STATS 790 Assignment 2

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The multivariate linear regression model is: $Y = \beta_0 + \beta_1 X_1 + \dots + \beta_k X_k + \epsilon$ In matrix format, with n observations and k predictors, $\vec{Y} = \mathbf{X} \vec{\beta} + \vec{\epsilon}$, where

$$\vec{Y} = \begin{pmatrix} Y_1 \\ \vdots \\ Y_n \end{pmatrix}$$

design matrix \mathbf{X}

$$\mathbf{X} = \begin{pmatrix} 1 & X_{11} & \cdots & X_{1k} \\ \vdots & \vdots & \ddots & \vdots \\ 1 & X_{n1} & \cdots & X_{nk} \end{pmatrix}$$

$$\vec{\beta} = \begin{pmatrix} \beta_0 \\ \vdots \\ \beta_k \end{pmatrix}$$

$$\vec{\epsilon} = \begin{pmatrix} \epsilon_1 \\ \vdots \\ \epsilon_n \end{pmatrix}$$

Compute $\vec{\beta}$ using naive linear algebra

using least squares estimation of $\vec{\beta}$:

$$L = \sum_{i=1}^n \epsilon_i^2 = \vec{\epsilon}^\top \vec{\epsilon} = (\vec{Y} - \mathbf{X} \vec{\beta})^\top (\vec{Y} - \mathbf{X} \vec{\beta})$$

To minimize L, we differentiate and set it to 0:

$$\frac{\partial (\vec{Y} - \mathbf{X}\vec{\beta})^{\top} (\vec{Y} - \mathbf{X}\vec{\beta})}{\partial \vec{Y}} = -2\mathbf{X}^{\top}\vec{Y} + 2\mathbf{X}^{\top}\mathbf{X}\vec{\beta}$$
$$= 0$$

Then we get the normal equation:

$$\Rightarrow \mathbf{X}^{\top} \mathbf{X} \vec{\beta} = \mathbf{X}^{\top} \vec{Y}$$

And thus,

$$\hat{\vec{\beta}} = (\mathbf{X}^{\top}\mathbf{X})^{-1}\mathbf{X}^{\top}\vec{Y}$$

Compute $\vec{\beta}$ using QR decomposition

In QR decomposition, \mathbf{X} can be decomposed as $\mathbf{X} = \mathbf{Q}\mathbf{R}$, where \mathbf{Q} is a nx(k+1) orthonormal matrix and \mathbf{R} is a (k+1)x(k+1) upper triangular matrix. By the property of orthogonality, $\mathbf{Q}^{\top}\mathbf{Q} = \mathbf{I}$, then using the normal equation, we have

$$\begin{split} \mathbf{X}^{\top}\mathbf{X}\vec{\beta} &= \mathbf{X}^{\top}\vec{Y} \\ \mathbf{Q}\mathbf{R}^{\top}\mathbf{Q}\mathbf{R}\vec{\beta} &= \mathbf{Q}\mathbf{R}^{\top}\vec{Y} \\ \mathbf{R}^{\top}\mathbf{Q}^{\top}\mathbf{Q}\mathbf{R}\vec{\beta} &= \mathbf{R}^{\top}\mathbf{Q}^{\top}\vec{Y} \\ \mathbf{R}^{\top}\mathbf{R}\vec{\beta} &= \mathbf{R}^{\top}\mathbf{Q}^{\top}\vec{Y} \\ \mathbf{R}\vec{\beta} &= \mathbf{Q}^{\top}\vec{Y} \end{split}$$

Thus,

$$\hat{\vec{\beta}} = \mathbf{R}^{-1} \mathbf{Q}^{\top} \vec{Y}$$

Compute $\vec{\beta}$ using SVD

In SVD, \mathbf{X} can be decomposed as $\mathbf{X} = \mathbf{U}\mathbf{S}\mathbf{V}^{\top}$, where \mathbf{U} is a nxn matrix, \mathbf{S} is a nx(k+1) matrix and \mathbf{V} is a (k+1)x(k+1) matrix. Both \mathbf{U} and \mathbf{V} are orthogonal, and so $\mathbf{U}^{\top}\mathbf{U} = \mathbf{V}^{\top}\mathbf{V} = \mathbf{I}$.

