Mathematical techniques in data science

Lecture 13: Shrinkage methods for linear regression

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- A short introduction to statistical learning theory
- Tree-based methods boosting and bootstrapping
- SVM the kernel trick
- Linear regression regularization and feature selection

The story so far...

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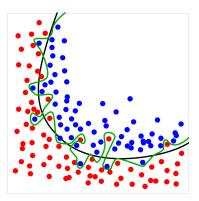
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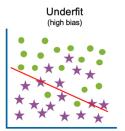
Failure of ERM

We hope that

$$R(\hat{h}_n) \approx R(h^*),$$

but in general, this is not be true if the hypothesis space ${\mathcal H}$ is too large

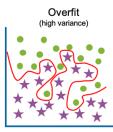




High training error High test error

Optimum

Low training error Low test error

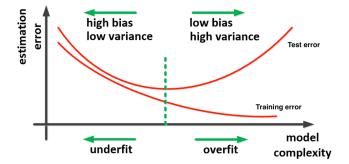


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Low training error
High test error
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(Source: IBM)

Underfiting/Overfitting



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If we want $\epsilon \to 0$ as $n \to \infty$:

 $dimension(\mathcal{H}) \ll n$

How do we get that?

- Model selection
- Feature selection
- Regularization:
 - Work for the case $dimension(\mathcal{H}) \gg n$
 - $\bullet\,$ Stabilize an estimator $\to\,$ force it to live in a neighborhood of a lower-dimensional surface

Regularization: LR

1.1.11.1. Binary Case

For notational ease, we assume that the target y_i takes values in the set $\{0, 1\}$ for data point *i*. Once fitted, the predict_proba method of LogisticRegression predicts the probability of the positive class $P(y_i = 1|X_i)$ as

$$\hat{p}(X_i)= ext{expit}(X_iw+w_0)=rac{1}{1+ ext{exp}(-X_iw-w_0)}$$

As an optimization problem, binary class logistic regression with regularization term r(w) minimizes the following cost function:

(1)

$$\min_w C \sum_{i=1}^n ig(-y_i \log(\hat{p}(X_i)) - (1-y_i) \log(1-\hat{p}(X_i))ig) + r(w).$$

We currently provide four choices for the regularization term r(w) via the penalty argument:

penalty	r(w)
None	0
ℓ_1	$\ w\ _1$
ℓ_2	$rac{1}{2}\ w\ _2^2 = rac{1}{2}w^Tw$
ElasticNet	$rac{1- ho}{2}w^Tw+ ho\ w\ _1$

Scaling the regularization parameter for SVCs

The following example illustrates the effect of scaling the regularization parameter when using Support Vector Machines for classification. For SVC classification, we are interested in a risk minimization for the equation:

$$C\sum_{i=1,n}\mathcal{L}(f(x_i),y_i)+\Omega(w)$$

where

- C is used to set the amount of regularization
- ${\cal L}$ is a loss function of our samples and our model parameters.
- Ω is a penalty function of our model parameters

MLP uses different loss functions depending on the problem type. The loss function for classification is Average Cross-Entropy, which in binary case is given as,

$$Loss(\hat{y},y,W) = -rac{1}{n}\sum_{i=0}^n (y_i \ln \hat{y_i} + (1-y_i) \ln (1-\hat{y_i})) + rac{lpha}{2n}||W||_2^2$$

where $\alpha ||W||_2^2$ is an L2-regularization term (aka penalty) that penalizes complex models; and $\alpha > 0$ is a non-negative hyperparameter that controls the magnitude of the penalty.

For regression, MLP uses the Mean Square Error loss function; written as,

$$Loss(\hat{y},y,W) = rac{1}{2n}\sum_{i=0}^n ||\hat{y}_i - y_i||_2^2 + rac{lpha}{2n}||W||_2^2$$

Starting from initial random weights, multi-layer perceptron (MLP) minimizes the loss function by repeatedly updating these weights. After computing the loss, a backward pass propagates it from the output layer to the previous layers, providing each weight parameter with an update value meant to decrease the loss.

