Linear models, validation and selection Formation ENSTA-ParisTech Conférence IA

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Model based approaches

Reminder on Supervised Learning

- ▶ input data $X \in \mathbb{R}^p$
- ► reponse *Y* to be predicted
- ▶ training set $(X_1, Y_1), \ldots, (X_n, Y_n)$

Discriminative models

Direct learning of P(Y|X), e.g. generalized linear models s.t.

- Linear regression
- ► Logistic regression (← classification tasks)

Outline

Linear Models

Linear regression
Linear model for classification: Logistic regression

Model Validation

Cross-Validation
Information Criterion

Model Selection

Subset selection
Regularization and shrinkage methods
Ridge regression
Lasso estimator

Applications

prostate data Heart diseases data

Linear Regression Problem

$$X_i = (X_{i,1}, \dots, X_{i,p})^T \in \mathbb{R}^p$$
,

for i = 1, ..., n (sized n training set)

 $ightharpoonup Y_i \in \mathbb{R}$,

Linear Regression Model

$$Y_i = \beta_0 + \sum_{j=1}^p \beta_j X_{i,j} + \sigma \epsilon_i, \quad \text{for } i = 1, \dots, n,$$

- $ightharpoonup \epsilon_i$ is a centered with unit variance $(E[\epsilon_i] = 0, \text{ var}(\epsilon_i) = 1)$ white noise
- $ightharpoonup eta_0$ is the "intercept" (reduces to the ordinate at the origin when p=1)
- $\beta \equiv (\beta_0, \dots, \beta_p) \in \mathbb{R}^{p+1}$ is the coefficient vector

Objective: estimation of $\beta \leftarrow$ supervised learning problem

Linear regression

Linear Regression Problem (Cont'd)

Linear Regression Model

$$Y_i = \beta_0 + \sum_{j=1}^p \beta_j X_{i,j} + \sigma \epsilon_i, \quad \text{for } i = 1, \dots, n,$$

Remark: model linear w.r.t. $\beta \equiv (\beta_0, \dots, \beta_p) \in \mathbb{R}^{p+1}$, but not necessarily linear w.r.t.

- ▶ the inputs X_i : we can add non linear predictors $h(X_1, ..., X_p)$ in the model, e.g. X_i^2 , $X_i X_j ...$
- ▶ the outputs Y_i : we can introduce a non linear link function \leftarrow generalized linear model, e.g. logistic regression

Linear model: Keep it simple!

Simple linear approach may seem overly simplistic

- true regression functions are never linear
- + extremly useful, both conceptually and practically

Practically

Gorge Box, 60': "Essentially, all models are wrong, but some are very useful"

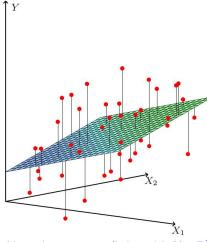
Simple is actually very good: works very well in a lot of situations by capturing the main effects (which are generally the most interesting)

Conceptually

Many concepts developed for the linear problem are important for a lot of the supervised learning techniques

Although it is nerver correct, a linear model serves as a good and interpretable approximation of the unknown true function f(X)

Least Squares (LS) Estimator



Linear least squares fitting with $X \in \mathbb{R}^2$

LS estimate defined by minimizing the Residual Sum of Squares (RSS) $\,$

$$\widehat{\beta} = \arg\min_{\beta} \underbrace{\sum_{i=1}^{n} \left(y_i - \beta_0 - \sum_{j=1}^{p} \beta_j x_{i,j} \right)^2}_{\text{RSS}(\beta)}$$

▶ $RSS(\beta)$ \propto training error rate for quadratic loss

Least Squares Estimator (Cont'd)

$$\widehat{\beta} = \arg\min_{\beta} \mathrm{RSS}(\beta), \quad \text{where } \mathrm{RSS}(\beta) = \sum_{i=1}^{n} \left(y_i - \beta_0 - \sum_{j=1}^{p} \beta_j x_{i,j} \right)^2$$

