

# Package ‘plotcli’

April 24, 2024

**Title** Command Line Interface Plotting

**Version** 0.1.0

**Date** 2024-04-01

**Description** The 'plotcli' package provides terminal-based plotting in R.

It supports colored scatter plots, line plots, bar plots, and box plots. The package allows users to customize plot appearance, add titles, labels, ticks, and legends, and output the plot as a text-based visualization.

**License** LGPL-3

**URL** <https://github.com/cheuerde/plotcli>

**Encoding** UTF-8

**VignetteBuilder** knitr

**Depends** R6, ggplot2

**Imports** utils, crayon, knitr, rmarkdown, stringr, rlang

**Suggests** testthat (>= 3.0.0)

**NeedsCompilation** yes

**RoxygenNote** 7.2.3

**Config/testthat/edition** 3

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**Repository** CRAN

**Date/Publication** 2024-04-24 14:10:07 UTC

## R topics documented:

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| +.plotcli | <i>Overload the "+" operator for plotcli objects</i> |
|-----------|--|

---

### Description

This function overloads the "+" operator to merge two plotcli objects.

### Usage

```
## S3 method for class 'plotcli'
plot1 + plot2
```

### Arguments

|       |                                |
|-------|--------------------------------|
| plot1 | A plotcli object to be merged. |
| plot2 | A plotcli object to be merged. |

### Value

A new plotcli object containing the combined data from both objects.

---

|           |                                   |
|-----------|-----------------------------------|
| bresenham | <i>Bresenham's line algorithm</i> |
|-----------|-----------------------------------|

---

**Description**

This function generates a list of points that form a line between two given points using Bresenham's line algorithm.

**Usage**

```
bresenham(x0, y0, x1, y1)
```

**Arguments**

|    |   |
|----|---|
| x0 | The x-coordinate of the starting point. |
| y0 | The y-coordinate of the starting point. |
| x1 | The x-coordinate of the ending point.   |
| y1 | The y-coordinate of the ending point.   |

**Value**

A list of points that form a line between the two given points.

**Examples**

```
bresenham(0, 0, 5, 5)
bresenham(0, 0, -5, -5)
```

---

|                 |                          |
|-----------------|--------------------------|
| cat_plot_matrix | <i>Print plot matrix</i> |
|-----------------|--------------------------|

---

**Description**

This function prints a plot matrix to the console.

**Usage**

```
cat_plot_matrix(plot_matrix)
```

**Arguments**

|             |                                |
|-------------|--------------------------------|
| plot_matrix | The plot matrix to be printed. |
|-------------|--------------------------------|

**Examples**

```
cat_plot_matrix(matrix(c("a", "b", "c", "d"), nrow = 2, ncol = 2))
```

---

`cbind.plotcli`*Generic function for combining plotcli objects horizontally*

---

**Description**

Generic function for combining plotcli objects horizontally

**Usage**

```
## S3 method for class 'plotcli'  
cbind(..., deparse.level = 1)
```

**Arguments**

`...` plotcli objects to be combined.  
`deparse.level` The deparsing level for the arguments.

**Value**

A combined plot matrix.

---

`cbind_plots`*Combine plot matrices horizontally*

---

**Description**

This function combines multiple plot matrices horizontally, centering them vertically.

**Usage**

```
cbind_plots(...)
```

**Arguments**

`...` A list of plot matrices to be combined.

**Value**

A combined plot matrix.

---

|                   |   |
|-------------------|---|
| format_four_chars | <i>Format number to four characters</i> |
|-------------------|---|

---

**Description**

This function formats a number to a string of exactly four characters.

**Usage**

```
format_four_chars(num)
```

**Arguments**

|     |                             |
|-----|-----------------------------|
| num | The number to be formatted. |
|-----|-----------------------------|

**Value**

A string representation of the number with exactly four characters.

**Examples**

```
format_four_chars(123)  
format_four_chars(-12.34)
```

---

|                 |  |
|-----------------|--|
| get_data_subset | <i>Get data subset for a specific geom</i> |
|-----------------|--|

---

**Description**

This function returns a subset of the data for a specific geom.

**Usage**

```
get_data_subset(geom_name, data, aes, p_build)
```

**Arguments**

|           |   |
|-----------|---|
| geom_name | The name of the geom for which the data subset is needed. |
| data      | The data to be subsetted.                                 |
| aes       | The aesthetic mappings for the geom.                      |
| p_build   | The ggplot build object.                                  |

**Value**

A list containing the data subset for the specified geom.

