Machine Learning Supervised learning 2

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Linear regression: probably the oldest machine learning technique



- Find linear correlation coefficient.
- Compute coefficients of the linear equation

$$\hat{y} = ax + b$$

Evaluate the model

• In multivariate case it is required to identify coefficients of the model

$$\hat{y} = a_1 x_1 + a_2 x_2 + \ldots + a_n x_n + b.$$

This leads the necessity to choose variables (perform model building).

Linear regression

• Correlation coefficient.

$$\rho = \frac{\sum_{i=1}^{n} (x_i - \bar{x})(y_i - \bar{y})}{\sqrt{\sum_{i=1}^{n} (x_i - \bar{x})^2 \sum_{i=1}^{n} (y_i - \bar{y})^2}}$$

where, n - is the sample size, x and y are the variable of interest.

- $-1 \le \rho \le 1$
- Assumption there are exist α and β such that for any $i = 1, \ldots, n$ $y_i = \alpha x_i + \beta + \varepsilon_i$ holds. Assumption: ε is sufficiently small normally distributed.
- The goal of regression is to find estimates of the coefficients α and $\beta,$ such that for a and b

$$y_i = ax_i + b + \hat{\varepsilon}_i$$

sum of squares of $\hat{\varepsilon}_i$ would be minimal. NB! notation $\hat{\alpha}$ and $\hat{\beta}$ is also widely use.

Least squares method

Least squares method:



For an arbitrary number of variables:

$$y = b_1 x_1 + \ldots + b_n x_n + b_0$$

then

$$\hat{b} = (X^T X)^{-1} X^T y.$$

where each row of matrix X is input vector with 1 in the first position.

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Machine Learning

Model validation

- Coefficient of determination R^2 and adjusted R^2 .
- Significance of the model and model coefficients.
- Verify assumption that residuals are normally distributed.
- Residual sum squares. RSS = $\sum_{i=1}^{N} (y_i x_i^T \beta)^2$.
- Sum squares of the regression $SSR = \sum_{i=1}^{N} (\hat{y}_i \bar{y})^2$.
- Total sum squares or sum of squares about the mean $SST = \sum_{i=1}^{N} (y_i \bar{y})^2$.
- R^2 computed as the ratio of Sum squares of the regression to total sum squares or one minus ratio of Residual sum squares to total sum squares whereas adjusted R^2 is one minus ratio of residual sum squares computed for n-1 to Total sum squares for n-pobservation points.

MLE for regression least squares I

• Linear regression is the model of the form

$$p(y|x,\theta) = \mathcal{N}(y|\beta^T x, \sigma^2)$$

where β are the coefficients of the linear model, σ is the standard deviation of x and $\theta=(\beta,\sigma^2)$

• Parameter estimation of a statistical model is usually performed by computing MLE $\hat{\theta} = \arg \max_{\theta} \log p(\mathcal{D}|\theta)$. remind that \mathcal{D} denotes the data set

MLE for regression least squares II

- Assumption: elements of the training set are independent and identically distributed.
- Then log likelihood is given by $\ell(\theta) = \log p(\mathcal{D}|\theta) = \sum_{i=1}^{N} \log p(y_i|x_i, \theta).$
- As usually instead of maximizing the log- likelihood one may minimize negative log likelihood.

$$\ell(\theta) = \sum_{i=1}^{N} \log\left[\left(\frac{1}{2\pi\sigma^2}\right) \exp\left(-\frac{1}{2\sigma^2}(y_i - \beta^T x_i)^2\right)\right]$$
$$= \frac{-1}{2\sigma^2} \operatorname{RSS}(\beta) - \frac{N}{2} \log(2\pi\sigma^2).$$

MLE for regression least squares II

• In order to minimize RSS differentiate its equation which lead

$$\nabla \theta = X^T X \beta - X^T y.$$

• Equate it to zero and solve for β

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

last equation is referred as normal equation.

Regularization

- Overfitting may be caused by the fact that chosen model structure and data are not conform on another.
- Regularization is the technique used to overcome overfitting.
- Regularization imposes cost or penalty on the cost function and prevent larger values of the coefficients.
- Loosely speaking, regularization shrinks the coefficients towards zero and towards one another.

Ridge regression

• Ridge regression shrinks the coefficients by penalizing their size.

$$\hat{\beta}^{\text{ridge}} = \operatorname{argmin}_{\beta} \left\{ \sum_{i=1}^{N} \left(y_i - \beta_0 - \sum_{j=1}^{p} x_{ij} \beta_j \right)^2 + \lambda \sum_{j=1}^{p} \beta_j^2 \right\}$$

 λ is the nonnegative shrinkage parameter, its large values correspond to the greater amount of shrinkage applied.

