

Package: itol.toolkit (via r-universe)

October 10, 2026

Title Helper Functions for 'Interactive Tree Of Life'

Version 1.2.2

URL <https://tongzhou2017.github.io/itol.toolkit/>

Description The 'Interactive Tree Of Life' <<https://itol.embl.de/>> online server can edit and annotate trees interactively. The 'itol.toolkit' package can support all types of annotation templates.

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Encoding UTF-8

Roxygen list(markdown = TRUE)

RoxygenNote 7.3.2

Imports dplyr, stringr, stats, seqinr, utils, methods, tidyr, ape, data.table, purrr, wesanderson, miniUI, shiny, rstudioapi, colourpicker, ggsci, RColorBrewer, stringdist, magrittr, ggplot2

Suggests knitr, rmarkdown, testthat (>= 3.0.0)

Config/testthat/edition 3

VignetteBuilder knitr

Depends R (>= 2.10)

LazyData true

Collate 'addins.R' 'color.R' 'data.R' 'globals.R' 'learn.R' 'object.R' 'utils.R' 'output.R' 'user.R' 'ops.R' 'tree.R' 'zzz.R'

Repository <https://tongzhou2017.r-universe.dev>

Date/Publication 2025-11-02 22:06:29 UTC

RemoteUrl <https://github.com/tongzhou2017/itol.toolkit>

RemoteRef HEAD

RemoteSha 9c5ecc85bd0c7adb0921e118df2cfcac78c97c89

Contents

<code>+,itol.hub,itol.unit-method</code>	3
<code>anno_col_group</code>	4
<code>binary_data_conversion</code>	5
<code>color_distance</code>	6
<code>complex_html_text</code>	6
<code>convert_01</code>	7
<code>convert_01_to_connect</code>	7
<code>convert_range_to_node</code>	8
<code>convert_to_binary</code>	8
<code>correct_get_color</code>	10
<code>correct_type</code>	11
<code>count_to_tree</code>	11
<code>create_hub</code>	12
<code>create_theme</code>	13
<code>create_unit</code>	13
<code>darken_color</code>	16
<code>df_merge</code>	17
<code>df_tree</code>	17
<code>dt_transpose_header</code>	18
<code>fa_read</code>	18
<code>fa_write</code>	19
<code>file_get_dir</code>	19
<code>file_get_name</code>	20
<code>file_to_unit</code>	20
<code>get_color</code>	21
<code>gradient_color</code>	21
<code>head_line</code>	22
<code>hub_to_unit</code>	22
<code>inbuilt_themes</code>	23
<code>itol.hub-class</code>	24
<code>itol.theme-class</code>	24
<code>itol.unit-class</code>	25
<code>learn_data</code>	25
<code>learn_data_from_file</code>	26
<code>learn_data_from_files</code>	26
<code>learn_data_from_unit</code>	27
<code>learn_data_from_unit_list</code>	27
<code>learn_df</code>	28
<code>learn_field</code>	28
<code>learn_legend</code>	29
<code>learn_line</code>	31
<code>learn_profile</code>	32
<code>learn_separator</code>	33
<code>learn_subdf</code>	34
<code>learn_theme_align</code>	34
<code>learn_theme_alignment</code>	35

learn_theme_bar	35
learn_theme_basic_plot	36
learn_theme_basic_theme	36
learn_theme_binary	37
learn_theme_border	37
learn_theme_common_themes	38
learn_theme_connection	38
learn_theme_domain	39
learn_theme_externalshape	39
learn_theme_heatmap	40
learn_theme_image	40
learn_theme_label	41
learn_theme_linechart	42
learn_theme_piechart	43
learn_theme_specific_themes	44
learn_theme_strip_label	44
learn_type	45
line_clean	46
line_split	47
longest_continuous_match	48
merge_unit	48
search_tree_file	49
show,itol.hub-method	49
sort_color	50
template_groups	50
template_parameters_count	51
train_theme	52
unite_rows	52
use.theme	53
vector_tree	54
write_hub	54
write_raw	56
write_unit	57

Index 59

+,itol.hub,itol.unit-method

plus method add method for S4 class itol.hub and itol.unit

Description

plus method add method for S4 class itol.hub and itol.unit

plus method add method for S4 class itol.unit and itol.unit

Usage

```
## S4 method for signature 'itol.hub,itol.unit'
e1 + e2

## S4 method for signature 'itol.unit,itol.unit'
e1 + e2
```

Arguments

- e1 An object of class itol.unit
- e2 An object of class itol.unit

Value

- a itol.hub object with new data from itol.unit object
- a itol.unit object with merged data

anno_col_group	<i>Annotate Column Groups</i>
----------------	-------------------------------

Description

Adds column grouping information as a hidden attribute to a data frame. This allows you to specify which columns belong to the same group without modifying the data structure itself.

Usage

```
anno_col_group(data, groups, attr_name = "col_groups")
```

Arguments

- data A data frame or data.table to annotate.
- groups A named list or vector where names are group names and values are column names (or indices) that belong to that group. Supports both explicit column names and regular expressions.
- attr_name The name of the attribute to store the grouping information. Default is "col_groups". This allows you to have separate grouping attributes for different purposes (e.g., "col_groups_color" for color grouping, "col_groups_shape" for shape grouping).

Details

The grouping information is stored as a hidden attribute, which does not affect normal data frame operations. This allows you to annotate data with grouping information without changing its structure.

By default, the grouping information is stored in an attribute named "col_groups". However, you can specify a different attribute name to allow for separate grouping configurations for color and shape (or other purposes).

Value

The input data frame with an additional attribute containing the grouping information.

Examples

```
## Not run:
# Create a data frame with multiple columns
data <- data.frame(
  ID = c("A", "B", "C"),
  Asia_1 = c(0.5, 0.6, 0.7),
  Asia_2 = c(0.4, 0.5, 0.6),
  North_1 = c(0.3, 0.4, 0.5),
  North_2 = c(0.2, 0.3, 0.4)
)

# Method 1: Explicit column names
data <- anno_col_group(data, list(
  Asia = c("Asia_1", "Asia_2"),
  North = c("North_1", "North_2")
))

# Method 2: Regular expression
data <- anno_col_group(data, list(
  Asia = "^Asia",
  North = "^North"
))

# Method 3: Separate grouping for color and shape
data <- anno_col_group(data, list(
  Group1 = c("Asia_1", "Asia_2")
), attr_name = "col_groups_color")

data <- anno_col_group(data, list(
  Group1 = c("North_1")
), attr_name = "col_groups_shape")

# Access grouping information
attr(data, "col_groups_color")
attr(data, "col_groups_shape")

## End(Not run)
```

binary_data_conversion

Binary Data Conversion

Description

Interactively convert data to binary format (-1, 0, 1) using range sliders. Provides three pairs of sliders for negative, zero, and positive ranges with automatic constraint handling to prevent overlaps.

Usage

binary_data_conversion()

color_distance	<i>Color distance</i>
----------------	-----------------------

Description

calculate distance between two color in hex or rgb format. rgb maxColorValue = 255.

Usage

color_distance(color_1, color_2)

Arguments

color_1 color 1 in hex or rgb format
color_2 color 2 in hex or rgb format

Value

a distance number

complex_html_text	<i>Complex HTML text</i>
-------------------	--------------------------

Description

Interactively combine columns by HTML styles and record workflow as reproducible code.

Usage

complex_html_text()

Details

When you're done, the code performing this operation will be emitted at the cursor position.

convert_01	<i>Convert character data to 0/1</i>
------------	--------------------------------------

Description

In data frame and list, convert character and numeric data to 0/1.

Usage

```
convert_01(object)
```

Arguments

object	data frame or list
--------	--------------------

Value

a data frame with 0/1 values

convert_01_to_connect	<i>Convert 0/1 data to connection pairs</i>
-----------------------	---

Description

If two column has more than 1 shared element then they have connection. Convert 0/1 data to connection pairs in long shape table. The 0-connection pairs are removed.

