Analyse Statistique avec Packages Explore

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```
if (!require(funModeling)) install.packages("funModeling")
```

```
## Le chargement a nécessité le package : funModeling
```

```
## Le chargement a nécessité le package : Hmisc
```

```
##
## Attachement du package : 'Hmisc'
```

```
## Les objets suivants sont masqués depuis 'package:base':
##
##     format.pval, units
```

```
## funModeling v.1.9.5 :)
## Examples and tutorials at livebook.datascienceheroes.com
##  / Now in Spanish: librovivodecienciadedatos.ai
```

```
library(funModeling)
data(heart_disease)
```

```
str(heart_disease)
```

```
## 'data.frame':    303 obs. of  16 variables:
##  $ age                   : int  63 67 67 37 41 56 62 57 63 53 ...
##  $ gender                : Factor w/ 2 levels "female","male": 2 2 2 2 1 2 1 1 2 2 ...
##  $ chest_pain            : Factor w/ 4 levels "1","2","3","4": 1 4 4 3 2 2 4 4 4 4 ...
##  $ resting_blood_pressure: int  145 160 120 130 130 120 140 120 130 140 ...
##  $ serum_cholestoral     : int  233 286 229 250 204 236 268 354 254 203 ...
##  $ fasting_blood_sugar   : Factor w/ 2 levels "0","1": 2 1 1 1 1 1 1 1 1 2 ...
##  $ resting_electro       : Factor w/ 3 levels "0","1","2": 3 3 3 1 3 1 3 1 3 3 ...
##  $ max_heart_rate        : int  150 108 129 187 172 178 160 163 147 155 ...
##  $ exer_angina           : int  0 1 1 0 0 0 0 1 0 1 ...
##  $ oldpeak               : num  2.3 1.5 2.6 3.5 1.4 0.8 3.6 0.6 1.4 3.1 ...
##  $ slope                 : int  3 2 2 3 1 1 3 1 2 3 ...
##  $ num_vessels_flour     : int  0 3 2 0 0 0 2 0 1 0 ...
##  $ thal                  : Factor w/ 3 levels "3","6","7": 2 1 3 1 1 1 1 1 3 3 ...
##  $ heart_disease_severity: int  0 2 1 0 0 0 3 0 2 1 ...
##  $ exter_angina          : Factor w/ 2 levels "0","1": 1 2 2 1 1 1 1 2 1 2 ...
##  $ has_heart_disease     : Factor w/ 2 levels "no","yes": 1 2 2 1 1 1 2 1 2 2 ...
```

```r
if (!require(dplyr)) install.packages("explore")
```

```
## Le chargement a nécessité le package : dplyr
```

```
##
## Attachement du package : 'dplyr'
```

```
## Les objets suivants sont masqués depuis 'package:Hmisc':
##
##     src, summarize
```

```
## Les objets suivants sont masqués depuis 'package:stats':
##
##     filter, lag
```

```
## Les objets suivants sont masqués depuis 'package:base':
##
##     intersect, setdiff, setequal, union
```

```r
#if (!require(dplyr)) install.packages("dyplr")
if (!require(here)) install.packages("here")
```

```
## Le chargement a nécessité le package : here
```

```
## here() starts at C:/Users/User/Documents/Apprendre_shiny/Analyse_pdf/Stat_descriptive
```

```r
# if (!require(dplyr)) install.packages("explore")
#install.packages("dyplr")
#install.packages("here")
```

