Statistical Computing with R: Masters in Data Sciences 503, S21 First Batch, SMS, TU, 2021

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Review Preview

- Simple Linear Regression
 - Gradient Decent fit
 - Model Accuracy
 - Model Prediction
 - Model Validation
 - Validation set, Leave one out cross validation, k-folds cross validation, repeated k-folds cross-validation etc.)

- Multiple Linear Regression
 - Simple linear regression +
 - Multicollinearity, its assessment and solutions
 - Regularization
 - Ridge
 - Lasso
 - Elastic Net (Ridge+Lasso)

Simple Linear Regression: Gradient Decent

https://towardsdatascience.com/linear-regression-using-gradient-descent-97a6c8700931

 Linear Regression can also be fitted with Gradient Decent algorithm instead of OLS

$$egin{align} D_m &= rac{1}{n} \sum_{i=0}^n 2(y_i - (mx_i + c))(-x_i) \ D_m &= rac{-2}{n} \sum_{i=0}^n x_i (y_i - ar{y}_i) \ \end{pmatrix}$$

 Here we minimize the loss function (E) to find m (slope b) and c (constant a)

$$E = rac{1}{n} \sum_{i=0}^n (y_i - (mx_i + c))^2$$

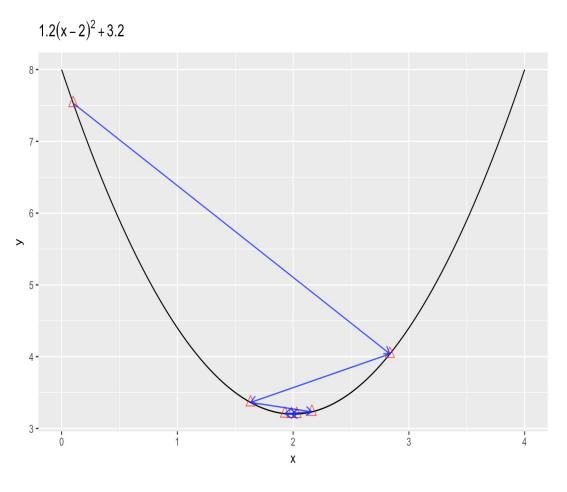
E = MSE = Mean Sum of Square

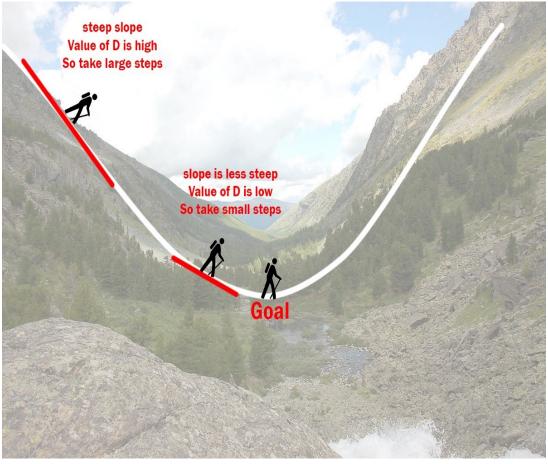
$$D_c = rac{-2}{n} \sum_{i=0}^n (y_i - ar{y}_i)$$

We can solve these two equations to get the value of m and c. This is an optimization problem as we need to MINIMIZE the loss function MSE using multiple iterations!

Illustration:

http://ethen8181.github.io/machine-learning/linear_regression/linear_regession.html





How to get MSE of Simple Linear Model? (OLS and Gradient Decent)

Using the residuals of linear model:
 lm1 <- lm(mpg ~ wt, data=mtcars)
 (mse <- mean(lm1\$residuals^2))

Saving predicted values:
 data <- data.frame(pred = predict(lm1), actual = mtcars\$mpg)
 head(data)
 mean((data\$actual - data\$pred)^2)

- > (mse <- mean(lm1\$residuals^2))
- [1] 8.697561

- > mean((data\$actual data\$pred)^2)
- [1] 8.697561

Model Accuracy of Linear Model:

- R-square Explained variance (higher is better!)
- RMSE Root of MSE (lower is better)
- MAE Mean Absolute Error (lower is better)
- MAPE Mean Absolute Percentage Error (lower is better)

$$MAPE = \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{100\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{1000\%}{n} \sum_{\text{Each residual is scaled against the actual value}} \frac{1000\%}{n} \sum_{\text{Each resi$$

#Better to use "caret" package

Install.packages("caret")

library(caret)

R2 <- R2(data\$pred, data\$actual)

• 0.7528328

RMSE <- RMSE(data\$pred, data\$actual)

• 2.949163

MAE <- MAE(data\$pred, data\$actual)

• 2.340642

Get MAPE in R as:

• 12.60733

If LINE is valid after BLUE then we can predict: (So, we will use "Im1<-Im(mpg~wt, data=mtcars)" model to do it!)

- We need to save independent variable value/values in a new data:
 new.wt <- data.frame(wt = 6)
- We can then use this data to predict the value of the dependent variable based on the fitted model as:

```
predict(lm1, newdata = new.wt)
```

Result = 5.218297

 Interpretation: Cars with 6000 lbs weight will (only) give 5.22 miles per gallon as per the linear regression algorithm!)

