Assignment 1

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DON'T FORGET TO UNCOMMENT Q4

Question 1

From our definitions of \tilde{X} and \tilde{Y}

$$\tilde{X} = X_{-1} - \mathbf{1}_n \bar{x}^T$$

$$\tilde{Y} = Y - \mathbf{1}_n^T \bar{Y},$$

we find

$$\begin{split} \hat{\beta}_{-1} &= \underset{\beta \in \mathbb{R}^{p-1}}{\min} \ \|\tilde{Y} - \tilde{X}\beta\|_{2}^{2} \\ &= \underset{\beta \in \mathbb{R}^{p-1}}{\min} \ \|Y - \mathbf{1}_{n}\bar{Y} - \left(X_{-1} - \mathbf{1}_{n}\bar{x}^{T}\right)\beta_{-1}\|_{2}^{2} \\ &= \underset{\beta \in \mathbb{R}^{p-1}}{\arg\min} \ \|Y - X_{-1}\beta_{-1} - \mathbf{1}_{n}\left(\bar{Y} - \bar{x}^{T}\beta_{-1}\right)\|_{2}^{2} \\ &= \underset{\beta \in \mathbb{R}^{p-1}}{\arg\min} \ \|Y - X_{-1}\beta_{-1} - \mathbf{1}_{n}\beta_{1}\|_{2}^{2} \quad \text{(by definition of } \beta_{1} \text{ above)} \\ &= \underset{\beta \in \mathbb{R}^{p-1}}{\arg\min} \ \|Y - \left[\mathbf{1}_{n}, X_{-1}\right] \left[\beta_{1}, \beta_{-1}\right]\|_{2}^{2} \\ &\equiv \underset{\beta \in \mathbb{R}^{p-1}}{\arg\min} \ \|Y - X\beta\|_{2}^{2}. \end{split}$$

Therefore, if $\hat{\beta} = \left(\hat{\beta}_1,\,\hat{\beta}_{-1}^T\right)^T \in \mathbb{R}^p$ and

$$\hat{\beta}_1 = \bar{Y} - \bar{x}^T \hat{\beta}_{-1},$$

then $\hat{\beta}$ also solves the uncentered problem

$$\hat{\beta} \equiv \left(\hat{\beta}_1, \, \hat{\beta}_{-1}^T\right)^T = \underset{\beta \in \mathbb{R}^p}{\operatorname{arg min}} \, \|Y - X\beta\|_2^2,$$

as desired.

Question 2

(a)

Define our objective function $f: \mathbb{R}^p \to \mathbb{R}$ by

$$\begin{split} f(\beta) &= \|\tilde{Y} - \tilde{X}\beta\|_2^2 + \lambda \|\beta\|_2^2 \\ &= \left(\tilde{Y} - \tilde{X}\beta\right)^T \left(\tilde{Y} - \tilde{X}\beta\right)^T + \lambda \beta^T \beta \\ &= \tilde{Y}^T \tilde{Y} - \tilde{Y}^T \tilde{X}\beta - \beta^T \tilde{X}^T \tilde{Y} + \beta^T \tilde{X}^T \tilde{X}\beta + \lambda \beta^T \beta \\ &= \tilde{Y}^T \tilde{Y} - 2\beta^T \tilde{X}^T \tilde{Y} + \beta^T \tilde{X}^T \tilde{X}\beta + \lambda \beta^T \beta. \end{split}$$

Therefore, by taking the gradient we find

$$\nabla f(\beta) = -2\tilde{X}^T \tilde{Y} + 2\tilde{X}^T \tilde{X}\beta + 2\lambda\beta,$$

as desired.

(b)

The Hessian $\nabla^2 f(\beta)$ is given by

$$\nabla^2 f(\beta) = 2\tilde{X}^T \tilde{X} + 2\lambda \mathbb{I}_{p-1},$$

where \mathbb{I}_{p-1} is the $(p-1)\times (p-1)$ identity matrix. Note that $2\tilde{X}^T\tilde{X}\in\mathbb{S}^{p-1}_+$ (positive semi-definite) and, for $\lambda>0$, we have $2\lambda\mathbb{I}_{p-1}\in\mathbb{S}^{p-1}_{++}$ (positive definite). Therefore, for all nonzero vectors $v\in\mathbb{R}^{p-1}$,

$$\begin{split} v^T \nabla^2 f(\beta) v &= v^T \left(2\tilde{X}^T \tilde{X} + 2\lambda \mathbb{I}_{p-1} \right) v \\ &= 2v^T \tilde{X}^T \tilde{X} v + 2\lambda v^T \mathbb{I}_{p-1} v \\ &= 2 \left(\underbrace{\|\tilde{X} v\|_2^2}_{\geq 0} + \underbrace{\lambda \|v\|_2^2}_{> 0 \text{ when } \lambda > 0} \right) \\ &> 0 \end{split}$$

Hence,

$$\nabla^2 f(\beta) = 2\tilde{X}^T \tilde{X} + 2\lambda \mathbb{I}_{p-1} \in \mathbb{S}_{++}^{p-1},$$

and so f must be strictly convex in β .

(c)

Suppose a strictly convex function f is globally minimized at distinct points x and y. By strict convexity

$$\forall t \in (0,1) \quad f(tx + (1-t)y) < tf(x) + (1-t)f(y).$$

Since f is minimized at both x and y we have f(x) = f(y), so

$$f(tx + (1-t)y) < tf(x) + (1-t)f(x) = f(x).$$

However, this implies that the point z = tx + (1-t)y yields a value of f even *smaller* than at x, contradicting our assumption that x is a global minimizer. Therefore, strict convexity implies that the global minimizer must be unique, and so for $\lambda > 0$, we are guaranteed that the above solution will be the unique solution to our penalized least squares problem.

