Mini Project in Statistical Machine Learning Spring 2021

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Abstract

- After testing three different algorithms on the data set we decided to proceed using the Logistic Regression algorithm. It was enhanced using Ridge Regularization.
- 3 (Will add final result here later)

4 1 Introduction

- 5 The goal of the project was to determine whether it is possible to tell if the lead actor of a movie is
- 6 female or male, based on up to 13 different variables of data, and if so, what algorithm is best suited
- 7 for doing so. The variables in the data were: Number words female, Total words, Number of words
- 8 lead, Difference in words lead and co-lead, Number of male actors, Year, Number of female actors,
- 9 Number words male, Gross, Mean Age Male, Mean Age Female, Age Lead, Age Co-Lead.

o 1.1 Cross Validation

- 11 The Cross Validation method is a way of testing a classification method against another. The idea
- behind cross validation is that the data set is split into different combinations of training and testing
- data, and the model is tested on each one of them. This leads to a better understanding of the model's
- 4 performance as we basically can generate different training and testing data sets from the same set of
- 15 data.

16 2 Methods

The following methods were implemented and tested. The implementation of each method in Python can be found in the appendix.

19 2.1 Linear Discriminant Analysis

- 20 The Linear Discriminant Analysis is a known method for dimensionality reduction, but can also
- 21 be very useful as a classification method. The goal of the method is to reduce the dimensions by
- focusing on optimizing separating the classes. The method does this by calculating the separability
- 23 between its classes, known as the between-class variance, which is the distance between the mean of
- its different classes [2]. This between-class variance can be calculated and stored in the so called
- between-class matrix with the following equation [2],

$$S_b = \sum_{i=1}^{g} N_i (\bar{x}_i - \bar{x}) (\bar{x}_i - \bar{x})^T$$
 (1)

where N_i represents the sample size of class i, the $\bar{x_i}$ is the sample mean of class i and the x_i is the overall mean. The next step in the method is to calculate the within-class variance, which is the distance between the mean and the sample of each class. This can be done with the following equation [2],

$$S_w = \sum_{i=1}^g (N_i - 1)S_i = \sum_{i=1}^g \sum_{j=1}^{N_i} (x_{i,j} - \bar{x})(\bar{x}_{i,j} - \bar{x})^T$$
 (2)

where the parameters represent the same things as in the previous equation with the addition of $x_{i,j}$ as the sample of each class. The final step in the LDA method is to create a lower-dimensional space that maximizes the separability between the classes and at the same time minimizes variance within the classes. This can be done in several different ways, some examples are by using a least square method, by using the eigenvalues or by using singular value decomposition [2]. The Scikit we used in the project does this by using the singular value decomposition if nothing else is specified. This is preferred since it does not require the computation of a covariance matrix which makes it more efficient for higher dimensions.

The between-class variance (the distance between the mean of different classes) along with the within-class variance (the distance between the mean and the sample of each class) can be seen as measures of how difficult the separation will be. Where a shorter distance in the between-class variance and a longer distance in the within-class variance indicate a more difficult separation.

42 2.1.1 Evaluation

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An important parameter when running the linear discriminant analysis is which solver to use. As 43 we mentioned before the linear discriminant analysis can be run using the least square method, eigenvalues or singular value decomposition which we will henceforth refer to as lsqr, eigen and svd respectively. We used Scikits model-selection package to compare the different solvers. The package include methods which allow you to run repeated k-fold and grid-searches which run the solvers 47 multiple times and then compare their averages. The best solver was svd with an accuracy of 0.865. 48 Another thing we explored with linear discriminant analysis was if adding a penalty in the form of a 49 shrinkage would increase performance. This is only possible for the lsqr and eigen solvers since svd 50 does not support this feature. The packages mentioned above includes options for testing shrinkage 51 aswell and implementing and testing the shrinkage where the shrinkage varied between 0 and 1 with 52 an increase of 0.01 in each step resulted in no enhancement in performance for the lsqr-solver but yielded a slight improvement for the eigen-solver when a shrinkage of 0.01 was used. However, the 54 mean accuracy for the eigen-solver was still less than the mean accuracy for the scd performance 55 and when taking these two tests in consideration we concluded that the svd-solver is the best one 56 for our particular problem. We decided to proceed with the svd-solver when comparing the linear 57 discriminant analysis with the other methods. 58

2.2 K-nearest neighbors

The k-nearest neighbors algorithm is a supervised classification algorithm that first was developed in the 1950's. It classifies an object based on the k closest objects from the training data set. The k is a positive, normally small integer that lets the model know how many neighbors the algorithm takes into consideration when classifying each object [?]. The choice of k is not clear and there is often data dependent. Different k values can result in different classification results, as seen in Figure 1 and Figure 2 below.

Since the KNN method relies on distance to assign class the data set is almost always normalized to handle different scaling and units in the data.

2.2.1 Evaluation

When testing for different k values from 1 to 23 on the entire normalized data set the KNN method failed to achieve an accuracy higher than in the low to mid seventies. This may seem like a good results, until the fact that the who data set is labeled 75.5 per cent male. The method's accuracy for

Classification with k = 1 v blue class v red class unlabeled

Figure 1: Classification of the yellow dot using k = 1 results in the blue class.

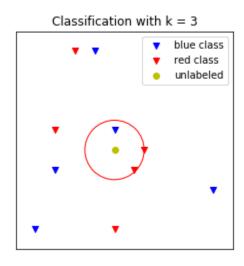


Figure 2: Classification of the yellow dot using k = 3 results in the red class.

predicting male actors was fairly good; ranging from around 0.85 to 0.95, but the number of female actors' accuracy predicted never exceeded 0.50.

As k increased the number of predicted female leads grew lower, this is a result of how skewed the data was towards male actors. For example: for a testing data set of size 159 and k=1 the model predicted 60 female leads (38 % of the total actors), but for k=11 on the same data set only 26 female leads were predicted (16 %). In Table 1 the accuracy and number of predicted females are shown.

When using k=3 and only one of the features in the data set the highest accuracy was achieved when only looking at the number of female actors. The model achieved an accuracy of 0.714, however the model only predicted that 18 of 259 lead actors were female, the lowest out of any single feature. The features Words Spoken by Males & Females, Year of Release and Gross resulted accuracies of 0.617,

Table 1: K-NN results for different K values.

K Value	Accuracy	Predicted Females
1	0.745	60
3	0.733	53
5	0.722	34
7	0.722	36
9	0.737	30
11	0.737	26
13	0.729	20
15	0.729	14
17	0.725	13
19	0.714	8
21	0.722	6
23	0.718	5

0.644, 0.629, each run predicting 53, 26, 46 female leads. When applying ten-fold sampling without replacement to the KNN model returned a lower accuracy and the number of predicted female leads was heavily reduced. We decided not to move forward with the KNN algorithm.

