S&DS 365 / 665 Intermediate Machine Learning

Sparse Regression

Friday, January 28

Topics

- Regression
- High dimensional regression
- Sparsity and the lasso

Regression

We observe pairs $(X_1, Y_1), \ldots, (X_n, Y_n)$.

 $\mathcal{D} = \{(X_1, Y_1), \dots, (X_n, Y_n)\}$ is called the *training data*

 $Y_i \in \mathbb{R}$ is the *response*; $X_i \in \mathbb{R}^p$ is the *covariate* (or feature) vector

For example, suppose we have n subjects and Y_i is the blood pressure of subject i and $X_i = (X_{i1}, \ldots, X_{ip})$ is a vector of p = 5,000 gene expression levels for subject i

Remember: $Y_i \in \mathbb{R}$ and $X_i \in \mathbb{R}^p$

Given a new pair (X, Y), we want to predict Y from X.

Regression

Let \hat{Y} be a prediction of Y. The *prediction error* or *risk* is

$$R = \mathbb{E}(Y - \widehat{Y})^2$$

where \mathbb{E} is the expected value (mean).

The best predictor is the *regression function*

$$m(x) = \mathbb{E}(Y|X=x) = \int y \, p(y|x) dy.$$

However, the true regression function m(x) is not known. We need to estimate m(x).

Regression

Given the training data $\mathcal{D} = \{(X_1, Y_1), \dots, (X_n, Y_n)\}$ we want to construct \widehat{m} to make

prediction risk =
$$R(\widehat{m}) = \mathbb{E}(Y - \widehat{m}(X))^2$$

small. Here, (X, Y) is a new pair.

Bias-variance decomposition

$$R(\widehat{m}) = \int bias^2(x)p(x)dx + \int var(x)p(x) + \sigma^2$$

where

bias(x) =
$$\mathbb{E}(\widehat{m}(x)) - m(x)$$

var(x) = Variance($\widehat{m}(x)$)
 $\sigma^2 = \mathbb{E}(Y - m(X))^2$

Bias-Variance Tradeoff

Prediction Risk = Bias² + Variance

Prediction methods with low bias tend to have high variance.

Prediction methods with low variance tend to have high bias.

For example, the predictor $\widehat{m}(x) \equiv 0$ has 0 variance but will be terribly biased.

To predict well, we need to balance the bias and the variance.

Bias-Variance Tradeoff

More generally, we need to tradeoff approximation error against estimation error:

$$R(\widehat{f}) - R^* = \underbrace{R(\widehat{f}) - \inf_{f \in \mathcal{F}} R(f)}_{\text{estimation error}} + \underbrace{\inf_{f \in \mathcal{F}} R(f) - R^*}_{\text{approximation error}}$$

where R^* is the smallest possible risk and $\inf_{f \in \mathcal{F}} R(f)$ is smallest possible risk using class of estimators \mathcal{F} .

- Approximation error is a generalization of squared bias
- Estimation error is a generalization of variance
- Decomposition holds more generally, even for classification

Linear Regression

Try to find the best linear predictor, that is, a predictor of the form:

$$m(x) = \beta_0 + \beta_1 x_1 + \cdots + \beta_p x_p.$$

Important: We do *not* assume the true regression function is linear.

Can always define $x_1 = 1$; then the intercept is β_1 and we can write

$$m(x) = \beta_1 x_1 + \cdots + \beta_p x_p = \beta^T x$$

where $\beta = (\beta_1, \dots, \beta_p)$ and $x = (x_1, \dots, x_p)$.

Low Dimensional Linear Regression

Assume for now that p (= length of each X_i) is small. To find a good linear predictor we choose β to minimize the *training error*.

training error =
$$\frac{1}{n} \sum_{i=1}^{n} (Y_i - \beta^T X_i)^2$$

The minimizer $\widehat{\beta} = (\widehat{\beta}_1, \dots, \widehat{\beta}_p)$ is called the *least squares estimator*.

