

# Package ‘bpa’

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**Type** Package

**Title** Basic Pattern Analysis

**Version** 0.1.1

**Date** 2016-04-03

**Description** Run basic pattern analyses on character sets, digits, or combined input containing both characters and numeric digits. Useful for data cleaning and for identifying columns containing multiple or nonstandard formats.

**Depends** base

**Imports** magrittr, plyr

**Suggests** testthat, knitr, rmarkdown

**License** GPL (>= 2)

**URL** <https://github.com/bgreenwell/bpa>

**BugReports** <https://github.com/bgreenwell/bpa/issues>

**RoxygenNote** 5.0.1

**VignetteBuilder** knitr

**NeedsCompilation** no

**Author** Brandon Greenwell [aut, cre]

**Maintainer** Brandon Greenwell <[greenwell.brandon@gmail.com](mailto:greenwell.brandon@gmail.com)>

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`get_pattern`*Basic Pattern Analysis*

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**Description**

Perform a basic pattern analysis

**Usage**

```
get_pattern(x, show_ws = TRUE, ws_char = "w")

basic_pattern_analysis(x, unique_only = FALSE, show_ws = TRUE,
  ws_char = "w", useNA = c("no", "ifany", "always"), ...)

## Default S3 method:
basic_pattern_analysis(x, unique_only = FALSE,
  show_ws = TRUE, ws_char = "w", useNA = c("no", "ifany", "always"), ...)

## S3 method for class 'data.frame'
basic_pattern_analysis(x, unique_only = FALSE,
  show_ws = TRUE, ws_char = "w", useNA = c("no", "ifany", "always"), ...)

bpa(x, ...)
```

**Arguments**

<code>x</code>	A data frame or character vector.
<code>show_ws</code>	Logical indicating whether or not to show whitespace using a special character. Default is TRUE.
<code>ws_char</code>	Character string to use to depict whitespace when <code>show_ws = TRUE</code> .
<code>unique_only</code>	Logical indicating whether or not to only show the unique patterns. Default is TRUE.
<code>useNA</code>	Logical indicating whether to include NA values in the table. See <a href="#">table</a> for details.
<code>...</code>	Additional optional arguments to be passed onto <code>l1ply</code> .

**Examples**

```
basic_pattern_analysis(iris)
basic_pattern_analysis(iris, unique_only = TRUE)
```

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match_pattern	<i>Pattern Matching</i>
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**Description**

Extract values from a vector that match a particular pattern.

**Usage**

```
match_pattern(x, pattern, unique_only = FALSE, ...)
```

**Arguments**

x	A vector, typically of class "character".
pattern	Character string specifying the particular pattern to match.
unique_only	Logical indicating whether or not to only return unique values. Default is FALSE.
...	Additional optional arguments to be passed onto <a href="#">get_pattern</a> .

**Details**

The pattern specified by the required argument pattern must be a valid pattern produced by the `get_pattern` function. That is, all digits should be represented by a "9", lowercase/uppercase letters by a "a"/"A", etc.

**Examples**

```
phone <- c("123-456-7890", "456-7890", "123-4567", "456-7890")
match_pattern(phone, pattern = "999-9999")
match_pattern(phone, pattern = "999-9999", unique_only = TRUE)
```

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messy	<i>Simulated Data</i>
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**Description**

Simulated (messy) data set to help illustrate some of the uses of basic pattern analysis.

**Format**

A data frame with 1000 rows and 3 variables

**Details**

- Gender Gender in various formats.
- Date Dates in various formats.
- Phone Phone numbers in various formats.

**Examples**

```
data(messy)
bpa(messy, unique_only = TRUE, ws_char = " ")
```

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`trim_ws`*Remove Leading/Trailing Whitespace*

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**Description**

Remove leading and/or trailing whitespace from character strings.

**Usage**

```
trim_ws(x, which = c("both", "left", "right"))
```

**Arguments**

<code>x</code>	A data frame or vector.
<code>which</code>	A character string specifying whether to remove both leading and trailing whitespace (default), or only leading ("left") or trailing ("right"). Can be abbreviated.

**Examples**

```
# Toy example
d <- data.frame(x = c(" a ", "b ", "c"),
               y = c(" 1 ", "2", " 3"),
               z = c(4, 5, 6))
print(d) # print data as is
trim_ws(d) # print data with whitespace trimmed off
sapply(trim_ws(d), class) # check that column types are preserved
```

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