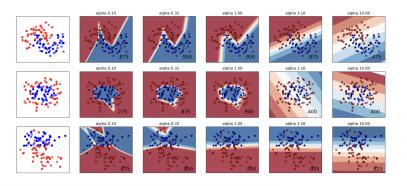
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Regularization: MLP

Varying regularization in Multi-layer Perceptron

A comparison of different values for regularization parameter 'alpha' on synthetic datasets. The plot shows that different alphas yield different decision functions.

Alpha is a parameter for regularization term, aka penalty term, that combats overfitting by constraining the size of the weights. Increasing alpha may fix high variance (a sign of overfitting) by encouraging smaller weights, resulting in a decision boundary plot that appears with lesser curvatures. Similarly, decreasing alpha may fix high bias (a sign of underfitting) by encouraging larger weights, potentially resulting in a more complicated decision boundary.



Linear regression

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$$Y = \beta^{(0)} + \beta^{(1)} X^{(1)} + \beta^{(2)} X^{(2)} + \dots \beta^{(p)} X^{(p)} + \epsilon$$

- *p*: number of variables $(X \in \mathbb{R}^p)$
- n: number of observations

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$$Y = \beta^{(0)} + \beta^{(1)} X^{(1)} + \beta^{(2)} X^{(2)} + \dots \beta^{(p)} X^{(p)}$$

- n ≫ p (n much larger than p). With enough observations, we hope to be able to build a good model
- even if the true relationship between the variables is not linear, we can include transformations of variables

$$X^{(p+1)} = [X^{(1)}]^2, \quad X^{(p+2)} = X^{(1)}X^{(3)}, \dots$$

• adding transformed variables can increase p significantly

$$Y = \beta^{(1)} X^{(1)} + \beta^{(2)} X^{(2)} + \dots \beta^{(p)} X^{(p)} + \epsilon$$

- Higher values of *p* lead to more complex model → increases prediction power/accuracy
- Higher values of *p* make it more difficult to interpret the model: It is often the case that some or many of the variables regression model are in fact not associated with the response

$$Y = \beta^{(0)} + \beta^{(1)} X^{(1)} + \beta^{(2)} X^{(2)} + \dots \beta^{(p)} X^{(p)} + \epsilon$$

- it is often the case that $n \ll p$
- requires supplementary assumptions (e.g. sparsity)
- can still build good models with very few observations.

Linear regression by least squares

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Settings

• $Y \in \mathbb{R}^{n \times 1}$, $X \in \mathbb{R}^{n \times (p+1)}$

$$Y = \begin{bmatrix} y_1 \\ y_2 \\ \dots \\ y_n \end{bmatrix} \qquad X = \begin{bmatrix} 1 & | & | & \dots & | \\ \dots & x^{(1)} & x^{(2)} & \dots & x^{(p)} \\ 1 & | & | & \dots & | \end{bmatrix}$$

where $x^{(1)}, x^{(2)}, \ldots, x^{(p)} \in \mathbb{R}^{n \times 1}$ are the observations of $X^{(1)}, X^{(2)}, \ldots, X^{(p)}$.

We want

$$Y = \beta^{(0)} + \beta^{(1)} X^{(1)} + \beta^{(2)} X^{(2)} + \dots \beta^{(p)} X^{(p)}$$

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• We want

$$Y = \beta^{(0)} + \beta^{(1)} X^{(1)} + \beta^{(2)} X^{(2)} + \dots \beta^{(p)} X^{(p)}$$

• Equivalent to

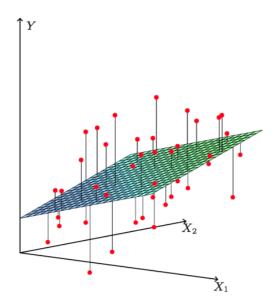
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$$Y = X\beta$$

