```
In [26]:
           import os, pandas as pd, numpy as np, matplotlib.pyplot as pl, pickle as pkl
           from sklearn import model_selection as modele, linear_model as lm, metrics as mes
           import matplotlib as plt
           import seaborn as sb
In [27]:
           os.getcwd()
           'C:\\Users\\HPPC\\regession logistic'
Out[27]:
In [28]:
           os.chdir("C:\\Users\\HPPC\\regession logistic")
In [29]:
           data = pd.read_excel("null.xlsx")
           data.head()
Out[29]:
             AGE
                     SEXE
                            TDT
                                 PAR CHOLESTEROL GAJ
                                                              ECG FCMAX ANGINE DEPRESSION
                                                                                                     PEN1
                                                 289
          0
               40
                   homme
                             AA
                                  140
                                                        0
                                                           Normal
                                                                       172
                                                                               Non
                                                                                              0.0
                                                                                                  Ascenda
          1
               49
                                  160
                                                 180
                                                                       156
                                                                                              1.0
                                                                                                        Ы
                   femme
                           DNA
                                                        0
                                                           Normal
                                                                               Non
          2
               37
                   homme
                             AA
                                  130
                                                283
                                                        0
                                                               ST
                                                                        98
                                                                                Non
                                                                                              0.0
                                                                                                  Ascenda
          3
                                                                                              1.5
                                                                                                        Ы
               48
                            ASY
                                  138
                                                214
                                                           Normal
                                                                       108
                                                                                Oui
                   femme
                                                        0
                           DNA
                                                 195
                                                                       122
                   homme
                                  150
                                                           Normal
                                                                                Non
                                                                                              0.0
                                                                                                  Ascenda
In [30]:
           df = data.copy()
           print(df)
                AGE
                                                                ECG FCMAX ANGINE
                       SEXE
                             TDT
                                   PAR
                                        CHOLESTEROL
                                                       GAJ
          0
                 40
                     homme
                                   140
                                                            Normal
                                                                        172
                              AA
                                                 289
                                                         0
                                                                                Non
                     femme
          1
                 49
                                   160
                                                 180
                                                             Normal
                                                                        156
                             DNA
                                                         0
                                                                                Non
          2
                 37
                     homme
                              AA
                                   130
                                                 283
                                                         0
                                                                 ST
                                                                         98
                                                                                Non
          3
                 48
                             ASY
                                   138
                                                 214
                                                                        108
                                                                                Oui
                     femme
                                                         0
                                                             Normal
          4
                                                 195
                 54
                     homme
                             DNA
                                   150
                                                         0
                                                             Normal
                                                                        122
                                                                                Non
                        . . .
                                                  . . .
                                                                        . . .
                                                                                . . .
                . . .
                                   . . .
                                                                . . .
          913
                 45
                     homme
                              ΑT
                                   110
                                                 264
                                                         0
                                                            Normal
                                                                        132
                                                                                Non
                             ASY
                                                 193
                                                         1
          914
                 68
                     homme
                                   144
                                                             Normal
                                                                        141
                                                                                Non
          915
                 57
                             ASY
                                   130
                                                 131
                                                             Normal
                                                                        115
                                                                                Oui
                     homme
          916
                 57
                     femme
                              AA
                                   130
                                                 236
                                                         0
                                                                LVH
                                                                        174
                                                                                Non
          917
                 38
                     homme
                             DNA
                                   138
                                                 175
                                                         a
                                                            Normal
                                                                        173
                                                                                Non
                DEPRESSION
                                           CŒUR
                                   PENTE
          0
                         0.0
                              Ascendant
                                              0
          1
                         1.0
                                    Plat
                                              1
          2
                         0.0
                              Ascendant
                                              0
          3
                         1.5
                                    Plat
                                              1
          4
                         0.0
                              Ascendant
                                              0
                         . . .
                                     . . .
          913
                         1.2
                                    Plat
                                              1
          914
                         3.4
                                    Plat
                                              1
                         1.2
          915
                                    Plat
                                              1
                                    Plat
                                              1
          916
                         0.0
          917
                         0.0 Ascendant
                                              0
```

