Package 'edm1.sequence'

July 29, 2024

Title Set of tools to manipulate time series

Version 2.0.0.0

Description Provides set of functions to manipulate time series dataset; create variables that designates the value of the individual at n-x , handle missing values...

License GPL (==3)

Encoding UTF-8

Roxygen list(markdown = TRUE)

RoxygenNote 7.3.1

Imports stringr

Contents

`historic_sequence1` *historic_sequence1*

Description

Allow to perform a pivot wider on a sequencial dataset (here the type is dataframe), each variable will be dupplicated in a column to show the value to this variable at n - 1 for each individual, see examples.

Usage

```
historic_sequence1(inpt_datf, bf_ = 1)
```

Arguments

`inpt_datf`	is the input dataframe
`bf_`	is the number of previous value of the individual it will search for, see examples

Examples

```
set.seed(123)
var1 <- round(runif(n = 14, min = 100, max = 122))
set.seed(123)
var2 <- round(runif(n = 14, min = 14, max = 20))

datf <- data.frame("ids" = c(20, 20, 20, 20, 19, 19, 19, 18, 18, 18, 18,
                             17, 17, 17),
                   "individual" = c("oui", "non", "peut1", "peut2",
                                    "oui", "peut1", "peut2"),
                   "var1" = var1,
                   "var2" = var2)
print(datf)
```

```
   ids individual var1 var2
1   20        oui  106   16
2   20        non  117   19
3   20      peut1  109   16
4   20      peut2  119   19
5   19        oui  121   20
6   19      peut1  101   14
7   19      peut2  112   17
8   18        oui  120   19
9   18        non  112   17
10  18      peut1  110   17
11  18      peut2  121   20
12  17        oui  110   17
13  17      peut1  115   18
14  17      peut2  113   17
```

```
historic_sequence1(inpt_datf = datf, bf_ = 2)
```

```
  id_seq individual var1-1 var1-2 var2-1 var2-2
1     20        oui    121    120     20     19
2     20        non     NA    112     NA     17
3     20      peut1    101    110     14     17
4     20      peut2    112    121     17     20
5     19        oui    120    110     19     17
6     19      peut1    110    115     17     18
7     19      peut2    121    113     20     17
```

```
historic_sequence1(inpt_datf = datf, bf_ = 3)
```

```
  id_seq individual var1-1 var1-2 var1-3 var2-1 var2-2 var2-3
1     20        oui    121    120    110     20     19     17
2     20        non     NA    112     NA     NA     17     NA
3     20      peut1    101    110    115     14     17     18
4     20      peut2    112    121    113     17     20     17
```

`historic_sequence2` *historic_sequence2*

Description

Allow to perform a pivot wider on a sequencial dataset (here the type is dataframe), each variable will be dupplicated in a column to show the value to this variable at n - 1 for each individual, see examples.

Usage

```
historic_sequence2(inpt_datf, bf_ = 1)
```

Arguments

`inpt_datf`	is the input dataframe
`bf_`	is the number of previous value of the individual it will search for, see examples

Examples

```
set.seed(123)
var1 <- round(runif(n = 14, min = 100, max = 122))
set.seed(123)
var2 <- round(runif(n = 14, min = 14, max = 20))

datf <- data.frame("ids" = c(20, 20, 20, 20, 19, 19, 19, 18, 18, 18, 18,
                             17, 17, 17),
                   "individual" = c("oui", "non", "peut1", "peut2",
                                    "oui", "peut1", "peut2"),
                   "var1" = var1,
                   "var2" = var2)
print(datf)
```

```
   ids individual var1 var2
1   20        oui  106   16
2   20        non  117   19
3   20      peut1  109   16
4   20      peut2  119   19
5   19        oui  121   20
6   19      peut1  101   14
7   19      peut2  112   17
8   18        oui  120   19
9   18        non  112   17
10  18      peut1  110   17
11  18      peut2  121   20
12  17        oui  110   17
13  17      peut1  115   18
14  17      peut2  113   17
```

```
print(historic_sequence2(inpt_datf = datf, bf_ = 2))
```

```
  id_seq individual var1-0 var1-1 var1-2 var2-0 var2-1 var2-2
1     20        oui    106    121    120     16     20     19
2     20        non    117     NA    112     19     NA     17
3     20      peut1    109    101    110     16     14     17
4     20      peut2    119    112    121     19     17     20
5     19        oui    121    120    110     20     19     17
6     19      peut1    101    110    115     14     17     18
7     19      peut2    112    121    113     17     20     17
```

```
print(historic_sequence2(inpt_datf = datf, bf_ = 3))

  id_seq individual var1-0 var1-1 var1-2 var1-3 var2-0 var2-1 var2-2 var2-3
1     20        oui    106    121    120    110     16     20     19     17
2     20        non    117     NA    112     NA     19     NA     17     NA
3     20      peut1    109    101    110    115     16     14     17     18
4     20      peut2    119    112    121    113     19     17     20     17
```