---

|                 |                            |
|-----------------|----------------------------|
| get_term_colors | <i>Get terminal colors</i> |
|-----------------|----------------------------|

---

**Description**

This function returns a vector of terminal colors.

**Usage**

```
get_term_colors(n = NULL)
```

**Arguments**

`n` The number of colors to return.

**Value**

A vector of terminal colors.

**Examples**

```
get_term_colors(5)
get_term_colors(10)
```

---

|           |  |
|-----------|--|
| ggplotcli | <i>ggplotcli - Render ggplot objects in the terminal</i> |
|-----------|--|

---

**Description**

This function takes a ggplot object and renders it in the terminal using ASCII or Braille characters.

**Usage**

```
ggplotcli(ggplot_obj, plot_width = 80, plot_height = 40, braille = TRUE)
```

**Arguments**

`ggplot_obj` A ggplot object to be rendered in the terminal.  
`plot_width` Width of the terminal plot in characters (default: 80).  
`plot_height` Height of the terminal plot in characters (default: 40).  
`braille` Use Braille characters for higher resolution (default: TRUE).

**Value**

A TerminalPlot object.

---

|            |  |
|------------|--|
| is_braille | <i>Check if a character is a Braille character</i> |
|------------|--|

---

**Description**

This function checks if a given character is a Braille character.

**Usage**

```
is_braille(char)
```

**Arguments**

|      |                              |
|------|------------------------------|
| char | The character to be checked. |
|------|------------------------------|

**Value**

A boolean value indicating whether the character is a Braille character or not.

**Examples**

```
is_braille("A")
```

---

|              |                          |
|--------------|--------------------------|
| make_colored | <i>Make colored text</i> |
|--------------|--------------------------|

---

**Description**

This function applies a specified color to a given text string.

**Usage**

```
make_colored(x, color = NULL)
```

**Arguments**

|       |  |
|-------|--|
| x     | The text string to be colored.   |
| color | The color to be applied to the text. If NULL, the color codes will be removed. |

**Value**

A colored text string or a text string with color codes removed.

**Examples**

```
make_colored("Hello, world!", "blue")  
make_colored("Hello, world!", NULL)
```

---

|                   |                          |
|-------------------|--------------------------|
| make_unique_names | <i>Make unique names</i> |
|-------------------|--------------------------|

---

**Description**

This function takes a vector of names and ensures that each name is unique by appending a number if necessary.

**Usage**

```
make_unique_names(names)
```

**Arguments**

|       |                              |
|-------|------------------------------|
| names | A character vector of names. |
|-------|------------------------------|

**Value**

A character vector of unique names.

**Examples**

```
make_unique_names(c("apple", "apple", "banana", "apple"))
```

---

|                |                       |
|----------------|-----------------------|
| normalize_data | <i>Normalize data</i> |
|----------------|-----------------------|

---

**Description**

This function normalizes the given data to a specified plot range.

**Usage**

```
normalize_data(data, data_min, data_max, plot_range)
```

**Arguments**

|            |                                     |
|------------|-------------------------------------|
| data       | The data to be normalized.          |
| data_min   | The minimum value of the data.      |
| data_max   | The maximum value of the data.      |
| plot_range | The range to normalize the data to. |

**Value**

The normalized data.



**Examples**

```
normalize_data(c(1, 2, 3, 4, 5), 1, 5, 10)
normalize_data(c(10, 20, 30, 40, 50), 10, 50, 100)
```

pclub

*Short version of plotcli\_bar***Description**

Short version of plotcli\_bar function.

**Usage**

```
pclub(
  y,
  x = NULL,
  plot_width = getOption("plotcli.plot_width", 80),
  plot_height = getOption("plotcli.plot_height", 40),
  x_label = "x",
  y_label = "y",
  color = NULL,
  braille = getOption("plotcli.braille", TRUE),
  name = "barplot",
  ...
)
```

**Arguments**

|             |  |
|-------------|--|
| y           | A numeric vector of values                                 |
| x           | A vector of categories                                     |
| plot_width  | Width of the plot (default: 80)                            |
| plot_height | Height of the plot (default: 40)                           |
| x_label     | Label for the x-axis (default: "x")                        |
| y_label     | Label for the y-axis (default: "y")                        |
| color       | Color of the plot elements (default: NULL)                 |
| braille     | Use Braille characters for the plot (default: TRUE)        |
| name        | Name of the plot element (default: "barplot")              |
| ...         | Additional arguments passed to the plotcli\$new() function |

**Examples**

```
x <- 1:5
y <- c(10, 15, 8, 12, 6)
pclub(x, y)
```

---

pplibx

*Short version of plotcli\_box*

---

## Description

Short version of plotcli\_box function.