(d)

To write our function computing the ridge coefficients we first set $\nabla f(\beta) = 0$

$$\hat{\beta}_{-1}^{(\lambda)} = \left(\tilde{X}^T \tilde{X} + \lambda \mathbb{I}_{p-1}\right)^{-1} \tilde{X}^T \tilde{Y}.$$

For the purpose of computational efficiency we make use of the singular value decomposition of \tilde{X}

$$\tilde{X} = UDV^T$$
.

for $U \in \mathbb{R}^{n \times n}$ and $V \in \mathbb{R}^{(p-1) \times (p-1)}$ both orthogonal matrices, $U^T U = \mathbb{I}_n$, $V^T V = \mathbb{I}_{p-1}$, and $D \in \mathbb{R}^{n \times (p-1)}$ a diagonal matrix with entries $\{d_j\}_{j=1}^{\min(n, p-1)}$ along the main diagonal and zero elsewhere. Hence,

$$\hat{\beta}_{-1}^{(\lambda)} = (\tilde{X}^T \tilde{X} + \lambda \mathbb{I}_{p-1})^{-1} \tilde{X}^T \tilde{Y}$$

$$= ((UDV^T)^T UDV^T + \lambda VV^T)^{-1} (UDV^T)^T \tilde{Y}$$

$$= (VD^T U^T UDV^T + \lambda VV^T)^{-1} VD^T U^T \tilde{Y}$$

$$= (V (D^T D + \lambda \mathbb{I}_{p-1}) V^T)^{-1} VD^T U^T \tilde{Y}$$

$$= V (D^T D + \lambda \mathbb{I}_{p-1})^{-1} V^T VD^T U^T \tilde{Y}$$

$$= V (D^T D + \lambda \mathbb{I}_{p-1})^{-1} D^T U^T \tilde{Y}.$$

Note that $D^TD + \lambda \mathbb{I}_{p-1}$ is a diagonal $(p-1) \times (p-1)$ matrix with entries $d_j^2 + \lambda$, j = 1, ..., p-1, and so the inverse $\left(D^TD + \lambda \mathbb{I}_{p-1}\right)^{-1}$ must also be diagonal with entries $\left(d_j^2 + \lambda\right)^{-1}$, j = 1, ..., p-1. We exploit this to avoid performing a matrix inversion in our function. For brevity, let

$$D^* = \left(D^T D + \lambda I_{p-1}\right)^{-1} D^T,$$

so that

$$\hat{\beta}^{(\lambda)} = V D^* U^T \tilde{Y}.$$

We present a function written in R performing such calculations below.

```
ridge_coef <- function(X, y, lam) {
   Xm1 <- X[,-1] # remove leading column of 1's marking the intercept

ytilde <- y - mean(y) # center response
   xbar <- colMeans(Xm1) # find predictor means
   Xtilde <- Xm1 - tcrossprod(rep(1, nrow(Xm1)), xbar) # center each predictor according to its mean
   # compute the SVD on the centered design matrix
   Xtilde_svd <- svd(Xtilde)</pre>
```

```
U <- Xtilde_svd$u
d <- Xtilde_svd$v

# compute the inverse (D^T D + lambda I_{p-1})^{-1} D^T
Dstar <- diag(d/(d^2 + lam))

# compute ridge coefficients
b <- V %*% (Dstar %*% crossprod(U, ytilde)) # slopes
b1 <- mean(y) - crossprod(xbar, b) # intercept
list(b1 = b1, b = b)
}</pre>
```

Note the choice to use V % % (Dstar %*% crossprod(U, ytilde)) to compute the matrix product $VD^*U^T\tilde{Y}$ as opposed to (the perhaps more intuitive) V % % Dstar %*% t(U) %*% ytilde. Such a choice is empirically justified in an appendix.

(e)

We first take the expectation of $\hat{\beta}_{-1}^{(\lambda)}$

$$\begin{split} \mathbb{E}\left[\hat{\beta}_{-1}^{(\lambda)}\right] &= \mathbb{E}\left[\left(\tilde{X}^T\tilde{X} + \lambda \mathbb{I}_{p-1}\right)^{-1}\tilde{X}^T\tilde{Y}\right] \\ &= \left(\tilde{X}^T\tilde{X} + \lambda \mathbb{I}_{p-1}\right)^{-1}\tilde{X}^T\mathbb{E}\left[\tilde{Y}\right] \\ &= \left(\tilde{X}^T\tilde{X} + \lambda \mathbb{I}_{p-1}\right)^{-1}\tilde{X}^T\tilde{X}\beta_{-1} \end{split}$$

If p >> n then using the SVD on \tilde{X} may yield some speed improvements, that is, with $\tilde{X} = UDV^T$ as above, we find

$$\begin{split} \mathbb{E}\left[\hat{\beta}_{-1}^{(\lambda)}\right] &= \left(\tilde{X}^T\tilde{X} + \lambda \mathbb{I}_{p-1}\right)^{-1}\tilde{X}^T\tilde{X}\beta_{-1} \\ &= V\left(D^TD + \lambda \mathbb{I}_{p-1}\right)^{-1}D^TDV^T\beta_{-1} \\ &= VD^*V^T\beta_{-1} \end{split}$$

where D^* is a diagonal min $(n, p-1) \times \min(n, p-1)$ matrix with diagonal entries $\left\{\frac{d_j^2}{d_j^2 + \lambda}\right\}_{j=1}^{\min(n, p-1)}$ and zero elsewhere.

We next compute the variance of our centered ridge estimates

$$\begin{aligned} \operatorname{Var}\left(\hat{\beta}_{-1}^{(\lambda)}\right) &= \operatorname{Var}\left(\left(\tilde{X}^T\tilde{X} + \lambda \mathbb{I}_{p-1}\right)^{-1}\tilde{X}^T\tilde{Y}\right) \\ &= \left(\tilde{X}^T\tilde{X} + \lambda \mathbb{I}_{p-1}\right)^{-1}\tilde{X}^T\operatorname{Var}\left(\tilde{Y}\right)\left(\left(\tilde{X}^T\tilde{X} + \lambda \mathbb{I}_{p-1}\right)^{-1}\tilde{X}^T\right)^T \\ &= \left(\tilde{X}^T\tilde{X} + \lambda \mathbb{I}_{p-1}\right)^{-1}\tilde{X}^T\operatorname{Var}\left(\tilde{Y}\right)\tilde{X}\left(\tilde{X}^T\tilde{X} + \lambda \mathbb{I}_{p-1}\right)^{-1} \\ &= \sigma_*^2\left(\tilde{X}^T\tilde{X} + \lambda \mathbb{I}_{p-1}\right)^{-1}\tilde{X}^T\tilde{X}\left(\tilde{X}^T\tilde{X} + \lambda \mathbb{I}_{p-1}\right)^{-1} \end{aligned}$$