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Low Dimensional Linear Regression

The least squares estimator is:

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

where

$$\mathbb{X}_{n \times p} = \begin{pmatrix} X_{11} & X_{12} & \cdots & X_{1p} \\ X_{21} & X_{22} & \cdots & X_{2p} \\ \vdots & \vdots & \vdots & \vdots \\ X_{n1} & X_{n2} & \cdots & X_{np} \end{pmatrix}$$

and

$$\mathbb{Y}=(Y_1,\ldots,Y_n)^T.$$

Exercise: Show this

Low Dimensional Linear Regression

Summary: the least squares estimator is $m(x) = \widehat{\beta}^T x = \sum_j \widehat{\beta}_j x_j$ where

$$\widehat{\beta} = (\mathbb{X}^T \mathbb{X})^{-1} \mathbb{X}^T \mathbb{Y}.$$

When we observe a new X, we predict Y to be

$$\widehat{\mathbf{Y}} = \widehat{\mathbf{m}}(\mathbf{X}) = \widehat{\boldsymbol{\beta}}^T \mathbf{X}.$$

We can try to improve this by:

- (i) dealing with high dimensions
- (ii) using something more flexible than linear predictors.

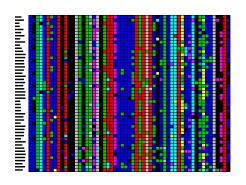
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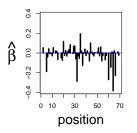
Example

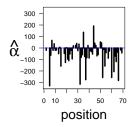
Y = HIV resistance

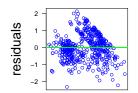
 X_j = amino acid in position j of the virus.

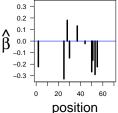
$$Y = \beta_0 + \beta_1 X_1 + \cdots + \beta_{100} X_{100} + \epsilon$$











Top left: $\widehat{\beta}$ fitted values
Top right: marginal regression coefficients (one-at-a-time)

Bottom left: $\widehat{Y}_i - Y_i$ versus \widehat{Y}_i

Bottom right: a sparse regression (coming up soon)

Topics

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High Dimensional Linear Regression

Now suppose p is large. We even might have p > n (more predictors than data points).

The least squares estimator is not defined since $\mathbb{X}^T\mathbb{X}$ is not invertible. The variance of the least squares prediction is huge.

Recall the bias-variance tradeoff:

Prediction Error = Bias² + Variance

We need to increase the bias so that we can decrease the variance.

Ridge Regression

Recall that the least squares estimator minimizes the training error $\frac{1}{n}\sum_{i=1}^{n}(Y_i-\beta^TX_i)^2$.

Instead, we can minimize the penalized training error.

Ridge regression

$$\widehat{\beta} = \underset{\beta}{\operatorname{arg\,min}} \frac{1}{n} \sum_{i=1}^{n} (Y_i - \beta^T X_i)^2 + \frac{\lambda}{n} \|\beta\|_2^2$$

where $\|\beta\|_2 = \sqrt{\sum_j \beta_j^2}$. The solution is:

$$\widehat{\beta} = (\mathbb{X}^T \mathbb{X} + \lambda I)^{-1} \mathbb{X}^T \mathbb{Y}$$

Ridge Regression

The tuning parameter λ controls the bias-variance tradeoff:

$$\lambda = 0 \implies \text{least squares.}$$
 $\lambda = \infty \implies \widehat{\beta} = 0.$

We choose λ to minimize $\widehat{R}(\lambda)$ where $\widehat{R}(\lambda)$ is an estimate of the prediction risk.

