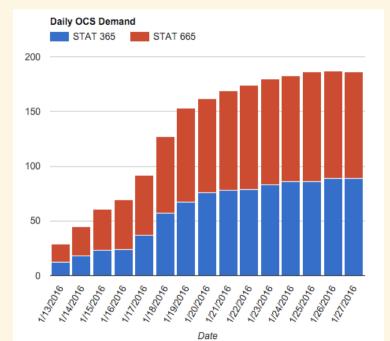
Lecture 04 Least squares and classification

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Yale

- Problem set notes:
 - ▶ brute force okay for implementation question
 - ▶ can use other libraries in the prediction and data analysis questions
 - ► consider FNN (R) or or sklearn.neighbors (Python)
- Office hours (more possibly to come):
 - ► Taylor Arnold Mondays, 13:00 14:15, HH 24, Office 206 (by appointment)
 - ▶ Yu Lu Tuesdays, 10:00-12:00, HH 24, Library
 - ► Jason Klusowski Thursdays, 19:00-20:30, HH 24



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Ordinary least squares

The multivariate linear regression model is given by:

$$y_i = x_{1,i}\beta_1 + x_{2,i}\beta_2 + \dots + x_{1,p}\beta_p + \epsilon_i$$

A sample can be re-written in terms of the vector x_i (the vector of covariates for a single observation):

$$y_i = x_i^t \beta + \epsilon_i$$

In matrix notation, we can write the linear model simultaneously for all observations:

$$\begin{pmatrix} y_1 \\ y_2 \\ \vdots \\ y_n \end{pmatrix} = \begin{pmatrix} x_{1,1} & x_{2,1} & \cdots & x_{p,1} \\ x_{1,2} & \ddots & & x_{p,2} \\ \vdots & & \ddots & \vdots \\ x_{1,n} & x_{2,n} & \cdots & x_{p,n} \end{pmatrix} \begin{pmatrix} \beta_1 \\ \beta_2 \\ \vdots \\ \beta_p \end{pmatrix} + \begin{pmatrix} \epsilon_1 \\ \epsilon_2 \\ \vdots \\ \epsilon_n \end{pmatrix}$$

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Which can be compactly written as:

 $y = X\beta + \epsilon$

For reference, note the following equation

$$y = X\beta + \epsilon$$

Yields these dimensions:

 $y \in \mathbb{R}^n$ $X \in \mathbb{R}^{n \times p}$ $\beta \in \mathbb{R}^p$ $\epsilon \in \mathbb{R}^n$

Least squares

To estimate the least squares solution, which is again the MLE for independent normal errors, we see that:

$$\widehat{\beta} \in \operatorname*{arg\,min}_{b \in \mathbb{R}^p} \left\{ ||y - Xb||_2^2 \right\}$$

$$||y - X\beta||_2^2 = (y - X\beta)^t (y - X\beta)$$

$$\begin{aligned} ||y - X\beta||_2^2 &= (y - X\beta)^t (y - X\beta) \\ &= (y^t - \beta^t X^t) (y - X\beta) \end{aligned}$$

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Normal Equations

In order to find the minimum of the sum of squares, we take the gradient with respect to β and set it equal to zero.

Recall that, for a vector *a* and symmetric matrix *A* :

 $\nabla_{\beta} a^{t} \beta = a$ $\nabla_{\beta} \beta^{t} A \beta = 2A\beta$

Normal Equations

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This gives the gradient of the sum of squares as:

$$\begin{aligned} \nabla_{\beta} ||y - X\beta||_{2}^{2} &= \nabla_{\beta} \left(y^{t}y - 2y^{t}X\beta + \beta^{t}X^{t}X\beta \right) \\ &= 2X^{t}X\beta - 2X^{t}y \end{aligned}$$

Setting this equal to zero gives a set of p equations called the normal equations:

 $X^t X \widehat{\beta} = X^t y$

Maximum or Minimum?

To determine whether the normal equations give a local minimum, maximum, or saddle point, we can calculate the Hessian matrix.

Maximum or Minimum?