¹Benchmarks are provided in an appendix for the cases of large n, large p, and $n \approx p$.

as desired. We once again may be interested in applying the SVD on \tilde{X} as we had done before. Such a decomposition gives us a more concise solution

$$\operatorname{Var}\left(\hat{\beta}_{-1}^{(\lambda)}\right) = V D^{**} V^T$$

where D^{**} is a diagonal min $(n, p-1) \times \min(n, p-1)$ matrix with diagonal entries $\left\{\frac{d_j^2}{\left(d_j^2 + \lambda\right)^2}\right\}_{j=1}^{\min(n, p-1)}$ and zero elsewhere.

We now wish to perform a simulation study to estimate our theoretical values $\mathbb{E}\left[\hat{\beta}_{-1}^{(\lambda)}\right]$ and $\operatorname{Var}\left(\hat{\beta}_{-1}^{(\lambda)}\right)$. For readability we first define functions computing the theoretical mean and variance according to our above expressions.

```
ridge_coef_params <- function(X, lam, beta, sigma) {</pre>
  n <- nrow(X); p <- ncol(X)</pre>
  betam1 <- beta[-1] # remove intercept term</pre>
  Xm1 \leftarrow X[,-1] # remove leading column of 1's in our design matrix
  xbar <- colMeans(Xm1) # find prector means</pre>
  Xtilde <- sweep(Xm1, 2, xbar) # center each predictor according to its mean
  if (n \ge p) {
    I \leftarrow diag(p - 1)
    inv <- solve(crossprod(Xtilde) + lam * I)</pre>
    b <- solve(crossprod(Xtilde) + lam * I) %*% (crossprod(Xtilde) %*% betam1)
    vcv <- sigma^2 * inv %*% crossprod(Xtilde) %*% inv</pre>
    list(b = b, vcv = vcv)
  } else {
    # compute SVD on the centered design matrix
    Xtilde_svd <- svd(Xtilde)</pre>
    d <- Xtilde_svd$d
    V <- Xtilde_svd$v
    Dstar \leftarrow diag(d^2/(d^2 + lam))
    Dstar2 \leftarrow diag(d^2/(d^2 + lam)^2)
    b <- V ** (Dstar ** crossprod(V, betam1))
    vcv <- V %*% tcrossprod(Dstar2, V)</pre>
    list(b = b, vcv = vcv)
  }
}
```

We may now perform our simulation.

```
# set parameters
nsims <- 1e3
n <- 25
p <- 7
lam <- 4
beta_star <- 1:p</pre>
```

```
sigma_star <- 1
# generate fixed design matrix
X \leftarrow cbind(1, matrix(rnorm(n * (p - 1)), nrow = n))
# compute theoretical mean and variance
par_true <- ridge_coef_params(X, lam, beta_star, sigma_star)</pre>
b_true <- as.vector(par_true$b)</pre>
vcv_true <- par_true$vcv
# simulate ridge coefficients nsims times
# outputs a matrix with rows corresponding to coefficients
# and columns correspond to simulation number
b_hat <- replicate(nsims, {</pre>
 y <- X ** beta_star + rnorm(n, 0, sigma_star)
 as.vector(ridge_coef(X, y, lam)$b)
# estimate variance of b1, ..., b_p estimates
vcv_hat <- var(t(b_hat))</pre>
# print estimated fused ridge coefficients vs. expected values
b <- rbind(rowMeans(b_hat), b_true)</pre>
rownames(b) <- c("b_hat", "b_true")</pre>
round(b, 4)
                    [,2] [,3]
            [,1]
                                  [,4]
                                        [,5]
## b_hat 0.7861 1.6595 3.2916 3.8786 4.2007 6.3650
## b_true 0.7797 1.6636 3.2936 3.8779 4.2025 6.3689
# print absolute error between estimated and true fused ridge variances
round(abs(vcv_true - vcv_hat), 4)
          [,1]
                 [,2]
                         [,3]
                                [,4]
                                        [,5]
## [1,] 0.0010 0.0008 0.0013 0.0012 0.0008 0.0009
## [2,] 0.0008 0.0008 0.0009 0.0017 0.0011 0.0003
## [3,] 0.0013 0.0009 0.0012 0.0006 0.0015 0.0015
## [4,] 0.0012 0.0017 0.0006 0.0014 0.0005 0.0001
## [5,] 0.0008 0.0011 0.0015 0.0005 0.0007 0.0012
## [6,] 0.0009 0.0003 0.0015 0.0001 0.0012 0.0013
```

We see that the empirical sample estimates are very close to their theoretical values, as expected.

Question 3

Prior to writing our cross-validation function we create some helper functions for the sake of readability

```
ridge_cv_lam <- function(X, y, lam, K) {
    # Helper function for ridge_cv()
    # perform K-fold cross-validation on the ridge regression
    # estimation problem over a single tuning parameter lam
    n <- nrow(X)

if (K > n) {
```

```
stop(paste0("K > ", n, "."))
  } else if (K < 2) {</pre>
    stop("K < 2.")
  # groups to cross-validate over
  folds <- cut(1:n, breaks = K, labels = F)</pre>
  # get indices of training subset
  train_idxs <- lapply(1:K, function(i) !(folds %in% i))</pre>
  cv_err <- sapply(train_idxs, function(tis) {</pre>
    # train our model, extract fitted coefficients
    b_train <- unlist(ridge_coef(X[tis,], y[tis], lam))</pre>
    # find observations needed for testing fits
    test_idx <- !((1:n) %in% tis)
    # fit data
    yhat <- X[test_idx,] %*% b_train</pre>
    # compute test error
    sum((y[test_idx] - yhat)^2)
  })
  # weighted average (according to group size, some groups may have
  # +/- 1 member depending on whether sizes divided unevenly) of
  # cross validation error for a fixed lambda
  sum((cv_err * table(folds)))/n
}
```