`sequence_na_mean1` *sequence_na_mean1*

Description

In a dataframe generated by the function historic_sequence1, convert all NA to the mean of the values at the same variable for the individual at the id where the NA occurs, see examples (only accepts numeric variables)

Usage

```
sequence_na_mean1(inpt_datf, bf_, step = 1)
```

Arguments

`inpt_datf`	is the input dataframe

Examples

```
set.seed(123)
var1 <- round(runif(n = 14, min = 100, max = 122))
set.seed(123)
var2 <- round(runif(n = 14, min = 14, max = 20))

datf <- data.frame("ids" = c(20, 20, 20, 20, 19, 19, 19, 18, 18, 18, 18,
17, 17, 17),
"individual" = c("oui", "non", "peut1", "peut2",
"oui", "peut1", "peut2"),
"var1" = var1,
"var2" = var2)
datf <- historic_sequence1(inpt_datf = datf, bf_ = 2)
datf[3, 4] <- NA
datf[6, 4] <- NA
datf[1, 3] <- NA
print(datf)

  id_seq individual var1-1 var1-2 var2-1 var2-2
1     20        oui     NA    120     20     19
2     20        non     NA    112     NA     17
3     20      peut1    101     NA     14     17
4     20      peut2    112    121     17     20
5     19        oui    120    110     19     17
6     19      peut1    110     NA     17     18
7     19      peut2    121    113     20     17
```

```
print(sequence_na_mean1(inpt_datf = datf, bf_ = 2))
```

```
  id_seq individual var1-1 var1-2 var2-1 var2-2
1     20        oui    115  120.0     20     19
2     20        non    112  112.0     17     17
3     20      peut1    101  105.5     14     17
4     20      peut2    112  121.0     17     20
5     19        oui    120  110.0     19     17
6     19      peut1    110  105.5     17     18
7     19      peut2    121  113.0     20     17
```

`sequence_na_mean2` *sequence_na_mean2*

Description

In a dataframe generated by the function historic_sequence1, convert all NA to the mean of the values at the same variable for the individual at the id where the NA occurs, see examples (only accepts numeric variables)

Usage

```
sequence_na_mean2(inpt_datf, bf_, step = 1)
```

Arguments

`inpt_datf`	is the input dataframe

Examples

```
set.seed(123)
var1 <- round(runif(n = 14, min = 100, max = 122))
set.seed(123)
var2 <- round(runif(n = 14, min = 14, max = 20))

datf <- data.frame("ids" = c(20, 20, 20, 20, 19, 19, 19, 18, 18, 18, 18,
17, 17, 17),
"individual" = c("oui", "non", "peut1", "peut2",
"oui", "peut1", "peut2"),
"var1" = var1,
"var2" = var2)
datf <- historic_sequence2(inpt_datf = datf, bf_ = 2)
datf[3, 4] <- NA
datf[6, 4] <- NA
datf[1, 3] <- NA
print(datf)
```

```
  id_seq individual var1-0 var1-1 var1-2 var2-0 var2-1 var2-2
1     20        oui     NA    121    120     16     NA     19
2     20        non    117     NA    112     19     NA     17
3     20      peut1    109     NA    110     16     14     17
4     20      peut2    119    112    121     19     17     20
5     19        oui    121    120    110     20     19     17
```

```
6     19      peut1    101     NA    115     14     17     18
7     19      peut2    112    121    113     17     20     17

print(sequence_na_mean2(inpt_datf = datf, bf_ = 2))

  id_seq individual var1-0   var1-1 var1-2 var2-0 var2-1 var2-2
1     20        oui    117 121.0000    120     16     18     19
2     20        non    117 114.5000    112     19     18     17
3     20      peut1    109 108.3333    110     16     14     17
4     20      peut2    119 112.0000    121     19     17     20
5     19        oui    121 120.0000    110     20     19     17
6     19      peut1    101 108.3333    115     14     17     18
7     19      peut2    112 121.0000    113     17     20     17
```

`sequence_na_med1` *sequence_na_med1*

Description

In a dataframe generated by the function historic_sequence1, convert all NA to the median of the values at the same variable for the individual at the id where the NA occurs, see examples (only accepts numeric variables)

Usage

```
sequence_na_med1(inpt_datf, bf_, step = 1)
```

Arguments

`inpt_datf`	is the input dataframe
`bf_`	is how at how many n - -1 we look for the value of the variables for the individual at time index n
`step`	is the base step for the time indexes, step of one year (or any time unit), two year (or any time unit)?