Usage

```
convert_01_to_connect(object)
```

Arguments

object	data frame with 0/1 data
--------	--------------------------

Value

a data frame with source and target connection information

convert_range_to_node *Convert range to node id*

Description

Convert the data frame with range id to node id by mrca method.

Usage

```
convert_range_to_node(df, tree)
```

Arguments

df	data frame with any type of id
tree	tree file path

Value

a data frame with converted id from range id

convert_to_binary *Convert data to binary format for DATASET_BINARY*

Description

Convert various data formats to binary format (-1, 0, 1) required by DATASET_BINARY. This function provides flexible data conversion options and can detect if data is already in the correct format.

Usage

```
convert_to_binary(
  data,
  force_convert = FALSE,
  verbose = TRUE,
  negative_range = NULL,
  zero_range = NULL,
  positive_range = NULL,
  lower_inclusive = TRUE,
  upper_inclusive = TRUE,
  auto_detect = TRUE
)
```

Arguments

data	A data frame with the first column as ID and subsequent columns as data to convert
force_convert	Logical, whether to force conversion even if data appears to be already in binary format
verbose	Logical, whether to print conversion messages
negative_range	Numeric vector of length 2, range for converting to -1 (e.g., c(0, 0.2))
zero_range	Numeric vector of length 2, range for converting to 0 (e.g., c(0.2, 0.8))
positive_range	Numeric vector of length 2, range for converting to 1 (e.g., c(0.8, 1))
lower_inclusive	Logical, whether middle range (zero_range) lower bound is inclusive (default: TRUE)
upper_inclusive	Logical, whether middle range (zero_range) upper bound is inclusive (default: TRUE)
auto_detect	Logical, whether to automatically detect data range and set ranges (default: TRUE)

Details

The function handles several data types with flexible range control:

- Boolean/logical data: TRUE->1, FALSE->0
- 0-1 range data: Custom ranges for -1, 0, 1 conversion
- Non-negative data: Custom ranges for -1, 0, 1 conversion
- Already binary data: No conversion if already in -1, 0, 1 format

Range parameters work as follows:

- Values in negative_range -> -1
- Values in zero_range -> 0 (with inclusive/exclusive boundary control)
- Values in positive_range -> 1
- Values outside all ranges -> NA (with warning)

The middle range (zero_range) boundary control:

- If lower_inclusive=TRUE: lower bound is inclusive
- If lower_inclusive=FALSE: lower bound is exclusive
- If upper_inclusive=TRUE: upper bound is inclusive
- If upper_inclusive=FALSE: upper bound is exclusive

This allows flexible scenarios like:

- Standard: negative->-1, middle->0, positive->1
- Inverted: negative->0, middle->-1, positive->1
- Custom: any range mapping to -1, 0, 1

Value

A data frame with data converted to binary format (-1, 0, 1)

Examples

```
# Convert 0-1 range data with default ranges
data <- data.frame(ID = c("A", "B", "C"),
                  Asia = c(0, 0.3, 0.8),
                  Europe = c(1, 0.7, 0.2))
convert_to_binary(data)

# Convert with custom ranges and boundary control
convert_to_binary(data,
                  negative_range = c(0, 0.2),
                  zero_range = c(0.2, 0.8),
                  positive_range = c(0.8, 1),
                  lower_inclusive = TRUE, # [0.2, ...]
                  upper_inclusive = TRUE) # [..., 0.8]

# Convert boolean data
data_bool <- data.frame(ID = c("A", "B", "C"),
                       Present = c(TRUE, FALSE, TRUE),
                       Absent = c(FALSE, TRUE, FALSE))
convert_to_binary(data_bool)
```

correct_get_color	<i>correct_get_color</i>
-------------------	--------------------------

Description

correct_get_color. (Version 0.0.0.9000)

Usage

```
correct_get_color(str)
```

Arguments

str	taxa string
-----	-------------

Value

a vector of colors

correct_type	<i>Correct type parameter</i>
--------------	-------------------------------

Description

Correct type parameter input by string similarity and synonym mapping.

Usage

```
correct_type(str)
```

Arguments

str	A character string representing a type name to correct.
-----	---

Value

A character string of the corrected type name.

Examples

```
correct_type("line")
```

count_to_tree	<i>Calculate tree based on count matrix</i>
---------------	---

Description

While we start analysis from count matrix not sequences alignment, we could use clustering methods to get main tree in phylo object class of output as Newick format file. If the samples or elements have group information, we could use weighted clustering method to get a clear grouped structure.

Usage

```
count_to_tree(count, group = NULL, weight = 0)
```

Arguments

count	a data frame containing numeric values of abundance or other count.
group	a vector of character containing the group information. The length of the vector should be same with the count columns number. If using unweighted clustering, should ignore this parameter.
weight	a number specifying the weight size of the group information. In most case, 1 is enough. If the value is between 0 and 1, it will make the weight of group information weak. If the value is more than 1, it will make the weight of group information strong.

Value

a phylo class object containing

edge	a vector of integers specifying edge id. The length of vector is double of node number
edge.length	a vector of numbers specifying edge length
tip.label	a vector of character specifying the tip label
Nnode	a number specifying the number of nodes
node.label	a vector of character specifying the node label. If the tree calculated from count matrix or other case, the node label will generated by ape::makeNodeLabel function. And the Most Recent Common Ancestors(MRCA) node will be named with weighted group information, if the parameter group is not null.

create_hub

Create itol.hub Object

Description

create a new object for itol.hub

Usage

```
create_hub(
  tree,
  field_tree = NULL,
  seq = NULL,
  abundance = NULL,
  taxonomy = NULL,
  node_data = NULL,
  tip_data = NULL
)
```

Arguments

tree	tree file
field_tree	phylogenetic tree for field data
seq	sequence data object
abundance	abundance data object
taxonomy	taxonomy data object
node_data	node-level metadata object
tip_data	tip-level metadata object

Value

Returns a itol.hub object

Examples

```
TREE <- system.file("extdata", "tree_of_itol_templates.tree", package = "itol.toolkit")
create_hub(tree = TREE)
```

create_theme	<i>Create itol.theme Object</i>
--------------	---------------------------------

Description

create a new object for itol.theme

Usage

```
create_theme(unit = NULL, file = NULL, tree = NULL, ...)
```

Arguments

unit	unit object
file	template file
tree	tree file
...	Further arguments to be passed to subsequent functions.

Value

Returns a itol.theme object

create_unit	<i>Create itol.unit</i>
-------------	-------------------------

Description

Create itol.unit from simple input in R environment.

Usage

```
create_unit(
  data,
  key,
  type,
  style = "default",
  subtype = NULL,
  color = NULL,
  line_type = NULL,
  font_type = NULL,
```

```

    size_factor = NULL,
    position = NULL,
    background_color = NULL,
    rotation = NULL,
    method = NULL,
    shape = NULL,
    fill = NULL,
    tree
)

```

Arguments

data	if type == "COLLAPSE", a vector of characters specifying the tips or node used for collapsing used for extracting.
key	a character specifying the output file name for hub object.
type	a character specifying the template type used for extracting. Following choices are possible: "COLLAPSE", "PRUNE", "SPACING", "TREE_COLORS", "DATASET_STYLE", "LABEL", "DATASET_BINARY", "DATASET_GRADIENT", "DATASET_HEATMAP", "DATASET_SYMBOL", "DATASET_EXTERNALSHAPE", "DATASET_DOMAINS", "DATASET_SIMPLEBAR", "DATASET_MULTIBAR", "DATASET_BOXPLOT", "DATASET_LINECHART", "DATASET_PIECHART", "DATASET_ALIGNMENT", "DATASET_CONNECTION", "DATASET_IMAGE", "POPUP_INFO."
style	a character specifying the specific version of template type used for extracting. The default value is "default" style for all types.
subtype	a character specifying the subtype under type. If the type is "TREE_COLORS", the following choices are possible: "range", "clade", "branch", "label", "label_background".
color	a character specifying the color pattern name. The following choices are possible: "table2itol", "RColorBrewer", "ggsci".
line_type	a character specifying the normal or dashed line type used in clade and branch subtype.
font_type	a character specifying the bold, italic, and bold-italic font type used in label and branch subtype.
size_factor	a number specifying the line width used in clade and branch subtype and size factor in label subtype.
position	If type == "DATASET_STYLE", a character specifying the position: The following choices are possible: "node" and "clade". If type == "DATASET_TEXT", a number specifying the position of the text on the tree: -1 = external label; a number between 0 and 1 = internal label positioned at the specified value along the node branch (for example, position 0 is exactly at the start of node branch, position 0.5 is in the middle, and position 1 is at the end)
background_color	Only used while type == "DATASET_STYLE" and subtype == "label". a character or a vector of character specifying the background color in hexadecimal, RGB or RGBA notation.
rotation	Only used while type == "DATASET_TEXT". a number or a vector of number specifying the rotation angle of the text.