$$\mathbf{X}^{\top}\mathbf{X}\vec{\beta} = \mathbf{X}^{\top}\vec{Y}$$
$$(\mathbf{U}\mathbf{S}\mathbf{V}^{\top})^{\top}(\mathbf{U}\mathbf{S}\mathbf{V}^{\top})\vec{\beta} = (\mathbf{U}\mathbf{S}\mathbf{V}^{\top})^{\top}\vec{Y}$$
$$\mathbf{V}\mathbf{S}^{\top}\mathbf{U}^{\top}\mathbf{U}\mathbf{S}\mathbf{V}^{\top}\vec{\beta} = \mathbf{V}\mathbf{S}^{\top}\mathbf{U}^{\top}\vec{Y}$$
$$\mathbf{V}\mathbf{S}^{\top}\mathbf{S}\mathbf{V}^{\top}\vec{\beta} = \mathbf{V}\mathbf{S}^{\top}\mathbf{U}^{\top}\vec{Y}$$
$$\mathbf{S}\mathbf{V}^{\top}\vec{\beta} = \mathbf{U}^{\top}\vec{Y}$$

Thus,

$$\hat{\vec{\beta}} = \mathbf{V}\mathbf{S}^{-1}\mathbf{U}^{\top}\vec{Y}$$

Compute $\vec{\beta}$ using Cholesky decomposition

In Cholesky decomposition, if $\mathbf{X}^{\top}\mathbf{X}$ is positive definite matrix, it can be decomposed as $\mathbf{X}^{\top}\mathbf{X} = \mathbf{L}\mathbf{L}^{\top}$, where \mathbf{L} is a lower triangular matrix and \mathbf{L}^{\top} is the conjugate transpose of \mathbf{L} .

Using the normal equation,

$$\mathbf{L}\mathbf{L}^{\top}\vec{\beta} = \mathbf{X}^{\top}\vec{Y}$$

Let $\mathbf{L}^{\top}\vec{\beta} = Z$ and $\mathbf{L}Z = \mathbf{X}^{\top}\vec{Y}$. Then we can solve for Z since \mathbf{L} , \mathbf{X}^{\top} , \vec{Y} are known. After solving Z, we can solve for $\vec{\beta}$ using $\mathbf{L}^{\top}\vec{\beta} = Z$, where \mathbf{L}^{\top} and Z are known.

So,

$$\hat{\vec{\beta}} = (\mathbf{L}^\top)^{-1} Z$$

Compare running time of Naive Linear Algebra, QR Decomposition and SVD

library(microbenchmark)
library(ggplot2)
library(dplyr)
library(pracma)

```
#simulate data
test_matrix <- function(n,p){</pre>
  X <- matrix(rnorm(n*p), n, p)</pre>
  y <- X %*% rnorm(p) + rnorm(p)
  list(X = X, y = y)
}
set.seed(1)
matrix1 <- test_matrix(1000,100)</pre>
# simply lm()
test_model <- function(X,y){</pre>
  b <- data.matrix(lm(y ~ X+0)$coef)</pre>
  b <- as.data.frame(b, row.names = 1:nrow(b))</pre>
  b
}
#naive linear algebra
linear_alg <- function(X,y){</pre>
  b <- data.matrix(solve(t(X) %*% X, t(X) %*% y))</pre>
  b <- as.data.frame(b, row.names = 1:nrow(b))</pre>
}
#QR decomposition
qr_decomp <- function(X,y){</pre>
  q_r \leftarrow qr(X)
  b <- data.matrix(backsolve(qr.R(q_r), t(qr.Q(q_r)) %*% y))</pre>
  b <- as.data.frame(b, row.names = 1:nrow(b))</pre>
  b
}
```

```
#SVD
sv_decomp <- function(X,y){</pre>
  s_v <- svd(X)
  b <- data.matrix(solve(diag(s_v$d) %*% t(s_v$v), t(s_v$u) %*% y))</pre>
 b <- as.data.frame(b, row.names = 1:nrow(b))</pre>
  b
}
#check
stopifnot(all.equal(test_model(matrix1$X,matrix1$y),
                     linear_alg(matrix1$X,matrix1$y)))
stopifnot(all.equal(test_model(matrix1$X,matrix1$y),
                     qr_decomp(matrix1$X,matrix1$y)))
stopifnot(all.equal(test_model(matrix1$X,matrix1$y),
                     sv_decomp(matrix1$X,matrix1$y)))
# benchmarking
bm <- microbenchmark(test_model(matrix1$X,matrix1$y)</pre>
                     , linear_alg(matrix1$X,matrix1$y)
                     , qr_decomp(matrix1$X,matrix1$y)
                     , sv_decomp(matrix1$X,matrix1$y))
result1 <- summary(bm)</pre>
#try more n
nvec \leftarrow round(10^seq(2, 5, by = 0.5))
results <- data.frame()
for (i in seq_along(nvec)) {
    s <- test_matrix(nvec[i], 100)
    mbm <- microbenchmark(test_model(matrix1$X,matrix1$y)</pre>
                     , linear_alg(matrix1$X,matrix1$y)
```