Matrix expression of RSS

$$RSS(\beta) = ||Y - X\beta||_2^2,$$

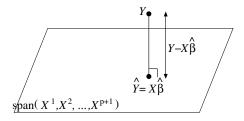
where
$$Y = \begin{pmatrix} y_1 \\ \vdots \\ y_n \end{pmatrix} \in \mathbb{R}^n$$
, $X = \begin{pmatrix} 1 & x_{1,1} & \dots & x_{1,p} \\ \vdots & \vdots & & \vdots \\ 1 & x_{n,1} & \dots & x_{n,p} \end{pmatrix} \in \mathbb{R}^{n \times (p+1)}$

LS Estimator derivation

 $\widehat{Y} = X\widehat{\beta}$ is the prediction in the space spanned by the column vectors of X such that the euclidean error norm $||Y - X\widehat{\beta}||_2$ is minimized

Orthogonality principle

Let X^{j} be the jth column of X



for
$$j = 1, ..., p + 1$$

$$\langle X^{j}, Y - X \widehat{\beta} \rangle = (X^{j})^{T} \left(Y - X \widehat{\beta} \right) = 0,$$

$$\Leftrightarrow \quad X^{T} \left(Y - X \widehat{\beta} \right) = 0,$$

$$\Leftrightarrow \quad \left(X^{T} X \right) \widehat{\beta} = X^{T} Y$$

Linear regression

LS Estimator computation

Assumption: rank X = p + 1, hence X^TX is invertible

Analytical expression

$$\widehat{\beta} = \left(X^T X \right)^{-1} X^T Y,$$

Numerical computation in high dimension

When $p > 10^3$ or $p > 10^4$, too expansive to compute $(X^T X)^{-1}$... More efficient to use a numerical procedure to minimize the criterion $J(\beta) \equiv \frac{1}{2} \mathrm{RSS}(\beta)$, e.g. steepest descent

$$\beta_{k+1} = \beta_k - \alpha_k \nabla_\beta J(\beta_k),$$

where step size $\alpha_k \in \mathbb{R}$ is the *learning rate*, and descent direction is computed as

- ▶ batch gradient $\nabla_{\beta} J(\beta) = X^T X \beta X^T Y$
- ▶ stochastic gradient $\nabla_{\beta} J(\beta) \approx X_i^T X_i \beta X_i^T Y_i$ for i = 1, ..., n (scan of the training set) ← cheaper than batch one for a single iteration
- mini-batch gradient: tradeoff between batch and stochastic gradients

Linear model for classification: Logistic regression

Classification problem $Y \in \mathcal{Y} \leftarrow \text{discrete set}$

Binary classification problem: $\mathcal{Y} = \{1, 2\}$

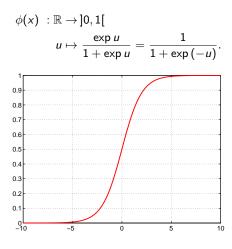
Consider the following model

$$\Pr(Y_i = 1 | X_i = x_i) = \phi(x_i^T \beta) = \frac{\exp(x_i^T \beta)}{1 + \exp(x_i^T \beta)},$$

where

- lacksquare $x_i = (1, x_{i,1}, \dots, x_{i,p})^T \in \mathbb{R}^{p+1} \leftarrow \text{intercept term included by default,}$
- $\phi: u \in \mathbb{R} \mapsto \frac{\exp(u)}{1+\exp(u)} \in (0,1)$ is the logistic function: maps a real value to a probability

Logistic function



Logit link function

Consider

- \triangleright $p_i \equiv \Pr(Y_i = 1 | X_i = x_i) = \phi(x_i^T \beta)$
- $\phi^{-1}: p \in (0,1) \mapsto \log \frac{p}{1-p} \in \mathbb{R}$ is the logit function

