

2303a52444-sml-project

October 24, 2024

```
[ ]: import pandas as pd
import numpy as np
d=pd.read_csv("/content/diabetes_data_upload.csv")
print(d.head())
```

	Age	Gender	Polyuria	Polydipsia	sudden weight loss	weakness	Polyphagia	\
0	40	Male	No	Yes		No	Yes	No
1	58	Male	No	No		No	Yes	No
2	41	Male	Yes	No		No	Yes	Yes
3	45	Male	No	No		Yes	Yes	Yes
4	60	Male	Yes	Yes		Yes	Yes	Yes

	Genital thrush	visual blurring	Itching	Irritability	delayed healing	\
0	No	No	Yes	No	Yes	
1	No	Yes	No	No	No	
2	No	No	Yes	No	Yes	
3	Yes	No	Yes	No	Yes	
4	No	Yes	Yes	Yes	Yes	

	partial paresis	muscle stiffness	Alopecia	Obesity	class
0	No	Yes	Yes	Yes	Positive
1	Yes	No	Yes	No	Positive
2	No	Yes	Yes	No	Positive
3	No	No	No	No	Positive
4	Yes	Yes	Yes	Yes	Positive

```
[ ]: X=d.drop('class',axis=1)
mle_params={}
y=d["class"]
print("target:")
print(y.head())
print("shape:",y.shape)
print("features:")
print(X.head())
```

target:

0	Positive
1	Positive

```
2    Positive
3    Positive
4    Positive
```

Name: class, dtype: object

shape: (520,)

features:

	Age	Gender	Polyuria	Polydipsia	sudden weight loss	weakness	Polyphagia \
0	40	Male	No	Yes	No	Yes	No
1	58	Male	No	No	No	Yes	No
2	41	Male	Yes	No	No	Yes	Yes
3	45	Male	No	No	Yes	Yes	Yes
4	60	Male	Yes	Yes	Yes	Yes	Yes

	Genital thrush	visual blurring	Itching	Irritability	delayed healing \
0	No	No	Yes	No	Yes
1	No	Yes	No	No	No
2	No	No	Yes	No	Yes
3	Yes	No	Yes	No	Yes
4	No	Yes	Yes	Yes	Yes

	partial paresis	muscle stiffness	Alopecia	Obesity
0	No	Yes	Yes	Yes
1	Yes	No	Yes	No
2	No	Yes	Yes	No
3	No	No	No	No
4	Yes	Yes	Yes	Yes

```
[ ]: S=X.replace({'Male':1,'Female':0,'Yes':1,'No':0})
print(S)
Y=y.replace({'Positive':1,'Negative':0})
print(Y)
```

	Age	Gender	Polyuria	Polydipsia	sudden weight loss	weakness \
0	40	1	0	1	0	1
1	58	1	0	0	0	1
2	41	1	1	0	0	1
3	45	1	0	0	1	1
4	60	1	1	1	1	1

..
515	39	0	1	1	1	0
516	48	0	1	1	1	1
517	58	0	1	1	1	1
518	32	0	0	0	0	1
519	42	1	0	0	0	0

	Polyphagia	Genital thrush	visual blurring	Itching	Irritability \
0	0	0	0	1	0
1	0	0	1	0	0

2	1	0	0	1	0
3	1	1	0	1	0
4	1	0	1	1	1
..
515	1	0	0	1	0
516	1	0	0	1	1
517	1	0	1	0	0
518	0	0	1	1	0
519	0	0	0	0	0

	delayed healing	partial paresis	muscle stiffness	Alopecia	Obesity
0	1	0	1	1	1
1	0	1	0	1	0
2	1	0	1	1	0
3	1	0	0	0	0
4	1	1	1	1	1
..
515	1	1	0	0	0
516	1	1	0	0	0
517	0	1	1	0	1
518	1	0	0	1	0
519	0	0	0	0	0

[520 rows x 16 columns]

