And Bayesian Machine Learning

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2/28

Previously in Machine Learning

► How to choose the right features if we have (too) many options

- Methods:
 - 1. Subset selection
 - 2. Regularization (shrinkage)
 - 3. Dimensionality reduction (next class)

Best Subset Selection

- Want to find a subset of p features
- ▶ The subset should be small and predict well
- ► Example: credit ~ rating + income + student + limit

```
\mathcal{M}_0 \leftarrow \textit{null model} (no features); 

for k = 1, 2, \dots, p do

| Fit all \binom{p}{k} models that contain k features;
| \mathcal{M}_k \leftarrow \text{best of } \binom{p}{k} models according to a metric (CV error, R^2, etc)

end

return Best of \mathcal{M}_0, \mathcal{M}_1, \dots, \mathcal{M}_p according to metric above

Algorithm 1: Best Subset Selection
```

Complexity of Best Subset Selection?

- ► Complexity of *Best Subset Selection*?
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- Heuristic approaches:
 - 1. **Stepwise selection**: Solve the problem approximately: greedy
 - 2. **Regularization**: Solve a different (easier) problem: <u>relaxation</u>

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 Direct error estimate: Cross validation, precise but computationally intensive

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- Direct error estimate: Cross validation, precise but computationally intensive
- 2. Indirect error estimate: Mellow's C_p :

$$C_p = \frac{1}{n} (\text{RSS} + 2d\hat{\sigma}^2) \text{ where } \hat{\sigma}^2 \approx \text{Var}[\epsilon]$$

Akaike information criterion, BIC, and many others. Theoretical foundations

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Interpretability Penalty: What is the cost of extra features

Regularization

1. Stepwise selection: Solve the problem approximately

- 2. **Regularization**: Solve a different (easier) problem: <u>relaxation</u>
 - ► Solve a machine learning problem, but penalize solutions that use "too much" of the features

Regularization

• Ridge regression (parameter λ), ℓ_2 penalty

$$\min_{\beta} RSS(\beta) + \lambda \sum_{j} \beta_{j}^{2} =$$

$$\min_{\beta} \sum_{i=1}^{n} \left(y_{i} - \beta_{0} - \sum_{j=1}^{p} \beta_{j} x_{ij} \right)^{2} + \lambda \sum_{j} \beta_{j}^{2}$$

Lasso (parameter λ), ℓ_1 penalty

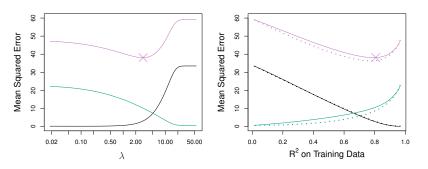
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• Approximations to the ℓ_0 solution

Why Lasso Works

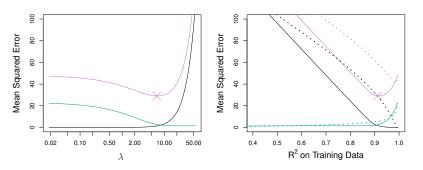
- Bias-variance trade-off
- Increasing λ increases bias
- Example: all features relevant



purple: test MSE, black: bias, green: variance dotted (ridge)

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Regularization: Constrained Formulation

• Ridge regression (parameter λ), ℓ_2 penalty

$$\min_{\beta} \sum_{i=1}^{n} \left(y_i - \beta_0 - \sum_{j=1}^{p} \beta_j x_{ij} \right)^2 \text{ subject to } \sum_{j} \beta_j^2 \leq s$$

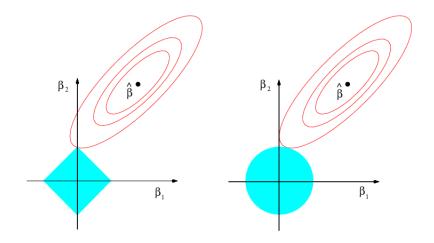
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Lasso Solutions are Sparse

Constrained Lasso (left) vs Constrained Ridge Regression (right)