88 2.3 Logistic Regression

The following section uses content from the course literature [1], in order to explain the logistic regression model as a classifier.

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92 In general, a classification model can be specified in terms of the conditional class probabilities

$$p(y = m|\mathbf{x}) \quad \text{for} \quad m \in 1, ..., M \tag{3}$$

which describes the probability for class y=1,...,M, given some known input variable(s) **x**. For binary classification problems, that is M=2 and $y\in\{-1,1\}$, we can learn a model $f(\mathbf{x})$ that describes the conditional class probability for the positive class y=1

$$f(\mathbf{x}) = p(y = 1|\mathbf{x}) \tag{4}$$

where $0 \le f(\mathbf{x}) \le 1$, since it is a model for probability. And given by the law of total probability, we can model the conditional class probability for the negative class y = -1 as

$$p(y = -1|\mathbf{x}) = 1 - f(\mathbf{x}) \tag{5}$$

One classification method using conditional probabilities is the *logistic regression model*. The idea of the logistic regression is based on the *linear regression model*, together with conditional probabilities for classification. The linear regression model is given by

$$z = \theta_0 + \theta_1 x_1 + \theta_2 x_2 + \dots + \theta_n x_n + \epsilon = \boldsymbol{\theta}^T \mathbf{x} + \epsilon \tag{6}$$

where the output variable z, a numerical value, is assumed to be a linear combination of the input variables ${\bf x}$ and unknown model parameters ${\boldsymbol \theta}$, including the intercept term θ_0 . The ϵ term accounts for random errors in the data, not explained by the model. The values assigned to the parameters ${\boldsymbol \theta}$ determine the relationship between the input and output variables described by the model. In order to interpreter the output z in terms of conditional class probabilities, the logistic function $h(z) = \frac{e^z}{1+e^z}$ is used, as it is restricted to the interval [0,1]. For binary classification problems, $y \in \{-1,1\}$, this results in the logistic regression model

$$p(y=1|\mathbf{x};\boldsymbol{\theta}) = f(\mathbf{x};\boldsymbol{\theta}) = \frac{e^{\boldsymbol{\theta}^T \mathbf{x}}}{1 + e^{\boldsymbol{\theta}^T \mathbf{x}}}$$
(7)

describing the conditional class probability for the positive class y = 1. Again, by the law of total probability, this also gives the logistic regression model for the negative class y = -1

$$p(y = -1|\mathbf{x}; \boldsymbol{\theta}) = 1 - f(\mathbf{x}; \boldsymbol{\theta}) = 1 - \frac{e^{\boldsymbol{\theta}^T \mathbf{x}}}{1 + e^{\boldsymbol{\theta}^T \mathbf{x}}} = \frac{e^{-\boldsymbol{\theta}^T \mathbf{x}}}{1 + e^{-\boldsymbol{\theta}^T \mathbf{x}}}$$
(8)

Note that there is no random error ϵ included the logistic regression model (7), as in the linear regression model (6), due to the randomness in classification that is statistically modeled by the conditional class probability $p(y|\mathbf{x})$, instead of an additional error.

So, by using the logistic function, the linear regression model - a model for solving regression problems - is modified into the logistic regression model - a model for classification problems. In order to use the logistic regression model for predicting class probabilities, or actual class predictions, we need to learn the parameters $\boldsymbol{\theta}$ from the training data $T = \{\mathbf{x_i}, y_i\}_{i=1}^n$. The parameters $\boldsymbol{\theta}$ can be learned by using the *likelihood function* $\ell(\boldsymbol{\theta})$, considering the (natural) logarithm of the likelihood function for numerical reasons,

$$\ln \ell(\boldsymbol{\theta}) = \sum_{i=1}^{n} \ln p(y_i \mid \mathbf{x_i}; \boldsymbol{\theta})$$
 (9)

to find the *maximum likelihood estimate* $\hat{\boldsymbol{\theta}}$ for $\boldsymbol{\theta}$

$$\hat{\boldsymbol{\theta}} = \underset{\boldsymbol{\theta}}{\operatorname{argmax}} \ p(\mathbf{y} \mid \mathbf{x}; \boldsymbol{\theta}) = \underset{\boldsymbol{\theta}}{\operatorname{argmax}} \ \sum_{i=1}^{n} \ln p(y_i \mid \mathbf{x}_i; \boldsymbol{\theta})$$
 (10)

Solving the problem, means estimating, or learning, the parameters $\hat{\boldsymbol{\theta}}$ that maximizes the likelihood that an input $\mathbf{x_i}$ belongs to class y_i . Generally, this maximization problem is turned into the dual minimization problem $\underset{\boldsymbol{\theta}}{\operatorname{argmin}} - \sum_{i=1}^n \ln p(y_i \mid \mathbf{x_i}; \boldsymbol{\theta})$, together with the statistical average, giving us $J(\boldsymbol{\theta}) = -\frac{1}{n} \sum_{i=1}^n \ln p(y_i \mid \mathbf{x_i}; \boldsymbol{\theta})$, commonly known as the *cost function*. From the logistic regression model given in (7) and (8), together with the cost function $J(\boldsymbol{\theta})$, this gives us

$$J(\boldsymbol{\theta}) = \frac{1}{n} \sum_{i=1}^{n} \begin{cases} -\ln f(\mathbf{x_i}; \boldsymbol{\theta}) & \text{if } y_i = 1\\ -\ln (1 - f(\mathbf{x_i}; \boldsymbol{\theta})) & \text{if } y_i = -1 \end{cases}$$
(11)

which can be written in a more compact way, using the binary class formulation of $y_i = \{-1, 1\}$, since the logistic regression model in (7) and (8) gives us

$$\begin{cases}
f(\mathbf{x}; \boldsymbol{\theta}) = \frac{e^{\theta^T \mathbf{x}}}{1 + e^{\theta^T \mathbf{x}}} = \frac{e^{y_i \theta^T \mathbf{x}}}{1 + e^{y_i \theta^T \mathbf{x}}} & \text{if } y_i = 1 \\
1 - f(\mathbf{x}; \boldsymbol{\theta}) = \frac{e^{-\theta^T \mathbf{x}}}{1 + e^{-\theta^T \mathbf{x}}} = \frac{e^{y_i \theta^T \mathbf{x}}}{1 + e^{y_i \theta^T \mathbf{x}}} & \text{if } y_i = -1
\end{cases}$$
(12)

the same expression on both cases, and thus we can formulate the cost function (11) as

$$J(\boldsymbol{\theta}) = \frac{1}{n} \sum_{i=1}^{n} -\ln \frac{e^{y_i \boldsymbol{\theta}^T \mathbf{x}}}{1 + e^{y_i \boldsymbol{\theta}^T \mathbf{x}}} = \frac{1}{n} \sum_{i=1}^{n} -\ln \frac{1}{1 + e^{-y_i \boldsymbol{\theta}^T \mathbf{x}}} = \frac{1}{n} \sum_{i=1}^{n} \ln(1 + e^{-y_i \boldsymbol{\theta}^T \mathbf{x}})$$
(13)

Hence, learning a logistic regression model, means solving

$$\hat{\boldsymbol{\theta}} = \underset{\boldsymbol{\theta}}{\operatorname{argmin}} \frac{1}{n} \sum_{i=1}^{n} \ln(1 + e^{-y_i \boldsymbol{\theta}^T \mathbf{x}})$$
 (14)

and this is equivalent to finding the maximum likelihood estimate $\hat{\theta}$ as written in (10). This is a nonlinear optimization problem that needs to be solved numerically, as there is no direct solution for finding $\hat{\theta}$.