## Usage

```
pplibx(  
  y,  
  plot_width = getOption("plotcli.plot_width", 80),  
  plot_height = getOption("plotcli.plot_height", 40),  
  x_label = "x",  
  y_label = "y",  
  color = NULL,  
  braille = getOption("plotcli.braille", TRUE),  
  name = "boxplot",  
  ...  
)
```

## Arguments

|             |  |
|-------------|--|
| y           | A list of numeric vectors of values                        |
| plot_width  | Width of the plot (default: 80)                            |
| plot_height | Height of the plot (default: 40)                           |
| x_label     | Label for the x-axis (default: "x")                        |
| y_label     | Label for the y-axis (default: "y")                        |
| color       | Color of the plot elements (default: NULL)                 |
| braille     | Use Braille characters for the plot (default: TRUE)        |
| name        | Name of the plot element (default: "boxplot")              |
| ...         | Additional arguments passed to the plotcli\$new() function |
| x           | A vector of categories                                     |

## Examples

```
y <- rnorm(50, mean = 0)  
pplib(y)
```

---

pclid *Short version of plotcli\_density*

---

## Description

Short version of plotcli\_density function.

## Usage

```
pclid(  
  x,  
  plot_width = getOption("plotcli.plot_width", 80),  
  plot_height = getOption("plotcli.plot_height", 40),  
  x_label = "x",  
  y_label = "Density",  
  color = NULL,  
  braille = getOption("plotcli.braille", TRUE),  
  name = "density",  
  ...  
)
```

## Arguments

|             |  |
|-------------|--|
| x           | A numeric vector of values                                 |
| plot_width  | Width of the plot (default: 80)                            |
| plot_height | Height of the plot (default: 40)                           |
| x_label     | Label for the x-axis (default: "x")                        |
| y_label     | Label for the y-axis (default: "Density")                  |
| color       | Color of the plot elements (default: NULL)                 |
| braille     | Use Braille characters for the plot (default: TRUE)        |
| name        | Name of the plot element (default: "density")              |
| ...         | Additional arguments passed to the plotcli\$new() function |

## Examples

```
x <- rnorm(100)  
pclid(x)
```

---

pcli

*Short version of plotcli\_histogram*

---

## Description

Short version of plotcli\_histogram function.

## Usage

```
pcli(
  x,
  plot_width = getOption("plotcli.plot_width", 80),
  plot_height = getOption("plotcli.plot_height", 40),
  x_label = "x",
  y_label = "Frequency",
  color = NULL,
  braille = getOption("plotcli.braille", TRUE),
  bin_width = NULL,
  ylim = NULL,
  name = "histogram",
  ...
)
```

## Arguments

|             |  |
|-------------|--|
| x           | A numeric vector of values                                 |
| plot_width  | Width of the plot (default: 80)                            |
| plot_height | Height of the plot (default: 40)                           |
| x_label     | Label for the x-axis (default: "x")                        |
| y_label     | Label for the y-axis (default: "Frequency")                |
| color       | Color of the plot elements (default: NULL)                 |
| braille     | Use Braille characters for the plot (default: TRUE)        |
| bin_width   | Width of the bins (default: NULL)                          |
| ylim        | y limits (default: NULL)                                   |
| name        | Name of the plot element (default: "histogram")            |
| ...         | Additional arguments passed to the plotcli\$new() function |

## Examples

```
x <- rnorm(100)
pcli(x)
```

---

`pclil`*Short version of plotcli\_line*

---

**Description**

Short version of plotcli\_line function.

**Usage**

```
pclil(  
  y,  
  x = NULL,  
  plot_width = getOption("plotcli.plot_width", 80),  
  plot_height = getOption("plotcli.plot_height", 40),  
  x_label = "x",  
  y_label = "y",  
  color = NULL,  
  braille = getOption("plotcli.braille", TRUE),  
  name = "line",  
  ...  
)
```

**Arguments**

|                          |  |
|--------------------------|--|
| <code>y</code>           | A numeric vector of y values                               |
| <code>x</code>           | A numeric vector of x values                               |
| <code>plot_width</code>  | Width of the plot (default: 80)                            |
| <code>plot_height</code> | Height of the plot (default: 40)                           |
| <code>x_label</code>     | Label for the x-axis (default: "x")                        |
| <code>y_label</code>     | Label for the y-axis (default: "y")                        |
| <code>color</code>       | Color of the plot elements (default: NULL)                 |
| <code>braille</code>     | Use Braille characters for the plot (default: TRUE)        |
| <code>name</code>        | Name of the plot element (default: "line")                 |
| <code>...</code>         | Additional arguments passed to the plotcli\$new() function |

**Examples**

```
x <- 1:10  
y <- x^2  
pclil(x, y)
```

---

pclis

*Short version of plotcli\_scatter*

---

## Description

Short version of plotcli\_scatter function.