To determine whether the normal equations give a local minimum, maximum, or saddle point, we can calculate the Hessian matrix. This is a $p \times p$ matrix giving every combination of the second partial derivatives:

$$Hf(\beta) = \begin{pmatrix} \frac{\partial^2 f}{\partial \beta_1 \partial \beta_1} & \frac{\partial^2 f}{\partial \beta_1 \partial \beta_2} & \cdots & \frac{\partial^2 f}{\partial \beta_1 \partial \beta_p} \\ \frac{\partial^2 f}{\partial \beta_2 \partial \beta_1} & \ddots & & \frac{\partial^2 f}{\partial \beta_2 \partial \beta_p} \\ \vdots & \ddots & \vdots \\ \frac{\partial^2 f}{\partial \beta_p \partial \beta_1} & \frac{\partial^2 f}{\partial \beta_p \partial \beta_2} & \cdots & \frac{\partial^2 f}{\partial \beta_p \partial \beta_p} \end{pmatrix}$$

If the Hessian is positive definite $(x^t H x \ge 0)$ at a critical point, then the critical point is a local minimum.

$$\nabla_{\beta}||y - X\beta||_2^2 = 2X^t X\beta - 2X^t y$$

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We can see that the Hessian is simply:

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We can see that the Hessian is simply:

$$H_{\beta}||y - X\beta||_2^2 = 2X^t \lambda$$

Why is this positive definite?

$$v^{t} (2X^{t}X) v = 2 (v^{t}X^{t}Xv)$$
$$= 2||Xv||_{2}^{2}$$
$$\geq 0$$

Back to the normal equations themselves, notice that if the matrix $X^{t}X$ is invertible, we can 'solve' the normal equations as:

 $X^{t}X\widehat{\beta} = X^{t}y$ $\widehat{\beta} = (X^{t}X)^{-1}X^{t}y$

Back to the normal equations themselves, notice that if the matrix $X^{t}X$ is invertible, we can 'solve' the normal equations as:

$$\mathcal{K}^t X \widehat{\beta} = X^t y$$

$$\widehat{\beta} = (X^t X)^{-1} X^t y$$

This is not a good way to solve the normal equations numerically, but is a useful theoretical form.

Ridge regression

The ridge regression estimator is the solution to the following modified least squares optimization problem for some value of $\lambda > 0$.

$$\widehat{\beta}_{ridge} = \operatorname*{arg\,min}_{b} \left\{ ||y - Xb||_{2}^{2} + \lambda ||b||_{2}^{2} \right\}$$

Why the ridge penalty?

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- 1. The equation shrinks the coefficients towards zero, adding some bias but reducing the variance of the estimator.
- 2. Using the ℓ_2 -norm keeps the equation rotationally invariant.
- 3. Ridge regression has an analytical solution.

To see this write the criterion as a matrix equation:

$$(y - Xb)^{t}(y - Xb) + \lambda b^{t}b = y^{t}y + b^{t}X^{t}Xb - 2y^{t}Xb + \lambda b^{t}b$$

To see this write the criterion as a matrix equation:

$$(y - Xb)^t(y - Xb) + \lambda b^t b = y^t y + b^t X^t Xb - 2y^t Xb + \lambda b^t b$$

And take its derivative:

$$\frac{\partial}{\partial\beta} \left(y^t y + b^t X^t X b - 2y^t X b + \lambda b^t b \right) = 2X^t X b - 2X^t y + 2\lambda b$$

Setting this to zero yields

$$2X^{t}X\widehat{\beta} + 2\lambda\widehat{\beta} = 2X^{t}y$$
$$(X^{t}X + I_{p}\lambda)\widehat{\beta} = X^{t}y$$
$$\widehat{\beta} = (X^{t}X + I_{p}\lambda)^{-1} \cdot X^{t}y$$

Setting this to zero yields

$$\begin{split} & 2X^t X \widehat{\beta} + 2\lambda \widehat{\beta} = 2X^t y \\ & (X^t X + I_p \lambda) \widehat{\beta} = X^t y \\ & \widehat{\beta} = (X^t X + I_p \lambda)^{-1} \cdot X^t y \end{split}$$

This is a useful analytical form, though as with least squares we would generally not invert the matrix directly but instead use a stable matrix decomposition.

Computational issues

How can we estimate the regression vector using a technique such as ordinary least squares

$$\widehat{\beta}_{ols} = \arg\min_{b} \left\{ \sum_{i=1}^{n} (y_i - x_i^t b)^2 \right\},\,$$

When we have a dataset size grows larger than the available memory?

At first glance this seems computationally very difficult as we are trying to minimize a summation with one component per observation.