Once we have learned the parameters $\hat{\theta}$, we can use the logistic regression model,(7) and (8), to predict class probabilities, given some test input data \mathbf{x}_* . The next step is to turn these predicted class probabilities into actual class predictions $\hat{\mathbf{y}}(\mathbf{x}_*) = \mathbf{1}$ or $\hat{\mathbf{y}}(\mathbf{x}_*) = -\mathbf{1}$. The most common approach is to let $\hat{\mathbf{y}}(\mathbf{x}_*)$ be the most probable class by

$$\hat{\mathbf{y}}(\mathbf{x}_*) = \begin{cases} 1 & \text{if } f(\mathbf{x}; \hat{\boldsymbol{\theta}}) > r, \\ -1 & \text{otherwise} \end{cases}$$
 (15)

with decision threshold r = 0.5, but in general the threshold is chosen by the user $0 \le r \le 1$. The input points where the prediction changes from one class to the other, that is either of the two classes

 $(y=1 \ {
m or} -1)$ is equally probable, is the decision boundary. This corresponds to solving the equation

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$$f(\mathbf{x}; \boldsymbol{\theta}) = 1 - f(\mathbf{x}; \boldsymbol{\theta}) \tag{16}$$

and for logistic regression this means

$$\frac{e^{\boldsymbol{\theta}^T \mathbf{x}}}{1 + e^{\boldsymbol{\theta}^T \mathbf{x}}} = 1 - \frac{e^{\boldsymbol{\theta}^T \mathbf{x}}}{1 + e^{\boldsymbol{\theta}^T \mathbf{x}}} \iff e^{\boldsymbol{\theta}^T \mathbf{x}} = 1 \iff \boldsymbol{\theta}^T \mathbf{x} = 0$$
 (17)

The equation $\theta^T \mathbf{x} = 0$ indicates a (linear) hyperplane, which means that the decision boundary for logistic regression is always linear.

143 2.4 Lasso and Ridge Regularization

To improve a regression model and avoid overfitting we can use regularization. The key concept 144 behind regularization is that we want to make sure that our parameters θ as small as possible. This is 145 done to ensure that the model complexity won't be too high and can be done in many ways. In Ridge, 146 or L^2 Regularization we alter the cost function by adding a penalty, defined as the sum of the square 147 of the θ terms. They are then scaled by a penalty term, c (often called λ). The size of c can vary, but as it approaches ∞ θ will approach zero. In LASSO, or L^1 Regularization, the penalty is defined as the 149 absolute value of the θ term. The difference between these methods is that in Ridge Regularization 150 the model will typically push its θ parameters towards small non-zero values, whereas in LASSO 151 Regularization the model will set θ values to zero, and therefore can act as a feature selection model. 152 [1] 153

2.4.1 Evaluation

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When plotting the validation errors from cross validation (e.i misclassification errors) in a box plot, 155 Ridge Regularization out-performed both LASSO, as well as logistic regression without regularization, 156 for our classification problem. Next step was to choose an appropriate value for the penalty term c. 157 The was done by testing a myriad of values between 1 and 2000 and comparing the validation errors 158 between these different models. Using cross validation (k = 10), the validation errors were calculated 159 and plotted in a box diagram. We were looking for a value with an error as small as possible, but also 160 with a small error variance too. Of the tested values of c, many produced similar mean errors, but 161 c = 1000 had the "smallest box", i.e. the lowest error variance. c was therefore set to 1000. 162

3 Our Choice of Method to Proceed With

After testing the three aforementioned methods, we decided to proceed with the Logistic Regression method, and try to tune it. This was done since the model performed best in the cross validation testing.

4 Conclusions

When using the logistic regression method with Ridge regularization, c=1000, we managed to achieve a validation error of between 0.08 and 0.18.

5 The Feature Importance Task

Unfortunately, we have not yet succeeded in evaluating the importance of the different features of the 171 data. So far, we have examine the parameters $\hat{\theta}$ of the logistic regression model with regularization, 172 both of the Ridge and LASSO regression, since the values of these parameters indicate which input 173 variables x that are most important (i.e those that are most predictive). The input variables with the 174 highest valued parameters, that is the ones hat indicate importance, where especially Number of male 175 actors, Number of female actors, followed by Mean Age Male. However, this is no intuitive to us, and 176 we will go on and examine the feature importance by the following actions; choosing a combination 177 of all input variables, including non-linear transformation of these, and measuring the AIC (Akaike Information Criterion), which will indicate the best fit.

80 References

- [1] Andreas Lindholm, Niklas Wahlström, Fredrik Lindsten, and Thomas B. Schön. Supervised
 Machine Learning, 2020.
- https://smlbook.org
- [2] C.R. Rao, Venu Govindaraju. Handbook of Statistics, Machinge Learning: Theory and Applications, 2013