## Usage

```
pclis(  
  y,  
  x = NULL,  
  plot_width = getOption("plotcli.plot_width", 80),  
  plot_height = getOption("plotcli.plot_height", 40),  
  x_label = "x",  
  y_label = "y",  
  color = NULL,  
  braille = getOption("plotcli.braille", TRUE),  
  name = "scatter",  
  ...  
)
```

## Arguments

|             |  |
|-------------|--|
| y           | A numeric vector of y values                               |
| x           | A numeric vector of x values                               |
| plot_width  | Width of the plot (default: 80)                            |
| plot_height | Height of the plot (default: 40)                           |
| x_label     | Label for the x-axis (default: "x")                        |
| y_label     | Label for the y-axis (default: "y")                        |
| color       | Color of the plot elements (default: NULL)                 |
| braille     | Use Braille characters for the plot (default: TRUE)        |
| name        | Name of the plot element (default: "scatter")              |
| ...         | Additional arguments passed to the plotcli\$new() function |

## Examples

```
x <- rnorm(100)  
y <- rnorm(100)  
pclis(x, y)
```

---

plotcli

*plotcli R6 Class*

---

## Description

plotcli R6 Class  
plotcli R6 Class

## Details

This class provides a set of methods to create and customize command-line plots using R6. It supports various plot types, such as scatter, line, bar, and box plots, and allows customization of plot elements, such as title, axis labels, ticks, and legend.

## Usage

```
plotcli <- plotcli$new()  
plotcli$add_data(data)  
plotcli$print_plot()
```

## Methods

**initialize()** Initializes the PlotCLI object with parameters.

**initialize\_plot\_matrix()** Initializes the plot matrix with the plot canvas.

**print()** Default print method for PlotCLI object.

**add\_row()** Adds a single row to the plot matrix.

**add\_col()** Adds a single column to the plot matrix.

**add\_borders()** Adds borders around the plot canvas.

**add\_row\_col\_index()** Adds row and column index to the plot matrix.

**add\_title()** Adds a title to the plot matrix.

**add\_y\_ticks()** Adds y-axis tick labels to the plot matrix.

**add\_y\_label()** Adds a y-axis label to the plot matrix.

**add\_x\_ticks()** Adds x-axis tick labels to the plot matrix.

**add\_x\_label()** Adds an x-axis label to the plot matrix.

**add\_legend()** Adds a legend to the plot matrix.

**add\_data()** Adds data to the object.

**get\_min\_max()** Gets minimum and maximum values for x and y.

**remove\_out\_of\_range\_data()** Removes out of range data points if xlim and ylim were given.

**draw\_scatter\_plot()** Draws a scatter plot on the plot canvas.

**draw\_line\_plot()** Draws a line plot on the plot canvas.

**draw\_barplot()** Draws a bar plot on the plot canvas.

**draw\_barplot\_braille()** Draws a bar plot with braille characters on the plot canvas.

**draw\_boxplot()** Draws a box plot on the plot canvas.

**print\_plot()** Assembles all plot elements and prints the plot to the console.

**Public fields**

plot\_width The width of the plot  
plot\_height The height of the plot  
plot\_canvas The canvas for drawing the plot  
plot\_matrix The matrix containing the entire plot, including borders, labels, and title  
data A list containing the data sets to be plotted  
title The title of the plot  
x\_label The label for the x-axis  
y\_label The label for the y-axis  
ylim The limits for the y-axis  
xlim The limits for the x-axis  
x\_min The minimum value of the x-axis  
x\_max The maximum value of the x-axis  
y\_min The minimum value of the y-axis  
y\_max The maximum value of the y-axis  
plot\_matrix\_canvas\_row\_start The starting row of the plot canvas within the plot matrix  
plot\_matrix\_canvas\_col\_start The starting column of the plot canvas within the plot matrix  
is\_boxplot A logical value indicating if the plot is a boxplot  
draw\_legend A logical value indicating if the legend should be drawn