method	a character specifying the numeric data summarise method. If type == "DATASET_BINARY", the following choices are possible: "mean", "sum", "binary" (for automatic binary conversion).
shape	a character or a vector of character specifying the symbol shape. If type == "DATASET_BINARY", the default is 2. If type == "DATASET_SYMBOL", the following choices are possible: 1 for rectangle, 2 for circle, 3 for star, 4 for left pointing triangle, 5 for right pointing triangle. If using NULL and there are data column, the functions will automatically help users to setup the shapes based on the levels of the data.
fill	If type == "DATASET_SYMBOL", 1/0 is specifying the shape outlier filled or not. If type == "DATASET_DOMAINS", the following choices are possible: "RE HH HV EL DI TR TL PL PR PU PD OC GP".
tree	a character specifying Newick format tree file path or a phylo object of main phylogenetic tree.

Value

a itol.unit object containing

type	This group holds information about the template type of the data only. This is a very critical piece of information. In many functions of the itol.toolkit package, the template type information is used to determine the different data processing and input/output methods.
sep	This group holds data separator information only. This is one of the most important parameters for data reading and output. It is a separate category because it is frequently used and is an input parameter for other subsequent parameters to be read.
profile	This group contains basic information about the dataset, such as the dataset name and a color label to distinguish the dataset. The dataset name is extremely important. This parameter is used almost throughout the data processing of the itol.toolkit package. With the content of this parameter as the key value, the data and theme information of the dataset are associated. In turn, high throughput learning and writing of large-scale data can be achieved. This parameter is not included in some template types with a particularly simple structure, so we choose a file name or a user-defined method as the key value.
field	This group contains information about each sample within the dataset, and this type of parameter exists only for multi-sample data. This information even includes the clustering tree between samples. This information is usually stored as part of the column names in the metadata part or abundance information of the itol.hub object.
common_themes	These themes are used at high frequency in different templates. These parameters are small in number but constitute some common features of iTOL visual style settings, such as legend, margin, etc.
specific_themes	These themes are used only in specific templates. The number of these parameters is very large. However, most of them are used in only one template to control the style details of the visualization. By unifying these parameters and

calling them according to the template type, users can perform secondary development and data processing with a high degree of parameter aggregation without worrying too much about the differences between different template types.

data This slot contains a list of two data frames with the nodes and tips data separately. The first column of the two data frames is the node or tip id. If the input data contains range id, it would be converted to node id by the `convert_range_to_node` function automatically.

Examples

```
tree <- system.file("extdata", "tree_of_itol_templates.tree", package = "itol.toolkit")
data("template_groups")
data("template_parameters_count")
# COLLAPSE
group_names <- unique(template_groups$group)
object <- create_hub(tree = tree)
unit <- create_unit(data = group_names, key = "E001_collapse_1",
  type = "COLLAPSE", tree = tree)
object <- learn_data_from_unit(object, unit)
# PRUNE
select_note = c("theme_style", "basic_plot")
unit <- create_unit(data = select_note, key = "E002_prune_1",
  type = "PRUNE", tree = tree)
object <- learn_data_from_unit(object, unit)
# SPACING
df_values = data.frame(id = row.names(template_parameters_count),
  values = rowSums(template_parameters_count))
unit <- create_unit(data = df_values, key = "E005_spacing_1",
  type = "SPACING", tree = tree)
object <- learn_data_from_unit(object, unit)
# TREE_COLORS
## range
unit <- create_unit(data = template_groups,
  key = "E006_tree_colors_1", type = "TREE_COLORS", subtype = "range",
  tree = tree)
object <- learn_data_from_unit(object, unit)
```

darken_color

Darken color while preserving hue and saturation

Description

Darken a color by proportionally reducing RGB values while maintaining the color's hue and saturation characteristics.

Usage

```
darken_color(color, factor = 0.4, method = "proportional")
```

Arguments

color	input color in hex format
factor	darkening factor (0-1), where 0 is no darkening and 1 is maximum darkening
method	method for darkening: "proportional" (default) or "gradient"

Value

darkened color in hex format

df_merge	<i>Merge two data frame</i>
----------	-----------------------------

Description

merge sub data frame into initial data frame

Usage

```
df_merge(df1, df2, by = "id")
```

Arguments

df1	initial data frame
df2	sub data frame
by	key column

Value

a data frame containing merged information

df_tree	<i>Data frame tree</i>
---------	------------------------

Description

build a tree by row and column names of a data frame.

Usage

```
df_tree(df, main = "col", order = "none", sep = "_")
```

Arguments

df	input data frame
main	col(default) or row
order	none or value
sep	default '_'

dt_transpose_header	<i>Transpose a data.table using the first column as new header</i>
---------------------	--

Description

Transpose a data.table such that the first column's values become the new column names, and the original column names (excluding the first) become a new 'Sample' column.

Usage

```
dt_transpose_header(dt)
```

Arguments

dt	A data.table to transpose. The first column contains the new column names.
----	--

Value

A transposed data.table with updated column names and a 'Sample' column.

fa_read	<i>Read fasta file</i>
---------	------------------------

Description

Read the fasta format sequences file into data.frame

Usage

```
fa_read(file)
```

Arguments

file	input file in fasta format
------	----------------------------

Value

a data frame with sequence id and sequence

fa_write	<i>Write fasta file</i>
----------	-------------------------

Description

Write the fasta format sequences file from data.frame. (Version 0.0.0.9000)

Usage

```
fa_write(object, file, id = "seq_name", seq = "sequence", append = FALSE)
```

Arguments

object	data.frame format data
file	input file in fasta format
id	id col
seq	seq col
append	append at the end of an already existing file

Value

No return value, only output a fasta file

file_get_dir	<i>Get file dir</i>
--------------	---------------------

Description

Get file dir from string

Usage

```
file_get_dir(str, up = FALSE)
```

Arguments

str	str
up	up dir

Value

a character specifying the dir path

file_get_name	<i>Get file name</i>
---------------	----------------------

Description

Get file name from string

Usage

```
file_get_name(str, with_ext = TRUE, keep_dir = FALSE)
```

Arguments

str	str
with_ext	with ext or not
keep_dir	keep file dir or not

Value

a character specifying the file name

file_to_unit	<i>Create itol.unit Object from file</i>
--------------	--

Description

create a new object for itol.unit

Usage

```
file_to_unit(file, tree, ...)
```

Arguments

file	template file
tree	tree file
...	Further arguments to be passed to subsequent functions.

Value

Returns a itol.unit object

get_color	<i>get_color</i>
-----------	------------------

Description

get color, support max length 40

Usage

```
get_color(n = 0, set = "table2itol")
```

Arguments

n	level length of a vector
set	a character specifying the palette set name. In default, table2itol is setted. The following choices are possible: wsanderson.

Value

a vector of colors

gradient_color	<i>Generate gradient colors</i>
----------------	---------------------------------

Description

generate a vector of gradient colors by start, mid, and end colors.

