Project Title

Comparison of Logistic Regression and SVM Method

Members

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Contents

- 1.Logistic Regression
- a. Setting up Logistic Regression Model
- b.The Main Goals of Logistictic Regression Model
- c.Setting up a Model in R
- 2.SVM Model
- a.Definition
- b.Equations of SVM Model
- c. SVM Kernels
- 3.ANALYSIS
- a.Logistic Regression Model Process
- b. SVM Model Process
- c. Chi-Square Method
- d. Chi-Square Method in R

Description of the project

In this project, we compare two methods that are logistic regression method and SVM method by using 1984 United Stated Congressional Voting Records in R language. We analyse that which method is better to predict the congressman belongs to which party according to first column of the data to the rest of them.

The methods to be used

There are two different methods that are logistic regression method and SVM method in this project. Logistic regression is a type of a prediction method that is used if the data has binary dependent variables and this method determines the relationship between the set of dependent variables and the independent set.

Support Vector Machines method is another prediction method that is used if the data is binary too. However, in this method two kinds of accumulation of dependent variables are determined and divided down the middle of the distance that is called as margin.

The data

The data is 1984 United Stated Congressional Voting Records that has 435 rows and 17 columns. This data consists of 435 congressmen that are 267 democrats and 168 republicans. In this data has binary dependent variables such as yes or no and two class of congressmen that are republican and democrat. There are 16 numbers of attributes that are categorical.

Code

As proof of work, you must run this notebook. Upload an HTML output of this notebook on your github account.

X<- read.csv("https://archive.ics.uci.edu/ml/machine-learning-databases/voting-records/house

head(X) V1 V2 V3 V4 V5 V6 V7 V8 V9 V10

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1.Logistic Regression

a. Setting up Logistic Regression Model

Logistic regression is a type of a prediction method that is used if the data has binary dependent variables and this method determines the relationship between the set of dependent variables and the independent set. Odds is the ratio of the probability of occurrence with probability of event that does not occur in this formula stated in [1].

$$odds = p/(1-p) \tag{1.1}$$

There is link between the Binomial distribution and the linear combination of independent variables and this function is called logit function presented in [1].

$$logitp = log(odds) = log(p/(1-p)) = \beta_0 + \beta_1 x_1 + \dots + \beta_k x_k$$
 (1.2)

This formula resembles a linear regression but logistic regression gets a best fitting by using maximum likelihood function that maximizes the probability.

$$p = \frac{e^{\beta_0 + \beta_1 x_1 + \dots + \beta_k x_k}}{1 + e^{\beta_0 + \beta_1 x_1 + \dots + \beta_k x_k}}$$
(1.3)

where β_0 is constant, $\beta_1, ..., \beta_k$ are the coefficients of predictor variables, e is a natural logarithm and p is the probability of a specific case.

Also we can write this equation which is emphasized in [1] as the following

$$e(logitp) = e^{(\beta_0) + (\beta_1 x_1) + \dots + (\beta_k x_k)}$$
 (1.4)

then

$$o = p/(1-p) = e^{(\beta_0)}e^{(\beta_1 x_1)}e^{(\beta_2 x_2)}....e^{(\beta_k x_k)}$$
(1.5)

where β'_k s are constant coefficients that determines the changes of logistic regression model when x_i is added. Also, adding or subtracting a unit alters the odds with constant amount.

b. The Main Goals of Logistictic Regression Model

There are two primary purposes of logistic regression models. First one of these purposes is to predict of group membership. This model uses the odds ratio; therefore, the conclusion of using this model has the the same form with the odds ratio. The second aim is suplying informations about connections and strongness between the variables.