[918 rows x 12 columns]

```
In [31]:
           df.duplicated().sum()
Out[31]:
In [32]:
           df.nunique()
          AGE
                            50
Out[32]:
          SEXE
                             2
          TDT
                             4
          PAR
                            67
          CHOLESTEROL
                           222
          GAJ
                             2
          ECG
                             3
          FCMAX
                           119
                             2
          ANGINE
          DEPRESSION
                            53
          PENTE
                             3
          CŒUR
                             2
          dtype: int64
In [33]:
           df.isna().sum()
          AGE
                           0
Out[33]:
          SEXE
                           0
          TDT
                           0
          PAR
                           0
          CHOLESTEROL
                           0
          GAJ
                           0
          ECG
                           0
          \mathsf{FCMAX}
                           0
          ANGINE
                           0
          DEPRESSION
                           0
                           0
          PENTE
          CŒUR
                           0
          dtype: int64
In [38]:
           quant = ['AGE', 'PAR', 'CHOLESTEROL', 'FCMAX', 'DEPRESSION ']
           #df.rename('DEPRESSION')
In [52]:
           def recoder(df):
                for i in df.select_dtypes('object').columns:
                    df[i]=df[i].astype('category').cat.codes
                return(df)
           recoder(df)
                                                          ECG FCMAX ANGINE DEPRESSION PENTE
                                                                                                      CO
Out[52]:
                AGE SEXE TDT PAR CHOLESTEROL GAJ
            0 0.75
                        1
                              0 1.06
                                               1.45
                                                       0
                                                             1
                                                                   1.26
                                                                              0
                                                                                         0.00
                                                                                                   0
               0.92
                                               0.91
                                                                                                   2
                        0
                              3 1.21
                                                       0
                                                             1
                                                                   1.14
                                                                              0
                                                                                         1.13
             2
               0.69
                        1
                              0 0.98
                                               1.42
                                                       0
                                                             2
                                                                   0.72
                                                                              0
                                                                                         0.00
                                                                                                   0
                0.90
                        0
                                               1.08
                                                       0
                                                             1
                                                                              1
                                                                                                   2
            3
                                 1.04
                                                                   0.79
                                                                                         1.69
                1.01
                        1
                              3
                                1.13
                                               0.98
                                                       0
                                                             1
                                                                   0.89
                                                                              0
                                                                                         0.00
                                                                                                   0
```

0.96

0

1.35

1

2 0.83

1.33

0

1

**913** 0.84

2

		AGE	SEXE	TDT	PAR	CHOLESTEROL	GAJ	ECG	FCMAX	ANGINE	DEPRESSION	PENTE	Ca
9	14	1.27	1	1	1.09	0.97	1	1	1.03	0	3.83	2	
9	15	1.07	1	1	0.98	0.66	0	1	0.84	1	1.35	2	
9	16	1.07	0	0	0.98	1.19	0	0	1.27	0	0.00	2	
9	17	0.71	1	3	1.04	0.88	0	1	1.26	0	0.00	0	