186 6 Appendix

```
1871 import pandas as pd
188 2 import numpy as np
189 3 import matplotlib.pyplot as plt
1915 import sklearn.preprocessing as skl_pre
1926 import sklearn.linear_model as skl_lm
1937 import sklearn.discriminant_analysis as skl_da
1948 import sklearn.neighbors as skl_nb
1959 import sklearn.model_selection as skl_ms
19610
19711 import os
19812
19913 #from IPython.display import set_matplotlib_formats
20014 #set_matplotlib_formats('png')
20115 from IPython.core.pylabtools import figsize
20216 figsize(10, 6) # Width and hight
20317 #plt.style.use('seaborn-white')
20418
20519
20620 # In [64]:
20721
20822
20923 # Set working directory and read in the csv file
21024 os.chdir("D:\Skola\System i teknik och samh lle\Statistisk
211 Maskininl rning\Project")
21225 cwd = os.getcwd()
21326 print ("The current working directory is: ", cwd)
21427
21528 trainData = pd.read_csv("train.csv")
21629
21831 # In [65]:
21932
22033
22134 #Make sure we can access the file and what it contains
22235 trainData.info()
22336 trainData.describe()
22437 print(trainData)
2258 pd.plotting.scatter_matrix(trainData)
22639
22740
22841
22942 # In [66]:
23043
23245 print(f"auto.shape: {trainData.shape}") #No. of rows, No. of columns
23447 #Split the data randomly into a training test and a test set of
      approximately same size
23648 #Set seed to get reproducible results
23749 np.random.seed(1)
23850
2331 trainI = np.random.choice(trainData.shape[0], size = 500, replace =
        False)
24152 trainIndex = trainData.index.isin(trainI)
2423 train = trainData.iloc[trainIndex]
24364 test = trainData.iloc[~trainIndex]
2456 #Size of Data ---> 1039
24657
24758
24859 # In [67]:
24960
```

```
25061
25162 #Set up train and test data
25263
25364 X_train = train[['Number words female', 'Total words', 'Number of
        words lead', 'Difference in words lead and co-lead',
254
                      'Number of male actors', 'Year', 'Number of female
25565
256
        actors', 'Number words male', 'Gross',
                      'Mean Age Male', 'Mean Age Female', 'Age Lead', 'Age
25766
       Co-Lead']]
258
25967 Y_train = train['Lead']
26088 X_test = test[['Number words female', 'Total words', 'Number of words
       lead', 'Difference in words lead and co-lead',
261
                    'Number of male actors', 'Year', 'Number of female
26269
        actors', 'Number words male', 'Gross',
263
                    'Mean Age Male', 'Mean Age Female', 'Age Lead', 'Age Co
26470
        -Lead']]
265
26671 Y_test = test['Lead']
26772
26873
26974 # In [68]:
27075
27176
27277 # set up the KNN-solver for comparison
27479 model = skl_nb.KNeighborsClassifier(n_neighbors = 1)
27580 model.fit(X_train, Y_train)
27681
27782 prediction = model.predict(X_test)
27883 print('Confusion Matrix :\n')
27984 print(pd.crosstab(prediction, Y_test), '\n')
2805 print(f'Accuracy: {np.mean(prediction == Y_test):.3f}')
28388 # In [69]:
28489
28590
28691 #Predict using LDA
288)3 model = skl_da.LinearDiscriminantAnalysis()
2894 model.fit(X_train, Y_train)
29196 predict_prob = model.predict_proba(X_test)
292)7 print('The class order in the model:')
29398 print(model.classes_)
29499
29500 print('Examples of predicted probabilities for the above classes:')
296) with np.printoptions(suppress = True, precision = 4): #Supress
       scientific notation, e.g. 1.0e-2.
297
        print(predict_prob[0:5], '\n \n Actual class:\n', trainData.loc
29802
        [0:5, 'Lead']) #Inspect the first five predictions
30003
30104
30205
30306 # In [70]:
30407
30609 prediction = np.empty(len(X_test), dtype = object)
30710 prediction = np.where(predict_prob[:,0]>=0.5, 'Female', 'Male')
30811 print ('First five predictions:')
30912 print(prediction[0:5], '\n') #Inspect the first five predictions after
        Labeling
31113 print('First five actual data:')
31214 print(trainData.loc[0:5, 'Lead'])
31416 #Confusion matrixt
```

```
31517 print(' \n Confusion matrix:')
31618 print(pd.crosstab(prediction, Y_test,),'\n')
31719
31820 #Accuracy
3191 print(f'Accuracy: {np.mean(prediction == Y_test):.3f}')
32123
32224 # In [71]:
32825
32426
32527 #Code for comparison of the different solvers for LDA
32628
32729 X_train2 = trainData[['Number words female', 'Total words', 'Number of
328
         words lead', 'Difference in words lead and co-lead',
                       'Number of male actors', 'Year', 'Number of female
32930
        actors', 'Number words male', 'Gross',
330
                       'Mean Age Male', 'Mean Age Female', 'Age Lead', 'Age
33131
        Co-Lead']]
332
33832 Y_train2 = trainData['Lead']
33433
33534 # set up the model
3365 model = skl_da.LinearDiscriminantAnalysis()
33736 # setup the evaluation method
3387 cv = skl_ms.RepeatedStratifiedKFold(n_splits=10, n_repeats=3,
339
       random_state=1)
34038 # set up grid
34139 grid = dict()
34240 grid['solver'] = ['svd', 'lsqr', 'eigen']
34841 # set up search
34412 search = skl_ms.GridSearchCV(model, grid, scoring='accuracy', cv=cv,
       n_{jobs=-1}
345
34643 # execute the search
34744 results = search.fit(X_train2, Y_train2)
34845 # summarize and evaluate which solver works best, we continue with the
349
         best one
35046 print('Mean Accuracy: %.3f' % results.best_score_)
35147 print('Config: %s' % results.best_params_)
35248 print('best param: %s' %results.best_params_)
35849
35450
35551 # In [72]:
35652
35753
35854 #Check how shrinkage affects the methods
35955
36066 X_train3 = trainData[['Number words female', 'Total words', 'Number of
         words lead', 'Difference in words lead and co-lead', 'Number of male actors', 'Year', 'Number of female
36257
        actors', 'Number words male', 'Gross',
363
                       'Mean Age Male', 'Mean Age Female', 'Age Lead', 'Age
        Co-Lead']]
36659 Y_train3 = trainData['Lead']
36760
36861
36962 # set up the model
3703 model = skl_da.LinearDiscriminantAnalysis(solver='eigen')
37164 # set up the evaluation method
37265 cv = skl_ms.RepeatedStratifiedKFold(n_splits=10, n_repeats=3,
       random_state=1)
37466 # set up the grid
37567 grid = dict()
37688 grid['shrinkage'] = np.arange(0, 1, 0.01)
37769 # set up the search
search = skl_ms.GridSearchCV(model, grid, scoring='accuracy', cv=cv,
   n_{jobs=-1}
```

```
38071 # execute the search
38#72 results = search.fit(X_train3, Y_train3)
38273 # summarize the affects of the shrinkage
38374 print('Mean Accuracy: %.3f' % results.best_score_)
38475 print('Config: %s' % results.best_params_)
38677
38778 # In [73]:
38879
38980
39081 #cross-validation comparisons of the different methods