**Methods****Public methods:**

- `plotcli$new()`
- `plotcli$initialize_plot_matrix()`
- `plotcli$print()`
- `plotcli$add_row()`
- `plotcli$add_col()`
- `plotcli$add_borders()`
- `plotcli$add_row_col_index()`
- `plotcli$add_title()`
- `plotcli$add_y_ticks()`
- `plotcli$add_y_label()`
- `plotcli$add_x_ticks()`
- `plotcli$add_x_label()`
- `plotcli$add_legend()`
- `plotcli$add_data()`
- `plotcli$get_min_max()`
- `plotcli$remove_out_of_range_data()`
- `plotcli$draw_scatter_plot()`



- `plotcli$draw_line_plot()`
- `plotcli$draw_barplot()`
- `plotcli$draw_barplot_braille()`
- `plotcli$draw_boxplot()`
- `plotcli$draw_colors()`
- `plotcli$draw_plot()`
- `plotcli$make_plot_matrix()`
- `plotcli$export_plot_matrix()`
- `plotcli$print_plot()`
- `plotcli$merge()`
- `plotcli$clone()`

**Method** `new()`: Initialize object

*Usage:*

```
plotcli$new(
  plot_width = 60,
  plot_height = 20,
  x_label = "x",
  y_label = "y",
  ylim = NULL,
  xlim = NULL,
  title = NULL,
  is_boxplot = FALSE,
  draw_legend = TRUE
)
```

*Arguments:*

`plot_width` integer, width of the plot canvas

`plot_height` integer, height of the plot canvas

`x_label` character, label for the x-axis

`y_label` character, label for the y-axis

`ylim` numeric vector, limits for the y-axis

`xlim` numeric vector, limits for the x-axis

`title` character, title of the plot

`is_boxplot` logical, whether the plot is a boxplot

`draw_legend` logical, whether to draw the legend This function initializes the plot matrix based on the plot canvas.

**Method** `initialize_plot_matrix()`: Initialize the plot matrix

*Usage:*

```
plotcli$initialize_plot_matrix()
```

*Arguments:*

`plot_width` The width of the plot

`plot_height` The height of the plot

*Returns:* A plot matrix object

**Method print():** Default print method for plotcli object

*Usage:*

```
plotcli$print(...)
```

*Arguments:*

... Additional arguments passed to the print method

*Returns:* The plotcli object, invisibly

**Method add\_row():** Add a single row to the plot matrix

*Usage:*

```
plotcli$add_row(bottom = FALSE)
```

*Arguments:*

bottom logical, if TRUE, add row to the bottom of the matrix, otherwise add to the top (default: FALSE)

**Method add\_col():** Add a single column to the plot matrix

*Usage:*

```
plotcli$add_col()
```

**Method add\_borders():** Add borders to the plot matrix

*Usage:*

```
plotcli$add_borders()
```

**Method add\_row\_col\_index():** Add row and column index to the plot matrix Add title to the plot matrix

*Usage:*

```
plotcli$add_row_col_index()
```

**Method add\_title():**

*Usage:*

```
plotcli$add_title()
```

*Arguments:*

title character, title of the plot Add y-ticks label to the plot matrix

**Method add\_y\_ticks():**

*Usage:*

```
plotcli$add_y_ticks(n_ticks = 5)
```

*Arguments:*

n\_ticks numeric, number of ticks Add y-axis label to the plot matrix

**Method add\_y\_label():** Add a y-axis label to the plot matrix

*Usage:*

```
plotcli$add_y_label(y_label = self$y_label)
```

*Arguments:*

y\_label character, the y-axis label to be added Add x-ticks label to the plot matrix

**Method** add\_x\_ticks():

*Usage:*

```
plotcli$add_x_ticks(n_ticks = 5)
```

*Arguments:*

n\_ticks numeric, number of ticks Add x-axis label to the plot matrix

**Method** add\_x\_label(): Add x-axis label to the plot matrix

*Usage:*

```
plotcli$add_x_label(x_label = self$x_label)
```

*Arguments:*

x\_label x label Add legend to the plot matrix

**Method** add\_legend(): Add legend to the plot matrix Add data to the object.

*Usage:*

```
plotcli$add_legend()
```

**Method** add\_data():

*Usage:*

```
plotcli$add_data(data)
```

*Arguments:*

data list, list with elements: x, y, type, color, braille, name Get minimum and maximum values for x and y

**Method** get\_min\_max(): Calculate the minimum and maximum values for x and y Function to remove out of range data points if xlim and ylim were given

*Usage:*

```
plotcli$get_min_max()
```

**Method** remove\_out\_of\_range\_data(): Remove data points that are outside the specified xlim and ylim Draw a scatter plot to the plot canvas.