918 rows × 12 columns

```
In [53]:
            df[quant] = round(df[quant]/df[quant].mean(), 2)
In [54]:
            df.head()
              AGE SEXE TDT PAR CHOLESTEROL GAJ
                                                           ECG FCMAX ANGINE DEPRESSION PENTE CŒU
Out[54]:
              0.75
                             0
                                1.06
                                               1.45
                                                        0
                                                                    1.26
                                                                                0
                                                                                           0.00
                                                                                                      0
                       1
                                                             1
              0.92
                                                                                                      2
                       0
                             3
                                1.21
                                               0.91
                                                        0
                                                             1
                                                                    1.14
                                                                                0
                                                                                           1.13
              0.69
                             0
                                0.98
                                               1.42
                                                             2
                                                                    0.72
                                                                                0
                                                                                           0.00
                                                                                                      0
           2
                       1
                                                        0
              0.90
                                1.04
                                               1.08
                                                        0
                                                             1
                                                                    0.79
                                                                                           1.69
                                                                                                      2
              1.01
                             3 1.13
                                               0.98
                                                        0
                                                                    0.89
                                                                                0
                                                                                           0.00
                                                                                                      0
                       1
                                                             1
In [55]:
            pl.figure(tight_layout = True, figsize = (14, 6))
            pl.suptitle("LES HISTOGRAMMES DES VARIABLES QUANTITATIVES")
            for y,x in enumerate(quant):
                pl.subplot(2,5,y+1)
                pl.hist(df[x])
                pl.title(f"{quant[y]}")
            pl.show()
                                            LES HISTOGRAMMES DES VARIABLES QUANTITATIVES
                                        PAR
                                                         CHOLESTEROL
                                                                               FCMAX
                                                                                                 DEPRESSION
                    AGE
                               400
                                                   300
                                                                                          350
                                                   250
                                                                       150
                                                                                          300
           150
                                                                                          250
                                                   200
                                                                       100
                                                                                          200
           100
                                                  150
                               200
                                                                                          150
                                                  100
                                                                       50
                                                                                          100
                               100
                                                   50
                0.75 1.00
                                                                             0.75 1.00 1.25 1.50
In [56]:
            qual = ['SEXE', 'TDT', 'GAJ', 'ECG', 'ANGINE', 'PENTE', 'CŒUR'] # La liste des varia
            pl.figure(tight_layout = True, figsize = (15,8))
            pl.suptitle("LES DIAGRAMMES CIRCULAIRE DES VARIABLES QUALITATIVES")
            for y,x in enumerate(qual):
                eff = df[x].value counts()
```

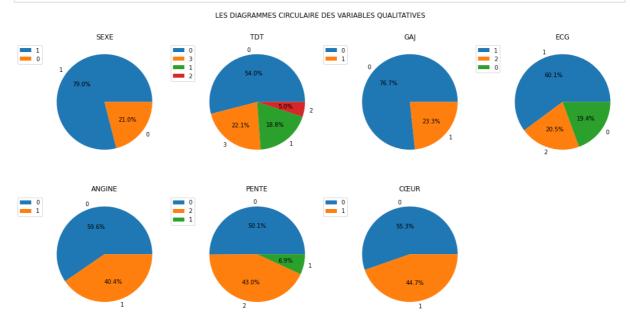
pl.subplot(2,4,y+1)

modalite = df[x].unique()

pl.legend(bbox\_to\_anchor = (0, 1))

pl.pie(eff, labels = modalite, autopct = '%1.1f%%')

```
pl.title(f"{qual[y]}")
pl.show()
```



```
In [57]: X = df.iloc[:, 0:11] X
```

Out[57]:		AGE	SEXE	TDT	PAR	CHOLESTEROL	GAJ	ECG	FCMAX	ANGINE	DEPRESSION	PENTE
	0	0.75	1	0	1.06	1.45	0	1	1.26	0	0.00	0
	1	0.92	0	3	1.21	0.91	0	1	1.14	0	1.13	2
	2	0.69	1	0	0.98	1.42	0	2	0.72	0	0.00	0
	3	0.90	0	1	1.04	1.08	0	1	0.79	1	1.69	2
	4	1.01	1	3	1.13	0.98	0	1	0.89	0	0.00	0
	•••											
	913	0.84	1	2	0.83	1.33	0	1	0.96	0	1.35	2
	914	1.27	1	1	1.09	0.97	1	1	1.03	0	3.83	2
	915	1.07	1	1	0.98	0.66	0	1	0.84	1	1.35	2
	916	1.07	0	0	0.98	1.19	0	0	1.27	0	0.00	2
	917	0.71	1	3	1.04	0.88	0	1	1.26	0	0.00	0