39182
39283 X_train4 = trainData[['Number words female', 'Total words', 'Number of
        words lead', 'Difference in words lead and co-lead',
393
                      'Number of male actors', 'Year', 'Number of female
39484
        actors', 'Number words male', 'Gross',
395
                      'Mean Age Male', 'Mean Age Female', 'Age Lead', 'Age
39685
        Co-Lead']]
397
3986 Y_train4 = trainData['Lead']
39987
40088 n_fold = 10
40189
40290 \text{ models} = []
4081 models.append(skl_da.LinearDiscriminantAnalysis()) #Set up LDA
404)2 models.append(skl_nb.KNeighborsClassifier(n_neighbors = 3)) #Set up K-
       nn with k = 2
4063 models.append(skl_lm.LogisticRegression(solver='liblinear')) #Set up
407
       Logistic Regression
40894
4095 missclassification = np.zeros((n_fold, len(models)))
410% cv = skl_ms.KFold(n_splits=n_fold, random_state = 1, shuffle = True)
41197
4128 for i, (train_index, val_index) in enumerate (cv.split(X_train4)):
        X_trainfold, X_val = X_train4.iloc[train_index], X_train4.iloc[
41899
414
        val index1
41500
        Y_trainfold, Y_val = Y_train4.iloc[train_index], Y_train4.iloc[
416
        val_index]
41701
        for m in range(np.shape(models)[0]): #try different models
41802
            model = models[m]
41903
42004
            model.fit(X_trainfold, Y_trainfold)
            prediction = model.predict(X_val)
42105
            missclassification[i,m] = np.mean(prediction != Y_val)
42206
            #print(missclassification[i,m])
42307
42408
42509
            modelLDA = models[0]
            X_lda = modelLDA.fit_transform(X_trainfold, Y_trainfold)
42610
            #print(modelLDA.explained_variance_ratio_)
42711
42812
43014 print(missclassification)
43115 plt.boxplot(missclassification)
43216 plt.title('cross validation error for different methods')
43317 plt.xticks(np.arange(4)+1, ('LDA', 'K-nn', 'Logistic Regression'))
43418 plt.ylabel('validation error')
43519 plt.show()
43620
43721 #model.explained_variance_ratio_
4381 #!/usr/bin/env python
439 2 # coding: utf-8
440 3
4414 # Load info from the csv file train.csv, located at 'C:\Users\vikin\
        Documents\STS\Statistisk maskininl rning\projects\1\train.csv'.
       The file consists of 13 datapoints and then the classifier.
443
```

```
444 5 #
4456 # Implement a KNN algorithm and use the training data as a basis for
446
447 7 #
448 8 #
4499 # Number of males:
45010 # 785
45111 #
45212 # Number of females:
45313 # 254
45414 #
45515 # -> predicting only males will result in an accuracy of
       0.7555341674687199
456
45716
45817 # In[11]:
45918
46019
46120 import matplotlib.pyplot as plt
4621 import numpy as np
46322 import csv
46423
46524 # KNN related functions
46625
46726 def eucledian_distance(p1, p2):
        '''returns the eucledian distance between the two tuples p1 and p2
46827
        , , ,
469
        d = 0
47028
        for i in range(len(p1)):
47129
             d += (p1[i] - p2[i])**2
47230
        return d**0.5
47331
47432
47533 def knn_classify_me(k, distance_and_class):
        ''returns the most common class from k points with shortest
47634
        distance
477
             in the unordered list of tuples (distance, class) in
47835
479
        distance_and_class','
48036
        distance_and_class.sort(key=lambda x: x[0])
48137
        class_dict = {}
48238
48339
        for i in range(k):
48440
             temp_class = distance_and_class[i][1]
48541
             if temp_class in class_dict:
48642
                 class_dict[temp_class] += 1
48743
             else:
48844
48945
                 class_dict[temp_class] = 1
49046
        return max(class_dict, key = class_dict.get)
49147
49248
49349 def knn_return_distance_and_class_vector(p, training_values,
        training_classes):
494
        ''returns a list of tuples (distance, class) distance_and_class
49550
             for every instance in the two training lists and their
49651
        distance to the point p'',
497
49852
        distance_and_class = []
49953
        for i in range(len(training_values)):
50054
             current_point = training_values[i]
50155
             distance = eucledian_distance(p, current_point)
50256
50357
             current_class = training_classes[i]
50458
             distance_and_class.append((distance, current_class))
50559
50660
        return distance_and_class
5082 def knn_p(k, point, training_values, training_classes):
```

```
''returns the class of point p accoring to the k nearest
50963
510
        neighbors
             in trainig_values and their classes in training_classes'',
51164
        distance_and_class = knn_return_distance_and_class_vector(point,
51265
513
        training_values, training_classes)
        return knn_classify_me(k, distance_and_class)
51466
51567
5168 def knn(k, testing_values, training_values, training_classes):
         ''returns a list of classes our testing_values get based on
51769
             our training_values and training_classes','
51870
51971
        classes = []
52072
        for point in testing_values:
52173
52274
             current_class = knn_p(k, point, training_values,
523
        training_classes)
             classes.append(current_class)
52475
        return classes
52576
52677
52778
52879 # In [72]:
52980
53081
53182 # general data related functions
53384
53485 def feasible(k, testing_values, training_values, training_classes):
53586
         '''ensures that the length of all lists and tuples match'''
        if len(training_values) != len(training_classes):
53687
             print("training data doesn't match")
53788
             return False
53889
53990
        if len(training_values[0]) != len(testing_values[0]):
54091
            print("values don't match")
54192
             return False
54293
        if k > len(training_values):
54394
             print("k is too big")
54495
54596
             return False
        return True
54697
54798
54899 def results(predicted, actual):
        ''returns accuracy and prints all sorts of stuff''
54900
        predicted_males_correct = 0
55001
        predicted_females_correct = 0
55102
55203
        predicted_males_false = 0
55804
55405
        predicted_females_false = 0
55506
        for i in range(len(predicted)):
55607
55708
             if predicted[i] == "Male":
                 if actual[i] == "Male":
55809
                     predicted_males_correct += 1
55910
                 else:
56011
                     predicted_males_false += 1
56112
56213
             else:
                 if actual[i] == "Female":
56814
56415