Ridge Regression

To estimate the prediction risk, do *not* use training error:

$$R_{\text{training}} = \frac{1}{n} \sum_{i=1}^{n} (Y_i - \widehat{Y}_i)^2, \qquad \widehat{Y}_i = X_i^T \widehat{\beta}$$

because it is biased: $\mathbb{E}(R_{\text{training}}) < R(\widehat{\beta})$

Instead, we use *leave-one-out cross-validation*:

- 1. leave out (X_i, Y_i)
- 2. find $\widehat{\beta}$
- 3. predict Y_i : $\widehat{Y}_{(-i)} = \widehat{\beta}^T X_i$
- 4. repeat for each i

Leave-one-out cross-validation

Can be shown that

$$\widehat{R}(\lambda) = \frac{1}{n} \sum_{i=1}^{n} (Y_i - \widehat{Y}_{(i)})^2 = \frac{1}{n} \sum_{i=1}^{n} \frac{(Y_i - \widehat{Y}_i)^2}{(1 - H_{ii})^2}$$

$$\approx \frac{R_{\text{training}}}{\left(1 - \frac{p_{\lambda}}{n}\right)^2}$$

$$\approx R_{\text{training}} + \frac{2 p_{\lambda} \widehat{\sigma}^2}{n}$$

where

$$H = \mathbb{X}(\mathbb{X}^T \mathbb{X} + \lambda I)^{-1} \mathbb{X}^T$$
$$p_{\lambda} = \operatorname{trace}(H)$$

Example

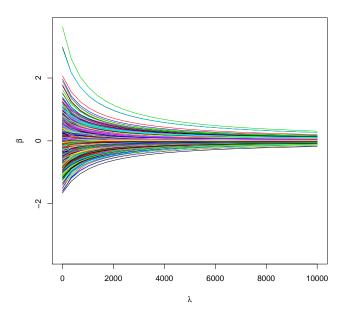
$$Y = 3X_1 + \dots + 3X_5 + 0X_6 + \dots + 0X_{1000} + \epsilon$$

$$n = 100, p = 1,000.$$

So there are 1000 covariates but only 5 are relevant.

What does ridge regression do in this case?

Ridge Regularization Paths



Sparse Linear Regression

Ridge regression does not take advantage of sparsity.

Maybe only a small number of covariates are important predictors. How do we find them?

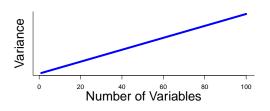
We could fit many submodels (with a small number of covariates) and choose the best one. This is called *model selection*.

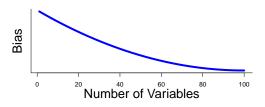
Now the inaccuracy is

prediction error = bias² + variance

The bias is the errors due to omitting important variables. The variance is the error due to having to estimate many parameters.

The Bias-Variance Tradeoff





The Bias-Variance Tradeoff

This is a Goldilocks problem: Can't use too few or too many variables.

Have to choose just the right variables.

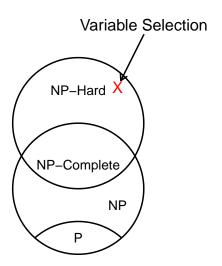
Have to try all models with one variable, two variables,...

If there are p variables then there are 2^p models.

Suppose we have 30,000 genes. We have to search through $2^{30,000}$ models. But $2^{30,000} >$ number of atoms in the universe.

This problem is NP-hard. This was a major bottleneck in statistics for many years.

You are Here



Two things that save us

Two key ideas to make this feasible are sparsity and convex relaxation.

Sparsity: probably only a few genes are needed to predict some disease Y. In other words, of $\beta_1, \ldots, \beta_{30,000}$ most $\beta_i \approx 0$.

But which ones? (Needle in a haystack.)

Convex Relaxation: Replace model search with something easier.

It is the marriage of these two concepts that makes it all work.

Topics

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- High dimensional regression
- Sparsity and the lasso

Sparsity

Consider:

$$\beta = (5, 5, 5, 0, 0, 0, \dots, 0).$$

This vector is high-dimensional but it is sparse.