Then, our cross-validation function is as follows:

```
ridge_cv <- function(X, y, lam.vec, K) {
    # perform K-fold cross-validation on the ridge regression
    # estimation problem over tuning parameters given in lam.vec
    n <- nrow(X); p <- ncol(X)

cv.error <- sapply(lam.vec, function(1) ridge_cv_lam(X, y, 1, K))

# extract best tuning parameter and corresponding coefficient estimates
best.lam <- lam.vec[cv.error == min(cv.error)]
best.fit <- ridge_coef(X, y, best.lam)
b1 <- best.fit$b1
b <- best.fit$b1
b <- best.fit$b

list(b1 = b1, b = b, best.lam = best.lam, cv.error = cv.error)
}</pre>
```

Question 4

Question 5

(a)

Taking the gradient of our objective function g with respect to coefficient vector β yields

$$\nabla_{\beta} g(\beta, \sigma^2) = \nabla_{\beta} \left(\frac{n}{2} \log \sigma^2 + \frac{1}{2\sigma^2} \|\tilde{Y} - \tilde{X}\beta\|_2^2 + \frac{\lambda}{2} \|\beta\|_2^2 \right)$$
$$= \frac{1}{\sigma^2} \left(-\tilde{X}^T \tilde{Y} + \tilde{X}^T \tilde{X}\beta \right) + \lambda \beta,$$

while the gradient of g with respect to σ^2 is given by

$$\begin{split} \nabla_{\sigma^2} g(\beta, \sigma^2) &= \nabla_{\beta} \left(\frac{n}{2} \log \sigma^2 + \frac{1}{2\sigma^2} \|\tilde{Y} - \tilde{X}\beta\|_2^2 + \frac{\lambda}{2} \|\beta\|_2^2 \right) \\ &= \frac{n}{2\sigma^2} - \frac{1}{2\sigma^4} \|\tilde{Y} - \tilde{X}\beta\|_2^2. \end{split}$$

as desired.

(b)

We first consider the objective function in terms of β . We find the Hessian with respect to β

$$\begin{split} \nabla_{\beta}^{2}g\left(\beta,\sigma^{2}\right) &= \nabla_{\beta}^{2}\left(\frac{n}{2}\log\sigma^{2} + \frac{1}{2\sigma^{2}}\|\tilde{Y} - \tilde{X}\beta\|_{2}^{2} + \frac{\lambda}{2}\|\beta\|_{2}^{2}\right) \\ &= \nabla_{\beta}\left(\frac{1}{\sigma^{2}}\tilde{X}^{T}\left(-\tilde{Y} + \tilde{X}\beta\right) + \lambda\beta\right) \\ &= \tilde{X}^{T}\tilde{X} + \lambda\mathbb{I}_{p-1}. \end{split}$$

The symmetric matrix $\tilde{X}^T\tilde{X}$ is always positive semi-definite, and for $\lambda \geq 0$, $\lambda \mathbb{I}_{p-1}$ will also be positive semi-definite (and strictly positive definite when $\lambda > 0$). Thus, the Hessian with respect to β must be positive semi-definite

$$\nabla^2_{\boldsymbol{\beta}} g\left(\boldsymbol{\beta}, \sigma^2\right) = \tilde{X}^T \tilde{X} + \lambda \mathbb{I}_{p-1} \in \mathbb{S}_+^{p-1},$$

and so our objective function $g(\beta, \sigma^2)$ is convex in β . Now, considering the Hessian with respect to σ^2 ,

$$\begin{split} \nabla_{\sigma^2}^2 g\left(\beta, \sigma^2\right) &= \nabla_{\sigma^2}^2 \left(\frac{n}{2} \log \sigma^2 + \frac{1}{2\sigma^2} \|\tilde{Y} - \tilde{X}\beta\|_2^2 + \frac{\lambda}{2} \|\beta\|_2^2\right) \\ &= \nabla_{\sigma^2} \left(\frac{n}{2\sigma^2} - \frac{1}{2\sigma^4} \|\tilde{Y} - \tilde{X}\beta\|_2^2\right) \\ &= -\frac{n}{2\sigma^4} + \frac{1}{\sigma^6} \|\tilde{Y} - \tilde{X}\beta\|_2^2 \end{split}$$

For g to be convex in σ^2 we require $\nabla^2_{\sigma^2}g\left(\beta,\sigma^2\right)\geq 0$. However, such a condition is equivalent to

$$n \ge \frac{2}{\sigma^2} \|\tilde{Y} - \tilde{X}\beta\|_2^2$$

As a counterexample consider the following data

```
set.seed(124)
n <- 20
p <- 100
beta <- rep(0.1, p)
sigma <- sqrt(2)

Xtilde <- matrix(rnorm(n * p), nrow = n)
eps <- rnorm(n, 0, sigma^2)
ytilde <- Xtilde %*% beta + eps

rhs <- as.numeric(2/sigma^2 * crossprod(ytilde - Xtilde %*% beta))
n >= rhs
```

[1] FALSE

and so it is not the case that $\nabla_{\sigma^2}^2 g\left(\beta, \sigma^2\right)$ is nonnegative, implying that our objective function $g\left(\beta, \sigma^2\right)$ is not convex in σ^2 .