                     predicted_females_correct += 1
                 else:
56516
                     predicted_females_false += 1
56617
        print('total predicted males {}'.format(predicted_males_correct +
56718
        predicted_males_false))
568
56919
        print('total predicted females {}'.format(
        predicted_females_correct + predicted_females_false))
570
57120
        print('predicted_males_correct / total males {}'.format(
        predicted_males_correct/(predicted_males_correct+
572
       predicted_females_false)))
573
```

```
print('predicted_females_correct / total females {}'.format(
57#21
        predicted_females_correct/(predicted_females_correct+
575
        predicted_males_false)))
576
        return (predicted_males_correct + predicted_females_correct) / len
57722
578
        (predicted)
57923
58024 def normalize(eg_array):
        ''', subtracts the mean and divides by the standard deviation'''
58125
        return (eg_array - eg_array.mean(axis=0)) / eg_array.std(axis=0)
58226
58327
58428 def split_into_k(arr, k = 10):
         ','returns a list of k subsections of arr','
58529
        part = len(arr) // k
58630
        list_of_matrices = [0] * k
58731
58832
58933
        for i in range(k):
             if i == k - 1:
59034
                 list_of_matrices[i] = arr[part * i : ]
59135
59236
                 list_of_matrices[i] = arr[part * i : part * (i + 1)]
59837
        return list_of_matrices
59438
59539
59640
59741 def combine_results(results_arrays):
        '''returns a list of the most common results for each index of the
59842
        results_arrays','
599
        results = []
60043
        for i in range(len(results_arrays[0])):
60144
             temp = {}
60245
             for j in range(len(results_arrays)):
60346
                 if results_arrays[j][i] in temp:
60447
                      temp[results_arrays[j][i]] = temp[results_arrays[j][i
60548
       ]] + 1
606
60749
                 else:
                      temp[results_arrays[j][i]] = 1
60850
             results.append(max(temp, key = temp.get))
60951
61052
        return results
61153
61254
6185 def make_predicted_csv(data, csv_link = "predictions.csv"):
        ''', takes the list data and pirnts its content on csv_link
61456
61557
             expects a list of either ints or female/male','
        with open(csv_link, 'w', newline = '') as f:
61658
             wr = csv.writer(f)
61759
             for prediction in data:
61860
61961
                  if type(prediction) == int:
                      wr.writerow(str(prediction))
62062
62163
                 else:
                      pred = prediction.lower()
62264
                      if pred == 'female':
62365
                          wr.writerow(str(1))
62466
                      else:
62567
                          wr.writerow(str(0))
62668
62769
             f.close()
62870
             return
62971
63072
63173 # In [22]:
63274
63375
63476 # testing of our algorithm
63577 # only really works in 2D
63678 ,,,
63779
```

```
6380 training_values = np.array([(-1, -10), (-3, -20), (-4, -40), (-1, -10) 

639 , (-5, -10), (10, 100), (11, 110), (10, 90), (9, 80), (20, 150)])

64081 training_classes = np.array([0, 0, 0, 0, 0, 1, 1, 1, 1, 1])
64182 testing_values = np.array([(-5, -5), (-2, -1), (4, 4), (9, 9)])
64283 k = 3
64384
64485
645% testing_classes = np.array(knn(k, testing_values, training_values,
     training_classes))
64787 print(testing_classes)
64888
64989
6500 plt.scatter(training_values[:, 0], training_values[:, 1], c =
       training_classes, label = "training")
65201 plt.scatter(testing_values[:, 0], testing_values[:, 1], c =
        testing_classes, marker = "v", label = "testing")
65#92 plt.legend()
65593 plt.show()
65694
65795
65896 print(training_values)
6597 training_values = normalize(training_values)
66098 print(training_values)','
66200
66301 # In [13]:
66402
66604 # we declare our data
66705 training_values = []
668)6 training_classes = []
66907 testing_values = []
67008 testing_actual = []
67209
67210 total_lines = 1039
67311
67412 number_of_training = 780
67513
67614
67715 link = r"C:\Users\vikin\Documents\STS\Statistisk maskininl rning\
       projects\1\train.csv"
67916 short = r"C:\Users\vikin\Documents\STS\Statistisk maskininl rning\
680
        projects\1\short.csv"
68117
68218 # we fill our data
68319 file = open(link)
68420 csv_reader = csv.reader(file)
6851 next(csv_reader) # skip the header
68622
68723 i = 0
68824 for row in csv_reader:
68925
         if i < number_of_training:</pre>
              training_values.append(tuple(row[0 : -1]))
69026
             training_classes.append(row[-1])
69127
69228
             testing_values.append(tuple(row[0 : -1]))
69329
              testing_actual.append(row[-1])
69430
         i += 1
69531
69733 # normalization of the data
6984 training_values = np.array(training_values).astype(np.float)
69935 training_values = normalize(training_values)
70036
70137 testing_values = np.array(testing_values).astype(np.float)
70238 testing_values = normalize(testing_values)
```

```
70339
70440
70541 # # Finding the Best K Value
70642 # We'll test K values in [1, 41] and pick a K value based on accuracy.
70844 # In [29]:
70945
71046
71147 for k in range(1, 42):
71248
        print("K:", k)
71349
        testing_classes = np.array(knn(k, testing_values, training_values,
712450
        training_classes))
715
71651
        print(results(testing_classes, testing_actual))
71752
71853
71954 # # About the Data
72055 # To test only certian attributes:
          modify the attributes list to contain the indices you want to
72156 #
722
        test and move it like below
72357 #
          altered_testing = testing_values[:, attributes]
72458 #
          altered_training = training_values[:, attributes]
72559 #
72660 #
72761 #
72862 # ## Key:
72963 # * Number words female: 0,
73064 # * Total words: 1,
73165 # * Number of words lead: 2,
73266 # * Difference in words lead and co-lead: 3,
73367 # * Number of male actors: 4,
73468 # * Year: 5,
73569 # * Number of female actors: 6,
73670 # * Number words male: 7,
737/1 # * Gross: 8,
73872 # * Mean Age Male: 9,
73973 # * Mean Age Female: 10,
74074 # * Age Lead: 11,
74175 # * Age Co-Lead: 12
74276
74377 # In [61]:
74478
74579
74680 # let's run this shit
74781 #for i in range(13):
74882 #print(i)
74983 attributes = [0,7] #[0,5,7,8]
75084
75185 k = 3
752% altered_testing = testing_values [:, attributes]
75387 altered_training = training_values [:, attributes]
75488
7559 testing_classes = np.array(knn(k, altered_testing, altered_training,
