Project 10 Structured sparsity in genetics

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Problem 28: Uniqueness of predictions from the lasso

We'll prove by contradiction.

Suppose $\mathbf{X}\hat{\beta}_1 \neq \mathbf{X}\hat{\beta}_2$.

Let the solution set be denoted as S, we have $\hat{\beta}_1, \hat{\beta}_2 \in S$.

Since S is convex, we also have $\alpha \hat{\beta}_1 + (1-\alpha) \hat{\beta}_2 \in S.$

So that we have:

$$\begin{split} &\frac{1}{2} \left\| \mathbf{y} - \mathbf{X} \left(\alpha \hat{\beta}_1 + (1 - \alpha) \hat{\beta}_2 \right) \right\|_2^2 + \lambda \left\| \alpha \hat{\beta}_1 + (1 - \alpha) \hat{\beta}_2 \right\|_1 = c^* \\ &\frac{1}{2} \left\| \mathbf{y} - \left(\alpha \mathbf{X} \hat{\beta}_1 + (1 - \alpha) \mathbf{X} \hat{\beta}_2 \right) \right\|_2^2 + \lambda \left\| \alpha \hat{\beta}_1 + (1 - \alpha) \hat{\beta}_2 \right\|_1 = c^* \end{split} \tag{1}$$

 $\&= c^* \end{aligned} \end{equation}$

Define the loss function as $f: S \mapsto \|\mathbf{y} - \mathbf{X}\beta\|_2^2$. We know that the linear loss function is strictly convex, so that we have:

$$f\left(\alpha\hat{\beta}_1 + (1-\alpha)\hat{\beta}_2\right) < \alpha f(\hat{\beta}_1) + (1-\alpha)f(\hat{\beta}_2) \tag{2}$$

Similarly, since l_1 -nrom is convex, we have:

$$\left\|\alpha\hat{\beta}_{1}+(1-\alpha)\hat{\beta}_{2}\right\|_{1}\leq\alpha\|\hat{\beta}_{1}\|_{1}+(1-\alpha)\|\hat{\beta}_{2}\|_{1}\tag{3}$$

Combining Equation 2 and Equation 3, we can start the following derivation:

$$\begin{split} &\frac{1}{2}f\left(\alpha\hat{\beta}_{1}+(1-\alpha)\hat{\beta}_{2}\right)+\lambda\left\|\alpha\hat{\beta}_{1}+(1-\alpha)\hat{\beta}_{2}\right\|_{1}\\ &<\alpha\left(f(\hat{\beta}_{1})+\lambda\|\hat{\beta}_{1}\|_{1}\right)+(1-\alpha)\left(f(\hat{\beta}_{2})+\lambda\|\hat{\beta}_{2}\|_{1}\right)\right)\\ &=\alpha c^{*}+(1-\alpha)c^{*}\\ &=c^{*} \end{split} \tag{4}$$

Just a re-write:

$$\frac{1}{2}f\left(\alpha\hat{\beta}_{1}+(1-\alpha)\hat{\beta}_{2}\right)+\lambda\left\|\alpha\hat{\beta}_{1}+(1-\alpha)\hat{\beta}_{2}\right\|_{1}< c^{*}$$

$$\frac{1}{2}\left\|\mathbf{y}-\left(\alpha\mathbf{X}\hat{\beta}_{1}+(1-\alpha)\mathbf{X}\hat{\beta}_{2}\right)\right\|_{2}^{2}+\lambda\left\|\alpha\hat{\beta}_{1}+(1-\alpha)\hat{\beta}_{2}\right\|_{1}< c^{*}$$
(5)

\end{equation}

Equation 5 contradicts with Equation 1, which suggests that our assumption $\mathbf{X}\hat{\beta}_1 \neq \mathbf{X}\hat{\beta}_2$ is wrong.

Problem 29: Ridge regression solution

Given $n \times p$ matrix **X**, response n-vector **y**, and the parameter p-vector β , consider augmenting **X** with rows corresponding to $\sqrt{\lambda}$ times a $p \times p$ identity matrix I:

$$\mathbf{X}_* = \begin{pmatrix} \mathbf{X} \\ \sqrt{\lambda} \mathbf{I} \end{pmatrix}$$

and \mathbf{y} is similarly augmented with p zeros at its end into \mathbf{y}_* .

Now, the least square objective function on the modified dataset turns out to be:

$$(\mathbf{y}_* - \mathbf{X}_* \beta)^T (\mathbf{y}_* - \mathbf{X}_* \beta)$$

$$= \mathbf{y}_*^T \mathbf{y}_* - 2\mathbf{y}_*^T \mathbf{X}_* \beta + \beta^T \mathbf{X}_*^T \mathbf{X}_* \beta$$
(6)

The solution to the linear regression on the modified dataset hence follows:

$$\hat{\beta} = (\mathbf{X}_*^T \mathbf{X}_*)^{-1} \mathbf{X}_*^T \mathbf{y} \tag{7}$$

Since we are just padding y with zeros, the augmented part of X_* will not have any effect when it is multiplied by y_*^T :

$$\mathbf{y}_*^T \mathbf{X}_* = \mathbf{y}^T \mathbf{X} \tag{8}$$

And finally with augmented dataset X_* we have:

$$\mathbf{X}_{*}^{T}\mathbf{X}_{*}$$

$$= (\mathbf{X}^{T} \quad \sqrt{\lambda}\mathbf{I}) \begin{pmatrix} \mathbf{X} \\ \sqrt{\lambda}\mathbf{I} \end{pmatrix}$$

$$= \mathbf{X}^{T}\mathbf{X} + \lambda\mathbf{I}$$
(9)

Plug Equation 8 and Equation 9 back into Equation 7, we delightfully sees that performing linear regression on the modified dataset gives:

$$\hat{\beta} = (\mathbf{X}^T \mathbf{X} + \lambda \mathbf{I})^{-1} \mathbf{X}^T \mathbf{y} \tag{10}$$

which is just the ridge regression solution.

