DecisionTree (1)

August 19, 2022

```
[1]: import pandas as pd
     import numpy as np
     data = pd.read_csv('horse.csv')
     data.head()
[1]:
       surgery
                        hospital_number
                                           rectal_temp pulse
                                                                respiratory_rate
                   age
     0
            no
                 adult
                                  530101
                                                   38.5
                                                          66.0
                                                                              28.0
     1
                                                   39.2
                                                          88.0
                                                                              20.0
                 adult
                                  534817
           yes
     2
                 adult
                                  530334
                                                   38.3
                                                          40.0
                                                                              24.0
            no
     3
                 young
                                 5290409
                                                   39.1 164.0
                                                                              84.0
           yes
                                                   37.3 104.0
     4
                 adult
                                  530255
                                                                              35.0
            no
       temp_of_extremities peripheral_pulse mucous_membrane capillary_refill_time
     0
                                      reduced
                                                                            more_3_sec
                       cool
                                                            NaN
     1
                        NaN
                                           NaN
                                                 pale cyanotic
                                                                            less 3 sec
     2
                                                      pale_pink
                     normal
                                       normal
                                                                             less_3_sec
     3
                       cold
                                       normal
                                                 dark_cyanotic
                                                                            more_3_sec
     4
                                                 dark_cyanotic
                        NaN
                                           NaN
                                                                            more_3_sec
        ... packed_cell_volume total_protein abdomo_appearance abdomo_protein
     0
                          45.0
                                          8.4
                                                             NaN
                                                                              NaN
     1
                          50.0
                                         85.0
                                                          cloudy
                                                                              2.0
     2
                          33.0
                                          6.7
                                                             NaN
                                                                              NaN
     3
                                          7.2
                          48.0
                                                   serosanguious
                                                                              5.3
                                          7.4
     4
                          74.0
                                                             NaN
                                                                              NaN
                     surgical_lesion lesion_1 lesion_2
                                                           lesion_3
           outcome
                                                                      cp_data
     0
               died
                                          11300
                                                        0
                                                                   0
                                   no
                                                                           no
        euthanized
                                           2208
                                                        0
                                                                   0
     1
                                   no
                                                                           no
     2
             lived
                                                        0
                                                                   0
                                              0
                                   no
                                                                          yes
     3
               died
                                           2208
                                                        0
                                                                   0
                                  yes
                                                                          yes
               died
                                           4300
     [5 rows x 28 columns]
[2]: data.info()
```

<class 'pandas.core.frame.DataFrame'> RangeIndex: 299 entries, 0 to 298 Data columns (total 28 columns):

#	Column	Non-Null Count	Dtype
0	surgery	299 non-null	object
1	age	299 non-null	object
2	hospital_number	299 non-null	int64
3	rectal_temp	239 non-null	float64
4	pulse	275 non-null	float64
5	respiratory_rate	241 non-null	float64
6	temp_of_extremities	243 non-null	object
7	peripheral_pulse	230 non-null	object
8	mucous_membrane	252 non-null	object
9	capillary_refill_time	267 non-null	object
10	pain	244 non-null	object
11	peristalsis	255 non-null	object
12	${\tt abdominal_distention}$	243 non-null	object
13	nasogastric_tube	195 non-null	object
14	nasogastric_reflux	193 non-null	object
15	nasogastric_reflux_ph	53 non-null	float64
16	rectal_exam_feces	197 non-null	object
17	abdomen	181 non-null	object
18	<pre>packed_cell_volume</pre>	270 non-null	float64
19	total_protein	266 non-null	float64
20	abdomo_appearance	134 non-null	object
21	abdomo_protein	101 non-null	float64
22	outcome	299 non-null	object
23	surgical_lesion	299 non-null	object
24	lesion_1	299 non-null	int64
25	lesion_2	299 non-null	int64
26	lesion_3	299 non-null	int64
27	cp_data	299 non-null	object
dtyp	es: float64(7), int64(4), object(17)	
memo	rv usage: 65.5+ KB		