Constraints are blue, red are contours of the objective

Today

- Dimension reduction methods
 - Principal component regression
 - Partial least squares
- Interpretation in high dimensions
- Bayesian view of ridge regression and lasso

- Different approach to model selection
- We have many features: X_1, X_2, \ldots, X_p
- ▶ Transform features to a *smaller* number $Z_1, ..., Z_M$

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- ▶ New features Z_m are **linear combinations** of X_j :

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▶ **Dimension reduction**: *M* is much smaller than *p*

Using Transformed Features

▶ New features Z_m are **linear combinations** of X_j :

$$Z_m = \sum_{j=1}^p \phi_{jm} X_j$$

► Fit linear regression model:

$$y_i = \theta_0 + \sum_{m=1}^{M} \theta_m z_{im} + \epsilon_i$$

 Run plain linear regression, logistic regression, LDA, or anything else

Prediction using transformed features

$$y_i = \theta_0 + \sum_{m=1}^{M} \theta_m z_{im} + \epsilon_i$$

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Dimension Reduction

1. Reduce dimensions of features Z from X

2. Fit prediction model to compute ϕ

3. Compute weights for the original features β

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How (and Why) Reduce Feaures?

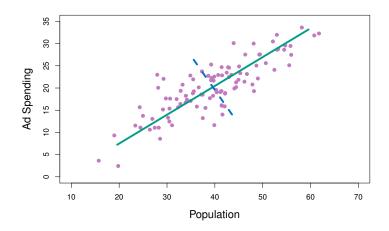
1. Principal Component Analysis (PCA)

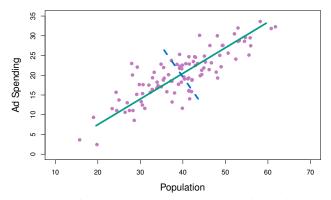
2. Partial least squares

3. Also: many other non-linear dimensionality reduction methods

Principal Component Analysis

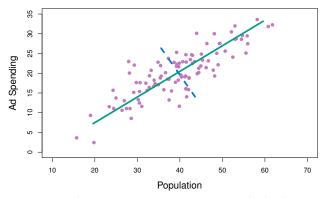
- Unsupervised dimensionality reduction methods
- ▶ Works with $n \times p$ data matrix **X** (no labels)
- Correlated features: pop and ad





▶ **1st Principal Component**: Direction with the largest variance

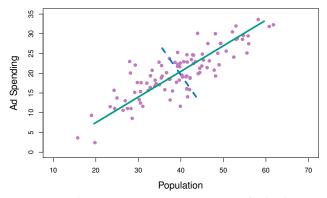
$$Z_1 = 0.839 \times (\mathsf{pop} - \overline{\mathsf{pop}}) + 0.544 \times (\mathsf{ad} - \overline{\mathsf{ad}})$$



▶ 1st Principal Component: Direction with the largest variance

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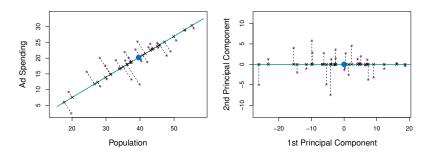
► Is this linear?



▶ 1st Principal Component: Direction with the largest variance

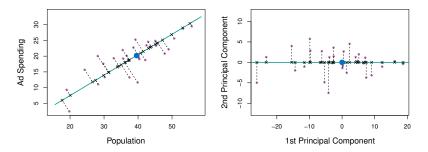
$$Z_1 = 0.839 \times (\mathsf{pop} - \overline{\mathsf{pop}}) + 0.544 \times (\mathsf{ad} - \overline{\mathsf{ad}})$$

Is this linear? Yes, after mean centering.



green line: 1st principal component, minimize distances to all points

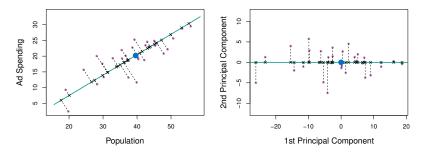
1st Principal Component



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Is this the same as linear regression?