*Usage:*

```
plotcli$remove_out_of_range_data()
```

**Method** draw\_scatter\_plot(): Draw a scatter plot of the specified data set on the plot canvas.

*Usage:*

```
plotcli$draw_scatter_plot(set_idx)
```

*Arguments:*

set\_idx numeric, the data element index to be drawn Draw a line plot to the plot canvas.

**Method** draw\_line\_plot():

*Usage:*

```
plotcli$draw_line_plot(set_idx)
```

*Arguments:*

set\_idx numeric, the data element index to be drawn Draw a barplot to the plot canvas.

**Method** draw\_barplot():*Usage:*

```
plotcli$draw_barplot(set_idx)
```

*Arguments:*

set\_idx numeric, the data element index to be drawn Draw a barplot to the plot canvas with braille characters.

**Method** draw\_barplot\_braille():*Usage:*

```
plotcli$draw_barplot_braille(set_idx)
```

*Arguments:*

set\_idx numeric, the data element index to be drawn Draw a boxplot to the plot canvas.

**Method** draw\_boxplot():*Usage:*

```
plotcli$draw_boxplot(set_idx)
```

*Arguments:*

set\_idx numeric, the data element index to be drawn Draw colors to the canvas

**Method** draw\_colors(): In the draw\_ functions we have been keeping track of the locations of the colored matrix elements. These are now being colored. Draw the different plots types from all data elements to the canvas

*Usage:*

```
plotcli$draw_colors()
```

**Method** draw\_plot(): This function iterates through all data elements and calls the appropriate draw\_ function based on the plot type (scatter, line, boxplot, or barplot). Make plot matrix: assembles all plot elements (canvas + borders + title + axes + legend)

*Usage:*

```
plotcli$draw_plot()
```

**Method** make\_plot\_matrix(): This function assembles all plot elements (canvas + borders + title + axes + legend) and creates the final plot matrix. Export plot matrix

*Usage:*

```
plotcli$make_plot_matrix()
```

**Method** export\_plot\_matrix(): This function exports the plot matrix.

*Usage:*

```
plotcli$export_plot_matrix()
```

*Returns:* The plot matrix. Main plotting function: assembles all plot elements (canvas + borders + title + axes + legend) and prints the plot by 'cat'ing the plot matrix to the console.

**Method** `print_plot()`: This function assembles all plot elements (canvas + borders + title + axes + legend) and prints the final plot by 'cat'ing the plot matrix to the console. Merge two plotcli objects

This method combines the data from two plotcli objects into a single plotcli object. It takes the maximum of the `plot_width` and `plot_height`, combines the titles, and sets the `xlim` and `ylim` to the minimum and maximum values of both objects.

*Usage:*

```
plotcli$print_plot()
```

**Method** `merge()`:

*Usage:*

```
plotcli$merge(other)
```

*Arguments:*

`other` A plotcli object to be merged with the current object.

*Returns:* A new plotcli object containing the combined data from both objects.

**Method** `clone()`: The objects of this class are cloneable with this method.

*Usage:*

```
plotcli$clone(deep = FALSE)
```

*Arguments:*

`deep` Whether to make a deep clone.

## Examples

```
# Create a new plotcli object
plotcli <- plotcli$new()

# Add data for a scatter plot
plotcli$add_data(list(x = 1:10, y = rnorm(10), type = "scatter", color = "red"))

# Print the plot
plotcli$print_plot()
```

---

plotcli\_bar

*Bar plot using plotcli*

---

## Description

Create a bar plot using plotcli. Short alias: `pclb`.