918 rows × 11 columns

```
In [58]: y = data.iloc[:,11:]
y
```

Out[58]:	CŒUR				
	0	0			
	1	1			
	2	0			
	3	1			

**CŒUR** 

0

4

```
913
         914
         915
         916
                   1
         917
         918 rows × 1 columns
In [59]:
          X_train, X_test, Y_train, Y_test = modele.train_test_split(X, y, test_size = 0.2, ra
In [60]:
          mod = lm.LogisticRegression(penalty = 'none', solver = 'newton-cg')
          mod.fit(X_train, Y_train)
         C:\ProgramData\Anaconda3\lib\site-packages\sklearn\utils\validation.py:63: DataConve
         rsionWarning: A column-vector y was passed when a 1d array was expected. Please chan
         ge the shape of y to (n_samples, ), for example using ravel().
           return f(*args, **kwargs)
         LogisticRegression(penalty='none', solver='newton-cg')
Out[60]:
In [61]:
          mod.predict_proba(X_test)
         array([[0.35496203, 0.64503797],
Out[61]:
                 [0.16960133, 0.83039867],
                 [0.03457301, 0.96542699],
                 [0.0764487 , 0.9235513 ],
                 [0.17768705, 0.82231295],
                 [0.74672317, 0.25327683],
                 [0.29597548, 0.70402452],
                 [0.51735064, 0.48264936],
                 [0.96541828, 0.03458172],
                 [0.2826058 , 0.7173942 ],
                 [0.88994162, 0.11005838],
                 [0.6539992 , 0.3460008 ],
                 [0.40417279, 0.59582721],
                 [0.16705882, 0.83294118],
                 [0.09203056, 0.90796944],
                 [0.88406903, 0.11593097],
                 [0.07255202, 0.92744798],
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                 [0.21131067, 0.78868933],
                 [0.07855208, 0.92144792],
                 [0.96654432, 0.03345568],
                 [0.03879931, 0.96120069],
                 [0.96237084, 0.03762916],
                 [0.41115221, 0.58884779],
                 [0.02625307, 0.97374693],
                 [0.86081418, 0.13918582],
                 [0.184074 , 0.815926 ],
```

[0.14306526, 0.85693474],

```
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         , 0.18157
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[0.07269278, 0.92730722],
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```

```
[0.44620256, 0.55379744],
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```

```
[0.04003012, 0.95996988],
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                 [0.81765148, 0.18234852],
                 [0.03583028, 0.96416972],
                 [0.54436461, 0.45563539],
                 [0.48625252, 0.51374748],
                 [0.97215613, 0.02784387],
                 [0.87130014, 0.12869986],
                 [0.58628658, 0.41371342]])
In [62]:
          pred = mod.predict(X_test)
          pred
         array([1, 1, 1, 1, 1, 0, 1, 0, 0, 1, 0, 0, 1, 1, 1, 0, 1, 0, 1, 1, 1, 0,
Out[62]:
                 1, 0, 1, 1, 0, 1, 1, 0, 0, 0, 0, 0, 1, 1, 1, 1, 0, 0, 1, 1, 1, 0,
                 0, 0, 1, 0, 0, 1, 0, 1, 1, 0, 1, 0, 1, 0, 1, 0, 1, 1, 1, 0, 0, 1,
                 1, 1, 0, 1, 1, 0, 1, 1, 0, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0,
                 0, 0, 1, 1, 0, 1, 1, 1, 1, 1, 0, 1, 0, 1, 1, 1, 0, 0, 0, 1, 0,
                 0, 1, 0, 0, 0, 0, 1, 0, 0, 1, 1, 1, 1, 0, 1, 0, 1, 0, 1,
                                                                          1,
                0, 1, 1, 1, 1, 0, 1, 1, 0, 0, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0,
                0, 1, 0, 1, 1, 1, 1, 1, 0, 1, 0, 0, 1, 1, 1, 0, 1, 1, 1, 1,
                 0, 0, 1, 0, 1, 0, 0, 0], dtype=int64)
In [63]:
          mes.confusion matrix(Y test, pred)
          array([[ 66,
                         8],
Out[63]:
                  8, 102]], dtype=int64)
In [64]:
          mes.accuracy score(Y test, pred)
          0.9130434782608695
Out[64]:
In [65]:
          mes.recall score(Y test, pred)
          0.9272727272727272
Out[65]:
In [66]:
          mes.precision_score(Y_test, pred)
          0.9272727272727272
Out[66]:
```

```
In [67]: # sauvegarde du modele de prediction mod
    pkl.dump(mod, open('mod.pkl', 'wb' ))
In []:
```