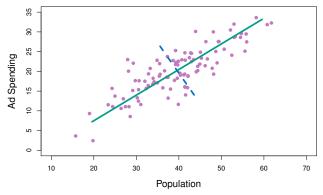
1st Principal Component



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Is this the same as linear regression? **No**, like *total least squares*.

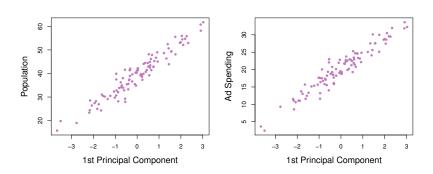
2nd Principal Component



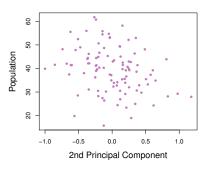
➤ **2nd Principal Component**: Orthogonal to 1st component, largest variance

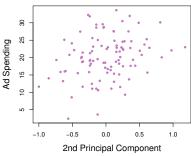
$$Z_2 = 0.544 \times (\mathsf{pop} - \overline{\mathsf{pop}}) - 0.839 \times (\mathsf{ad} - \overline{\mathsf{ad}})$$

1st Principal Component



2nd Principal Component





Properties of PCA

No more principal components than features

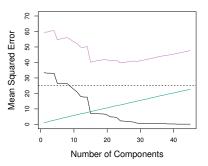
- Principal components are perpendicular
- Principal components are eigenvalues of X^TX

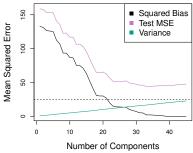
Assumes normality, can break with heavy tails

PCA depends on the scale of features

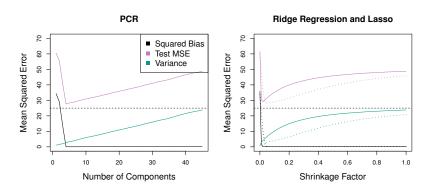
Principal Component Regression

- Use PCA to reduce features to a small number of principal components
- 2. Fit regression using principal components



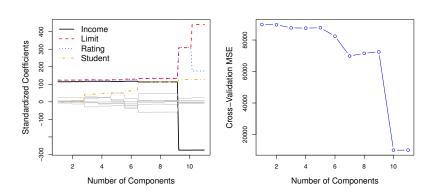


PCR vs Ridge Regression & Lasso



- PCR selects combinations of all features (not feature selection)
- PCR is closely related to ridge regression

PCR Application



Standardizing Features

- Regularization and PCR depend on scales of features
- ► Good practice is to *standardize* features to have **same variance**

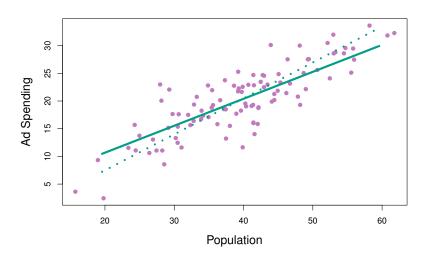
$$\tilde{x}_{ij} = \frac{x_{ij}}{\sqrt{\frac{1}{n} \sum_{i=1}^{n} (x_{ij} - \bar{x}_j)^2}}$$

- Do not standardize features when they have the same units
- PCA needs mean-centered features

$$\tilde{x}_{ij} = x_{ij} - \bar{x}_j$$

Partial Least Squares

Supervised version of PCR



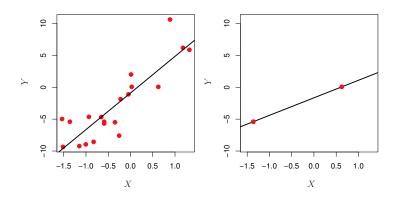
High-dimensional Data

1. Predict blood pressure from DNA: $n=200, p=500\,000$

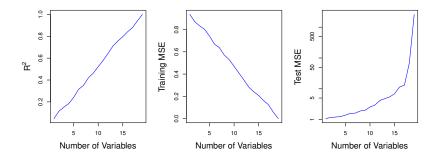
2. Predicting user behavior online: $n=10\,000, p=200\,000$