Examples

```
set.seed(123)
var1 <- round(runif(n = 14, min = 100, max = 122))
set.seed(123)
var2 <- round(runif(n = 14, min = 14, max = 20))

datf <- data.frame("ids" = c(20, 20, 20, 20, 19, 19, 19, 18, 18, 18, 18,
17, 17, 17),
"individual" = c("oui", "non", "peut1", "peut2",
"oui", "peut1", "peut2"),
"var1" = var1,
"var2" = var2)
datf <- historic_sequence1(inpt_datf = datf, bf_ = 2)
datf[3, 4] <- NA
datf[6, 4] <- NA
datf[1, 3] <- NA
```

```
print(datf)
```

```
  id_seq individual var1-1 var1-2 var2-1 var2-2
1     20        oui     NA    120     20     19
2     20        non     NA    112     NA     17
3     20      peut1    101     NA     14     17
4     20      peut2    112    121     17     20
5     19        oui    120    110     19     17
6     19      peut1    110     NA     17     18
7     19      peut2    121    113     20     17
```

```
print(sequence_na_med1(inpt_datf = datf, bf_ = 2))
```

```
  id_seq individual var1-1 var1-2 var2-1 var2-2
1     20        oui    115  120.0     20     19
2     20        non    112  112.0     17     17
3     20      peut1    101  105.5     14     17
4     20      peut2    112  121.0     17     20
5     19        oui    120  110.0     19     17
6     19      peut1    110  105.5     17     18
7     19      peut2    121  113.0     20     17
```

`sequence_na_med2` *sequence_na_med2*

Description

In a dataframe generated by the function historic_sequence2, convert all NA to the median of the values at the same variable for the individual at the id where the NA occurs, see examples (only accepts numeric variables)

Usage

```
sequence_na_med2(inpt_datf, bf_, step = 1)
```

Arguments

`inpt_datf`	is the input dataframe
`bf_`	is how at how many n - -1 we look for the value of the variables for the individual at time index n
`step`	is the base step for the time indexes, step of one year (or any time unit), two year (or any time unit)?

Examples

```
set.seed(123)
var1 <- round(runif(n = 14, min = 100, max = 122))
set.seed(123)
var2 <- round(runif(n = 14, min = 14, max = 20))
datf <- data.frame("ids" = c(20, 20, 20, 20, 19, 19, 19, 18, 18, 18, 18,
17, 17, 17),
"individual" = c("oui", "non", "peut1", "peut2",
```

```
"oui", "peut1", "peut2"),
"var1" = var1,
"var2" = var2)
datf <- historic_sequence2(inpt_datf = datf, bf_ = 2)
datf[3, 4] <- NA
datf[6, 4] <- NA
datf[1, 3] <- NA
print(datf)
```

```
  id_seq individual var1-0 var1-1 var1-2 var2-0 var2-1 var2-2
1     20        oui     NA    121    120     16     20     19
2     20        non    117     NA    112     19     NA     17
3     20      peut1    109     NA    110     16     14     17
4     20      peut2    119    112    121     19     17     20
5     19        oui    121    120    110     20     19     17
6     19      peut1    101     NA    115     14     17     18
7     19      peut2    112    121    113     17     20     17
```

```
print(sequence_na_med2(inpt_datf = datf, bf_ = 2))
```

```
  id_seq individual var1-0 var1-1 var1-2 var2-0 var2-1 var2-2
1     20        oui    120  121.0    120     16     20     19
2     20        non    117  114.5    112     19     18     17
3     20      peut1    109  109.0    110     16     14     17
4     20      peut2    119  112.0    121     19     17     20
5     19        oui    121  120.0    110     20     19     17
6     19      peut1    101  109.0    115     14     17     18
7     19      peut2    112  121.0    113     17     20     17
```

Index