

1 Introduction and penalized regression

Sub-task 1:

Consider a balanced regression problem where for each input \mathbf{x}_i , $i = 1, \dots, N$, one has J repeated outputs y_{ij} , $j = 1, \dots, J$ and one fits a parameterized model $f_\theta(\mathbf{x})$ by least squares.

- Show that the fit can be obtained from a least squares problem involving only \mathbf{x}_i and the average values $\bar{y}_i = \frac{1}{J} \sum_{j=1}^J y_{ij}$.
- Explain how the least squares problem changes if the design is not balanced, i.e., one has different number of repetitions for each input \mathbf{x}_i .

Sub-task 2:

Assume a set of training data $(\mathbf{x}_1, y_1), \dots, (\mathbf{x}_N, y_N)$ drawn at random from a population as well as some test data $(\tilde{\mathbf{x}}_1, \tilde{y}_1), \dots, (\tilde{\mathbf{x}}_M, \tilde{y}_M)$ also drawn at random from the same population as the training data are given.

Show that

$$\mathbb{E} \left[\frac{1}{N} \sum_{i=1}^N (y_i - \mathbf{x}_i^\top \hat{\beta})^2 \right] \leq \mathbb{E} \left[\frac{1}{M} \sum_{i=1}^M (\tilde{y}_i - \tilde{\mathbf{x}}_i^\top \hat{\beta})^2 \right],$$

where $\hat{\beta}$ is the ordinary least squares estimate obtained for the training data assuming a linear regression model with p regression coefficients and the expectations are over all that is random in each expression.

Sub-task 3:

For a regression model with N observations, the coefficient of determination R^2 is defined as

$$R^2 = 1 - \frac{RSS}{TSS} = 1 - \frac{\sum_{i=1}^N (y_i - \hat{y}_i)^2}{\sum_{i=1}^N (y_i - \bar{y})^2},$$

where $\bar{y} = \frac{1}{N} \sum_{i=1}^N y_i$ and \hat{y}_i is the predicted value for the i th observation.

Show that the coefficient of determination R^2 is equal to the square of the correlation between X and Y in the simple linear regression case, where

$$Y = \beta_0 + \beta_1 X + \epsilon$$

and the regression coefficients $(\beta_0, \beta_1)^\top$ are estimated using ordinary least squares.

Sub-task 4:

Assume a linear regression model with

$$\mathbf{y} = \mathbf{X}\beta + \epsilon$$

where $\mathbf{y} \in \mathbb{R}^N$, $\mathbf{X} \in \mathbb{R}^{N \times p}$, $\beta \in \mathbb{R}^p$, $\epsilon \in \mathbb{R}^N$ and

1. $E(\epsilon) = \mathbf{0}$
2. $\text{Var}(\epsilon) = \sigma^2 \mathbf{I}$
3. \mathbf{X} deterministic with full column rank.

Show that in the case the design matrix $\frac{1}{N}\mathbf{X}$ is orthonormal (i.e., $\frac{1}{N}\mathbf{X}^\top \mathbf{X} = \mathbf{I}$) and $\beta \neq 0$, that there always exists a value of the penalty parameter λ such that the ridge estimator

$$\hat{\beta}^{\text{ridge}}(\lambda) = \left(\frac{1}{N}\mathbf{X}^\top \mathbf{X} + \lambda \mathbf{I} \right)^{-1} \frac{1}{N}\mathbf{X}^\top \mathbf{y}$$

has lower mean squared error than the OLS estimator.

Note that the mean squared error is given by:

$$E[(\tilde{\beta} - \beta)^\top (\tilde{\beta} - \beta)]$$

with $\tilde{\beta}$ either the OLS estimator or the ridge estimator for a given λ .

Sub-task 5:

Assume the following data generating process with X drawn from some distribution and

$$\begin{aligned} Y|X &\sim f(X) + \epsilon, \\ \epsilon &\sim N(0, \sigma^2). \end{aligned}$$

A sample of N pairs (x_i, y_i) independently drawn from the data generating process are given.

Construct an estimator for f linear in y_i ,

$$\hat{f}(x_0) = \sum_{i=1}^N \ell_i(x_0, \mathcal{X}) y_i,$$

where the weights $\ell_i(x_0, \mathcal{X})$ do not depend on y_i , but depend on the entire training sequence x_i , denoted here by \mathcal{X} .

- Show that k -nearest-neighbor regression is a member of this class of estimators and explicitly describe the weights $\ell_i(x_0, \mathcal{X})$.
- Show that linear regression given by

$$\hat{f}(x) = \hat{\beta}_0 + \hat{\beta}_1 x$$

is a member of this class of estimators and explicitly describe the weights $\ell_i(x_0, \mathcal{X})$.

Sub-task 6:

Assume that

$$Y = f(X) + \epsilon, \quad f(X) = X_1$$

with $\epsilon \sim N(0, \sigma^2)$ and X is independent and uniformly distributed on $[-1, 1]^p$.

