June 2, 2023

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1 Setup

Importiamo le librerie qui usate

```
import numpy as np
import pandas as pd
import pylbmisc as lb

# Per la visualizzazione utilizziamo seaborn. un po di setup
import matplotlib.pyplot as plt
import seaborn as sns
sns.set_theme()
iris = sns.load_dataset('iris')
```

Queste importazioni doppie servono per gli ambienti pyconsole, isolati dal resto:

```
>>> import numpy as np
>>> import pandas as pd
```

2 Info generali

```
>>> df = pd.DataFrame(np.random.randn(1000, 5), columns=["a", "b", "c", "d", "e"])
>>> df[::2] = np.nan
>>> df.info()
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1000 entries, 0 to 999
Data columns (total 5 columns):
    Column Non-Null Count Dtype
            500 non-null
 1
            500 non-null float64
 2
            500 non-null float64
 3
            500 non-null float64
 4
            500 non-null
                            float64
dtypes: float64(5)
memory usage: 39.2 KB
>>> df.head()
          а
                   b
                             С
                                       d
                                                 е
0
        NaN
                 NaN
                           NaN
                                     NaN
                                               NaN
  1.570055 -0.393051 0.880055 -0.032189 -0.161610
       NaN
                 NaN
                           NaN
                                     NaN
3 -1.392931 -0.274947
                      0.719258 0.093073 -1.661034
       NaN
                 NaN
                           NaN
                                     NaN
                                               NaN
```

3 Univariate

3.1 Variabili numeriche

3.1.1 Statistiche numeriche

Usare il metodo describe e fare trasposizione.

```
>>> df = pd.DataFrame(np.random.randn(1000, 5), columns=["a", "b", "c", "d", "e"])
>>> df.describe().transpose() # count sono i valori non mancanti (qui tutti)
                                             25%
    count
              mean
                         std
                                   min
                                                       50%
                                                                75%
  1000.0 0.004130 1.062286 -3.621185 -0.643718 -0.040742
                                                           0.693380
                                                                     3.206745
  1000.0 0.000216 1.007270 -3.297063 -0.666956 0.054856
                                                           0.671322
                                                                     3.084294
  1000.0 -0.002650 0.994217 -3.316414 -0.623244 -0.020238 0.614408
                                                                     3.665185
  1000.0 -0.053436 1.023828 -3.017280 -0.745594 -0.025781 0.628180 3.064176
  1000.0 -0.038803 1.016394 -3.472263 -0.726240 -0.046140 0.633126 2.721213
```

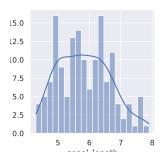


Figure 1: Istogramma

3.1.2 Istogramma

```
In figura 1.
```

```
# plot = sns.histplot(data=iris, x="sepal_length", kde=True, bins=20)
plot = sns.histplot(x=iris.sepal_length, kde=True, bins=20)
fig = plot.get_figure()
lb.fig.dump(fig, label="sns_histo", caption = 'Istogramma', scale = 0.5)
plt.clf() # pulire figura se no la ritroviamo sovrapposta
```

3.2 Categoriche

3.2.1 Frequenze

```
>>> df = pd.DataFrame({"x": ["a", "a", "a", "a", np.nan, "b", "b"],
                        "y": ["1", "2", "1", "2", "2", "1", "2"]})
>>> df.x.value_counts()
\mathbb{X}
а
     2
b
Name: count, dtype: int64
>>> df.x.value_counts(dropna=False)
X
       4
b
       2
       1
NaN
Name: count, dtype: int64
```

3.2.2 Diagramma a barre

```
Figura ??
```

```
tips = sns.load_dataset("tips")
plot = sns.barplot(
```

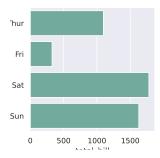


Figure 2: Diagramma a barre

```
x="total_bill",
y="day",
data=tips,
estimator=sum,
ci=None,
color='#69b3a2');
fig = plot.get_figure()
lb.fig.dump(fig, label="sns_barre", caption = 'Diagramma a barre', scale = 0.5)
plt.clf()
```