Generalized linear model

▶ Linear equation w.r.t. β ,

$$\operatorname{logit}(p_i) = x_i^T \beta,$$

additional nonlinear constraint:

$$\Pr(Y_i = 2 | X_i = x_i) = 1 - \underbrace{\Pr(Y_i = 1 | X_i = x_i)}_{\rho_i} = \frac{1}{1 + \exp(x_i^T \beta)}$$

Maximum Likelihood Estimates

$$\widehat{\beta} = \arg\min_{\beta} -\ell(\beta)$$

where $\ell(\beta)$ is the log-likelihood (here logistic, but normal model yields LSE for linear regression)

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Limitations of LS estimator

Problem

when rank X , or when <math>X has singular values close to zero, then X^TX is no more invertible, or ill conditioned (eigenvalues close to zero)...

Causes

- redundant or nearly-collinear predictors, e.g. $X^k \approx aX^l + b$, where X^j is the jth column of X
- ▶ high dimensional problem where $p \approx n$ (or p > n)

Effects

no single, or stable, solution for $\widehat{\beta}$

- ▶ high variance of $\widehat{\beta}$ as an eigenvalue λ_i of X^TX is close to zero $(||\widehat{\beta}|| \to +\infty \text{ as } \lambda_i \to 0)$,
- true error rate explodes since a small perturbation in the training set yields a substantially different estimate $\hat{\beta}$ and prediction rule $\hat{y} = x^T \hat{\beta}$
- over-fitting problem

Instability of LSE: Deconvolution illustration

- $y \in \mathbb{R}^n$ with $n = 256^2$, $\beta \in \mathbb{R}^p$ with $p = 256^2$,
- ▶ $X \in \mathbb{R}^{n \times p} \leftarrow \text{sized } (256^2) \times (256^2) \text{ matrix...}$



 $\beta \leftarrow \mathsf{original\ image}$



 $y = X\beta \leftarrow \text{blurred image}$

Instability of LSE: Deconvolution illustration

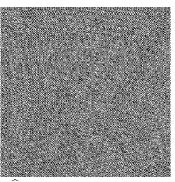
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 $\widehat{\beta} = (X^T X)^{-1} X^T y \leftarrow \mathsf{LS}$ estimate

Reminder on Train and Test Errors

- Loss-function
 - ► Classification: $L(y, \hat{y}) = 0$ if $y = \hat{y}$ else $1 \leftarrow 0$ -1 loss
 - ▶ Regression: $L(y, \hat{y}) = (y \hat{y})^2 \leftarrow \text{quadratic loss}$
- ▶ **Train error**: average loss over the training sample

$$\mathsf{Err}_{\mathsf{train}} = \frac{1}{n} \sum_{i=1}^{n} L(y_i, \hat{y}_i)$$

- ► Test/Prediction error: average loss over a new test sample → Generalization error
- General picture:

$$\mathsf{Err}_{\mathsf{test}} \approx \mathsf{Err}_{\mathsf{train}} + O$$

O would be the average optimism (overfitting problem!)

Model Selection vs Model Validation

Model selection

- Estimate the best set of hyperparameters
- ▶ Estimate the performance of differents models

Model Validation

Estimate the generalization error on unseen/test sample $% \label{eq:control} % \begin{center} \begin{center}$

Model Selection vs Model Validation

Model selection

- Estimate the best set of hyperparameters
- Estimate the performance of differents models

Model Validation

Estimate the generalization error on unseen/test sample



Cross-Validation

K-fold Cross-Validation (CV): Principle

- ▶ Method to estimate prediction error using the training sample
- ▶ Based on splitting the data in K-folds, here K = 5:

$Err_1(\hat{f_1},\lambda)$	Validation	Train	Train	Train	Train
$Err_2(\hat{f}_2,\lambda)$	Train	Validation	Train	Train	Train
$Err_3(\hat{f}_3,\lambda)$	Train	Train	Validation	Train	Train
$Err_4(\hat{f}_4,\lambda)$	Train	Train	Train	Validation	Train
$Err_5(\hat{f}_5,\lambda)$	Train	Train	Train	Train	Validation

where λ are some hyperparameters of the model/method

► Estimate of Test error:

$$\mathrm{CV}(\hat{f},\lambda) = \sum_{k=1}^K \mathsf{Err}_k(\hat{f}_k,\lambda)$$

Cross-Validation

K-fold Cross-Validation (CV): Algorithm

Input: input variables X (dimension $n \times p$), responses y (dim. n), number of folds k

Divide randomly the set $\{1, 2, ..., n\}$ in k subsets (i.e., folds) of roughly equal sizes (e.g., size equals to the integer part of n/k with a little smaller last part if n is not a multiple of k) denoted as $F_1, ..., F_k$

for i = 1 to k:

- Form the validation set (X_{val}, y_{val}) where the indexes of the X and y variables belongs to the ith fold F_i
- Form the training set (X_{train}, y_{train}) where the indexes of the X and y variables belongs to all the folds except F_i
- Train the algorithm/model on the training set (X_{train}, y_{train})
- \triangleright Apply the resulting prediction rule on the input X_{val} of the validation set
- ightharpoonup Compute the error rate on the validation set based on the predictions and the true responses y_{val}

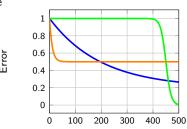
Output: the average error rate computed over all the k folds

Pratical advices

► K? Usually K=5 or 10 is a good trade-off (K=n is called leave-one-out)

	Bias	Variance
K low K high	High Low	Low High
K = n	Low	Very High

Be careful to the learning curve



Number of samples

- Model should be trained completely for each fold (i.e., data normalization, optimization, etc . . .)
- Notebooks: N1_validation_and_model_selection.ipynb, N2_nested_cross_validation_iris.ipynb

Information Criterion

Penalized log-likelihood criterion

$$C(K) = -\hat{\ell}(x; \lambda) + \text{pen}(\lambda, n)$$

- $\hat{\ell}(x;\lambda) \equiv \ell(x;\hat{\beta}_{\lambda})$ with $\hat{\beta}_{\lambda}$ the MLE of the model parameters
- $\hat{\ell}(x;\lambda) \propto \mathrm{RSS}(\hat{\beta}_{\lambda})$ for linear regression (normal model)
- λ are the hyperparameters of the model that drive its complexity (e.g. number k of variables that enter in the model)

Trade-off between two terms (to minimize)

- $-\hat{\ell}(x;\lambda)$: fidelity to the data (likelihood)
- $ightharpoonup \operatorname{pen}(\lambda, n)$: low complexity of the model

Bayesian Information Criterion (BIC)

Asymptotic $(n \gg k_{\lambda})$ criterion for Bayesian models (i.e. with a prior on the model parameters)

$$pen(k_{\lambda}, n) = \frac{1}{2}k_{\lambda}\log(n)$$

- n is the size of the dataset
- \blacktriangleright k_{λ} is the *effective* number of parameters for the λ hyperparameter

$$BIC(\lambda) = -2\hat{\ell}(x;\lambda) + k_{\lambda}\log(n)$$

- Model comparison: select model $\hat{\ell}_j(x;\lambda_j)$ with minimal BIC
- lacktriangleq Hyperparmeter tuning: choose value $\widehat{\lambda}$ that minimizes BIC

Akaïke Information Criterion (AIC)

Other popular asymptotic $(n \gg k_{\lambda})$ criterion

$$AIC(\lambda) = -2\hat{\ell}(x;\lambda) + 2k_{\lambda}$$

- \blacktriangleright k_{λ} is the *effective* number of parameters for the λ hyperparameter
- ► Comparison with BIC: $\log(n) k_{\lambda}$ replaced by $2 k_{\lambda}$
- Variant less aggressive (as long as log(n) > 2, where n is the sample size)