X_1 denotes the first element of the p -dimensional vector X .

- Use simulations to approximate the EPE given by

$$\begin{aligned} \text{EPE}_{\hat{f}_{\mathcal{T}}}(x_0) &= E[(Y - \hat{f}_{\mathcal{T}}(x_0))^2 | X = x_0] \\ &= \text{Var}(Y|X = x_0) + E_{\mathcal{T}}[(E(Y|X = x_0) - \hat{f}_{\mathcal{T}}(x_0))^2 | X = x_0] \\ &= \text{Var}(Y|X = x_0) + [\text{Bias}_{\mathcal{T}}^2(\hat{f}_{\mathcal{T}}(x_0)) + \text{Var}_{\mathcal{T}}(\hat{f}_{\mathcal{T}}(x_0))] \end{aligned}$$

with \mathcal{T} the training sample at $x_0 = \mathbf{0}$ based on $m = 1000$ repetitions

- for a training sample size of $N = 500$,
 - the number of dimensions p varying from 1 to 10,
 - with a standard deviation σ of either zero or one,
 - using linear models (with an intercept) as well as 1-nearest neighbors to estimate $f(X)$.
- Assess and interpret the impact of the dimension on the EPE in dependence of the method used as well as the value of the standard deviation.

Sub-task 7:

Use the diabetes data set to fit different linear models. The data set is available in the R package **lars** and can be loaded using:

```
> data("diabetes", package = "lars")
```

The dependent variable is contained in `diabetes$y`, the model matrix in `diabetes$x` for the linear regression.

- Set a random seed and split the data set into a training and test data set such that 400 observations are used for training and the remaining ones for testing. Explain why it might be good to randomly select 400 observations from the available data set instead of using the first 400.
- The covariates are only available in standardized form. Explain if this is an issue for the subsequent analysis.
- Analyze the pairwise correlation structure between the covariates as well as the covariates and the dependent variable. Interpret the results and explain how these correlations impact model selection.
- Fit a linear regression model containing all explanatory variables. Inspect the model and evaluate the in-sample fit as well as the performance on the test data based on the mean squared error (MSE).
- Fit a smaller model where only the covariates are contained which according to a t -test are significant at the 5% significance level conditional on all other variables being included. Evaluate the performance in-sample as well as on the test data. Compare this model to the full model using an F -test. data and compare this model to the full model using an F -test.

Sub-task 8:

Use again the diabetes data set available in the R package **lars** as in the previous sub-task. Split again the data set into a training and test data set such that 400 observations are used for training and the remaining ones for testing after setting a random seed.

- Use backward-stepwise regression based on the AIC to select a suitable model. Evaluate the performance in-sample as well as on the test data and compare this model to the full model using an F -test.
- Use best subset selection to select a suitable model based on the AIC. Evaluate the performance in-sample as well as on the test data and compare this model to the full model using an F -test.

Summarize the results in a table containing the regression coefficients of the full model as well as the model selected with stepwise regression and best subset selection as well as the in-sample and the test data performance and interpret the results.

Sub-task 9:

Use the wage data set to fit different linear models. The data set is available in the R package **ISLR** and can be loaded using:

```
> data("Wage", package = "ISLR")
```

- Fit a linear regression model to predict **Wage**. Omit the variable **logwage** as well as all other variables where inclusion in the model is problematic before analysis, specify linear, quadratic and cubic effects for the variable **age** and use suitable contrasts for the variable **education**.
- Use best subset selection to determine a suitable model. Interpret the selected model.
- Assess and explain if it makes a difference if you include the polynomial of the original variable **age** or orthogonal polynomials constructed using **poly(age, k)**.

Sub-task 10:

Implement a function

```
> forwardstagewise <- function(x, y, tol = sqrt(.Machine$double.eps)) { ... }
```

with **x** the model matrix (without intercept), **y** the response and **tol** a tolerance to stop the iterative procedure, which performs forward stagewise regression assuming that an intercept is also included.

- The function should return a matrix of the regression coefficients for **x** containing the estimates for each step in the rows.
- Check that the input arguments are suitably specified and issue an error otherwise.
- Ensure that the variables in **x** as well as **y** are centered before performing the analysis, i.e., subtract the empirical mean from each of the variables.

Compare forward stagewise regression to best subset regression on a simulated linear regression problem using the following setting:

- There are $N = 300$ observations.
- There are $p = 31$ covariates drawn from a standard Gaussian distribution, with pairwise correlations all equal to 0.85.
- For 10 of the variables, the coefficients are drawn at random from a $N(0, 0.6^2)$ distribution; the rest are zero.
- The noise is distributed $\epsilon \sim N(0, 2.5^2)$.
- Use package **leaps** for best subset selection and the self-implemented function for forward stagewise regression.
- Estimate $E(\|\hat{\beta}_k - \beta\|_2^2)$ based on 50 simulations with k the subset size for best subset selection and step number in the case of forward stagewise selection.
- Visualize the results.