c.Setting up a Model in R

Initially, we divide the data to constitute our model and we divide 30% of the data randomly. To do this we use this code.

```
sample(1:nrow(X), 0.3*nrow(X))
test <-sample(1:nrow(X), 0.3*nrow(X))</pre>
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130 members of this data is selected randomly by using this code in R but these numbers are not useful for testing and forming our model. Therefore, we convert them to a matrix by using the code below.

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We use the rest of the model data to test our model; consequently, it is assigned to testdata variable.

testdata<-X[-test,] nrow(testdata)</pre>

305

У

Logistic regression model is formed by using glm function in R language. The meaning of glm is fitting generalized linear models.

 $\label{logitmodel} $$\log itmodel <- glm(V1 \sim V4+V7+V9+V12+V15+V17, \ family=binomial(link="logit"), \ data= \ modeldata)$$ logitmodel$

Call: $glm(formula = V1 \sim V4 + V7 + V9 + V12 + V15 + V17, family = binomial(link = "logit") data = modeldata)$

Coefficients:

OCCITIONOD.					
(Intercept)	V4n	V4y	V7n	V7y	V9n
3.2776	3.3063	-0.2505	-1.3029	-2.8804	-2.2424
V9y	V12n	V12y	V15n	V15y	V17n
-4.9442	1.2610	-1.8200	-18.1931	-0.0705	0.1346
V17y					
0.5301					

Degrees of Freedom: 129 Total (i.e. Null); 117 Residual

Null Deviance: 163.6

Residual Deviance: 42.88 AIC: 68.88

After the model is generated, we use summary code to get a short information about the model that we generate.

```
summary(logitmodel)
anova(logitmodel, test="Chisq")

Call:
glm(formula = V1 ~ V4 + V7 + V9 + V12 + V15 + V17, family = binomial(link = "logit"),
```

Deviance Residuals:

data = modeldata)

Min 1Q Median 3Q Max -2.53865 -0.06169 -0.00002 0.28523 2.49880

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	3.2776	4.5160	0.726	0.468
V4n	3.3064	4.5938	0.720	0.472
V4y	-0.2505	4.5355	-0.055	0.956
V7n	-1.3029	4.5629	-0.286	0.775
V7y	-2.8804	4.4843	-0.642	0.521
V9n	-2.2424	3.4889	-0.643	0.520
V9y	-4.9442	3.5722	-1.384	0.166
V12n	1.2610	2.0928	0.603	0.547
V12y	-1.8200	2.1078	-0.863	0.388
V15n	-18.1931	2167.5919	-0.008	0.993
V15y	-0.0705	3.1309	-0.023	0.982
V17n	0.1346	1.4207	0.095	0.925
V17y	0.5301	1.1543	0.459	0.646

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 163.583 on 129 degrees of freedom Residual deviance: 42.883 on 117 degrees of freedom

AIC: 68.883

Number of Fisher Scoring iterations: 19

Df

Deviance

Resid. Df

Resid. Dev

 $\Pr(>\text{Chi})$

NULL

NA

NA

129

163.58343

NA

V4

2

79.347659

127

84.23577

5.886740 e-18

V7

2

3.490466

125

80.74531

1.746043e-01

V9

2

15.116268

123

65.62904

5.218482 e-04

V12

2

17.154162

```
48.47488

1.883740e-04

V15

2

5.312743

119

43.16213

7.020250e-02

V17

2

0.279193

117

42.88294

8.697091e-01
```

After the model is formed, we use predict function to estimate the model for testdata variable which is the rest of the modeldata.

```
prelogitmodel<-predict(logitmodel,testdata, type = "response")</pre>
```

Now, prelogitmodel contains the probability that given MP is a republican or a democrat. We need to set a threshold for the probability. If it is greater than 0.6then we will predict that MP is a republican

```
a<-ifelse(prelogitmodel>0.6, "republican", "democrat")
table(a)
a
  democrat republican
  206 99
```

To compare logitmodel that we constitute and prelogitmodel that predict our model, we use table function. We need to compare the number of democrat congressmen and republican congressmen. For this reason, we take V1 column of the testdata that consists of the class of the congressmen which are democrat and republican. Also, the table(a) that is formed by using prelogitmodel consists of the number of the congressmen that belongs to two class.

```
logistic_table1<-table(real=testdata$V1,predicted=a)</pre>
```

logistic_table1

real democrat republican democrat 173 6 republican 33 93

When we look at the anova function that gives informations about prelogitmodel, we take out laws that their residual deviance has a small differences. Therefore, V9 and V17 are taken out from the code to change our model.