Plot

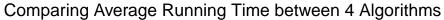
```
#compare using log-log plot

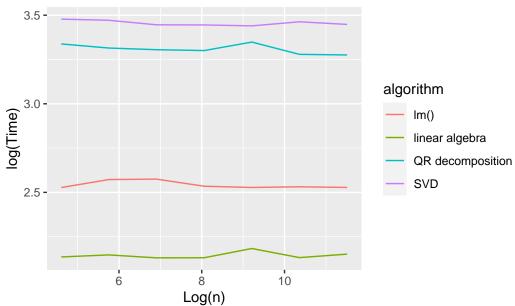
ggplot(data = results_new, aes(x=lognvec, y=logmean, group=expr)) +

geom_line(aes(color=expr)) +

labs(color='algorithm', x="Log(n)", y="log(Time)") +

ggtitle("Comparing Average Running Time between 4 Algorithms")
```





The plot shows that native linear algebra runs fastest among the four algorithms, even faster than the native implementation of linear regression. In contrast, the SVD algorithm runs the slowest of the three other algorithms.

Model

```
#fit log-log model
logmodel <- lm(logmean~lognvec, data = results_new)
summary(logmodel)</pre>
```

Call:

lm(formula = logmean ~ lognvec, data = results_new)

Residuals:

Coefficients:

Estimate Std. Error t value Pr(>|t|)

(Intercept) 2.886811 0.386208 7.475 6.17e-08 ***
lognvec -0.002978 0.046079 -0.065 0.949