Problem 30: Variable selection under various norms

Load data and split into training and test set

```
# load data
set.seed(42)
load("yeastStorey.rda")

sample <- createDataPartition(data$Marker, p = 0.7, list = FALSE)

train <- data[sample, ]
test <- data[-sample, ]

X_train <- as.matrix(train[, -1])
y_train <- train[, 1]

X_test <- as.matrix(test[, -1])
y_test <- test[, 1]</pre>
```

Do 10-fold cross validation

colnames(df) <- c("alpha", "lambda", "cvm")</pre>

```
alphas <- seq(0, 1, 0.1)
foldid <- sample(1:10, size = length(sample), replace = TRUE)</pre>
num_cores <- detectCores()</pre>
registerDoParallel(num_cores)
start <- Sys.time()</pre>
res <- foreach (
  alpha = alphas,
  .combine = c,
  .options.RNG = 42,
  .packages = c('glmnet')
) %dorng% {
  cv_res <- cv.glmnet(</pre>
   X_{train}
    y_train,
    alpha = alpha,
   family = "binomial",
   foldid = foldid,
    type.measure = "mse"
  list(cv_res)
end <- Sys.time()</pre>
end - start
## Time difference of 4.363132 secs
stopImplicitCluster()
df <- do.call(rbind, lapply(1:length(alphas), function(x) {</pre>
  cbind.data.frame(alphas[x], res[[x]]$lambda, res[[x]]$cvm)
```

```
param <- filter(df, cvm == min(cvm))
param

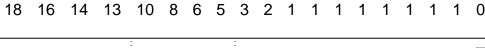
## alpha lambda cvm
## 1 1 0.0126508 0.04302968</pre>
```

For the optimal α (here is 1)...

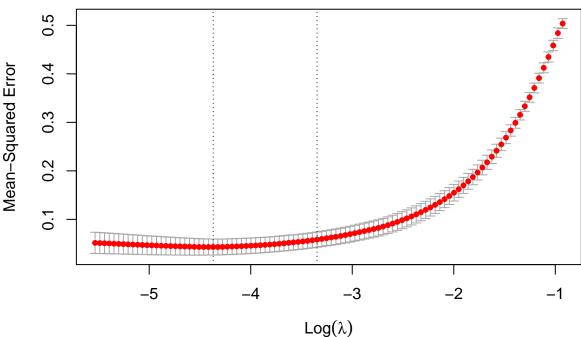
```
opt_idx <- which(alphas == param$alpha)
res.opt_alpha <- res[[opt_idx]]</pre>
```

Plot the mean cross-validated error as a function of $\log \lambda$.

plot(res.opt_alpha)

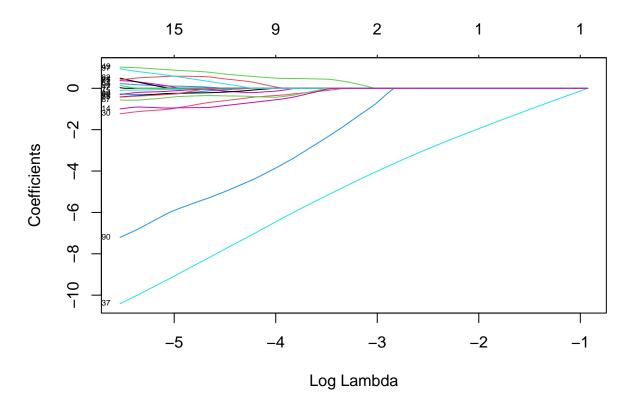


Plot



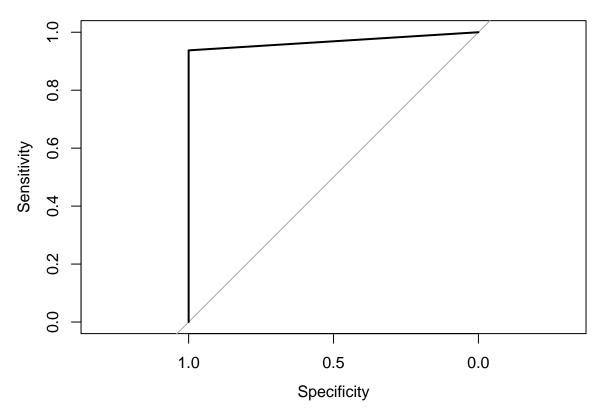
the trace curve of coefficients as a function of log λ .

plot(res.opt_alpha\$glmnet.fit, "lambda", label=TRUE)



Fit the final model and predict the response

```
model <- glmnet(</pre>
  X_train,
  y_train,
  alpha = param$alpha,
  family = "binomial",
  lambda = param$lambda
)
prob <- predict(model, newx = X_test, type = "response")</pre>
y_pred <- ifelse(prob > 0.5, 1, 0)
Report the varaibles selected.
selected <- coef(res.opt_alpha, param$lambda)</pre>
selected@Dimnames[[1]][selected@i]
   [1] "YDL180W" "YEL007W" "YGR046W" "YHL018W" "YIR016W" "YKR096W" "YLR012C"
   [8] "YLR281C" "YNL149C" "YNL213C" "YOL057W" "YPL066W"
Plot ROC...
test_roc <- roc(y_test, as.vector(y_pred))</pre>
## Setting levels: control = 0, case = 1
## Setting direction: controls < cases
plot(test_roc)
```



... and report the corresponding AUC.

test_roc\$auc

Area under the curve: 0.9688