Package 'washi'

September 8, 2023

```
Title Washington Soil Health Initiative Branding
```

Version 0.2.0

Description Create plots and tables in a consistent style with WaSHI (Washington Soil Health Initiative) branding. Use 'washi' to easily style your 'ggplot2' plots and 'flextable' tables.

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```
URL https://github.com/WA-Department-of-Agriculture/washi,
    https://wa-department-of-agriculture.github.io/washi/
```

BugReports https://github.com/WA-Department-of-Agriculture/washi/issues/

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example_data_long

Example data in long (tidy) format

Description

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A subset WaSHI State of the Soils Assessment dataset that has been anonymized. This dataset is tidied, so each measurement is in its own row.

Usage

```
example_data_long
```

Format

```
example_data_long A data frame with 1,800 rows and 14 columns::

year Year of sample
sampleId, producerId, fieldId Anonymized IDs
farmName, producerName, fieldName Anonymized names
longitude, latitude Truncated coordinates
texture Measured soil texture
measurement Measurement name with units
value Measurement result ...
```

Source

https://agr.wa.gov/departments/land-and-water/natural-resources/soil-health/state-of-the-soils

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example_data_wide

Example data in wide format

Description

A subset WaSHI State of the Soils Assessment dataset that has been anonymized. This dataset presents each sample in its own row, with columns for each measurement.

Usage

```
example_data_wide
```

Format

```
example_data_wide A data frame with 30 rows and 72:
columns:
year Year of sample
sampleId, producerId, fieldId Anonymized IDs
farmName, producerName, fieldName Anonymized names
longitude, latitude Truncated coordinates
texture Measured soil texture
other columns Column name includes measurement and units; value is the measurement results
```

Source

https://agr.wa.gov/departments/land-and-water/natural-resources/soil-health/state-of-the-soils

washi_flextable

WaSHI flextable style

Description

Creates a flextable in WaSHI's style.

Usage

```
washi_flextable(
  data,
  cols_bold = NULL,
  header_font = "Lato",
  body_font = "Poppins",
  header_font_color = "white",
  header_bg_color = washi_pal[["standard"]][["green"]],
  border_color = washi_pal[["standard"]][["tan"]]
)
```

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Arguments

data Dataframe for the table.

cols_bold Numeric indices of columns to bold. Defaults to NULL.

header_font Font family for header text. Defaults to "Lato".

body_font Font family for table body text. Defaults to "Poppins".

header_font_color

Hexcode color for header font. Defaults to white.

header_bg_color

Hexcode color for header background. Defaults to WaSHI green.

border_color Hexcode color for horizontal borders. Defaults to WaSHI tan.

Value

A flextable formatted in WaSHI's style.

Examples

```
subset(
  example_data_wide,
  select = c(
    "sampleId",
    "county",
    "crop",
    "totalN_%",
    "totalC_%"
  )
) |>
  washi_flextable(cols_bold = 1)
```

washi_install_fonts

Install Lato and Poppins fonts on your system

Description

Opens the font files to install on your computer.

Usage

```
washi_install_fonts()
```

Value

Opens a directory containing font files.

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washi_pal

WaSHI color palettes

Description

Color palettes are stored in a named list.

Usage

```
washi_pal
```

Format

An object of class list of length 11.

Value

List of available washi color palettes

See Also

Other color palette functions: washi_pal_setup(), washi_pal_view()

Examples

```
# List names of available palettes
names(washi_pal)

# Get hex codes from a palette using dollar `$name` or
# double bracket

# `[["name"]]` operators for extracting list elements
washi_pal$standard

washi_pal[["green_gradient"]]

# Extract a color from the standard WaSHI palette
washi_pal[["standard"]][["green"]]
```

washi_pal_setup

Setup a color palette

Description

Choose desired number of colors and whether the colors are reversed.

Usage

```
washi_pal_setup(palette = "standard", n, reverse = FALSE)
```

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Arguments

palette Character name of palette in washi_pal. See names(washi_pal) for a list of

available palettes.

n Number of colors in palette.

reverse Boolean indicating whether the palette should be reversed. Default is FALSE.

Value

A vector of color hex codes.

See Also

Other color palette functions: washi_pal_view(), washi_pal

Examples

```
washi_pal_setup("color_blind")
washi_pal_setup("green_gradient", 12)
```

washi_pal_view

View a WaSHI palette

Description

Show the colors within a palette in a plot.

Usage

```
washi_pal_view(palette = "color_blind", n, reverse = FALSE)
```

Arguments

palette Character name of palette in washi_pal. See names(washi_pal) for a list of

available palettes.

n Number of colors in palette.

reverse Boolean indicating whether the palette should be reversed. Default is FALSE.

Value

A plot with each color displayed.

See Also

Other color palette functions: washi_pal_setup(), washi_pal

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Examples

```
washi_pal_view("standard")
washi_pal_view("color_blind")
washi_pal_view("blue_gradient", 4, reverse = TRUE)
```

washi_scale

Scales for plotting with WaSHI palettes

Description

Provides compatibility with ggplot2.

Usage