(c)

Let $\bar{\beta}$ be a solution to our maximum likelihood ridge estimation problem such that, for $\lambda > 0$, we have

$$\tilde{Y} - \tilde{X}\bar{\beta} = 0.$$

Since $\bar{\beta}$ is a solution it must satisfy our first order condition

$$\nabla_{\beta} g(\beta, \sigma^2) = \frac{1}{\sigma^2} \left(-\tilde{X}^T \tilde{Y} + \tilde{X}^T \tilde{X} \beta \right) + \lambda \beta = 0 \iff \frac{1}{\sigma^2} \left(\tilde{X}^T \left(-\tilde{Y} + \tilde{X} \beta \right) \right) + \lambda \beta = 0.$$

Thus, for such a solution $\bar{\beta}$ and $\lambda > 0$,

$$0 = \frac{1}{\sigma^2} \left(\tilde{X}^T \left(-\tilde{Y} + \tilde{X}\bar{\beta} \right) \right) + \lambda \bar{\beta}$$
$$= \frac{1}{\sigma^2} \left(\tilde{X}^T \left(-\tilde{Y} + \tilde{Y} \right) \right) + \lambda \bar{\beta}$$
$$= \lambda \bar{\beta}$$
$$\iff \bar{\beta} = 0.$$

Similarly, using our second first order condition $\nabla_{\sigma^2} g(\beta, \sigma^2) = 0$, at $\beta = \bar{\beta}$,

$$\begin{split} \nabla_{\sigma^2} g(\beta, \sigma^2) &= \frac{n}{2\sigma^2} - \frac{1}{2\sigma^4} \|\tilde{Y} - \tilde{X}\beta\|_2^2 \\ &= \frac{n}{2\sigma^2} - \frac{1}{2\sigma^4} \|\tilde{Y} - \tilde{X}\bar{\beta}\|_2^2 \\ &= \frac{n}{2\sigma^2} - \frac{1}{2\sigma^4} \|\tilde{Y} - \tilde{Y}\|_2^2 \\ &= \frac{n}{2\sigma^2} = 0 \end{split}$$

This conditions implies that either n=0 or $\sigma^2\to\infty$. Thus, no such global minimizer could exist.

(d)

Solving our first order conditions

$$\frac{1}{\sigma^2} \left(\tilde{X}^T \left(-\tilde{Y} + \tilde{X}\bar{\beta} \right) \right) + \lambda \bar{\beta} = 0$$
$$\frac{n}{2\sigma^2} - \frac{1}{2\sigma^4} \|\tilde{Y} - \tilde{X}\beta\|_2^2 = 0,$$

we find the maximum likelihood estimate $\hat{\beta}^{(\lambda, ML)}$ to be

$$\hat{\beta}^{(\lambda, ML)} = (\tilde{X}^T \tilde{X} + \sigma^2 \lambda \mathbb{I}_{p-1})^{-1} \tilde{X}^T \tilde{Y}.$$

and the maximum likelihood estimate $\hat{\sigma}^{2(\lambda, ML)}$ to be

$$\hat{\sigma}^{2(\lambda, ML)} = \frac{1}{n} \|\tilde{Y} - \tilde{X}\hat{\beta}^{(\lambda, ML)}\|_2^2$$

To compute such estimates we may use the following algorithm: Consider some fixed data set $\mathcal{D} = \{X, Y\}$ and a fixed tuning parameter λ .

- (1) Center the data: Center each predictor by its mean $X \mapsto \tilde{X}$, center the response vector by its mean $Y \mapsto \tilde{Y}$.
- (2) Have some initial proposal for the estimate $\hat{\sigma}_0^{2(\lambda, ML)} \in \mathbb{R}^+$.
- (3) Compute an initial proposal for $\hat{\beta}_0^{(\lambda,ML)}$ based on $\hat{\sigma}_0^{2\,(\lambda,ML)}$.
- (4) Update our variance estimate $\hat{\sigma}_i^{2(\lambda, ML)}$ using the previous estimate of $\hat{\beta}_{i-1}^{(\lambda, ML)}$.
- (5) Update our coefficient estimate $\hat{\beta}_i^{(\lambda, ML)}$ using the new estimate of $\hat{\sigma}_i^{2(\lambda, ML)}$.
- (6) Repeat steps (5)-(6) until some convergence criteria is met, say $\|\hat{\sigma}_i^{2\,(\lambda,\,ML)} \hat{\sigma}_{i-1}^{2\,(\lambda,\,ML)}\|$, is small.

(e)

Our function is as follows

```
ridge_coef_mle <- function(X, y, lam, tol = 1e-16) {
   Xm1 <- X[,-1] # remove leading column of 1's marking the intercept

ytilde <- y - mean(y) # center response
   xbar <- colMeans(Xm1) # find predictor means
   Xtilde <- sweep(Xm1, 2, xbar) # center each predictor according to its mean

# compute the SVD on the centered design matrix

Xtilde_svd <- svd(Xtilde)
   U <- Xtilde_svd$u
   d <- Xtilde_svd$d
   V <- Xtilde_svd$v

## generate some initial guess for sigma and beta
   sig0 <- rexp(1)
   Dstar <- diag(d/(d^2 + sig0^2 * lam))</pre>
```

```
b0 <- V ** (Dstar ** crossprod(U, ytilde))
  i <- 1
  repeat {
    # update sigma and beta
    sig_new <- sqrt(1/n * crossprod(ytilde - Xtilde %*% b0))</pre>
    Dstar \leftarrow diag(d/(d^2 + sig_new^2 * lam))
    b_new <- V %*% (Dstar %*% crossprod(U, ytilde))</pre>
    if (abs(sig_new^2 - sig0^2) < tol)</pre>
      break
    sig0 <- sig_new
    b0 <- b_new
    i <- i + 1
  list(niter = i, sigma = as.numeric(sig_new), b = b_new)
grad_mle <- function(X, y, lam, b, s) {</pre>
  n \leftarrow nrow(X)
  Xm1 <- X[,-1] # remove leading column of 1's marking the intercept
  ytilde <- y - mean(y) # center response</pre>
  xbar <- colMeans(Xm1) # find predictor means</pre>
 Xtilde <- sweep(Xm1, 2, xbar) # center each predictor according to its mean</pre>
  gb <- 1/s^2 * crossprod(Xtilde, Xtilde %*% b - ytilde) + lam * b
  gs \langle -n/(2 * s^2) - 1/(2 * s^4) * crossprod(ytilde - Xtilde %*% b)
  c(grad_b = gb, grad_s = gs)
```

(f)

```
set.seed(124)
n <- 100
p <- 5
lam <- 1
beta_star <- (-1)^(1:p) * rep(5, p)
sigma_star <- sqrt(1/2)
X \leftarrow cbind(1, matrix(rnorm(n * (p - 1)), nrow = n))
y <- X %*% beta_star + rnorm(n, 0, sigma_star)
rcm <- ridge_coef_mle(X, y, lam)</pre>
rcm
## $niter
## [1] 9
##
## $sigma
## [1] 0.6559084
##
## $b
```

```
## [,1]
## [1,] 4.976904
## [2,] -5.000078
## [3,] 4.888082
## [4,] -5.017066
grad_mle(X, y, lam, rcm$b, rcm$sigma)

## grad_b1 grad_b2 grad_b3 grad_b4 grad_s
## 5.178080e-13 -1.419309e-12 4.849454e-13 -9.281464e-13 1.421085e-14
```

Question 6

as desired.