Here is a less obvious example:

$$\beta = (50, 12, 6, 3, 2, 1.4, 1, 0.8, 0.6, 0.5, \ldots)$$

It turns out that, if the β_j 's die off fairly quickly, then β behaves a like a sparse vector.

Sparsity

We measure the (lack of) sparsity of $\beta = (\beta_1, \dots, \beta_p)$ with the *q*-norm

$$\|\beta\|_q = (|\beta_1|^q + \dots + |\beta_p|^q)^{1/q} = (\sum_j |\beta_j|^q)^{1/q}.$$

Which values of q measure (lack of) sparsity?

sparse:
$$a = 1 \quad 0 \quad 0 \quad \cdots \quad 0$$
 not sparse: $b = 1/\sqrt{p} \quad 1/\sqrt{p} \quad 1/\sqrt{p} \quad \cdots \quad 1/\sqrt{p}$
$$\frac{\sqrt{\qquad \qquad }}{q = 0 \quad q = 1 \quad q = 2} = \frac{\|a\|_q}{\|b\|_q} = \frac{1}{p} \quad \frac{1}{\sqrt{p}} = \frac{1}{\sqrt{p}} = \frac{1}{\sqrt{p}}$$

Lesson: Need to use $q \le 1$ to measure sparsity.

Sparsity

So we estimate $\beta = (\beta_1, \dots, \beta_p)$ by minimizing

$$\sum_{i=1}^{n} \left(Y_i - (\beta_0 + \beta_1 X_{i1} + \cdots + \beta_p X_{ip}) \right)^2$$

subject to the constraint that β is sparse i.e. $\|\beta\|_q \leq \text{small}$.

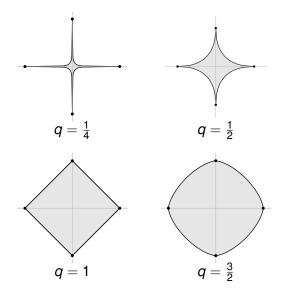
Can we do this minimization?

If we use q = 0 this is same as searching through all 2^p models.

What about other values of *q*?

What does the set $\{\beta : \|\beta\|_q \le \text{small}\}\$ look like?

The set $\|\beta\|_q \le 1$ when p = 2



Sparsity Meets Convexity

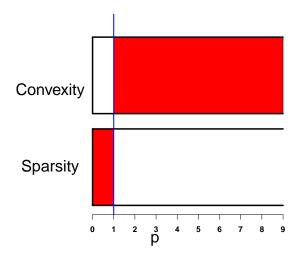
We need these sets to have a nice shape (convex). If so, the minimization is no longer NP-hard. In fact, it is easy.

Sensitivity to sparsity: $q \le 1$ (actually, q < 2 suffices)

Convexity (niceness): $q \ge 1$

This means we should use q = 1.

Where Sparsity and Convexity Meet



Sparsity Meets Convexity

Lasso regression

$$\widehat{\beta} = \operatorname*{arg\,min}_{\beta} \frac{1}{n} \sum_{i=1}^{n} (Y_i - \beta^T X_i)^2 + \lambda \|\beta\|_1$$

where $\|\beta\|_1 = \sum_j |\beta_j|$.

Invented by Rob Tibshirani in 1996. (Related work by Donoho and others around the same time).

Lasso

The result is an estimated vector

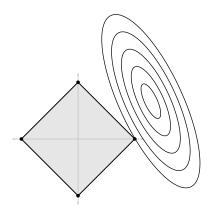
$$\widehat{\beta}_1,\ldots,\widehat{\beta}_p$$

Most are 0!

Magically, we have done model selection without searching (thanks to sparsity plus convexity).

The next picture explains why some $\widehat{\beta}_j = 0$.