Usage

```
gradient_color(n, start, mid = NULL, end = "#FFFFFF")
```

Arguments

n	the length of vector
start	start color in hex format
mid	mid color in hex format, default is null.
end	end color in hex format, default is white.

Value

a vector of gradient colors

head_line	<i>head line</i>
-----------	------------------

Description

Head line for templates

Usage

head_line(function_name)

Arguments

function_name parent function name

Value

a character specifying the template type

hub_to_unit	<i>Create itol.unit Object from object</i>
-------------	--

Description

create a new object for itol.unit

Usage

hub_to_unit(object, theme, key)

Arguments

object	itol.hub object
theme	itol.theme object
key	key id of dataset name

Value

Returns a itol.unit object

inbuilt_themes

inbuilt themes

Description

Default themes learned from iTOL official template examples.

Usage

```
inbuilt_themes
```

Format

```
inbuilt_themes:
```

A list with 23 template themes:

COLLAPSE Default theme of collapse template

PRUNE Default theme of prune template

SPACING Default theme of spacing template

TREE_COLORS Default theme of tree colors template

DATASET_STYLE Default theme of style template

LABELS Default theme of labels template

DATASET_TEXT Default theme of text template

DATASET_COLORSTRIP Default theme of colorstrip template

DATASET_BINARY Default theme of binary template

DATASET_GRADIENT Default theme of gradient template

DATASET_HEATMAP Default theme of heatmap template

DATASET_SYMBOL Default theme of symbol template

DATASET_EXTERNALSHAPE Default theme of externalshape template

DATASET_DOMAINS Default theme of domains template

DATASET_SIMPLEBAR Default theme of simple bar template

DATASET_MULTIBAR Default theme of multi bar template

DATASET_BOXPLOT Default theme of box plot template

DATASET_LINECHART Default theme of line chart template

DATASET_PIECHART Default theme of pie chart template

DATASET_ALIGNMENT Default theme of alignment template

DATASET_CONNECTION Default theme of connection template

DATASET_IMAGE Default theme of image template

POPUP_INFO Default theme of popup info template ...

<code>itol.hub-class</code>	<i>The itol.hub Class</i>
-----------------------------	---------------------------

Description

The itol.hub object is an intermediate storage container used internally throughout the integration procedure to hold bits of data that are useful downstream.

Slots

- `tree` a list of meta data table, usually raw, full, and analyze
- `seq` identity of the active assay
- `abundance` abundance
- `taxonomy` taxonomy
- `meta.data` other meta.data
- `theme` itol theme

<code>itol.theme-class</code>	<i>The itol.theme Class</i>
-------------------------------	-----------------------------

Description

The itol.theme object is an intermediate storage container used internally throughout the integration procedure to hold bits of data that are useful downstream.

Slots

- `type` a list of meta data table, usually raw, full, and analyze
- `sep` identity of the active assay
- `profile` abundance
- `field` taxonomy
- `common_themes` other meta.data
- `specific_themes` itol theme

itol.unit-class	<i>The itol.unit Class</i>
-----------------	----------------------------

Description

The itol.unit object is an intermediate storage container used internally throughout the integration procedure to hold bits of data that are useful downstream.

Slots

type a list of meta data table, usually raw, full, and analyze
sep identity of the active assay
profile abundance
field taxonomy
common_themes other meta.data
specific_themes itol theme
data data

learn_data	<i>Learn data from template file</i>
------------	--------------------------------------

Description

Learn data from template file into data frame

Usage

```
learn_data(df1 = NULL, file, tree = NULL, ...)
```

Arguments

df1	initial data frame
file	template file
tree	tree file
...	Further arguments to be passed to subsequent functions.

Value

a list with two data frame of node and tip annotation data

`learn_data_from_file` *Learn object data from file*

Description

Learn itol.hub object data from template file.

Usage

```
learn_data_from_file(object, file)
```

Arguments

<code>object</code>	itol.hub object
<code>file</code>	template file

Value

a itol.hub object with new data from template file

`learn_data_from_files` *Learn object data from files*

Description

Learn itol.hub object data from template file.

Usage

```
learn_data_from_files(object, files = NULL, dir = NULL, pattern = ".", ...)
```

Arguments

<code>object</code>	itol.hub object
<code>files</code>	template files path
<code>dir</code>	files path
<code>pattern</code>	file name pattern in regex
<code>...</code>	Further arguments to be passed to subsequent functions.

Value

a itol.hub object with new data from template files

learn_data_from_unit	<i>Learn object data from unit</i>
----------------------	------------------------------------

Description

Learn itol.hub object data from unit object.

Usage

```
learn_data_from_unit(object, unit)
```

Arguments

object	itol.hub object
unit	itol.unit object

Value

a itol.hub object containing new data from itol.unit object

learn_data_from_unit_list	<i>Learn object data from units</i>
---------------------------	-------------------------------------

Description

Learn itol.hub object data from list of unit object.

Usage

```
learn_data_from_unit_list(object, units)
```

Arguments

object	itol.hub object
units	itol.unit object list

Value

a itol.hub object with new data from a list of itol.unit objects

learn_df	<i>Learn from tree</i>
----------	------------------------

Description

Learn initial data frame from Newick format tree leaves.

Usage

```
learn_df(tree, node = FALSE, tip = TRUE)
```

Arguments

tree	Newick tree file or phylo object.
node	a logical to control output with node label or not. The default value is FALSE.
tip	a logical to control output tip label or not. The default value is TRUE.

Value

a list containing	
node	a data frame with id column. The id information is from the node label in Newick format tree file or phylo object. If the node parameter set as FALSE, the node information will be NULL.
tip	a data frame with id column. The id information is from the tip label in Newick format tree file or phylo object. If the tip parameter set as FALSE, the tip information will be NULL.

Examples

```
tree <- system.file("extdata",
                    "tree_of_itol_templates.tree",
                    package = "itol.toolkit")
sub_df <- learn_df(tree, node=TRUE, tip=TRUE)
```

learn_field	<i>Learn field</i>
-------------	--------------------

Description

learn field paramters as list

Usage

```
learn_field(lines, sep)
```

Arguments

`lines` a vector of character strings from template file.

`sep` a character specifying the separator.

Value

a list of field parameters containing

`labels` a vector of characters specifying the field name. In DATASET_HEATMAP, the labels are shown as heatmap column names.

`colors` define colors for each individual field column (use hexadecimal, RGB or RGBA notation; if using RGB/RGBA, COMMA cannot be used as SEPARATOR)

`shapes` Shape should be a number between 1 and 6, or any protein domain shape definition. 1-square, 2-circle, 3-star, 4-right pointing triangle, 5-left pointing triangle, 6-checkmark

Examples

```
tree <- system.file("extdata",
                    "tree_of_itol_templates.tree",
                    package = "itol.toolkit")
df_frequence <- data.table::fread(system.file("extdata",
                                              "templates_frequence.txt",
                                              package = "itol.toolkit"))

## create unit
unit <- create_unit(data = df_frequence,
                   key = "Quickstart",
                   type = "DATASET_HEATMAP",
                   tree = tree)

## write unit
file <- tempfile()
write_unit(unit, file)
## Learn legend parameters
lines <- line_clean(file=file)
sep = learn_separator(file = file)
learn_field(lines = lines, sep = sep)
```

learn_legend

Learn legend

Description

learn legend parameters as list

Usage

```
learn_legend(lines, sep)
```

Arguments

`lines` a vector of character strings from template file.

`sep` a character specifying the separator.

Value

a list of legned parameters containing

`title` a character specifying the title of legend. There should not be the character same with separator within.

`position_x` a number specifying the x axis px value of the legend.

`position_y` a number specifying the y axis px value of the legend.

`horizontal` To order legend entries horizontally instead of vertically, set this parameter to 1

`shapes` Shape should be a number between 1 and 6, or any protein domain shape definition. 1-square, 2-circle, 3-star, 4-right pointing triangle, 5-left pointing triangle, 6-checkmark

`colors` define colors for each legend element (use hexadecimal, RGB or RGBA notation; if using RGB/RGBA, COMMA cannot be used as SEPARATOR)

`labels` The legend element label. There should not be the character same with separator within.