4 Bivariate

4.1 Numeriche stratificate

```
>>> df = pd.DataFrame({"x": np.random.randn(7),
                       "y": np.random.randn(7),
. . .
                       "z": np.random.randn(7),
. . .
                       "g": ["trt", "ctrl", "trt", "ctrl", "trt", "ctrl", "trt"]})
>>> spl = df.groupby("g")
>>> spl.describe().transpose() # descrizione complesiva
             ctrl
x count 3.000000 4.000000
 mean 0.232905 0.207106
        1.053657 1.007551
 std
       -0.790423 -1.142858
 min
  25%
       -0.307880 -0.093417
 50%
        0.174663 0.340478
 75%
        0.744569 0.641000
        1.314474 1.290324
 max
y count 3.000000 4.000000
```

```
0.287371 -0.938949
  mean
        0.986902 0.778552
  std
        -0.794455 -1.824555
 min
       -0.138180 -1.445004
  25%
  50%
        0.518094 -0.918370
        0.828283 -0.412315
  75%
        1.138472 -0.094499
 max
z count 3.000000 4.000000
        0.564759 -0.103995
 mean
        1.168875 1.536222
  std
        -0.693046 -2.195190
 min
  25%
        0.038344 -0.768697
 50%
        0.769734 0.277340
        1.193661 0.942042
 75%
 max
        1.617589 1.224528
>>> sel = ["x", "z"] # descrizione di solo alcune colonne
>>> spl[sel].describe().transpose()
             ctrl
                        trt
x count 3.000000 4.000000
 mean
        0.232905 0.207106
        1.053657 1.007551
  std
       -0.790423 -1.142858
 min
  25%
       -0.307880 -0.093417
  50%
        0.174663 0.340478
 75%
        0.744569 0.641000
 max
        1.314474 1.290324
z count 3.000000 4.000000
        0.564759 -0.103995
 mean
        1.168875 1.536222
  std
       -0.693046 -2.195190
 min
  25%
        0.038344 -0.768697
  50%
        0.769734 0.277340
  75%
        1.193661 0.942042
 max
        1.617589 1.224528
4.2
     Tabelle di contingenza
>>> df = pd.DataFrame({"x": ["a", "a", "a", "a", "a", "b", "b"],
                       "y": ["1", "2", "2", "2", "2", "1", "2"],
                       "g": ["trt", "ctrl", "trt", "ctrl", "trt", "ctrl", "trt"]})
. . .
>>> pd.crosstab(df.x, df.y, margins=True) # frequenze schiette con totali
    1 2 All
У
            5
    1 4
```

1 1

```
All 2 5 7
>>> pd.crosstab(df.x, df.y, margins=True, normalize = 'columns') # percentuali di colonna
y 1 2 All
x
a 0.5 0.8 0.714286
b 0.5 0.2 0.285714
```

4.3 Tabella trial

Utilizzare la libreria tableone.

```
>>> import tableone
>>> df = tableone.load_dataset('pn2012')
>>> df.info()
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1000 entries, 0 to 999
Data columns (total 8 columns):
# Column Non-Null Count Dtype
             1000 non-null int64
0
    Age
    SysABP
             709 non-null float64
1
2 Height
             525 non-null float64
3
   Weight
             698 non-null float64
             1000 non-null object
4
    ICU
5
   MechVent 1000 non-null int64
   LOS
            1000 non-null int64
             1000 non-null int64
7
   death
dtypes: float64(3), int64(4), object(1)
memory usage: 62.6+ KB
>>> df.head()
  Age SysABP Height Weight ICU MechVent LOS death
   54
        NaN
                NaN
                       NaN SICU
                                         0
                                             5
1
   76
      105.0
              175.3
                      80.6 CSRU
                                          1
                                             8
                                                     0
2
   44
      148.0
                NaN
                      56.7 MICU
                                          0
                                            19
                                                     0
3
   68
               180.3
                        84.6 MICU
        NaN
                                          0
                                             9
                                                     0
   88
          NaN
                 NaN
                       NaN MICU
>>> ft = {0: "alive", 1: "dead"}
>>> df["group"] = pd.Categorical(df.death.map(ft))
>>> select = ['Age', 'SysABP', 'Height', 'Weight', 'ICU', 'group']
>>> categ = ['ICU', 'group']
>>> groupby = ['group']
>>> nonnormal = ['Age']
>>> labels={'death': 'mortality'}
>>> tab1 = tableone.TableOne(df,
                           columns=select,
. . .
                           categorical=categ,
. . .
```

```
groupby=groupby,
                              nonnormal=nonnormal,
                              rename=labels,
                              pval=False)
>>> tab1
                           Grouped by group
                                                                             alive
                                     Missing
                                                        Overall
                                                           1000
                                                                               864
Age, median [Q1,Q3]
                                              68.0 [53.0,79.0]
                                                                 66.0 [52.8,78.0]
                                                                                    75.0 [62.0,8
SysABP, mean (SD)
                                         291
                                                   114.3 (40.2)
                                                                      115.4 (38.3)
                                                                                         107.6 (4
Height, mean (SD)
                                         475
                                                   170.1 (22.1)
                                                                      170.3 (23.2)
                                                                                         168.5 (
Weight, mean (SD)
                                         302
                                                    82.9 (23.8)
                                                                                          82.3 (2
                                                                       83.0 (23.6)
ICU, n (%)
                     CCU
                                           0
                                                     162 (16.2)
                                                                        137 (15.9)
                                                                                            25 (
                     CSRU
                                                     202 (20.2)
                                                                        194 (22.5)
                                                                                              8
                     MICU
                                                     380 (38.0)
                                                                        318 (36.8)
                                                                                            62 (4
                     SICU
                                                     256 (25.6)
                                                                        215 (24.9)
                                                                                            41 (3
group, n (%)
                     alive
                                           0
                                                     864 (86.4)
                                                                       864 (100.0)
                     dead
                                                     136 (13.6)
                                                                                          136 (10
```