- In general, the system has no solution ($n \gg p$) or infinitely many solutions ($n \ll p$)
- The most popular estimation method is least squares, in which we pick the coefficients to minimize the residual sum of squares

$$RSS(\beta) = \sum_{i=1}^{n} (y_i - f(x_i))^2$$



• Minimize the residual sum of squares

$$RSS(\beta) = \sum_{i=1}^{n} (y_i - f(x_i))^2$$

• Or alternatively,

$$\hat{\beta} = \min_{\beta} \|Y - X\beta\|_2^2$$

• Minimize the residual sum of squares

$$RSS(\beta) = \sum_{i=1}^{n} (y_i - f(x_i))^2$$

=
$$\sum_{i=1}^{n} \left(y_i - \beta^{(0)} - \beta^{(1)} x_i^{(1)} - \beta^{(2)} x_i^{(2)} - \dots - \beta^{(p)} x_i^{(p)} \right)^2$$

• Taking derivative

$$\frac{\partial RSS}{\partial \beta^{(j)}} = \sum_{i=1}^{n} 2(y_i - x_i\beta) x_i^{(j)} = 2[x^{(j)}]^T (Y - X\beta)$$

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• Set derivatives to zero

$$X^{\mathsf{T}}(Y-X\beta)=0$$

• If $X^T X$ is invertible

$$\hat{\beta} = (X^T X)^{-1} X^T Y$$

Predicted values

$$\hat{Y} = X_{test} \hat{\beta} = X_{test} (X_{train}^{T} X_{train})^{-1} X_{train}^{T} Y_{train}$$

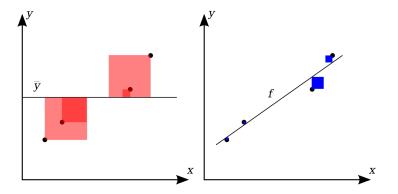
• The coefficient of determination, called "R squared" and denoted by

$$R^{2} = 1 - \frac{\sum_{i=1}^{n} (y_{i} - \hat{y}_{i})^{2}}{\sum_{i=1}^{n} (y_{i} - \bar{y})^{2}}$$

where \bar{y} is the average of y_1, \ldots, y_n

- Often used to measure the quality of a linear model
- A model with a R^2 close to 1 fit the data well.

The coefficient of determination



In some sense, the R^2 measures how much better is the prediction compared to a constant prediction

- It is desirable to adjust R^2 to take account of the fact that its value may be quite high just because many predictors were used relative to the amount of data
- The adjusted coefficient of multiple determination

$$R_a^2 = 1 - \frac{\frac{1}{n-p-1}\sum_{i=1}^n (y_i - \hat{y}_i)^2}{\frac{1}{n-1}\sum_{i=1}^n (y_i - \bar{y})^2}$$

where \bar{y} is the average of y_1, \ldots, y_n

```
>>> import numpy as np
>>> from sklearn.linear_model import LinearRegression
>>> X = np.array([[1, 1], [1, 2], [2, 2], [2, 3]])
>>> # y = 1 * x_0 + 2 * x_1 + 3
>>> y = np.dot(X, np.array([1, 2])) + 3
>>> reg = LinearRegression().fit(X, y)
>>> reg.score(X, y)
1.0
>>> reg.coef_
array([1, 2.])
>>> reg.intercept_
3.0000...
>>> reg.predict(np.array([[3, 5]]))
array([16.])
```

```
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>>> X = np.arange(6).reshape(3, 2)
>>> X
array([[0, 1],
         [2, 3],
[4, 5]])
>>> poly = PolynomialFeatures(2)
>>> poly.fit_transform(X)
array([[ 1., 0., 1., 0., 0., 1.],
[ 1., 2., 3., 4., 6., 9.],
[ 1., 4., 5., 16., 20., 25.]])
>>> poly = PolynomialFeatures(interaction_only=True)
>>> poly.fit_transform(X)
array([[ 1., 0., 1., 0.],
[ 1., 2., 3., 6.],
[ 1., 4., 5., 20.]])
```

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- Is at least one of the predictors X_1, X_2, \ldots, X_p useful in predicting the response?
- Do all the predictors help to explain *Y*, or is only a subset of the predictors useful?