memory usage: 65.5+ KB

[3]: data.isna().sum()

```
[3]: surgery
                                0
     age
                                0
    hospital_number
                                0
     rectal_temp
                                60
     pulse
                                24
     respiratory_rate
                                58
     temp_of_extremities
                                56
    peripheral_pulse
                                69
     mucous_membrane
                                47
```

```
capillary_refill_time
                                55
     pain
                                44
     peristalsis
     abdominal_distention
                                56
     nasogastric_tube
                               104
     nasogastric_reflux
                               106
    nasogastric_reflux_ph
                               246
     rectal_exam_feces
                               102
     abdomen
                               118
     packed_cell_volume
                                29
     total_protein
                                33
     abdomo_appearance
                               165
     abdomo_protein
                               198
     outcome
                                 0
     surgical_lesion
                                 0
                                 0
     lesion_1
     lesion_2
                                 0
     lesion_3
                                 0
                                 0
     cp_data
     dtype: int64
[4]: #Target Class
     data.outcome.value_counts()
[4]: lived
                   178
                    77
     died
     euthanized
                    44
     Name: outcome, dtype: int64
[5]: features = data.drop(['outcome'], axis = 1)
     target = data[['outcome']]
[6]: features.shape,target.shape
[6]: ((299, 27), (299, 1))
[7]: features.dtypes
[7]: surgery
                                object
                                object
     age
     hospital_number
                                 int64
                               float64
     rectal_temp
     pulse
                               float64
     respiratory_rate
                               float64
     temp_of_extremities
                                object
     peripheral_pulse
                                object
    mucous_membrane
                                object
```

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```
capillary_refill_time
                                object
      pain
                                object
      peristalsis
                                object
      abdominal_distention
                                object
     nasogastric_tube
                                object
     nasogastric_reflux
                                object
     nasogastric_reflux_ph
                               float64
      rectal_exam_feces
                                object
      abdomen
                                object
      packed_cell_volume
                               float64
      total_protein
                               float64
      abdomo_appearance
                                object
      abdomo_protein
                               float64
      surgical_lesion
                                object
      lesion 1
                                 int64
      lesion_2
                                 int64
      lesion_3
                                 int64
      cp_data
                                object
      dtype: object
 [8]: features_transformed = pd.get_dummies(features)
 [9]: from sklearn.model_selection import train_test_split
      from sklearn.tree import DecisionTreeClassifier
      from sklearn.ensemble import RandomForestClassifier
[10]: X_train , X_test, y_train, y_test = train_test_split(features_transformed,__
       →target, random_state = 10)
[11]: print(X train.shape)
      print(X_test.shape)
      print(y_train.shape)
      print(y_test.shape)
     (224, 67)
     (75, 67)
     (224, 1)
     (75, 1)
[12]: from sklearn.impute import SimpleImputer
[13]: imputer = SimpleImputer(missing_values=np.nan, strategy='most_frequent')
[14]: X_train = imputer.fit_transform(X_train)
      X_test = imputer.fit_transform(X_test)
```

```
[15]: my_DT_model = DecisionTreeClassifier(criterion='entropy', random_state=2,_u
       \rightarrowmax_depth = 1)
[16]: my_DT_model.fit(X_train,y_train)
[16]: DecisionTreeClassifier(criterion='entropy', max_depth=1, random_state=2)
     0.0.1 Using GridSearchCV to find best params
[17]: from sklearn.model selection import GridSearchCV
[18]: params = {'criterion':['gini', 'entropy'], 'max_depth':[1,2,3,10], 'splitter':
       →['best', 'random']}
[19]: grid_search = GridSearchCV(my_DT_model, params, cv = 3, n_jobs = -1)
[20]: grid_search.fit(X_train, y_train)
[20]: GridSearchCV(cv=3,
                   estimator=DecisionTreeClassifier(criterion='entropy', max_depth=1,
                                                    random_state=2),
                   n_jobs=-1,
                   param_grid={'criterion': ['gini', 'entropy'],
                               'max_depth': [1, 2, 3, 10],
                               'splitter': ['best', 'random']})
[21]: grid_search.best_params_
[21]: {'criterion': 'gini', 'max_depth': 3, 'splitter': 'best'}
[22]: my_DT_model = DecisionTreeClassifier(criterion='gini', random_state=2,__
       →max_depth = 3, splitter = 'best')
[23]: my_DT_model.fit(X_train, y_train)
[23]: DecisionTreeClassifier(max_depth=3, random_state=2)
[24]: my_preds = my_DT_model.predict(X_test)
[25]: from sklearn.metrics import accuracy_score, confusion_matrix,_
       →classification report
[26]: accuracy_score(y_test, my_preds)
[26]: 0.666666666666666
```

```
[27]: print(confusion_matrix(y_test, my_preds, ))

[[ 3     0     12]
       [ 1     1     9]
       [ 3     0     46]]

[28]: print(classification_report(y_test, my_preds))
```

support	f1-score	recall	precision	
15	0.27	0.20	0.43	died
11	0.17	0.09	1.00	euthanized
49	0.79	0.94	0.69	lived
75	0.67			accuracy
75	0.41	0.41	0.71	macro avg
75	0.60	0.67	0.68	weighted avg

1 Voting Classifiers

```
[29]: from sklearn.linear_model import LogisticRegression from sklearn.svm import SVC from sklearn.ensemble import VotingClassifier
```

```
[30]: rf_clf = RandomForestClassifier()
log_clf = LogisticRegression()
svm_clf = SVC()
```

```
[32]: voting_clf.fit(X_train, y_train)
```

/usr/local/lib/python3.7/site-packages/sklearn/utils/validation.py:63: DataConversionWarning: A column-vector y was passed when a 1d array was expected. Please change the shape of y to (n_samples,), for example using ravel().

