Machine learning Part 2 Dimensionality reduction & cross-validation

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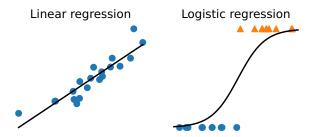
QLS course 2021-07-30





Supervised learning

- Regression: least-squares linear regression
- Classification: logistic regression



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Regularization

• ℓ_2 a.k.a. ridge regularization

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Model evaluation and selection

- Out-of-sample generalization; independent test set
- Performance metrics:
 - regression: mean squared error
 - · classification: accuracy, ROC curve
- Cross-validation

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Don't remember? watch Part 1 again!

$$Y = f(X) + E \tag{1}$$

• $Y \in \mathbb{R}$: output (a.k.a. target, dependent variable) to predict

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- $E \in \mathbb{R}$: unmodelled noise

Y = f(X) + E

- independent variables)
- $E \in \mathbb{R}$: unmodelled noise • f: the function we try to approximate

$$Y = \beta_0 +$$

$$Y = \beta_0 + \langle X, \beta \rangle + E$$

$$1 = \beta 0 + \langle X, \beta \rangle + C$$

$$= \rho \cdot + \sum_{i=1}^{p} V_{ii} \rho_{ii}$$

$$= \beta_0 + \langle X, \beta \rangle + C$$

$$= \beta_0 + \sum_{j=1}^p X_j \beta_j + C$$

(2)

(1)

$$-\beta_0 + \sum_{j=1}^{N_j} N_j \beta_j + C$$

How to estimate parameters: Empirical Risk Minimization

Given \mathfrak{n} training examples $X \in \mathbb{R}^{\mathfrak{n} \times \mathfrak{p}}$, $\mathfrak{y} \in \mathbb{R}^{\mathfrak{n}}$, find $\hat{\beta}_0 \in \mathbb{R}, \hat{\beta} \in \mathbb{R}^{\mathfrak{p}}$ that minimize the empirical risk:

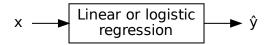
$$\|\mathbf{y} - \hat{\mathbf{y}}\|_{2}^{2} = \|\mathbf{y} - \hat{\boldsymbol{\beta}}_{0} - \mathbf{X}\,\hat{\boldsymbol{\beta}}\|_{2}^{2}$$

$$= \sum_{i=1}^{n} (\mathbf{y}_{i} - \hat{\boldsymbol{\beta}}_{0} - \sum_{j=1}^{p} \mathbf{X}_{ij}\,\hat{\boldsymbol{\beta}}_{j})^{2}$$
(5)

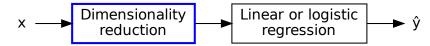
"Fitting" the parameters to X, y.

Dimensionality reduction

Until now



Add a step in the pipeline: simplifying the inputs



Dimensionality reduction

$$\hat{y} = \hat{\beta}_0 + \hat{\beta}_1 X_{:,1} + \hat{\beta}_2 X_{:,1} + \dots + \hat{\beta}_p X_{:,p}$$
 (6)

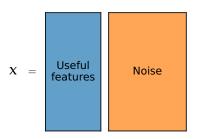
Problems when the number of features p becomes large

- Bigger errors on test data (larger variance of predictions)
- Numerical stability issues
- · Computational cost and memory usage

Simulated data for linear regression

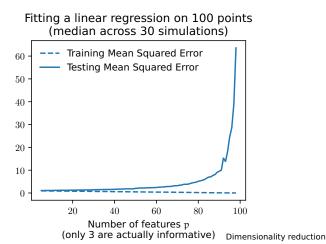
- Generate $X \in \mathbb{R}^{n \times 3}$, $\beta \in \mathbb{R}^3$, $e \in \mathbb{R}^n$ and $y = X\beta + e \in \mathbb{R}^n$
- Append columns containing random noise to X
- Now $X \in \mathbb{R}^{n \times p}$, with $p \geqslant 3$, but only the first 3 columns are linked with y
- Split into training and testing tests and evaluate a linear regression model: what happens when p becomes large?

See sklearn.datasets.make_regression for generating data

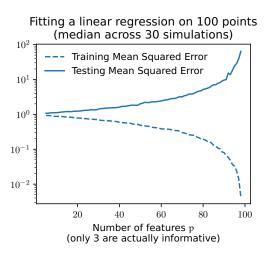


Model complexity: overfitting

- Model complexity increases with dimension.
- Example: a linear model in dimension $\mathfrak p$ can fit exactly (0 training error) any set of $\mathfrak p+1$ points.
- Risk of overfitting: fitting exactly training data but failing on test data

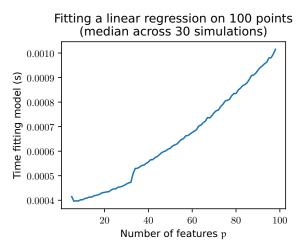


Same plot in log scale



Cost of fitting many parameters

- Many algorithms require polynomial time in p
- Implementations often make copies of the design matrix (e.g. for centering & rescaling)