0	1
1	1
2	1
3	1
4	1
..	
515	1
516	1
517	1
518	0
519	0

Name: class, Length: 520, dtype: int64

<ipython-input-4-e79e090bab8f>:1: FutureWarning: Downcasting behavior in `replace` is deprecated and will be removed in a future version. To retain the old behavior, explicitly call `result.infer_objects(copy=False)`. To opt-in to the future behavior, set `pd.set_option('future.no_silent_downcasting', True)`

S=X.replace({'Male':1,'Female':0,'Yes':1,'No':0})

<ipython-input-4-e79e090bab8f>:3: FutureWarning: Downcasting behavior in `replace` is deprecated and will be removed in a future version. To retain the old behavior, explicitly call `result.infer_objects(copy=False)`. To opt-in to the future behavior, set `pd.set_option('future.no_silent_downcasting', True)`

Y=y.replace({'Positive':1,'Negative':0})

```
[ ]: from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(S, Y , test_size=0.2,
↳random_state=30)
print("X_train:",X_train)
print("X_test:",X_test)
print("Y_train:",y_train)
print("Y_test:",y_test)
```

X_train:	Age	Gender	Polyuria	Polydipsia	sudden weight loss	weakness \
195	31	1	1	0	0	0
349	37	1	0	0	0	0
257	48	0	1	1	0	1
157	48	1	1	1	1	0
459	57	1	1	1	1	1
..
430	32	1	0	0	0	0
145	61	1	1	0	0	1
140	47	1	1	1	0	0
500	66	1	1	0	1	0
421	61	0	1	0	0	0

	Polyphagia	Genital thrush	visual blurring	Itching	Irritability \
195	1	0	1	0	0
349	0	0	0	0	0
257	0	0	1	1	0
157	1	1	0	0	0
459	1	0	1	0	0
..
430	0	1	0	0	1
145	1	1	1	1	1
140	0	0	0	0	0
500	0	1	0	1	1
421	1	0	0	0	1

	delayed healing	partial paresis	muscle stiffness	Alopecia	Obesity
195	0	1	0	1	0
349	0	0	0	0	0
257	1	1	0	0	0
157	1	0	0	0	0
459	0	1	0	0	0
..
430	1	0	0	0	1
145	1	1	0	0	0
140	0	0	1	0	0
500	0	0	0	1	0
421	0	0	0	1	0

[416 rows x 16 columns]

X_test:	Age	Gender	Polyuria	Polydipsia	sudden weight loss	weakness	\
248	42	1	0	0	0	0	
60	65	0	1	1	0	1	
162	35	0	0	0	0	0	
391	58	1	0	1	0	0	
493	44	1	1	0	1	1	
..	
188	70	1	1	0	1	1	
158	56	1	1	0	1	1	
363	68	0	1	1	0	1	
164	48	0	1	1	1	1	
42	50	0	0	1	0	1	

	Polyphagia	Genital thrush	visual blurring	Itching	Irritability	\
248	0	0	0	0	0	
60	1	0	0	1	0	
162	0	0	0	0	0	
391	0	0	1	1	0	
493	0	1	0	1	0	
..	
188	1	1	0	0	1	
158	0	1	0	1	1	
363	1	0	1	1	0	
164	1	0	1	1	1	
42	1	0	1	1	1	

	delayed healing	partial paresis	muscle stiffness	Alopecia	Obesity
248	0	0	0	1	0
60	0	1	1	0	0
162	0	0	0	0	0
391	0	0	1	0	0
493	1	0	0	1	1
..
188	1	1	1	1	0
158	0	0	0	1	0
363	1	1	0	0	0
164	1	1	1	0	0
42	1	1	1	0	0

[104 rows x 16 columns]

Y_train:	195	1
349	0	
257	1	
157	1	
459	1	
..		
430	1	

```

145    1
140    1
500    1
421    1
Name: class, Length: 416, dtype: int64
Y_train: 248    0
60     1
162    1
391    0
493    0
..
188    1
158    1
363    1
164    1
42     1
Name: class, Length: 104, dtype: int64

```

```

[ ]: from sklearn.linear_model import LogisticRegression
LR = LogisticRegression()
lr=LR.fit(X_train,y_train)
print("coefficient:",lr.coef_)
print("intercept:",lr.intercept_)