```r
library(explore)
```

```
##
## Attachement du package : 'explore'
```

```
## L'objet suivant est masqué depuis 'package:Hmisc':
##
##     describe
```

```r
#library(dyplr)
library(here)
```

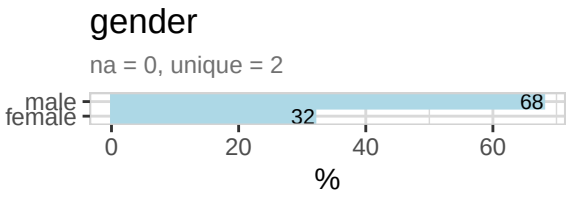
```r
explore(heart_disease)
```

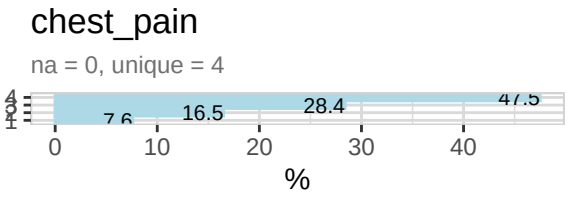
```
## Warning in explore_shiny(data, ...): This function can only be used in an
## interactive R session
```

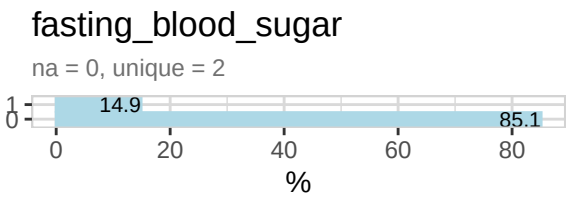
```
heart_disease %>%
  select(where(is.numeric)) %>%
  explore_all()
```

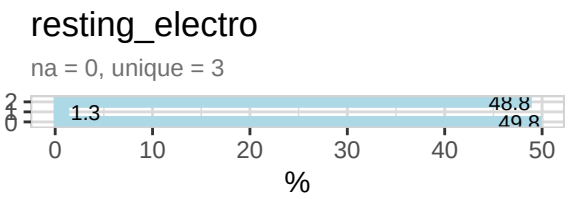
heart_disease %>% # sélection des variables de type facteur select(where(is.factor)) %>% explore_all()

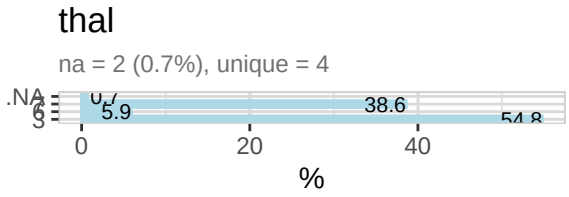
```
heart_disease %>%
  # sélection des variables de type facteur
  select(where(is.factor)) %>%
  explore_all()
```

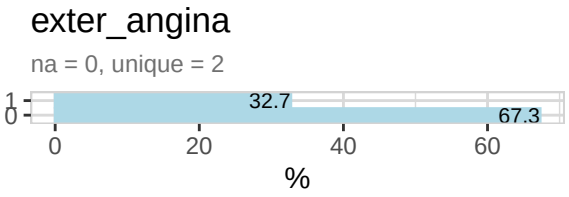

gender
na = 0, unique = 2
male
female
68
32
0
20
40
60
%

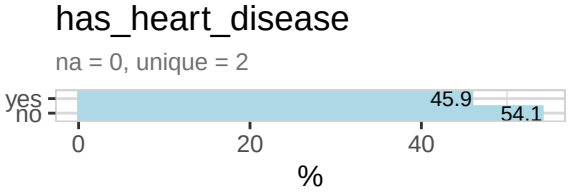

chest_pain
na = 0, unique = 4
4
3
2
1
47.5
28.4
16.5
7.6
0
10
20
30
40
%


fasting_blood_sugar
na = 0, unique = 2
1
0
14.9
85.1
0
20
40
60
80
%


resting_electro
na = 0, unique = 3
2
1
0
48.8
1.3
49.8
0
10
20
30
40
50
%


thal
na = 2 (0.7%), unique = 4
.NA
7
6
3
5.9
38.6
0
20
40
%


exter_angina
na = 0, unique = 2
1
0
32.7
67.3
0
20
40
60
%


has_heart_disease
na = 0, unique = 2
yes
no
45.9
54.1
0
20
40
%