Subset selection

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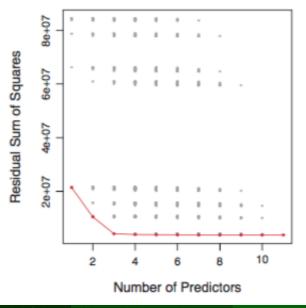
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$$Y = \beta^{(1)} X^{(1)} + \beta^{(2)} X^{(2)} + \dots \beta^{(p)} X^{(p)} + \epsilon$$

- Higher values of p lead to more complex model → increases prediction power/accuracy
- Higher values of p make it more difficult to interpret the model
- Ideally, we would like to try out a lot of different models, each containing a different subset of the predictors, then select the best model
- Problem: there are 2^p models that contain subsets of p variables

Best subset selection



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Algorithm 6.2 Forward stepwise selection

- 1. Let \mathcal{M}_0 denote the *null* model, which contains no predictors.
- 2. For $k = 0, \ldots, p 1$:
 - (a) Consider all p k models that augment the predictors in \mathcal{M}_k with one additional predictor.
 - (b) Choose the *best* among these p k models, and call it \mathcal{M}_{k+1} . Here *best* is defined as having smallest RSS or highest R^2 .
- 3. Select a single best model from among $\mathcal{M}_0, \ldots, \mathcal{M}_p$ using cross-validated prediction error, C_p (AIC), BIC, or adjusted R^2 .

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Algorithm 6.3 Backward stepwise selection

- 1. Let \mathcal{M}_p denote the *full* model, which contains all p predictors.
- 2. For $k = p, p 1, \dots, 1$:
 - (a) Consider all k models that contain all but one of the predictors in \mathcal{M}_k , for a total of k-1 predictors.
 - (b) Choose the *best* among these k models, and call it \mathcal{M}_{k-1} . Here *best* is defined as having smallest RSS or highest R^2 .
- 3. Select a single best model from among $\mathcal{M}_0, \ldots, \mathcal{M}_p$ using cross-validated prediction error, C_p (AIC), BIC, or adjusted \mathbb{R}^2 .

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- Hybrid versions of forward and backward stepwise selection are available
- variables are added to the model sequentially
- after adding each new variable, the method may also remove any variables that no longer provide an improvement in the model fit

Adjusted training errors

- Adjusted R^2
- Mallow's Cp

$$C_p = \frac{1}{n} (RSS + 2d\hat{\sigma}^2)$$

where $\hat{\sigma}^2$ is an estimate of the variance of the error, d is the number of predictors

• AIC (Akaike information criterion)

$$AIC = \frac{1}{n\hat{\sigma}^2}(RSS + 2d\hat{\sigma}^2)$$

• BIC (Bayesian information criterion)

$$BIC = \frac{1}{n\hat{\sigma}^2}(RSS + \log(n)\hat{\sigma}^2)$$

sklearn.feature_selection.RFE

class sklearn.feature_selection. RFE (estimator, n_features_to_select=None, step=1, verbose=0) [source]

Feature ranking with recursive feature elimination.

Given an external estimator that assigns weights to features (e.g., the coefficients of a linear model), the goal of recursive feature elimination (RFE) is to select features by recursively considering smaller and smaller sets of features. First, the estimator is trained on the initial set of features and the importance of each feature is obtained either through a coef_ attribute or through a feature_importances_ attribute. Then, the least important features are pruned from current set of features. That procedure is recursively repeated on the pruned set until the desired number of features to select is eventually reached.