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prostate data Heart diseases data

Subset selection

Motivation

- interpretation: with a large number of predictors, we often would like to determine a smaller subset that exhibit the main (strongest) effects.
- prediction accuracy: can always be improved by shrinking or setting some coefficients to zero, thus preventing from overfitting

Principle

- ▶ Retain only a subset of the variables, and eliminate the rest from the model
- different strategies for choosing the subset

Subset selection

Forward-stepwise selection

Greedy algorithm producing an increasing nested sequence of models:

- starts with the intercept
- then sequentially adds into the model the predictor that most improves the fit

Produces a sequence of models indexed by k, the subset size, which must be determined.

choosing k? cross-validation, AIC/BIC, 'significance' criterion, ...

Remarks

- suboptimal procedure, but we can always compute the forward stepwise sequence (even when $p \gg n$)
- similar to orthogonal matching pursuit in signal processing

Subset selection

Backward-stepwise selection

Greedy algorithm producing a decreasing nested sequence of models:

- starts with the full model,
- deletes the predictor that has the least impact on the fit.

How to choose the candidate for dropping: variable X_i with the smallest absolute Z-score

$$Z_j \equiv \frac{\hat{eta}_j}{\widehat{\mathrm{sd}}(\hat{eta}_j)},$$

where $\widehat{\mathrm{sd}}(\hat{eta}_j)$ is the approx. standard error for \hat{eta}_j

Remarks

b Backward selection can only be used when n > p, while forward stepwise can always be used.

Random Forests

Subset selection

Variable importance measure

- ▶ For bagged/RF regression trees, we record the total amount that the RSS is decreased due to splits over a given predictor, averaged over all *B* trees. A large value indicates an important predictor
- Similarly, for bagged/RF classification trees, we add up the total amount that the Gini index is decreased by splits overa given predictor, averaged over all B trees.
- Greedy selection procedures, e.g. Backward-stepwise selection

Regularization: shrinkage

Idea: introducing a little bias in the estimation of β may lead to a substantial decrease in variance and, hence, in the true error rate

Penalized regression

Regularize the estimation problem by introducing a penalization term for β

$$\widetilde{\beta} = \arg\min_{\beta} \left[\text{RSS}(\beta) + \lambda \operatorname{Pen}(\beta) \right]$$

- ▶ RSS(β) is the *fidelity term* to the training set (replace with the opposite log-likehihood $-\ell(\beta)$ for generalized linear model, e.g. logistic regression)
- $ightharpoonup \operatorname{Pen}(\beta)$ is the *a priori* to regularize the solution,
- \triangleright $\lambda > 0$ is the penalization coefficient

Choosing λ : tradeoff between overfitting (small λ) and underfitting (large λ)

standard practice is to use cross-validation to estimate an optimal λ for the test error rate (but AIC/BIC can also be used)

Ridge regression

Penalization in the (squared) ℓ_2 sense:

$$\operatorname{Pen}(\beta) \equiv \beta^T \beta = ||\beta||_2^2, \leftarrow \operatorname{Tychonov regularization}$$

 \widetilde{eta} is thus obtained by minimizing

$$\begin{split} \mathrm{RSS}(\beta) + \lambda \operatorname{Pen}(\beta) &= (Y - X\beta)^T (Y - X\beta) + \lambda \beta^T \beta, \\ &= \Big(\beta - (X^T X + \lambda I)^{-1} X^T Y\Big)^T \Big(X^T X + \lambda I\Big) \Big(\beta - (X^T X + \lambda I)^{-1} X^T Y\Big) + \mathrm{Cst}, \end{split}$$