        training_classes))
756
75790 print(results(testing_classes, testing_actual))
75891
75992
76093 # # Sampling Without Replacement
762)5 # We split our matrix into k subsets and predict based on that. We the
763
        choose the most common prediction.
76496
76597 # In [35]:
76698
76799
```

```
76800 k_split = 10
76901 \text{ k_neighbors} = 3
77.002
77303 #print(len(training_values))
77204 #print(training_values)
77406 #print(len(training_classes))
77507 #print(training_classes)
77608
77709 split_training_values = split_into_k(training_values, k_split)
77810 #print(split_training_values)
779 || split_training_classes = split_into_k(training_classes, k_split)
78012 #print(split_training_classes)
78313
78214 sampling_results = [0] * k_split
78315
78416 #print(split_training_values[i])
78517 #print(split_training_classes[i])
78618
78719 for i in range(k_split):
        sampling_results[i] = knn(k_neighbors, altered_testing,
78820
        split_training_values[i], split_training_classes[i])
789
79.021
79223 #print(cross_validaton_results)
79324 sampling_predictions = combine_results(sampling_results)
79425
79526
7967 print(results(sampling_predictions, testing_actual))
79728
79829
79930 # # Making a CSV File
80132 #
80233
80334 # In [71]:
80435
80536
80637 import os
80738 print(os.getcwd())
80940 d = [1,0,1,0,1,0,1,0,2,3,1,1]
81041 #d = ["male", "female", "MAle", "Female"]
81342
81243
81344 make_predicted_csv(d)
81345
81546
81647 # In [64]:
81748
81950 # 2D plot
82051
82152
82253
82354 blue_class = np.array([(1,1),(4,10),(5,6),(10, 3),(2,4)])
8245 red_class = np.array([(5,1),(6,4),(2,6),(3, 10),(6.5,5)])
8256 yellow_class = np.array([(5, 5)])
82657
82758
82859
82960 fig, ax = plt.subplots()
83061 #plt.xticks([])
83162 #plt.yticks([])
83263
```

```
83364 ax.set(xlim=(0, 11), ylim = (0, 11))
83465 ax.set_aspect('equal')
83566 ax.axes.xaxis.set_visible(False)
83667 ax.axes.yaxis.set_visible(False)
8389 ax.scatter(blue_class[:, 0], blue_class[:, 1], c = "b", marker = "v",
       label = "blue class")
84070 ax.scatter(red_class[:, 0], red_class[:, 1], c = "r", marker = "v",
       label = "red class")
842/1 ax.scatter(yellow_class[:, 0], yellow_class[:, 1], c = "y", label = "
       unlabeled")
84472 #ax.scatter([x^2 + y^2 = 1])
84573 cir1 = plt.Circle((5, 5), 1, color='b',fill=False)
84674 #ax.add_artist(cir1)
847/5 cir2 = plt.Circle((5, 5), 1.5, color='r',fill=False)
84876 ax.add_artist(cir2)
84977 plt.title('Classification with k = 3')
85078
85179
85280
85381 ax.legend()
85482
85583 fig.savefig('plot2.png')
85785 #ax.show()
85886
85987 ,,,
86088
86189 training_values = np.array([(-1, -10), (-3, -20), (-4, -40), (-1, -10)
       , (-5, -10), (10, 100), (11, 110), (10, 90), (9, 80), (20, 150)])
86390 training_classes = np.array([0, 0, 0, 0, 0, 1, 1, 1, 1, 1])
86%) testing_values = np.array([(-5, -5), (-2, -1), (4, 4), (9, 9)])
86592 k = 3
86693
86794
868)5 testing_classes = np.array(knn(k, testing_values, training_values,
869
        training_classes))
87096 print(testing_classes)
87397
87298
8739 plt.scatter(training_values[:, 0], training_values[:, 1], c =
       training_classes, label = "training")
87500 plt.scatter(testing_values[:, 0], testing_values[:, 1], c =
       testing_classes, marker = "v", label = "testing")
87701 plt.legend()
87802 plt.show()
87903
88004
88405 print(training_values)
882)6 training_values = normalize(training_values)
88307 print(training_values)'',
884 | #!/usr/bin/env python
885 2 # coding: utf-8
886 3
887 4 # In[]:
888 5
889 6
890 7 import pandas as pd
8918 import numpy as np
892 9 import matplotlib.pyplot as plt
89310 import csv
89411
89512 #scikit
89613 import sklearn.preprocessing as skl_pre
```

```
89714 import sklearn.linear_model as skl_lm
89815 import sklearn.model_selection as skl_ms
90017 from IPython.core.pylabtools import figsize
90118 figsize(10, 6)
90320
90421 # In[]:
90522
90623
90724 # Read data
90825 directory = 'directory of dataset'
90926 traindata = pd.read_csv(directory, na_values='?', dtype={'Lead': str})
       .dropna().reset_index()
91127 #traindata.info()
91228
91329 # Preprocessing data
91430 np.random.seed(1)
91531
91622 # Splitting up dataset in training and test data: 538 training, 500
917
9183 trainIndex = np.random.choice(traindata.shape[0], size=538, replace=
919
       False)
92135 trainIndexBool = traindata.index.isin(trainIndex) # generates a np
     array with True/False if corresponding index in in trainIndex
9236 train = traindata.iloc[trainIndexBool] # training set of 538
9247 test = traindata.iloc[~trainIndexBool] # test set of 500 (1038-538)
92538
92639 # Feature selection
92740 features = ['Number words female', 'Total words', 'Number of words
       lead', 'Difference in words lead and co-lead', 'Number of male
       actors', 'Year', 'Number of female actors', 'Number words male',
       Gross', 'Mean Age Male', 'Mean Age Female', 'Age Lead', 'Age Co-
930
       Lead']
931
93241
93342 # Training set
93443 X_train = train[features] # input
93544 Y_train = train['Lead'] # output
93645
93746 # Test set
93847 X_test = test[features]
93948 Y_test = test['Lead']
94049
94150
94251 # In[]:
94352
94453
94554 # Learn model
9465 log_reg_model_ridge = skl_lm.LogisticRegression(solver='liblinear',
       penalty='12', C=1000) # Ridge Regression
94856 log_reg_model_ridge.fit(X_train,Y_train)
9497 #print('Model coefficients, Ridge Logistic Regression: n',
       log_reg_model_ridge.coef_, '\n')
950
95158
95259 # Prediction
95360 r = 0.5
95461 prediction = np.where(prob[:,0] >= r, 'Female', 'Male')
95663 # Confusion matrix
95%4 print('Confusion matrix:\n', pd.crosstab(prediction, Y_test), '\n')
95865
95966 # Accuracy
9607 print(f'Accuracy: {np.mean(prediction == Y_test):.3f}', '\n') # rounds
up to 3 decimals
```

```
96268
96470 # In[]:
96571
96672
96773 # Compare Logistic regression models with/without regularization
96874 log_reg_model_lasso = skl_lm.LogisticRegression(solver='liblinear',
        penalty='11', C=1000) # Lasso Regression
97075 log_reg_model_ridge = skl_lm.LogisticRegression(solver='liblinear',
        penalty='12', C=1000) # Ridge Regression
97276 log_reg_model_none = skl_lm.LogisticRegression(solver='lbfgs', penalty
        ='none', max_iter=2500) # no regularization
973
97477
97578 \text{ models} = []
97679 models.append(log_reg_model_lasso)
97780 models.append(log_reg_model_ridge)
9781 models.append(log_reg_model_none)
97982
98083 # learn model from training data
98184 for model in models:
         model.fit(X_train,Y_train)
98285
98386