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

Residual standard error: 0.5615 on 26 degrees of freedom

Multiple R-squared: 0.0001606, Adjusted R-squared: -0.03829

F-statistic: 0.004177 on 1 and 26 DF, $\,$ p-value: 0.949 $\,$

The log-log model is log(mean) = 2.9133 - 0.0019*log(n)

```
library(glmnet)
  #read table
  prostate <- read.delim("prostate.data", row.names=1)</pre>
  #split to training and test sets
  prostate_train <- filter(prostate, train == "TRUE")</pre>
  prostate_test <- filter(prostate, train == "FALSE")</pre>
  #remove last column "train"
  prostate_train <- prostate_train[1:9]</pre>
  prostate train matrix <- as.matrix(subset(prostate train, select = -lpsa))</pre>
  prostate_test_x <- as.matrix(subset(prostate_test, select = -c(train, lpsa)))</pre>
  prostate_test1 <- subset(prostate_test, select = -train)</pre>
  #native implementation of ridge regression
  ridge <- function(x,y){</pre>
    set.seed(1)
    cv <- cv.glmnet(x, y, alpha = 0, standardize=TRUE)</pre>
    lambda <- cv$lambda.min</pre>
    rid <- glmnet(x, y, alpha = 0, standardize=TRUE, lambda = lambda)</pre>
    list("model" = rid, "opt_lambda" = lambda)
  }
  RR <- ridge(prostate_train_matrix,prostate_train$lpsa)</pre>
  RR
$model
Call: glmnet(x = x, y = y, alpha = 0, lambda = lambda, standardize = TRUE)
```

```
Df %Dev Lambda
1 8 68.87 0.08789
$opt_lambda
[1] 0.08788804
  # ridge regression by data augmentation
  # augment data by adding gaussian noise
  ridge_aug <- function(x){</pre>
    set.seed(1)
    len <- dim(x)[1] * dim(x)[2]
    noise <- matrix(rnorm(len, mean=0, sd = 1), dim(x)[1])</pre>
    aug <- as.matrix(x) + noise</pre>
    aug <- rbind(aug, as.matrix(x))</pre>
    aug <- as.data.frame(aug)</pre>
    osl <- lm(lpsa~., data = aug)</pre>
  }
  RR_aug <- ridge_aug(prostate_train)</pre>
  RR_aug
Call:
lm(formula = lpsa ~ ., data = aug)
Coefficients:
(Intercept)
                   lcavol
                                lweight
                                                                lbph
                                                                               svi
                                                  age
  -0.589414
                 0.431696
                               0.123882
                                             0.009317
                                                           0.186757
                                                                         0.354539
        lcp
                  gleason
                                  pgg45
   0.141671
                 0.205649
                              -0.001791
```

```
#R^2 from ridge regression by data augmentation
  pred <- predict(RR_aug, newdata = prostate_test[,-9])</pre>
  sst <- sum((prostate_test$lpsa - mean(prostate_test$lpsa))^2)</pre>
  ssr <- sum((prostate_test$lpsa - pred)^2)</pre>
  rsq <- 1 - ssr/sst
  rsq
[1] 0.514647
  #R^2 from ridge regression
  RRpred <- predict(RR$model, s=RR$opt_lambda, newx = prostate_test_x)</pre>
  RRsst <- sum((prostate_test$lpsa - mean(prostate_test$lpsa))^2)</pre>
  RRssr <- sum((prostate test$1psa - RRpred)^2)</pre>
  RRrsq <- 1 - RRssr/RRsst
  RRrsq
[1] 0.5289966
  set.seed(1)
  bm_ridge <- microbenchmark(ridge(prostate_train_matrix,prostate_train$lpsa)</pre>
                        , ridge_aug(prostate_train))
  result_ridge <- summary(bm_ridge)</pre>
  data.frame(result_ridge$expr, result_ridge$mean)
                                    result_ridge.expr result_ridge.mean
```

The R^2 of ridge regression without data augmentation is 52.90%, while the R^2 of ridge regression by data augmentation is 51.46%. This indicates that the model using ridge regression without data augmentation explains more variations compared to the model using ridge regression with data augmentation. However, the algorithm of ridge regression by data augmentation is much faster than the native ridge regression.

ridge_aug(prostate_train)

53.168495

1.454887

1 ridge(prostate_train_matrix, prostate_train\$lpsa)

2

By ESL Equation 3.44, we know that

$$\hat{\beta}^{ridge} = (\mathbf{X}^{\top}\mathbf{X} + \lambda \mathbf{I})^{-1}\mathbf{X}^{\top}\mathbf{y}$$

By the definition of posterior distribution,

$$p(\beta|\mathbf{y}) \propto p(\mathbf{y}|\beta)p(\beta)$$

where $p(\mathbf{y}|\beta)$ is the likelihood, with $\mathbf{y} \sim N(\mathbf{X}\beta, \sigma^2 \mathbf{I})$ and $p(\beta)$ is the prior distribution with $\beta \sim N(0, \tau \mathbf{I})$. Since the product of two gaussians is also a gaussian, the mean of the posterior distribution is the same value as it's mode.

Let β be the mode and thus the negative log-posterior density of β is

$$\begin{split} \hat{\beta} &= \underset{\beta}{argmax} & log(p(\mathbf{y}|\beta)) + log(p(\beta)) \\ &= \underset{\beta}{argmin} & C + \frac{(\mathbf{y} - \mathbf{X}\beta)^{\top}(\mathbf{y} - \mathbf{X}\beta)}{2\sigma^2} + \frac{\beta^{\top}\beta}{2\tau} \\ &= \underset{\beta}{argmin} & (\mathbf{y} - \mathbf{X}\beta)^{\top}(\mathbf{y} - \mathbf{X}\beta) + \frac{\sigma^2}{\tau}\beta^{\top}\beta \end{split}$$

By ESL Equation 3.43, both equation is equavelent when $\lambda = \frac{\sigma^2}{\tau}$, so

$$\hat{\beta} = (\mathbf{X}^{\top}\mathbf{X} + \frac{\sigma^2}{\tau}\mathbf{I})^{-1}\mathbf{X}^{\top}\mathbf{y}$$

 $\hat{\beta}$ is the estimated mean and mode of posterior distribution, and thus, $\hat{\beta} = \hat{\beta}^{ridge}$ when $\lambda = \frac{\sigma^2}{\tau}$.