Validation & Cross-validation for Predictive Modelling including Linear Model:

- In statistics, we normally use the "full" data to do predictions
- In machine learning, we use validation/cross-validation sets to do the predictions
- Validation/Cross-validation can be done with:
 - Validation set (data split)
 - Leave one out cross-validation (LOOCV)
 - K-fold cross-validation
 - Repeated k-fold validation

Validation: Validation set (widely used!)

- Here the full data is "randomly" divided into two sets:
 - Training set
 - Testing set (validation set)
- Then model is fitted in the training set
- The model fit is then validated in the testing set using prediction
- This is the most widely used cross-validation method in the supervised machine learning

Lets do it for "mtcars" data:

test.data <- data[ind==2,]

```
#Define the mtcars data as "data":
data <- mtcars
#Use random seed to replicate the result
set.seed(1234)
#Do random sampling to divide the cases into two independent samples
ind <- sample(2, nrow(mtcars), replace = T, prob = c(0.7, 0.3))
#Data partition
train.data <- data[ind==1,]
```

Model Fit, Prediction and Cross-Validation: Validation set approach

```
lm4 <- lm(mpg~wt, data = train.data)</pre>
library(dplyr)
library(caret)
predictions <- Im4 %>%
predict(test.data)
data.frame(R2 = R2(predictions,
test.data$mpg),
      RMSE = RMSE(predictions,
      test.data$mpg),
      MAE = MAE(predictions,
      test.data$mpg))
```

Model Accuracy of Training dataset:

summary(lm4)

- Multiple R-squared: 0.7013
- MSE = 9.526359 (How?)
- RMSE = SQRT(MSE) = 3.08648

Model Accuracy of Testing dataset:

R2 RMSE MAE

0.90310852.2793031.698583

Model Fit, Prediction and Cross-Validation: Leave-One-Out Cross-Validation approach:

```
#Leave one out CV
library(caret)
# Define training control
train.control <- trainControl(method = "LOOCV")</pre>
# Train the model
model1 <- train(mpg ~wt, data = mtcars, method =
"lm",
        trControl = train.control)
# Summarize the results
print(model1)
```

Linear Regression

32 samples

1 predictor

No pre-processing

Resampling: Leave-One-Out Cross-Validation

Summary of sample sizes: 31, 31, 31, 31, 31, 31, ...

Resampling results:

RMSE Rsquared MAE

3.201673 0.7104641 2.517436

Tuning parameter 'intercept' was held constant at a value of TRUE

Prediction with LOOCV:

```
predictions1 <- model1 %>%
predict(test.data)

data.frame(R2 = R2(predictions1,
test.data$mpg),

    RMSE = RMSE(predictions1,
test.data$mpg),

    MAE = MAE(predictions1,
test.data$mpg))
```

R2 RMSE MAE

0.9031085 2.244232 1.714515

Model Fit, Prediction and Cross-Validation: K-folds Cross-Validation approach

```
#k-fold cross validation
library(caret)
# Define training control
set.seed(123)
train.control <- trainControl(method = "cv", number
= 10)
# Train the model
model2 <- train(mpg ~ wt, data = mtcars, method =
"lm",
        trControl = train.control)
# Summarize the results
print(model2)
```

```
Linear Regression
32 samples
1 predictor
```

No pre-processing

Resampling: Cross-Validated (10 fold)

Summary of sample sizes: 28, 28, 29, 29, 29, 30, ...

Resampling results:

RMSE Rsquared MAE

2.85133 0.7346939 2.375068

Tuning parameter 'intercept' was held constant at a value of TRUE

Predictions with k-folds CV:

```
predictions2 <- model2 %>%
predict(test.data)

data.frame(R2 = R2(predictions2,
test.data$mpg),

    RMSE = RMSE(predictions2,
test.data$mpg),

    MAE = MAE(predictions2,
test.data$mpg))
```

```
R2 RMSE MAE

• 0.9031085 2.244232 1.714515
```

Model Fit, Prediction and Cross-Validation: Repeated K-folds Cross-Validation approach

```
#repeated k-fold cross validation
library(caret)
# Define training control
set.seed(123)
train.control <- trainControl(method = "repeatedcv",
                 number = 10, repeats = 3)
# Train the model
model <- train(mpg ~wt, data = mtcars, method =
"lm",
        trControl = train.control)
# Summarize the results
print(model)
```

```
Linear Regression
32 samples
1 predictor
No pre-processing
Resampling: Cross-Validated (10 fold, repeated 3 times)
Summary of sample sizes: 28, 28, 29, 29, 29, 30, ...
Resampling results:
 RMSE
                     Rsquared
                                           MAE
 2.975392
                     0.8351572
                                           2.539797
Tuning parameter 'intercept' was held constant at a value of
TRUE
```

Prediction with repeated k-folds CV:

```
predictions3 <- model3 %>%
predict(test.data)
data.frame(R2 = R2(predictions3,
test.data$mpg),
      RMSE = RMSE(predictions3,
test.data$mpg),
      MAE = MAE(predictions3,
test.data$mpg))
```

R2 RMSE MAE

0.9031085 2.244232 1.714515

Summary: Which one should be used based on R-squared values of "Im" model?