98487 print('Model coefficients, Lasso Logistic Regression: \n',
        log_reg_model_lasso.coef_, '\n')
986% print ('Model coefficients, Ridge Logistic Regression: \n',
        log_reg_model_ridge.coef_, '\n')
987
9889 print('Model coefficients, Logistic Regression: \n',
        log_reg_model_ridge.coef_, '\n')
99090
99191
99292 # In[]:
99393
9955 pred_prob_lasso = log_reg_model_lasso.predict_proba(X_test)
9986 pred_prob_ridge = log_reg_model_ridge.predict_proba(X_test)
99797 pred_prob_none = log_reg_model_none.predict_proba(X_test)
99999 models_prob = {}
100000 models_prob[log_reg_model_lasso] = pred_prob_lasso
100101 models_prob[log_reg_model_ridge] = pred_prob_ridge
1002)2 models_prob[log_reg_model_none] = pred_prob_none
100303
100404 # Descision
100505 r = 0.5
100606
100707 for model, prob in models_prob.items():
             print('Class order in the model: ', model.classes_)
100808
100909
101010
             # Print first 5 predictions
             print('Examples of predicted probabilities for the above
101111
        classes: \n', item[1][0:5], '\n')
1012
             \# print(f'Average probability for Female lead is \{100*(sum(
101812
        prob[:,0])/len(prob)):.0f} %')
1014
        # print(f'Average probability for Male lead is {100*(sum(prob
[:,1])/len(prob)):.0f} %', '\n')
101513
1016
101714
             # Prediction
101815
101916
             prediction = np.where(prob[:,0] >= r, 'Female','Male')
102017
102118
             # Confusion matrix
102219
             print('Confusion matrix:\n', pd.crosstab(prediction, Y_test),
        '\n')
1023
102420
102521
             # Accuracy
```

```
print(f'Accuracy: {np.mean(prediction == Y_test):.3f}', '\n')
102622
1027
         # rounds up to 3 decimals
              # print('Accuracy: ', np.mean(prediction == Y_test))
102823
102924
103025
              # Precision
103126
              # Recall
103227
103828
              # F1
103429
103530
103631
103732 # In[]:
103833
103934
104035 # decide positive resp negative class
104136 pos_class = 'Male'
104237 neg_class = 'Female'
104838
104439 # count num of positive resp negative samples in test
104540 P = np.sum(Y_test == pos_class) # num of positive samples in test
104641 N = np.sum(Y_test == neg_class) # num of negative samples in test
104843 # find indices with male/positive class in data
104944 pos_class_index = np.argwhere(log_reg_model.classes_ == pos_class).
1050
         squeeze()
105145
105246 # lists to append TP / FN rate for different r
105847 true_pos_rate = []
105#48 false_pos_rate = []
105549
105600 \text{ # r } [0,1], \text{ dvs } r= 0.00, 0.01, \dots, 0.1, 0.11, \dots, 0.99, 1.0
105%1 threshold = np.linspace(0.00, 1, 101) # start=0, stop=1, 101 num
         evenly spaced samples
105952
106053 for model, prob in models_prob.items():
         for r in threshold:
106154
106255
              prediction = np.where(prob[:,pos_class_index] > r, pos_class,
         neg_class)
1063
              TP = np.sum((prediction == pos_class) & (Y_test == pos_class))
106456
          #True Positive
1065
              true_pos_rate.append(TP/P)
106657
106758
              FP = np.sum((prediction == pos_class)&(Y_test == neg_class)) #
         False Positive
1068
             false_pos_rate.append(FP/N)
106959
107060
107161
         plt.plot(false_pos_rate, true_pos_rate);
         for i in [0,1,10,50,98,100]:
107262
              plt.text(false_pos_rate[i], true_pos_rate[i], f'r={threshold[i
107863
         ]:.2f}')
1074
         plt.xlim([0, 1])
107564
107665
         plt.ylim([0, 1.1])
         plt.xlabel('False positive rate')
107766
         plt.ylabel('True positive rate');
107867
107968
108069 # print('false_pos_rate',false_pos_rate)
108170 # print('true_pos_rate', true_pos_rate)
108271
108872
108#73 # In[]:
108574
108675
1087/6 # Cross validation
108877
108978 \text{ n_fold} = 10
109079
```

```
109180 models = []
109281 models.append(skl_lm.LogisticRegression(solver='liblinear', penalty='
        11', C=1000)) # Lasso Regression
109482 models.append(skl_lm.LogisticRegression(solver='liblinear', penalty='
1095
        12', C=1000)) # Ridge Regression
10963 models.append(skl_lm.LogisticRegression(solver='lbfgs', penalty='none'
1097
        , max_iter=2500)) # no regularization
109884
10985 missclassification = np.zeros((n_fold, len(models)))
1100% cv = skl_ms.KFold(n_splits=n_fold, random_state = 1, shuffle = True)
110288 for i, (train_index, val_index) in enumerate (cv.split(X_train)):
        X_trainfold, X_val = X_train.iloc[train_index], X_train.iloc[
110389
1104
        val_index]
        Y_trainfold, Y_val = Y_train.iloc[train_index], Y_train.iloc[
110590
        val_index]
1106
110791
        for m in range(np.shape(models)[0]): #try different models
110892
             model = models[m]
110993
             model.fit(X_trainfold, Y_trainfold)
111094
             prediction = model.predict(X_val)
111195
             missclassification[i,m] = np.mean(prediction != Y_val)
111296
111397
111#98 print(missclassification)
111599 plt.boxplot(missclassification)
111600 plt.title('cross validation error for different methods')
111701 plt.xticks(np.arange(4)+1, ('Logistic with Lasso', 'Logistic with
       Ridge', 'Logistic no regularization'))
111902 plt.ylabel('validation error')
112003 plt.show()
112104
112205
112306 # In[]:
112407
112508
1126)9 # Cross validation, changing pentalty parameter C (or lambda)
112811 n_fold = 10
[112912] C = [1,10,50,100,400,999,1000,2000,5000]
11314 missclassification = np.zeros((n_fold, len(C)))
113215 cv = skl_ms.KFold(n_splits=n_fold, random_state = 1, shuffle = True)
113316
113417 for i, (train_index, val_index) in enumerate (cv.split(X_train)):
113518
        X_trainfold, X_val = X_train.iloc[train_index], X_train.iloc[
1136
        val_index]
        Y_trainfold, Y_val = Y_train.iloc[train_index], Y_train.iloc[
113719
        val_index]
1138
113920
        for j in range(len(C)): #try different models/C
114021
             model = skl_lm.LogisticRegression(solver='liblinear', penalty=
114322
        '12', C=C[j]) # Ridge Regression
1142
             model.fit(X_trainfold, Y_trainfold)
114323
             prediction = model.predict(X_val)
114224
114525
             missclassification[i,j] = np.mean(prediction != Y_val)
114626
#print(missclassification,'\n')
114828 plt.boxplot(missclassification)
11499 plt.title('cross validation error for different methods')
115080 plt.xticks(np.arange(9)+1, ('c=1', 'c=10', 'c=50', 'c=100', 'c=400', '
     c=999','c=1000', 'c=1001', 'c=5000'))
plt.ylabel('validation error')
115332 plt.show()
```