## Package 'edm1.sequence'

July 29, 2024

<b>Title</b> Set of tools to manipulate time series
<b>Version</b> 2.0.0.0
<b>Description</b> Provides set of functions to manipulate time series dataset; create variables that designates the value of the individual at n-x, handle missing values
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Encoding UTF-8
<b>Roxygen</b> list(markdown = TRUE)
RoxygenNote 7.3.1
Imports stringr
Contents  historic_sequence
historic_sequence historic_sequence
<b>Description</b> Allow to perform a pivot wider on a sequencial dataset (here the type is dataframe), each variable

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_sequence(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

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#### **Examples**

```
set.seed(123)
var1 < - round(runif(n = 14, min = 100, max = 122))
set.seed(123)
var2 \leftarrow 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
  20 oui 106 16
2 20
           non 117
                      19
3 20
         peut1 109
                     16
4
  20
         peut2 119 19
  19
           oui 121
5
                      20
  19
6
         peut1 101
                      14
  19
7
         peut2 112
                      17
  18
8
          oui 120
                      19
          non 112
peut1 110
peut2 121
oui 110
peut1 115
9 18
10 18
11 18
                      17
                      17
                      20
12 17
                      17
13 17
                      18
          peut2 113
14 17
                     17
historic_sequence(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
                    NA
                          112
                                 NA
             non
3
     20
                   101
                          110
            peut1
                                 14
                                        17
     20
            peut2 112
                          121
                                 17
                                        20
5
     19
             oui
                   120
                         110
                                 19
                                       17
6
     19
            peut1
                    110
                         115
                                 17
                                       18
7
                   121
                          113
                                 20
     19
            peut2
                                        17
historic_sequence(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
2
     20
                     NA
                           112
                                 NA
                                         NA
                                               17
              non
                                                     NA
          peut1
3
     20
                    101
                           110
                                 115
                                        14
                                               17
                                                     18
                          121
     20
            peut2
                    112
                                 113
                                        17
                                               20
                                                    17
```

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