• R-square for training set: 0.7013

• R-square for testing set: 0.9031085

- R-square for training with LOOCV:
 0.7104641
- R-square for testing with LOOCV: 0.9031085

- R-square for training with k-folds CV: 0.7346939
- R-square for testing with k-folds CV: 0.9031085

- R-square for training with repeated k-folds CV: 0.8351572
- R-square for testing with repeated k-folds CV: 0.9031085

Summary: Which one should be used based on RMSE value?

• RMSE for training set: 3.08648

• RMSE for testing test: 2.279303

• RMSE for training with LOOCV: 3.201673

RMSE for testing with LOOCV:
 2.244232

RMSE for training with k-folds CV:
 2.85133

• RMSE for testing with k-folds CV: 2.244232

 RMSE for training with repeated kfolds CV: 2.975392 RMSE for testing with repeated kfolds CV: 2.244232

Quick Think!

 Which model must be selected: Based on R-square or based on RMSE? If BLUE and LINE test is a must then which training model should be checked?

- Do we need to check the BLUE and LINE assumptions for the fit done with the training data?
- Validation set model?
- LOOCV set model?
- K-fold CV set model?
- Repeated K-fold CV set model?

Question/queries so far?

Multiple linear regression:

- It is an extension of the simple linear regression
- Multiple linear regression have more than one (two or more) independent variables
- Multiple linear regression has one

 (1) continuous dependent variable
 so it is a supervised learning

- All the assumptions of the simple linear regression are also applicable here
- There is one more condition:

 Multicollinearity must not be present i.e. correlations between independent variables must not be "high"

Assessing multicollinearity:

- Pearson correlation coefficients can be used
- Variance Inflation Factor (VIF) is most commonly used to assess multicollinearity
- We need to get a correlation matrix and flag the correlations with more than 0.75
- We can get the VIF for each independent variable

 These pair/s of independent variables influences the linear model coefficients Multicollinearity will be confirmed for an independent variable with VIF > 10 for linear models

Fitting multiple linear regression model using "mtcars" data:

- mlr <- lm(mpg ~., data = mtcars)
- summary(mlr)
- library(car)
- vif(mlr)
- We need to drop the independent variable with highest VIF and run the model again until all the VIF <10!

- None of the variables used in the model are statistically significant!
- > vif(mlr)

7.908747

cyl	disp	hp
15.373833	21.620241	9.832037
drat	wt	qsec
3.374620	15.164887	7.527958
VS	am	gear
4.965873	4.648487	5.357452
carb		

Fitting multiple linear regression using "mtcars" data:

#Removing "disp" variable:

• The "wt" variable is significant

mlr1 <- lm(mpg ~ cyl+hp+drat+wt+qsec+vs+am+gear+c arb, data = mtcars) summary(mlr1) vif(mlr1)

 We need to drop the independent variable with highest VIF and run the model again until all the VIF <10! • > vif(mlr1) hp drat cyl 14.284737 7.123361 3.329298 qsec wt VS 6.189050 6.914423 4.916053 carb gear am 4.645108 5.324402 4.310597

Fitting multiple linear regression using "mtcars" data:

#Removing "cyl" variable:

```
mlr2 <- lm(mpg ~
hp+drat+wt+qsec+vs+am+gear+carb,
data = mtcars)
summary(mlr1)
vif(mlr1)
```

- We need to drop the independent variable with highest VIF and run the model again until VIF <10!
- If all the VIF < 10 then we can interpret the model and do the predictions

• The "wt" variable is significant, b = -2.60968 (1 unit increase in wt reduces the mpg by 2.61 unit controlling for other independent variables)

• > vif(mlr2)

hp	drat	wt
6.015788	3.111501	6.051127
qsec	VS	am
5.918682	4.270956	4.285815
gear carb		
4.690187	4.290468	

Assignment:

- Use the validation and crossvalidation methods for the multiple linear regression (mlr2) model
- Which model is the best model?
- Why?
- Predict the weight of the cars based on the best model identified using the test.data

- Change all the variables (except mpg) as standardized variable using "scale" command in R/R Studio
- Fit the multiple linear regression model with these standardized variables
- Does it solve the multicollinearity issue?
- Why? Write conclusions.

Alternative way to deal with multicollinearity in data science/machine learning:

 We can use the "regularization" methods

- The most common ones are:
 - Ridge regression
 - Lasso regression
 - Elastic net regression

 Once the "multicollinearity" problem is fixed then we can do the predictions and use the validation indices to select the best model for our data!

• I will post a YouTube video link and you can fit these models for the "mtcars" data and learn from them!

Question/queries?

Next class

- Other regression models used in the supervised learning: polynomial regression, KNN algorithm etc.
- Logistic regression
- Other classification models used in the supervised learning: Naïve Bayes, Decision Trees, SVM etc.

Thank you!

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