dbg	debug (default: FALSE)

Value

data.table with imported xls(x) files

read_bwb_header2 *Import: read_bwb_header2*

Description

Import: read_bwb_header2

Usage

```
read_bwb_header2(
  file,
  skip = 2,
  keep_pattern = column_pattern_gather_ignore(),
  site_id_pattern = "[0-9]{1,4}",
  dbg = TRUE
)
```

Arguments

file file path(s) to EXCEL spreadsheet
 skip number of rows to skip in each sheet (default: 2)
 keep_pattern (default: column_pattern_gather_ignore())
 site_id_pattern
 (default: "[0-9]1,4")
 dbg debug (default: TRUE)

Value

data.table with imported xls(x) files

read_bwb_header4 *Import: read_bwb_header4*

Description

Import: read_bwb_header4

Usage

```
read_bwb_header4(
  file,
  skip = 4,
  keep_pattern = column_pattern_gather_ignore(),
  site_id_pattern = "[0-9]{1,4}",
  dbg = TRUE
)
```

Arguments

file file path(s) to EXCEL spreadsheet
 skip number of rows to skip in each sheet (default: 4)
 keep_pattern (default: column_pattern_gather_ignore())
 site_id_pattern
 (default: "[0-9]1,4")
 dbg debug (default: TRUE)

Value

data.table with imported xls(x) files

read_emshoff91_ods	<i>Reads a Single EMSHOFF 91 ODS File into Tibble</i>
--------------------	---

Description

Reads a Single EMSHOFF 91 ODS File into Tibble

Usage

```
read_emshoff91_ods(emshoff91_import_selected)
```

Arguments

emshoff91_import_selected
a row as retrieved by [create_emshoff91_import](#)

Value

imports ods file into tibble

read_isotopes	<i>Read Isotopes</i>
---------------	----------------------

Description

Read Isotopes

Usage

```
read_isotopes(path)
```

Arguments

path path to Isotopes delim (field separator ";")

Value

imported isotopes data

read_lab_bwb	<i>Read Lab BWB</i>
--------------	---------------------

Description

Read Lab BWB

Usage

```
read_lab_bwb(path, sheet = "Analysen")
```

Arguments

path	path to file with lab BWB data
sheet	name of sheet containing analysis data. Default: "Analysen"

Value

cleaned data frame with master data and lab values for all samples but only for selected parameters (columns A-BA and HB-HC)

read_master_data	<i>Read Master Data</i>
------------------	-------------------------

Description

Read Master Data

Usage

```
read_master_data(path)
```

Arguments

path	path to file with master data (currently in file: lab BWB data)
------	---

Value

imported master data contained in sheet "Stammdaten"

read_measurementchains_data

Measurement Chains: read csv data from multiple files

Description

Measurement Chains: read csv data from multiple files

Usage

```
read_measurementchains_data(  
  csv_files,  
  datetime_installation = as_gmt_plus_one("2022-09-27 11:00:00"),  
  run_parallel = TRUE,  
  debug = FALSE  
)
```

Arguments

csv_files	vector of paths as retrieved by download_measurementchains_data
datetime_installation	datetime of first logger installation in well K10. Used to filter out older measurement data! Default: <code>kwb.geosalz::as_gmt_plus_one("2022-09-27 11:00:00")</code>
run_parallel	default: TRUE
debug	show debug messages (default: FALSE)

Value

data frame with imported data from csv files

Examples

```
## Not run:  
mc_files <- kwb.geosalz::get_measurementchains_files()  
target_directory <- tempdir()  
csv_files <- kwb.geosalz::download_measurementchains_data(  
  sftp_paths = mc_files$sftp_path,  
  target_directory  
)  
mc_data <- kwb.geosalz::read_measurementchains_data(csv_files)  
  
## End(Not run)
```

```
read_multiple_emshoff91_ods
```

Reads Multiple EMSHOFF 91 ODS Files into List

Description

Reads Multiple EMSHOFF 91 ODS Files into List

Usage

```
read_multiple_emshoff91_ods(emshoff91_import)
```

Arguments

```
emshoff91_import
```

a tibble as retrieved by [create_emshoff91_import](#)

Value

imports multiple ods files into a list of tibbles

Examples

```
## Not run:
ods_dir <- "<replace-with-path-to-files>/emshoff91/emshoff91/converted_ods"
ods_dir <- "C:/users/mrustl/Downloads/emshoff91/emshoff91/converted_ods"
emshoff91_import <- create_emshoff91_import(ods_dir)
read_multiple_emshoff91_ods(emshoff91_import)

## End(Not run)
```

```
replace_nanb_with_na
```

Replace n.a. (not available) and n.b. (not determined) from lab data with NA

Description

Replace n.a. (not available) and n.b. (not determined) from lab data with NA

Usage

```
replace_nanb_with_na(string)
```

Arguments

```
string
```

string

Value

string with NA instead of "n.a." or "n.b." (including 0-10 spaces between "n" and "a"/"b")

Examples

```
string <- c("19.2", "n.b.", "n. b.", "n. b.", "n.a.", "n. a.", "n. a.")
replace_nanb_with_na(string)
```

safe_office_folder *Helper function: safe_office_folder*

Description

Helper function: safe_office_folder

Usage

```
safe_office_folder(office_path = "C:/Program Files (x86)/Microsoft Office")
```

Arguments

office_path office folder path (default: "C:/Program Files (x86)/Microsoft Office")

Value

path of office folder (if existing)

stop_if_duplicated_samples_found
Helper function: stop if duplicated sample ids are found

Description

Helper function: stop if duplicated sample ids are found

Usage

```
stop_if_duplicated_samples_found(df, col_sampleid, path, sheet = "")
```

Arguments

df data frame with samples in wide format
col_sampleid column name of sample id
path path to file from which df was read (for information only)
sheet optional in case EXCEL is used (default: "")

Value

error in case duplicated samples were found

stop_on_missing_or_inform_on_extra_sheets

Helper function: stop_on_missing_or_inform_on_extra_sheets

Description

Helper function: stop_on_missing_or_inform_on_extra_sheets

Usage

stop_on_missing_or_inform_on_extra_sheets(has_site_id, file, sheets)

Arguments

has_site_id	has_site_id
file	file
sheets	sheets

to_full_metadata_2

Helper function: to_full_metadata2

Description

Helper function: to_full_metadata2

Usage

to_full_metadata_2(header, file, sheet)

Arguments

header	header
file	file
sheet	sheet

Value

data frame with metadata for header2 (EXCEL) files

to_full_metadata_4 *Helper function: to_full_metadata_4*

Description

Helper function: to_full_metadata_4

Usage

```
to_full_metadata_4(header, file, sheet)
```

Arguments

header	header
file	file
sheet	sheet

Value

data frame with metadata for header4 (EXCEL) files

write_measurementchains_data
Measurement Chains: write csv data

Description

Measurement Chains: write csv data

Usage

```
write_measurementchains_data(
  mc_data,
  target_directory,
  to_zip = FALSE,
  debug = FALSE
)
```

Arguments

mc_data	measurement chains data as retrieved by read_measurementchains_data
target_directory	target directory
to_zip	should data be zipped? (default: FALSE), if TRUE only a temporary csv file is created which will be subsequently zipped and deleted
debug	print debug messages (default: FALSE)

Value

writes csv data to path

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