Ridge estimator:
$$\widetilde{\beta} = (X^T X + \lambda I)^{-1} X^T Y$$

Remark

similar to LS estimator, with an additional 'ridge' on the diagonal of X^TX

- $X^TX + \lambda I$ has all its eigenvalues greater than $\lambda > 0$, \leftarrow ensures that $\widetilde{\beta}$ is always defined, and stable for large enough λ
- when $\lambda \to 0$, then $\widetilde{\beta} \to \widehat{\beta}$,
- when $\lambda \to +\infty$, then $\widetilde{\beta} \to 0$

Ridge Regression: deconvolution illustration

- $y \in \mathbb{R}^n$ with $n = 256^2$, $\beta \in \mathbb{R}^p$ with $p = 256^2$,
- ▶ $X \in \mathbb{R}^{n \times p} \leftarrow \text{sized } (256^2) \times (256^2) \text{ matrix...}$



 $\beta \leftarrow \mathsf{original\ image}$



 $y = X\beta \leftarrow \mathsf{blurred\ image}$

Regularization and shrinkage methods

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 $\beta \leftarrow \mathsf{original\ image}$



 $y = X\beta \leftarrow \mathsf{blurred} \mathsf{image}$



$$\widehat{\beta} = (X^T X)^{-1} X^T y \leftarrow \mathsf{LS}$$
 estimate

Ridge Regression: deconvolution illustration

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- ▶ $X \in \mathbb{R}^{n \times p} \leftarrow \text{sized } (256^2) \times (256^2) \text{ matrix...}$



 $\beta \leftarrow \text{original image}$



 $\widetilde{\beta} = (x^\mathsf{T} x + \lambda \mathbf{I})^{-1} x^\mathsf{T} \mathbf{y} \leftarrow \mathsf{ridge} \ \mathsf{estimate}$



 $y = X\beta \leftarrow \text{blurred image}$



 $\widehat{\beta} = (X^T X)^{-1} X^T y \leftarrow \mathsf{LS}$ estimate

Regularization by promoting sparsity

Sparse representations/approximations

A representation, or an approximation, is said to be sparse when most of the coefficients (in a given basis) are zero

'Bet on Sparsity' principle

Sparsity is a good option in high dimension!

- if the sparsity assumption does not hold, no method will be able to recover the underlying model in high dimension where $p \approx n$ or p > n
- but if the sparsity assumption holds true, then the parameters can be efficiently estimated by a method that promotes sparsity
- Occam's razor or KISS (keep it simple, stupid) principles: same idea that simpler models are preferable than more complex ones

Application to the regression problem

choosing a penalization function $\operatorname{Pen}(\beta)$ that promotes the sparsity of β (i.e. with many components $\beta_j = 0$ for $j = 1, \dots, p+1$) \leftarrow Lasso estimator

Lasso ('least absolute shrinkage and selection operator') estimator

Definition

$$\widetilde{\beta}_{\mathrm{lasso}} = \arg\min_{\beta} \left[\mathrm{RSS}(\beta) + \lambda \, ||\beta||_1 \right],$$

where $||\beta||_1 = \sum_{j=1}^{p+1} |\beta_j|$ is the ℓ_1 norm

- ightharpoonup no analytical expression of $\widetilde{eta}_{\mathrm{lasso}}$
- but convex optimization problem where very efficient numerical procedures are available to compute \widetilde{eta}_{lasso}

Lasso advantages

Converges to a generally sparse solution, i.e. such that $\beta_k=0$ for a subset of index k

- the less significant variables are explicitly discarded
- similar stability than ridge estimator + variable selection
- ► Tensorflow Playground: https://playground.tensorflow.org/
- ► Notebook: N3_curse_dimensionality.ipynb

