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## ▼ Perceptron

```
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt

df = pd.read_csv('Diabetes.csv')

print(df.shape)
df.head()
```

(768, 9)

•	- /								
	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	${\tt DiabetesPedigreeFunction}$	Age	Outcome
0	6	148	72	35	0	33.6	0.627	50	1
1	1	85	66	29	0	26.6	0.351	31	0
2	8	183	64	0	0	23.3	0.672	32	1
3	1	89	66	23	94	28.1	0.167	21	0
4	0	137	40	35	168	43.1	2.288	33	1

```
warnings.warn(
    <AxesSubplot:xlabel='Pregnancies', ylabel='Glucose'>
      200
      175
                                     25
      150
                                    • 75
• 100
Insulin
                                       75
100
      125
      100
                                    •
                                      150
       75
       50
                                       450
x = df.iloc[:,0:2]
y = df.iloc[:,-1]
                                   * 7
print (x)
        Pregnancies Glucose
    0
                6
                      148
    1
                1
                       85
    2
                8
                      183
    3
                1
                       89
                0
                      137
    763
               10
                      101
    764
                2
                      122
    765
                5
                      121
    766
                1
                      126
    767
                1
                       93
    [768 rows x 2 columns]
                                   • 29
print(y)
    1
         0
    2
         1
    3
         0
    4
         1
    763
         0
    764
         0
    765
         0
    766
    767
    Name: Outcome, Length: 768, dtype: int64
from sklearn.linear_model import Perceptron
p = Perceptron()
                                   • 52
p.fit(x,y)
    ▼ Perceptron
    Perceptron()
p.coef_
```

array([[212., 91.]])

from mlxtend.plotting import plot\_decision\_regions

plot\_decision\_regions(x.values, y.values, clf=p, legend=2)

array([-281.])

p.intercept\_

C:\Users\acer\anaconda3\lib\site-packages\sklearn\base.py:465: UserWarning: X does not have valid feature names, but Perceptron was warnings.warn(

<AxesSubplot:>