• Alternatively the following notation is widely used:

$$\hat{\beta}^{\text{ridge}} = \operatorname{argmin}_{\beta} \sum_{i=1}^{N} (y_i - \beta_0 - \sum_{j=1}^{p} x_{ij} \beta_j)^2$$

subject to
$$\sum_{j=1}^{N} \beta_j^2 \le t$$

The Lasso

• Ridge regression shrinks the coefficients by penalizing their size.

$$\hat{\beta}^{\text{lasso}} = \operatorname{argmin}_{\beta} \frac{1}{2} \left\{ \sum_{i=1}^{N} \left(y_i - \beta_0 - \sum_{j=1}^{p} x_{ij} \beta_j \right)^2 + \lambda \sum_{j=1}^{p} |\beta_j| \right\}$$

 λ is the nonnegative shrinkage parameter, its large values correspond to the greater amount of shrinkage applied.

• Alternatively the following notation is widely used:

$$\hat{\phi}^{\text{ridge}} = \operatorname{argmin}_{\beta} \sum_{i=1}^{N} (y_i - \beta_0 - \sum_{j=1}^{p} x_{ij}\beta_j)^2$$

subject to $\sum_{j=1}^{N} |\beta_j| \le t$

• Computing the lasso solution is a quadratic programming problem.

Statistical hypothesis testing (brief reminder I)

- Assumption about a parameter of population is a statistical hypothesis.
- Usually a pair of hypothesis is stated (H_0, H_1) , notation (H_0, H_a) .
 - ► *H*⁰ the null hypothesis usually states that there is no statistically significant relationship between two phenomena.
 - H_1 the alternative hypothesis usually states the opposite to the H_0 .
- Choose and compute test statistic and rejection rule.
- Interpret the results.
- What can possibly go wrong?

Statistical hypothesis testing (brief reminder II)

	Accept H_0	Reject H_0
H_0 is true	Correct	Type 1 Error
H_0 is false	Type II Error	Correct

Model building (feature selection)

Let us suppose that observed process has p independent variables x_1, \ldots, x_p and one dependent variable y. Should one build the regression equation using all p variables or not?

- Are all the variables x_1, \ldots, x_p uncorrelated?
- Which subset of variables result in a "better" model?
- How to prove that as a result of adding or deleting a variable model quality has improved?

"Butler tracking company" example

- Independent variables: Distance to drive and number of parcels to deliver. Dependent variable: time.
- Distances to drive for each assignment: 100, 50, 100, 100, 50, 80, 75, 65, 90, 90.
- Number of parcels to deliver: 4, 3, 4, 2, 2, 2, 3, 4, 3, 2
- Time in hours: 9.3, 4.8, 8.9, 6.5, 4.2, 6.2, 7.4, 6, 7.6, 6.1.
- Pearson correlation coefficient between distance and time is 0.81.

"Butler tracking company" example continued



Model 1

Is significant p = 0.004, F = 15.1846 whereas $R^2 = 0.6641$.

Model 2

Is significant p = 0.000276, F = 32.9 whereas adjusted $R^2 = 0.87$.

Is it enough to say that model 2 is more precise?

Quality comparison

- To compare different models residual sum of squares (RSS) is used.
- Hypothesis statements: $H_0: RSS_s \leq RSS_c H_1: RSS_s > RSS_c$.
- Test statistic (empirical parameter) for ANOVA:

$$F_{stat} = \left(\frac{\text{RSS}_s - \text{RSS}_c}{m}\right) \left(\frac{\text{RSS}_c}{n - p - 1}\right)^{-1}$$

where RSS_c is the residuals sum squares of model with more variables, RSS_s - is the residuals sum squares of model with less variables, m number of variables added or removed, n is the number of observation points, p - is the number of variables in more complicated model.

- Rejection rule for α (significance level), degrees of freedom: first is the number of variables added or removed, second is n - p - 1.
 Desirion:
- Decision:
 - (if adding variables) rejected null hypothesis proves that adding variables caused model quality to increase significantly.
 - (if deleting variables) rejected alternative hypothesis proves that deleting variables did not cause model quality to significant decrease.

"Butler tracking company" example continued

- $RSS_1 = 15.8713$, $RSS_2 = 2.2994$ NB! Observe that corresponding MATLAB notation is SSE!!!
- choose $\alpha = 0.05$ degrees of freedom: first will be 1 (one variable (number of parcels)) were added, second 7 (n = 10, p = 2).
- Rejection rule: reject H_0 if $F_{stat} > 5.5914$
- Compute $F_{stat} = 17.4411$. (use table, or MATLAB or EXCEL)
- Reject H_0 . Adding the variable has increased the model quality.

- Choose or determine all the hyperparameters. Possible order limitations, backward elimination / forward selection/ batch processing, set the level of significance and threshold for correlation. These parameters also define stopping criteria.
- Stop when: model is significant, and goodness parameters as expected OR no more variables to add or delete OR maximal or minimal order is reached etc.
- Investigate if available explanatory variables (predictors) are linearly independent. Strong dependencies between variables chosen as "independent" lead problems with inverting matrix X. Compute <u>multicollinearity</u> matrix where element in *i*th row and *j*th column is Pearson correlation coefficients computed for variables *i* and *j*. Based on this table determine subset(s) of variables which are linearly independent.