`shape_scales` For each shape, you can define a scaling factor between 0 and 1.

Examples

```
tree <- system.file("extdata",
                    "tree_of_itol_templates.tree",
                    package = "itol.toolkit")
df_frequence <- data.table::fread(system.file("extdata",
                                              "templates_frequence.txt",
                                              package = "itol.toolkit"))

## create unit
unit <- create_unit(data = df_frequence,
                    key = "Quickstart",
                    type = "DATASET_SIMPLEBAR",
                    method = "mean",
                    tree = tree)

## write unit
file <- tempfile()
write_unit(unit, file)
## Learn legend parameters
lines <- line_clean(file=file)
sep = learn_separator(file = file)
learn_legend(lines = lines, sep = sep)
```

learn_line	<i>Learn paramter</i>
------------	-----------------------

Description

learn paramter name and values based on the key name in the front of line.

Usage

```
learn_line(lines, param, sep)
```

Arguments

lines	a vector of character strings from template file.
param	a charactor string of paramter key name. The key name should be uppercse letters or '_' without spacing.
sep	a charactor specifying the separator.

Value

a charactor string containing parameter value.

Examples

```
tree <- system.file("extdata",
                    "tree_of_itol_templates.tree",
                    package = "itol.toolkit")
data("template_groups")
df_group <- data.frame(id = unique(template_groups$group),
                      data = unique(template_groups$group))

## create unit
unit <- create_unit(data = df_group,
                   key = "Quickstart",
                   type = "DATASET_COLORSTRIP",
                   tree = tree)

## write unit
file <- tempfile()
write_unit(unit, file)
## Learn parameter
lines <- line_clean(file=file)
sep = learn_separator(file = file)
learn_line(lines = lines, param = "STRIP_WIDTH", sep = sep)
```

learn_profile	<i>Learn profile</i>
---------------	----------------------

Description

learn profile paramters as list

Usage

```
learn_profile(lines, sep)
```

Arguments

lines	a vector of character strings from template file.
sep	a character specifying the separator.

Value

a list of profile parameters containing

name	a character specifying label, which is used in the legend table
color	dataset color in the legend (use hexadecimal, RGB or RGBA notation; if using RGB/RGBA, COMMA cannot be used as SEPARATOR)

Examples

```
tree <- system.file("extdata",
                    "tree_of_itol_templates.tree",
                    package = "itol.toolkit")
df_frequence <- data.table::fread(system.file("extdata",
                                              "templates_frequence.txt",
                                              package = "itol.toolkit"))

## create unit
unit <- create_unit(data = df_frequence,
                    key = "Quickstart",
                    type = "DATASET_HEATMAP",
                    tree = tree)

## write unit
file <- tempfile()
write_unit(unit, file)
## Learn legend parameters
lines <- line_clean(file=file)
sep = learn_separator(file = file)
learn_profile(lines = lines, sep = sep)
```

learn_separator	<i>Learn separator</i>
-----------------	------------------------

Description

Learn 3 types of separators: tab, space, and comma.

Usage

```
learn_separator(lines = NULL, file = NULL)
```

Arguments

lines	a vector of character strings from template file. If the file parameter is NULL, this parameter should be set.
file	a character specifying the template file path. If this parameter is setted, the lines parameter will be replaced.

Value

a character specifying the separator

Examples

```
tree <- system.file("extdata",
                    "tree_of_itol_templates.tree",
                    package = "itol.toolkit")
data("template_groups")
df_group <- data.frame(id = unique(template_groups$group),
                      data = unique(template_groups$group))

## create unit
unit <- create_unit(data = df_group,
                   key = "Quickstart",
                   type = "DATASET_COLORSTRIP",
                   tree = tree)

## write unit
file <- tempfile()
write_unit(unit, file)
## Learn template type
learn_separator(file = file)
```

learn_subdf	<i>Learn sub data frame</i>
-------------	-----------------------------

Description

Learn sub data frame from template file

Usage

```
learn_subdf(lines, type, sep, dataset_name = NULL, field_labels = NULL)
```

Arguments

- lines a vector of character strings from template file.
- type template type
- sep a character specifying the separator.
- dataset_name label in template file
- field_labels sample ids for binary, heatmap, and other multi-column value templates

Value

a data frame containing the data learned from template file

learn_theme_align	<i>Learn align</i>
-------------------	--------------------

Description

learn connection paramters as list

Usage

```
learn_theme_align(lines, sep)
```

Arguments

- lines a vector of character strings from template file.
- sep a character specifying the separator.

Value

a list of align parameters containing

learn_theme_alignment *Learn alignment*

Description

learn alignment paramters as list

Usage

```
learn_theme_alignment(lines, sep)
```

Arguments

lines	a vector of character strings from template file.
sep	a character specifying the separator.

Value

a list of alignment parameters containing

learn_theme_bar *Learn bar*

Description

learn bar paramters as list

Usage

```
learn_theme_bar(lines, sep)
```

Arguments

lines	file lines
sep	a character specifying the separator.

Value

a list of bar parameters containing

learn_theme_basic_plot

Learn basic plot

Description

learn basic plot paramters as list

Usage

```
learn_theme_basic_plot(lines, sep)
```

Arguments

lines	a vector of character strings from template file.
sep	a character specifying the separator.

Value

a list of basic plot parameters containing

learn_theme_basic_theme

Learn basic theme

Description

learn basic theme paramters as list

Usage

```
learn_theme_basic_theme(lines, sep)
```

Arguments

lines	a vector of character strings from template file.
sep	a character specifying the separator.

Value

a list of basic theme parameters containing

learn_theme_binary	<i>Learn binary</i>
--------------------	---------------------

Description

learn binary paramters as list

Usage

```
learn_theme_binary(lines, sep)
```

Arguments

lines	a vector of character strings from template file.
sep	a character specifying the separator.

Value

a list of binary chart parameters containing

learn_theme_border	<i>Learn border</i>
--------------------	---------------------

Description

learn border paramters as list

Usage

```
learn_theme_border(lines, sep)
```

Arguments

lines	a vector of character strings from template file.
sep	a character specifying the separator.

Value

a list of border parameters containing

`learn_theme_common_themes`*Learn common themes*

Description

learn common theme paramters as list

Usage

```
learn_theme_common_themes(lines, sep)
```

Arguments

<code>lines</code>	a vector of character strings from template file.
<code>sep</code>	a character specifying the separator.

Value

a list of common theme parameters containing

`learn_theme_connection`*Learn connection*

Description

learn connection paramters as list

Usage

```
learn_theme_connection(lines, sep)
```

Arguments

<code>lines</code>	a vector of character strings from template file.
<code>sep</code>	a character specifying the separator.

Value

a list of connection parameters containing

learn_theme_domain	<i>Learn domain</i>
--------------------	---------------------

Description

learn domain paramters as list

Usage

```
learn_theme_domain(lines, sep)
```

Arguments

lines	a vector of character strings from template file.
sep	a character specifying the separator.

Value

a list of domain parameters containing

learn_theme_externalshape	<i>Learn externalshape</i>
---------------------------	----------------------------

Description

learn connection paramters as list

Usage

```
learn_theme_externalshape(lines, sep)
```

Arguments

lines	a vector of character strings from template file.
sep	a character specifying the separator.

Value

a list of external shape parameters containing

learn_theme_heatmap	<i>Learn heatmap</i>
---------------------	----------------------

Description

learn heatmap paramters as list

Usage

```
learn_theme_heatmap(lines, sep)
```

Arguments

lines	a vector of character strings from template file.
sep	a character specifying the separator.

Value

a list of heatmap parameters containing

learn_theme_image	<i>Learn image</i>
-------------------	--------------------

Description

learn connection paramters as list

Usage

```
learn_theme_image(lines, sep)
```

Arguments

lines	a vector of character strings from template file.
sep	a character specifying the separator.