```
logitmodel<- glm(V1 ~ V4+V9+V12, family=binomial(link="logit"), data= modeldata)
logitmodel</pre>
```

Call: glm(formula = V1 ~ V4 + V9 + V12, family = binomial(link = "logit"),
 data = modeldata)

Coefficients:

```
(Intercept) V4n V4y V9n V9y V12n
2.178 2.601 -1.217 -2.634 -5.433 1.087
V12y
-2.217
```

Degrees of Freedom: 129 Total (i.e. Null); 123 Residual

Null Deviance: 163.6

Residual Deviance: 49.99 AIC: 63.99

```
summary(logitmodel)
anova(logitmodel, test="Chisq")
```

Call:

```
glm(formula = V1 ~ V4 + V9 + V12, family = binomial(link = "logit"),
    data = modeldata)
```

Deviance Residuals:

```
Min 1Q Median 3Q Max -2.55772 -0.25809 -0.04986 0.27824 2.61483
```

Coefficients:

```
Estimate Std. Error z value Pr(>|z|)
(Intercept) 2.178 2.580 0.844 0.39861
V4n 2.601 2.807 0.926 0.35419
```

```
V4y -1.217 2.769 -0.440 0.66026

V9n -2.634 1.880 -1.401 0.16121

V9y -5.433 1.965 -2.765 0.00569 **

V12n 1.087 1.995 0.545 0.58590

V12y -2.217 2.000 -1.109 0.26761
```

Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 163.58 on 129 degrees of freedom Residual deviance: 49.99 on 123 degrees of freedom

AIC: 63.99

Number of Fisher Scoring iterations: 7

Df

Deviance

Resid. Df

Resid. Dev

Pr(>Chi)

NULL

NA

NA

129

163.58343

NA

V4

2

79.34766

127

84.23577

5.886740 e-18

V9

2

17.44960

```
125
66.78617
1.625051e-04
V12
2
16.79647
123
49.98970
2.252650e-04
When we look at the table above, the differences between the residual deviances
are close to each other. Consequently, we constitute our logitmodel again without
V7, V9 and V17 columns.
prelogitmodel<-predict(logitmodel,testdata, type = "response")</pre>
a<-ifelse(prelogitmodel>0.6, "republican", "democrat")
table(a)
  democrat republican
                    100
        205
```

```
predicted
real democrat republican
democrat 173 6
republican 32 94
```

logistic_table2

According to table functions, the total number of congressmen that is known wrongly is 49; therefore, these two table functions are compared by using Pearson's Chi-squared test. To use Pearson's Chi-squared test, chisq.test(matrix()) is used and compared it X-squared values. The logitmodel is statistically good when it has a big X-squared value.

logistic_table2<-table(real=testdata\$V1,predicted=a)</pre>

