

# Package ‘washi’

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**Title** Washington Soil Health Initiative Branding

**Version** 0.2.1

**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/>

**Depends** R (>= 4.1.0)

**Imports** cli, flextable, ggplot2, graphics, grDevices, officer, scales,  
systemfonts

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**Config/testthat/edition** 3

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example_data_long	<i>Example data in long (tidy) format</i>
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### Description

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

**sample\_id, producer\_id, field\_id** Anonymized IDs

**farm\_name, producer\_name, field\_name** 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	<i>Example data in wide format</i>
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### 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

**sample\_id, producer\_id, field\_id** Anonymized IDs

**farm\_name, producer\_name, field\_name** 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>

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washi_flextable	<i>WaSHI flextable style</i>
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### 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"]]  
)
```

**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(
    "sample_id",
    "county",
    "crop",
    "total_n_percent",
    "total_c_percent"
  )
) |>
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.

---

washi_pal	<i>WaSHI color palettes</i>
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**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	<i>Setup a color palette</i>
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---

**Description**

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

**Usage**

```
washi_pal_setup(palette = "standard", 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 vector of color hex codes.

**See Also**

Other color palette functions: [washi\\_pal](#), [washi\\_pal\\_view\(\)](#)

**Examples**

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

---

washi_pal_view	<i>View a WaSHI palette</i>
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**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](#), [washi\\_pal\\_setup\(\)](#)

**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\(\)](#)

**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 = total_c_percent, 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".

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.
gridline_y	Boolean indicating whether major gridlines are displayed for the y axis. Default is TRUE.
gridline_x	Boolean indicating whether major gridlines are displayed for the x axis. Default 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**

```
library(ggplot2)
# NOTE: These examples do not use Poppins or Lato in order to pass
# automated checks on computers without these fonts installed.

# 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
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("total_c_percent", "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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