Solution 1: univariate feature selection

- a.k.a. feature screening, filtering . . .
- Check features (columns of X) one by one for association with the output y
- Keep only a fixed number or percentage of the features

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Simple (linear) association criteria

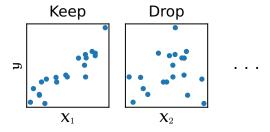
- for regression: correlation
- for classification: ANalysis Of VAriance

Read more in the scikit-learn user guide

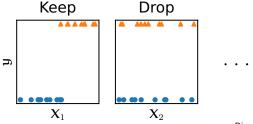
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https://scikit-learn.org/stable/modules/feature_selection.
html#feature-selection
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Simple selection criteria

· Regression: correlation

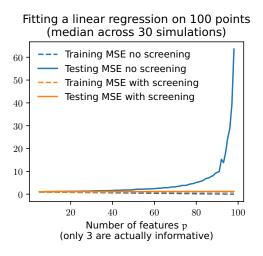


· Classification: ANOVA

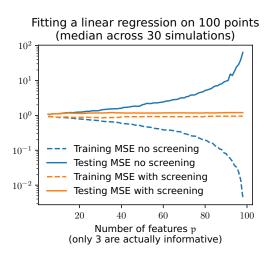


Univariate feature selection

Keeping only the 10 best features (most correlated with y)

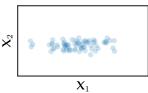


Same plot in log scale

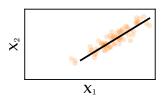


Solution 2: linear decomposition methods

Maybe OK to drop X_2 :



Data low-dimensional but no feature can be dropped:

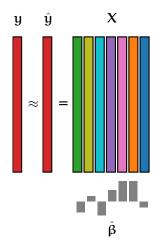


Find a better referential in which to represent the data

Linear regression: projection on the column

space of X

$$\hat{y} = X \hat{\beta}$$
 (7)

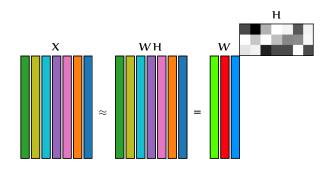


- Too many features: high variance & unstable solution
- Feature selection: drop some columns of X
- Other ways to build a family of k vectors on which to regress y?

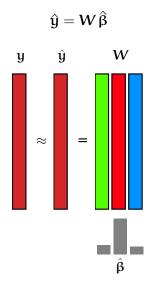
Linear decomposition: low-rank approximation of \mathbf{X}

Minimize

$$\|X - WH\|_F^2 = \sum_{i,j} (X_{i,j} - (WH)_{i,j})^2$$
 (8)



Linear regression after dimensionality reduction



(9)

Prediction for a new data point $x \in \mathbb{R}^p$

- Find the combination of rows of H that is closest to x: regress x on H^T
- Multiply by $\hat{\beta}$

$$x \in \mathbb{R}^p o \mathsf{projection} o w \in \mathbb{R}^k o \langle \cdot \,, \, \hat{eta}
angle o \hat{y} \in \mathbb{R}$$
 (10)

Principal Component Analysis

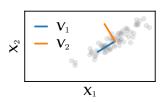
• Singular Value Decomposition of X:

$$X = \mathbf{U} \, \mathbf{S} \, \mathbf{V}^{\mathsf{T}} \tag{11}$$

with $X \in \mathbb{R}^{n \times p}$, $U \in \mathbb{R}^{n \times r}$, $S \in \mathbb{R}^{r \times r}$, $V \in \mathbb{R}^{r \times p}$

- r = min(n, p)
- $\textbf{S}\succeq \textbf{0}$ diagonal with decreasing values s_j along the diagonal
- $\mathbf{u}^\mathsf{T} \mathbf{u} = \mathbf{I}_r$
- $V^T V = I_r$

Truncating the SVD to keep only the first k components gives the best rank-k approximation of \boldsymbol{X}



Singular Value Decomposition

$$X = \mathbf{U} \, \mathbf{S} \, \mathbf{V}^{\mathsf{T}}$$

$$\mathbf{v}_{1}$$

$$\mathbf{u}_{1}$$

$$\mathbf{v}_{2}$$

$$\mathbf{v}_{3}$$

$$\mathbf{v}_{4}$$

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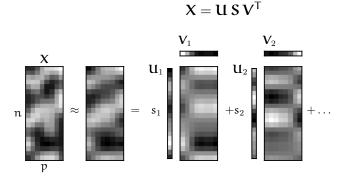
$$\mathbf{v}_{8}$$

$$\begin{aligned} \boldsymbol{u}^\mathsf{T}\,\boldsymbol{u} &= \mathrm{I}_{p} \\ \boldsymbol{V}^\mathsf{T}\,\boldsymbol{V} &= \mathrm{I}_{p} \end{aligned}$$

$$\mathbf{V}^\mathsf{T} \mathbf{V} = \mathbf{I}_n$$

(12)