Penalization with ℓ_1 and ℓ_2 norms: geometrical interpretation

- ▶ Least Squares estimator: $\widehat{\beta} = \arg \min \operatorname{RSS}(\beta)$,
- Regularized estimator: $\widetilde{\beta} = \arg\min \left(\operatorname{RSS}(\beta) + \lambda \operatorname{Pen}(\beta) \right)$ $\Leftrightarrow \widetilde{\beta} = \arg\min \ \operatorname{RSS}(\beta) \text{ under the constraint } \operatorname{Pen} \left(\widetilde{\beta} \right) \leq s(\lambda).$

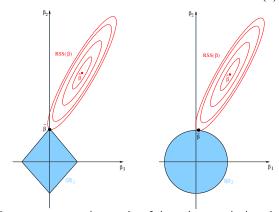


Illustration in dimension p=2: $\beta=(\beta_1,\beta_2)^T$

- ▶ Lasso $(\ell_1 \text{ norm})$: Pen $(\beta) = ||\beta||_1 = \sum_{k=1}^{p} |\beta_k|$
- ► Ridge regression (ℓ_2 squared norm) : Pen (β) = $||\beta||_2^2 = \beta^T \beta$

 ℓ_1 norm promotes the sparsity of the estimator: the less significant predictors are explicitly discarded (coeffs β_k are zero) \leftarrow model selection

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prostate data Heart diseases data prostate data

Application: prostate data

Stamey et al. (1989) study to examine the association between prostate specific antigen (PSA) and several clinical measures that are potentially associated with PSA in men. Objective is to predict the Log PSA from eight variables

- lcavol: Log cancer volume
- ► Iweight: Log prostate weight
- ▶ age: The man's age
- lbph: Log of the amount of benign hyperplasia
- ▶ svi: Seminal vesicle invasion; 1=Yes, 0=No
- lcp: Log of capsular penetration
- ▶ gleason: Gleason score
- pgg45: Percent of Gleason scores 4 or 5

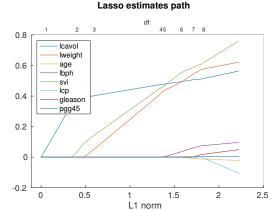
Application: prostate data

Model selection: ℓ_1 penalization (Lasso)

$$\widetilde{\beta}(\lambda) = \arg\min_{\beta} \mathrm{RSS}(\beta) + \lambda ||\beta||_1,$$

 \rightarrow function of λ where less significant variables are explicitly discarded

Path of the ℓ_1 -penalized coefficients vs $||\widetilde{\beta}(\lambda)||_1$



Choosing λ

- ▶ large $||\widetilde{\beta}(\lambda)||_1$ (small λ) → overfitting
- ▶ small $||\widetilde{\beta}(\lambda)||_1$ (large λ) \rightarrow underfitting
- ▶ $0.48 \le ||\widetilde{\beta}(\lambda)||_1 \le 1.43 \to 3$ predictors (lcavol,svi,lweight)

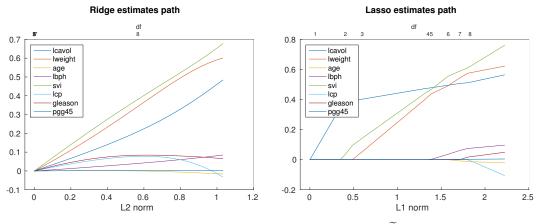
 $||\widetilde{\beta}(\lambda)||_1 = 1.06$ ($\lambda = 0.21$) estimated by cross validation

Notebook: N4_lasso_model_selection.ipynb

prostate data

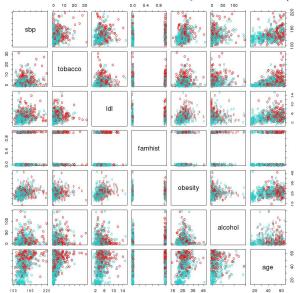
Application: prostate data

Comparison of ridge and lasso estimators



Path of the penalized coefficients as a function of $||\widehat{\beta}(\lambda)||$

Application: South African coronary heart disease (CHD)