```
chisq.test(logistic_table1)
chisq.test(logistic_table2)
```

Pearson's Chi-squared test with Yates' continuity correction

data: logistic_table1
X-squared = 164.25, df = 1, p-value < 2.2e-16</pre>

Pearson's Chi-squared test with Yates' continuity correction

data: logistic_table2
X-squared = 167.14, df = 1, p-value < 2.2e-16</pre>

The second modal that is used has bigger X-squared value and it has less independent variables; consequently, using this model is statistically good. However, we will choose the second logistic model if we have small X-squared number because the second logistic model has a less independent variables than the first logistic model.

2.SVM Model

a.Definition

SVM is the contraction of Support Vector Machines. Cortes & Vapnik developed SVM's for binary classification. In this model, support vectors are determined by using the closest points that are the elements of the classes. Support vectors pass on the closest points and the distance between the two of support vectors are called margin. Also margin is maximized in this model. There is a hyperplane that seperates the support vectors and margin in the middle. In this model, each of classes shows a tendency to -1 or 1. Therefore, there are two hyperplanes that are $w.x_i >= 1$ when $y_i = 1$ and $w.x_i <= -1$ when $y_i = -1$ for x_i 's are the set of input, y_i is set of output corresponding to x_i and w is the weight vector that predicts the y_i value. H_1 and H_2 are planes such that

$$H_1: w.x_i = 1 (2.1)$$

and

$$H_2: w.x_i = -1. (2.2)$$

Also there is a plane in the middle of this planes such that

$$H_0: w.x_i = 0. (2.3)$$

These equations of hyperplanes are stated in [3]. The shortest distance between H_0 and H_1 is called +d and the shortest distance between H_0 and H_2 is called -d.

b.Equations of SVM Model

Considering the movement of the support vector changes the boundary; concequently, the form of equation of hyperplanes occurs such that $w^T.x + b = 0$ where w is a weight vector, x is an input vector and b is bias. The distance between H_0 and H_1 is (w.x + b)/||w|| = 1/||w||, then the margin is 2/||w||. To maximize margin,w can be minimized. When hyperplanes that we mentioned are gathered up, the equation $y_i(w.x_i) >= 1$ is formed. Suppose that we have a problem and we need to minimize w,

$$minf(x) = (1/2)||w||^2$$
 (2.4)

and $g(x) = y_i(w.x_i) + b = 1$ or g(x) can be written as $g(x) = (y_i(w.x_i) + b) - 1 = 0$. It can be solved by Langrange Multiplier and we have two constraints that one of them is g(x) = 0 and the solution maximum or minimum according to the gradient of f or g. When these two constraints are considered the Langrange Equation which is analysed in [3] must be like this L(x, a) = f(x) - ag(x) and the derivative of L(x, a) is zero. In general

$$L(x,a) = f(x) - \sum_{i} a_{i}g_{i}$$
 (2.5)

In this case given f(x) and g(x) we have Langrangian

$$minLangrangian = (1/2)||w||^2 - \sum_{i} a_i((y_i(w.x_i) + b) - 1)$$
(2.6)

and the last form of the equation is

$$minLangrangian = (1/2)||w||^2 - \sum_{i} a_i((y_i(w.x_i+b) + \sum_{i} a_i))$$
 (2.7)

In SVM model the Langrangian stated in is

$$(1/2)||w||^2 - \sum_{i=1}^l a_i((y_i(w.x_i + b) + \sum_{i=1}^l a_i)$$
(2.8)

where $i, a_i > 0$ and l is the number of training points. The derivatives of Langrangian with respect to w and b are zero therefore, we get

$$w = \sum_{i=1}^{l} a_i y_i x_i \qquad and \qquad \sum_{i=1}^{l} a_i y_i = 0$$
 (2.9)

When we rewrite the minimum Langrangian by putting w and $b = \sum_{i=1}^{l} a_i y_i = 0$ then we have maximum Langrangian

$$maxL = \sum_{i=1}^{l} a_i - (1/2) \sum_{i=1}^{l} a_i a_j y_i y_j (x_i \cdot x_j)$$
 (2.10)

where a_i 's are support vectors and positive. The above-stated equations about SVM model emphasized in [3] and these equations informed about the occurrence of this model.

In this equation that is emphasized in [4] we learned the margin is maximized or not by using the inner product of x_i and x_j and owing to Kernel function, this computation is easier because Kernel function

$$K(x_i, x_i) = \phi(x_i).\phi(x_i)$$
 (2.11)

describes the inner product or resembels in transformed space.

c. SVM Kernels

There are kernels that are used in SVM model. When the SVM model increases the dimensions of transformed space by using the data, the model determines the proper kernel. There are three type of kernels that are most known and the equations of kernels are determined in [3].

1. Polinomial Kernel

$$K(x,y) = (x.y+1)^p (2.12)$$

2. Radial Basis Kernel

$$K(x,y) = \exp(-(|x-y|^2)/2\sigma^2)$$
 (2.13)

3.Sigmoid Kernel

$$K(x,y) = \tanh(\kappa x.y - \delta) \tag{2.14}$$

First of all, we need to download the library of SVM for using the model in R. Therefore, we use this code below

```
library (e1071)
```

After we downloaded the library, we construct our SVM model by using modeldata and classes that are the first column of this data.

```
svmmodel<- svm(V1 ~ . , data=modeldata)
svmmodel

Call:
svm(formula = V1 ~ ., data = modeldata)

Parameters:
   SVM-Type: C-classification
SVM-Kernel: radial
        cost: 1
        gamma: 0.03030303</pre>
Number of Support Vectors: 43
```

After the symmodel is constructed, to take a short information about symmodel, summary() function can be used as the following.

```
summary(svmmodel)

Call:
svm(formula = V1 ~ ., data = modeldata)

Parameters:
   SVM-Type: C-classification
SVM-Kernel: radial
   cost: 1
   gamma: 0.03030303

Number of Support Vectors: 43

( 21 22 )
```

```
Number of Classes: 2
```

Levels:

democrat republican

As indicated above the C-classification is one of the dual representation of SVM and in this type of classification the error function minimized. The minimized error function is $(1/2)ww^T + C\sum i = 1^n\xi_i$ subject to constraints: $y_i(w^T\phi(x_i) + b) >= 1 - \xi_i$ where $\xi_i > 0$ and i = 1, 2, ..., N. In error function C is the capasity constant, w is weight vector or vector of coefficient, b is constant, i is the number of training, i is the sign of class and i is the set of independent variables. Besides the kernel $\phi(x_i)$ converts from the input data(independent variables) to feature space.

We need to predict that the symmodel after we set up model by using testdata. Also predict() function is used to predict symmodel and it assigns sympredict variable.