```
washi_scale(
  palette = "color_blind",
  aesthetics = c("color", "fill"),
  alpha = 1,
  reverse = FALSE,
  discrete = TRUE,
  ...
)
```

Arguments

palette	Character name of palette in washi_pal. See names(washi_pal) for a list of available palettes.
aesthetics	Character string or vector of character strings listing the name(s) of the aesthetic(s) that this scale works with. Defaults to c("color", "fill"), which applies the palette to both the color and fill aesthetics at the same time.
alpha	Numeric transparency level of the color from 0 to 1. Default is 1 (not transparent).
reverse	Boolean indicating whether the palette should be reversed. Default is FALSE.
discrete	Boolean indicating whether color aesthetic is discrete or not. Default is TRUE.
•••	Additional arguments passed to discrete_scale() or scale_color_gradientn(), used respectively when discrete is TRUE or FALSE

Value

A ScaleContinuous or ScaleDiscrete object that can be added to a ggplot object.

See Also

```
Other ggplot2 functions: washi_theme()
```

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Examples

```
library(ggplot2)

# Discrete scale
example_data_wide |>
    subset(crop %in% c("Apple", "Cherry", "Potato")) |>
    ggplot(aes(x = pH, y = Mn_mg.kg, color = crop)) +
    geom_point(size = 2.5) +
    theme_minimal() +
    washi_scale()

# Continuous scale
example_data_wide |>
    ggplot(aes(x = `totalC_%`, y = poxC_mg.kg, color = poxC_mg.kg)) +
    geom_point(size = 2.5) +
    theme_minimal() +
    washi_scale("green_gradient", reverse = TRUE, discrete = FALSE)
```

washi_theme

Create standard WaSHI plots

Description

All changed defaults from this function can be overridden by another call to ggplot2::theme() with the desired changes.

Usage

```
washi_theme(
  header_font = "Lato Black",
  header_color = "#151414",
  body_font = "Poppins",
  body_color = "#151414",
  text_scale = 1,
  legend_position = "top",
  facet_space = 2,
  color_gridline = washi_pal[["standard"]][["tan"]],
  gridline_y = TRUE,
  gridline_x = TRUE,
  ...
)
```

Arguments

header_font Font family for title and subtitle. Defaults to "Lato Black".

header_color Font color for title and subtitle. Defaults to almost black.

body_font Font family for all other text Defaults to "Poppins".

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body_color Font color for all other text Defaults to almost black. text_scale Scalar that will grow/shrink all text defined within. legend_position Position of legend ("none", "left", "right", "bottom", "top", or two-element numeric vector). Defaults to "top". facet_space Controls how far apart facets are from each other. color_gridline Gridline color. Defaults to WaSHI tan. Boolean indicating whether major gridlines are displayed for the y axis. Default gridline_y is TRUE. Boolean indicating whether major gridlines are displayed for the x axis. Default gridline_x is TRUE. Pass any parameters from theme that are not already defined within. . . .

Value

ggplot2 object

Source

```
Adapted from glitr::si_style().
```

See Also

Other ggplot2 functions: washi_scale()

Examples

```
# NOTE: These examples do not use Poppins or Lato in order to pass
# automated checks on computers without these fonts installed.
library(ggplot2)
# Single geom_point plot
example_data_wide |>
 subset(crop %in% c("Apple", "Cherry", "Potato")) |>
 ggplot(aes(x = pH, y = Mn_mg.kg, color = crop)) +
 labs(
    title = "Scatter plot of pH and Mn (mg/kg)",
   subtitle = "Example with geom_point().",
   caption = "This is a caption."
 ) +
 geom_point(size = 2.5) +
 washi_theme(
   header_font = "sans",
   body_font = "sans"
 washi_scale()
# Bar plot
```

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```
if (requireNamespace("forcats")) {
 example_data_wide |>
   ggplot(aes(x = forcats::fct_rev(forcats::fct_infreq(crop)))) +
    geom_bar(fill = washi_pal[["standard"]][["blue"]]) +
   geom_text(
     aes(
       y = after_stat(count),
       label = after_stat(count)
     ),
     stat = "count",
     hjust = 2.5,
     color = "white"
    # Flip coordinates to accommodate long crop names
    coord_flip() +
   labs(
     title = "Number of samples in each crop",
     subtitle = "Example plot with geom_bar() without gridlines.",
     y = NULL,
     x = NULL
   ) +
    # Turn gridlines off
   washi_theme(
     gridline_y = FALSE,
     gridline_x = FALSE,
     header_font = "sans",
     body_font = "sans"
   ) +
    # Remove x-axis
    theme(axis.text.x = element_blank())
}
# Facetted geom_density plots
example_data_long |>
 subset(measurement %in% c("totalC_%", "poxC_mg.kg") &
    !texture == "Loamy Sand") |>
 ggplot(aes(x = value, fill = texture, color = texture)) +
 labs(
   title = "Distribution of POXC (mg/kg) and Total C (%)",
    subtitle = "Example with geom_density() and facet_wrap()."
 geom\_density(alpha = 0.4) +
 facet_wrap(. ~ measurement, scales = "free") +
 washi_theme(
   legend_position = "bottom",
   header_font = "sans",
   body_font = "sans"
 ) +
 washi_scale() +
 xlab(NULL) +
 guides(col = guide_legend(nrow = 2, byrow = TRUE))
```

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