Correlazione 4.4

Si usa il metodo corr

. . .

```
>>> df = pd.DataFrame(np.random.randn(1000, 5), columns=["a", "b", "c", "d", "e"])
>>> df.corr() # correlazione di pearson
         а
                   b
                             С
 1.000000 -0.011736 -0.020552
                               0.047175 0.035546
b -0.011736 1.000000 0.009603
                               0.002779 -0.011595
c -0.020552  0.009603  1.000000  0.016710 -0.022531
d 0.047175 0.002779 0.016710
                               1.000000
                                         0.020769
  0.035546 -0.011595 -0.022531
                                0.020769
                                          1.000000
>>> df.corr(method='spearman')
                   b
  1.000000 0.004511 -0.027011
                               0.052206 0.027015
  0.004511 1.000000 -0.007078
                                0.014060 -0.001029
c -0.027011 -0.007078 1.000000 0.018606 -0.012641
d 0.052206 0.014060 0.018606 1.000000 0.019836
  0.027015 -0.001029 -0.012641 0.019836 1.000000
```

4.5 Grafici

Scatterplot

Uno con colorazione condizionale e alpha shading in figura 3

```
# data
group1 = pd.DataFrame({'x': np.random.normal(10, 1.2, 2000),
```

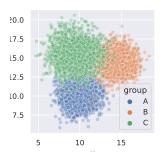


Figure 3: Scatterplot

```
'y': np.random.normal(10, 1.2, 2000),
                        'group': np.repeat('A',2000) })
group2 = pd.DataFrame({'x': np.random.normal(14.5, 1.2, 2000),
                        'y': np.random.normal(14.5, 1.2, 2000),
                        'group': np.repeat('B',2000) })
group3 = pd.DataFrame({'x': np.random.normal(9.5, 1.5, 2000),
                        'y': np.random.normal(15.5, 1.5, 2000),
                        'group': np.repeat('C',2000) })
df = pd.concat([group1, group2, group3])
# Plot
plot = sns.scatterplot(x='x', y='y', data=df, hue='group', alpha = 0.30)
fig = plot.get_figure()
lb.fig.dump(fig, label="sns_scatter", caption = 'Scatterplot', scale = 0.5)
plt.clf()
  Invece per la matrice di scatterplot ne mettiamo senza con regressione (??
e con colorazioni??
# # primo
# plot = sns.pairplot(iris, kind="reg")
# fig = plot.get_figure()
# lb.fig.dump(fig, label="sns_pair1", caption = 'Pairplot 1', scale = 0.5)
# plt.clf()
# # secondo
# plot = sns.pairplot(iris, kind="scatter", hue="species", markers=["o", "s", "D"], palette
# fiq = plot.get_figure()
# lb.fig.dump(fig, label="sns_pair2", caption = 'Pairplot 2', scale = 0.5)
# plt.clf()
```

4.5.2 Boxplot

In figura 4.

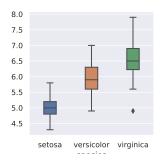


Figure 4: Boxplot

```
plot = sns.boxplot(x=iris.species, y=iris.sepal_length, width=0.3)
fig = plot.get_figure()
lb.fig.dump(fig, label="sns_boxplot", caption = 'Boxplot', scale = 0.5)
plt.clf()
```