lab1-asn2

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Assignment 2. Linear regression and ridge regression

1

```
rawdata <- read.csv("parkinsons.csv")</pre>
library(dplyr)
##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
##
       filter, lag
## The following objects are masked from 'package:base':
##
##
       intersect, setdiff, setequal, union
rawdata <- rawdata %>% select(-c(1:4,6))
n <- nrow(rawdata)</pre>
set.seed(12345)
id <- sample(1:n,floor(n*0.6))</pre>
train <- rawdata[id,]</pre>
test <- rawdata[-id,]</pre>
library(caret)
## Loading required package: ggplot2
## Loading required package: lattice
param <- preProcess(train)</pre>
train_scaled <- predict(param,train)</pre>
test_scaled <- predict(param,test)</pre>
```

 $\mathbf{2}$

```
mse <- function(true_value, predict_value){</pre>
  mean((true_value - predict_value)^2)
}
model <- lm(motor_UPDRS ~ ., data = train_scaled)</pre>
train_mse <- mse(train_scaled$motor_UPDRS, predict(model, train_scaled))</pre>
cat(paste0("Training MSE is: ",train_mse,".\n"))
## Training MSE is: 0.878543102826275.
test_mse <- mse(test_scaled$motor_UPDRS, predict(model, test_scaled))</pre>
cat(paste0("Test MSE is: ",test_mse,".\n"))
## Test MSE is: 0.935447712156712.
summary(model)
##
## Call:
## lm(formula = motor_UPDRS ~ ., data = train_scaled)
## Residuals:
##
      Min
               1Q Median
                               3Q
                                      Max
## -3.0255 -0.7363 -0.1087 0.7333 2.1960
##
## Coefficients:
##
                  Estimate Std. Error t value Pr(>|t|)
                 6.575e-15 1.583e-02 0.000 1.000000
## (Intercept)
## Jitter...
                 1.869e-01 1.496e-01
                                       1.250 0.211496
## Jitter.Abs.
                -1.696e-01 4.081e-02 -4.156 3.32e-05 ***
## Jitter.RAP
                -5.270e+00 1.884e+01 -0.280 0.779688
## Jitter.PPQ5
                -7.457e-02 8.778e-02 -0.850 0.395659
## Jitter.DDP
               5.250e+00 1.884e+01 0.279 0.780541
## Shimmer
                5.924e-01 2.060e-01 2.876 0.004055 **
## Shimmer.dB.
                -1.727e-01 1.393e-01 -1.239 0.215380
## Shimmer.APQ3 3.207e+01 7.717e+01 0.416 0.677738
## Shimmer.APQ5 -3.875e-01 1.138e-01 -3.405 0.000669 ***
## Shimmer.APQ11 3.055e-01 6.124e-02 4.989 6.37e-07 ***
                -3.239e+01 7.717e+01 -0.420 0.674739
## Shimmer.DDA
                -1.854e-01 4.557e-02 -4.068 4.85e-05 ***
## NHR
## HNR
                -2.385e-01 3.640e-02 -6.553 6.45e-11 ***
## RPDE
                                       0.179 0.857576
                 4.068e-03 2.267e-02
## DFA
                -2.803e-01 2.014e-02 -13.919 < 2e-16 ***
## PPE
                 2.265e-01 3.289e-02
                                       6.886 6.75e-12 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.9396 on 3508 degrees of freedom
## Multiple R-squared: 0.1212, Adjusted R-squared: 0.1172
## F-statistic: 30.24 on 16 and 3508 DF, p-value: < 2.2e-16
```

The significant contributors are Jitter. Abs., Shimmer. APQ5, Shimmer. APQ11, NHR, HNR, DFA, and PPE, which are marked *** in summary output.

3

```
Loglikelihood <- function(theta_vec, sigma){</pre>
  y <- train_scaled[,1]</pre>
  x <- as.matrix(train_scaled[,-1])</pre>
  n <- nrow(train_scaled)</pre>
  value <-(n/2*\log(sigma^2*2*pi)+sum((y-x%*%theta_vec)^2)/(2*sigma^2))
  return(value)
}
Ridge <- function(theta_vec, sigma, lambda){</pre>
  -Loglikelihood(theta_vec, sigma)+lambda*sum(theta_vec^2)
RidgeOpt <- function(lambda){</pre>
  init_theta <- rep(0, ncol(train_scaled)-1)</pre>
  init_sigma <- 0.9</pre>
  objective_function <- function(params) {</pre>
    theta_vec <- params[-length(params)]</pre>
    sigma <- params[length(params)]</pre>
    return(Ridge(theta_vec, sigma, lambda))
  }
  rlt <- optim(c(init_theta, init_sigma), fn = objective_function, method = "BFGS")</pre>
  optimal_theta_vec <- rlt$par[-length(rlt$par)]</pre>
  optimal_sigma <- rlt$par[length(rlt$par)]</pre>
  return(list(theta_vec = optimal_theta_vec, sigma = optimal_sigma))
}
DF <- function(lambda){</pre>
  x <- as.matrix(train_scaled[,-1])</pre>
  df \leftarrow sum(diag(x%*%solve(t(x)%*%x+lambda*diag(ncol(x)))%*%t(x)))
  return(df)
}
```

4

```
train_x <- as.matrix(train_scaled[,-1])
train_y <- train_scaled[,1]
test_x <- as.matrix(test_scaled[,-1])
test_y <- test_scaled[,1]</pre>
```

```
Lambda <- c(1,100,1000)
store_list <- list()
```

```
for (i in 1:length(Lambda)){
   store_list$lambda[i] <- Lambda[i]
   optimal_theta_vec <- RidgeOpt(Lambda[i])$theta_vec
   store_list$train_MSE[i] <- mse(train_y, train_x%*%optimal_theta_vec)
   store_list$test_MSE[i] <- mse(test_y, test_x%*%optimal_theta_vec)
   store_list$DoF[i] <- DF(Lambda[i])
   store_list$sigma[i] <- RidgeOpt(Lambda[i])$sigma
}
store_df <- as.data.frame(store_list)
print(store_df)</pre>
```

As the table shown, train_MSE increases slightly as λ increases, while test_MSE reaches minimum at $\lambda = 100$, which is regarded as the optimal penalty parameter among these three.

Larger λ results to smaller DoF. Therefore, effects of DoF on models can be viewed as the opposite of effects of λ on them. When λ increases (DoF decreases), model becomes simpler; when λ decreases (DoF increases), model becomes more complex. Since there is a trade-off of model complexity for the best model, there exists optimal values of λ and DoF.

```
# Just for showing the inversely proportional relationship between lambda and DoF.
lam <- seq(0,1000,by=50)
dof <- c()
for (i in 1:length(lam)){
   dof[i] <- DF(lam[i])
}
plot(lam,dof)</pre>
```