However, recall that the ordinary least squares solution can be computed by:

$$\widehat{\beta}_{ols} = (X^t X)^{-1} X^t y.$$

Now, assume that the data matrix *X* is broken by rows into *K* different chunks:

$$X = \begin{pmatrix} X_{B_1} \\ X_{B_2} \\ \vdots \\ X_{B_K} \end{pmatrix}$$

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$$X = \begin{pmatrix} X_{B_1} \\ X_{B_2} \\ \vdots \\ X_{B_K} \end{pmatrix}$$

The Gram matrix $X^t X$ can then be computed by summing up the Gram matrices of the individual chunks:

$$X^t X = \sum_{i=1}^K X^t_{B_i} X_{B_i}$$

If the vector y is broken in to the same set of chunks, and corresponding blocks are stored next to one another, such as:

$$X = \begin{pmatrix} X_{B_1} \\ X_{B_2} \\ \vdots \\ X_{B_K} \end{pmatrix}, \quad y = \begin{pmatrix} y_{B_1} \\ y_{B_2} \\ \vdots \\ y_{B_K} \end{pmatrix}$$

The exact same technique works for computing the correlations $X^t y$.

$$X^t y = \sum_{i=1}^K X^t_{B_i} y_{B_i}.$$

So, we can compute the least squares solution:

$$\widehat{\beta}_{ols} = (X^t X)^{-1} X^t y$$

by only working with chunks of the data. Specifically:

$$\widehat{\beta}_{ols} = \left(\sum_{i=1}^{K} X_{B_i}^t X_{B_i}\right)^{-1} \cdot \left(\sum_{i=1}^{K} X_{B_i}^t y_{B_i}\right)$$

This means that we can either work in parallel or with a single process that only reads a small chunk of the data into memory at any given time.

The whole operation only requires, at most, memory for and transmission of $K \cdot (p^2 + p)$ values.

By applying the summation iteratively via *folds*, this can be done by only holding $2(p^2 + p)$ values in memory.

More generally, the following 6 quantities can be easily computed over chunks of the data set:

Gram matrix = $X^t X$ correlation vector = $X^t y$ column sums = $X^t \mathbf{1}_n$ response sums = $y^t \mathbf{1}_n$ response variance = $y^t y$ sample size = $\mathbf{1}_n^t \mathbf{1}_n$

It turns out that these alone are sufficient to calculate many classical and modern estimation techniques.

Ordinary least squares

Not only can we solve ordinary least squares,

$$||y - X\beta||_2^2 = y^t y + \beta^t X^t X\beta + 2y^t X\beta,$$

But we can also calculate an estimator of the noise variance:

$$\widehat{\sigma^2} = \frac{1}{n-p} \left(y^t y + \beta^t X^t X \beta + 2 y^t X \beta \right)$$

And compute standard errors:

S.E(
$$\widehat{\beta}_j$$
) = $\sqrt{\widehat{\sigma^2}(\underline{X^t X})_{jj}^{-1}}$

Ridge regression

The objective function in Ridge regression is also easily written in terms of these quantities:

$$||y - X\beta||_2^2 + \lambda ||\beta||_2^2 = y^t y + \beta^t X^t X\beta + 2y^t X\beta + \lambda \beta^t \beta$$

With similar formulas for standard errors.

Classification problems

Many (most?) of the tasks we'll consider this semester are actually classification tasks. That is, the values y_i that we are trying to predict are class labels rather than a continuous response.

When we have two classes, we can encode these with numerical values. For example, either

$$y_i = \begin{cases} 0\\ 1 \end{cases}$$

Or,

$$y_i = \begin{cases} -1\\ +1 \end{cases}$$



Methods for continuous responses can be used as is (whether the theory extends to discrete data is a different matter entirely). The values of \hat{y} can be interpreted as probabilities. If we need to estimate the actually class label of y, we can use some threshold:

$$\widehat{y}_i^{class} = \begin{cases} 0, & \widehat{y}_i < \alpha \\ 1, & \text{else} \end{cases}$$

For some cutoff value α .

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Some classification methods such as knn can directly produce class estimates; for example, simply using whichever class label is most common near x_{new} .

How can we evaluate how well a classification algorithm works, say when doing cross validation? Mean squared error on the predicted probabilities can work well in many cases; using statistical deviance (based on the log-likelihood) can also make sense in certain contexts.

Another method, which we will largely use in this course, is to instead evaluate the actually class predictions themselves using the **misclassification rate**. On the validation set this can be written as:

$$MCR(k) = \frac{\#\left\{\widehat{y}_i^{class} \neq y_i\right\}_{i \in V}}{\#V}$$

Multiclass classification

It will also be common that we will have values of y_i that can come from more than two classes. For example, tagging a word as a part of speech or labeling an image in the CIFAR-10 corpus.

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Like classification, there are several techniques for evaluating the fit of a multiclass problem. Misclassification rate is also again the simplest and can be defined exactly as it was before.