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Shrinkage methods

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$$Y \in \mathbb{R}^{n \times 1}, \quad X \in \mathbb{R}^{n \times (p+1)}$$
$$Y = \begin{bmatrix} y_1 \\ y_2 \\ \cdots \\ y_n \end{bmatrix} \quad X = \begin{bmatrix} 1 & | & | & \cdots & | \\ \cdots & x^{(1)} & x^{(2)} & \cdots & x^{(p)} \\ 1 & | & | & \cdots & | \end{bmatrix}$$

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• Linear model

$$Y = \beta^{(0)} + \beta^{(1)} X^{(1)} + \beta^{(2)} X^{(2)} + \dots \beta^{(p)} X^{(p)} + \epsilon$$

• Equivalent to

$$\mathbf{Y} = \mathbf{X}\beta, \qquad \beta = \begin{bmatrix} \beta^{(0)} \\ \beta^{(1)} \\ \vdots \\ \beta^{(p)} \end{bmatrix}$$

• Least squares regression

$$\hat{\beta}^{LS} = \min_{\beta} \|\mathbf{Y} - \mathbf{X}\beta\|_2^2$$

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• ℓ_0 regularization

$$\hat{\beta}^0 = \min_{\beta} \|\mathbf{Y} - \mathbf{X}\beta\|_2^2 + \lambda \sum_{i=1}^p \mathbf{1}_{\beta^{(i)} \neq \mathbf{0}}$$

where $\lambda > 0$ is a parameter

- pay a fixed price λ for including a given variable into the model
- variables that do not significantly contribute to reducing the error are excluded from the model (i.e., $\beta_i = 0$)
- problem: difficult to solve (combinatorial optimization). Cannot be solved efficiently for a large number of variables.

• Ridge regression/ Tikhonov regularization

$$\hat{\beta}^{RIDGE} = \min_{\beta} \|\mathbf{Y} - \mathbf{X}\beta\|_2^2 + \lambda \sum_{j=1}^{p} [\beta^{(j)}]^2$$

where $\lambda > 0$ is a parameter

- shrinks the coefficients by imposing a penalty on their size
- penalty is a smooth function.
- easy to solve (solution can be written in closed form)
- can be used to regularize a rank deficient problem (n < p)

ℓ_2 (Tikhonov) regularization

$$\frac{\partial \left(\|\mathbf{Y} - \mathbf{X}\beta\|_2^2 + \lambda \|\beta\|^2 \right)}{\partial \beta} = 2\mathbf{X}^{\mathsf{T}} (\mathbf{Y} - \mathbf{X}\beta) + 2\lambda\beta$$

• The critical point satisfies

$$(\mathbf{X}^{\mathsf{T}}\mathbf{X} + \lambda \mathbf{I})\beta = \mathbf{X}^{\mathsf{T}}\mathbf{Y}$$

- Note: $(\mathbf{X}^T \mathbf{X} + \lambda \mathbf{I})$ is positive definite, and thus invertible
- Thus

$$\hat{\beta}^{RIDGE} = (\mathbf{X}^T \mathbf{X} + \lambda \mathbf{I})^{-1} \mathbf{X}^T \mathbf{Y}$$

$$\hat{\beta}^{\text{RIDGE}} = (\mathbf{X}^T \mathbf{X} + \lambda \mathbf{I})^{-1} \mathbf{X}^T \mathbf{Y}$$

- When $\lambda > 0$, the estimator is defined even when n < p
- When $\lambda = 0$ and n > p, we recover the usual least squares solution

The Lasso

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• The Lasso (Least Absolute Shrinkage and Selection Operator)

$$\hat{eta}^{\textit{lasso}} = \min_{eta} \|\mathbf{Y} - \mathbf{X}eta\|_2^2 + \lambda \sum_{j=1}^p |eta^{(j)}|$$

- As with ridge regression, the lasso shrinks the coefficient estimates towards zero
- However, the ℓ_1 penalty has the effect of forcing some of the coefficient estimates to be exactly equal to zero when λ is sufficiently large
- ullet the lasso performs variable selection \rightarrow models are easier to interpret

Alternative form of lasso (using the Lagrangian and min-max argument)

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Lasso: alternative form

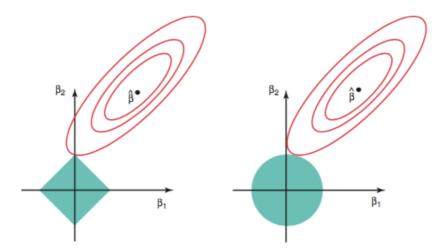


FIGURE 6.7. Contours of the error and constraint functions for the lasso (left) and ridge regression (right). The solid blue areas are the constraint regions, $|\beta_1| + |\beta_2| \le s$ and $\beta_1^2 + \beta_2^2 \le s$, while the red ellipses are the contours of the RSS.