(a)

Consider our objective function

$$f(\beta) = \frac{1}{2} \|\tilde{Y} - \tilde{X}\beta\|_2^2 + \frac{\lambda_1}{2} \|\beta\|_2^2 + \frac{\lambda_2}{2} \sum_{j=2}^p (\beta_j - \beta_{j-1})^2$$

To show convexity we wish to show $\nabla^2 f(\beta) \in \mathbb{S}^{p-1}_+$. However, it's not immediately obvious how to take such a gradient with our fused sum terms $(b_j - \beta_{j-1})^2$. One way to get around this is to define vector $B \in \mathbb{R}^{p-1}$ given by

$$B = \begin{bmatrix} \beta_2 - \beta_1 \\ \vdots \\ \beta_p - \beta_{p-1} \end{bmatrix}$$

Then

$$\sum_{j=2}^{p} (\beta_j - \beta_{j-1})^2 = B^T B$$

In order to achieve our task of expressing the fused sum in terms of the vector β we must next decompose B into a product of β and some matrix. To this end we define matrix $A \in \mathbb{R}^{(p-2)\times (p-1)}$ with entries -1 along the main diagonal and 1 along the upper diagonal, i.e.,

$$A = \begin{bmatrix} -1 & 1 & 0 & \cdots & 0 & 0 \\ 0 & -1 & 1 & \cdots & 0 & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & 0 & 0 & \cdots & -1 & 1 \end{bmatrix}$$

Then

$$\sum_{j=2}^{p} (\beta_j - \beta_{j-1})^2 = B^T B$$
$$= \beta^T A^T A \beta$$
$$\equiv ||A\beta||_2^2$$

Therefore, our objective function can be expressed as

$$\begin{split} f(\beta) &= \frac{1}{2} \|\tilde{Y} - \tilde{X}\beta\|_2^2 + \frac{\lambda_1}{2} \|\beta\|_2^2 + \frac{\lambda_2}{2} \|A\beta\|_2^2 \\ &\equiv \frac{1}{2} \tilde{Y}^T \tilde{Y} - \beta^T \tilde{X}^T \tilde{Y} + \frac{1}{2} \beta^T \tilde{X}^T \tilde{X}\beta + \frac{\lambda_1}{2} \beta^T \beta + \frac{\lambda_2}{2} \beta^T A^T A\beta \end{split}$$

Hence

$$\nabla f(\beta) = -\tilde{X}^T \tilde{Y} + \tilde{X}^T \tilde{X} \beta + \lambda_1 \beta + \lambda_2 A^T A \beta$$

admitting the Hessian

$$\nabla^2 f(\beta) = \tilde{X}^T \tilde{X} + \lambda_1 \mathbb{I}_{p-1} + \lambda_2 A^T A$$

Recalling that a matrix multiplied with its transpose must always be positive semi-definite, we find \tilde{X}^TX and A^TA must be positive semi-definite. Thus, since $\lambda_1 > 0$, we find that our sum $\tilde{X}^T\tilde{X} + \lambda_1\mathbb{I}_{p-1} + \lambda_2A^TA = \nabla^2 f(\beta)$ is positive semi-definite, and so $f(\beta)$ must be strictly convex, as desired.

(b)

We first solve for $\hat{\beta}_{-1}^{(\lambda_1,\,\lambda_2)}$ in (a) by setting $\nabla f(\beta)=0$

$$0 = -\tilde{X}^T \tilde{Y} + \tilde{X}^T \tilde{X}\beta + \lambda_1 \beta + \lambda_2 A^T A \beta$$
$$\tilde{X}^T \tilde{Y} = \left(\tilde{X}^T \tilde{X} + \lambda_1 \mathbb{I}_{p-1} + \lambda_2 A^T A\right) \beta$$
$$\implies \hat{\beta}_{-1}^{(\lambda_1, \lambda_2)} = M \tilde{X}^T \tilde{Y}$$

where we have set $M = (\tilde{X}^T \tilde{X} + \lambda_1 \mathbb{I}_{p-1} + \lambda_2 A^T A)^{-1}$ for brevity. Therefore

$$\begin{split} \mathbb{E}\left[\hat{\beta}_{-1}^{(\lambda_1,\,\lambda_2)}\right] &= \mathbb{E}\left[M\tilde{X}^T\tilde{Y}\right] \\ &= M\tilde{X}^T\mathbb{E}\left[\tilde{Y}\right] \\ &= M\tilde{X}^T\beta_{*,\,-1} \end{split}$$

and

$$\operatorname{Var}\left(\hat{\beta}_{-1}^{(\lambda_{1}, \lambda_{2})}\right) = \operatorname{Var}\left(M\tilde{X}^{T}Y\right)$$
$$= M\tilde{X}^{T}\operatorname{Var}\left(\tilde{Y}\right)\tilde{X}M^{T}$$
$$= \sigma_{*}^{2}M\tilde{X}^{T}\tilde{X}M^{T}$$

as desired. We now perform our fused ridge simulation study to test the theoretical values with some empirical estimates. We first define our fused ridge coefficient estimation function (as well as functions permitting us to easily compute the theoretical means and variances of the fused ridge problem)