- Repeat
- Apply mean squares (or other technique) to build the model from selected variables.
- Evaluate significance- and quality- of the model. For quality observe determination coefficient and error. For significance use *F* test and *t*-test variable wise.
- If model fail goodness or significance check then return to the previous model and choose another set of variables to add/delete.
- Starting from second iteration prove, using *F* test, that as a result of adding/deliting variables model quality has improved/did not decreased significantly.
- If adding/deliting variables was not successful return to the previous model and if possible chose another variable(s) to add /delete or report the model from previous step.
- If goodness criteria (quality and significance) is met stop and return the model.
- If goodness criteria was not met but adding deleting variables proved to be successful chose the set of variables to be added or deleted (*t*-test) on the next step.
- Until stopping criteria is reached.
- Report the results.

Reminder p - is the number of variables n is the sample size.

- F -test of overall significance in regression analysis.
- Test for model significance. $H_0: b_1 = \ldots = b_p = 0, H_1:$ $\exists i: 1 \le i \le p \& b_i \ne 0.$
- Test statistic:

$$F = \frac{\frac{\sum_{i=1}^{n} (\hat{y}_i - \bar{y})^2}{p-1}}{\frac{\sum_{i=1}^{n} (y_i - \hat{y}_i)^2}{n-p}}$$

• Rejection rule: Determine using F-table or corresponding software function with chosen significance level, n degrees of freedom in denominator and p degrees of freedom in nominator.

- F -test to determine significance of change in model quality caused by adding variables
 - H_0 : $RSS_S \leq RSS_C$, H_1 : $RSS_S > RSS_C$.
 - Test statistic:

$$F = \frac{\frac{RSS_S - RSS_C}{m}}{\frac{RSS_C}{n - p - 1}}$$

- Rejection rule: Determine using F-table or corresponding software function with chosen significance level, n p 1 degrees of freedom in denominator and m degrees of freedom in nominator.
- t test on individual regression coefficients
 - $H_0: b_i = 0, H_1: b_i \neq 0.$
 - Test statistic: $t = \hat{b_i}/se(\hat{b_i})$
 - Use t table or corresponding function to find rejection rule for chosen significance and n-2 degrees of freedom.



- By replacing independent variables X with a nonlinear mapping $\phi(X)$.
- This will lead

$$f_{\theta}(X) = \theta^T \phi(X)$$

- This process is referred as basis function expansion.
- Example: Polynomial regression has basis function $\phi(X) = [1, x, x^2, \dots, x^d]$. The model remains linear in the parameters.

Polynomial regression 1

- Higher degree polynomial models tend to over fit. The coefficients become relatively large, which causes the regression curve to "wiggle".
- In order to achieve "encourage" smaller weight values introduce zero-mean Gaussian prior:

$$p(\theta) = \prod_{j} \mathcal{N}(\theta_j | 0, \tau^2)$$

where $1/\tau^2$ controls the strength of prior.

• This lead following log-likelihood estimate

$$\ell = \sum_{i=1}^{N} \log \mathcal{N}(y_i | \theta^T x_i, \sigma^2) + \sum_{j=1}^{p} \log \mathcal{N}(\theta_j | 0, \tau^2)$$

• The solution is given by:

$$\hat{\theta_r} = (\lambda I + X^T X)^{-1} X^T y$$

Logistic regression

• Remind that linear regression may be written in the following form:

$$p(y|x,\theta) = \mathcal{N}(y|\mu(x),\sigma^2(x))$$

• This may be generalized to the binary setting as follows:

$$p(y|x, \theta) = \text{Ber}(y|\text{sigm}(\theta^T x))$$

where sigm $(\eta) = (1 + e^{-\eta})^{-1}$. Will be referred as <u>logistic regression</u>. • Fitting is usually done by maximum likelihood

$$\ell(\theta) = \sum_{i=1}^{N} \log p(g_i)(x_i|\theta) = \sum_{i=1}^{N} \left\{ y_i \beta^T x_i - \log(1 + e^{\beta^T x_i}) \right\}$$

• Solving the last one is done by means of iterative algorithm.

$$b^{\text{new}} = \arg\min_{b} (z - Xb)^{T} W(z - Xb)$$
$$z = Xb + W^{-1}(y - p)$$

where W is a $N\times N$ diagonal matrix with ith element $p(x_i,|b)(1-p(x_i|b))$

k-nn regression

The value of the response (dependent variable) defined as the average of its k nearest neighbours from the training set.



Regression trees

- Partition the feature space into the set of rectangles.
- Fit a simple model (for example constant) in each rectangle.
- Fitting the model is similar to the case of classification trees.