Sparsity: How corners create sparse estimators



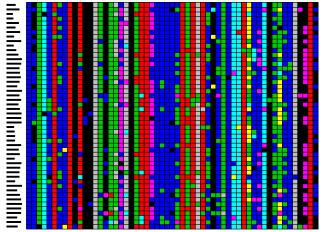
Standardization

Note that, for both lasso and ridge regression, it's important to standarize the features — scale so that the standard deviation of each feature (column) is a constant.

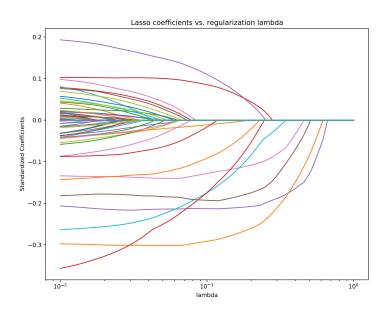
Let's go to the notebook!

The lasso: HIV example again

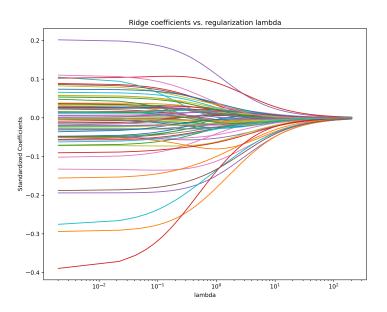
- Y is resistance to HIV drug.
- X_j = amino acid in position j of the virus.
- p = 99, $n \approx 100$.



The lasso: HIV example

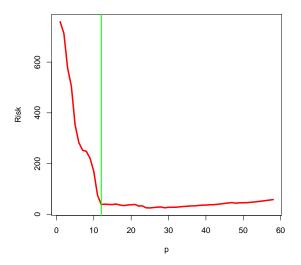


Contrast with ridge regression

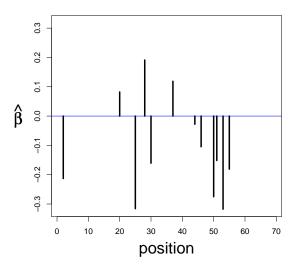


Selecting λ

We choose the sparsity level by estimating prediction error.



The lasso: An example



Sparsity and convexity

To summarize: we penalize the sums of squares with

$$\|\beta\|_q = \left(\sum_j |\beta_j|^q\right)^{1/q}.$$

To get a sparse answer: q < 2.

To get a convex problem: $q \ge 1$.

So q = 1 works.

The marriage of sparsity and convexity is one of the biggest developments in statistics and machine learning.

The lasso

• $\widehat{\beta}(\lambda)$ is called the lasso estimator. Then define

$$\widehat{S}(\lambda) = \left\{ j : \ \widehat{\beta}_j(\lambda) \neq 0 \right\}.$$

• After you find $\widehat{S}(\lambda)$, you should re-fit the model by doing least squares on the sub-model $\widehat{S}(\lambda)$.

The lasso

Choose λ by risk estimation.

Re-fit the model with the non-zero coefficients. Then apply leave-one-out cross-validation:

$$\widehat{R}(\lambda) = \frac{1}{n} \sum_{i=1}^{n} (Y_i - \widehat{Y}_{(i)})^2 = \frac{1}{n} \sum_{i=1}^{n} \frac{(Y_i - \widehat{Y}_i)^2}{(1 - H_{ii})^2} \approx \frac{1}{n} \frac{RSS}{(1 - \frac{s}{n})^2}$$

where *H* is the hat matrix and $s = \#\{j : \widehat{\beta}_j \neq 0\}$.

Choose $\hat{\lambda}$ to minimize $\hat{R}(\lambda)$.

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The lasso

The complete steps are:

- **1** Find $\widehat{\beta}(\lambda)$ and $\widehat{S}(\lambda)$ for each λ .
- **2** Choose $\hat{\lambda}$ to minimize estimated risk.
- 3 Let \hat{S} be the selected variables.
- **4** Let $\widehat{\beta}$ be the least squares estimator using only \widehat{S} .
- **5** Prediction: $\widehat{Y} = X^T \widehat{\beta}$.