Therefore, the ridge regression estimate is the mean and mode of the posterior distribution, under a Gaussian prior $\beta \sim N(0, \tau \mathbf{I})$, and Gaussian sampling model $\mathbf{y} \sim N(\mathbf{X}\beta, \sigma^2 \mathbf{I})$, when $\lambda = \frac{\sigma^2}{\tau}$.

Using SVD, let $\mathbf{X} = \mathbf{U}\mathbf{D}\mathbf{V}^{\top}$, by ESL Equation 3.47,

$$\hat{\beta}^{ridge} = \mathbf{U}\mathbf{D}(\mathbf{D}^2 + \lambda \mathbf{I})^{-1}\mathbf{D}\mathbf{U}^{\top}\mathbf{y} = \sum_{i=1}^{p} u_j \frac{d_j^2}{d_j^2 + \lambda} u_j^{\top}\mathbf{y}$$

where u_j are the column of ${\bf U}$ and d_j are the diagonal entries of diagonal matrix ${\bf D}$. So, $\hat{\beta}^{ridge}$ is

$$\begin{split} \hat{\beta}^{ridge} &= \mathbf{X}^{-1} \mathbf{U} \mathbf{D} (\mathbf{D}^2 + \lambda \mathbf{I})^{-1} \mathbf{D} \mathbf{U}^{\top} \mathbf{y} \\ &= \mathbf{V} \mathbf{D}^{-1} \mathbf{U}^{-1} \mathbf{U} \mathbf{D} (\mathbf{D}^2 + \lambda \mathbf{I})^{-1} \mathbf{D} \mathbf{U}^{\top} \mathbf{y} \\ &= \mathbf{V} (\mathbf{D}^2 + \lambda \mathbf{I})^{-1} \mathbf{D} \mathbf{U}^{\top} \mathbf{y} \end{split}$$

Thus,

$$\begin{split} \| \hat{\beta}^{ridge} \| &= (\mathbf{V} (\mathbf{D}^2 + \lambda \mathbf{I})^{-1} \mathbf{D} \mathbf{U}^{\top} \mathbf{y})^{\top} \mathbf{V} (\mathbf{D}^2 + \lambda \mathbf{I})^{-1} \mathbf{D} \mathbf{U}^{\top} \mathbf{y} \\ &= \mathbf{y}^{\top} \mathbf{U} \mathbf{D} (\mathbf{D}^2 + \lambda \mathbf{I})^{-1} \mathbf{V}^{\top} \mathbf{V} (\mathbf{D}^2 + \lambda \mathbf{I})^{-1} \mathbf{D} \mathbf{U}^{\top} \mathbf{y} \\ &= \mathbf{y}^{\top} \mathbf{U} \mathbf{D} (\mathbf{D}^2 + \lambda \mathbf{I})^{-2} \mathbf{U}^{\top} \mathbf{y} \\ &= \sum_{j=1}^{p} \mathbf{y} u_j \frac{d_j^2}{(d_j^2 + \lambda)^2} u_j^{\top} \mathbf{y} \end{split}$$

So, as $\lambda \to 0$, $\frac{d_j^2}{(d_j^2 + \lambda)^2}$ increases, and so $\|\hat{\beta}^{ridge}\|$ increases.

Also, by ESL 3.41 and 4.42 the penality term is $\lambda \sum_{i=1}^{p} \beta_{j}^{2}$. To keep the formula constant, $\sum_{i=1}^{p} \beta_{j}^{2}$ should increase as $\lambda \to 0$. So the value of t increases as λ decreases.