Value

a list of image parameters containing

learn_theme_label	<i>Learn label</i>
-------------------	--------------------

Description

learn label paramters as list

Usage

```
learn_theme_label(lines, sep)
```

Arguments

lines	a vector of character strings from template file.
sep	a character specifying the separator.

Value

a list of label parameters containing

display	1/0 specifying display or hide the text labels above each field column
size	a number specifying the size factor for the text labels
top	1/0 specifying the labels position. If 0, label text which does not fit into the shape will be hidden
below	1/0 specifying the labels position. By default, internal labels will be placed above the branches. If 1, labels will be below the branches
rotation	a number specifying text label rotation angle
straight	1/0 specifying tree rotation. If set to 1, tree rotation will not influence the individual label rotation
vertical	a number specifying the label vertical shift. Shift internal labels vertically by this amount of pixels (positive or negative)
shift	a number specifying the label shift. text label shift in pixels (positive or negative)
external_shift	1/0 specifying label external shift that add extra horizontal shift to the external labels. Useful in unrooted display mode to shift text labels further away from the node labels.

Examples

```
library(dplyr)
tree <- system.file("extdata",
  "tree_of_itol_templates.tree",
  package = "itol.toolkit")
tab_tmp <- data.table::fread(system.file("extdata",
  "parameter_groups.txt",
  package = "itol.toolkit"))
```

```

tab_id_group <- tab_tmp[,c(1,2)]
tab_tmp <- tab_tmp[,-c(1,2)]
tab_tmp_01 <- convert_01(object = tab_tmp)
tab_tmp_01 <- cbind(tab_id_group,tab_tmp_01)
order <- c("type","separator","profile","field","common themes",
  "specific themes","data")
tab_tmp_01_long <- tab_tmp_01 %>%
  tidyr::gather(key = "variable",
    value = "value",
    c(-parameter,-group))
template_start_group <- tab_tmp_01_long %>%
  group_by(group,variable) %>%
  summarise(sublen = sum(value)) %>%
  tidyr::spread(key=variable,
    value=sublen)
template_start_group$group <- factor(template_start_group$group,
  levels = order)
template_start_group <- template_start_group %>% arrange(group)
start_group <- data.frame(Var1 = template_start_group$group,
  Freq = apply(template_start_group[,-1], 1, max))
start_group$start <- 0
for (i in 2:nrow(start_group)) {
  start_group$start[i] <- sum(start_group$Freq[1:(i-1)])
}
template_start_group[template_start_group == 0] <- NA
template_end_group <- template_start_group[,2:(ncol(template_start_group)-1)] + start_group$start
template_end_group <- data.frame(group = order,template_end_group)
template_end_group_long <- template_end_group %>%
  tidyr::gather(key = "variable",
    value = "value",
    -group)
names(template_end_group_long)[3] <- "end"
template_end_group_long$start <- rep(start_group$start,
  length(unique(template_end_group_long$variable)))
template_end_group_long <- template_end_group_long %>% na.omit()
template_end_group_long$length <- sum(start_group$Freq)
template_end_group_long <- template_end_group_long[,c(2,5,4,3,1)]
template_end_group_long$group <- factor(template_end_group_long$group,levels = order)
unit <- create_unit(data = template_end_group_long,
  key = "Quickstart",
  type = "DATASET_DOMAINS",
  tree = tree)
file <- tempfile()
write_unit(unit,file)
lines <- line_clean(file=file)
sep = learn_separator(file = file)
learn_theme_label(lines,sep)

```

Description

learn linechart paramters as list

Usage

```
learn_theme_linechart(lines, sep)
```

Arguments

lines	a vector of character strings from template file.
sep	a character specifying the separator.

Value

a list of line chart parameters containing

learn_theme_piechart	<i>Learn piechart</i>
----------------------	-----------------------

Description

learn piechart paramters as list

Usage

```
learn_theme_piechart(lines, sep)
```

Arguments

lines	a vector of character strings from template file.
sep	a character specifying the separator.

Value

a list of pie chart parameters containing

`learn_theme_specific_themes`*Learn specific themes*

Description

learn specific theme paramters as list

Usage

```
learn_theme_specific_themes(lines, sep, type)
```

Arguments

<code>lines</code>	a vector of character strings from template file.
<code>sep</code>	a character specifying the separator.
<code>type</code>	template type

Value

a list of specific theme parameters containing

`learn_theme_strip_label`*Learn strip label*

Description

learn strip label paramters as list

Usage

```
learn_theme_strip_label(lines, sep)
```

Arguments

<code>lines</code>	a vector of character strings from template file.
<code>sep</code>	a charactor specifying the separator.

Value

a list of strip label parameters containing

display	0/1 specifying display or hide the individual label inside each colored strip (when defined in the data below)
width	a number specifying width of the colored strip
size	a number specifying strip label size factor (relative to the tree leaf labels)
color	define colors for each strip label element (use hexadecimal, RGB or RGBA notation; if using RGB/RGBA, COMMA cannot be used as SEPARATOR)
color_branches	1/0 specifying branches of the tree will or not be colored according to the colors of the strips above the leaves. When all children of a node have the same color, it will be colored the same, ie. the color will propagate inwards towards the root.
position	a character specifying position of the strip label within the box; 'top', 'center' or 'bottom'
shift	a number specifying strip label shift in pixels (positive or negative)
rotation	a number specifying rotation of the strip labels; used only in rectangular tree display mode
outline_width	a number specifying draw a black outline around the text (width in pixels)

Examples

```
tree <- system.file("extdata",
                    "tree_of_itol_templates.tree",
                    package = "itol.toolkit")
data("template_groups")
df_group <- data.frame(id = unique(template_groups$group),
                      data = unique(template_groups$group))

## create unit
unit <- create_unit(data = df_group,
                   key = "Quickstart",
                   type = "DATASET_COLORSTRIP",
                   tree = tree)

## write unit
file <- tempfile()
write_unit(unit, file)
## Learn parameter
lines <- line_clean(file=file)
sep = learn_separator(file = file)
learn_theme_strip_label(lines = lines, sep = sep)
```

learn_type

Learn template type

Description

Extract first line of template to learn type information.

Usage

```
learn_type(file)
```

Arguments

file template file. All the template files should follow the format rules as same with iTOL official template files. The files should start with the following headers: "COLLAPSE", "PRUNE", "SPACING", "TREE_COLORS", "DATASET_STYLE", "LABELS", "DATASET_TEXT", "DATASET_COLORSTRIP", "DATASET_BINARY", "DATASET_GRADIENT", "DATASET_HEATMAP", "DATASET_SYMBOL", "DATASET_EXTERNALSHAPE", "DATASET_DOMAINS", "DATASET_SIMPLEBAR", "DATASET_MULTIBAR", "DATASET_BOXPLOT", "DATASET_LINECHART", "DATASET_PIECHART", "DATASET_ALIGNMENT", "DATASET_CONNECTION", "DATASET_IMAGE", "POPOPUP_INFO".

Value

a character specifying header information

Examples

```
tree <- system.file("extdata",
                    "tree_of_itol_templates.tree",
                    package = "itol.toolkit")
data("template_groups")
df_group <- data.frame(id = unique(template_groups$group),
                      data = unique(template_groups$group))

## create unit
unit <- create_unit(data = df_group,
                   key = "Quickstart",
                   type = "DATASET_COLORSTRIP",
                   tree = tree)

## write unit
file <- tempfile()
write_unit(unit, file)
## Learn template type
learn_type(file)
```

line_clean

Filter out comments and empty lines

Description

Remove the lines start with # or without any information.

Usage

```
line_clean(lines = NULL, file = NULL)
```

Arguments

- lines a vector of character strings. The strings are containing the lines of template file. If the file parameter is NULL, this parameter should be set.
- file a character specifying the template file path. If this parameter is setted, the lines parameter will be replaced.