Matrix of the predictor scatterplots

- ightharpoonup each plot \equiv pair of risk factors
- ► 160 cases / 302 controls
- ► Idl: ~ cholesterol, sbp: systolic blood pressure

Application: South African CHD (Cont'd)

Logistic regression fit

	Coefficient	Std. Error	Z score
(Intercept)	-4.130	0.964	-4.285
sbp	0.006	0.006	1.023
tobacco	0.080	0.026	3.034
ldl	0.185	0.057	3.219
famhist	0.939	0.225	4.178
obesity	-0.035	0.029	-1.187
alcohol	0.001	0.004	0.136
age	0.043	0.010	4.184

▶ A Z score (\equiv Coeff / Std. Error) > 2 in absolute value is significant at the 5% level.

Must be interpreted with caution!

- systolic blood pressure (sbp) is not significant!
- ▶ nor is obesity (conversely, < 0 coefficient)!
- $\boldsymbol{\rightarrow}$ result of the strong correlations between the predictors

Application: South African CHD (Cont'd)

Model selection: greedy backward procedure

Find the variables that are sufficient for explaining the CHD outputs

- drop the least significant predictor, and refit the model
- ightharpoonup repeat until no further terms can be dropped \leftarrow backward selection

Logistic regression fit with backward model selection procedure

	Coefficient	Std. Error	Z score
(Intercept)	-4.204	0.498	-8.45
tobacco	0.081	0.026	3.16
ldl	0.168	0.054	3.09
famhist	0.924	0.223	4.14
age	0.044	0.010	4.52

Interpretations

- ▶ Tobacco is measured in total lifetime usage in kilograms, with a median of 1kg for the controls and 4.1kg for the cases
- ▶ An increase of 1kg \Rightarrow increase of the CHD proba of exp (0.081) = 1.084 or 8.4% (confidence interval at 95% [1.03, 1.14])

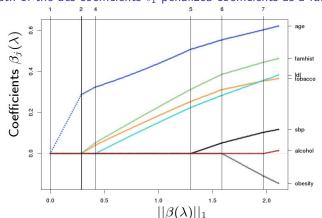
Application: South African CHD (Cont'd)

Model selection: ℓ_1 penalization (Lasso type method)

$$\widetilde{\beta}(\lambda) = \arg\min_{\beta} -\ell(\beta) + \lambda ||\beta||_1,$$

ightarrow function of λ where less significant variables are explicitly discarded

Path of the des coefficients ℓ_1 -penalized coefficients as a function of $||\hat{\beta}(\lambda)||_1$



Choosing λ

- ▶ large $||\widetilde{\beta}(\lambda)||_1$ (small λ) → overfitting
- small $||\widetilde{eta}(\lambda)||_1$ (large $\lambda)
 ightarrow$ underfitting
- ▶ 0.43 $\leq ||\widetilde{\beta}(\lambda)||_1 \leq 1.3 \rightarrow 4$ same predictors than backward selection procedure

Notebook:

N5_LR_heart_diseases_SA.ipynb

Heart diseases data

Conclusions

Generalized Linear Models

Learning of the prediction rule based on a model of Y given X

Linear regression, Logistic regression

Properties

- ► Simplicity: useful to capture the main effects
- Interpretability
- Shrinkage and Selection procedures
- extends to non-linear methods

Notebook: N6_select_features_boston.ipynb

Heart diseases data

Conclusions (Cont'd)

Model Selection

Model selection methods are an essential tool for data analysis, especially for big datasets involving many predictors

- Information criterion (AIC, BIC) make an adjustment to the training error to account for the overfitting bias and can be used to select among models with different numbers of variables,
- ▶ AIC/BIC are simple, but are known to be inacurate in high dimension (e.g. when the number of predictors is comparable or greater than the sample size)
- ► Cross-validation has an advantage relative to AIC, BIC in that it provides a direct estimate of the test error. Can be used for model selection and/or hyperparameter tuning