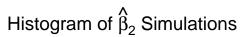
```
fused_ridge_coef <- function(X, y, lam1, lam2) {</pre>
  n \leftarrow nrow(X); p \leftarrow ncol(X)
  Xm1 \leftarrow X[,-1] # remove leading column of 1's marking the intercept
  ytilde <- y - mean(y) # center response</pre>
  xbar <- colMeans(Xm1) # find predictor means</pre>
  Xtilde <- sweep(Xm1, 2, xbar) # center each predictor according to its mean
  I \leftarrow diag(p - 1)
  UD <- cbind(rep(0, p - 2), diag(p - 2)) # upper diagonal matrix
  J \leftarrow -1 * cbind(diag(p - 2), rep(0, p - 2)) # diag(p - 2)*(p - 1) matrix
  A \leftarrow J + UD
  M <- solve(crossprod(Xtilde) + lam1 * I + lam2 * crossprod(A))</pre>
  b <- M %*% crossprod(Xtilde, y)
  b0 <- mean(y) - crossprod(xbar, b)
  return(list(b0 = b0, b = b))
fused_ridge_coef_params <- function(X, lam1, lam2, beta, sigma) {</pre>
  # omits intercept term b0
  # returns theoretical means and variances for the fused ridge problem
  n <- nrow(X); p <- ncol(X)</pre>
  Xm1 \leftarrow X[,-1] # remove leading column of 1's marking the intercept
  betam1 <- beta[-1] # remove intercept term</pre>
  xbar <- colMeans(Xm1) # find predictor means</pre>
  Xtilde <- sweep(Xm1, 2, xbar) # center each predictor according to its mean
  I \leftarrow diag(p - 1)
  UD <- cbind(rep(0, p - 2), diag(p - 2)) # upper diagonal matrix
  J \leftarrow -1 * cbind(diag(p - 2), rep(0, p - 2)) # diag(p - 2)*(p - 1) matrix
  A <- J + UD
  M <- solve(crossprod(Xtilde) + lam1 * I + lam2 * crossprod(A))
  b <- M <pre>%*% crossprod(Xtilde, (Xtilde %*% betam1))
  vcv \leftarrow matrix(0, nrow = p - 1, ncol = p - 1)
  if (n > p) { # when n > p this matrix multiplication routine is quicker
    vcv <- sigma^2 * M %*% tcrossprod(crossprod(Xtilde), M)</pre>
  \} else { # when p > n this matrix multiplication routine is quicker
   vcv <- sigma^2 * tcrossprod(M, Xtilde) %*% tcrossprod(Xtilde, M)</pre>
  return (list(b = b, vcv = vcv))
```

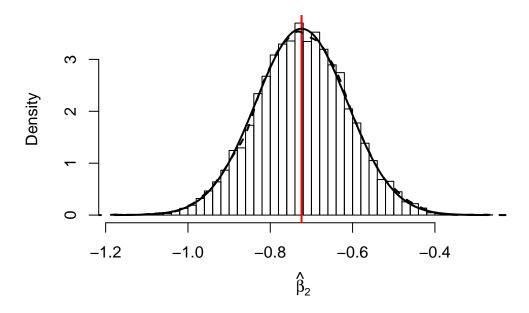
}

We now simulate some data to test our estimates:

```
set.seed(124)
# set parameters
nsims <- 1e4
n <- 1e2
p <- 5
lam1 <- 1
lam2 <- 1
sigma star <- 1
beta_star <- rnorm(p)</pre>
# generate (fixed) design matrix
X \leftarrow cbind(rep(1, n), matrix(rnorm(n * (p - 1)), nrow = n, ncol = p - 1))
# compute expected parameter values
par_true <- fused_ridge_coef_params(X, lam1, lam2, beta_star, sigma_star)</pre>
b_true <- as.vector(par_true$b)</pre>
vcv_true <- par_true$vcv
# simulate our fused ridge coefficients nsims times
# outputs a matrix with rows corresponding to coefficients
# and columns correspond to simulation number
pt <- proc.time()</pre>
b_hat <- replicate(nsims, {</pre>
 y <- X %*% beta_star + rnorm(n, 0, sigma_star) # generate response
 return (as.vector(fused_ridge_coef(X, y, lam1, lam2)$b))
})
proc.time() - pt
##
      user system elapsed
##
     1.862
           0.023
                    1.927
# estimate variance of b2, ..., b_p estimates
vcv_hat <- var(t(b_hat))</pre>
# print estimated fused ridge coefficients vs. expected values
b <- rbind(rowMeans(b_hat), b_true)</pre>
rownames(b) <- c("b_hat", "b_true")</pre>
round(b, 4)
##
            [,1]
                    [,2]
                            [,3]
                                   [,4]
## b hat 0.0316 -0.7226 0.2226 1.3899
## b_true 0.0313 -0.7240 0.2235 1.3920
# print absolute error between estimated and true fused ridge variances
round(abs(vcv_true - vcv_hat), 4)
##
         [,1] [,2] [,3] [,4]
## [1,] 2e-04 1e-04 1e-04 1e-04
## [2,] 1e-04 1e-04 1e-04 2e-04
## [3,] 1e-04 1e-04 0e+00 1e-04
## [4,] 1e-04 2e-04 1e-04 3e-04
```

As a case study, we may look at the simulations of $\hat{\beta}_2^{(\lambda_1,\lambda_2)}$ and compare it with it's theoretical distribution. Note that the estimates $\hat{\beta}^{(\lambda_1,\lambda_2)} = M\tilde{X}^T\tilde{Y}$ are normally distributed because they are a linear combination of $\tilde{Y} \sim \mathcal{N}(\tilde{X}\beta,\sigma^2)$ (when our noise terms $\epsilon \sim \mathcal{N}(0,\sigma^2)$). We visualize the histogram of the $\hat{\beta}_2^{(\lambda_1,\lambda_2)}$ simulations with its empirical and theoretical densities overlaid (dashed, solid), along with its expected value (vertical line) below.