Value

a vector of character strings

Examples

```
strs <- c("#comment","DATA")
line_clean(lines=strs)
```

line_split	<i>Split lines into two parts</i>
------------	-----------------------------------

Description

Split lines based on the data block marker

Usage

```
line_split(lines, param = "data")
```

Arguments

- lines a vector of character strings from template file.
- param "theme" or "data" for the theme paramters or the data lines

Value

a vector of character strings containing data or theme information

longest_continuous_match

Calculate longest continuous match between input and target strings

Description

This function calculates the longest continuous matching substring between an input string and a target string.

Usage

```
longest_continuous_match(input_str, target_str)
```

Arguments

input_str	A character string to match against the target string.
target_str	A character string to compare with the input string.

Value

An integer value representing the longest continuous match.

merge_unit

Merge units

Description

Merge two itol.unit with same type. The second unit data will be added into the first one.

Usage

```
merge_unit(obj1, obj2)
```

Arguments

obj1	a itol.unit object specifying the first unit
obj2	a itol.unit object specifying the second unit

Value

a itol.unit object with merged data

search_tree_file	<i>Search tree file</i>
------------------	-------------------------

Description

Search Newick format tree file in dir

Usage

```
search_tree_file(  
  dir = getwd(),  
  n = "first",  
  method = "mtime",  
  max_size = 10240  
)
```

Arguments

dir	a path with tree file and other template files
n	'first', 'last', 'all'
method	sort by 'mtime', 'ctime', 'atime', 'character'
max_size	limit file size to accelerate searching

Value

a vector of characters specifying the file name

show, itol.hub-method	<i>show method for S4 class itol.hub</i>
-----------------------	--

Description

show method for S4 class itol.hub

Usage

```
## S4 method for signature 'itol.hub'  
show(object)
```

Arguments

object	An object of class itol.hub
--------	-----------------------------

Value

a stdout screen information about itol.hub object

sort_color	<i>Sort colors</i>
------------	--------------------

Description

sort colors by similarity

Usage

```
sort_color(colors, root = NULL, rev = FALSE, plot = FALSE)
```

Arguments

- colors a vector of colors in hex format
- root sort from root color
- rev sort order by similarity(default) or difference
- plot preview color order

Value

a vector of sorted colors

template_groups	<i>template groups</i>
-----------------	------------------------

Description

Templates were clustered into 5 groups by parameter similarity.

Usage

```
template_groups
```

Format

template_groups:
A data frame with template group clustering reslut:
template All the 23 template types of iTOL
group 5 clustring gourps: Tree structure: This group only controls the topology of tree branch merging, filtering, and spacing. There are no style and rich annotation data, even though most of the annotation data only include single-column id information and do not contain any dataset base information, sample information, or common and specific style information. It is a particularly simple type of template. Theme style: This does not change any topology or add any text information but only changes the color scheme, line type and width,

and font style and size of existing information. This is an extremely comprehensive and diverse type of annotation information. Text: This group contains any templates with added text information. With super flexible and convenient annotation methods, users can modify even a single character's style in HTML. Users can also modify the text annotation style of nodes and branches in batch based on matching conditions in `itol.hub` objects, which require regular expression replacement and precise data filtering. This high-frequency data processing is difficult to achieve and retain the workflow in the EXCEL-based editor. Basic plot: This group contains basic visualization methods. From a functional point of view, this is the most feature-rich class of templates. The similarity of the parameters within this part is very high. The structured and uniform organization of these templates can greatly reduce code redundancy and the user workload of data organizing. Moreover, boxplot, which is not a regular enough data annotation template, can be automatedly manipulated in R. The lack of template data structure makes using frequency unbalanced among research. Hence, the frequency of using these low-frequency templates can be increased. Advanced plot: Compared with the basic visualization methods, these visualization methods contain more comprehensive data types and often require third-party tools for input data processing. But they are the most extensible type of visualization methods for iTOL.

...

template_parameters_count

template parameters count

Description

Template types and parameters count matrix. The row names are template types. The column names are parameters short ids. The parameters are including the themes parameters and data column names. All the details are introduced in the full-page Excel file on GitHub.

Usage

```
template_parameters_count
```

Format

```
template_parameters_count:
```

A data frame with template types and parameters 0/1 count matrix:

V1 head. file type head notice

V2 separator. select the separator which is used to delimit the data below (TAB,SPACE or COMMA).This separator must be used throughout this file.

V3 dataset name. label is used in the legend table ...

train_theme	<i>Train inbuilt theme</i>
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Description

The inbuilt theme is the template of all output file and unit. Using this function can train the inbuilt theme object by custom files.

Usage

```
train_theme(dir = getwd())
```

Arguments

dir the path of tree file and template files

Value

replace the global variable inbuilt_themes

unite_rows	<i>Paste rows</i>
------------	-------------------

Description

Paste rows group by key column

Usage

```
unite_rows(df)
```

Arguments

df input data frame

Value

a data frame with pasted row by same id

use.theme	<i>Extract theme from inbuilt_themes</i>
-----------	--

Description

Extract theme from 23 template types in inbuilt_themes data in package.

Usage

```
use.theme(type, style = "default")
```

Arguments

type	a character specifying the template type used for extracting. Following choices are possible: "COLLAPSE", "PRUNE", "SPACING", "TREE_COLORS", "DATASET_STYLE", "LABEL", "DATASET_BINARY", "DATASET_GRADIENT", "DATASET_HEATMAP", "DATASET_SYMBOL", "DATASET_EXTERNALSHAPE", "DATASET_DOMAINS", "DATASET_SIMPLEBAR", "DATASET_MULTIBAR", "DATASET_BOXPLOT", "DATASET_LINECHART", "DATASET_PIECHART", "DATASET_ALIGNMENT", "DATASET_CONNECTION", "DATASET_IMAGE", "POPUP_INFO.
style	a character specifying the specific version of template type used for extracting. The default value is "default" style for all types.

Value

	a itol.theme object containing
type	This group holds information about the template type of the data only. This is a very critical piece of information. In many functions of the itol.toolkit package, the template type information is used to determine the different data processing and input/output methods.
sep	This group holds data separator information only. This is one of the most important parameters for data reading and output. It is a separate category because it is frequently used and is an input parameter for other subsequent parameters to be read.
profile	This group contains basic information about the dataset, such as the dataset name and a color label to distinguish the dataset. The dataset name is extremely important. This parameter is used almost throughout the data processing of the itol.toolkit package. With the content of this parameter as the key value, the data and theme information of the dataset are associated. In turn, high throughput learning and writing of large-scale data can be achieved. This parameter is not included in some template types with a particularly simple structure, so we choose a file name or a user-defined method as the key value.
field	This group contains information about each sample within the dataset, and this type of parameter exists only for multi-sample data. This information even includes the clustering tree between samples. This information is usually stored as part of the column names in the metadata part or abundance information of the itol.hub object.

- common_themes

These themes are used at high frequency in different templates. These parameters are small in number but constitute some common features of iTOL visual style settings, such as legend, margin, etc.
- specific_themes

These themes are used only in specific templates. The number of these parameters is very large. However, most of them are used in only one template to control the style details of the visualization. By unifying these parameters and calling them according to the template type, users can perform secondary development and data processing with a high degree of parameter aggregation without worrying too much about the differences between different template types.

Examples

```
theme <- use.theme("COLLAPSE")
```

vector_tree	<i>Vector tree</i>
-------------	--------------------

Description

build a tree by a vector of characters.

Usage

```
vector_tree(vector, branch_length = 1)
```

Arguments

- vector

input vector
- branch_length

branch length, default 1.

write_hub	<i>Write all data object into files</i>
-----------	---

Description

Write itol.hub object into template files.