**Usage**

```

plotcli_bar(
  y,
  x = NULL,
  plot_width = getOption("plotcli.plot_width", 80),
  plot_height = getOption("plotcli.plot_height", 40),
  x_label = "x",
  y_label = "y",
  color = NULL,
  braille = getOption("plotcli.braille", TRUE),
  name = "barplot",
  ...
)

```

**Arguments**

|                          |   |
|--------------------------|---|
| <code>y</code>           | A numeric vector of values  |
| <code>x</code>           | A vector of categories  |
| <code>plot_width</code>  | Width of the plot (default: 80)   |
| <code>plot_height</code> | Height of the plot (default: 40)  |
| <code>x_label</code>     | Label for the x-axis (default: "x")                                     |
| <code>y_label</code>     | Label for the y-axis (default: "y")                                     |
| <code>color</code>       | Color of the plot elements (default: NULL)                              |
| <code>braille</code>     | Use Braille characters for the plot (default: TRUE)                     |
| <code>name</code>        | Name of the plot element (default: "barplot")                           |
| <code>...</code>         | Additional arguments passed to the <code>plotcli\$new()</code> function |

**Examples**

```

x <- 1:5
y <- c(10, 15, 8, 12, 6)
plotcli_bar(x, y)

```

---

plotcli\_box

*Box plot using plotcli*


---

**Description**

Create a box plot using plotcli. Short alias: `pc1bx`.

**Usage**

```
plotcli_box(  
  y,  
  plot_width = getOption("plotcli.plot_width", 80),  
  plot_height = getOption("plotcli.plot_height", 40),  
  x_label = "x",  
  y_label = "y",  
  color = NULL,  
  braille = getOption("plotcli.braille", TRUE),  
  name = "boxplot",  
  ...  
)
```

**Arguments**

|             |  |
|-------------|--|
| y           | A list of numeric vectors of values                        |
| plot_width  | Width of the plot (default: 80)                            |
| plot_height | Height of the plot (default: 40)                           |
| x_label     | Label for the x-axis (default: "x")                        |
| y_label     | Label for the y-axis (default: "y")                        |
| color       | Color of the plot elements (default: NULL)                 |
| braille     | Use Braille characters for the plot (default: TRUE)        |
| name        | Name of the plot element (default: "boxplot")              |
| ...         | Additional arguments passed to the plotcli\$new() function |
| x           | A vector of categories                                     |

**Examples**

```
y <- rnorm(50, mean = 0)  
plotcli_box(y)
```

---

|                 |                                   |
|-----------------|-----------------------------------|
| plotcli_density | <i>Density plot using plotcli</i> |
|-----------------|-----------------------------------|

---

**Description**

Create a density plot using plotcli. Short alias: pcld.

**Usage**

```
plotcli_density(
  x,
  plot_width = getOption("plotcli.plot_width", 80),
  plot_height = getOption("plotcli.plot_height", 40),
  x_label = "x",
  y_label = "Density",
  color = NULL,
  braille = getOption("plotcli.braille", TRUE),
  name = "density",
  ...
)
```

**Arguments**

|             |  |
|-------------|--|
| x           | A numeric vector of values                                 |
| plot_width  | Width of the plot (default: 80)                            |
| plot_height | Height of the plot (default: 40)                           |
| x_label     | Label for the x-axis (default: "x")                        |
| y_label     | Label for the y-axis (default: "Density")                  |
| color       | Color of the plot elements (default: NULL)                 |
| braille     | Use Braille characters for the plot (default: TRUE)        |
| name        | Name of the plot element (default: "density")              |
| ...         | Additional arguments passed to the plotcli\$new() function |

**Examples**

```
x <- rnorm(100)
plotcli_density(x)
```

---

plotcli\_histogram      *Histogram plot using plotcli*

---

**Description**

Create a histogram plot using plotcli. Short alias: pcli.h.

**Usage**

```
plotcli_histogram(
  x,
  plot_width = getOption("plotcli.plot_width", 80),
  plot_height = getOption("plotcli.plot_height", 40),
  x_label = "x",
```



```

    y_label = "Frequency",
    color = NULL,
    braille = getOption("plotcli.braille", TRUE),
    bin_width = NULL,
    ylim = NULL,
    name = "histogram",
    ...
)

```

### Arguments

|             |  |
|-------------|--|
| x           | A numeric vector of values                                 |
| plot_width  | Width of the plot (default: 80)                            |
| plot_height | Height of the plot (default: 40)                           |
| x_label     | Label for the x-axis (default: "x")                        |
| y_label     | Label for the y-axis (default: "Frequency")                |
| color       | Color of the plot elements (default: NULL)                 |
| braille     | Use Braille characters for the plot (default: TRUE)        |
| bin_width   | Width of the bins (default: NULL)                          |
| ylim        | y limits (default: NULL)                                   |
| name        | Name of the plot element (default: "histogram")            |
| ...         | Additional arguments passed to the plotcli\$new() function |

### Examples

```

x <- rnorm(100)
plotcli_histogram(x)

```

---

|              |                                |
|--------------|--------------------------------|
| plotcli_line | <i>Line plot using plotcli</i> |
|--------------|--------------------------------|

---

### Description

Create a line plot using plotcli. Short alias: pcli.