Some convexity theory for the lasso

Consider a simpler model than regression: Suppose $Y \sim N(\mu, 1)$. Let $\widehat{\mu}$ minimize

$$A(\mu) = \frac{1}{2}(Y - \mu)^2 + \lambda |\mu|.$$

How do we minimize $A(\mu)$?

• Since A is convex, we set the subderivative = 0. Recall that c is a subderivative of f(x) at x_0 if

$$f(x)-f(x_0)\geq c(x-x_0).$$

• The *subdifferential* $\partial f(x_0)$ is the set of subderivatives. Also, x_0 minimizes f if and only if $0 \in \partial f$.

ℓ_1 and soft thresholding

• If $f(\mu) = |\mu|$ then

$$\partial f = \begin{cases} \{-1\} & \mu < 0 \\ [-1, 1] & \mu = 0 \\ \{+1\} & \mu > 0. \end{cases}$$

Hence,

$$\partial A = \begin{cases} \{\mu - Y - \lambda\} & \mu < 0 \\ \{\mu - Y + \lambda z : -1 \le z \le 1\} & \mu = 0 \\ \{\mu - Y + \lambda\} & \mu > 0. \end{cases}$$

ℓ_1 and soft thresholding

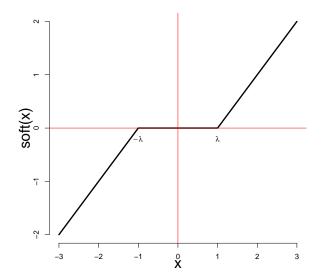
- $\widehat{\mu}$ minimizes $A(\mu)$ if and only if $0 \in \partial A$.
- So

$$\widehat{\mu} = \left\{ \begin{array}{ll} Y + \lambda & Y < -\lambda \\ 0 & -\lambda \le Y \le \lambda \\ Y - \lambda & Y > \lambda. \end{array} \right.$$

This can be written as

$$\widehat{\mu} = \mathsf{soft}(Y, \lambda) \equiv \mathsf{sign}(Y) (|Y| - \lambda)_{+}.$$

ℓ_1 and soft thresholding



The lasso: Computing $\widehat{\beta}$

To minimize $\sum_{i} (Y_i - \beta^T X_i)^2 + \lambda \|\beta\|_1$ by coordinate descent:

- Set $\widehat{\beta} = (0, \dots, 0)$ then iterate the following
- for j = 1, ..., p:
 - set $R_i = Y_i \sum_{s \neq j} \widehat{\beta}_s X_{si}$
 - ▶ Set $\widehat{\beta}_j$ to be least squares fit of R_i 's on X_j .
 - $ightharpoonup \widehat{eta}_j \leftarrow \operatorname{soft}(\widehat{eta}_j, \lambda/\sum_i X_{ij}^2)$
- Then use least squares $\widehat{\beta}$ on selected subset S.

Variations on the lasso

"Elastic net": minimize

$$\sum_{i=1}^{n} (Y_i - \beta^T X_i)^2 + \lambda_1 \|\beta\|_1 + \lambda_2 \|\beta\|_2^2$$

Group lasso:

$$\beta = (\underbrace{\beta_1, \ldots, \beta_k}_{V_1}, \ldots, \underbrace{\beta_t, \ldots, \beta_p}_{V_m})$$

minimize:

$$\sum_{i=1}^{n} (Y_{i} - \beta^{T} X_{i})^{2} + \lambda \sum_{j=1}^{m} ||v_{j}||$$

Summary

- For low dimensional (linear) prediction, we can use least squares.
- For high dimensional linear regression, we face a bias-variance tradeoff: omitting too many variables causes bias while including too many variables causes high variance.
- The key is to select a good subset of variables.
- The lasso (l1-regularized least squares) is a fast way to select variables.
- If there are good, sparse linear predictors, lasso will work well.