Appendix

Computing $\mathbb{E}\left[\hat{eta}^{(\lambda)}\right]$

```
Consider the case of n >> p
library(microbenchmark)
set.seed(124)
#==== Large n case =====#
# parameters
n < - 1e2
p <- 1e1
lam <- 1
# generate data
beta <- rnorm(p)</pre>
X <- matrix(rnorm(n * p), nrow = n)</pre>
I <- diag(p)</pre>
# define functions
f1 <- function() solve(crossprod(X) + lam * I) %*% (crossprod(X) %*% beta)
f2 <- function() {
  X_svd <- svd(X)</pre>
  V <- X_svd$v
  d \leftarrow X_svd$d
  Dstar \leftarrow diag(d^2/(d^2 + lam))
  V %*% (Dstar %*% crossprod(V, beta))
# test speed
microbenchmark(f1(), f2(), times = 1e3, unit = "us")
## Unit: microseconds
## expr min
                                 mean median
                        lq
                                                      uq
                                                               max neval
## f1() 40.372 44.2995 61.40828 49.5850 65.5915 437.949 1000
## f2() 133.436 142.1035 192.80813 150.4875 192.9895 1808.623 1000
and the case for p >> n
#==== Large p case =====#
# parameters
n <- 1e1
p <- 1e2
lam <- 1
# generate data
beta <- rnorm(p)</pre>
X <- matrix(rnorm(n * p), nrow = n)</pre>
I \leftarrow diag(p)
# define functions
f1 <- function() solve(crossprod(X) + lam * I) %*% (crossprod(X) %*% beta)
f2 <- function() {</pre>
 X_svd <- svd(X)</pre>
```

```
V <- X_svd$v
  d \leftarrow X_svd$d
  Dstar \leftarrow diag(d^2/(d^2 + lam))
  V %*% (Dstar %*% crossprod(V, beta))
# test speed
microbenchmark(f1(), f2(), times = 1e3, unit = "us")
## Unit: microseconds
## expr
                                         median
              min
                          lq
                                  mean
                                                        uq
                                                                  max neval
## f1() 2508.989 2793.9725 3235.0076 3014.476 3376.9035 47197.359 1000
## f2() 144.755 168.1655 251.1069 222.317 272.2155 2082.819 1000
and n \approx p
#==== n ~ p case ====#
# parameters
n <- 1e2
p <- 1e2
lam <- 1
# generate data
beta <- rnorm(p)
X <- matrix(rnorm(n * p), nrow = n)</pre>
I \leftarrow diag(p)
# define functions
f1 <- function() solve(crossprod(X) + lam * I) %*% (crossprod(X) %*% beta)
f2 <- function() {
 X_svd <- svd(X)</pre>
  V <- X_svd$v
 d <- X_svd$d
 Dstar \leftarrow diag(d^2/(d^2 + lam))
  V %*% (Dstar %*% crossprod(V, beta))
# test speed
microbenchmark(f1(), f2(), times = 1e3, unit = "us")
## Unit: microseconds
## expr
              min
                         lq
                                mean
                                       median
                                                              max neval
                                                     uq
## f1() 3297.960 3592.409 4541.548 3931.411 4524.395 61611.97 1000
## f2() 6325.234 6986.127 8254.252 7667.305 8528.119 62919.38 1000
```

Matrix Multiplication Timing

Consider the following matrix multiplication benchmarks (for the cases of n >> p and p >> n).

```
set.seed(124)
#==== Large n case ====#

# set parameters
n <- 1e3
p <- 1e2</pre>
```

```
lam < -1
# generate data
X <- matrix(rnorm(n * p), nrow = n)</pre>
beta <- rnorm(p)
eps <- rnorm(n)
y <- X %*% beta + eps
ytilde <- y - mean(y)</pre>
xbar <- colMeans(X)</pre>
Xtilde <- sweep(X, 2, xbar)</pre>
# compute decomposition
Xtilde_svd <- svd(Xtilde)</pre>
U <- Xtilde_svd$u
d <- Xtilde_svd$d
V <- Xtilde_svd$v</pre>
Dstar \leftarrow diag(d/(d^2 + lam))
# define multiplication functions
f1 <- function() V ** Dstar ** t(U) ** ytilde
f2 <- function() V %*% Dstar %*% (t(U) %*% ytilde)
f3 <- function() V ** (Dstar ** (t(U) ** ytilde))
f4 <- function() V ** (Dstar ** crossprod(U, ytilde))
f5 <- function() V ** crossprod(Dstar, crossprod(U, ytilde))
# test speed
microbenchmark(f1(), f2(), f3(), f4(), f5(), times = 100, unit = "us")
## Unit: microseconds
## expr
                                  mean
                                                                    max neval
              min
                         lq
                                            median
                                                           uq
## f1() 8583.400 9425.3555 10699.1601 10104.0675 10990.2105 45616.497
                                                                           100
## f2() 1106.389 1240.4895 2028.6171 1396.7560 2073.7475 37677.972
                                                                           100
## f3() 366.287 418.3830
                             688.5485
                                        494.5440
                                                    658.9570 2238.895
                                                                           100
                                                     168.7945 1051.026
## f4() 129.725 137.0815
                              171.1268 151.6555
                                                                          100
## f5() 126.731 131.0305
                              157.9678
                                        148.3220
                                                     162.3690 413.771
                                                                           100
#==== Large p case ====#
set.seed(124)
# set parameters
n < - 1e2
p < -1e3
lam < -1
# generate data
X <- matrix(rnorm(n * p), nrow = n)</pre>
beta <- rnorm(p)
eps <- rnorm(n)
y <- X %*% beta + eps
# define multiplication functions
f1 <- function() V %*% Dstar %*% t(U) %*% ytilde
f2 <- function() V ** Dstar ** (t(U) ** ytilde)
```

```
f3 <- function() V %*% (Dstar %*% (t(U) %*% ytilde))
f4 <- function() V ** (Dstar ** crossprod(U, ytilde))
f5 <- function() V %*% crossprod(Dstar, crossprod(U, ytilde))
# test speed
microbenchmark(f1(), f2(), f3(), f4(), f5(), times = 100, unit = "us")
## Unit: microseconds
## expr
             min
                       lq
                                mean
                                       median
                                                              max neval
                                                     uq
## f1() 8543.140 9405.967 10143.3669 10144.878 10907.813 12869.605
                                                                    100
## f2() 1101.534 1183.899 2147.7061 1276.919 1573.379 35488.036
                                                                    100
## f3() 369.646 414.184
                                                                    100
                           926.5855
                                      487.932
                                                561.038 35247.893
## f4() 129.664 137.932
                            157.4483
                                      149.375
                                                165.022
                                                          363.137
                                                                    100
## f5() 126.205 131.436
                            157.5801
                                      146.959
                                                159.629
                                                          764.477
                                                                    100
```