### Usage

```

plotcli_line(
  y,
  x = NULL,
  plot_width = getOption("plotcli.plot_width", 80),
  plot_height = getOption("plotcli.plot_height", 40),
  x_label = "x",
  y_label = "y",
)

```

```

    color = NULL,
    braille = getOption("plotcli.braille", TRUE),
    name = "line",
    ...
)

```

### Arguments

|                          |   |
|--------------------------|---|
| <code>y</code>           | A numeric vector of y values  |
| <code>x</code>           | A numeric vector of x values  |
| <code>plot_width</code>  | Width of the plot (default: 80)   |
| <code>plot_height</code> | Height of the plot (default: 40)  |
| <code>x_label</code>     | Label for the x-axis (default: "x")                                     |
| <code>y_label</code>     | Label for the y-axis (default: "y")                                     |
| <code>color</code>       | Color of the plot elements (default: NULL)                              |
| <code>braille</code>     | Use Braille characters for the plot (default: TRUE)                     |
| <code>name</code>        | Name of the plot element (default: "line")                              |
| <code>...</code>         | Additional arguments passed to the <code>plotcli\$new()</code> function |

### Examples

```

x <- 1:10
y <- x^2
plotcli_line(x, y)

```

---

`plotcli_options`      *Set global options for plotcli*

---

### Description

Set global options for plotcli

### Usage

```
plotcli_options(plot_width = 60, plot_height = 20, braille = FALSE)
```

### Arguments

|                          |  |
|--------------------------|--|
| <code>plot_width</code>  | Default plot width (default: 60)         |
| <code>plot_height</code> | Default plot height (default: 20)        |
| <code>braille</code>     | Default braille setting (default: FALSE) |

---

plotcli\_scatter      *Scatter plot using plotcli*

---

### Description

Create a scatter plot using plotcli. Short alias: pclis.

### Usage

```
plotcli_scatter(  
  y,  
  x = NULL,  
  plot_width = getOption("plotcli.plot_width", 80),  
  plot_height = getOption("plotcli.plot_height", 40),  
  x_label = "x",  
  y_label = "y",  
  color = NULL,  
  braille = getOption("plotcli.braille", TRUE),  
  name = "scatter",  
  ...  
)
```

### Arguments

|             |  |
|-------------|--|
| y           | A numeric vector of y values                               |
| x           | A numeric vector of x values                               |
| plot_width  | Width of the plot (default: 80)                            |
| plot_height | Height of the plot (default: 40)                           |
| x_label     | Label for the x-axis (default: "x")                        |
| y_label     | Label for the y-axis (default: "y")                        |
| color       | Color of the plot elements (default: NULL)                 |
| braille     | Use Braille characters for the plot (default: TRUE)        |
| name        | Name of the plot element (default: "scatter")              |
| ...         | Additional arguments passed to the plotcli\$new() function |

### Examples

```
x <- rnorm(100)  
y <- rnorm(100)  
plotcli_scatter(x, y)
```

---

|               |  |
|---------------|--|
| rbind.plotcli | <i>Generic function for combining plotcli objects vertically</i> |
|---------------|--|

---

**Description**

Generic function for combining plotcli objects vertically

**Usage**

```
## S3 method for class 'plotcli'  
rbind(..., deparse.level = 1)
```

**Arguments**

... plotcli objects to be combined.  
deparse.level The deparsing level for the arguments.

**Value**

A combined plot matrix.

---

|             |   |
|-------------|---|
| rbind_plots | <i>Combine plot matrices vertically</i> |
|-------------|---|

---

**Description**

This function combines multiple plot matrices vertically, centering them horizontally.

**Usage**

```
rbind_plots(...)
```

**Arguments**

... A list of plot matrices to be combined.

**Value**

A combined plot matrix.

---

|                    |   |
|--------------------|---|
| remove_color_codes | <i>Remove color codes from a string</i> |
|--------------------|---|

---

**Description**

This function removes ANSI color codes from a given text string.

**Usage**

```
remove_color_codes(s)
```

**Arguments**

`s`                    The text string containing ANSI color codes.

**Value**

A text string with ANSI color codes removed.

**Examples**

```
colored_text <- make_colored("Hello, world!", "blue")
remove_color_codes(colored_text)
```

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