```
return f(*args, **kwargs)
```

/usr/local/lib/python3.7/site-packages/sklearn/linear_model/_logistic.py:765: 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
       extra_warning_msg=_LOGISTIC_SOLVER_CONVERGENCE_MSG)
[32]: VotingClassifier(estimators=[('lr', LogisticRegression()),
                                  ('rf', RandomForestClassifier()), ('svc', SVC())])
[33]: from sklearn.metrics import accuracy_score
[34]: for clf in (log_clf, rf_clf, svm_clf, voting_clf):
          clf.fit(X_train, y_train)
         y_pred = clf.predict(X_test)
         print(clf.__class__.__name__, accuracy_score(y_test, y_pred))
     /usr/local/lib/python3.7/site-packages/sklearn/utils/validation.py:63:
     DataConversionWarning: A column-vector y was passed when a 1d array was
     expected. Please change the shape of y to (n_samples, ), for example using
     ravel().
       return f(*args, **kwargs)
     /usr/local/lib/python3.7/site-packages/sklearn/linear_model/_logistic.py:765:
     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
       extra_warning_msg=_LOGISTIC_SOLVER_CONVERGENCE_MSG)
     /usr/local/lib/python3.7/site-packages/ipykernel_launcher.py:2:
     DataConversionWarning: A column-vector y was passed when a 1d array was
     expected. Please change the shape of y to (n_samples,), for example using
     ravel().
     LogisticRegression 0.626666666666667
     SVC 0.65333333333333333
     /usr/local/lib/python3.7/site-packages/sklearn/utils/validation.py:63:
     DataConversionWarning: A column-vector y was passed when a 1d array was
     expected. Please change the shape of y to (n_samples, ), for example using
     ravel().
       return f(*args, **kwargs)
     /usr/local/lib/python3.7/site-packages/sklearn/utils/validation.py:63:
     DataConversionWarning: A column-vector y was passed when a 1d array was
     expected. Please change the shape of y to (n_samples, ), for example using
     ravel().
       return f(*args, **kwargs)
```

```
/usr/local/lib/python3.7/site-packages/sklearn/linear_model/_logistic.py:765:
     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
       extra_warning_msg=_LOGISTIC_SOLVER_CONVERGENCE_MSG)
     VotingClassifier 0.64
[35]: from sklearn.ensemble import BaggingClassifier
[36]: bag_clf = BaggingClassifier(DecisionTreeClassifier(), n_estimators=100)
[37]: bag_clf.fit(X_train,y_train)
     /usr/local/lib/python3.7/site-packages/sklearn/utils/validation.py:63:
     DataConversionWarning: A column-vector y was passed when a 1d array was
     expected. Please change the shape of y to (n_samples, ), for example using
     ravel().
       return f(*args, **kwargs)
[37]: BaggingClassifier(base_estimator=DecisionTreeClassifier(), n_estimators=100)
[38]: y_pred = bag_clf.predict(X_test)
      accuracy_score(y_pred, y_test)
[38]: 0.7333333333333333
[39]: my_rf_classifier = RandomForestClassifier()
[40]: my_rf_classifier.fit(X_train, y_train)
     /usr/local/lib/python3.7/site-packages/ipykernel_launcher.py:1:
     DataConversionWarning: A column-vector y was passed when a 1d array was
     expected. Please change the shape of y to (n_samples,), for example using
     ravel().
       """Entry point for launching an IPython kernel.
[40]: RandomForestClassifier()
[41]: my_predictions = my_rf_classifier.predict(X_test)
[42]: print(accuracy_score(y_test, my_predictions))
```