• The Lasso:

$$\hat{eta}^{\textit{lasso}} = \min_{eta} \|\mathbf{Y} - \mathbf{X}eta\|_2^2 + \lambda \sum_{j=1}^{p} |eta^{(j)}|$$

- More "global" approach to selecting variables compared to previously discussed greedy approaches
- Can be seen as a convex relaxation of the $\hat{\beta}^0$ problem
- No closed form solution, but can solved efficiently using convex optimization methods.
- Performs well in practice
- Very popular. Active area of research

Other shrinkage methods

• ℓ_q regularization $(q \ge 0)$:

$$\hat{\beta} = \min_{\beta} \|\mathbf{Y} - \mathbf{X}\beta\|_2^2 + \lambda \sum_{j=1}^{p} [\beta^{(j)}]^q$$



FIGURE 3.12. Contours of constant value of $\sum_{j} |\beta_j|^q$ for given values of q.

Other shrinkage methods

Elastic net

$$\lambda \sum_{j=1}^{p} \alpha [\beta^{(j)}]^2 + (1-\alpha) |\beta^{(j)}|$$

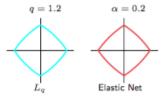


FIGURE 3.13. Contours of constant value of $\sum_j |\beta_j|^q$ for q = 1.2 (left plot), and the elastic-net penalty $\sum_j (\alpha \beta_j^2 + (1-\alpha)|\beta_j|)$ for $\alpha = 0.2$ (right plot). Although visually very similar, the elastic-net has sharp (non-differentiable) corners, while the q = 1.2 penalty does not.

Alternative form of lasso (using the Lagrangian and min-max argument)

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Lasso: alternative form

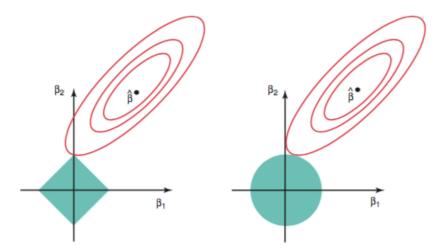


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• Linear model

$$Y = \beta^{(0)} + \beta^{(1)} X^{(1)} + \beta^{(2)} X^{(2)} + \dots \beta^{(p)} X^{(p)} + \epsilon$$

• Equivalent to

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Linear model

$$Y = \beta^{(1)} X^{(1)} + \beta^{(2)} X^{(2)} + \dots \beta^{(p)} X^{(p)} + \epsilon$$

- Higher values of p lead to more complex model → increases prediction power/accuracy
- Higher values of p make it more difficult to interpret the model

 $\bullet \ \ell_0$ regularization

$$\hat{\beta}^0 = \min_{\beta} \|\mathbf{Y} - \mathbf{X}\beta\|_2^2 + \lambda \sum_{i=1}^{p} \mathbf{1}_{\beta^{(i)} \neq 0}$$

• Ridge regression/Tikhonov regularization

$$\hat{\beta}^{\textit{RIDGE}} = \min_{\beta} \|\mathbf{Y} - \mathbf{X}\beta\|_2^2 + \lambda \sum_{j=1}^{p} [\beta^{(j)}]^2$$

Lasso

$$\hat{eta}^{\textit{lasso}} = \min_{eta} \|\mathbf{Y} - \mathbf{X}eta\|_2^2 + \lambda \sum_{j=1}^{p} |eta^{(j)}|$$

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- ℓ_0 , ridge, lasso have regularization parameters λ
- We obtain a family of estimators as we vary the parameter(s)
- optimal parameters needs to be chosen in a principled way
- cross-validation is a popular approach for rigorously choosing parameters.

Demo: Cross-validation with Lasso

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