```

```

coefficient: [[-0.02698604 -2.60136808  2.48594712  2.86268139  0.60790669
 0.34334826
 1.04831723  1.13493086  0.6559378  -1.38534348  1.47220207 -0.43748972
 1.04402217 -0.42196691 -0.00355869 -0.21565816]]
intercept: [1.15463711]

```

```

/usr/local/lib/python3.10/dist-packages/sklearn/linear_model/_logistic.py:469:
ConvergenceWarning: lbfgs failed to converge (status=1):
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

```

Increase the number of iterations (max_iter) or scale the data as shown in:
<https://scikit-learn.org/stable/modules/preprocessing.html>
Please also refer to the documentation for alternative solver options:
https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression
n_iter_i = _check_optimize_result(

```

[ ]: from sklearn.metrics import
      accuracy_score, confusion_matrix, classification_report
y_pred = lr.predict(X_train)
print("accuracy:",accuracy_score(y_train, y_pred))
print("confusion matrix:")
print(confusion_matrix(y_train,y_pred))
print("classification report:")

```

```
print(classification_report(y_train,y_pred))
```

accuracy: 0.9471153846153846

confusion matrix:

```
[[152   7]
 [ 15 242]]
```

classification report:

	precision	recall	f1-score	support
0	0.91	0.96	0.93	159
1	0.97	0.94	0.96	257
accuracy			0.95	416
macro avg	0.94	0.95	0.94	416
weighted avg	0.95	0.95	0.95	416

```
[ ]: from sklearn.svm import SVC
SV= SVC()
sv=SV.fit(X_train,y_train)
print("intercept:",sv.intercept_)
```

intercept: [0.73657188]

```
[ ]: from sklearn.metrics import accuracy_score,confusion_matrix,classification_report
y_pred = sv.predict(X_train)
print("accuracy:",accuracy_score(y_train, y_pred))
print("confusion matrix:")
print(confusion_matrix(y_train,y_pred))
print("classification report:")
print(classification_report(y_train,y_pred))
```

accuracy: 0.6177884615384616

confusion matrix:

```
[[ 0 159]
 [ 0 257]]
```

classification report:

	precision	recall	f1-score	support
0	0.00	0.00	0.00	159
1	0.62	1.00	0.76	257
accuracy			0.62	416
macro avg	0.31	0.50	0.38	416
weighted avg	0.38	0.62	0.47	416

```
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_classification.py:1531:
UndefinedMetricWarning: Precision is ill-defined and being set to 0.0 in labels
with no predicted samples. Use `zero_division` parameter to control this
behavior.
```

```
_warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_classification.py:1531:
UndefinedMetricWarning: Precision is ill-defined and being set to 0.0 in labels
with no predicted samples. Use `zero_division` parameter to control this
behavior.
```

```
_warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_classification.py:1531:
UndefinedMetricWarning: Precision is ill-defined and being set to 0.0 in labels
with no predicted samples. Use `zero_division` parameter to control this
behavior.
```

```
_warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
```

```
[ ]: from sklearn.linear_model import Perceptron
PR= Perceptron()
pr=PR.fit(X_train,y_train)
print("intercept:",pr.intercept_)
print("coefficient:",pr.coef_)
```

```
intercept: [-87.]
coefficient: [[ 17. -539.  718.  713.  359.   90.  287.  198.  202. -183.  310.
-81.
 363. -28. -259.  35.]]
```

```
[ ]: from sklearn.metrics import accuracy_score
y_pred = pr.predict(X_train)
print("accuracy:",accuracy_score(y_train, y_pred))
print("confusion matrix:")
print(confusion_matrix(y_train,y_pred))
print("classification report:")
print(classification_report(y_train,y_pred))
```

```
accuracy: 0.7451923076923077
```

```
confusion matrix:
```

```
[[ 56 103]
```

```
 [  3 254]]
```

```
classification report:
```

	precision	recall	f1-score	support
0	0.95	0.35	0.51	159
1	0.71	0.99	0.83	257
accuracy			0.75	416
macro avg	0.83	0.67	0.67	416
weighted avg	0.80	0.75	0.71	416