Singular Value Decomposition



$$\mathbf{U}^\mathsf{T}\,\mathbf{U} = \mathrm{I}_\mathrm{p} \tag{16}$$

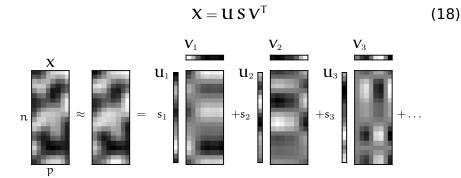
$$\mathbf{V}^\mathsf{T}\,\mathbf{V} = \mathrm{I}_\mathrm{p} \tag{17}$$

$$\mathbf{V}^\mathsf{T} \mathbf{V} = \mathbf{I}_n$$

(17)

(15)

Singular Value Decomposition



Explained variance: 0.97

$$\mathbf{U}^{\mathsf{T}} \, \mathbf{U} = \mathrm{I}_{\mathrm{p}} \tag{19}$$

$$\mathbf{V}^{\mathsf{T}} \, \mathbf{V} = \mathrm{I}_{\mathrm{p}} \tag{20}$$

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Other decomposition methods

Many other methods use the same objective (sum of squared reconstruction errors), but add penalties or constraints on the factors

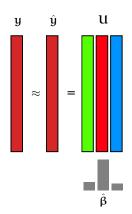
- · Dictionary Learning
- Non-negative Matrix Factorization
- K-means clustering
- ...

What about u?

- PCA is an example of unsupervised learning: it does not use y
- Some other methods take it into account: e.g. Partial Least Squares

Ridge regression and PCA

- Both ridge regression and PC regression compute the coordinates of y in the basis given by the SVD of X
- Ridge shrinks the coordinate along U_j by a factor $s_j^2/(s_j^2+\lambda)$
- PC regression sets the coordinates to 0 except for those corresponding to the k largest s_j: shrinks by a factor 1_{j≤k}



Setting hyperparameters

How can we choose:

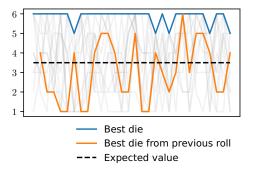
- Number of features or PCA components k?
- The ridge hyperparameter λ?

Try a few and pick the best one...
But measure its performance on separate data!

Need for fresh test data

When you hear "best", "maximum", "select", ... think "bias"

- I have 4 dice and want to find one that rolls high numbers
- I roll them all once and select the die that gives the highest number
- The selected die rolled a 5. Is 5 a good estimate of that die's average result? What if I had 1,000 dice?
- I need to roll it again to get an unbiased estimate



Nested cross-validation

When you hear "best", "maximum", "select", ... think "bias"

Nested cross-validation

When you hear "best", "maximum", "select", ... think "bias" Setting the parameters

- Select β that gives the **best** prediction on training data
- The prediction score for $\hat{\beta}$ is biased: compute a new score on unseen test data.

Nested cross-validation

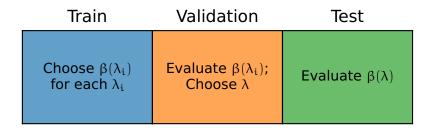
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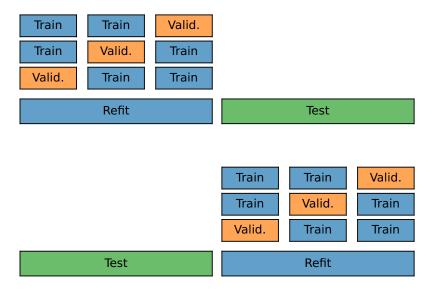
Setting the hyperparameters

- Repeat step 1 for a few values of λ, k, etc.., fitting and testing several models
- Select the hyperparameter that obtains the best prediction on test data
- The prediction score of that model on *test* data is biased: evaluate it again on unseen data

One split



Nested cross-validation



See sklearn.model_selection.GridSearchCV

- e.g. fit PCA on all data, then do cross-validation on dim-reduced dataset
- USE sklearn.pipeline.Pipeline

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Ignoring dependencies between samples

- · Multiple datapoints per participant
- Time series

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 Training sets overlap: cross-validation scores of different splits are not independent

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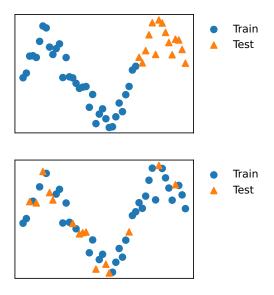
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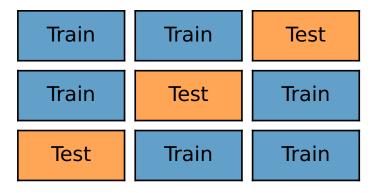
Over-interpreting good CV scores

 Good CV scores on one dataset do not mean the model will always perform well on a new dataset

Split choice example: time series Which is easier?



Remember that CV training sets overlap



So the scores are not independent! Their variance can be underestimated.

Supervised learning with fMRI

 Predict in which site / with which scanner a resting-state fMRI sequence was acquired

The prediction pipeline

- Masking: extracting voxels that are inside the brain
- Connectivity: measuring correlations between brain regions to build a feature vector for each participant
- Univariate feature selection with ANalysis Of VAriance
- Classifier: logistic regression

Implementation: in class