Problem With High Dimension

- Computational complexity
- Overfitting is a problem



Overfitting with Many Variables



Interpreting Feature Selection

- 1. Solutions may not be unique
- 2. Must be careful about how we report solutions
- 3. Just because one combination of features predicts well, does not mean others will not

Bayesian Machine Learning

- Maximum likelihood
- What if we have prior knowledge?
- Improve on maximum likelihood

Estimating Coefficients: Maximum Likelihood

▶ **Likelihood**: Probability that data is generated from a model

$$\ell(\text{model}) = \Pr[\text{data} \mid \text{model}]$$

Find the most likely model:

$$\max_{\text{model}} \ell(\text{model}) = \max_{\text{model}} \Pr[\text{data} \mid \text{model}]$$

- Likelihood function is difficult to maximize
- ► Transform it using log (strictly increasing)

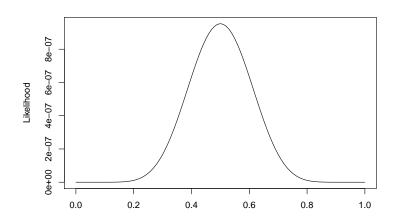
$$\max_{model} \log \ell(model)$$

Strictly increasing transformation does not change maximum

Example: Maximum Likelihood

- Assume a coin with p as the probability of heads
- ▶ **Data**: *h* heads, *t* tails
- ► The likelihood function is:

$$\ell(p) = \binom{h+t}{h} p^h (1-p)^t \approx p^h (1-p)^t.$$



Problems with Maximum Likelihood

► Example!

Bayes Theorem

Classification from label distributions:

$$\Pr[Y = k \mid X = x] = \frac{\Pr[X = x \mid Y = k] \Pr[Y = k]}{\Pr[X = x]}$$

Example:

$$\frac{\Pr[\mathsf{default} = yes \mid \mathsf{balance} = \$100] =}{\Pr[\mathsf{balance} = \$100 \mid \mathsf{default} = yes] \Pr[\mathsf{default} = yes]}{\Pr[\mathsf{balance} = \$100]}$$

▶ Notation:

$$\Pr[Y = k \mid X = x] = \frac{\pi_k f_k(x)}{\sum_{l=1}^K \pi_l f_l(x)}$$

Better Options

1. Maximum likelihood

$$\max_{\substack{\text{model}}} \Pr[\text{data} \mid \text{model}]$$

2. Maximum a posteriori estimate (MAP)

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\max_{\text{model}} \Pr[\text{model} \mid \text{data}]
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Identical when the prior is normal

Maximum a Posteriori Estimate

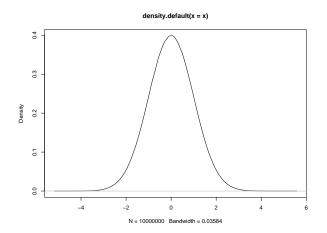
$$\Pr[\beta \mid X,Y] = \alpha f(Y \mid X,\beta) p(\beta \mid X) = \alpha f(Y \mid X,\beta) p(\beta)$$

- Prior: $p(\beta)$
- Likelihood: $f(Y \mid X, \beta)$

Normal Distribution

► Normal density function:

$$f(x) = \frac{1}{\sigma\sqrt{2\pi}}e^{-\frac{(x-\mu)^2}{2\sigma^2}}$$



Maximum Likelihood For OLS

- Assume $\epsilon_i \sim \mathcal{N}(0,1)$
- Likelihood of a single data point

$$f(y_i) = \frac{1}{\sqrt{2\pi}} e^{-\frac{(y_i - \hat{y}_i)^2}{2}}$$

Recall

$$\hat{y}_i = \beta_0 + \beta_1 x_i$$

Likelihood of all data

$$\prod_{i=1}^n f(y_i)$$