Statistical Learning weekly assignment 5

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```
[32]: import feather
import pandas as pd
import numpy as np
from statsmodels.stats.outliers_influence import variance_inflation_factor
from sklearn.linear_model import Ridge, Lasso, ElasticNet
from sklearn.model_selection import train_test_split, KFold
from sklearn.metrics import accuracy_score, confusion_matrix
```

```
[]: train = feather.read_dataframe("masq_train.feather").replace('NA', np.nan).

dropna()

test = feather.read_dataframe("masq_test.feather").replace('NA', np.nan).dropna()

train['GENDER.f'] = train.apply(lambda r: 0 if r['GENDER'] == 'm' else 1, axis=1)

test['GENDER.f'] = test.apply(lambda r: 0 if r['GENDER'] == 'm' else 1, axis=1)
```

We create a new gender column with binary values since we cannot run predictive models on string variables.

```
parameter VIF
20 MASQ22 4.037796
77 MASQ79 3.917666
14 MASQ16 3.864820
45 MASQ47 3.691897
75 MASQ77 3.642254
```

```
6 MASQ07 1.488875
83 MASQ85 1.367481
10 MASQ12 1.331534
35 MASQ37 1.320590
30 MASQ32 1.196895
[89 rows x 2 columns]
```

We sorted the VIF values in descending order in order to spot the parameters with the highest multicollinearity, but it looks like none of them really show bad multicollinearity.

```
[10]: outcome_counts = train['D_DEPDYS'].value_counts()
  outcome_prob = outcome_counts/outcome_counts.sum()
  outcome_prob
```

After checking the proportions of our response variable, it looks like our data set is pretty well balanced so we can just use a regular KFold:

```
[1]: def predictRidge(X_train, y_train, X_test, alpha):
    ridge = Ridge(alpha)
    ridge.fit(X_train, y_train)
    y_hat = ridge.predict(X_test)
    return y_hat

def predictLasso(X_train, y_train, X_test, alpha):
    lasso = Lasso(alpha)
    lasso.fit(X_train, y_train)
    y_hat = lasso.predict(X_test)
    return y_hat

def predictNet(X_train, y_train, X_test, alpha):
    elastic = ElasticNet(alpha)
    elastic.fit(X_train, y_train)
    y_hat = elastic.predict(X_test)
    return y_hat
```

```
[31]: X_train, y_train = train.drop(['D_DEPDYS', 'GENDER'], axis=1), train['D_DEPDYS']

kf = KFold(n_splits=10, shuffle=True, random_state=0)
df_accuracies = pd.DataFrame(columns=['ridge', 'lasso', 'elastic net'])

for train_ix, val_ix in kf.split(X_train, y_train):
```

```
fold_X_train, fold_X_val = X_train.iloc[train_ix], X_train.iloc[val_ix]
    fold_y_train, fold_y_val = y_train.iloc[train_ix], y_train.iloc[val_ix]
    # we need to convert the y preds into binary variables:
    y_pred_ridge = (predictRidge(fold_X_train, fold_y_train, fold_X_val, alpha=0.
 \rightarrow1) > 0.5).astype(int)
    y_pred_lasso = (predictLasso(fold_X_train, fold_y_train, fold_X_val, alpha=0.
 \rightarrow1) > 0.5).astype(int)
    y_pred_net = (predictNet(fold_X_train, fold_y_train, fold_X_val, alpha=0.1)_u
 \rightarrow 0.5).astype(int)
    df_accuracies = pd.concat([
        df_accuracies
        , pd.DataFrame({
             'ridge': [accuracy_score(y_true=fold_y_val, y_pred=y_pred_ridge)]
             , 'lasso': [accuracy_score(y_true=fold_y_val, y_pred=y_pred_lasso)]
             , 'elastic net': [accuracy_score(y_true=fold_y_val,__

    y_pred=y_pred_net)]

        })
    ])
df_accuracies = df_accuracies.mean(axis=0)
df accuracies
```

C:\Windows\Temp\ipykernel_26400\1077609598.py:15: FutureWarning: The behavior of DataFrame concatenation with empty or all-NA entries is deprecated. In a future version, this will no longer exclude empty or all-NA columns when determining the result dtypes. To retain the old behavior, exclude the relevant entries before the concat operation.

df_accuracies = pd.concat([

[31]: ridge 0.741684 lasso 0.746607 elastic net 0.754575 dtype: float64

Elastic net stands out with the highest accuracy. We will now test it on the holdout set we've kept aside until now:

```
[41]: X_test, y_test = test.drop(['D_DEPDYS', 'GENDER'], axis=1), test['D_DEPDYS']

elastic = ElasticNet(alpha=0.1)
  elastic.fit(X_test, y_test)
  y_hat = (elastic.predict(X_test) > 0.5).astype(int)
```

```
[42]: cf_mat = confusion_matrix(y_true=y_test, y_pred=y_hat)
MCR = round((cf_mat[0,1] + cf_mat[1,0]) / len(y_test), 4)
MCR
```

[42]: np.float64(0.2286) We obtain an MCR of 0.229 on the test set. Let's extract the coefficient from the elastic net model: [50]: coefs = pd.Series(elastic.coef_, index=X_test.columns) coefs = coefs[coefs !=0] coefs [50]: Leeftijd 0.001300 MASQ01 0.013209 MASQ16 0.080116 MASQ22 0.004418 MASQ29 0.002592 MASQ30 0.055978 MASQ33 0.017074 MASQ43 0.003388 MASQ46 0.000583 MASQ51 0.012450 MASQ53 0.009676 MASQ56 0.013934 MASQ60 0.000414 MASQ62 0.016268 MASQ83 0.003872 MASQ89 0.007496 dtype: float64 [46]: masq_subscales = { "Anhedonic Depression": [1, 14, 18, 21, 23, 26, 27, 30, 33, 35, 36, 39, 40, __ 44, 49, 53, 58, 66, 72, 78, 86, 89], "Anxious Arousal": [3, 19, 25, 45, 48, 52, 55, 57, 61, 67, 69, 73, 75, 79, 11 485, 87, 88], "General Distress Depression": [6, 8, 10, 13, 16, 22, 24, 42, 47, 56, 64, ... \hookrightarrow 74], "General Distress Anxiety": [2, 9, 12, 15, 20, 59, 63, 65, 77, 81, 82], "General Distress Mixed": [4, 5, 17, 29, 31, 34, 37, 50, 51, 70, 76, 80, 83, __ <u>→</u>84, 90] [47]: coefs.index [47]: Index(['Leeftijd', 'MASQ01', 'MASQ16', 'MASQ22', 'MASQ29', 'MASQ30', 'MASQ33', 'MASQ43', 'MASQ46', 'MASQ51', 'MASQ53', 'MASQ56', 'MASQ60', 'MASQ62', 'MASQ83', 'MASQ89'], dtype='object') [51]: coef_ids = list(map(lambda item: int(item.split('MASQ')[1]), coefs.index[1:]))

coef_ids

```
[51]: [1, 16, 22, 29, 30, 33, 43, 46, 51, 53, 56, 60, 62, 83, 89]
```

```
[52]: subscales = {}
for key, value in masq_subscales.items():
    counter = 0
    for i in coef_ids:
        if i in value:
            counter += 1
        subscales[key] = counter
    subscales
```

```
[52]: {'Anhedonic Depression': 5,
    'Anxious Arousal': 0,
    'General Distress Depression': 3,
    'General Distress Anxiety': 0,
    'General Distress Mixed': 3}
```

It looks like the subscale with the most selected items is anhedonic depression.