0.746666666666667

```
[43]: print(confusion_matrix(y_test, my_predictions))
     [[ 8 0 7]
      [ 1 2 8]
      [ 3 0 46]]
[44]: print(classification_report(y_test, my_predictions))
                   precision
                                recall f1-score
                                                    support
             died
                        0.67
                                  0.53
                                             0.59
                                                         15
       euthanized
                        1.00
                                  0.18
                                             0.31
                                                         11
                        0.75
                                  0.94
                                             0.84
                                                         49
            lived
                                             0.75
                                                         75
         accuracy
                                             0.58
                                                         75
        macro avg
                        0.81
                                  0.55
     weighted avg
                        0.77
                                  0.75
                                             0.71
                                                         75
[45]: from sklearn.ensemble import VotingClassifier, BaggingClassifier
      from sklearn.linear_model import LogisticRegression
[46]: my_logreg_clf = LogisticRegression()
[47]: my_vt_clf = VotingClassifier(estimators=[('lr', my_logreg_clf), ('rf', __
       →my_rf_classifier)],
                                  voting = 'hard')
[48]: my_vt_clf.fit(X_train, y_train)
     /usr/local/lib/python3.7/site-packages/sklearn/utils/validation.py:63:
     DataConversionWarning: A column-vector y was passed when a 1d array was
     expected. Please change the shape of y to (n_samples, ), for example using
     ravel().
       return f(*args, **kwargs)
     /usr/local/lib/python3.7/site-packages/sklearn/linear_model/_logistic.py:765:
     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
       extra_warning_msg=_LOGISTIC_SOLVER_CONVERGENCE_MSG)
```

```
[48]: VotingClassifier(estimators=[('lr', LogisticRegression()),
                                   ('rf', RandomForestClassifier())])
[49]: my_bagging_clf = BaggingClassifier(DecisionTreeClassifier(), n_estimators=100,
                                            max_samples=100, bootstrap=True)
      my_bagging_clf.fit(X_train, y_train)
     /usr/local/lib/python3.7/site-packages/sklearn/utils/validation.py:63:
     DataConversionWarning: A column-vector y was passed when a 1d array was
     expected. Please change the shape of y to (n_samples, ), for example using
     ravel().
       return f(*args, **kwargs)
[49]: BaggingClassifier(base_estimator=DecisionTreeClassifier(), max_samples=100,
                        n_estimators=100)
[50]: from sklearn.ensemble import AdaBoostClassifier
      ada_clf = AdaBoostClassifier(DecisionTreeClassifier(), n_estimators= 100)
[51]: ada_clf.fit(X_train, y_train)
     /usr/local/lib/python3.7/site-packages/sklearn/utils/validation.py:63:
     DataConversionWarning: A column-vector y was passed when a 1d array was
     expected. Please change the shape of y to (n_samples, ), for example using
     ravel().
       return f(*args, **kwargs)
[51]: AdaBoostClassifier(base_estimator=DecisionTreeClassifier(), n_estimators=100)
[52]: from sklearn.ensemble import GradientBoostingClassifier
[53]: | #learning_rate = 0.3, max_depth=5, n_estimators=1100, n_iter_no_change=10
      gbc clf = GradientBoostingClassifier()
[54]: gbc_clf.fit(X_train, y_train)
     /usr/local/lib/python3.7/site-packages/sklearn/utils/validation.py:63:
     DataConversionWarning: A column-vector y was passed when a 1d array was
     expected. Please change the shape of y to (n samples, ), for example using
     ravel().
       return f(*args, **kwargs)
[54]: GradientBoostingClassifier()
[55]: gbc_clf.n_estimators_
```

```
[55]: 100
[56]: import xgboost, time
[57]: xgb_clf = xgboost.XGBClassifier()
[58]: start = time.time()
      xgb_clf.fit(X_train, y_train)
      end = time.time()
      time elapsed = end - start
      print(time_elapsed)
     /usr/local/lib/python3.7/site-packages/sklearn/utils/validation.py:63:
     DataConversionWarning: A column-vector y was passed when a 1d array was
     expected. Please change the shape of y to (n_samples, ), for example using
     ravel().
       return f(*args, **kwargs)
     88.5109612941742
[59]: y_pred = xgb_clf.predict(X_test)
[60]: accuracy_score(y_pred, y_test)
[60]: 0.746666666666667
 []: params = {'n_estimators':[100, 200, 400, 800], 'max depth':[1,2,3,6,10],__

→'learning_rate': [0.1, 0.2, 0.3, 0.5], 'min_child_weight': [1, 2, 3, 4, 5],

       \rightarrow 'subsample' : [0.5, 0.6, 0.7, 0.8, 1.0]}
      grid_search = GridSearchCV(xgb_clf, params, cv = 3, n_jobs = -1)
      grid_search.fit(X_train, y_train)
 []: grid_search.best_params_
 []: xgb_clf = xgboost.XGBClassifier(learning_rate = 0.1, max_depth = 3,__
       →min_child_weight = 5, n_estimators = 200, subsample = 0.6)
 []: xgb_clf.fit(X_train, y_train)
 []: y_pred = xgb_clf.predict(X_test)
      accuracy_score(y_pred, y_test)
 []:
```