In lasso, the penality is replaced by $\sum_{i=1}^{p} |\beta_j|$, but the relationship between t and λ still holds. So in lasso, as $\lambda \to 0$, $\|\hat{\beta}^{lasso}\|$ increases, as well.

By ESL Equation 3.51, we have

$$\begin{split} \hat{\beta}^{lasso} &= argmin \quad \sum_{i=1}^{N} (y_i - \beta_0 - \sum_{m=1}^{p} x_{im} \beta_m)^2 \\ &\text{subject to} \quad \sum_{m=1}^{p} |\beta_m| \leq t \end{split}$$

Then suppose we augment \mathbf{X} with an identical copy $X_j^* = X_j$. Let $\beta_j = \tilde{\beta}_j + \tilde{\beta}_j^*$, where $\tilde{\beta}_j$ is the coefficient of \mathbf{X}_j and $\tilde{\beta}_j^*$ is the coefficient of \mathbf{X}_j^* in new problem.

The new lasso problem becomes

$$\begin{split} \hat{\beta}^{lassoNew} &= argmin \quad \sum_{i=1}^{N} (y_i - \beta_0 - \sum_{m \neq j}^{p} x_{im} \beta_m - x_{ij} \beta_j)^2 \\ &\text{subject to} \quad \sum_{m \neq j}^{p} |\beta_m| + |\beta_j| \leq t \end{split}$$

The penalty term has been adjusted to include the penalty on β_j . To let this becomes the original lasso problem, we need to find the optimal values of $\tilde{\beta}_j$ and $\tilde{\beta}_j^*$ so that

$$\sum_{m\neq j}^p |\beta_m| + |\beta_j| = \sum_{m\neq j}^p |\beta_m| + |\tilde{\beta}_j| + |\tilde{\beta}_j^*| \le t$$

By the property of absolute value, $|\beta_j| = |\tilde{\beta}_j + \tilde{\beta}_j^*| \le |\tilde{\beta}_j| + |\tilde{\beta}_j^*|$.

So only when $\tilde{\beta}_j$ and $\tilde{\beta}_j^*$ have the same sign i.e. $\tilde{\beta}_j \tilde{\beta}_j^* > 0$, the condition holds.

Since the value of t remains the same, $\beta_j=\hat{\beta}_j=\tilde{\beta}_j+\tilde{\beta}_j^*=a$

Therefore, when the exact collinearity happens, the sum of coefficients of two identical predictors will be the original coefficient of one predictor. Moreover, the coefficients of two identical predictors must have the same sign.

Let augment the matrix X with p additional rows and augment y with p zeros. Then

$$ilde{\mathbf{X}} = egin{pmatrix} \mathbf{X} \ \gamma \mathbf{I}_p \end{pmatrix}$$

$$ilde{\mathbf{y}} = egin{pmatrix} \mathbf{y} \\ 0_p \end{pmatrix}$$

Then the residual sum of squares is

$$\begin{split} RSS(\tilde{\beta}) &= \|\tilde{\mathbf{y}} - \tilde{\mathbf{X}}\beta\|_2^2 \\ &= \|\begin{pmatrix} \mathbf{y} - \mathbf{X}\beta \\ \gamma\beta \end{pmatrix}\|_2^2 \\ &= \|\mathbf{y} - \mathbf{X}\beta\|_2^2 + \gamma^2 \|\beta\|_2^2 \end{split}$$

Then turn this into a lasso problem, we add the penalty term and minimize it,

$$\Rightarrow \mathop{argmin}_{\beta} \quad \|\mathbf{y} - \mathbf{X}\beta\|_2^2 + \gamma^2 \|\beta\|_2^2 + \tilde{\lambda} \|\beta\|_1$$

Thus, when $\gamma^2 = \lambda \alpha$ and $\tilde{\lambda} = \lambda (1 - \alpha)$, the elastic-net optimization problem

$$\underset{\boldsymbol{\beta}}{argmin} \quad \|\mathbf{y} - \mathbf{X}\boldsymbol{\beta}\|_2^2 + \lambda[\boldsymbol{\alpha}\|\boldsymbol{\beta}\|_2^2 + (1-\boldsymbol{\alpha})\|\boldsymbol{\beta}\|_1]$$

turns into a lasso problem.

Reference

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