```
svmpredict<- predict(svmmodel,testdata)
svmpredict</pre>
```

4

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democrat

7

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10

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democrat

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 ${\rm democrat}$

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democrat

421

republican

422

 ${\rm democrat}$

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424

 ${\rm democrat}$

429

democrat

democrat

431

republican

433

republican

summary(svmpredict)

democrat

182

republican

123

As noted above summary() code gives a short brief summary about sympredict variable.

Finally, we have to compare between the real data and the model that we set up by using SVM. When this model is compared with real data, the only column that needed to be taken is the first column because it has the classes of congressmen that are democrat and republican. Besides, we use table()code to make comparison between the real data and symmodel and we assign the SVM_table variable because we will not write the table data later.

```
SVM_table<-table(testdata$V1,svmpredict)</pre>
```

SVM_table

sympredict democrat republican 172 republican 10 116

3.ANALYSIS

democrat

a.Logistic Regression Model Process

When we formed the first logistic regression model we used

$$glm(V1 .., family = binomial(link = "logit"), data = modeldata)$$

code but it did not converge. For this reason we examined the data set and we tried to figure out the relationship between the class of congressmen and statues. Besides, we swithced the code to

$$glm(V1\ V4+V7+V9+V12+V15+V17, family = binomial(link = "logit"), data = model data)$$

then this code converges but still it is not a best form of our logistic regression model because we have many independent variables in this code. Therefore, we use

$$anova(logitmodel, test =' Chisq')$$

code to state which of columns are used or not. When the colums are determined, the table of anova function is researched. The residual deviance of anova table tells the dependency of the columns and the class of congressmen by using the difference between the residual deviance for each of columns. When the difference of residual deviance is greater than the other one's, this means that the column has a stronger relationship than the others. Hence these columns are choosed by using method that is mentioned. Later we constituted the second logistic regression model by using the columns that are choosed again. After the prediction function of logitmodel formed, constructing a table for logitmodel we took a threshold for the probability. Then we got a table for comparing the real data. Besides we compared which model is better by using Chi-Square method and the second model is determined. The reason of this detection is having less independent variables although the second logistic regression model has a small chi-square number for some cases.

b. SVM Model Process

To begin with we download a library for using SVM model. Then we constructed our SVM model simply because it did not get a warning for not converging. Moreover, we predicted the SVM model that we formed and we made a comparison between the prediction of this model and the real data by using table function. According to table data, there are 18 congressmen are known wrongly which means that there are congressmen that are democrat but they pretend republican or vise versa. It is significant because if the republican or democrat congressmen attract supporter for themselves then they will need to find this off diagonal terms.

c. Chi-Square Method

Chi-Square method is one of the tests that detemines the existence of the relationship between two variables that are numerical or not in statistic. This method makes a comparison between the observed values and the expected values theoritically. To begin with, there is a subtraction between observed values and expected values. Then this test squares the difference that it found and divided by expected values. The formula of Chi-Square stated in [5] is as the following

$$\chi_c^2 = \sum_{i=1}^k \frac{(O_i - E_i)^2}{E_i} \tag{3.1}$$

where the O_i is the observed values, E_i is the expected values and c is the degree of freedom. Chi- Square method tests whether there is a dependency in data set. Also if the calculated value of this method is bigger than the value of table data then there will be a relationship between the variables.

d. Chi-Square Method in R

When we use this method we use chisq.test(matrix()) code and we determine which model is statistically good according to the results of the chisq.test()code.

```
chisq.test(logistic_table2)
chisq.test(SVM_table)
```

Pearson's Chi-squared test with Yates' continuity correction

```
data: logistic_table2
X-squared = 167.14, df = 1, p-value < 2.2e-16</pre>
```

Pearson's Chi-squared test with Yates' continuity correction

```
data: SVM_table
X-squared = 235.14, df = 1, p-value < 2.2e-16</pre>
```

According to the results of these codes, if we compare of two model that we constituted the SVM model is better than the Logistic Regression Model because the X-squared value of SVM model is greater than Logistic Regression model and the SVM model is easier to form.

References

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- [5]: S.Deviant, The Practically Cheating Statistics Handbook, Retrieved from http://www.statisticshowto.com/probability-and-statistics/chi-square/