Usage

```
write_hub(object, dir = getwd(), with_tree = FALSE)
```

Arguments

object	itol.hub object holds the complete data and theme information. This is an all-in-one object that collects all the information. Based on this object, it is possible to export template files directly. It can also be converted to an operation unit object for the detailed processing of individual datasets. The object can also be saved locally for reproducible visualization to share. This object contains species or sample clustering trees, sequence alignment, species abundance or gene expression table, multi-level taxonomic information, metadata, and a list of custom themes. Each element name in the theme list is prefixed with the column name of the metadata and is used to establish the association between the theme and the data. For some special dataset types, the storage location is not in the metadata, but it also conforms to the association with themes. The program automatically decides where to read the data according to the different output template types. The user only needs to explicitly define the theme name to be output consistent with the data name prefix.
dir	output dir path. Define the output files location using absolute or relative path. The template files will output by the key information from theme name in the hub object.
with_tree	output with tree file in newick format.

Value

No return value, only output template files

Examples

```
tree <- system.file("extdata",
                    "tree_of_itol_templates.tree",
                    package = "itol.toolkit")
hub <- create_hub(tree = tree)
data("template_groups")
df_group <- data.frame(id = unique(template_groups$group),
                      data = unique(template_groups$group))

## create unit
unit_1 <- create_unit(data = df_group,
                     key = "Quickstart_1",
                     type = "TREE_COLORS",
                     subtype = "clade",
                     line_type = c(rep("normal",4),"dashed"),
                     size_factor = 5,
                     tree = tree)
unit_2 <- create_unit(data = df_group,
                     key = "Quickstart_2",
                     type = "DATASET_COLORSTRIP",
                     tree = tree)

## write hub
hub <- hub + unit_1 + unit_2
write_hub(hub,tempdir())
```

write_raw	<i>Write raw data into files</i>
-----------	----------------------------------

Description

Write raw data in itol.hub object into files

Usage

```
write_raw(object, dir, title)
```

Arguments

object	itol.hub object holds the complete data and theme information. This is an all-in-one object that collects all the information. Based on this object, it is possible to export template files directly. It can also be converted to an operation unit object for the detailed processing of individual datasets. The object can also be saved locally for reproducible visualization to share. This object contains species or sample clustering trees, sequence alignment, species abundance or gene expression table, multi-level taxonomic information, metadata, and a list of custom themes. Each element name in the theme list is prefixed with the column name of the metadata and is used to establish the association between the theme and the data. For some special dataset types, the storage location is not in the metadata, but it also conforms to the association with themes. The program automatically decides where to read the data according to the different output template types. The user only needs to explicitly define the theme name to be output consistent with the data name prefix.
dir	output dir path. Define the output files location using absolute or relative path. The raw data will write into files. The following raw data will be outputted: main tree, sample tree, alignment sequences, abundance count table, taxonomy table, metadata on nodes and tips.
title	files name title string. This character specified the prefix of raw data files.

Value

No return value, only output raw data files

Examples

```
tree <- system.file("extdata",
                    "tree_of_itol_templates.tree",
                    package = "itol.toolkit")
hub <- create_hub(tree = tree)
df_values <- data.table::fread(system.file("extdata",
                                          "templates_frequence.txt",
                                          package = "itol.toolkit"))
unit <- create_unit(data = df_values,
```

```

      key = "Quickstart",
      type = "DATASET_HEATMAP",
      tree = tree)
hub <- hub + unit
write_raw(hub, tempdir(), "Quickstart")

```

write_unit

*Write unit object into file***Description**

Write itol.unit object into template file. This function will using the type information in unit object to decide different output methods for the template formats.

Usage

```
write_unit(unit, file = getwd())
```

Arguments

unit	unit object. The unit object holds the data and theme of a single dataset. This is the smallest data operation unit. At this level, individual data can be fine-tuned. It is also possible to extract the style of a unit for use in other units. It is also possible to use many units to learn a complete itol.hub object. Almost all specific data operations behind the itol.toolkit package are performed at the unit level. Because itol.hub objects have comprehensive information, but to ensure that the correspondence with phylogenetic branches or nodes remains consistent when different data types are saved, many complex data aggregations are saved, which does not facilitate data processing. Therefore, in the actual data processing process, unit objects are generated from the itol.hub object and then processed.
file	output file path. Define the output file location and file name using absolute or relative path.

Value

No return value, only output a template file

Examples

```

tree <- system.file("extdata",
                    "tree_of_itol_templates.tree",
                    package = "itol.toolkit")
data("template_groups")
df_group <- data.frame(id = unique(template_groups$group),
                      data = unique(template_groups$group))
## create unit
unit <- create_unit(data = df_group,

```

```
        key = "Quickstart",  
        type = "DATASET_COLORSTRIP",  
        tree = tree)  
## write unit  
write_unit(unit,tempfile())
```

Index

- * **datasets**
 - inbuilt_themes, [23](#)
 - template_groups, [50](#)
 - template_parameters_count, [51](#)
- * **objects**
 - itol.hub-class, [24](#)
 - itol.theme-class, [24](#)
 - itol.unit-class, [25](#)
- * **object**
 - create_hub, [12](#)
 - create_theme, [13](#)
 - file_to_unit, [20](#)
 - hub_to_unit, [22](#)
- +, itol.hub, itol.unit-method, [3](#)
- +, itol.unit, itol.unit-method
 - (+, itol.hub, itol.unit-method), [3](#)
- anno_col_group, [4](#)
- binary_data_conversion, [5](#)
- color_distance, [6](#)
- complex_html_text, [6](#)
- convert_01, [7](#)
- convert_01_to_connect, [7](#)
- convert_range_to_node, [8](#)
- convert_to_binary, [8](#)
- correct_get_color, [10](#)
- correct_type, [11](#)
- count_to_tree, [11](#)
- create_hub, [12](#)
- create_theme, [13](#)
- create_unit, [13](#)
- darken_color, [16](#)
- df_merge, [17](#)
- df_tree, [17](#)
- dt_transpose_header, [18](#)
- fa_read, [18](#)
- fa_write, [19](#)
- file_get_dir, [19](#)
- file_get_name, [20](#)
- file_to_unit, [20](#)
- get_color, [21](#)
- gradient_color, [21](#)
- head_line, [22](#)
- hub_to_unit, [22](#)
- inbuilt_themes, [23](#)
- itol.hub (itol.hub-class), [24](#)
- itol.hub-class, [24](#)
- itol.theme (itol.theme-class), [24](#)
- itol.theme-class, [24](#)
- itol.unit (itol.unit-class), [25](#)
- itol.unit-class, [25](#)
- learn_data, [25](#)
- learn_data_from_file, [26](#)
- learn_data_from_files, [26](#)
- learn_data_from_unit, [27](#)
- learn_data_from_unit_list, [27](#)
- learn_df, [28](#)
- learn_field, [28](#)
- learn_legend, [29](#)
- learn_line, [31](#)
- learn_profile, [32](#)
- learn_separator, [33](#)
- learn_subdf, [34](#)
- learn_theme_align, [34](#)
- learn_theme_alignment, [35](#)
- learn_theme_bar, [35](#)
- learn_theme_basic_plot, [36](#)
- learn_theme_basic_theme, [36](#)
- learn_theme_binary, [37](#)
- learn_theme_border, [37](#)
- learn_theme_common_themes, [38](#)
- learn_theme_connection, [38](#)

`learn_theme_domain`, [39](#)
`learn_theme_externalshape`, [39](#)
`learn_theme_heatmap`, [40](#)
`learn_theme_image`, [40](#)
`learn_theme_label`, [41](#)
`learn_theme_linechart`, [42](#)
`learn_theme_piechart`, [43](#)
`learn_theme_specific_themes`, [44](#)
`learn_theme_strip_label`, [44](#)
`learn_type`, [45](#)
`line_clean`, [46](#)
`line_split`, [47](#)
`longest_continuous_match`, [48](#)

`merge_unit`, [48](#)

`search_tree_file`, [49](#)
`show`, `itol.hub-method`, [49](#)
`sort_color`, [50](#)

`template_groups`, [50](#)
`template_parameters_count`, [51](#)
`train_theme`, [52](#)

`unite_rows`, [52](#)
`use.theme`, [53](#)

`vector_tree`, [54](#)

`write_hub`, [54](#)
`write_raw`, [56](#)
`write_unit`, [57](#)